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*Loyola University Chicago*

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QUANTITATIVE ANALYSIS OF RESPIRATORY CELL ACTIVITY

by

Charles Lewis Webber, Jr.

A Dissertation Submitted to the Faculty of the Graduate School  
of Loyola University of Chicago in Partial Fulfillment  
of the Requirements for the Degree of  
Doctor of Philosophy

February, 1974

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## DEDICATION

I take great pleasure in dedicating this dissertation to my wife, Connie, in appreciation for her constant encouragement and positive attitude throughout the period of my graduate training.

## ACKNOWLEDGEMENTS

The preparation of this dissertation would not have been possible without the help and encouragement of many individuals. I am greatly indebted to Dr. Walter C. Randall, Professor and Chairman of the Department of Physiology, for initially introducing me to physiological research four years ago. My sincerest appreciation is expressed to Dr. Clarence N. Peiss, Professor of Physiology and Associate Dean of the Graduate School, for directing this study and overseeing my training as a scientist. It is also a pleasure to acknowledge the expert technical assistance of Dr. Robert D. Wurster in neurophysiological problems and Dr. Robert D. McCook in computer software programming.

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## BIOGRAPHY

Charles Lewis Webber, Jr., was born on July 26, 1947, in Bay Shore, Long Island, New York. He completed primary and secondary education requirements of the Sayville Public School system, graduating from high school in June, 1965, with a double diploma from the local district and New York State Board of Regents. He studied at Taylor University, Upland, Indiana, for the next four years and received the Bachelor of Arts degree, cum laude, in June, 1969, with a chemistry major and zoology minor.

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BIOGRAPHY  
(continued)

In June, 1970, Charles married Constance Anne Folkers, a 1969 Taylor graduate, from Minonk, Illinois. During his last three years of graduate training, his wife taught seventh grade language arts at Gurrie Junior High School in LaGrange, Illinois. The couple resided in Broadview, Illinois.

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## PUBLICATIONS

1. Webber, C.L., Jr. and C.N. Peiss. Computer analysis of activity patterns in single respiratory cells. Abstract, 1st Annual Society for Neuroscience, Washington, D.C., October, 1971.
2. Webber, C.L., Jr. and C.N. Peiss. Barbiturate induced apneusis in the cat. The Physiologist 15: 299, 1972.

## TABLE OF CONTENTS

Chapter	Page
I. INTRODUCTION.....	1
II. LITERATURE REVIEW	
A. Respiratory Areas of the Medulla.....	3
B. Apneusis.....	15
C. Medullary Outputs.....	29
D. Discharge Patterns.....	45
III. METHODS	
A. Surgical Preparation.....	60
B. Data Acquisition.....	63
C. Experimental Manipulations.....	70
IV. DATA ANALYSIS	
A. PDP-12 Overview.....	73
B. Machine Language Programs.....	75
C. FOCAL-12 Programs.....	99
D. FOCALPL Programs.....	114
V. RESULTS	
A. Data Presentation.....	117
B. Population Data.....	118
C. Single Cell Data.....	167
D. Barbiturate Apneusis.....	218



TABLE OF CONTENTS  
(continued)

Chapter	Page
VI. DISCUSSION	
A. Population Modeling.....	242
B. Single Cell Elements.....	266
C. Respiratory Complex Organization.....	272
D. Information Transfer.....	279
VII. CONCLUSION.....	284
VIII. BIBLIOGRAPHY.....	287
IX. APPENDIX.....	305

## LIST OF TABLES

Table	Page
I. PROGRAM DATA COMPUTER PRINTOUT.....	96
II. PROGRAM \$READ COMPUTER PRINTOUT.....	107
III. GENERAL STATISTICS OF RESPIRATORY CELLS.....	119
IV. LIST OF PARAMETER ABBREVIATIONS.....	125
V. RANK ORDERING OF CORRELATION COEFFICIENTS FOR THE REGRESSION OF 24 PARAMETERS ON RESPIRATORY RATE.....	154
VI. RANK ORDERING OF CORRELATION COEFFICIENTS FOR THE REGRESSION OF 24 PARAMETERS ON MODE.....	155
VII. CORE MEMORY ASSIGNMENTS FOR PROGRAM DATA.....	306
VIII. REGRESSION EQUATIONS OF 24 PARAMETERS ON RESPIRATION RATE FOR I-VO CELLS.....	389
IX. REGRESSION EQUATIONS OF 24 PARAMETERS ON RESPIRATION RATE FOR I-VI CELLS.....	390
X. REGRESSION EQUATIONS OF 24 PARAMETERS ON RESPIRATION RATE FOR E-VI CELLS.....	391
XI. REGRESSION EQUATIONS OF 24 PARAMETERS ON MODE FOR I-VO CELLS.....	392
XII. REGRESSION EQUATIONS OF 24 PARAMETERS ON MODE FOR I-VI CELLS.....	393

LIST OF TABLES  
(continued)

Table	Page
XIII. REGRESSION EQUATIONS FOR 24 PARAMETERS ON MODE FOR E-VI CELLS.....	394
XIV. BEST FITTING REGRESSION EQUATIONS FOR 24 PARAMETERS ON RESPIRATION RATE AND MODE.....	395

## LIST OF FIGURES

Figure	Page
1. INSTRUMENTAL SCHEMATIC.....	64
2. PARALLEL T NOTCH FILTER.....	68
3. PROGRAM DATA FLOW CHART.....	77
4. REPRESENTATIVE EXAMPLES OF SINGLE CELL ANALOG DATA.....	78
5. TECHNIQUE FOR DETECTION OF ACTION POTENTIALS.....	81
6. INTERSPIKE INTERVAL HISTOGRAM.....	84
7. OGIVE CURVE.....	86
8. GENERATION OF IDEALIZED INTERSPIKE INTERVAL MODULATION CURVE.....	88
9. EXPERIMENTAL INTERSPIKE INTERVAL MODULATION CURVES.....	90
10. IDEALIZED REPRESENTATION OF AIRFLOW VELOCITY, INTRAPLEURAL PRESSURE AND ACTION POTENTIAL DISCHARGE.....	121
11. REGRESSION OF T50 AND T70 ON RR.....	126
12. REGRESSION OF T90 AND T100 ON RR.....	127
13. REGRESSION OF PL AND TL ON RR.....	128
14. REGRESSION OF FI AND MI ON RR.....	129
15. REGRESSION OF LI AND MEN ON RR.....	130

LIST OF FIGURES  
(continued)

Figure	Page
16.	REGRESSION OF MED AND MOD ON RR.....131
17.	REGRESSION OF N50 AND N70 ON RR.....132
18.	REGRESSION OF N90 AND N100 ON RR.....133
19.	REGRESSION OF ST AND SL ON RR.....134
20.	REGRESSION OF T5P AND T7P ON RR.....135
21.	REGRESSION OF T9P AND T10P ON RR.....136
22.	REGRESSION OF PLP AND TLP ON RR.....137
23.	REGRESSION OF T50 AND T70 ON MOD.....138
24.	REGRESSION OF T90 AND T100 ON MOD.....139
25.	REGRESSION OF PL AND TL ON MOD.....140
26.	REGRESSION OF FI AND MI ON MOD.....141
27.	REGRESSION OF LI AND MEN ON MOD.....142
28.	REGRESSION OF MED AND RR ON MOD.....143
29.	REGRESSION OF N50 AND N70 ON MOD.....144
30.	REGRESSION OF N90 AND N100 ON MOD.....145
31.	REGRESSION OF ST AND SL ON MOD.....146
32.	REGRESSION OF T5P AND T7P ON MOD.....147

LIST OF FIGURES  
(continued)

Figure	Page
33.	REGRESSION OF T9P AND T10P ON MOD.....148
34.	REGRESSION OF PLP AND TLP ON MOD.....149
35.	REGRESSION OF T100, T90, T70, AND T50 ON RR AND MOD.....157
36.	REGRESSION OF TL AND PL ON RR AND MOD.....158
37.	REGRESSION OF FI, LI AND MI ON RR AND MOD.....159
38.	REGRESSION OF MEN, MED AND MOD ON RR AND MOD....160
39.	REGRESSION OF N100, N90, N70 AND N50 ON RR AND MOD.....161
40.	REGRESSION OF ST AND RR ON RR AND MOD.....162
41.	REGRESSION OF SL ON RR AND MOD.....163
42.	REGRESSION OF T10P, T9P, T7P AND T5P ON RR AND MOD.....164
43.	REGRESSION OF TLP AND PLP ON RR AND MOD.....165
44.	EFFECT OF SPONTANEOUS INCREASE IN RESPIRATORY DEPTH ON HISTOGRAM AND INTERSPIKE INTERVAL MODULATION CURVE.....169
45.	EFFECT OF SPONTANEOUS DECREASE IN RESPIRATORY DEPTH ON HISTOGRAM AND INTERSPIKE INTERVAL MODULATION CURVE.....171

LIST OF FIGURES  
(continued)

Figure	Page
46.	TWO SIMULTANEOUSLY RECORDED INSPIRATORY CELLS FROM RIGHT AND LEFT MEDULLA, THEIR HISTOGRAMS AND INTERSPIKE INTERVAL MODULATION CURVES.....173
47.	EFFECT OF SPONTANEOUS INCREASE IN RESPIRATORY RATE ON HISTOGRAM AND INTERSPIKE INTERVAL MODULATION CURVE.....175
48.	EFFECT OF DOXAPRAM ON INTERSPIKE INTERVAL MODULATION CURVE.....177
49.	EFFECT OF DOXAPRAM ON PARAMETERS OF INSPIRATORY CELL DISCHARGE.....178-180
50.	EFFECT OF CUMULATIVE DOSES OF PENTOBARBITAL ON HISTOGRAM AND INTERSPIKE INTERVAL MODULATION CURVE.....184
51.	EFFECT OF PENTOBARBITAL ON PARAMETERS OF INSPIRATORY CELL DISCHARGE.....185-187
52.	EFFECT OF CUMULATIVE DOSES OF PENTOBARBITAL ON HISTOGRAM AND INTERSPIKE INTERVAL MODULATION CURVE.....190
53.	EFFECT OF PENTOBARBITAL ON PARAMETERS OF INSPIRATORY CELL DISCHARGE.....191-193
54.	EFFECT OF CUMULATIVE DOSES OF PENTOBARBITAL ON HISTOGRAM AND INTERSPIKE INTERVAL MODULATION CURVE.....195

LIST OF FIGURES  
(continued)

Figure	Page
55.	EFFECT OF PENTOBARBITAL ON PARAMETERS OF INSPIRATORY CELL DISCHARGE.....196-198
56.	EFFECT OF CUMULATIVE DOSES OF PENTOBARBITAL ON HISTOGRAM AND INTERSPIKE INTERVAL MODULATION CURVE.....201
57.	EFFECT OF PENTOBARBITAL ON PARAMETERS OF EXPIRATORY CELL DISCHARGE.....202-204
58.	EFFECT OF CUMULATIVE DOSES OF PENTOBARBITAL ON HISTOGRAM AND INTERSPIKE INTERVAL MODULATION CURVE.....206
59.	EFFECT OF PENTOBARBITAL ON PARAMETERS OF EXPIRATORY CELL DISCHARGE.....207-209
60.	EFFECT OF CUMULATIVE DOSES OF MORPHINE SULPHATE ON HISTOGRAM AND INTERSPIKE INTERVAL MODULATION CURVE.....212
61.	EFFECT OF MORPHINE ON PARAMETERS OF EXPIRATORY CELL DISCHARGE.....213-215
62.	HISTOGRAM AND INTERSPIKE INTERVAL MODULATION CURVES BEFORE AND AFTER UNILATERAL LEFT VAGOTOMY.....217
63.	GENESIS OF CLASSICAL APNEUSIS IN THE CAT.....219
64.	PENTOBARBITAL APNEUSIS.....221



LIST OF FIGURES  
(continued)

Figure	Page
65.	APNEUSTIC PATTERN PRODUCED BY ACUTE CEREBELLECTOMY.....222
66.	MEAN AND SEM OF BARBITURATE DOSAGE AT APNEUSTIC THRESHOLD.....224
67.	EFFECT OF SEQUENTIAL LEFT AND RIGHT VAGOTOMY ON RESPIRATORY PATTERNS.....227
68.	SINGLE INSPIRATORY CELL RECORDING DURING PENTOBARBITAL APNEUSIS.....229
69.	SINGLE EXPIRATORY CELL RECORDING DURING PENTOBARBITAL APNEUSIS.....230
70.	CHANGES IN SINGLE INSPIRATORY CELL HISTOGRAM DURING THIOPIENTAL DEPRESSION OF RESPIRATION RATE.....231
71.	CHANGES IN MEAN, MEDIAN AND MODE AS A FUNCTION OF THIOPIENTAL DOSAGE AND RESPIRATION RATE.....233
72.	EFFECT OF PENTOBARBITAL ON DISCHARGE FREQUENCY OF THREE NON-RESPIRATORY PONTINE CELLS.....235
73.	CHANGES IN RESPIRATORY PATTERNS DURING BILATERAL CAROTID OCCLUSION.....237
74.	CHANGES IN RESPIRATORY PATTERN PRODUCED BY HISTAMINE PHOSPHATE.....239

LIST OF FIGURES  
(continued)

Figure		Page
75.	CHANGES IN RESPIRATORY PATTERN PRODUCED BY DOXAPRAM.....	241
76.	COMPARISON OF CORRELATION COEFFICIENTS FOR 24 PARAMETERS ON RESPIRATION RATE.....	250
77.	COMPARISON OF CORRELATION COEFFICIENTS OF 24 PARAMETERS ON MODE.....	251
78.	COMPARISON OF CORRELATION COEFFICIENTS FOR 24 PARAMETERS ON RESPIRATION RATE AND MODE FOR I-VO CELLS.....	252
79.	COMPARISON OF CORRELATION COEFFICIENTS FOR 24 PARAMETERS ON RESPIRATION RATE AND MODE FOR I-VI CELLS.....	253
80.	COMPARISON OF CORRELATION COEFFICIENTS FOR 24 PARAMETERS ON RESPIRATION RATE AND MODE FOR E-VI CELLS.....	254
81.	THEORETICAL RECONSTRUCTION OF INTERSPIKE INTERVAL MODULATION CURVES FOR I-VO CELLS.....	259
82.	THEORETICAL RECONSTRUCTION OF INTERSPIKE INTERVAL MODULATION CURVES FOR I-VI CELLS.....	260
83.	THEORETICAL RECONSTRUCTION OF INTERSPIKE INTERVAL MODULATION CURVES FOR E-VI CELLS.....	261
84.	ATYPICAL INSPIRATORY CELL RESPONSE DURING INDUCTION OF APNEUSTIC BREATH AFTER ADMINISTRATION OF PENTOBARBITAL.....	277

LIST OF FIGURES  
(continued)

Figure	Page
85.    INTERSPIKE INTERVAL MODULATION CURVES FOR SINGLE TRAIN OF VAGAL AFFERENT AND PHRENIC MOTONEURON.....	281

## LIST OF PROGRAMS

Program	Page
A. MACHINE LANGUAGE PROGRAMS	
1. ERASE.....	307
2. DATA.....	308
B. FOCAL-12 PROGRAMS	
1. \$CALC1.....	366
2. \$CALC2.....	367
3. \$READ.....	368
4. \$SUM1.....	369
5. \$SUM2.....	369
6. \$SUM3.....	370
7. \$SUM4.....	370
8. \$REORD.....	371
9. \$IVO.....	372
10. \$IVI.....	373
11. \$EVO.....	374
12. \$EVI.....	374
13. \$ROWM.....	375
14. \$RLIN.....	376
15. \$REXP.....	377
16. \$RLOG.....	378
17. \$RLLG.....	379
18. \$REGSD.....	380
19. \$RSIGD.....	381
20. \$PVALR.....	382
21. \$STTEST.....	383
22. \$TIMFOL.....	384
C. FOCALPL PROGRAMS	
1. %PLOT.....	385

LIST OF PROGRAMS  
(continued)

Program	Page
2. %REGP1.....	386
3. %REGP2.....	387
4. %LINEP.....	388

CHAPTER I  
INTRODUCTION

Information transfer along central, efferent and afferent arcs of the respiratory system are coded in terms of interspike intervals of single unit spike discharges. At the output stage, respiratory muscles decode the neuronal signals into meaningful outputs of respiratory rate and depth, the two major determinants of alveolar ventilation. However, to follow dynamic changes in information flow in a control systems approach, it is necessary to evaluate mathematically the discharge patterns from various respiratory sub-systems. Comparison of such patterns can yield important information on transfer functions between various respiratory components. The plasticity of these functions during different ventilation demands can also be evaluated.

This study was designed to examine the discharge patterns of single respiratory neurons in the medulla over wide ranges of rate and depth. The combination of

new mathematical criteria and advanced computer techniques has revealed high neuronal correlates of respiratory rate and depth, identified as medullary outputs. This has been achieved by examination of thousands of interspike intervals, their distribution and sequencing.

CHAPTER II  
LITERATURE REVIEW

A. Respiratory Areas of the Medulla

Numerous studies during the last century and a half have implicated the medulla as the primary control area responsible for genesis of the respiratory act. Many technical advances have been made in the study of this system, but differences in opinion concerning medullary organization still exist today. In general, opposing views can be grouped into two broad categories irrespective of the experimental technique or animal used. One theory maintains that respiratory neurons can be assigned specific anatomical localizations within the medullary architecture, with a definite segregation of inspiratory and expiratory neurons. The alternate theory maintains that respiratory cells are freely scattered and intermingled throughout the reticular formation and are functionally tied together via extensive synaptic interconnections. It is important to consider the



evidence for each of these views.

The discrete localization or center theory was the first to gain prominence. Lorry in 1760 (111) confirmed the work of Galen (65) by showing that section of the spinal cord at the first cervical segment immediately arrested respiration in the expiratory position of apnea. This indirectly demonstrated that the genesis of rhythmic respiratory movements was rostral to the cord. Legallois in 1812 (107) supplied the first direct evidence that the medulla possessed respiratory function by showing that rhythmic breathing persisted in young rabbits with removal of the cerebellum and part of the upper bulb. Flourens in 1842 (59) further localized this active lower bulbar region to a one millimeter area at the apex of the calamus scriptorius. Destruction of this "vital node" halted all respiratory movements. Flourens (60) later relocated the "vital node" 2.5 millimeters on either side of the midline at the level of the obex. In 1880, Marckwald and Kronecker (118) attempted to divide the respiratory center into inspiratory and expiratory portions, and Lumsden in 1923 (112, 113, 114) distinguished between an expiratory and gasping center in the medulla of the cat by means of transection procedures.

Active expiratory and inspiratory regions were

mapped with stereotaxic placement of stimulating electrodes. Pitts, Magoun and Ranson (145) explored the brainstem of the cat from the hypothalamus to the cervical cord with a bipolar stimulating electrode and were able to elicit maintained inspiratory or expiratory responses from the medulla. Since maximal responses could not be evoked either above or below the medulla, the authors claimed that their stimulations were selective for cell bodies and not fiber tracts. Expiratory responses were concentrated dorsal, rostral and medial to inspiratory responses, each occupying approximately 30 cubic millimeters of the reticular formation bilaterally, although some overlapping was observed. Pitts (140) reconfirmed functional subdivisions of the respiratory center into inspiratory and expiratory portions in a subsequent publication. Monnier (124) arrived at similar conclusions from his own work.

In the monkey, Beaton and Magoun (23) reported the existence of two discrete regions which when activated produced either inspiratory apneusis or expiratory apnea. The inspiratory activity was located dorsal and medial to the rostral half of the inferior olive. The expiratory activity surrounded the inspiratory field rostrally, caudally, laterally and dorsally. The authors concluded

that this mapping in the monkey corresponded to that in the cat as reported by Pitts et al. (145), but this is questionable. Amoroso, Bell and Rosenberg (6) completed a similar study in the sheep and found that respiratory responses were aggregated in the reticular formation of the medulla from the aroustic tubercles rostrally, to just behind the obex caudally. The expiratory region was located dorsal, anterior and lateral to the inspiratory region and both had bilateral representation.

Ondina, Yamamoto and Masland (136) prepared stimulation maps of the rat brainstem. Respiratory regions identified by sustained inspiratory or expiratory apnea were clearly separated. The inspiratory center was found bilaterally above the rostral one-third of the inferior olive and the somewhat more diffuse expiratory center was located dorsal and caudal to the inspiratory center.

Localization of specific structures in the brainstem by the use of stimulating electrodes met with much criticism since such a large volume of tissue was activated. This technique was defended by Pitts (140) and Magoun and Beaton (115). The latter identified active respiratory sites in the medulla by stimulation, placed a lesion at the same locus and then restimulated. Lesions which extended no more than 0.7 millimeter from

the electrode tip abolished the respiratory response to the second stimulation. The authors used this as a measure of current spread during stimulation procedures.

Comroe (46) attempted to identify respiratory areas of the brainstem by microinjection of CO<sub>2</sub>-bicarbonate solutions into the brain tissue, but met with serious problems localizing the stimulus to a small area. Nevertheless, general areas which yielded immediate hyperpneic responses corresponded to the inspiratory region of Pitts (140, 145). No expiratory responses were seen.

Evidence for separation of inspiratory and expiratory regions in the medulla has also come from extracellular recordings of single respiratory cell discharges. Nelson (127) found 110 single units in the cat and concluded that the expiratory area was dorsal to the inspiratory area. He was careful to record only from cell bodies which could be distinguished from fiber pathways by a technique of wave form identification. Haber, Kohn, Ngai, Holaday and Wang (79), however, suggested a new location for the expiratory center. Recording from single cells, these workers found that the inspiratory region in the cat corresponded to the stimulation maps of Pitts et al. (140, 145), but the expiratory region was located

more caudal with no difference in dorsal-ventral distribution of inspiratory and expiratory areas.

Woldring and Dirken (180) explored the brainstem of the rabbit with extracellular microelectrodes from the caudal border of the corpora quadrigemina to the exit level of the first spinal roots. They found respiratory activity only in an area extending from the obex to a plane 3 millimeters rostral to the obex. In this area, inspiratory and expiratory cells had a definite anatomical distribution. The inspiratory area was located 2.5 millimeters from the dorsal surface and was associated with the ventromedial reticular substance at the level of the vagal rootlets. The expiratory area was located 2.0 millimeters from the dorsal surface and was associated with the dorsolateral reticular formation. The expiratory area appeared to be connected with the spinal trigeminal root laterally and coursed parallel to the solitary tract medially. Batsel (14) published similar results from microelectrode recordings in the cat, dog and monkey. The medulla was explored from 6 millimeters rostral to 4 millimeters caudal to the obex and two bilateral groups of bulbar respiratory cells were identified. The larger area was located in and along the motor nuclei of the ninth and tenth cranial nerves.

The smaller area was located ventrolateral to the fasciculus solitarius, just rostral to the obex. Inspiratory and expiratory cells could not be separated in the dorsal-ventral direction, but the latter were found more frequently in the caudal medulla. Batsel (14) suggested that the recorded electrical activity came from closely packed cells indicative of specialized respiratory centers.

Merril (122) was able to distinguish between two major concentrations of respiratory cells in the medulla by mapping the data from each cat separately. One group was near the nucleus ambiguus and the other was associated with the solitary tract. Achard and Bucher (1) suggested that the lateral respiratory cells were not reticular cells, but rather nucleus ambiguus motoneurons. Von Baumgarten and Kanzow (19) recorded inspiratory potentials from the reticular formation just ventral to the tractus solitarius and 1-3 millimeters rostral to the obex. Nesland, Plum, Nelson and Siedler (129) occasionally recorded from expiratory and inspiratory cells at the same location, but reported that cells with expiratory phasings were localized more caudal and dorsal to the inspiratory cells.

In contrast to the foregoing, other experimental evidence suggests that medullary neurons are diffusely

organized and do not exhibit anatomical localization of inspiratory and expiratory areas. Longet in 1847 (110) showed that respiration was not affected by localized destruction of the pyramids and restiform bodies. However, destruction of the reticular formation at the same level immediately halted respiration. Gad and Marinesco in 1892 (64) confirmed this observation by cauterizing certain areas of the medulla. Only when the reticular substance of the floor of the fourth ventricle was destroyed did breathing stop. Arnheim (8) was also of the opinion that the bulbar neurons responsible for the generation of respiration were located diffusely in the reticular gray matter.

The stimulation technique has been used to show diffuse organization. Brookhart (34) stimulated the brainstem of dogs from 9 millimeters above to 6 millimeters below the obex with microelectrodes. With low voltage stimulations he got variable respiratory rate changes which were not correlated with any structure. The results failed to confirm the existence of compact inspiratory and expiratory centers described by Pitts et al. (140, 145) in the cat. Brookhart (34) suggested that the reticular formation was the seat of primary respiratory cells without detectable anatomical

differentiation of function. Liljestrand (108) studied in depth the respiratory movements which could be induced by stimulation of the medulla both electrically and chemically. He stated that inspiratory and expiratory areas were not separated and in his own words added, "Until further evidence of a respiratory centre has accumulated the term should be strictly defined when used, or otherwise it should be avoided."

Kim and Carpenter (100) used a different stimulation technique to induce respiratory movements. Ventilation changes comparable to electrical activation of the same area could be produced by injections of isotonic solutions of  $\text{HCO}_3^-$ ,  $\text{HPC}_4^-$  or citrate into the medullary reticular formation of the cat. The apneustic responses observed could be evoked from random reticular formation sites and the authors concluded that chemical stimulation of respiratory neurons represents a functionally non-specific phenomenon.

The extracellular recording technique has also been used to demonstrate diffuse medullary organization. Gesell, Bricker and Magee (70) published the first report of single respiratory cell recordings. The medulla, upper cervical cord, thalamus, hypothalamus, midbrain and pons of the dog were explored with bipolar needle



electrodes. Areas yielding rhythmic respiratory discharges were lesioned for histological identification. In all the regions examined, respiratory potentials were found most frequently in the reticular gray in the region of the obex, but no correlation could be made associating cell type with any one particular brainstem structure, either nucleus or tract. The organization of inspiratory and expiratory regions appeared to be diffuse. The authors commented that the reticular gray is influenced by every type of sensory signal, which may explain activation of the respiratory system during different situations. Amoroso, Bainbridge, Bell, Lawn and Rosenberg (5) used metal microelectrodes to study single respiratory cell discharges in the dog, cat and rat. These electrodes recorded from a brain tissue volume which was one thousandth of that activated by a small stimulating electrode. Because of this high specificity, respiratory potentials were difficult to locate. The authors wrote, "The scarcity of respiratory potentials in the reticular formation of the medulla suggests that generally this structure does not function as a whole but only in parts. It seems to act as a trigger, periodically firing spinal, phylogenetically older, neuronal aggregates which are ultimately responsible for the

coordinated respiratory movements."

Hukuhara, Nakayama and Okada (91) recorded respiratory potentials from the striae acousticae level in the lateral reticular formation of the dog and cat brain and suggested that this was the normal respiratory center. Although this localization did not agree with that of Gesell et al. (70), no differentiation of inspiratory and expiratory areas was possible. Hukuhara et al. (91) also discussed the coursing of centrifugal respiratory pathways through the lateral reticular formation of the medulla. Salmoiraghi and Burns (155) prepared a map of the brainstem localizing respiratory discharges in the cat. While no respiratory cells were found in the pons, potentials in phase with the respiratory cycle were localized in the medulla 2-4 millimeters below the dorsal surface on either side of the obex. Inspiratory and expiratory cells could not be anatomically distinguished and were sometimes found within 100 microns of each other. Brodie (31) reported that one electrode could record activity from both inspiratory and expiratory cells at the same time and concluded that these cell types were intermingled. Harris (81) came to the same conclusion.

Finally, von Baumgarten and Salmoiraghi (21)

studied the localization of respiratory neurons in the goldfish via electrical recording. They found that respiratory cells were bilaterally organized in narrow bands extending rostrally from the mid-vagal lobes. The cells with opposite phasing patterns corresponding to inspiratory and expiratory units were not segregated into different areas. The cells laid close to the segmental columns of the motor nuclei from the fifth to twelfth cranial nerves.

Conclusions defining strict medullary organization of the respiratory system are difficult to make. From the clinical work of Baker, Matzke and Brown (11), respiratory failure during bulbar poliomyelitis definitely involves pathologic alterations in cells of the nucleus ambiguus although similar structural changes are also seen among other large and small reticular cells, many of which must also subserve a respiratory function. Weighing the evidence presented, it can be suggested that any concept of medullary organization must simultaneously explain the following two characteristics. First, inspiratory cells are anatomically scattered throughout the medullary reticular formation apparently without any logical ordering. Second, inspiratory and expiratory divisions of the respiratory system each function in a

highly integrated fashion. Obviously, the respiratory representation in the medulla poses interesting structure-function problems.

B. Apneusis

Knowledge concerning the organization of the brain-stem respiratory system has been extended by many studies into the phenomenon of apneusis. It is now clear that medullary structures alone are not responsible for the coordinated act of respiration. The pons appears to contain two general populations of neurons which influence medullary respiratory cell discharge to respiratory motoneurons.

It is interesting that the definition of apneusis has slowly changed in the literature. When the concept was first introduced (116, 117) it was associated with maximal inspiratory cramps which were maintained until the preparation died from asphxia. Later, apneusis was described as an abnormal respiratory pattern characterized by periodic escape from sustained inspiratory efforts (26, 87, 112). Workers attributed this pattern to a specific cyclic function of the respiratory system which could be maintained for long periods of time. Other investigators (112) redefined apneusis as cyclic submaximal inspiratory holds. This, along with the

observation that apneusis was not a permanent pattern, prompted the suggestion that apneusis was the result of non-specific facilitation of medullary respiratory areas (26, 87). These differences in length and depth of inspiratory holds and duration of apneusis are a reflection of the various techniques used to induce this unusual breathing pattern.

Marckwald in 1887 (116) and 1890 (117) observed long inspiratory cramps in vagotomized rabbits if the brainstem was transected just behind the posterior colliculi. These inspiratory holds continued uninterrupted until asphyxial death if the trigeminal nerves were damaged. Vagotomy in preparations with transections placed just in front of the posterior colliculi failed to develop the characteristic inspiratory movement. Breathing simply became deeper and slower as seen after vagotomy in animals with intact brainstems. Marckwald (116, 117) concluded that the posterior colliculi contained a center inhibitory to inspiration. Loewy in 1888 (109) confirmed this observation.

Lumsden (112, 113, 114), using the transection technique, made major contributions on pontine organization. In one paper (112) he showed that vagotomy in cats with brainstem sectioned immediately behind the

posterior colliculi did not produce the inspiratory cramps described by Marckwald (116, 117). The inspiratory holds or apneusis, as Lumsden (112) called them (Greek for a holding of the breath), occurred only in cats with transections a few millimeters below the upper border of the pons when the vagi were cut. This apneusis was converted into a gasping respiration when another transection was introduced below the striae acousticae. The conclusions were obvious. That is, the lower pons contained an apneustic center which drove the gasping center of the medulla continuously when inhibition from the pneumotaxic center in the upper pons and vagi were removed. In another article, Lumsden (113) discussed his concept of an expiratory center in the medulla in order to explain active forced expirations and summarized his thoughts in a third publication (114) relating vagal impulses to central respiratory organization and rhythmic breathing.

An alternate view of apneusis was presented by Henderson and Sweet (83) and Hess (85) who suggested that the phenomenon was a manifestation of decerebrate rigidity. The authors explained that the apneusis they saw in mid-pontine cats with cold blocked vagi was due to tonic activity of medullary centers released from red nucleus inhibition by severing of the rubro-spinal

tracts. The vagi were believed to inhibit inspiration via direct inhibition of the general brainstem facilitatory mechanism. Hoff and Breckenridge (26, 87) also concluded that apneusis was best regarded as decerebrate rigidity of the respiratory system. They made the following major observations in the dog and cat. First, apneusis was not permanent as suggested by Marckwald (116, 117), but was an oscillating respiratory pattern. Second, apneusis was not total since alternating inspiration-expiration efforts were often seen superimposed on apneustic breaths. Third, as preparations deteriorated, apneusis was replaced by relatively normal respiratory patterns. Last, apneusis could be abolished by separation of the medulla from the pons as described by Lumsden (112). Breckenridge, Hoff and Smith (28) also showed that intravenously administered myanesin, a drug which blocks decerebrate rigidity, also converted apneustic breathing into normal respiratory patterns.

Stella (167) refuted the conclusions of Henderson and Sweet (83), Hess (85) and Hoff and Breckenridge (26, 87) and maintained that apneusis was a specific respiratory event. He showed that elimination of the red nucleus and/or posterior colliculi did not alter normal respiratory patterns. Apneusis resulted only when the upper pons

was removed in vagotomized preparations. This confirmed the inhibitory function of the upper pons (pneumotaxic area) first introduced by Lumsden (112). Stella (167) also demonstrated that apneusis was not dependent on proprioceptive reflexes from the respiratory muscles by showing that apneusis persisted in a deafferented brainstem preparation. This, of course, differentiated apneusis from decerebrate rigidity, since the latter could be terminated by dorsal root section.

Wang, Ngai and Frumin (176) reexamined the results of Breckenridge, Hoff, and Smith (28) who abolished both decerebrate rigidity and apneusis with myanesin. By using low doses of this drug, Wang et al. (176) were able to differentially block body rigidity without altering the apneustic breathing pattern. The authors interpreted these results as inferring that apneusis and decerebrate rigidity were two separate phenomena. Ondina, Yamamoto and Masland (136) qualified these results by suggesting that if apneusis were a phenomenon of bulbar facilitation, the sources of this facilitation did not lie in the same cells as those which facilitated the postural musculature. They demonstrated that in the rat, an animal which rarely displays decerebrate rigidity, apneusis can readily be induced by brainstem transection



at the mid-pons level and vagotomy without the manifestation of body rigidity.

Electrophysiological evidence supports the notion of a pontine apneustic area. Ngai and Wang (135) used the stimulation technique in cats with intact brainstems. They found that stimulation from a level 3 millimeters behind the rostral border of the pons extending caudally to the trapezoid body interrupted rhythmic respiration with a maximal inspiratory movement. The active brainstem area corresponded to Lumsden's (112) apneustic center. Kahn and Wang (96) recorded from single inspiratory and expiratory cells in midpontine cats and monitored the phrenic discharge. After vagotomy, inspiratory cells and phrenic patterns became continuous as apneustic breathing was initiated. Expiratory cells discharged only during the brief interruptions of inspiratory holds. In a central vagal stimulation study, Kerr, Dunlop, Best and Mullner (99) assigned the apneustic area to a supramedullary level at the origin of the eighth cranial nerve just lateral to the midline. The authors suggested that the apneustic area may have some involvement with the vestibular system.

The removal of pneumotaxic and vagal inhibitions is important for the unmasking of apneustic drive.

Consequently, apneusis has been used as an endpoint in vagotomized preparations to locate pneumotaxic areas in the upper pons by tissue destruction. Pitts, Magoun and Ranson (147) made bilateral electrolytic lesions in the cat brainstem with stereotaxic placement of electrodes and localized the pneumotaxic area to the tegmentum of the upper few millimeters of the pons. Lesions in the inferior colliculi and red nuclei did not result in apneusis thereby refuting the work of Marckwald (116, 117) and Henderson and Sweet (83) respectively. Tang and Ruch (172) localized the pneumotaxic area in the dorso-lateral portion of the rostral pons by suction destruction or electrolysis. Tang (170) later refined the location to the extreme dorsolateral portion of the anterior pontine tegmentum, the isthmus. Ngai and Wang (134) associated the pneumotaxic region with the dorso-lateral reticular formation since stimulation in this area accelerated respiratory rate and destruction led to apneusis when the vagi were cut. Johnson (93) found similar results and placed the pneumotaxic area in the latero-dorsal tegmental nucleus or locus coeruleus. Baxter and Olszewski (22) also suggested that the cells of the locus coeruleus nucleus formed a major part of the pneumotaxic area.

Using the above localization data for the pneumotaxic center, St. John, Glasser and King (151) placed chronic lesions in the dorsolateral pontine regions of the cat. Rhythmic respiration was observed in all cats after recovery from anesthesia, but when the vagi were severed up to thirteen weeks later, cats with pneumotaxic center lesions immediately developed apneustic patterns. Cats with control lesions a few millimeters anterior or posterior to the pneumotaxic level did not develop apneusis. The authors concluded that their lesions chronically knocked out pneumotaxic participation in the genesis of eupnea. St. John, Glasser and King (152) later showed that vagotomized cats with chronic pneumotaxic lesions were capable of normal respiratory rhythmicity when awake. This demonstrated that higher regions of the central nervous system could maintain respiratory rhythmicity in the absence of pneumotaxic or vagi inhibitions.

From these accumulated data it has been concluded that the pneumotaxic and apneustic areas of the brainstem are directly involved with respiratory mechanisms. Unlike medullary inspiratory and expiratory areas, the pontine regions appear to be anatomically discrete.

The length and depth of apneustic breaths depend

on a number of factors besides the removal of pneumo-  
taxic and vagal afferent inhibition. Additional factors  
to be considered are the chemical status of the prepara-  
tion, integrity of cerebellar structures and the anes-  
thetic drug type and level.

Stella (168) produced apneustic breathing in cats  
by transecting at the mid-pontine region and cold-blocking  
the vagus nerves. When the  $\text{CO}_2$  content of the inhaled air  
was elevated to 8-10%, apneustic breaths became much  
deeper and prolonged. These results could be duplicated  
by using a rebreathing procedure. Hoff and Breckenridge  
(87) found that peripheral chemoreceptor stimulation by  
carotid occlusion prolonged or intensified the apneustic  
pattern in dogs. Also, vagotomy failed to induce apneu-  
sis in mid-pontine dogs with carotid denervation. The  
authors suggested from these results that apneusis de-  
rives from the interplay of peripheral carotid body  
and central lower pontine factors.

Ngai (130) determined the  $\text{O}_2$  consumption, arterial  
blood pH,  $\text{PaCO}_2$ ,  $\text{PaO}_2$  and  $\text{O}_2$  saturation during apneustic  
breathing in cats. Compared with control respiratory  
values, apneusis induced a respiratory acidosis with no  
change in the  $\text{PaO}_2$  or  $\text{O}_2$  consumption (cats breathed oxygen  
rich mixtures). Administration of 10%  $\text{CO}_2$  increased the

depth of each apneustic breath and accelerated the apneustic cycle. The latter finding is in disagreement with that reported by Stella (168). Since this response persisted after carotid denervation, Ngai (130) concluded that apneustic cycling was probably due to the elevated arterial  $\text{CO}_2$  tensions and was independent of the peripheral chemoreceptors.

Katz, Ngai, Nahas and Wang (98) studied the effect of pH alterations on apneusis. Two agents which elevate the pH (THAM, an organic buffer, and sodium bicarbonate) were infused intravenously in mid-pontine vagotomized cats displaying apneustic respiration. Neither agent altered the depth of apneusis, but THAM decreased the apneustic cycling while sodium bicarbonate increased the cycling. These apparently conflicting results were explained on the basis of possible differential distribution of the two agents across cell membranes. The authors suggested that if this were the case, the intracellular pH may be the critical factor for modulation of apneusis.

The effect of cyanide on apneusis was studied by Brodie and Borison (32). They showed that cyanide can convert apneustic breathing into a gasping pattern, a response which was independent of peripheral chemoreceptor

denervation. Ngai (130) reported that intracarotid injection of sodium cyanide can accelerate apneustic cycling thus demonstrating that cyanide does exert a peripheral influence also.

Although it is not commonly recognized, the cerebellum appears to have some connections with the brainstem respiratory system. Henderson and Sweet (83) mentioned that cerebellectomy could produce apneusis which disappeared after a short period. Using mid-pontine cats, Glasser, Tippett and Davidian (76) studied respiratory pattern changes during cerebellar depression induced by several techniques: occlusion of the cerebellar arteries, procaine or xylocaine application to the surface or sub-cortex of the cerebellum, or removal of part or all of the cerebellum by suction. Irrespective of the method used, cerebellar depression in mid-pontine cats with intact vagi led to a fall in respiratory rate and an increase in respiratory depth to the point of apneusis in many cases. The authors concluded that the cerebellum exerted a tonic inhibitory influence on the brainstem respiratory mechanism. The inhibition might have arisen in the anterior and posterior lobes of the corpus cerebelli, but brainstem projections were unknown. Another interesting report from Gesell, Bricker and Magee (70)

suggested that the cerebellum may contribute to the integration of the respiratory act. They were able to locate three single cell potentials in the region of the brachium conjunctivum (superior cerebellar penduncle) which had respiratory discharge patterns. It is possible that these were efferent fiber tracts from the cerebellum to the pons or medulla mediating respiratory inhibition.

There is a considerable body of evidence which indicates that high levels of barbiturate can convert normal respiratory patterns into apneustic ones even when all brainstem structures and vagi are intact. Harris and Borison (82) anesthetized cats with 36 mg/Kg pentobarbital and stimulated inspiratory regions of the medulla during pentobarbital titration until the spontaneous respiration was arrested. With progressively higher doses, the threshold to produce apneusis via medullary stimulation consistently decreased. At an accumulated dose of 45 mg/Kg pentobarbital, the respiratory cycle was lengthened with a progressive increase in the expiratory phase. With higher doses approaching the respiratory depression level of pentobarbital (mean 50 mg/Kg, range 33-70 mg/Kg) inspiratory holds of apneusis were observed in preparations with intact brainstems and vagus nerves. These results showed that pentobarbital selectively depressed systems

which were inhibitory to inspiration. The inspiratory apparatus was presumably depressed to some extent since apneustic breaths were of low amplitude.

A similar study was done by Ngai (131) who extended stimulation sites to the pons during pentobarbital accumulation in the midcollicular, vagotomized cat. At doses from 24 to 30 mg/Kg the inspiratory movements became apneustic in type except for a greatly reduced depth. At this same dose level, stimulation of the pneumotaxic region failed to accelerate respiratory rate as in control cases, but central vagal stimulation was still effective. The author concluded that pentobarbital produced apneusis in the decerebrate vagotomized cat by blocking pneumotaxic inhibition of the apneustic areas. Comparison of Ngai's (131) data with those of Harris and Borison (82) reveals that the pneumotaxic mechanism is more sensitive to pentobarbital depression than is vagal afferent feedback. St. John, Glasser and King (151) reported that cats with chronic pneumotaxic lesions, anesthetized with 35 mg/Kg pentobarbital, repeatedly had bouts of apneusis following small additional doses of pentobarbital and bilateral vagotomy. A similar prolongation of the inspiratory phase with thiopental has been published by Brodie (31).

Korczyn, Leibowitz and Bergmann (104) studied the



effect of pentobarbitone dose level on the respiratory response evoked by sciatic nerve stimulation. Control responses ranging from acceleration at low frequency stimulation to deceleration of breathing at high frequency stimulation could all be blocked by high doses of the barbiturate. These data parallel the blockade of vagal inhibition in the experiments of Harris and Borison (82) who observed apneusis in vagi intact cats with high doses of pentobarbital.

Robson, Houseley and Solis-Quiroga (148) studied barbiturate effects on respiration at the single neuron level. Successive increments of sodium pentobarbital or sodium thiopental in cats caused either the inspiratory or expiratory cell discharges to fire continuously. The inspiratory cell pattern resembled that of apneusis. The authors explained that the elevated excitatory firing of expiratory cells was due to barbiturate arrest of mutually inhibitory pathways between inspiratory and expiratory neurons.

Finally, Naifeh, Huggins and Hoff (125) studied respiratory pattern changes in crocodiles during pentobarbital accumulation. The authors termed this technique "anesthetic dissection," but were not able to induce apneusis as observed in other animal preparations under

similar conditions. These results were explained by Naifeh, Huggins and Hoff (126) in another publication in which serial transections were made in the crocodile brainstem. No evidence was found for an apneustic area in this animal. This is a good example of species differentiation in brainstem organization of the respiratory control system.

### C. Medullary Outputs

The primary function of respiration is to maintain arterial blood oxygen, carbon dioxide and hydrogen ion concentrations in homeostatic proportions. To do this, metabolic drives determine the proper alveolar ventilation for the situation at hand by adjustment of respiratory rate and tidal volume, the two outputs from the brainstem respiratory complex. Although there are numerous respiration rate - tidal volume combinations which can generate a specific alveolar ventilation, workers such as Rohrer (149) and Otis, Fenn and Rahn (137) have suggested that the respiratory frequency is selected so as to minimize the amount of work expended by the respiratory musculature. Alternately, Mead (120, 121) has proposed that breathing is adjusted to minimize the muscle force instead of the work. In either case, any acute or chronic change in the mechanical properties of the chest or lung should manifest itself by a different optimal

breathing frequency. Also, removal of major compliance feedback via vagotomy should increase the work of breathing. Zechman, Salzano and Hall (184) confirmed this by showing that the regulation of rate and depth of breathing was less efficient when pulmonary afferents were blocked. Salzano and Hall (157) later reported that vagal reflexes did not influence the work of breathing during severe hypoxia or hypocapnia as much as they did during normal or obstructive breathing.

It is difficult to rank medullary outputs in order of primary (active) or secondary (passive) control. Many experiments have shown that rate and depth components are highly integrated and only a few have succeeded in separating the outputs into two component systems. The following discussion will consider system interactions responsible for the genesis of rate and depth outputs.

The respiratory rate output presents itself as a two-fold problem. First, generation of the basis rhythmicity must be explained and second, basic mechanisms which modulate this rhythm need to be examined. Hoff and Breckenridge (88) have classified breathing into three normal patterns (eupnea, sighing and panting) and three abnormal patterns (Cheyne-Stokes, Biots and apneusis). Each pattern is characterized by a specific rate-depth

combination which derives from medullary, pontine and vagal afferent interactions.

A large body of experimental evidence suggests that respiratory systems of the medulla are spontaneously rhythmic. One of the first observations came from Lumsden (112) who showed that medullary cats gasped rhythmically. Stella (167) reported that regular and smooth respiration was seen in the cat after thorough deafferentation of the medullary centers and concluded that the pattern was due to automaticity of the centers involved. Hoff and Breckenridge (87) showed that a dog can continue to breathe for 5 to 6 hours in an approximately normal fashion after having been deprived of all pontine and vagal connections. The authors suggested that the rhythm was maintained via reciprocal innervation of inspiratory and expiratory neurons. In another paper, Breckenridge and Hoff (26) repeated these experiments in the cat. Again, medullary preparations showed rates and depths similar to those seen in midcollicular preparations before vagotomy. The authors concluded that medullary centers were automatic and were regulated only secondarily by facilitatory and inhibitory areas of the brain. Breckenridge and Hoff (27) later showed that breathing could be maintained reflexly when medullary centers were depressed to apnea

following drug administration or anoxia.

Other workers have agreed that the medulla possesses inherent rhythmicity. Ngai, Frumin and Wang (133) wrote, "The medullary respiratory centers have an autonomous rhythm when released from the influence of pneumotaxic and other pontine centers and afferent vagal impulses." Tang (170) wrote, "It thus appears that vagal and pneumotaxic afferent influx is not the sole determinant of respiratory periodicity for rhythmic breathing persists after elimination of such influx."

Brodie and Borison (33) attempted to study the relationship between gasping mechanisms and the generation of other rhythmic forms of respiration. In decerebrate cats, gasping could be induced by stimulation of the floor of the fourth ventricle a few millimeters rostral to the obex. Since ablation of Pitts' (145) expiratory region did not abolish this gasping response, the workers suggested that the gasping rhythm was not due to reciprocal connections between inspiratory and expiratory neurons. An alternate interpretation is that all expiratory cells were not destroyed.

Ondina, Yamamoto and Masland (136) observed the medullary rat for a prolonged time and described the development of respiratory incoordination or ataxia

consisting of a dissociation of inspiration-expiration sequencing. The authors suggested that this might have been due to altered linkage between loosely associated inspiratory and expiratory systems in the cat medulla.

In a series of stimulation experiments, Borison (25) described a spasmodic respiratory response which could be evoked from activation of the dorsolateral region of the myelencephalon. The response consisted of a series of strong inspiratory and expiratory movements every one to four seconds. Borison (25) suggested that medullary cats without pontine and vagal connections may rhythmically breathe or gasp because of oscillatory drives from this spasmodic center.

Other workers have suggested that gasping is not a normal respiratory pattern. Hukuhara and Nakayama (90) showed that eupneic-like gasping did not develop immediately following isolation of the medulla, but depended on deterioration of the animal. Barcroft (13) described gasping as an all or none type of breathing in which inspiratory depth appeared to be maximum. Brodie and Borison (32) showed that gasping rate was not increased by chemoreceptor activation at low  $\text{PaO}_2$ . Ngai (130) showed that gasping provided inadequate ventilation for the cat, resulting in respiratory acidosis. From this

evidence it is concluded that medullary areas are spontaneously rhythmic, but the isolated medulla is incapable of providing adequate ventilation.

Kahn and Wang (96) emphasized that pontine circuits must be accounted for in any explanation of the genesis of eupneic respiration. Coordinated oscillations of pontine and medullary structures was demonstrated by Adrian and Buytendijk (4) in isolated goldfish brains. They recorded gross potential discharges from the vagal lobes which oscillated at frequencies close to the gill movement rhythm in the intact fish. This paper demonstrated that rhythmicity can be present in the absence of afferent input, although the frequency did depend on the  $O_2$  and  $CO_2$  tensions in the surrounding fluid.

Mechanical afferent input can be eliminated by pharmacological blockade of the respiratory musculature. Von Baumgarten and Salmoiraghi (21) recorded rhythmic cell discharges from respiratory neurons in goldfish treated with succinylcholine to produce myoneural block. Gesell, Atkinson and Brown (69) showed that curare-induced motor paralysis in the dog did not block rhythmic phrenic discharges. Kahn and Wang (94) reported the same results in gallamine-blocked cats. Joels and Samueloff (92) found that the recurrent laryngeal nerve still

exhibited rhythmic bursts of action potentials in the dog and cat whose respiratory movements had been blocked with succinylcholine.

While there is general agreement that midcollicular-decerebrate and vagotomized preparations exhibit rhythmic respiratory movements, the mechanism for the genesis of these oscillations is still disputed, mainly due to differences of opinion on pontine-medullary organization. Lumsden (113) attributed rhythmicity in the mammal to an interaction between his apneustic and expiratory centers which were both under the control of the higher pneumotaxic center. He suggested that the gasping center was overridden by pontine influence when both pons and medulla were intact.

Pitts, Magoun and Ranson (147) suggested that medullary centers were not inherently rhythmic since their medullary cats displayed apneusis. Rhythmic breathing was believed to be due to periodic inhibition of the tonically active inspiratory areas via pneumotaxic or vagal afferent feedback. Although rhythmicity could be explained on the basis that the pneumotaxic system was either spontaneously rhythmic or reflexly activated via proprioceptive afferents like the vagal mechanism, the authors favored the concept that the inspiratory area



sent collaterals up to the pneumotaxic region to activate inhibitory outflow. The reciprocal interchange between these two areas could produce rhythmic respiration even when the vagi were severed.

Wang, Ngai and Frumin (176) modified the schema of Pitts et al. (147), in whose experiments a portion of the lower pontine area had been left intact in the preparation of medullary animals. Wang et al. (176) made a definite distinction between the apneustic region of the lower pons and the medullary inspiratory area. Rhythmicity was believed to result from periodic inhibition of the apneustic region by vagal afferents and by the pneumotaxic area which was activated via apneustic center output. The authors suggested that inspiratory areas of the medulla, passive by themselves, were driven by apneustic outflow.

The pneumotaxic region may exert some modulatory control on respiratory rate since both Ngai and Wang (134) and Johnson (93) have shown that stimulation of the dorsolateral reticular formation of the pons accelerated respiratory rate. Kahn and Wang (95) studied the role of the pneumotaxic area in the establishment of rhythmicity by examining phrenic nerve discharges. Bilateral vagotomy had only slight effect on the phrenic discharge

pattern. Mid-pontine transection, however, greatly modified the discharge which became semi-continuous, yet still oscillatory. It was concluded that the pneumotaxic center provided important information for respiratory manifestation of eupnea and rate-setting control. Cohen and Gootman (44) suggested that the pneumotaxic region may be involved in reverberating mechanisms of respiratory rhythmicity in the brainstem.

Further evidence of pneumotaxic function has come from two different laboratories although both substantiate each other. First, Bertrand and Hugelin (24) delivered single or double stimuli, given at random within the respiratory cycle, to the nucleus parabrachialis medialis of the upper lateral pons. They found that the phrenic discharge could be synchronized by this pneumotaxic stimulation suggesting direct excitation of a self-reexciting system acting as a pacemaker. Cohen (43) brought the evidence one step further and was able to differentiate between two pneumotaxic functions: (1) stimulation of the dorsolateral pons facilitated phrenic discharge with earlier burst termination and decreased the expiratory phase; (2) stimulation of the ventrolateral pons reduced the phrenic discharge with an even earlier burst termination and prolonged the expiratory phase. The phase

switching response was shown to be dependent on stimulus current, frequency, total number of stimuli, and the timing of stimulation in relation to the respiratory cycle. Cohen (43) concluded that the pneumotaxic region may function to smooth transitions from inspiration to expiration and vice versa.

Tang (171) reported that destruction of the pneumotaxic center in cats with intact vagi depressed the depth response to  $\text{CO}_2$ . The respiration rate increase evoked by  $\text{CO}_2$  was essentially unaffected. Tang (171) suggested that the pneumotaxic mechanism was responsible for depth modulation and could maintain constant rates in absence of vagal feedback. The pneumotaxic control was believed to operate through the apneustic system. St. John (150) confirmed this observation in conscious cats with either unilateral or bilateral chronic lesions placed in the pneumotaxic region. These cats displayed diminished tidal volume responses to hypercapnia. The tidal volume response to hypoxia was unaffected, however, and St. John (150) concluded that other regions of the central nervous system had tidal volume regulating capability in the conscious animal.

For a long time it has been recognized that bilateral vagotomy results in slower and deeper breathing.

As early as 1868, Hering and Breuer (84) and Breuer (29) demonstrated that inflation and deflation reflexes were mediated via vagal afferents and concluded that the oscillatory nature of breathing was regulated by pulmonary reflexes. Widdicomb (178, 179) has reviewed a multiplicity of reflexes which are carried in the vagu nerves, but one, the inflation reflex, has prime importance in directly modifying respiratory rate and depth. Consequently, much evidence has been compiled on this reflex concerning receptor site, afferent path, anatomical and functional central projections and involvement in rate and depth control.

Receptors mediating the inflation reflex have been found in the small and large air passages in the lungs. Whitteridge and Bülbring (177) assigned a bronchial or bronchiolar location to the receptors and Hammouda and Wilson (80) showed reflex inhibition of inspiration by tracheal distension. Davis, Fowler and Lambert (49) found that many pulmonary afferents exhibited discharge patterns which paralleled airway pressures more closely than lung volume changes and concluded that the receptors were located in the air passages. An interesting confirmation of this came from Aviado and Schmidt (10) who showed that steam inhalation blocked the inflation reflex

without any apparent alveolar damage.

The afferent nervous pathway for the inflation reflex has been established as vagal. Hering and Breuer (84) and Steffensen, Brookhart and Gesell (166) showed that the inflation reflex could be blocked in the dog, cat and rabbit by cooling the vagi to 8°C while other vagal reflexes were left unimpaired. Paintal (138) identified fibers mediating the inflation reflex as belonging to the A $\alpha$  and A $\beta$  groups. Einthoven (52) in 1908, studied vagal afferent action currents from the cut peripheral vagus nerve with the string galvanometer and showed that discharge frequency increased with inflation of the lungs. Adrian (2) recorded from single inflation afferents using the new fiber splitting technique. Discharges were identified by lung inflation and receptors were found to be slowly-adapting to maintained lung inflations. Knowlton and Larrabee (103) confirmed this observation, but also described a second rapidly-adapting receptor discharge.

Inflation afferents project to the brainstem in the rostral vagal rootlets and distribute to the vagal nucleus and solitary tract of the cat as shown by Foley and DuBois (61). Wyss (182) showed that central vagal stimulation responses could be blocked by localized lesions in the tractus solitarius. He concluded that the

tractus solitarius is probably the locus proper of vagal respiratory reflex centers. Harris (81) applied single shocks to the vagi and recorded maximal evoked potentials in the medullary tractus solitarius, nucleus solitarius and nucleus ambiguus. Wyss, Anderegg and Oberholzer (183) suggested that the afferent fibers synapse in the solitary nucleus and then travel caudally in the solitary tract. Culberson and Kimmel (48) found degenerated vagal afferents which coursed dorsomedially through the medulla, passed through the spinal trigeminal tract and entered the tractus solitarius. These connections may interface respiratory reflex afferents with inspiratory and/or expiratory areas of the medulla.

Other workers believe that vagal afferent fibers project to the apneustic region in the lower pons. Kerr, Dunlop, Best and Mullner (99) observed that apneustic patterns in mid-pontine vagotomized rabbits could easily be modified by central vagal stimulation. Wang, Ngai and Frumin (176) found that vagotomy failed to alter gasping patterns in medullary cats. However, Hoff and Breckenridge (88) showed that vagotomy could release eupnea by increasing rate and depth, provided vagal inflow was initially intact in the medullary dog. Finally, Ngai and Wang (135) showed that upper pontine transection or removal of the

pneumotaxic area had very little effect on activity of the inflation reflex. From all of this evidence, it is believed that respiratory vagal afferents have an influence confined to medullary and lower pontine brainstem structures.

Evidence has accumulated that vagal afferents affect both the inspiratory and expiratory phases of respiration. Wyss (181) was able to produce opposite respiratory responses in the monkey by stimulating the central vagus at different frequencies. Low frequency stimulations (30-80 Hz) caused slight but distinct inspiratory effects with rate acceleration. High frequency stimulations (100-400 Hz) caused marked respiratory slowing with shifts favoring expiratory movements. These results were ascribed to selective activation of the inflation and deflation reflexes respectively. Pitts, Magoun and Ranson (146), using constant stimulating frequency of 240 Hz were able to evoke inspiratory or expiratory movements, respectively, by low or high intensity stimulation of the central vagus of the cat. In either case, stimulation of the inspiratory or expiratory center would override vagal stimulation effects. Pitts et al. (146) discussed results in terms of reciprocal inhibition between inspiratory and expiratory systems.

Gesell and Worzniak (74) made the comment that either inspiration or expiration may coincide with pulmonary inflation by a pump. Dirken and Woldring (51) investigated this further by occlusion of the airways during either the inspiratory or expiratory phase of spontaneous respiration in the rabbit. Occlusion during inspiration led to a gradual decrease in lung volume with a subsequent decrease in vagal afferent discharge frequency. Occlusion during expiration produced just the opposite results. Also, inspiratory occlusion tended to elevate the discharge frequency of expiratory cells, but had no effect on inspiratory cell frequency. Blockade of the vagus by anelectrotonus depressed expiratory discharges but enhanced inspiratory discharges. Stimulation of the central vagus at 9 Hz depressed inspiratory cells and at 200 Hz inspiratory cells were completely inhibited. Expiratory cells tended to go into continuous patterns during vagal stimulation at both frequencies. Dirken et al. (51) came to the conclusion that vagal afferents activate expiratory cells, while they reciprocally inhibit inspiratory cells.

Finally, Gesell, Atkinson and Brown (69) suggested that the act of respiration may be coordinated by vagal afferents that shift their drive to the quiescent medullary



center. This could explain the smooth transition between inspiratory and expiratory phases.

The vagal inflation reflex is of prime importance for control of rate and depth mechanisms. Gesell, Steffensen and Brookhart (73) concluded that, "The pulmonary vagi exercise two diametrically opposite functions: curtailment of the central respiratory discharge which reduces the depth of breathing, and acceleration of the inspiratory act, which allows increased depth of breathing." Tang (171) found that CO<sub>2</sub> administration failed to induce a rate change if the vagi were severed. Only changes in depth were observed. In the absence of pneumotaxic circuits, vagal afferents maintained a relatively constant depth during respiratory stimulation. Von Euler, Herrero and Wexler (55) also reported that the increase in respiratory rate during CO<sub>2</sub> stimulation depended on intact vagal mechanisms. The conclusions of Nesland, Plum, Nelson and Siedler (129) were identical.

Shannon, Zechman and Frazier (162) studied the first-breath response of inspiratory cells during mechanical loading in cats. In preparations with intact vagi, elevation of the resistance or elastance led to an increased depth and decreased rate of respiration, which were paralleled by increased frequency and increased

duration of single cell discharges. Vagotomy abolished the single cell changes during loading, but depth increases were still present. The authors concluded that the vagi were the only source of sensory information for activation of medullary inspiratory neurons during mechanical loading. Increase in depth in vagotomized preparations was believed to be due to modulation at the spinal cord level involving facilitation of external intercostal motor activity by their muscle spindles during loading. This conclusion was confirmed by Shannon and Zechman (161).

Finally, Fallert and Mühlemann (57) studied entrainment of respiratory centers during pump inflation with different tidal volumes and frequencies in the intact rabbit. Elevation of inflation volume increased the frequency range to which the rabbit would entrain. Small frequency ranges correlated with small volume inflations. The authors concluded that the Hering-Breuer reflex was important for determination of respiratory rates and depths.

#### D. Discharge Patterns

The ability to monitor single cell activity from the central nervous system has been of great value in the study of respiratory control. Various testing

procedures have been introduced whereby primary respiratory neurons can be distinguished from other cell types. Qualitative and quantitative assessment of respiratory cell discharge patterns have yielded satisfactory neural correlates of respiratory rate and tidal volume outflows from medullary structures. Consequently, numerous respiratory cell subtypes have been described and new theories on respiratory rhythmicity have been introduced.

Single respiratory potentials can be recorded from the medulla by using glass or metal microelectrodes (12, 20, 62, 78, 89). Gesell, Bricker and Magee (70) first reported that finding extracellular respiratory discharges was slow and tedious work. Von Baumgarten (16) found only 23 cells with 900 penetrations using 25 microelectrodes. Salmoiraghi and von Baumgarten (154) worked for a year and were successful in penetrating only eight respiratory cells for intracellular recording. Once located, extracellular respiratory potentials can be recorded for extended periods of time. Brodie (31) reported that he could usually hold cells for 30-60 minutes. Occasionally, cells were observed for more than seven hours.

Positive identification of primary respiratory neurons is based on phasing and spiking characteristics of the discharges and modification of patterns during

various testing procedures. Although both tonic and phasic cells may function in respiratory systems, the phasic cells have been classified as primary respiratory cell candidates since they fire during one phase of respiration and are quiescent during the alternate phase. Classically, respiratory neurons are grouped into inspiratory or expiratory categories. Tonic cells can show a frequency peaking in phase with the respiratory cycle, but these cells never become quiescent.

Care must be taken to show that phasic discharges do not arise from movement artifacts. Various techniques have been used to eliminate mechanical movement. Von Baumgarten, Kanzow, Koepchen and Timm (20) covered the brainstem with a thick layer of stiff agar jelly before recording respiratory potentials. Batsel (15) reported that cisternal drainage resulting from compression of the abdomen and chest was equal to or superior to the use of agar to arrest movements. Batsel (14) also found that cells responding to fluctuations in the blood pressure, which has a slow oscillating component in phase with respiration, could not be recorded from for a long time. Artifacts from this source were eliminated because of their instability. Cohen (39, 40, 41) used cats with neuromuscular blockade to avoid movement potentials. Movements

due to pump inflations were minimized by using preparations with a pneumothorax. Finally, von Baumgarten (17) introduced a floating electrode technique which permitted recording of respiratory potentials from moving brainstems.

Probably the best way to screen out rhythmic discharges due to movement is by examination of the spike potentials. Dirken and Woldring (51) and Hukuhara, Nakayama and Okada (91) used constant spike amplitude as their criterion. Any modulation of amplitude in phase with the respiratory cycle must be due to cell movement in relation to electrode tip position in the potential fields around active cells. Also, differences in spike amplitude can be used as a check against recording from more than one cell at the same time.

The phasing of periodic discharges is important in respiratory cell identification. Amoroso, Bainbridge, Bell, Lawn and Rosenberg (5) accepted as primary respiratory neurons only those discharges which preceded muscular electrical activity and which fired throughout the inspiratory or expiratory phase. Special care must be taken in identification when rhythmic potentials are found in the region of the solitary tract or nucleus since they may arise from vagal afferents or vagal motor-neurons. The former are easily identified by their phase

spanning patterns as shown by Adrian (2), but the latter are more difficult to distinguish. Eyzaguirre and Taylor (56) showed that vagal motoneurons had discharge patterns paralleling phrenic bursts. Some had expiratory phasing patterns. In addition, von Baumgarten and Kanzow (19) described two respiratory cell types in the region of the tractus solitarius, neither of which were vagal afferents or vagal motoneurons. Both had inspiratory patterns, but lung inflation led to an inhibition of one and an excitation of the other.

When recording respiratory discharges it is important that the potentials arise from cell bodies. It is possible that ascending afferent and descending efferent traffic may carry respiratory potentials along fiber tracts which traverse medullary regions. These, of course, cannot be considered as primary respiratory potentials. Various techniques have been used to differentiate between cell body and fiber tract recordings. Due to differences in diameters, Cohen and Wang (45) suggested that cell bodies have a higher current output than single fibers. For this reason, potentials from the former have larger amplitudes (47) and longer durations (30, 79) than the latter and are found more frequently. Since the recording volume of steel microelectrodes with tip diameters

of 10-40 microns is only  $5 \times 10^{-3}$  cubic millimeters (5), fiber potentials are more susceptible to electrode movement than cell body potentials. For example, von Baumgarten and Salmoiraghi (21) concluded that they were recording from a cell body if the microelectrode could be moved 30-100 microns without losing the potential. Haber, Kohn, Ngai, Holaday and Wang (79) could not record respiratory activity from the caudal medulla and suggested that their microelectrodes were sensitive only to cell body potentials.

Nelson (127) and Cohen and Wang (45) introduced a new criterion and claimed that sources of unit activity could be distinguished on the basis of spike polarity. Pure negative or positive-negative waves were believed to arise from cell bodies. These were the most common. Pure positive or positive-negative-positive triphasic spikes were believed to arise from axons or fiber tracts.

Cells with respiratory characteristics may not always function in the respiratory complex and are therefore probably not primary respiratory neurons. Scheibel and Scheibel (158) observed reticular cells for many hours and reported that some non-respiratory cells assumed activity characteristics resembling respiratory patterns. The authors concluded that such neurons may function in different

neural nets at different periods of time. Gesell, Bricker and Magee (70) found that respiratory discharges may differ considerably in frequency and may vanish and reappear under modifying conditions. This indicated that the reticular formation could command recruitment and adjustment of respiratory cell patterns. Cohen (38) described periodic discharges recorded in the isolated pons which resembled respiratory discharges in both burst duration and firing frequency. On close examination, however, these cells were not locked to the respiratory cycle, but tended to drift in random patterns. This demonstrated that cells in the reticular formation could exhibit similar respiratory patterns without having any known respiratory function.

Elementary descriptions of respiratory cell discharge patterns have most commonly included measurements of the number of action potentials per burst (spikes per train), the discharge duration (train length) and the average discharge frequency. It has been implied that increases in these parameters for inspiratory cells correlate with deeper respiratory movements. Pitts (141) stimulated the inspiratory region of the medulla with various intensity and frequency combinations and found increased firing frequency of single motor units and motor unit



recruitment from the subliminal fringe of the phrenic population. Adrian and Bronk (3) found a linear relationship between interpleural pressure and stimulation frequency of the phrenic from 20-50 Hz. Integrated total phrenic nerve activity has subsequently been shown to be proportional to the tidal volume (39, 50, 58). Eldridge (53) recently demonstrated that integration of the phrenic signals during the 100 milliseconds coincident with the peak of inspiration is the best neural analog of tidal volume over wide ranges of tidal volume and respiratory rate.

Nesland, Plum, Nelson and Siedler (129), studying inspiratory cell patterns, introduced the product of spikes per train times respiration rate and showed that this spike output increased proportionally with elevation in minute ventilation. For example, CO<sub>2</sub> administration in intact cats led to increases in depth and rate of respiration, increase in mean frequency discharge, and decrease in train length. Spikes per train remained constant. Parameter changes due to alteration of respiratory rate were prevented by vagotomy. In vagotomized cats, CO<sub>2</sub> administration induced pure depth changes with corresponding increases in mean frequency discharge and spikes per train. Train length did not change. It was also shown that hypoxia produced less consistent responses than CO<sub>2</sub>

administration. Hyperoxia depressed all parameters and some cells were completely inhibited. Nesland et al.

(129) concluded, "It appears that the medullary respiratory neuron population regulates the magnitude of ventilation primarily by changes in the impulse frequency and total number of discharges of already active cells rather than by increasing or decreasing the number of cells actively discharging during each breath."

Nelson (127) recognized the importance of sampling from many respiratory cells to "resynthesize" characteristics of the whole population. Discharge parameters from inspiratory and expiratory cells were examined at different respiratory rates and the following conclusions were made. The train length is directly proportional to the respiratory cycle time. The spikes per train are proportional to the train length. The spike frequency is proportional to the spikes per train. Dirken and Woldring (51) confirmed the train length-spikes per train length relationship for expiratory cells at a constant discharge frequency. Cohen (41) confirmed the spikes per train-inspiratory phase duration relationship for inspiratory cells in lung inflation studies. Brodie (31) studied discharge patterns in inspiratory cells during modification of the spontaneous respiratory rate with drugs.

Thiopental, a respiratory depressant to both rate and amplitude, increased the spikes per train and burst duration but slightly decreased the discharge frequency. Cyanide, a respiratory stimulant, reversed all the responses. Brodie (31) concluded that respiration rate and train length were reciprocally related while respiration rate and discharge frequency were directly related.

Wang and Ngai (175) have stated that respiratory rate is probably determined by the rate of depolarization of respiratory neurons. In this case, measurements of spikes per train, train length and mean discharge frequencies fail to account for dynamic changes that take place in burst discharges during depolarization of respiratory cells. To quantitate these dynamic events, frequency modulation curves have been constructed where the instantaneous spike frequency is plotted as a function of time. Salmoiraghi and Burns (155) showed that respiratory cells fired in non-regular patterns, necessitating curve fitting for single burst discharges.

Gesell, Magee and Bricker (72) reported that inspiratory cells exhibited a slowly augmenting pattern in which the spike frequency consistently increased as the train progressed. The frequency discharge rapidly decreased just prior to the off-phase for the cell. This

frequency pattern correlated with tidal movement of air. Two expiratory patterns were described. One was a steady state discharge in which expiratory cells fired with a constant frequency throughout the whole train. The other pattern showed a rapid increase in frequency which slowly declined before the cells turned off. Similar discharge patterns were found at all points along the respiratory arc from the respiratory muscles through the central nervous system and back along sensory nerves. These parallel patterns suggested serial linking between stations of the respiratory mechanism. Gesell, Atkinson and Brown (69) restudied respiratory discharge patterns in muscle-blocked dogs. No alteration in inspiratory patterns were seen, suggesting that the frequency modulation of these cells was of central origin. Expiratory cell patterns were all of the steady state type. The authors suggested that the rapidly augmenting expiratory patterns might have been due to reflex modification of the basic steady state pattern. Gesell and Worzniak (74) supported this suggestion by showing that steady state expiratory cell patterns could be converted into rapidly augmenting patterns during spontaneous respiration or into slowly augmenting patterns during pump inflation.

Dirken and Woldring (51) studied the frequency

curves of inspiratory cells, expiratory cells and vagal afferents. Decreasing lung volume by tracheal blockade in mid-inspiration caused inhibition of vagal afferent frequency discharge and drop in the respiratory rate. The time course of frequency modulation in inspiratory cells was unaltered except for the duration of inspiration, which was increased. Vagal blockade by anelectrotonus, however, depressed the initial rate of change in frequency in the same cells. Assuming that rate of frequency change depends on the rate of depolarization in respiratory cells, a depression in the latter can be associated with low respiratory rates after vagotomy.

Hukuhara, Nakayama and Okada (91) described frequency modulation patterns of inspiratory and expiratory cells in cats and dogs. The basic patterns observed agreed with those of Gesell et al. (72) and up to four variations were noted for both cell types. Variations ranged from slowly and rapidly augmenting discharges to steady state patterns. The authors attributed variations in starting time, spike frequency and number of spikes per train to differences in thresholds throughout both inspiratory and expiratory populations.

Von Baumgarten, Balthasar and Koepchen (18) and Salmoiraghi and von Baumgarten (154) studied intracellular

discharge patterns from inspiratory and expiratory neurons. Burst discharges were initiated by a spontaneous depolarization of the resting membrane potential which continued to depolarize slowly until the burst was terminated. Inspiratory cells showed a frequency increase paralleling the degree of membrane depolarization. The authors suggested that the high depolarization level at the end of each spike train reflected a maximum threshold, above which the cells could not generate additional action potentials. This self-limiting theory quickly replaced the suggestion of Burns (35) that spike discharges terminated because of fatigue. Batsel (15) argued against the latter view by pointing out that respiratory cells could be driven at higher than normal rates for long periods of time without any sign of fatigue. Also, Robson, Houseley and Solis-Quiroga (148) showed that respiratory cells could fire continuously after barbiturate administration.

Salmoiraghi (153) proposed a theory for rhythmicity of breathing based on three excitatory mechanisms: (1) increased firing frequency of inspiratory cells, as the cycle progressed, depended on self-reexciting mechanisms within the inspiratory population; (2) concurrent decreased probability of expiratory cell firing was attributed

to reciprocal inhibition; (3) increased firing thresholds of inspiratory neurons explained transition from inspiration to expiration. This cyclic exchange between inspiratory and expiratory populations was expected to continue as long as the three postulated mechanisms were intact.

Although Waldron (174) has reported that distinct categories of activity patterns could not be distinguished within the inspiratory group, other investigators have described many respiratory cell subtypes. Cells have been classified into no less than eight categories based on phasing pattern differences with the respiratory cycle (14, 15, 40, 41, 42, 45, 128, 169, 173). It is interesting that cells do not change their phasing during CO<sub>2</sub> administration, lung inflation or other procedures. Most workers, led by Cohen (43), have assumed that these different respiratory cell patterns reflected differences in function. Elaborate theories on the genesis of respiratory rhythm have subsequently been devised. Batsel (15) concluded that the bulbar respiratory center may exhibit rhythmicity due to the differences in onset time of discharges of the various early inspiratory cells. Temporal overlapping of different subgroups was believed to smooth the transition between respiratory phases.

Nesland and Plum (128) suggested that any hypothesis

on respiratory oscillation derived from the interrelationships among functionally different subgroups within the medullary inspiratory and expiratory populations required three assumptions. First, one active subgroup must activate another quiescent subgroup. Second, one active subgroup must inhibit another active subgroup. Third, activity must be found within at least one subgroup at any point in time. Cohen (42) used these criteria and proposed a model for genesis of respiratory rhythmicity based on respiratory cell subtype response to CO<sub>2</sub>, lung inflation and brainstem electrical stimulation. The schema was constructed around a master oscillatory loop which paralleled the inspiratory and expiratory neurons of Salmoiraghi and Burns (36, 155, 156). Three other loops had direct influence on the master loop. One, consisting of expiratory-inspiratory neurons, initiated inspiration. The other two loops inhibited inspiration and expiration respectively. Inspiratory units activated by lung inflation might function in the inspiratory inhibitory loop, but oscillation did not depend on vagal innervation.



## CHAPTER III

### METHODS

#### A. Surgical Preparation

Mongrel cats anesthetized intraperitoneally with 30 mg/Kg sodium pentobarbital (Nembutal, Abbott Laboratories) were used exclusively in this study. The cats had a mean weight of 2.7 Kg (range 1.6 to 4.9 Kg) and there was no sex discrimination. Cats were placed on a Gorman-Rupp Industries, Inc., Model M11 warm water circulated pad which was driven by a Model K-1-3 water heater-pump. Rectal temperatures, which were monitored with a Yellow Springs Instrument Company, Inc., Model 46 TUC Tele-thermometer, were maintained at  $37 \pm 2^{\circ}\text{C}$ . Surgical preparations and instrument calibrations were usually completed within 2 hours. Most cats remained remarkably stable for more than 8 hours and breathed spontaneously throughout the duration of the experiment.

Four surgical incisions were made. First, the femoral triangle was exposed to permit catheterization of the

femoral artery and vein with PE100 polyethelene tubing. Catheter tips were advanced to the abdominal aorta for blood pressure recording and inferior vena cava for drug injections, respectively.

Second, a four or five centimeter midline incision was made through the ventral cervical neck skin and sternomastoid muscle. Reflection of the sternohyoid muscle exposed the trachea for intubation with a quarter-inch polypropylene (Nalgene) Y-tube. The cervical vagosympathetic nerves were isolated and loosely ligatured with umbilical tape. In some experiments loose triple zero silk ligatures were placed around each cervical carotid artery for bilateral carotid occlusions or tugs.

Third, after mounting the cat in David Kopf 1530 Stereotaxic Frame Assembly, a small incision was made through the skin of the left lateral thoracic cage and the pectoralis minor muscle was separated by blunt dissection. A transthoracic needle was inserted through the serratus ventralis and internal intercostal muscles at the fourth or fifth rib interspace. This probe was used for intrapleural pressure recording.

The last surgical procedure involved exposure of medullary brainstem structures. A dorsal midline incision was made through the skin of the head extending three to

four centimeters rostral and caudal to the occipital crest. The lambdoidal ridge was exposed by separation of the cut skin interfaces with retractors. A National cautery was used to separate neck muscles (clavotrapezius, levator auris longus, epicranium occipitalis, auricularis superior, abductor auris longus, abductor auris brevis and temporalis muscles). Some muscles were removed and other were retracted until the caudal interparietal and occipital bone surfaces were laid bare. The exposed dura mater between the occipital bone and the atlas vertebra was ruptured resulting in a free flow of cerebral spinal fluid. Portions of the occipital bone were removed with Rongeur forceps carefully inserted above but never touching the dorsal surface of the medulla. Medullary brain-stem structures from the first cervical nerve to the cerebellar border were exposed. The posterior lobes of the cerebellum were also uncovered for suction removal of the whole cerebellum if necessary. In most experiments, however, the entire cerebellum was left intact. Frequent application of 0.9% saline solution prevented the brain tissue from drying out. In some cases, the spontaneous flow of cerebrospinal fluid provided satisfactory irrigation of the fourth ventricle.

## B. Data Acquisition

The interconnections of the electronic equipment used to monitor and store on-line analog data are presented schematically in Figure 1. A Grass Model 7 Polygraph strip chart recorder was used to record four physiological parameters. In the first channel, airflow velocity was measured with a Fleisch pneumotachograph connected in series with one arm of the tracheal Y-cannula. The opposite arm was blocked with U-screw clamp occlusion of a rubber tube extension. During movement of air through the pneumotachograph the small pressure gradient produced across a metal honeycomb grid was detected by a Grass Model PT5A differential volumetric pressure transducer. The device was calibrated and found to be linear with airflow velocities up to 110 cc/second in accordance with Poiseuille's Law. This was the upper limit for airflows in most experimental cases.

Intrapleural pressure was monitored on the second polygraph channel. A 15 gauge needle beveled at the tip with sideports along its shaft was inserted across the lateral thoracic wall at the fourth or fifth rib interspace. An elliptical copper plate attached to the needle shaft at a 45° angle limited the probe's insertion distance to 2.5 centimeters. Skin flaps were pulled

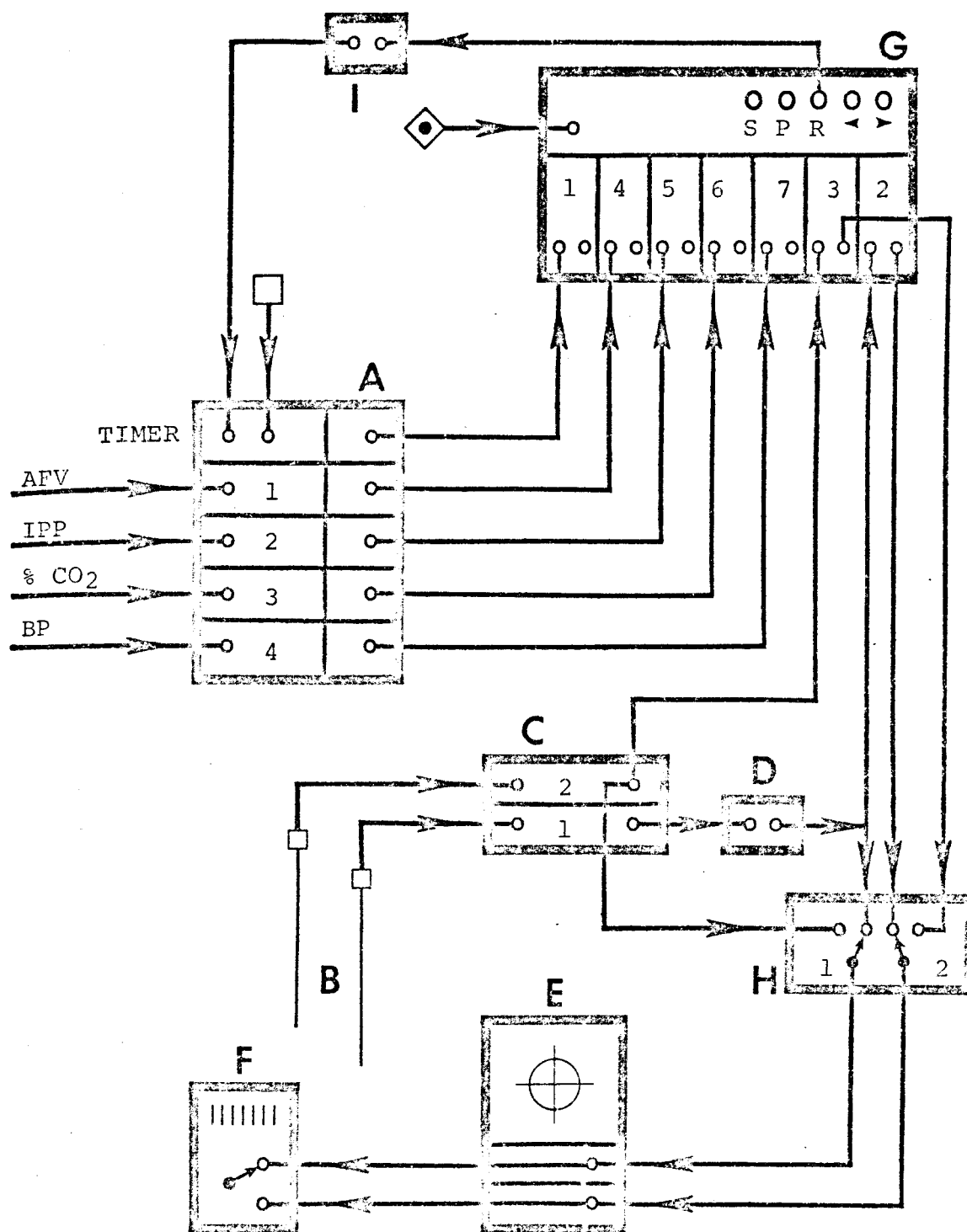


Figure 1. Instrumental schematic: A. polygraph; B. microelectrodes; C. preamplifier; D. 60 Hz filter; E. oscilloscope; F. audioamplifier; G. tape recorder; H. switching network; I. current amplifier.

together over this plate to seal any air leaks and maintain the needle in an approximate parallel apposition with the lung. Air pressure fluctuations were recorded with a Statham P23Dc pressure transducer which was initially calibrated with a water manometer.

The third polygraph channel monitored the end expiratory % CO<sub>2</sub>. A short length of tygon tubing was attached to the exit port of the pneumotachograph to support a PE260 CO<sub>2</sub> sampling catheter in the center of the airflow profile. A Beckman microcatheter sample pump continuously drew a small fraction of the expired air through the detector head of a Beckman LB-1 Medical Gas Analyzer. Using an infrared absorption technique, a voltage signal related to the % CO<sub>2</sub> of the sample was obtained. This curvilinear signal was processed with a Beckman linearizer before going to the polygraph. At the beginning of each experiment the CO<sub>2</sub> channel was calibrated with a CO<sub>2</sub> reference gas (Matheson Gas Products). The linearizer was periodically adjusted for linearity.

Finally, the blood pressure was recorded on the fourth polygraph channel. A Statham PE23Gb pressure transducer attached to an intra-aortic PE100 catheter filled with 0.9% non-heparinized saline was used for this measurement. The catheter was kept clot-free by frequent

flushings with saline. The blood pressure transducer was initially calibrated with a mercury manometer.

The last physiologic parameter examined in this study was the extracellular electrical activity of single respiratory neurons in the medulla. As shown in Figure 1, potentials were recorded with microelectrodes, AC amplified (2 channel capacity) and filtered. In early experiments, glass micropipettes filled with 3 molar sodium chloride solution were used. With the exception of one cell, however, all single cell potentials analyzed in this study were recorded with Green's (78) metal microelectrodes. Steel insect pins were etched to fine tip diameters in 12 molar hydrochloric acid using a 6.3 volt A.C. source. The electrodes were insulated with clear Insl-X E33 solution and baked in an oven overnight. Tip dimensions were measured with a Carl Zeiss microscope (X10) and Vickers A.E.I. image splitting eyepiece (X10) and ranged from 1-5 microns. Corresponding electrical impedances, checked on a General Radio Company Type 1650-A impedance bridge, ranged from 80-5 megohms respectively.

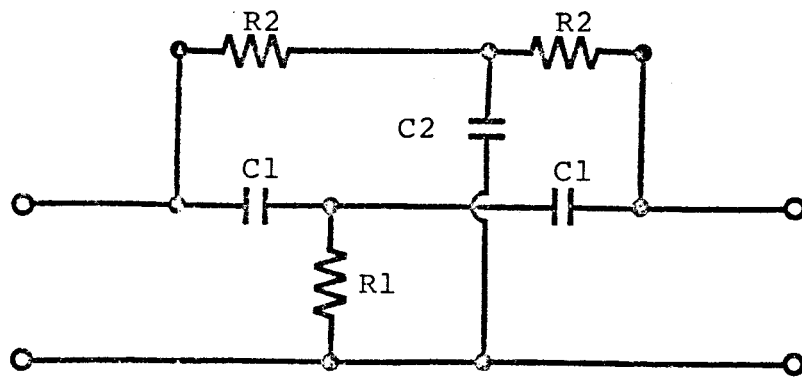
Microelectrodes were mounted on a Kopf 1460 electrode carrier fitted with a Kopf 1206 reduction drive and were slowly advanced through medullary brain tissue in 10 micron steps. No attempt was made to cut away the pia

mater. Single cell potentials picked up by the microelectrodes were in the 200-800 microvolt range. These potentials were amplified about 2000 times with a Grass DP9B dual channel AC preamplifier (single-ended input, cathode follower mode). The signal-to-noise ratio was maximized by keeping input leads short, use of shielding, establishment of common grounds between the amplifier and cat, and signal filtering. The low and high half amplitude frequency filters on the amplifier were set at 0.1 and 40 K Hz respectively. A 60 Hz notch filter, shown in Figure 2A, was constructed to eliminate line noise. The frequency response curve of this filter, which is plotted on a log scale in Figure 2B, shows maximum attenuation at 60 Hz. Experimental tests demonstrated that peak-to-peak voltages of single respiratory spike potentials were not attenuated by filtering. Filtered waveforms were more biphasic than nonfiltered signals and the former displayed a small leading phase shift (less than 0.2 millisecond). Neither of these changes affected the validity of data collection or spike train analysis used in this study.

Amplified and filtered single cell discharges were monitored on-line. Potentials were displayed on a Tetrnix type 502A dual-beam oscilloscope and were audio-amplified on a Grass AM7 audio-monitor as shown in Figure 1. Both



A



$$R1 = 3.92 \text{ K}$$

$$R2 = 7.50 \text{ K}$$

$$C1 = 0.350 \text{ } \mu\text{fd}$$

$$C2 = 0.689 \text{ } \mu\text{fd}$$

$$R2 = 2R1$$

$$C2 = 2C1$$

$$f = [2\pi (R2) (C1)]^{-1}$$

B

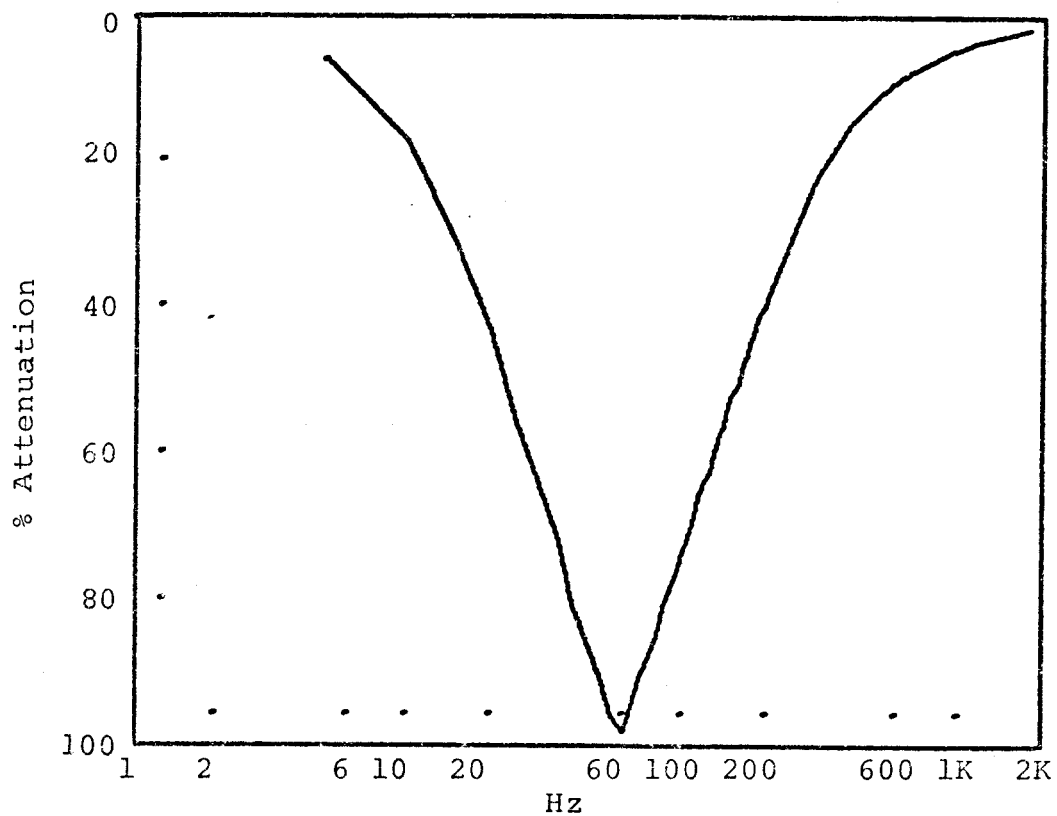


Figure 2. Parallel T notch filter: A. schematic diagram; B. frequency response curve.

of these instruments played important roles in the positive identification of single respiratory cell discharges.

Whenever a single respiratory cell potential of good amplitude was found, on line data from both this cell and from the polygraph were stored on a seven channel Philips ANA-LOG 7 tape recorder as shown in Figure 1. The tape recorder was operated at 3-3/4 inches per second in the FM mode. During a recording procedure, single cell data were immediately played back to the oscilloscope and audio-monitor to determine the quality of the recorded signal. A switching network permitted various combinations of data review when recordings from two different cells were obtained at the same time.

Finally, a current amplifier was constructed which sensed current flow in the "record light" logic circuit of the tape recorder. Whenever the recorder was storing analog data, this device closed a relay which caused a downward identification mark to be placed on the polygraph strip chart. An upward deflection on the polygraph event marker channel was activated by a remote foot switch to identify experimental manipulations. Other features including voice description on the tape (channel 8) and digital tape counter were also useful in locating single cell discharges during data analysis.

### C. Experimental Manipulations

After a cat was surgically prepared and all polygraph channels were calibrated, a microelectrode search for respiratory potentials was started. Strict stereotaxic coordinates were not used, but by following the map of Salmoiraghi and Burns (155) with the obex as a reference point, respiratory cells could be found. Care was taken in the identification of cells as primary respiratory neurons. Many types of potential discharges were found in the medulla, but only those showing an on-off firing pattern in phase with the respiratory cycle (airflow, intrapleural pressure and % CO<sub>2</sub> oscillations) were considered as primary respiratory cell candidates. Some of these potentials were rejected for various reasons. Potentials whose amplitude fluctuated as a function of the respiratory cycle were eliminated because of the possibility that they were generated by movement artifacts. Potentials with uniform spike height showing highly irregular or high frequency discharges (instantaneous frequencies over 200 Hz) were rejected since more than one cell was probably present. Finally, potentials showing phase-spanning patterns were rejected since they may have been vagal afferent in origin. Only those potentials of constant amplitude exhibiting pure inspiratory or pure expiratory phasings were recorded

as primary respiratory neurons. These potentials were believed to be derived from cell bodies since no respiratory discharges could be found from the lower medulla to the upper cord where respiratory efferents are known to pass.

When a respiratory cell was found and positively identified, a control record was stored on magnetic tape. Various manipulations of the respiratory system were then performed to induce alterations in the discharge pattern. In general, procedures were used which evoked changes in respiratory rate and depth. Single cell data were obtained in both vagi-intact and vagotomized preparations.

Increases in respiratory rate were induced by intravenous injection of doxapram hydrochloride (Dopram, A.H. Robins Company). Respiratory rate was readily depressed by sodium pentobarbital or sodium pentothal (Pentothal, Abbott Laboratories). In a few cases, high respiratory rates arose spontaneously during exposure of the medulla. This acceleration was probably due to warming of brainstem structures. Other procedures, including CO<sub>2</sub> administration, anterior hypothalamic heating and intravenous injections of morphine sulfate (U.S.P.) were tried in several cats, but associated rate changes, if any, were not consistent.

Respiration depth was increased by increasing airflow

resistance. For this procedure, the cross-sectional area of the pneumotachograph exit port was decreased by known fractions with calibrated resistance plugs. It was very difficult to retain single unit recordings during induced depth changes. Either the cell would be lost because of movement problems or other cells would be recruited, thus masking the original discharge. For this reason, data obtained with airflow resistance plugs will not be presented.

Finally, other studies were designed to examine the correlation between apneustic breaths and barbiturate level in the spontaneously breathing cat. This phenomenon was studied at the single cell level and apneustic thresholds were evaluated by cerebellectomy and bilateral carotid occlusion. Data obtained from this work was important for the interpretation of single cell discharge pattern changes during barbiturate titration.

CHAPTER IV  
DATA ANALYSIS

A. PDP-12 Overview

The general objective of this study was to investigate interspike interval distribution and sequencing of single respiratory cell discharges at various steady state breathing frequencies and depths. A Digital Equipment Corporation PDP-12 (Programmed Data Processor-12) general purpose laboratory computer was programmed to process off-line analog tape data of single cell activity. For each steady state observation, thousands of interspike intervals were examined and manipulated to construct histogram and interspike interval modulation curves. The latter involved a new mathematical technique devised by the author to quantitate ordering of interspike intervals during burst discharges. In addition to powerful software capabilities, a full array of PDP-12 output options was available to the user: CRT display; LA-30 Decwriter (Digital Equipment Corporation) hard copy printing; DP-1 Complot

(Houston Instrument, Baush and Lomb) digital increment plotting; LINC magnetic tape storage.

Communication with the PDP-12 computer was accomplished with essentially three programming languages. The first two are machine language in format and actually represent the two mode operation capacity of the PDP-12. The first or LINC-8 (Laboratory INstrument Computer) mode activates peripherals including analog to digital conversion, sense switch lines, relay buffers, CRT display and auxiliary LINC tape storage. The second or PDP-8 mode provides for extended arithmetic element (EAE) operations, real-time clock programming, digital incremental plotting and Decwriter input-output. While each mode has its own complete instruction set, both modes are given equal status by the single central processor. This permits rapid switching between modes under program control. Also, the 1.6 micro-second memory access time of the PDP-12 permits fast execution of individual instructions. This speed is advantageous for rapid analog to digital conversions.

FOCAL (Formulating On-line Calculations in Algebraic Language) was the third programming language used. This is a conversational language that permits on-line user-machine interaction and resembles FORTRAN IV (FORMula TRANslation version IV) in power for calculation of complex

mathematical expressions. Two FOCAL systems, FOCAL-12 and FOCALPL (plot), were utilized respectively to retrieve and manipulate data stored on magnetic tape from machine language programs and to graphically plot calculated parameters located on digital tape.

Two major disadvantages are associated with FOCAL. First, FOCAL programs are limited in length since the FOCAL interpreter, a machine language program itself, takes up a large portion of the PDP-12 8K core memory. Second, execution of individual FOCAL program instructions is slow since each mnemonic code command must be interpreted into machine language binary each time it is performed. Under the conditions of this study, these disadvantages were by no means limiting.

#### B. Machine Language Programs

##### 1. ERASE

For convenience and speed of data handling, data were stored on LINC magnetic tape of standard format. Each tape was marked off into  $1000_8$  ( $512_{10}$ ) blocks, each block containing  $400_8$  ( $256_{10}$ ) 12-bit words. Prior to data storage, program ERASE was run to store a zero in each of the  $131,072_{10}$  ( $512_{10} \times 256_{10}$ ) 12-bit words. By this procedure, new and old tapes could be cleared of all digital values before proceeding with other programs. Program ERASE along



with the programs discussed below are listed in the Appendix for reference.

## 2. DATA

Program DATA was written in machine language making free use of both LINC and PDP-8 modes. As outlined in the Appendix (Table VII), program instructions were located in the lower 4K memory of the PDP-12 and data were stored in the upper 4K memory. An abbreviated logic flow diagram of the program instructions is presented in Figure 3. The logic is represented by three major loops (A,B,C) and calculation (CALC) procedures between loops B and C. To facilitate understanding of data handling by program DATA, each of these four program sections will be discussed in relation to a real physiological input signal. The reader should be able to trace data processing of the analog respiratory cell discharge input to the final digitalized output before proceeding to higher levels of data analysis.

Inspiratory and expiratory cell discharges were examined in this study, and two representative cells are shown in Figure 4A and 4B respectively (35 mm records from a Grass Kymograph Model C4N Camera). Both cells were obtained from different cats, each of which was breathing at a rate of 30 breaths per minute. It can be seen that the inspiratory cell is in phase with the fall in the intrapleural pressure

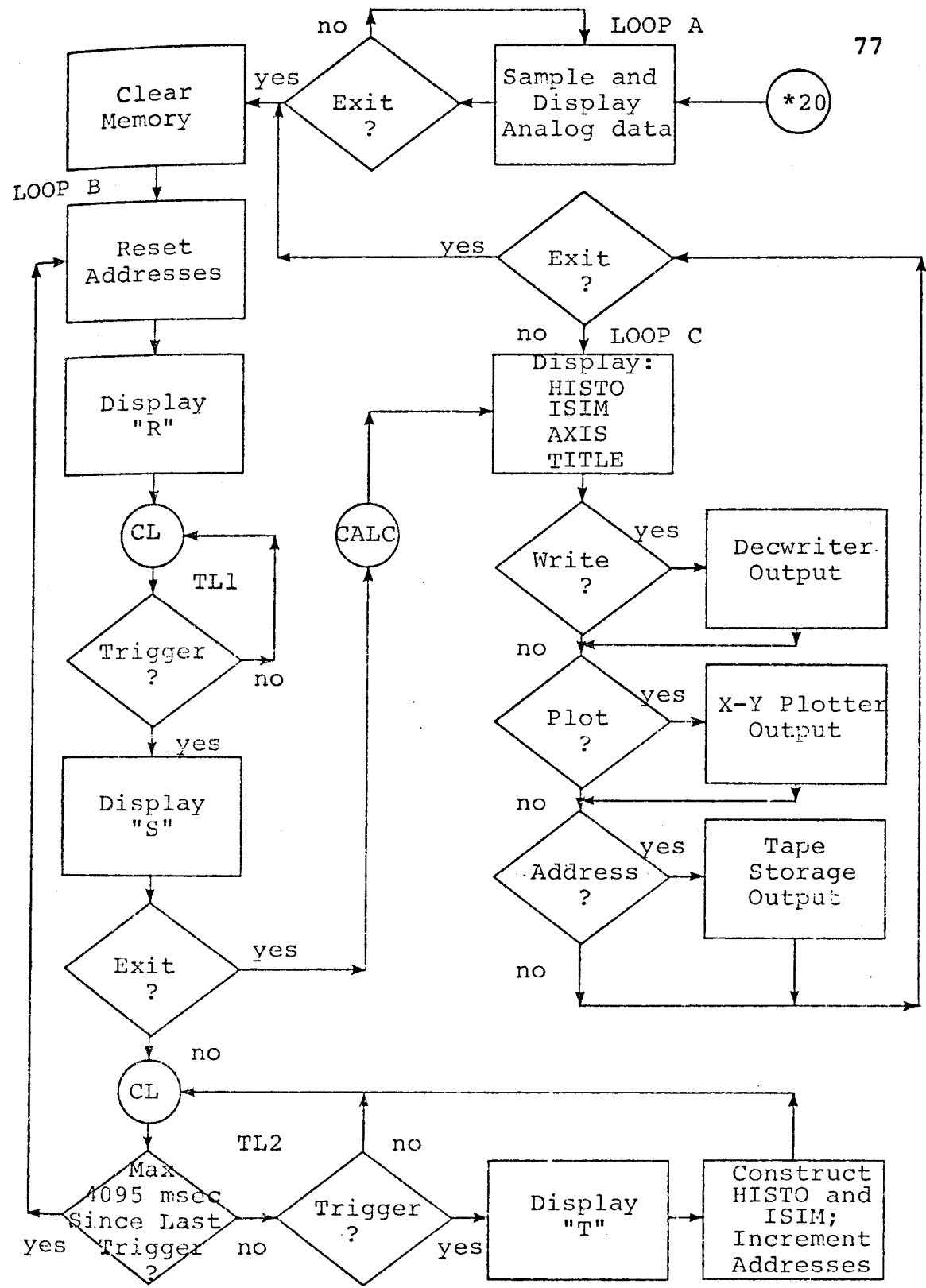


Figure 3. Program DATA flow chart. See text for details.

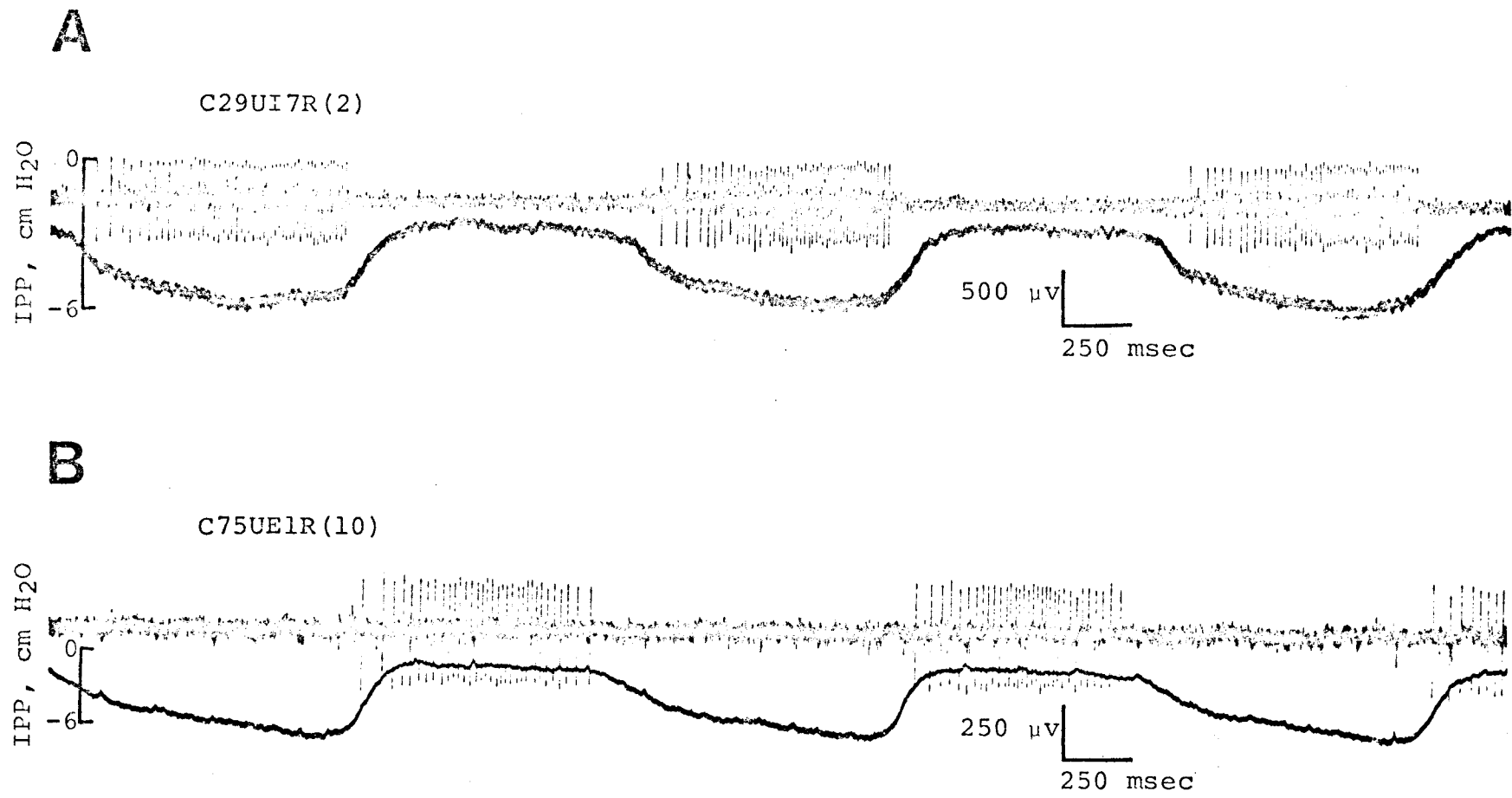


Figure 4. Representative examples of single cell analog data: A. inspiratory cell; B. expiratory cell. Lower tracing in each panel is intrapleural pressure (IPP).

while the expiratory cell is  $180^\circ$  out of phase. The inspiratory signal C29UI7R(2) of Figure 4A will be processed step by step in the discussion that follows.

The analog respiratory discharge was played back to the PDP-12 computer four times slower than record speed. In loop A of program DATA (Figure 3) the signal was digitalized and displayed on the CRT for assessment of signal quality. The sample rate was determined by the clock logic which halted program flow until a predetermined amount of time had elapsed. The KW12 real time clock, a crystal-controlled pulse generator oscillating at 400K Hz, was programmed to give a basic clock overflow count rate of 0.4 millisecond. This rate, coupled with the fourfold slow-down of the tape transport, established a simulated sampling rate of 0.1 millisecond per sample or 10 samples per millisecond.

Loop A was programmed to sample and store  $1000_8$  points before displaying the stored data on the CRT. The display therefore corresponded to a 51.2 millisecond segment ( $1000_8$  samples  $\times$  ( $512_{10}/1000_8$ )  $\times$  0.1 msec/sample) of the respiratory cell discharge. The memory was constantly updated causing the CRT to flash at about 5 Hz ( $[0.0512 \text{ sec} \times 4]^{-1}$ ) although any segment could be frozen on the CRT for close analysis via sense switch control. After a signal

was judged to be satisfactory for computer analysis, exit from loop A was manually controlled under sense switch selection. Between loops A and B, the upper 4K of core memory was cleared along with various constants. Certain addresses were also reset to appropriate values.

The overall responsibility of loop B was to construct histogram and interspike interval modulation curves from the analog signal. The logic centered around detection of action potentials, measurement of interspike interval times and proper memory storage of interval times in various combinations for histogram and interval sequence analysis.

The PDP-12 computer is equipped with a Schmitt trigger. This device senses potentials whose amplitude exceed a manually set threshold and hence would appear ideal for action potential detection. The Schmitt trigger, however, has several disadvantages when used to detect the action potentials recorded in this study. For this reason a software rate-of-rise trigger was designed which proved highly reliable in action potential identification. The rate-of-rise trigger diagrammed in Figure 5 operated as follows. Analog sampling in loop B occurred at a constant simulated rate of 0.1 millisecond per sample as determined by the clock logic (CL). At this rate, action potentials

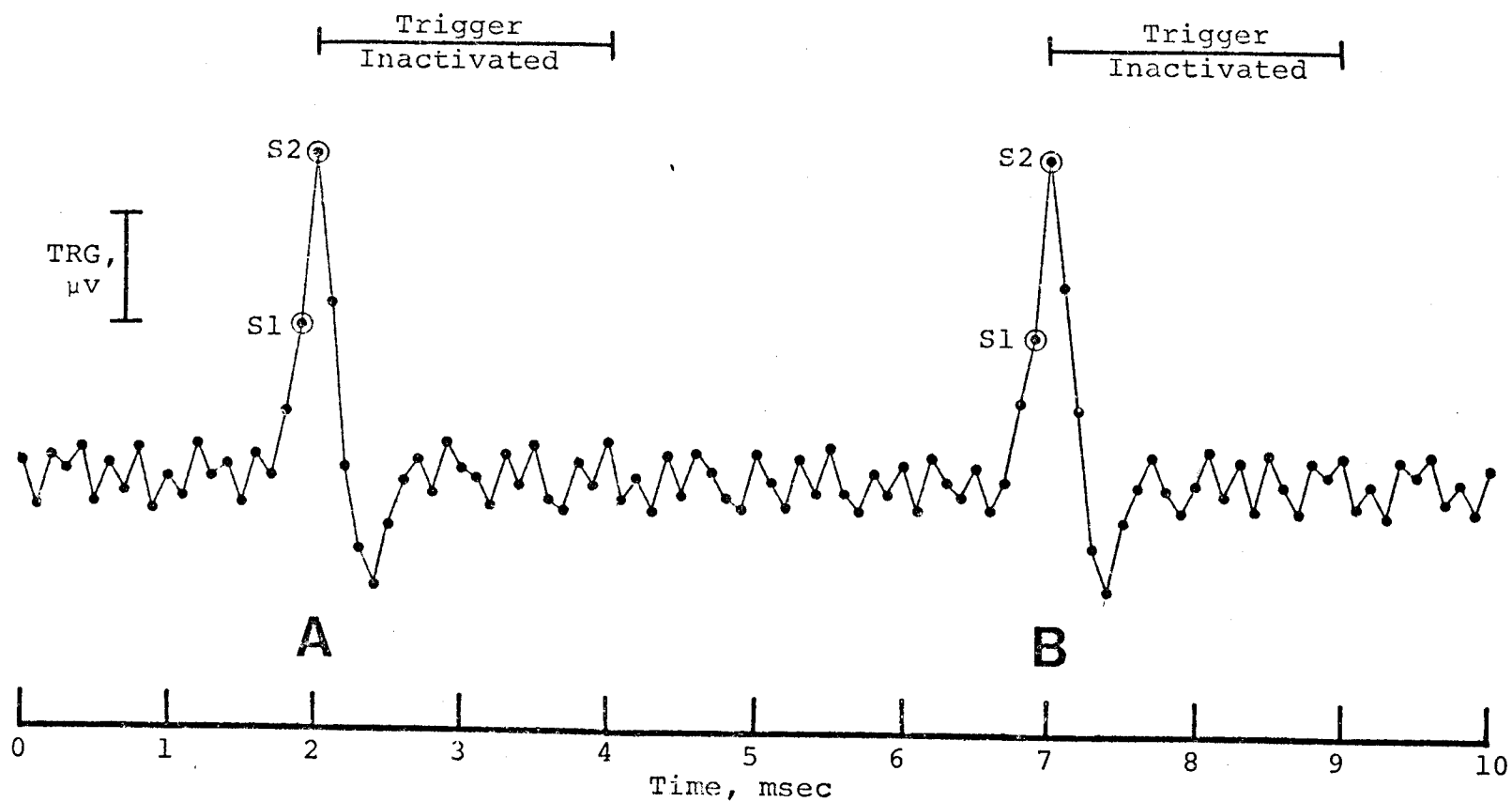


Figure 5. Technique for detection of action potentials. Dots represent digitalized 0.1 msec samples of analog signal. The change in voltage ( $\Delta S = S_2 - S_1$ ) per unit time represents a signal exceeding the trigger (TRG) level ( $\Delta S > TRG$ ). This identifies an action potential (A or B) and inactivates the trigger logic for the next 2 msec.

1 millisecond in duration could be sampled 10 times. Since the sample rate was constant, the vertical distance between samples was proportional to the rate of potential change. Therefore, by testing for vertical separation between successive samples, action potentials could easily be detected even when the signal to noise ratio was as low as 2:1 and when slow baseline shifts were present. The trigger level (TRG) was manually adjusted until a "T" flashed on the CRT in synchrony with the audio amplified action potentials. The trigger operated only on the rising phase of action potentials. After a trigger occurred, 2 milliseconds elapsed before the trigger was "activated" again. This eliminated the possibility of generating two trigger pulses from a single biphasic action potential.

Interspike interval times were calculated to the nearest 0.1 millisecond by counting the elapsed time between action potential triggers. In short, a trigger caused the accumulated time of the previous interval to be stored with subsequent resetting of the interval counter. In Figure 5, for example, action potential A resets the counter to 0.0 milliseconds. Detection of action potential B 5.0 milliseconds later terminates the accumulation of interval time. The A-B interspike interval is stored as 5.0 milliseconds and the counter is reset for the next interval.

Histogram analysis has been a popular approach to quantitation of electrophysiological data from single neurons (66, 67). In this study, a 100 millisecond interval histogram was constructed by accumulating 500-4000 intervals, the number under manual selection. The histogram had 125 bins each of 0.8 millisecond width. The histogram plot for inspiratory cell C29UI7R(2) is presented in Figure 6A in which the terminal 30 milliseconds have been deleted. In Figure 6B the same data are plotted on an expanded scale. As consistent with all units studied, the distribution of interspike intervals for this cell shows unimodal peaking and is skewed to the right, favoring longer intervals. Parrot and Fleming (139) attempted to associate this characteristic shape of the respiratory interspike interval histogram with random Poisson processes. Although this has been done for non-respiratory discharges of truly random patterns (68, 106, 165), it is difficult to assign such random mechanisms to respiratory cells which possess predictable discharges. Against the random approach, Marczynski and Sherry (119) suggested that time-locked patterns which repeat over and over again with a sequential arrangement of adjacent intervals can best be described by the autocorrelation function. High correlations for respiratory cells (R.A. Mitchell - unpublished



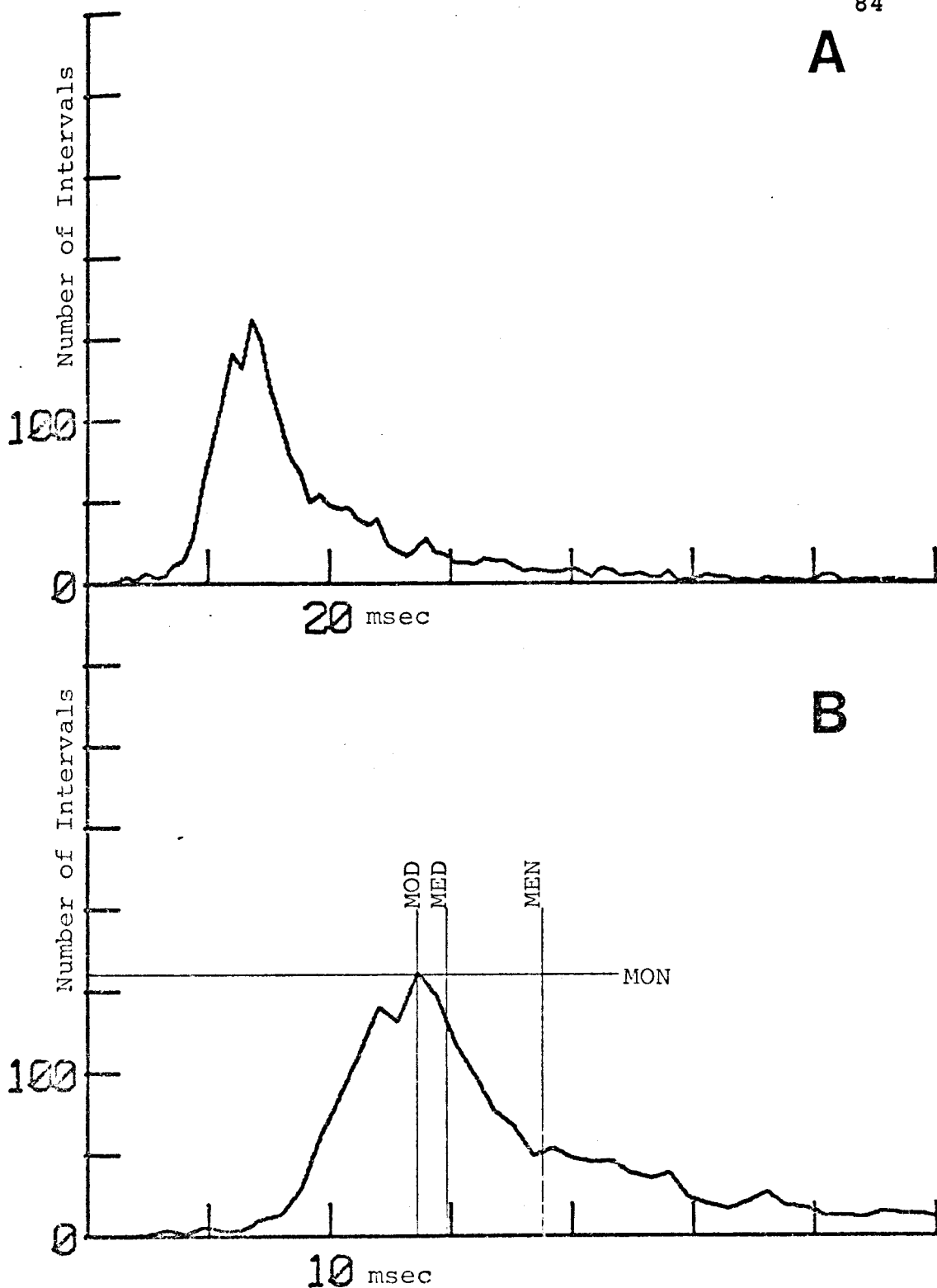


Figure 6. A. Interspike interval histogram derived from inspiratory cell C29UI7R(2) in Figure 4A. B. Same histogram on an expanded time scale showing mean (MEN), median (MED), mode (MOD) and number of intervals at the mode (MON). The histogram is made up of 2000 intervals.

observations) are certainly not indicative of random ordering of intervals. The interdependence series matrix method introduced by Sherry, Marczynski and Wolf (163) may also be useful in establishing a relationship between a specific interval and an interval removed by a fixed number of other intervals for respiratory train analysis.

A cumulative histogram or ogive curve is plotted in Figure 7 for inspiratory cell C29UI7R(2). The plot is linear on a log scale indicative of exponential processes operating in interval distribution. Similar plots are found for both Poisson and normal distribution functions. A horizontal line has been drawn at the 50% level and a vertical line intercepting the curve at this same level has been extended to the time scale. By definition, the interpolated time of 14.8 milliseconds corresponds to the median interval time. From Figure 6B, it can be seen that the median time is longer than the mode time. Also, the mean interval time is longer than median time. These relationships establish the skewing characteristics of the interspike interval histogram to the right.

For discharges that have random arrangement of intervals, quantification of cell output is satisfactorily completed by histogram analysis and measurement of standard

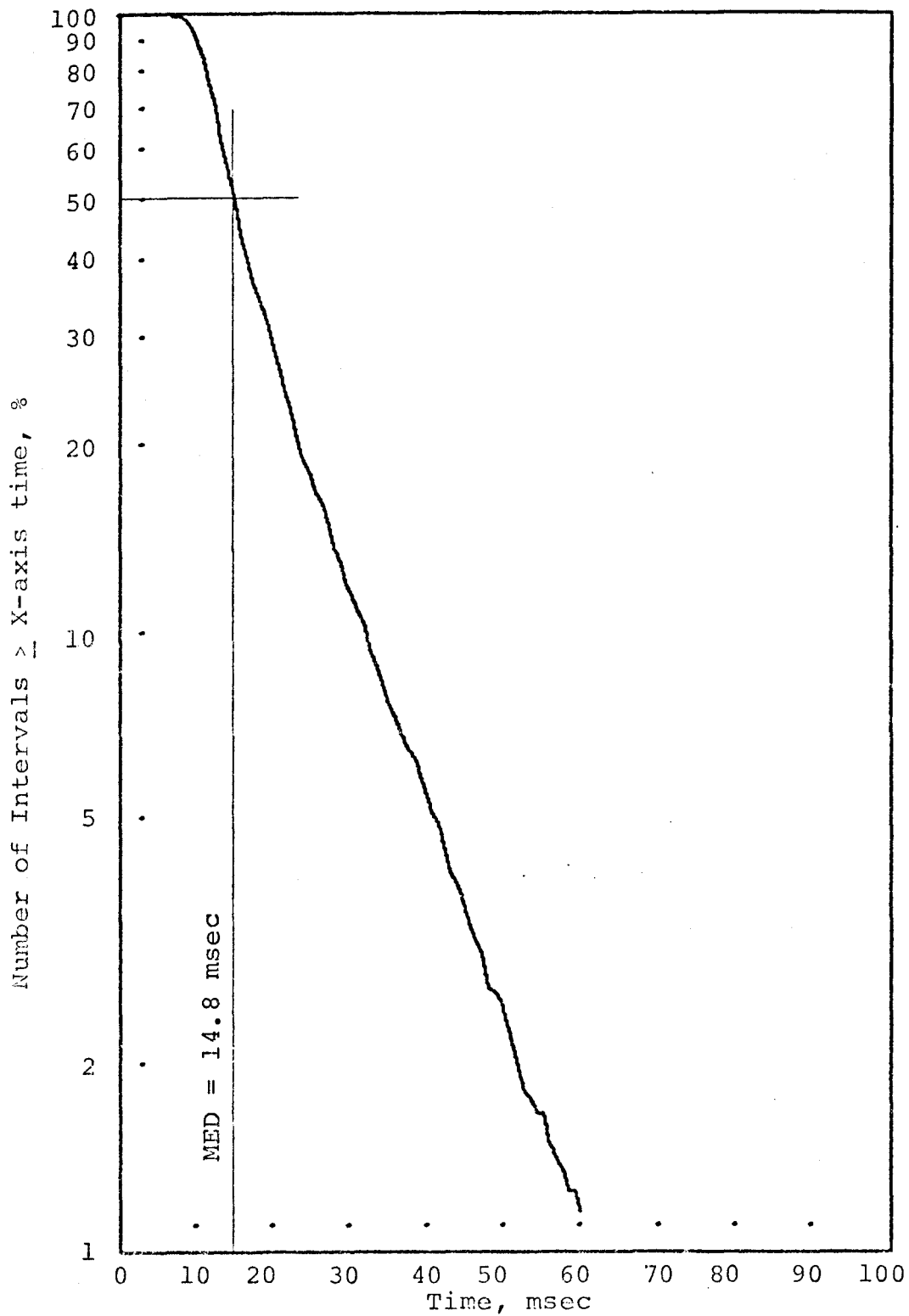


Figure 7. Ogive curve derived from histogram in Figure 6. Median (MED) interval time interpolated from 50% level.

statistical parameters such as the mean, median, mode and standard deviation (119). As reviewed in the Literature, however, respiratory discharges undergo characteristic frequency increase, plateauing and frequency decrease during their "on" phase. For this reason, assessment of respiratory cell output must involve description of sequential interval ordering, information that cannot be extracted from histogram interval distributions. Cohen (40) and Bertrand and Hugelin (24) approached this problem by using phase-triggered time histograms and cycle-triggered spike-density histograms for analysis of respiratory discharges, respectively. Although qualitative assessment was achieved, these methods failed to quantitate dynamic changes in the interspike interval during respiratory cell discharge.

To study dynamic changes in the interspike interval during respiratory burst activity, a new statistical averaging technique was devised as diagrammed in Figure 8 with hypothetical data. Intervals were numbered consecutively (Figure 8A) and corresponding intervals were averaged to construct an idealized respiratory train (Figure 8B). Plotting the sequential interspike intervals of the idealized train as a function of time resulted in a smooth "U"-shaped interspike interval modulation curve (Figure 8C).

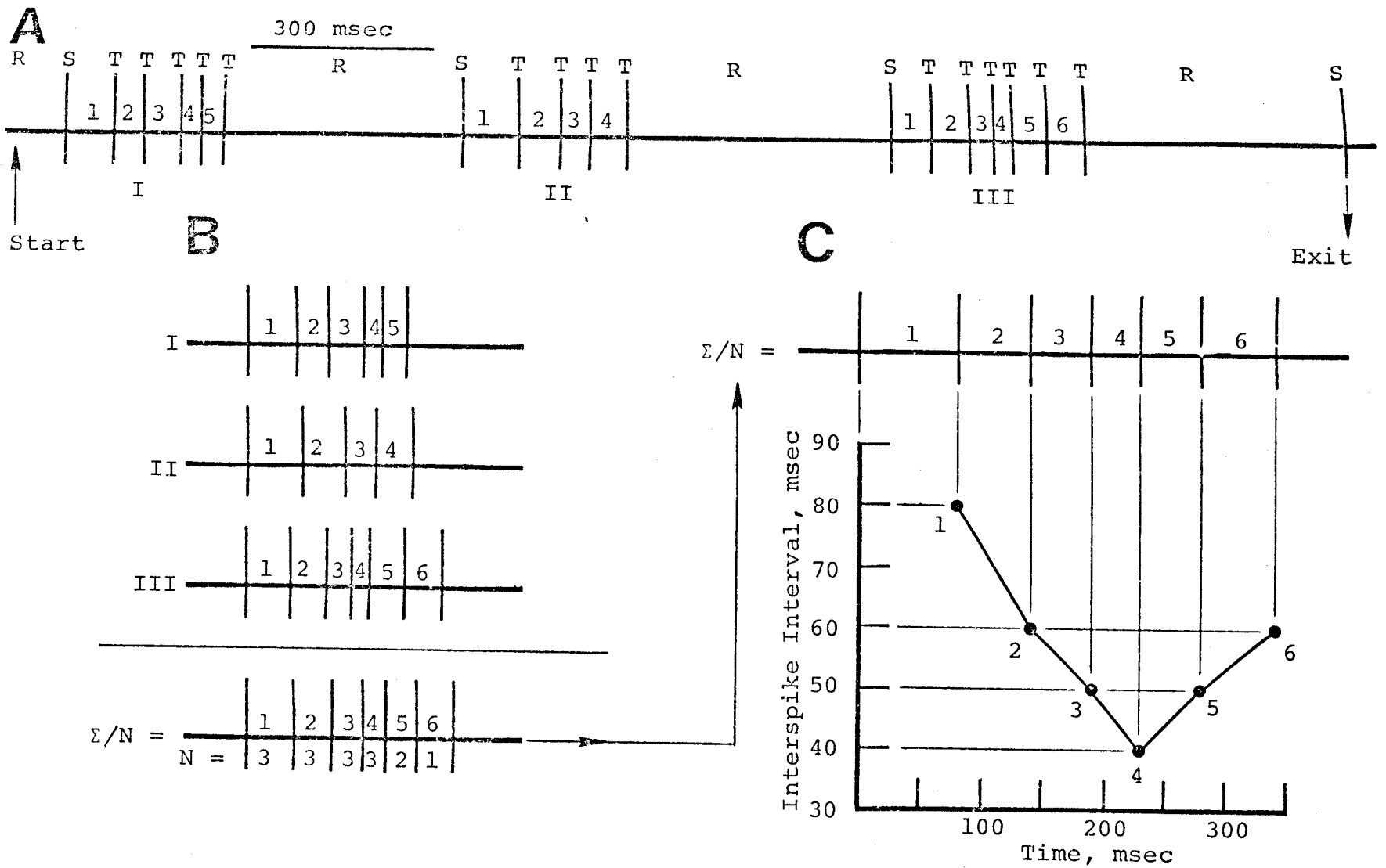


Figure 8. Generation of idealized interspike interval modulation curve from raw data: A. analog data; B. interval averaging; C. graphic display of averaged intervals. See text for details.

In Figure 9, similar plots for cell C29UI7R(2) are presented. For a single burst discharge (NT=1) a very rough curve was described reflecting non-smooth transition between consecutive interspike intervals as the train progressed (Figure 9A). An average of 44 sequential burst discharges (NT=44) eliminated the variability between adjacent intervals and resulted in a smooth interspike interval modulation curve (Figure 9B), readily accessible for mathematical analysis. This mathematical method differed from standard signal averaging techniques (164) which depend on external time references to recover repetitive waveforms from random noise. The analysis worked for both inspiratory and expiratory discharges (Figure 4) and was independent of cycle cues from peripheral respiratory parameters.

The computer techniques employed to generate interspike interval modulation curves can best be understood by paralleling the "S" (start of train), "T" (trigger on successive action potentials) and "R" (reset) designations over the action potentials in Figure 8A with the display "S", display "T" and display "R" commands of loop B in Figure 3. For example, loop B logic was manually entered from either loop A or loop C during the quiescent period of the respiratory discharge (interburst interval). This

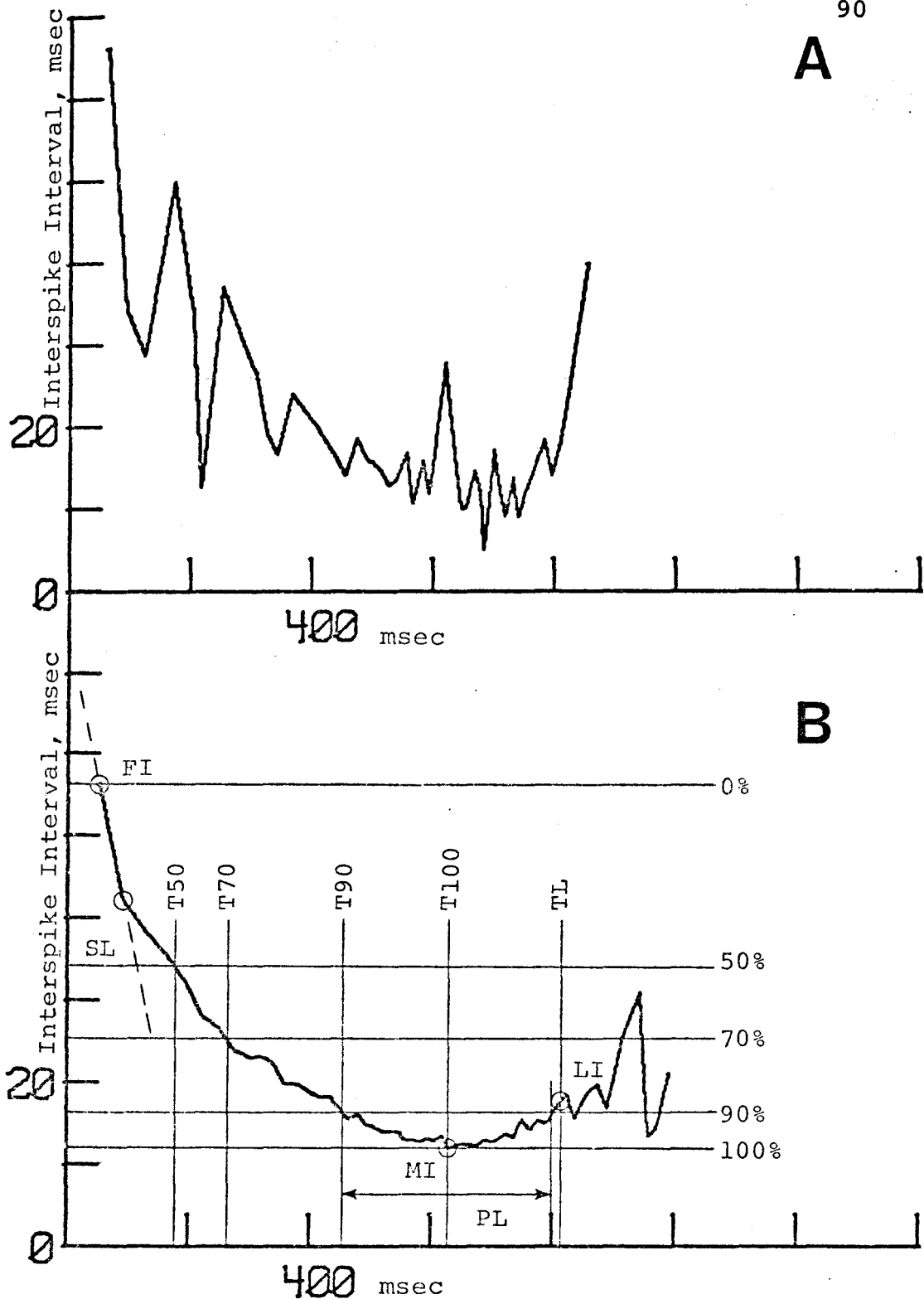


Figure 9. Interspike interval modulation curves derived from inspiratory cell C29UI7R(2) in Figure 4A: A. single train; B. average of 44 trains. Derivation of parameters shown in B are described in text, pages 99-106.

caused an "R" to be flashed on the CRT indicating that appropriate addresses had been reset. Program flow was then delayed in the first trigger loop (TL1) until the first action potential was detected. At this time, an "S" was flashed on the CRT and program flow was then delayed in the second trigger loop (TL2). Starting with the second action potential, all the remaining action potentials in the train caused a "T" to be displayed on the CRT. In conjunction with the clock logic (CL), each of these successive action potentials defined an interspike interval which was stored in an incremental address. Thus, consecutive intervals were stored in consecutive addresses. When a predetermined amount of time (maximum of 4095 msec scaled time) elapsed without any action potentials being detected, it was assumed that the train discharge was terminated for that breath and reset procedures were automatically initiated as revealed by CRT display of the "R". The sequence started all over again when the first action potential (display "S") of the next burst discharge was detected.

By this iterative technique, the PDP-12 memory accumulated the sum of the first intervals, the sum of the second intervals and so forth. To handle large sums, double precision arithmetic was used. That is, two 12-bit registers were daisy-chained together to form a 24-bit register



(maximum value accepted =  $2^{23} = 8,388,608$ ). The idealized train was recovered after exit from loop B by dividing each accumulated interval sum by the number of intervals that contributed to that sum. Since the divisor N was variable, the analysis was not limited by the shortest train, a problem described by other workers (24, 40). However, commencing with the first interval after the shortest train and continuing until the last interval of the longest train, the divisor N decreased as fewer and fewer trains contributed to the idealized interspike interval average (Figure 8B). For this reason, a breakdown in the statistical averaging and hence poor curve smoothing was observed towards the end of the train (Figure 9B).

In addition to histogram and interspike interval modulation curve genesis, loop B also stored consecutive cycle times and first interval times during the complete analysis. Cycle time (CT) was defined as the time between first action potentials of consecutive burst discharges. The first interval (FI) was defined as the time between the first two action potentials of each train. This parameter is circled in Figure 9B at the beginning of the train.

Exit from loop B was under both manual and automatic control. During trial runs, output procedures could be initiated by sense switch selection. Also, if more than 510

burst discharges were accumulated, automatic transfer to loop C was performed. For final runs, however, exit from loop B occurred after the histogram N had been filled. If the histogram was filled during the middle of a burst discharge, intervals were still calculated to complete the interspike interval modulation curve for that burst, but the intervals were deleted from histogram construction. Exit proceeded with the first action potential of the very next train (Figure 8A). This permitted accurate measurement of the last cycle time.

Between loops B and C, a series of calculation (CALC) steps were performed (Figure 3). As already discussed, mean interspike intervals were recovered to generate the idealized train. Individual instantaneous respiratory rates were calculated from the reciprocal of each cycle time measurement and these rates were averaged to give a mean respiratory rate (RR) for the analysis period. Mean rates calculated in this manner were slightly higher than averages taken by dividing the number of respiratory cycles observed by the total analysis period. This was due to the hyperbolic relationship between the period and the reciprocal of the period. High and low rates were isolated to reveal the degree of steady-stateness of instantaneous frequencies.

The number of intervals in each train were calculated and the mean spikes per train (ST) were found by adding one to the mean number of intervals per train. By examination of each idealized interval, the minimum interval (MI) in the respiratory train was localized by a search technique. The minimum interval (MI) is circled in Figure 9B. Also, the mean (MEN) interval time was calculated from the idealized train taking into account the variable divisor  $N$ , for each interval. The mode (MOD) and median (MED) interval times were extracted from the histogram data and the number of intervals in the bin at the mode (MON) was found. The mean (MEN), median (MED), mode (MOD) and number of intervals at the mode (MON) are diagrammed in the histogram data of cell C29UI7R(2) in Figure 6B.

Finally, the following eleven values were stored in a specific portion of core memory to facilitate transfer to LINC magnetic tape: mean respiration rate (RR), number of trains averaged (NT), spikes per train (ST), first interval time (FI), minimum interval time (MI), number of intervals in the interspike interval modulation curve (FMN), number of intervals in the histogram (HSN), mean (MEN), median (MED) and mode (MOD) interval times, and number of intervals at the mode (MON).

After completion of the calculation procedures,

program flow proceeded to loop C instructions. Loop C (Figure 3) was an output loop responsible for retrieving calculated data. The CRT was used to continuously display the histogram or interspike interval modulation curve (sense switch selection) along with axes and appropriate scaling integers. Titles could be placed on the CRT from the Decwriter keyboard for photographic purposes.

Selection of sense switch W (write) caused the Decwriter to print a hard copy list of digitized values stored in core memory. Table I is the output from cell C29UI7R(2). In A, the mean, high and low respiration rates (breaths/10 minutes) and instantaneous rates for each breath are given. The number of breaths averaged ( $N=NT$ ) was 44. In B, the mean spikes per train and the number of intervals for each individual train are given. Concerning the latter, one train was 37 intervals long, six trains were 43 intervals long, five trains were 48 intervals long, etc. The N is repeated as an internal program check. In C, the individual first interval times for each breath are printed to nearest 0.1 millisecond and in D, the digitized values for the idealized interspike interval curve are listed to nearest 0.1 millisecond in sequential order (e.g. first interval, second interval, etc.). An average of data in C equals the first value in D. Also in D, the mean (MEN),



minimum interval (MI) and number of intervals contributing to interspike interval modulation curve ( $N=FMN$ ) are retrieved. Finally, in E, the contents of each of the 125 - 0.8 millisecond bins of the histogram are printed. The mode (MOD), number of intervals at the mode (MON), median (MED) and number of intervals in the histogram ( $N=HSN$ ) are also given. It can be seen that the 17th bin contains the largest number of counts ( $MON=161$ ). This by definition is the mode (e.g. 17 bins x 0.8 msec per bin = 13.6 msec or to place the peak at center of 0.8 msec bin,  $MOD = 13.6 \text{ msec} - 0.4 \text{ msec} = 13.2 \text{ msec}$ ). The reader should be able to correlate the digitalized values of Table 1-D with Figure 9B and Table 1-E with Figure 6.

Selection of sense switch P (plot) caused the Calcomp X-Y digital plotter to plot the histogram and interspike interval modulation data on paper from Table 1-E and 1-D data, respectively. These plots along with axes and scalings were identical to the CRT displays of the same data. Figures 6 and 9 are computer plots for cell C29UI7R(2) obtained from this output procedure.

Finally, selection of sense switch A (address) transferred data in core memory to a specified address of the LINC magnetic tape. Included in this transfer were the individual number of intervals per train and individual first

intervals, histogram and interspike interval modulation data, and the eleven constants mentioned previously (RR, NT, ST, FI, MI, FMN, HSN, MEN, MED, MOD, MON). A CRT display of "F" indicated that the tape contents at the selected address were full. This prohibited overwriting of stored data by mistake, although contents could be replaced if so desired. The tape-stored data were accessed via FOCAL programs for advanced mathematical analysis. In addition, other machine language programs could be run to replot histogram and interspike interval modulation data from the magnetic tape.

Exit from loop C (sense switch control) cleared the core memory and initiated loop B procedures (Figure 3). To examine the single cell input signal it was necessary to manually halt program flow and restart the program at \*20.

### 3. REPLT

Program REPLT permitted replotting of either 100 or 50 millisecond histogram and interspike interval modulation data from the LINC magnetic tape. Data from several steady state observations on the same cell or different cells could be superimposed for visual comparison. Since this program was a modification of the plot routine in program DATA, it is not listed in the Appendix.

C. FOCAL-12 Programs

1. \$CALC1

Data stored on the LINC magnetic tape were accessed, manipulated and restored in columnar arrangement by this first calculation program. The following new eleven parameters were derived and added to the list of eleven parameters previously calculated by program DATA. The data from cell C29UI7R(2) found in Table I are used for all examples, which can be substantiated via hand calculations. This program along with all other FOCAL-12 programs are listed in the Appendix.

The train length (TL) was determined to the nearest millisecond by summing consecutive intervals of the idealized interspike interval modulation curve data until the mean number of intervals per train was reached. Mathematically,

$$n = \text{integer}(ST) - 1$$

$$f = ST - 1 - n$$

$$TL(\text{msec}) = \left[ \sum_{i=1}^n \text{int}_i \right] + f[\text{int}_{n+1}]$$

where ST is the mean spikes per train;  $\text{int}_n$  is the last whole interval of the intervals contributing to the train length (TL) and "f" corrects for the fractional contribution from the very next interval,  $\text{int}_{n+1}$ , for trains terminating



between intervals  $int_n$  and  $int_{n+1}$ . For cell C29UI7R(2),  $n = 45$ , "f" = 0.7 and  $TL = 845.9 + 0.7(18.5) = 859$  milliseconds. A vertical train length (TL) line has been drawn on the interspike interval modulation curve in Figure 9B corresponding to the above measurement for cell C29UI7R(2). It can be observed that the mean train length (TL) was reached before the curve lost the smooth transitions between intervals toward the end of the train. No measurements or interpolations were made past the mean train length (TL). This cutoff was observed for most cells.

The mean cycle time (CT) was calculated to the nearest 10 milliseconds by computing the reciprocal of the mean respiratory rate (RR). That is,

$$CT(\text{msec}) = 10 \cdot \text{integer}[(RR^{-1} \cdot 60000 \text{ msec/min})/10]$$

For cell C29UI7R(2),  $CT = 10 \cdot \text{integer}(6000/30.1) = 1990$  milliseconds.

From program DATA, the number of intervals per individual train and individual first interval times contributing to single steady state averages were stored on LINC magnetic tape. These values were retrieved by program \$CALC1 for calculation of the standard deviation (SD) and standard error (SE) for each. These standard statistical measurements were made as follows:

$$SD = \sqrt{[\Sigma X^2 - (\Sigma X)^2/N]/[N - 1]}$$

$$SE = SD/\sqrt{N}$$

where X represents the individual intervals for either case and N is the number of respiratory bursts averaged. For cell C29UI7R(2), N = 44 and  $SD_{ST} = 3.9$ ,  $SE_{ST} = 0.58$ ,  $SD_{FI} = 11.1$  and  $SE_{FI} = 1.68$  milliseconds.

The plateau level (PL), defined as the length of time the interspike interval modulation curve was within 10% of the minimum interval (FI - MI difference = 100%), was measured to the nearest millisecond. That is,

$$n = \text{integer}(ST) - 1$$

$$K = 0.10(FI - MI) + MI$$

$$PL(\text{msec}) = \left[ \sum_{i=1}^n \text{int}_i < K \right]$$

where ST is the mean spikes per train, FI is the first interval, MI is the minimum interval and  $\text{int}_n$  is the last whole interval of the intervals contributing to the mean train length. The plateau level (PL) is drawn in Figure 9B as a horizontal line extending between two vertical lines which intercept the experimental interspike interval modulation curve at the 90% level. Intervals meeting the requirements, but falling after the mean train length (TL), were disregarded. The plateau level (PL) for cell

C29UI7R(2) was 368 milliseconds.

The last interval (LI) was defined as the interspike interval time of the experimental curve at the end of the measured mean train length. This parameter was calculated to the nearest 0.1 millisecond and is circled in Figure 9B at the end of the train. Mathematically,

$$n = \text{integer}(\text{ST}) - 1$$

$$f = \text{ST} - 1 - n$$

$$\text{LI}(0.1 \text{ msec}) = f[\text{int}_{n+1} - \text{int}_n] + \text{int}_n$$

where ST is the mean spikes per train,  $\text{int}_n$  is the last whole interval of the intervals contributing to the mean train length (TL) and  $\text{int}_{n+1}$  is the very next interval. The "f" measurement computes a linearly interpolated last interval (LI) correction for train lengths (TL) terminating between intervals  $\text{int}_n$  and  $\text{int}_{n+1}$ . For cell C29UI7R(2),  $\text{LI}(0.1 \text{ msec}) = 0.7(18.5 - 16.7) + 16.7 = 18.0$  milliseconds with the train terminating between the 45th and 46th intervals.

The slope (SL) was a measure of the initial rate of change in the interval time of the interspike interval modulation curve. The slope was defined as

$$\text{SL} = (\text{SI} - \text{FI})/\text{SI}$$

where FI is the first interval and SI is the second interval. The slope of the dashed lined connecting the first

two circled intervals in Figure 9B corresponds to this measurement. For cell C29UI7R(2),  $SL = (41.9 - 56.1)/41.9 = -0.34$ .

The plateau level percent (PLP) was defined as the ratio of the plateau level (PL) to the mean train length (TL). For example,

$$PLP(\%) = (PL/TL)100$$

This parameter was a normalized measure of the fraction of the cell's "on" time occupied by intervals within 10% of the minimum interval (MI). For cell C29UI7R(2),  $PLP = (368/859)100 = 42.8\%$ .

The last measurement made by program \$CALC1 was the train length percent (TLP). This parameter was defined as the ratio of the mean train length (TL) to the cycle time (CT) or

$$TLP(\%) = (TL/CT)100$$

This measurement normalized the fraction of the respiratory cycle occupied by respiratory cell activity (e.g. percent "on" time). For cell C29UI7R(2),  $TLP = (859/1990)100 = 43.2\%$ .

## 2. \$CALC2

Similar to program \$CALC1, this second calculation program derived the final twelve parameters from the LINC magnetic tape. These parameters were added to the columnar

data previously calculated, thus completing the number of parameters measured for each steady state observation to thirty-four. The following new parameters were calculated. All examples are taken from cell C29UI7R(2) data listed in Table I permitting verification via hand calculations.

As illustrated by the horizontal lines traversing the interspike interval modulation curve in Figure 9B, the interspike interval times were normalized into percent levels. The range extended from the first interval (FI = 0%) to the minimum interval (MI = 100%) with additional horizontals drawn at the 50, 70 and 90% levels. Where these imaginary lines crossed the experimental curve, vertical lines corresponding to times to reach 50, 70, 90 and 100% levels (T50, T70, T90, T100) were constructed. Measurements of T values were carried out to the nearest millisecond. Linear interpolations were performed in cases where 50, 70, and 90% horizontal levels crossed the curve between two intervals. Associated with these four measurements (T50, T70, T90, T100) were four parameters (N50, N70, N90, N100) representing the number of intervals elapsed before each level was reached. The mathematical expressions used to calculate the T and N values are given below:

$$K = (1 - P/100)(FI - MI) + MI$$

$$T(\text{msec}) = \left[ \sum_{i=1}^n \text{int}_i \right] + \left[ \frac{K - \text{int}_n}{(\text{int}_{n+1} - \text{int}_n)/\text{int}_{n+1}} \right]$$

$$N = n - (\text{int}_n - K)/(\text{int}_{n+1} - \text{int}_n)$$

where P is the percent level and n is the sequence position of the interval preceeding the first interval less than K. For cell C29UI7R(2),

$$K_{50} = (1 - 50/100)(56.1 - 12.0) + 12.0 = 34.05$$

$$T_{50}(\text{msec}) = (56.1 + 41.9 + 38.1 + 35.4) + \frac{(34.05 - 35.4)}{(31.9 - 35.4)/31.9} = 184 \text{ msec}$$

$$N_{50} = 4 - (35.4 - 34.05)/(31.9 - 35.4) =$$

4.4 intervals

Also for cell C29UI7R(2), T70 = 270, T90 = 467 and T100 = 655 milliseconds, and N70 = 7.5, N90 = 17.4 and N100 = 31.0 intervals. From Table 1-D, it can be seen that the minimum interval (MI = 12.0 msec) is the thirty-first interval in the train and corresponds to N100.

Four last parameters were derived by taking the ratio of the T values to the mean train length. That is,

$$T5P(\%) = (T50/TL)100$$

$$T7P(\%) = (T70/TL)100$$

$$T9P(\%) = (T90/TL)100$$

$$T10P(\%) = (T100/TL)100$$

For example, for cell C29UI7R(2),  $T5P = (184/859)100 = 21.4\%$ . Also,  $T7P = 31.4\%$ ,  $T9P = 54.4\%$  and  $T10P = 76.2\%$ .

3. \$READ

The thirty-four parameters calculated via programs DATA, \$CALC1 and \$CALC2 and stored on LINC magnetic tape were retrieved and printed out by program \$READ for each steady state respiratory discharge. The computer listing of these parameters for cell C29UI7R(2) is found in Table II. All time measurements ( $TL$ ,  $CT$ ,  $SD_{ST}$ ,  $SE_{ST}$ ,  $FI$ ,  $SD_{FI}$ ,  $SE_{FI}$ ,  $MI$ ,  $MEN$ ,  $MED$ ,  $MOD$ ,  $PL$ ,  $LI$ ,  $T50$ ,  $T70$ ,  $T90$  and  $T100$ ) are standardized in milliseconds. Parameters that count events ( $NT$ ,  $ST$ ,  $FMN$ ,  $HSN$ ,  $MON$ ,  $N50$ ,  $N70$ ,  $N90$  and  $N100$ ) and parameters that are derived from the ratio of time measurements ( $SL$ ,  $T5P$ ,  $T7P$ ,  $T10P$ ,  $PLP$  and  $TLP$ ) are all dimensionless. The respiratory rate ( $RR$ ) is given in breaths per minute.

- 4. \$SUM1
- 5. \$SUM2
- 6. \$SUM3
- 7. \$SUM4

Similarly to program \$READ, these four programs printed out the thirty-four parameters calculated for each steady state observation, but in a modified format. That is, instead of listing the different parameters in one vertical column as in Table II, parameters were typed out in horizontal rows so that similar measurements from

Table II. Program \$READ computer printout derived from inspiratory cell C29UI7R(2) in Figure 4A.

DATA POINT 20	
TL	859
CT	1990
RR	30.1
NT	44
ST	46.7
SD	3.9
SE	0.58
FI	56.1
SD	11.1
SE	1.68
MI	12.0
FMW	2011
HSN	2000
REN	18.8
MED	14.8
MOD	13.2
MON	161
PL	368
LI	18.0
SL	- 0.34
T50	184
T70	270
T90	467
T100	655
N50	4.4
N70	7.5
N90	17.4
N100	31.0
T5P	21.4
T7P	31.4
T9P	54.4
T10P	75.2
PLP	42.8
TLP	43.2



consecutive steady state observations formed vertical columns. For example, parameters listed by programs \$SUM1 (RR, NT, ST, SD<sub>ST</sub>, SE<sub>ST</sub>, FI, SD<sub>FI</sub>, SE<sub>FI</sub>, MI), \$SUM2 (FMN, HSN, MEN, MED, MOD, MON, PL, LI), \$SUM3 (SL, T50, T70, T90, T100, N50, N70, N90, N100) and \$SUM4 (TL, CT, T5P, T7P, T9P, T10P, PLP, TLP) formed four tables of data. By following down any one of the thirty-four columns, alterations in a single parameter for any one cell could be traced and correlated with induced modifications of the breathing pattern.

8. \$REORD

The thirty-four parameters associated with any one steady state observation were located in sequential addresses on LINC magnetic tape. Since this was an inconvenient format for accessing similar parameter types in certain plotting procedures, program \$REORD was used to reorder all the parameters on new LINC magnetic tape. In effect, rows and columns were interchanged which placed similar parameter types in sequential addresses. Two recorded tapes containing inspiratory and expiratory cell data, respectively, were obtained by this procedure.

- 9. \$IVO
- 10. \$IVI
- 11. \$EVO
- 12. \$EVI

The reordered inspiratory and expiratory tapes that

were constructed via program \$REORD contained steady state measurements from both vagotomized and vagi-intact preparations. The responsibility of these four programs was to divide vagi-intact (VI) from vagi-out (VO) data points for both inspiratory (I) and expiratory (E) cells. This data manipulation of the two reordered types produced four new LINC tapes including inspiratory vagi-out (I-VO), inspiratory vagi-intact (I-VI), expiratory vagi-out (E-VO) and expiratory vagi-intact (E-VI) data groups. These four tapes were formatted as the reordered tapes and were used for statistical computations and regression plotting of raw data.

13. \$ROWM

This program calculated the mean, standard deviation (SD), standard error (SE), high and low values and range for each parameter group stored on LINC tapes I-VO, I-VI, E-VO, and E-VI. The standard deviation (SD) and standard error (SE) calculations were identical to those described in program \$CALC1. One hundred and thirty-six sets of statistical data (4 tapes · 34 parameters types/tape) were printed out by program \$ROWM.

- 14. \$RLIN
- 15. \$REXP
- 16. \$RLOG
- 17. \$RLLG

Linear and curvilinear regressions were obtained

between groups of similar parameter types taking two groups at a time. Excluding eight groups ( $SD_{ST}$ ,  $SE_{ST}$ ,  $SD_{FI}$ ,  $SE_{FI}$ , NT, FMN, HSN and MON), all possible combinations for the remaining twenty-six groups were examined using four different curve fits for each of the four tapes I-VO, I-VI, E-VO and E-VI ( $26^2 \cdot 4 \cdot 4 = 10,816$  regression comparisons). Coefficients for the four equations,

$$\text{fit 1} \quad Y = B + X \cdot M \pm E \quad (\text{linear})$$

$$\text{fit 2} \quad Y = \text{EXP}(B + X \cdot M \pm E) \quad (\text{exponential})$$

$$\text{fit 3} \quad Y = B + \ln(X) \cdot M \pm E \quad (\text{logarithmic})$$

$$\text{fit 4} \quad Y = \text{EXP}(B + \ln(X) \cdot M \pm E) \quad (\log_e - \log_e)$$

along with the correlation coefficient "r" and other statistical computations were derived by transforming data for curvilinear regressions,

$$Y = \ln(Y) \text{ for exponential fit}$$

$$X = \ln(X) \text{ for logarithmic fit}$$

$$Y = \ln(Y) \text{ and } X = \ln(X) \text{ for } \log_e - \log_e \text{ fit}$$

and performing the following linear regression analysis:

$$DX = \Sigma X^2 - (\Sigma X)^2/N$$

$$DY = \Sigma Y^2 - (\Sigma Y)^2/N$$

$$M = [\Sigma XY - (\Sigma X)(\Sigma Y)/N]/DX$$

$$B = [\Sigma Y - (\Sigma X)M]/N$$

$$r = \frac{M \cdot DX}{\sqrt{(DX)(DY)}} = M(SD_X/SD_Y)$$

$$S = DY - M^2 \cdot DX$$

$$E = \sqrt{S/(N - 2)}$$

$$V = r^2$$

where X and Y represent the independent and dependent variables, respectively, of the paired groups, N is the number of parameters in either group, M is the slope, B is the Y intercept, and "r" is the correlation coefficient ( $r \leq 1.00$ ). Note that "r" can be defined as slope (M) times the ratio of X to Y standard deviations. The standard error of estimate (E) defined a range of regression curves including 68% of the individual observations of Y on X. Assuming that the distribution about the regression line was normal and of equal variance, this error term corresponded to one standard deviation unit on either side of the mean regression line. The V value ( $r^2$ ) described the proportion of the variance of Y that could be attributed to its linear regression on X.

#### 18. \$REGSD

The significant difference between two regression slopes ( $M_1, M_2$ ) was tested by this program. The following calculations applied only to regressions of the same fit (i.e. linear, exponential, logarithmic or log-log):

$$S = (S_1 + S_2)/(N_1 + N_2 - 4)$$

$$DI = \sqrt{S/(DX_1 + DX_2)}$$

$$t = (M_1 - M_2)/DI$$

where S, N, DX and M values were obtained from the regression programs and "t" is the standard Student "t" value with degrees of freedom equal to  $(N_1 + N_2 - 4)$ . The "t" values were converted into P coefficients in the program, eliminating the use of standard statistical tables.

#### 19. \$RSIGD

This program calculated the significant difference between the regression coefficients of two regression equations independent of fit. The following calculations were made:

$$Z_1 = [\ln(1 + r_1) - \ln(1 - r_1)]/2$$

$$Z_2 = [\ln(1 + r_2) - \ln(1 - r_2)]/2$$

$$C = \frac{|Z_1 - Z_2|}{\sqrt{\frac{1}{N_1 - 3} + \frac{1}{N_2 - 3}}}$$

where  $r_1$  and  $r_2$  are the two regression coefficients being compared,  $N_1$  and  $N_2$  are the number of X-Y pairs in fit<sub>1</sub> and fit<sub>2</sub>, respectively, and the C value is correlated with P levels of significance from the normal distribution curve (not the "t" distribution). The values for P were computed for the following levels:  $>0.05$ ,  $\leq 0.05$ ,  $\leq 0.04$ ,

$\leq 0.03$ ,  $\leq 0.02$  and  $\leq 0.01$ .

20. \$PVALR

Correlation coefficients were converted into P levels of significance by first computing Student "t" values.

For example,

$$t = r \sqrt{N - 2} / \sqrt{1 - r^2}$$

where r is the correlation coefficient and N is the number of X-Y pairs correlated. As in program \$REGSD, "t" values were converted directly into P coefficients.

21. \$STTEST

The standard Student "t" test was programmed to calculate P values for miscellaneous paired or unpaired data. The "t" value was calculated as follows:

$$DM = \sqrt{(SE_1)^2 + (SE_2)^2 - 2r(SE_1)(SE_2)}$$

$$t = |MN_1 - MN_2| / DM$$

$$DF \text{ paired} = (N_1 + N_2 - 2) / 2 = N - 1$$

$$DF \text{ unpaired} = N_1 + N_2 - 2$$

where  $SE_1$  and  $SE_2$  are the standard errors of the two sets of data, "r" is the regression coefficient for paired data ( $r = 0$  for unpaired data),  $MN_1$  and  $MN_2$  are the means for each data set and N is the number of observations in each set ( $N_1 = N_2$  for paired data). The P values were computed automatically from the "t" values.

## 22. \$TIMFOL

This program was designed to compute the time elapsed between two digital tape counter values on the tape recorder. Since the tape counter was driven from the take-up reel, the digital read-out was a non-linear function of time. For this reason, a calibration curve was constructed in which the tape counter value was plotted as a function of time (0 - 100 minutes). This curvilinear plot was broken up into nine linear segments from which accurate time durations could be interpolated from two tape counter values. With this program, the length of time cells were recorded and the time between events could be computed, eliminating the need for hand computations from polygraph records.

D. FOCALPL Programs

## 1. %PLOT

Groups of data, correlated via regression calculations (programs \$RLIN, \$REXP, \$LOG and \$LLG) from LINC magnetic tape data (I-VO, I-VI, E-VO and E-VI), were plotted in scattergram format on the Calcomp digital plotter by program %PLOT. The independent (X) and dependent (Y) variables plotted were under manual selection and scaling factors were determined on the basis of program \$ROWM range values for each data set. All regression

plots were transformed to linear coordinates and six plots were placed on each page for space conservation. Program %PLOT is listed in the Appendix along with the other FOCALPL programs.

2. %REGP1

Program %REGP1 superimposed mean regression lines and standard error of estimate lines on the scattergram plots of program %PLOT. For any one plot, scaling factors and type of fit were identical. Coefficients for the linear, exponential, logarithmic and  $\log_e$ - $\log_e$  equations (fit 1-fit 4) were obtained from the regression data of programs \$RLIN, \$REXP, \$RLOG and \$RLLG respectively. All equations were plotted on a linear scale thus rendering exponential, logarithmic and  $\log_e$ - $\log_e$  fits curvilinear. Because of this, error lines above and below the mean regression line were not always parallel with or symmetrical to the latter as with linear regressions.

3. %REGP2

This program was used to superimpose mean regression lines on the same graph for visual comparison. To avoid confusion, scattergram points of program %PLOT and error lines of program %REGP1 were omitted. Six sets of multiple graphs were condensed to one page. This plotting procedure facilitated data reduction and was useful for



data summarization.

4. %LINEP

Induced changes in single unit parameters followed digitally by programs \$SUM1, \$SUM2, \$SUM3 and \$SUM4 were plotted in time sequence by program %LINEP (line plot) for each respiratory cell that was examined for more than one steady state period. With the exception of cycle time (CT), the twenty-five parameters from the regression analysis (T50, T70, T90, T100, PL, TL, FI, MI, LI, MEN, MED, MOD, RR, N50, N70, N90, N100, ST, SL, T5P, T7P, T9P, T10P, PLP and TLP) were plotted on nine graphs (three pages) for each cell. The X scaling was proportioned according to the number of different steady state observations made for each cell and was a non-linear function of time.

## CHAPTER V

### RESULTS

#### A. Data Presentation

Single cell studies on the neural control of respiration have been organized into three categories which are presented in the following order. First, parameter measurements from similar cell types have been grouped from many cells for regression analysis. This was an attempt to describe general population characteristics of discharge patterns at different respiratory rates and depths. Second, changes in single unit parameters have been followed during induced modification of the spontaneous breathing pattern. Comparison of these single cell data with population regression data revealed the degree of discharge pattern heterogeneity among similar cell types. Third, the correlation between high barbiturate levels and apneusis has been examined at the single cell level. These data were important for interpretation of population and single cell data obtained after barbiturate

modification of the spontaneous breathing pattern.

B. Population Data

The general statistics of the single cell recordings attempted in 60 out of 85 cats used in this study are summarized in Table III. On the average, approximately three respiratory cells were recorded from each cat in a ratio of two inspiratory cells to one expiratory cell. Of these recorded cells, however, over fifty percent were rejected from final data compilation due to generation of uneven interspike interval modulation curves from poor signal-to-noise ratios or non-uniform breathing patterns.

Steady state computer averages were divided into four categories of inspiratory vagi-intact (I-VI), inspiratory vagi-out (I-VO), expiratory vagi-intact (E-VI) and expiratory vagi-out (E-VO) observations. Cells recorded during unilateral vagotomy were categorized with vagi-intact data, since no significant respiratory rate or depth changes were manifest after removal of one vagus nerve.

Each steady state average was identified by a data point number. There was a total of 333 data points each of which contributed 34 parameter measurements ( $333 \times 34 = 11322$  parameters) for regression analysis. An average of four data points (range = 1-17) was obtained from each

Table III. General statistics of respiratory cells analyzed by computer techniques.

<u>Cell Type</u>	<u># Cells Recorded</u>	<u># Cells Analyzed</u>	<u>State of Vagi</u>	<u># Data Points</u>		<u>Analysis Time (Min)</u>		<u># Trains (NT)</u>		<u># Intervals (FMN)</u>	
				<u>Total</u>	<u>Mean</u>	<u>Total</u>	<u>Mean</u>	<u>Total</u>	<u>Mean</u>	<u>Total</u>	<u>Mean</u>
I	125	57	VI	175	3.4	188.7	1.1	6079	34.7	256111	1463
			VO	50	5.0	78.0	1.6	1159	23.2	61329	1227
E	57	26	VI	105	4.6	107.7	1.0	3776	36.0	119825	1141
			VO	3	1.0	3.5	1.2	64	21.3	2543	848
Sum Total	182	83		333		377.9		11078		439808	
Grand Mean					4.0		1.1		33.3		1321

respiratory cell. Respiratory cells were recorded for a period of time ranging from 30 seconds to three hours. Steady state averages were derived from mean breathing segments of 1.1 minutes. For all the cells, this analysis time represented more than six hours of spontaneous breathing or more than 25 hours of computer time taking into account the time scaling factor of 4. Derivation of the 333 data points involved over eleven thousand individual breaths and close to one half million interspike intervals. Average interspike interval modulation curves were generated from 33 burst discharges and more than thirteen hundred intervals.

As described in Data Analysis, similar parameter types were grouped together and all possible regression comparisons taking two groups at a time were performed. Interest centered around two regression types in which respiration rate (RR) and mode (MOD) parameters were selected as the independent variables. The respiration rate (RR) parameter, derived from the reciprocal of the cycle time, was a direct measurement of the spontaneous breathing frequency.

The mode (MOD) parameter reflected changes in respiratory depth (or tidal volume assuming constant airflow resistance) as illustrated in Figure 10 for an inspiratory

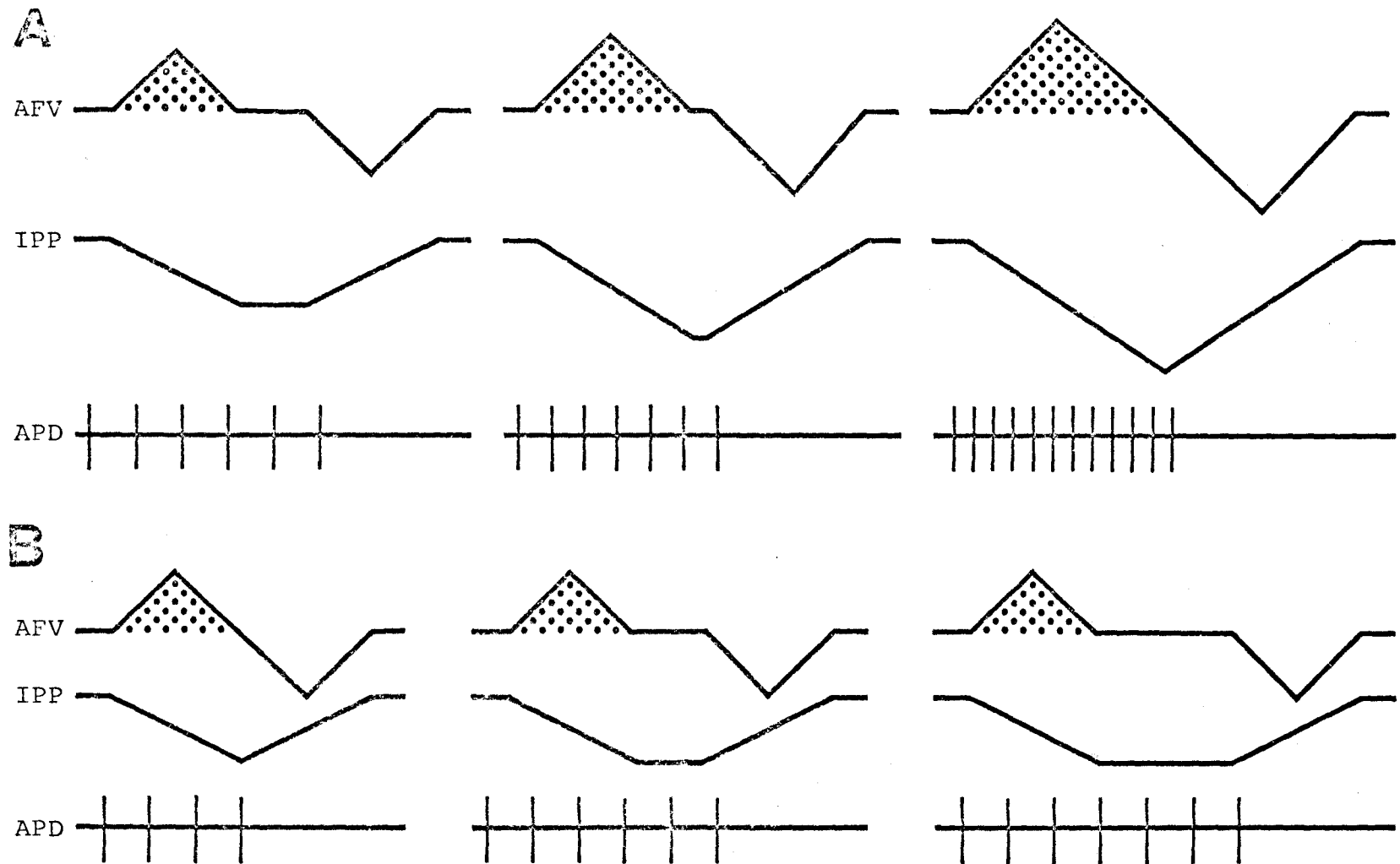


Figure 10. Idealized representation of airflow velocity (AFV), intrapleural pressure (IPP) and action potential discharge (APD) during: A. constant respiratory rate, increasing depth; B. constant depth, decreasing respiratory rate.

cell. In Figure 10A, for example, a decrease in mode (MOD) time is associated with an increased intrapleural pressure and increased tidal volume (shaded area under airflow curve) with no change in respiratory rate ( $RR = k$ ). Figure 10B diagrams a hypothetical situation where a rate (RR) change is observed in the absence of any depth or tidal volume change. In such a case, the mode (MOD) parameter can be shown to be independent of rate (RR) changes.

This reciprocal relationship between mode (MOD) time and depth or tidal volume ( $TV \propto 1/MOD$ ) is not unusual since it has been established for both phrenic (39, 50, 53, 58) and central inspiratory cells (72, 129), but does make the assumption that no recruitment of units takes place, at least at the central level. Nesland et al. (129) deny that respiratory neurons are recruited, but this can be debated. The mode (MOD) time was selected over the mean (MEN) or median (MED) times since the former falls within the 100 millisecond segment of the phrenic traffic coincident with the peak of inspiration, integration of which yields the best neural correlate of tidal volume (53). The mode (MOD) is also by definition the most common or frequent interval in the average respiratory train.

For expiratory cells, the reciprocal relationship

between mode (MOD) and tidal volume is more involved. Nevertheless, higher frequency discharges (lower mode times) are expected to be found at larger tidal volumes due to elevated afferent feedback, increased active expiratory movements and postulated higher inhibitory thresholds on inspiratory neurons.

The thirty four parameters defined and measured in this study successfully quantitated all phases of a respiratory neuronal spike discharge. The respiratory train or interspike interval modulation curve (Figure 9B) was segmented into initial (frequency increase), middle (frequency plateau) and terminal phases (frequency decrease). Examination of parameter changes during induced alterations in the spontaneous breathing pattern revealed selective modification of certain phases over others. For example, changes in SL, FI, T50, T70, N50, N70, T5P and T7P reflected changes in the initial phase. Changes in MI, T90, T100, N90, N100, T9P, T10P, PL and PLP reflected changes in the middle phase. Changes in LI, TL and TLP reflected changes in the terminal phase of spike activity. The reader should keep the parameters in this type of perspective when following parameter modification at different spontaneous respiratory rates (RR) and depths (1/MOD). The thirty four parameters have been



tabulated in Table IV for convenient reader reference. Both code identification and definition as used throughout the dissertation are listed for each parameter.

Regression plots of 25 dependent variables (\* Table IV) on respiration rate (RR) and mode (MOD) are presented in Figures 11-22 and Figures 23-34 respectively. Three plots for two dependent variables are organized from left to right in each figure for inspiratory vari-out (I-VO), inspiratory vagi-intact (I-VI) and expiratory vagi-intact (E-VI) data respectively. Expiratory vagi-out data (E-VO) have been deleted since only three data points from three expiratory cells were located in vagotomized preparations (cf. Table III). All scaling factors are given for X and Y axes which extend from border to border. The respiration rate (RR) scaling of Figures 11-22 and the mode (MOD) scaling of Figures 23-34 are consistent from plot to plot (X-axis). For any one figure, the scalings of dependent variables (Y-axis) are identical for each of the six plots. Exceptions are found in Figures 19, 28, and 31 where the different dependent parameters have different scaling factors. In these cases, the I-VO, I-VI and E-VI scalings are the same for any one parameter, however. Each axis is labeled with parameter type and appropriate dimension

Table IV. List of parameter abbreviations.

1	CT	Cycle Time
2	T50*	Time to 50%
3	T70*	Time to 70%
4	T90*	Time to 90%
5	T100*	Time to 100%
6	PL*	Plateau Level
7	TL*	Train Length
8	RR*	Respiration Rate
9	ST*	Spikes per Train
10	FI*	First Interval
11	MI*	Minimum Interval
12	LI*	Last Interval
13	MEN*	Mean
14	MED*	Median
15	MOD*	Mode
16	N50*	# of Intervals to 50%
17	N70*	# of Intervals to 70%
18	N90*	# of Intervals to 90%
19	N100*	# of Intervals to 100%
20	T5P*	T50 Percent
21	T7P*	T70 Percent
22	T9P*	T90 Percent
23	T10P*	T100 Percent
24	PLP*	Plateau Level Percent
25	TLP*	Train Length Percent
26	SL*	Slope
27	SEST	Standard Error (ST)
28	SEFI	Standard Error (FI)
29	SDST	Standard Deviation (ST)
30	SDFI	Standard Deviation (FI)
31	NT	# of Trains
32	FMN	# of Intervals in Interspike Interval Modulation Curve
33	HSN	# of Intervals in Histogram
34	MON	# of Intervals at Mode

\*Parameter plotted in Figures 11-43.

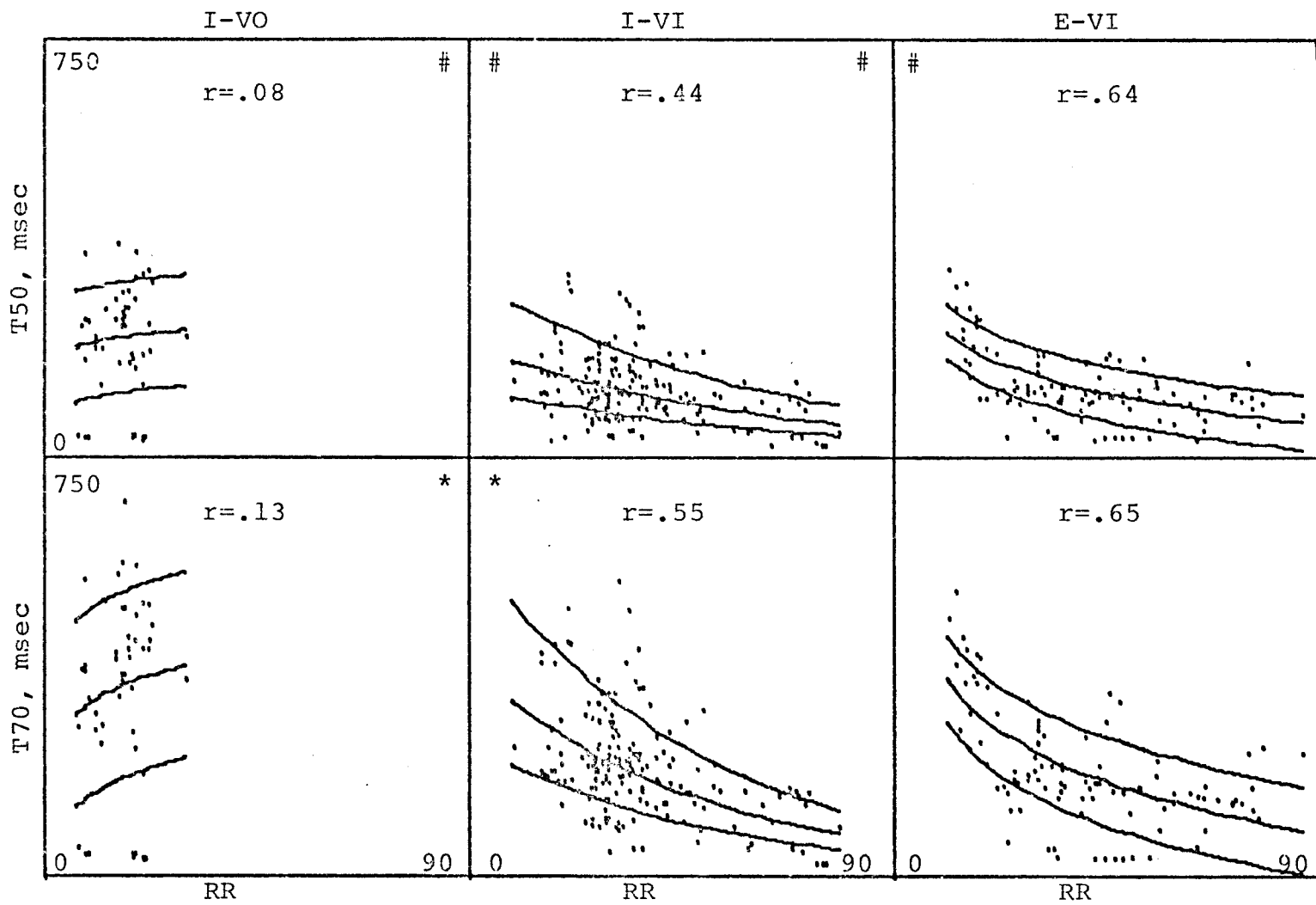


Figure 11. Regression of T50 and T70 on RR for I-VO, I-VI and E-VI cells.

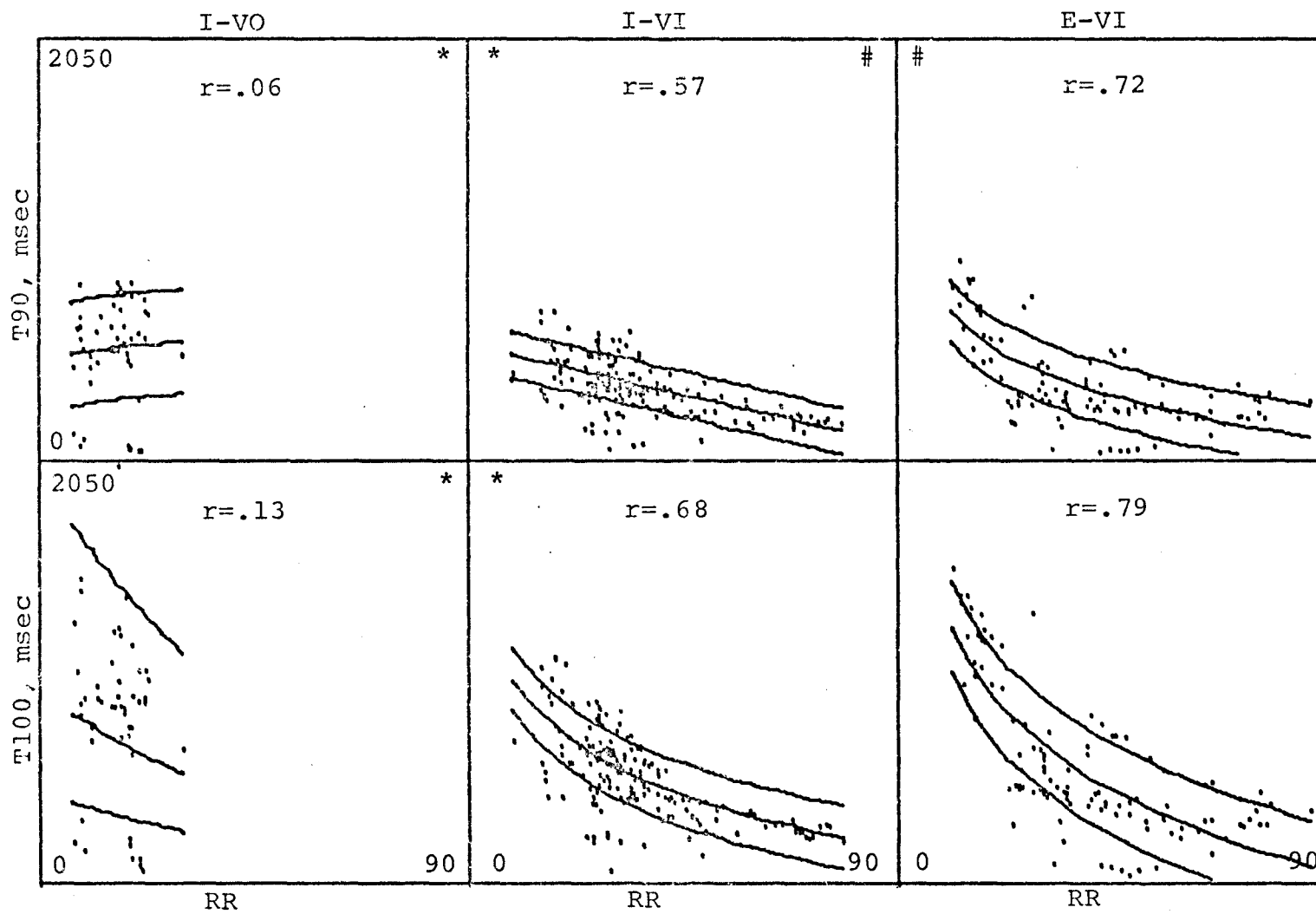


Figure 12. Regression of T90 and T100 on RR for I-VO, I-VI and E-VI cells.

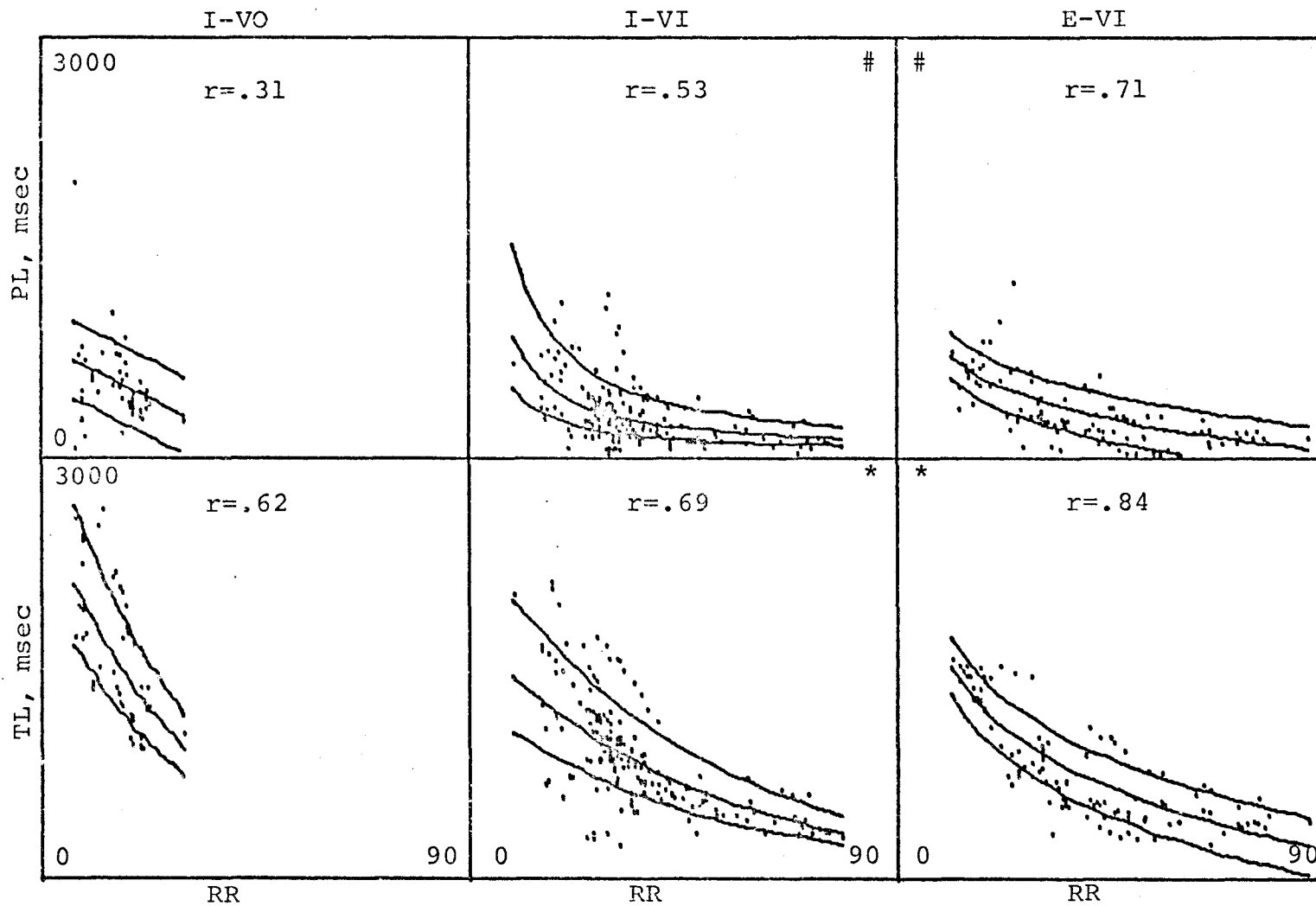


Figure 13. Regression of PL and TL on RR for I-VO, I-VI and E-VI cells.

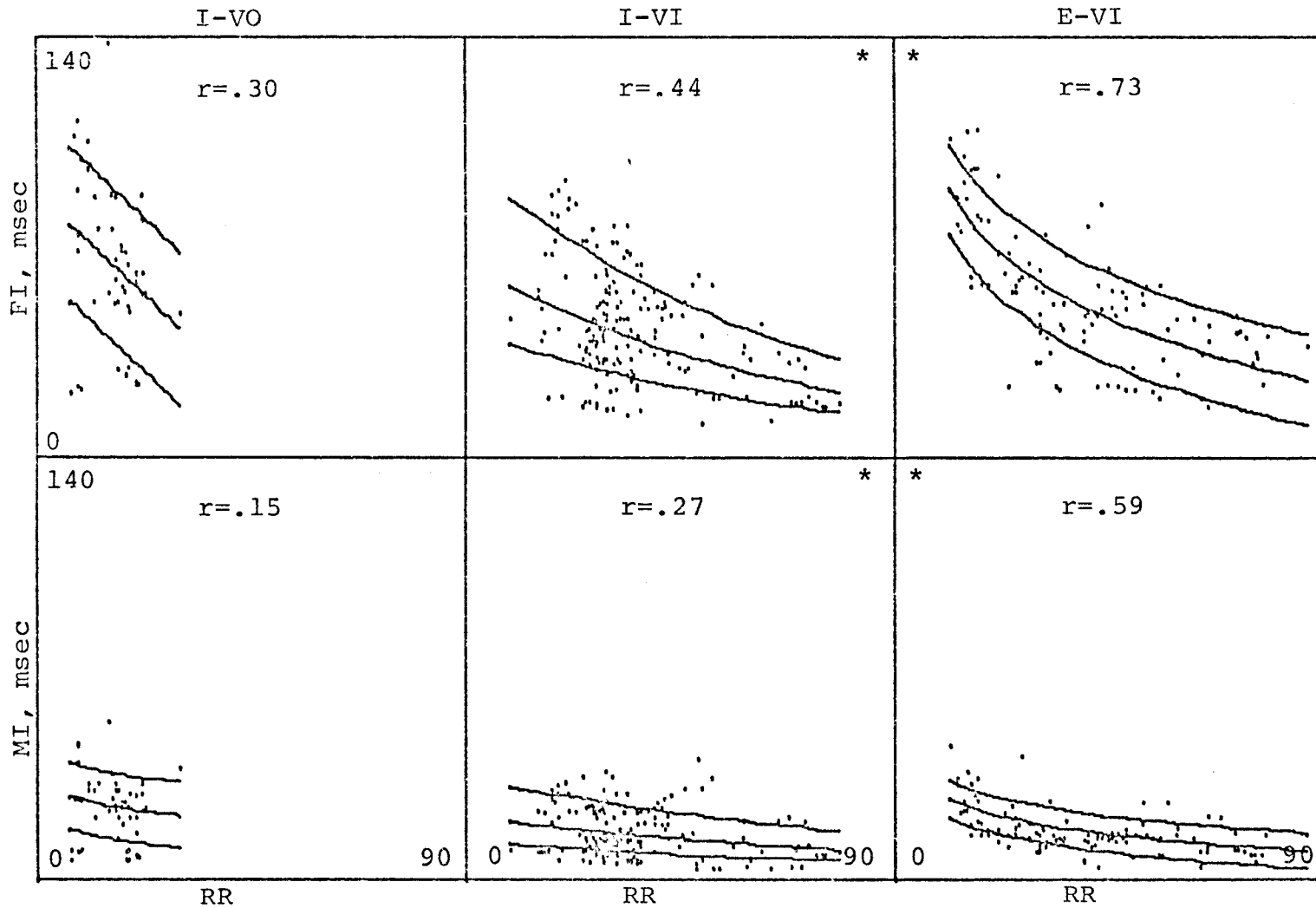


Figure 14. Regression of FI and MI on RR for I-VO, I-VI and E-VI cells.

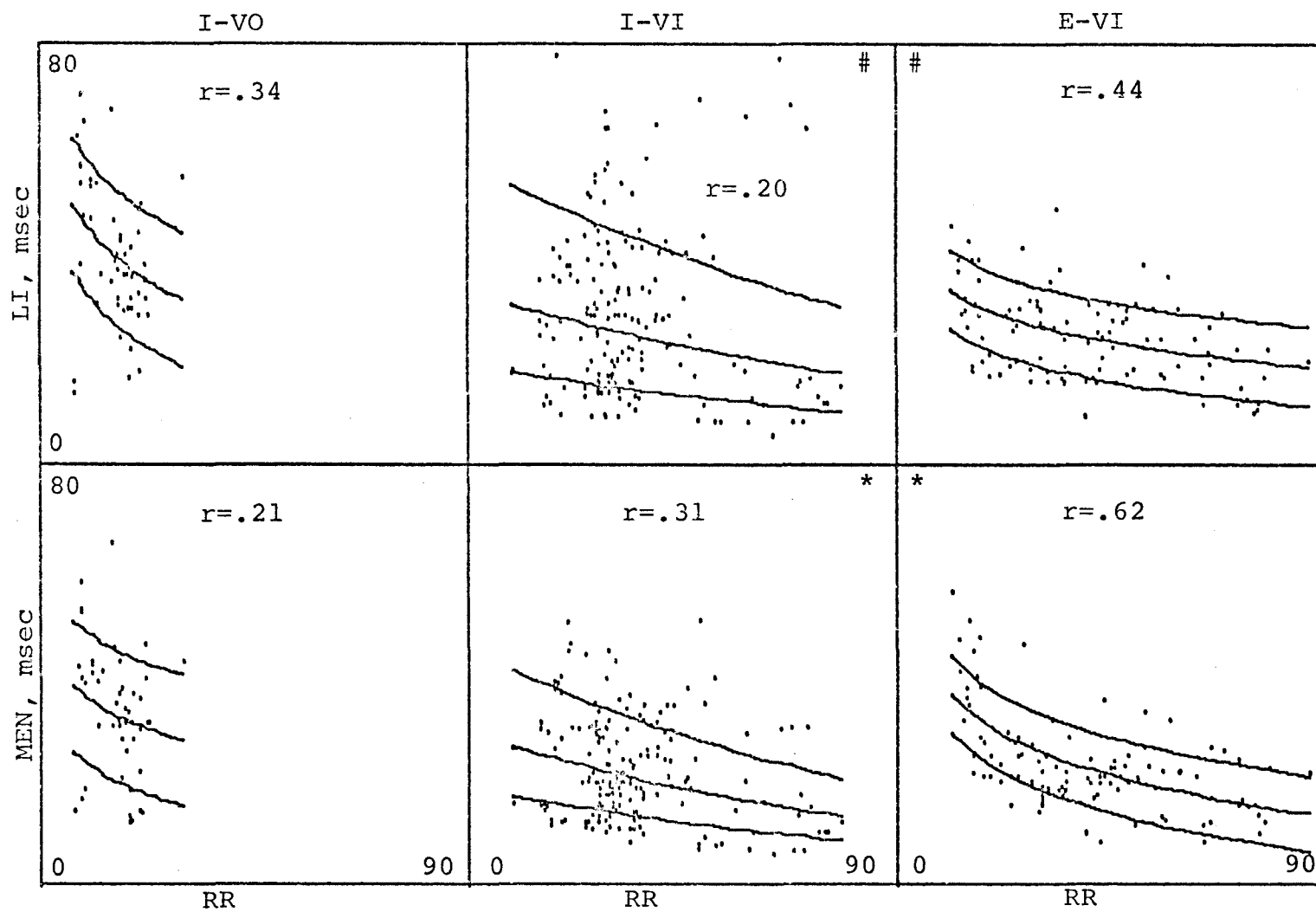


Figure 15. Regression of LI and MEN on RR for I-VO, I-VI and E-VI cells.

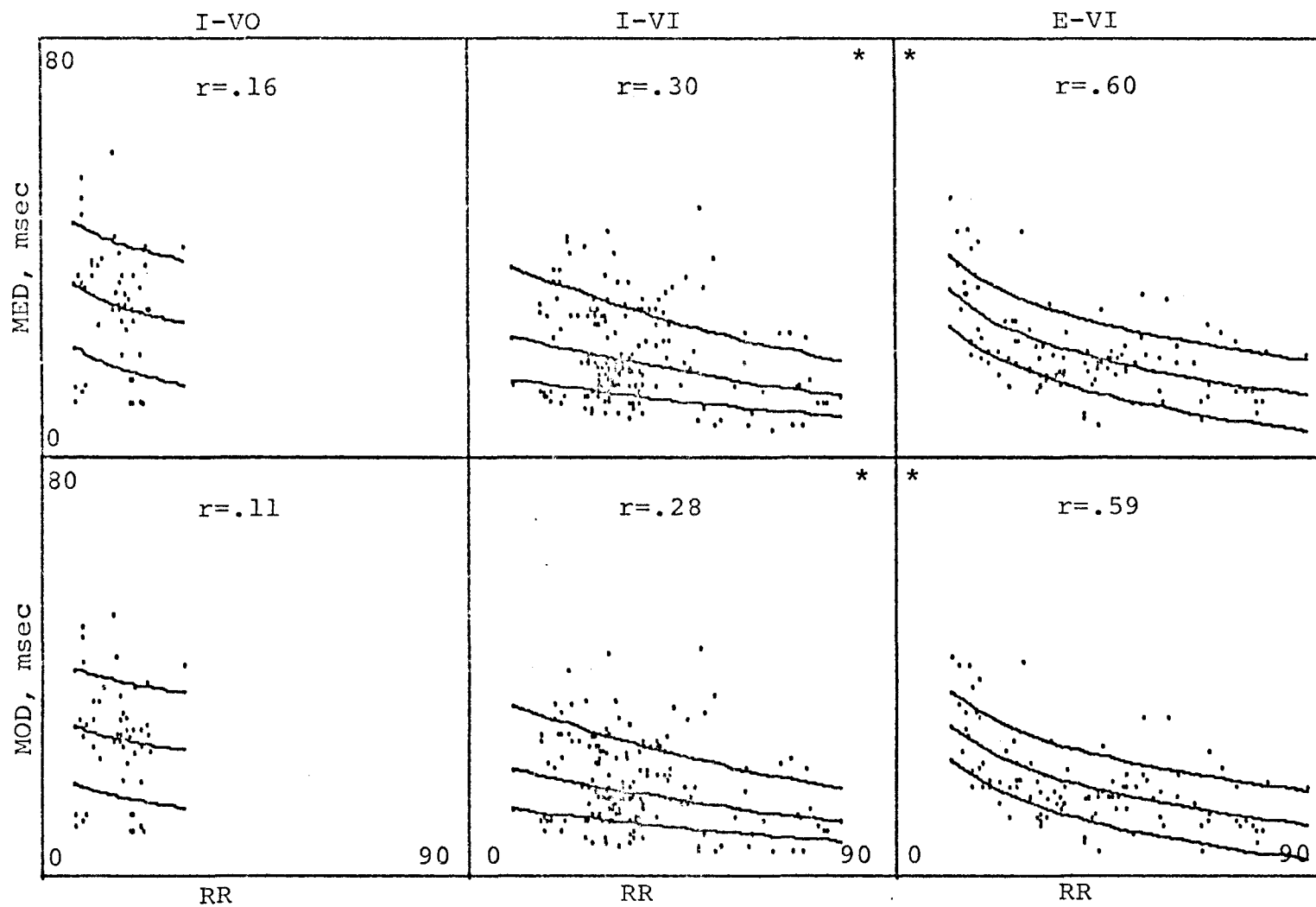


Figure 16. Regression of MED and MOD on RR for I-VO, I-VI and E-VI cells.



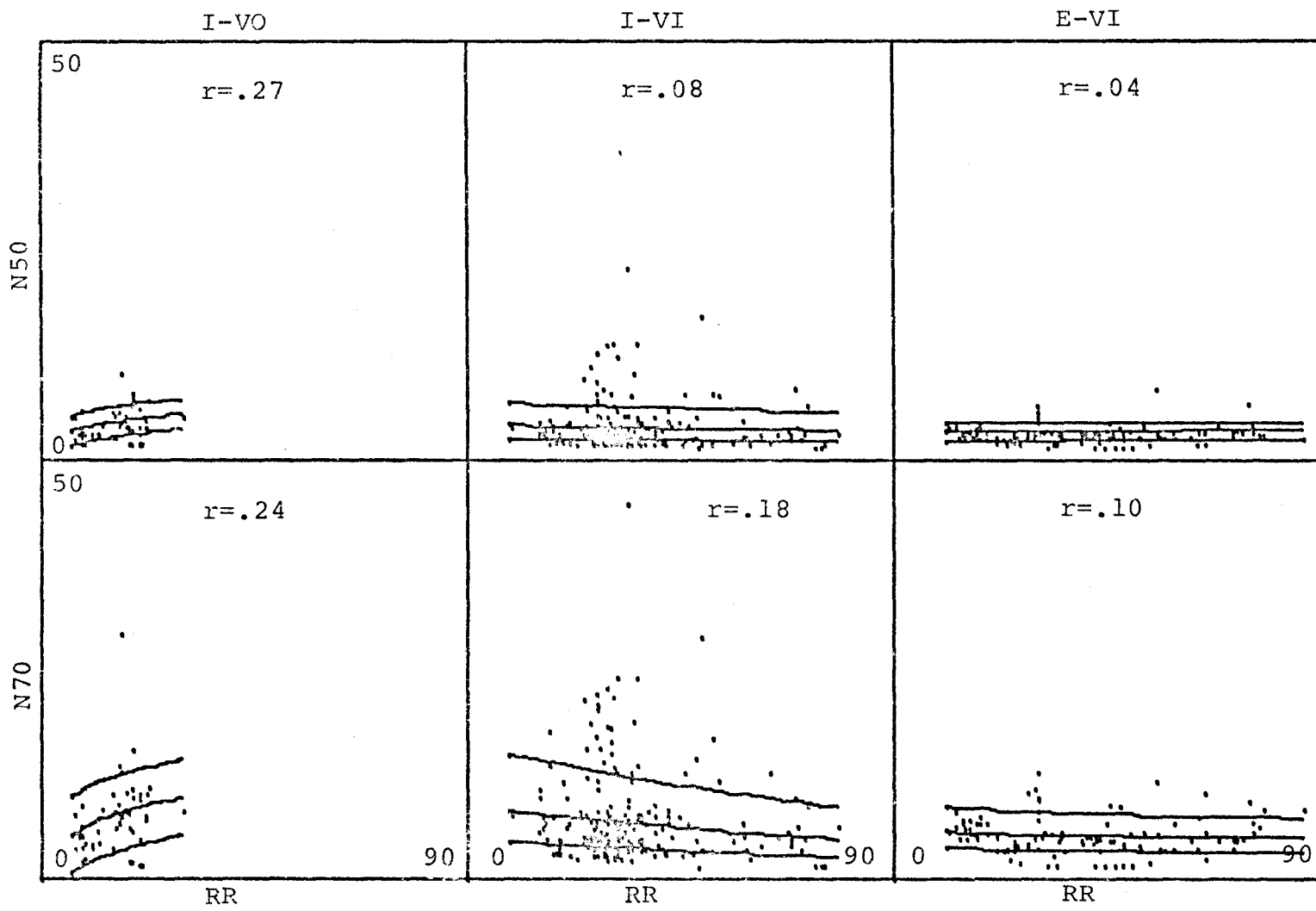


Figure 17. Regression of N50 and N70 on RR for I-VO, I-VI and E-VI cells.

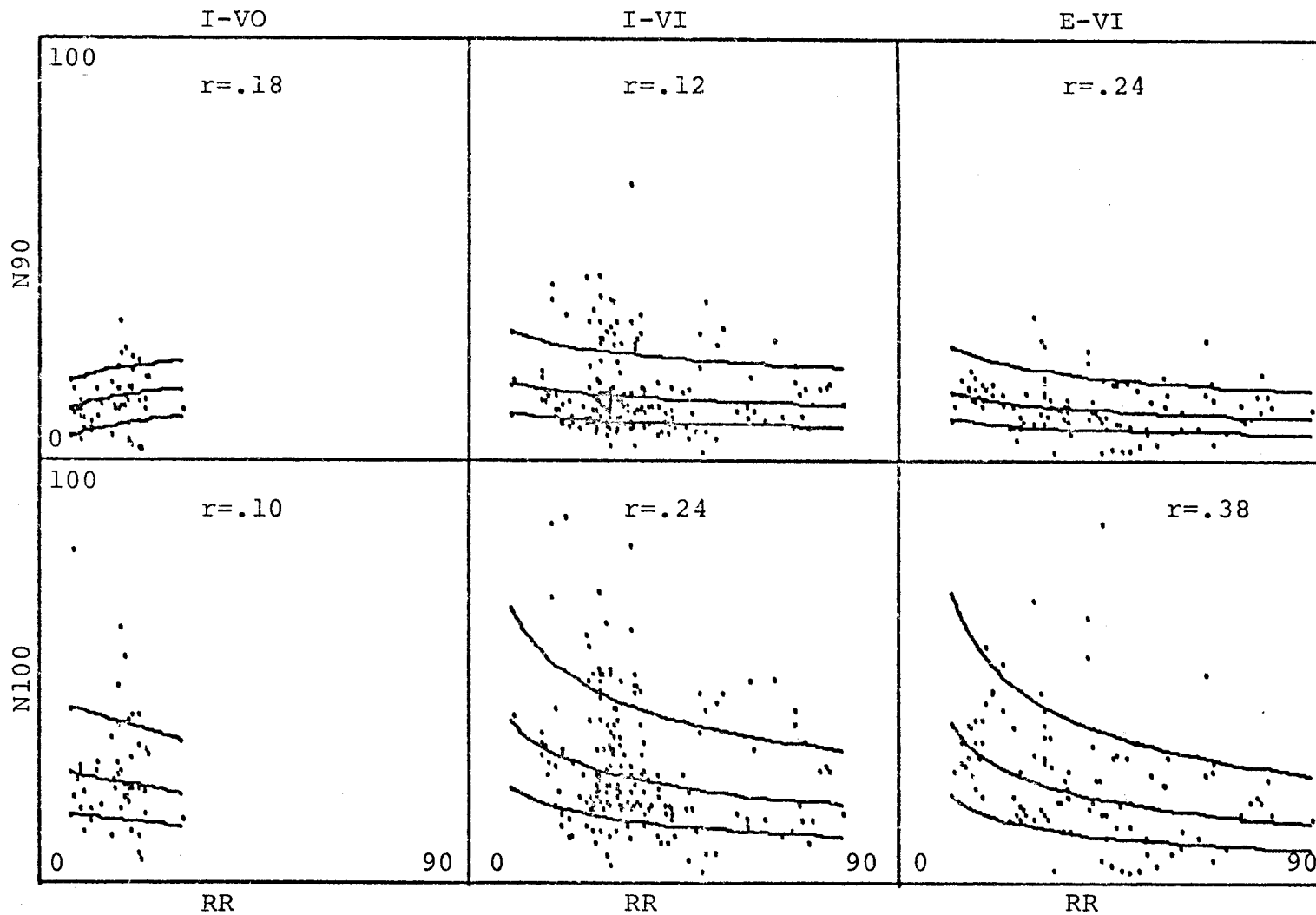


Figure 18. Regression of N90 and N100 on RR for I-VO, I-VI and E-VI cells.

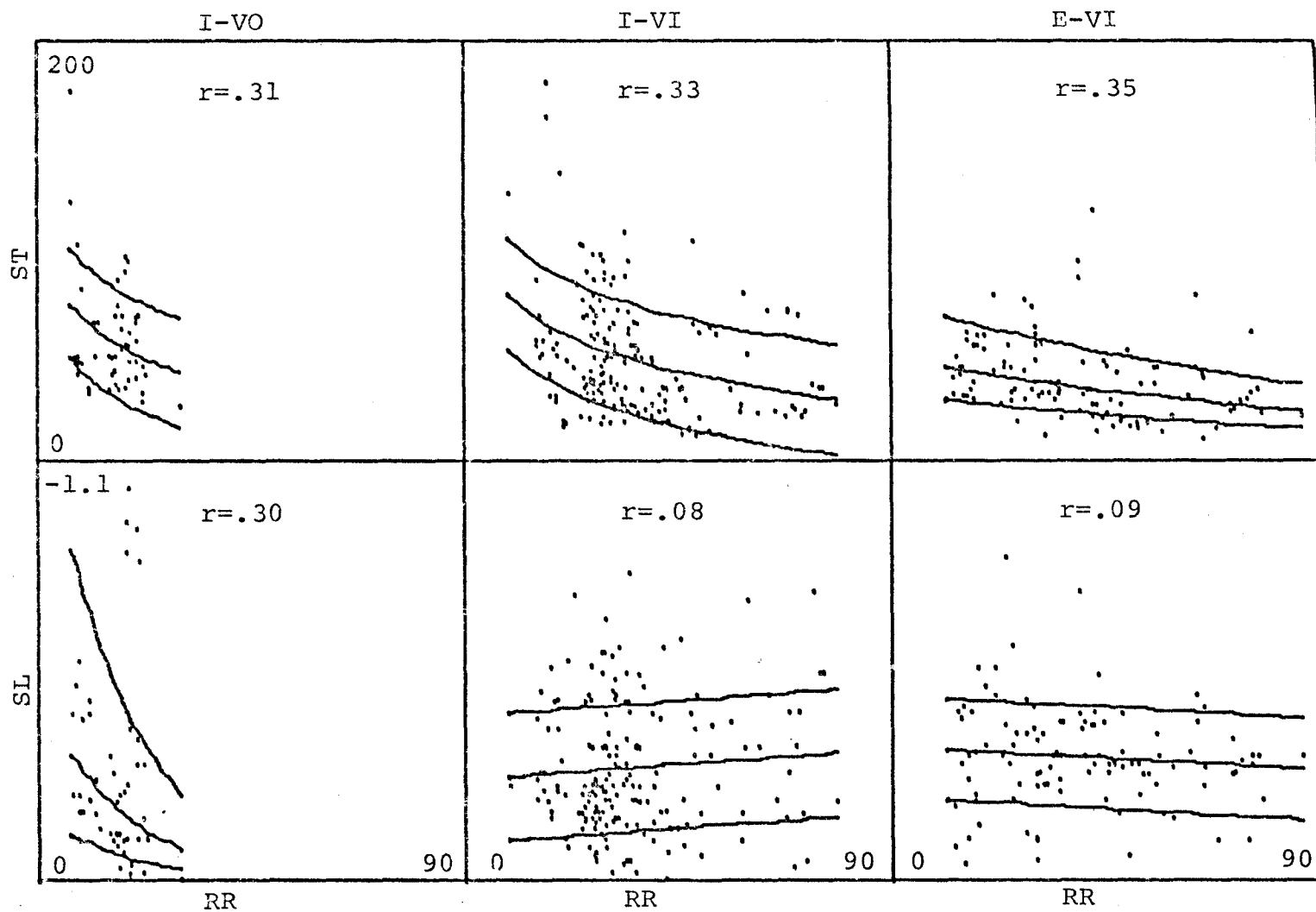


Figure 19. Regression of ST and SL on RR for I-VO, I-VI and E-VI cells.

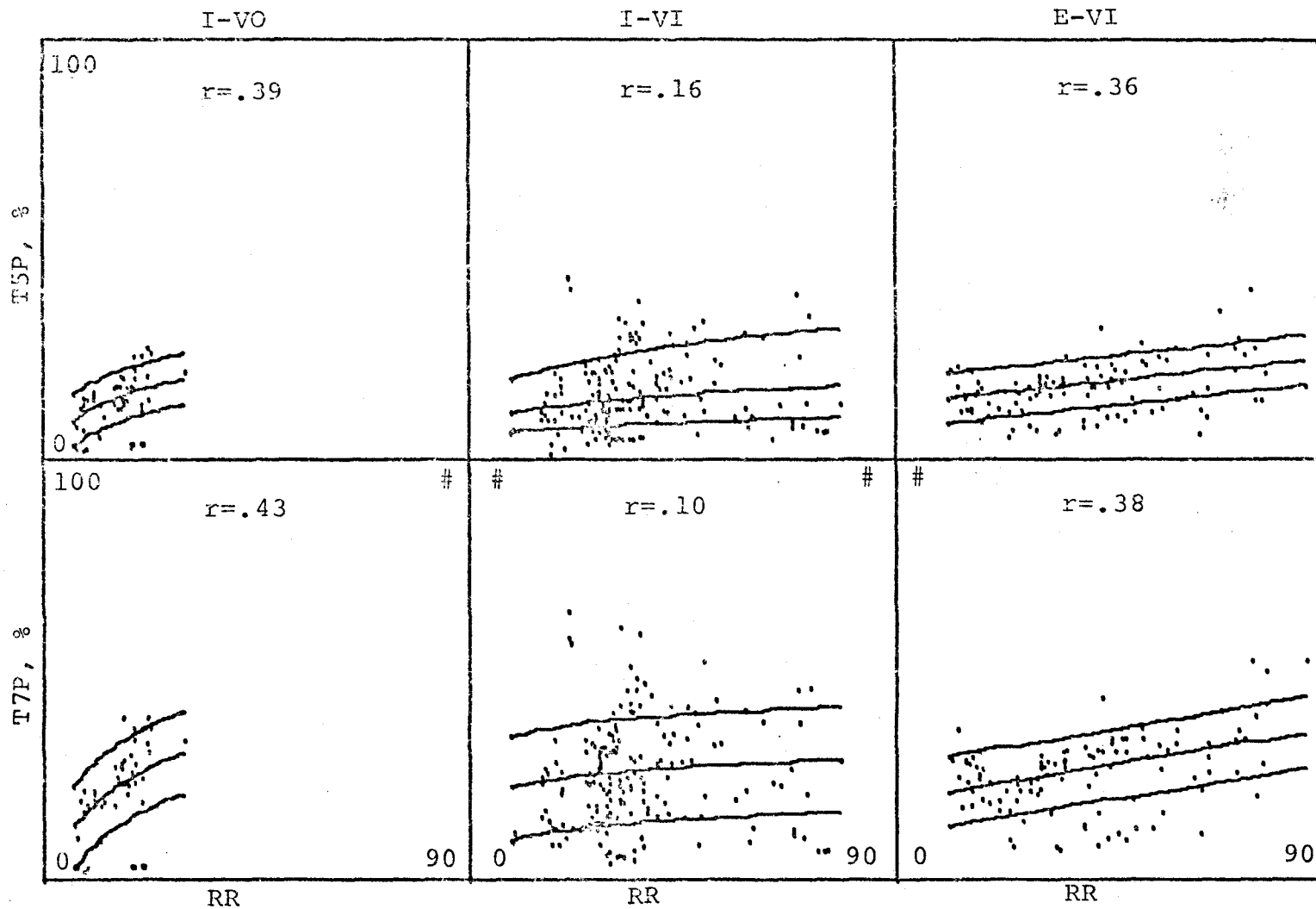


Figure 20. Regression of T5P and T7P on RR for I-VO, I-VI and E-VI cells.

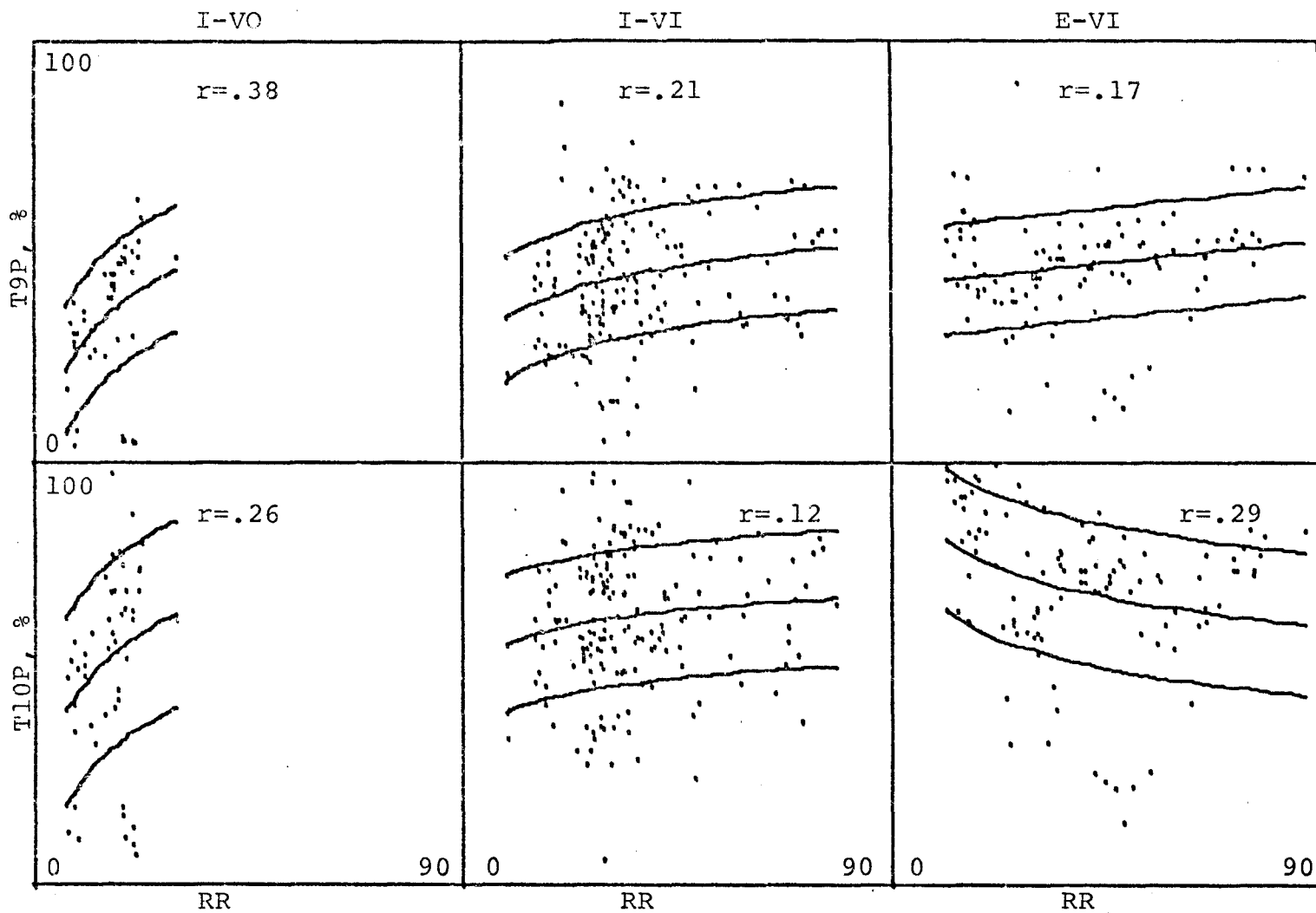


Figure 21. Regression of T9P and T10P on RR for I-VO, I-VI and E-VI cells.

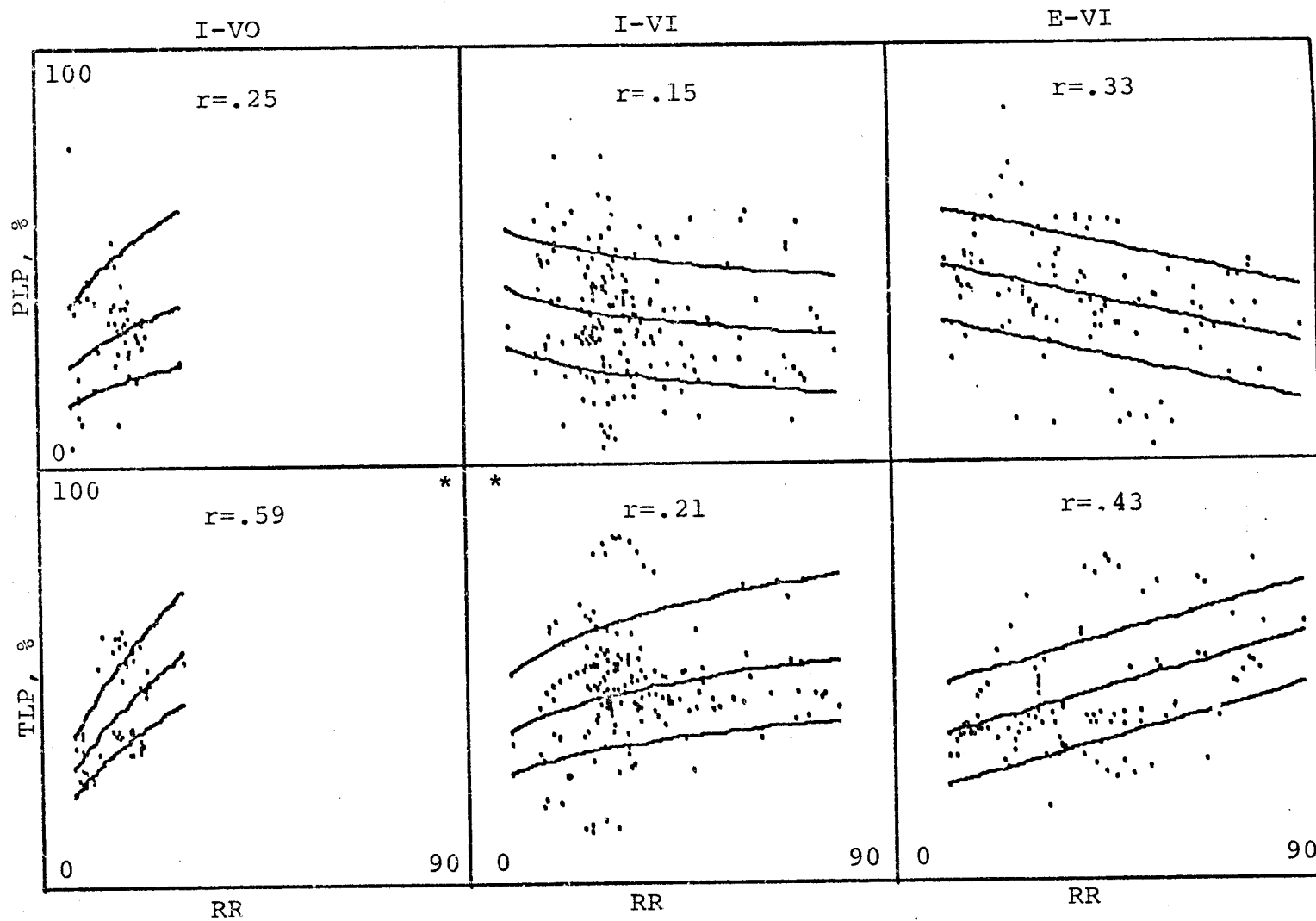


Figure 22. Regression of PLP and TLP on RR for I-VO, I-VI and E-VI cells.

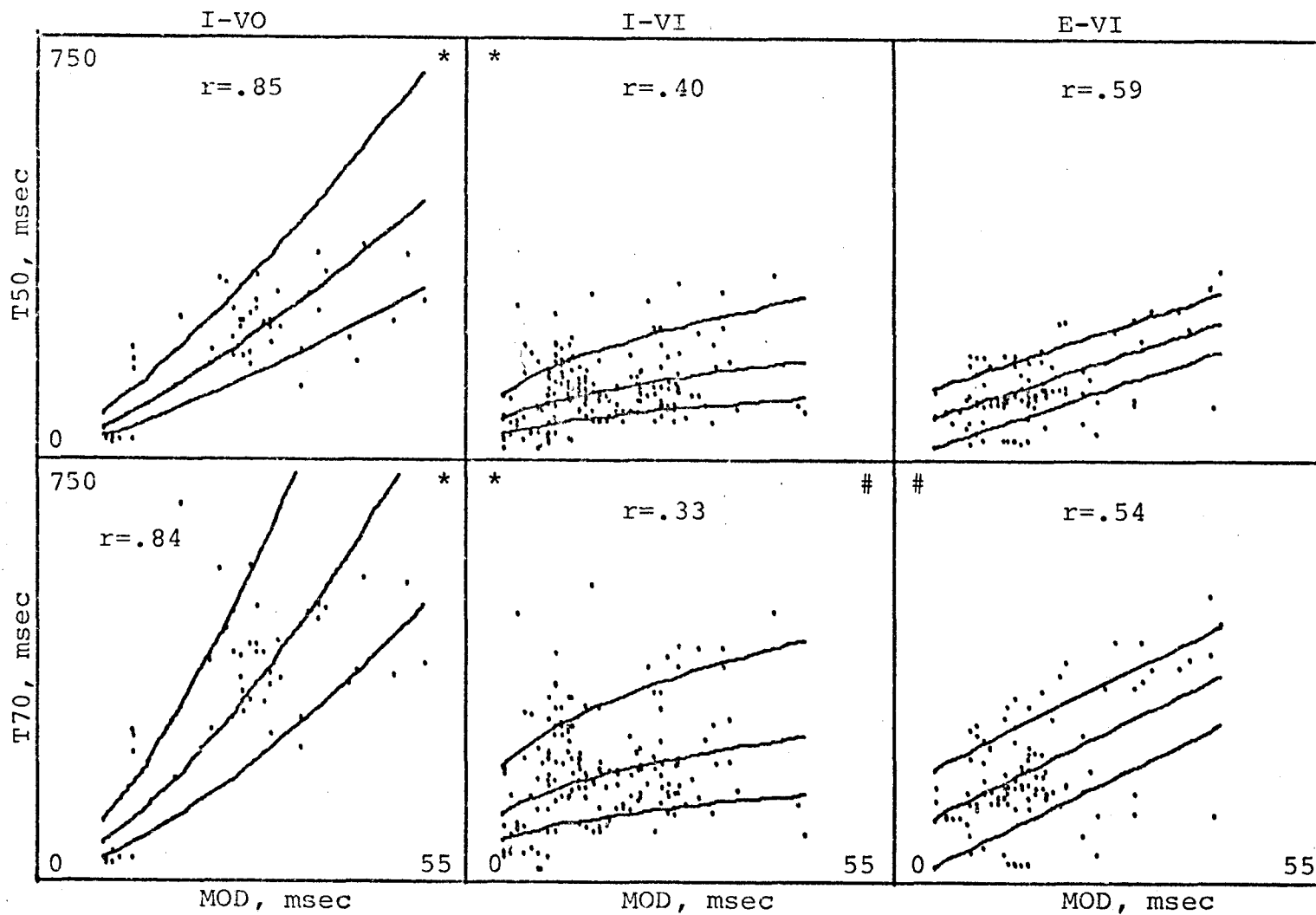


Figure 23. Regression of T50 and T70 on MOD for I-VO, I-VI and E-VI cells.

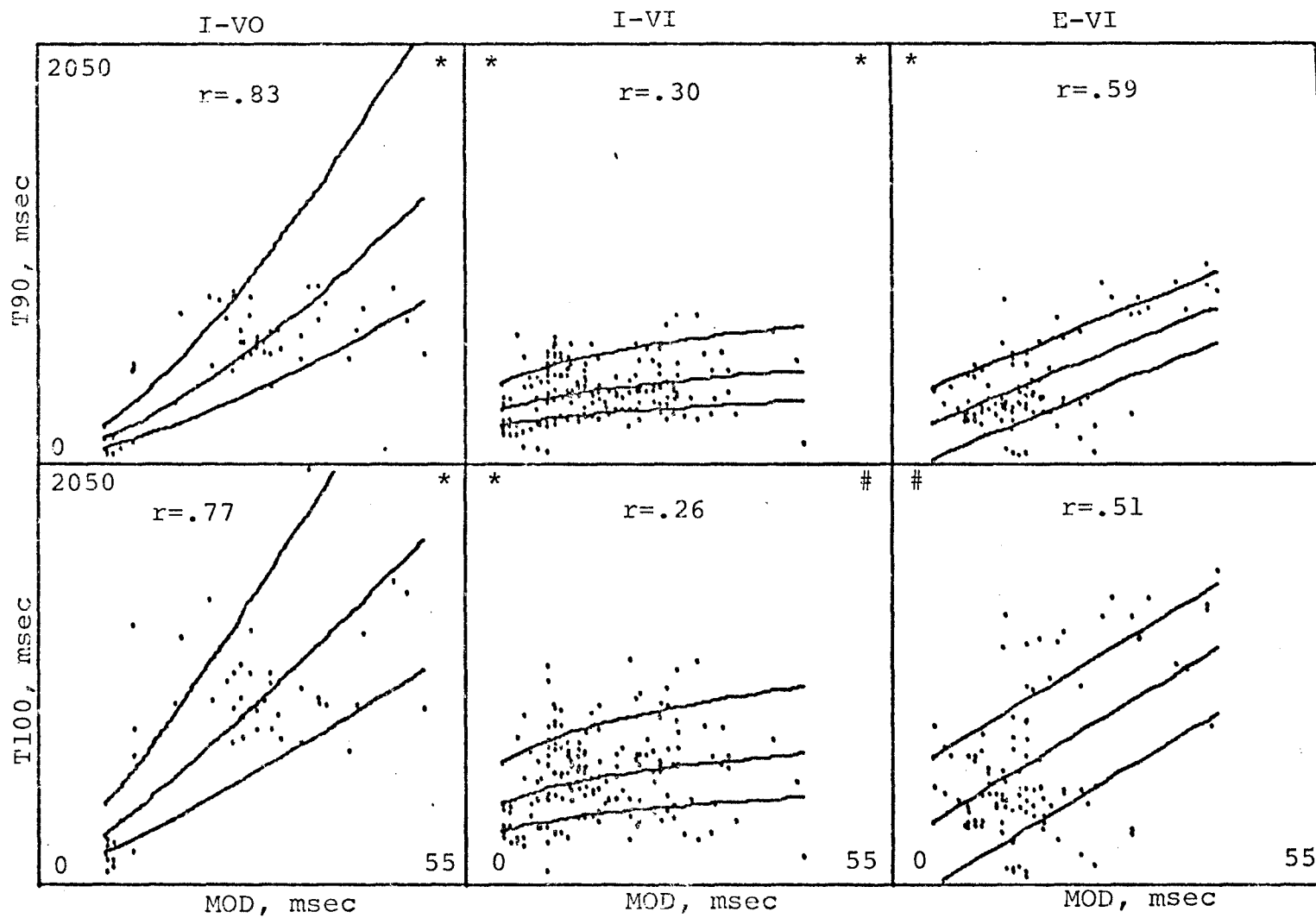


Figure 24. Regression of T90 and T100 on MOD for I-VO, I-VI and E-VI cells.



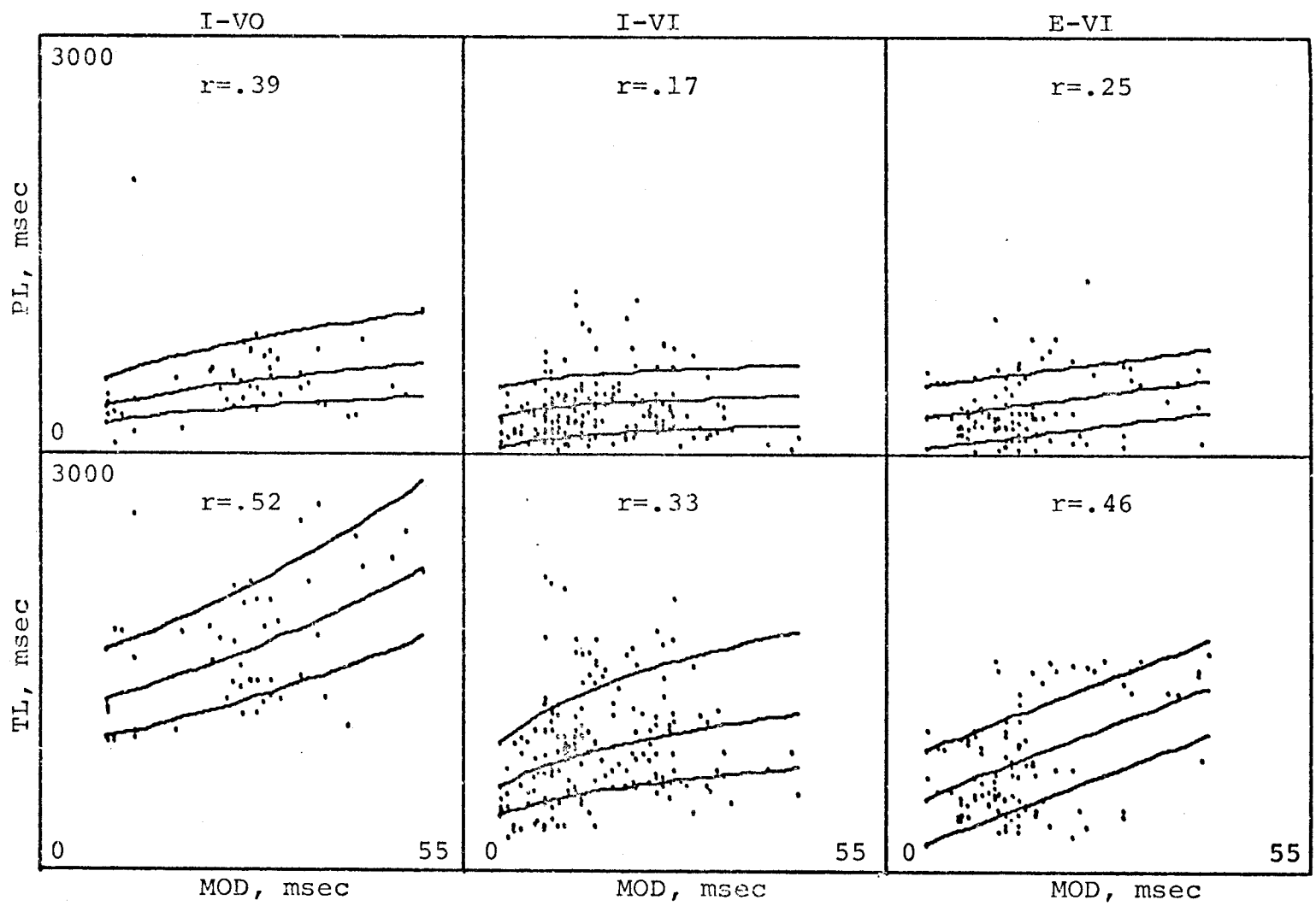


Figure 25. Regression of PL and TL on MOD for I-VO, I-VI and E-VI cells.

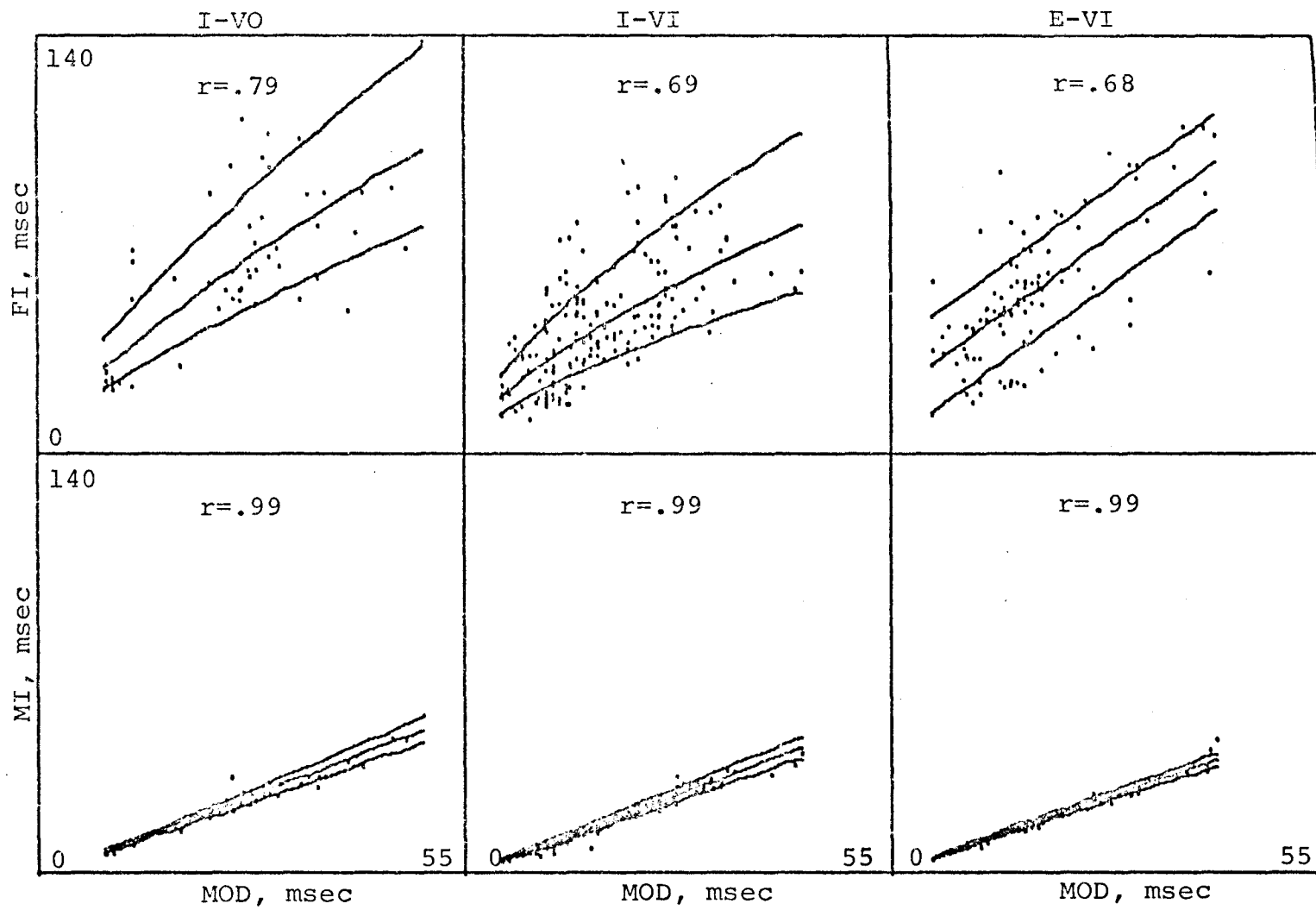


Figure 26. Regression of FI and MI on MOD for I-VO, I-VI and E-VI cells.

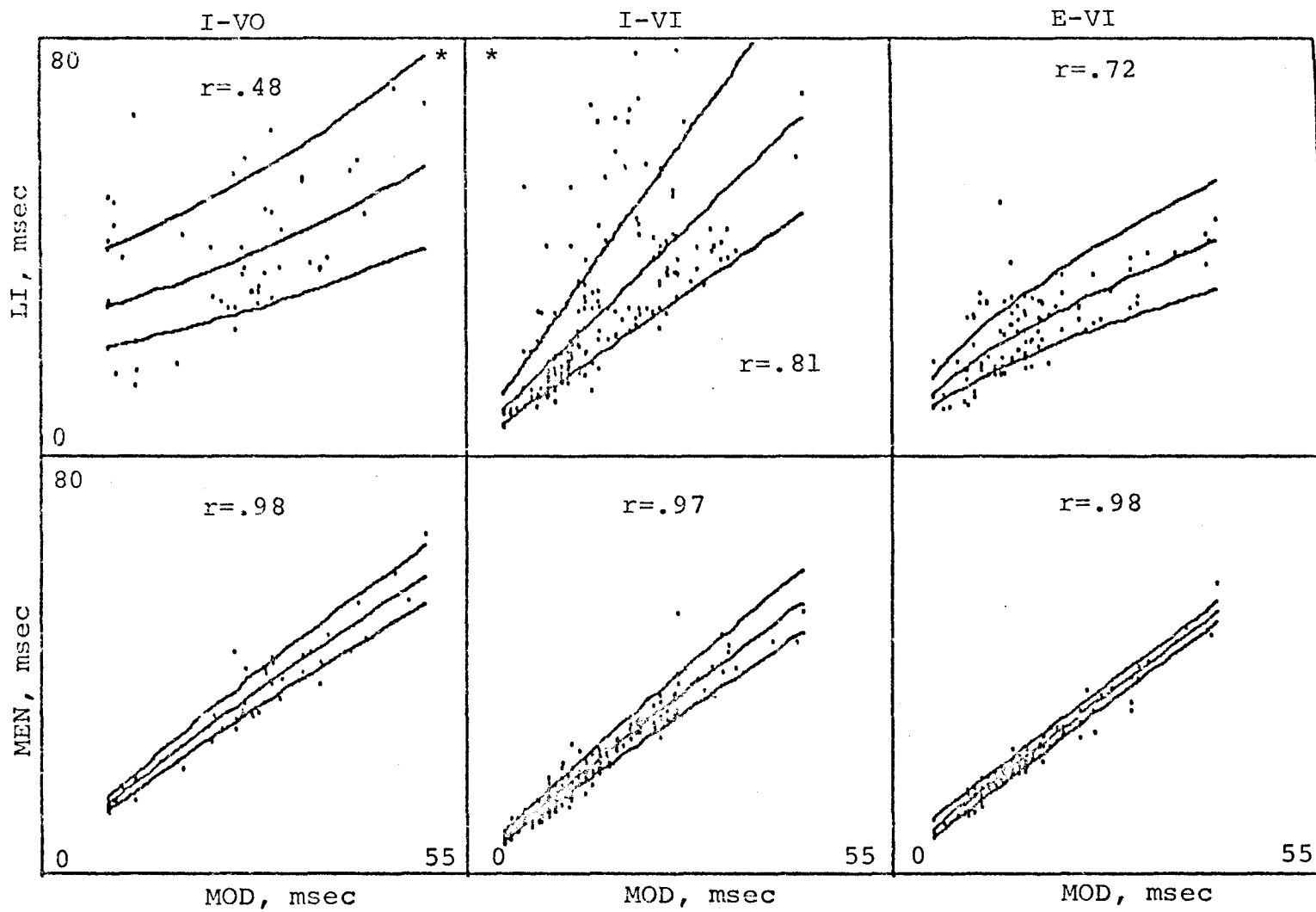


Figure 27. Regression of LI and MEN on MOD for I-VO, I-VI and E-VI cells.

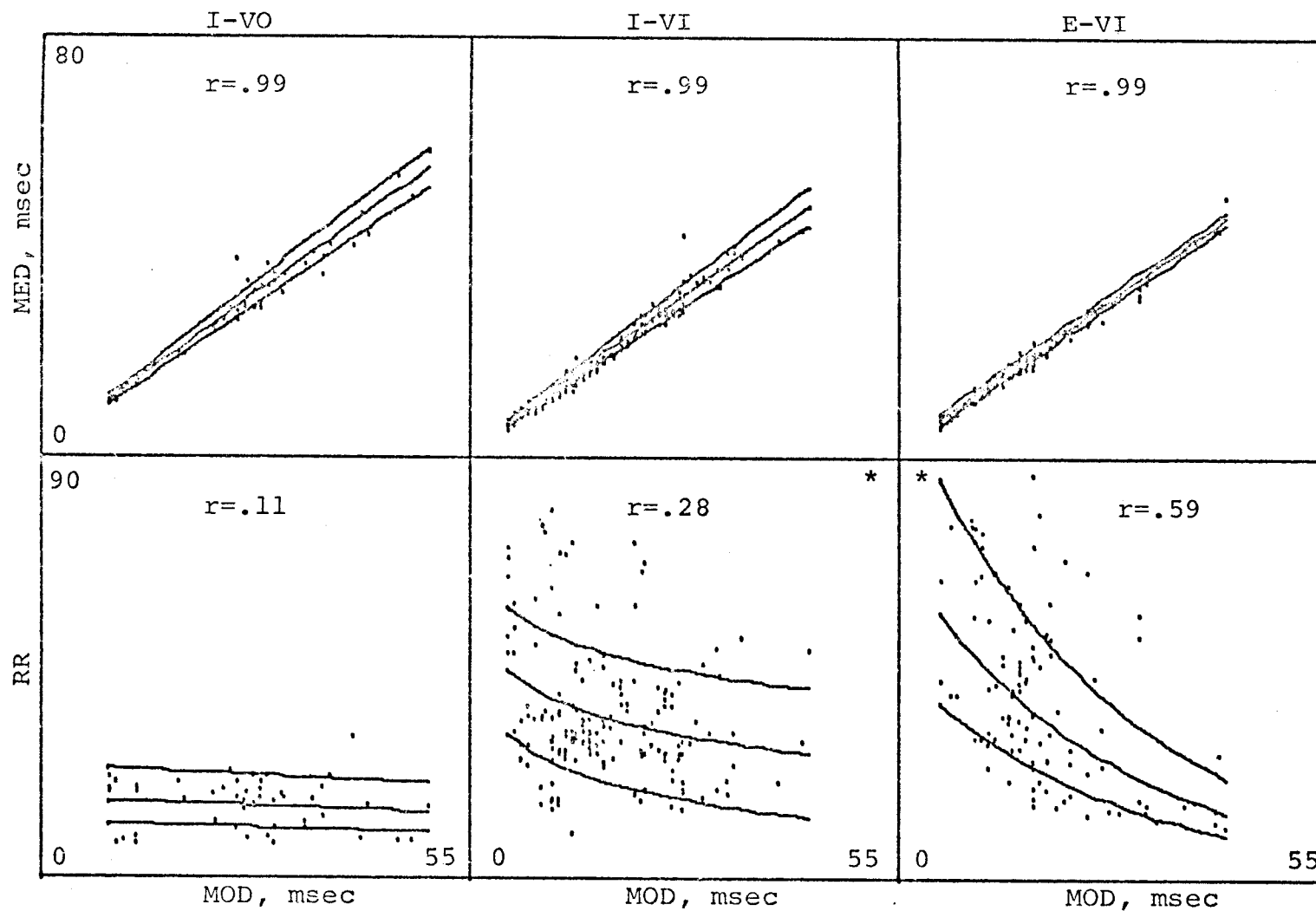


Figure 28. Regression of MED and RR on MOD for I-VO, I-VI and E-VI cells.

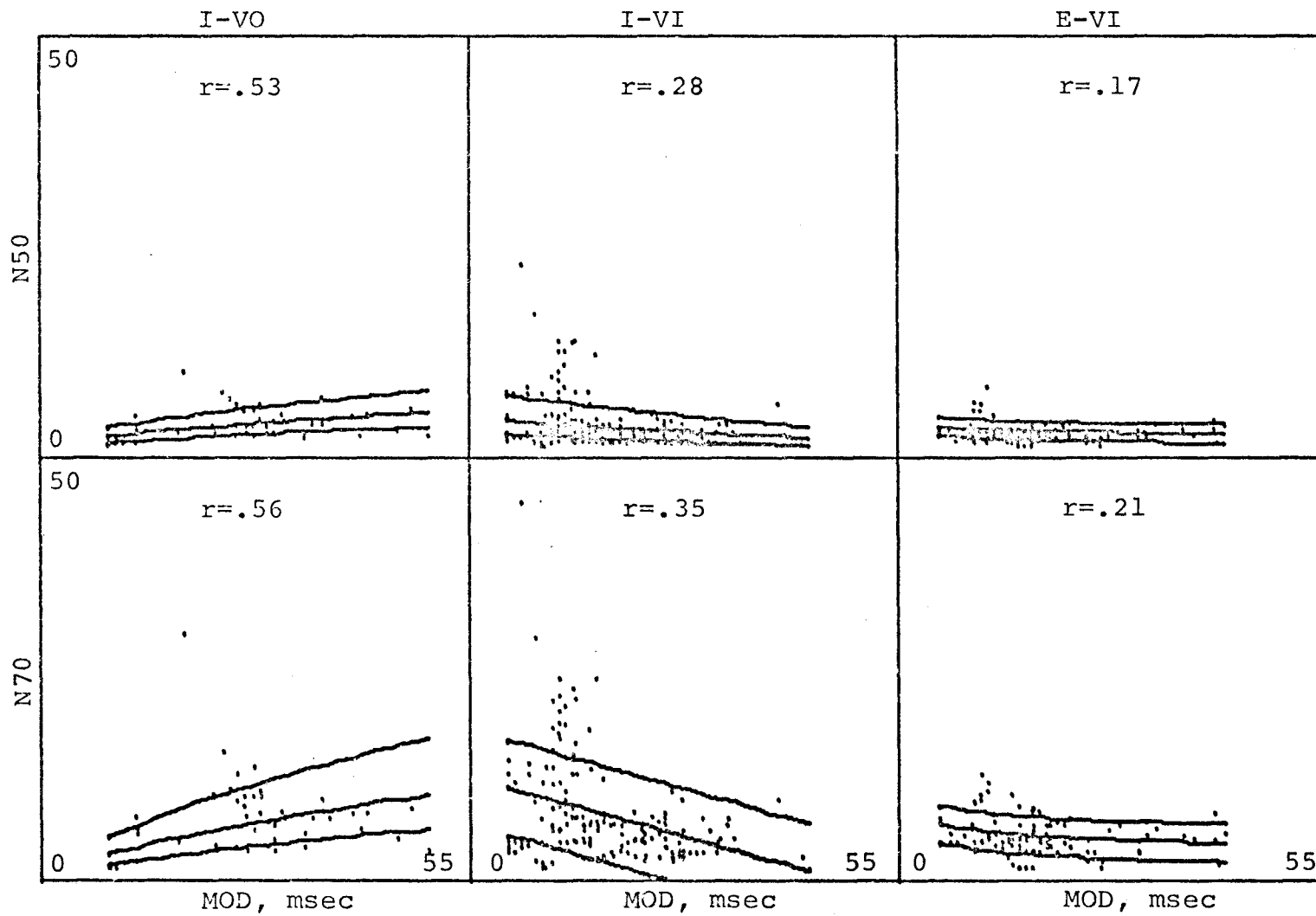


Figure 29. Regression of N50 and N70 on MOD for I-VO, I-VI and E-VI cells.

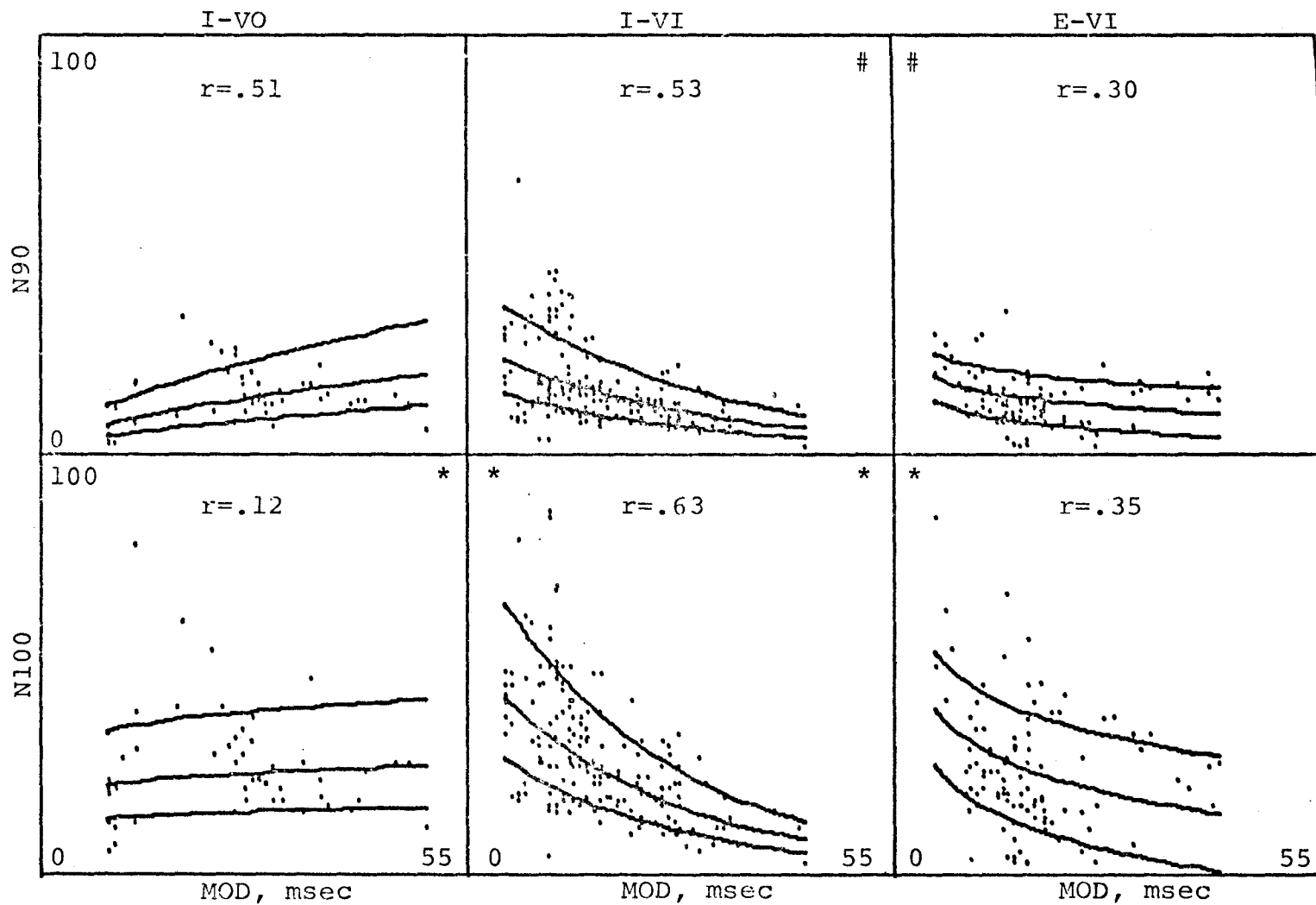


Figure 30. Regression of N90 and N100 on MOD for I-VO, I-VI and E-VI cells.

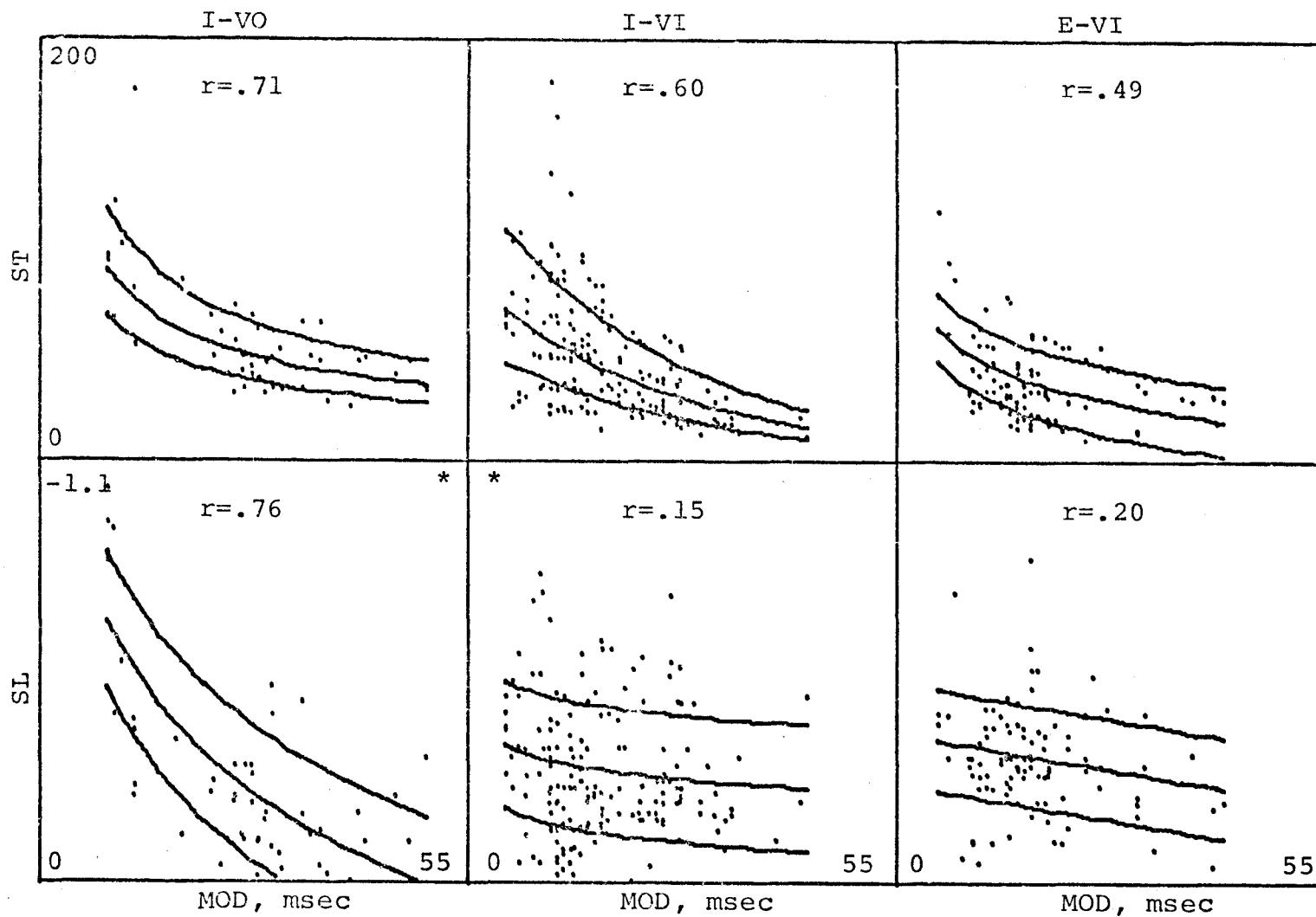


Figure 31. Regression of ST and SL on MOD for I-VO, I-VI and E-VI cells.

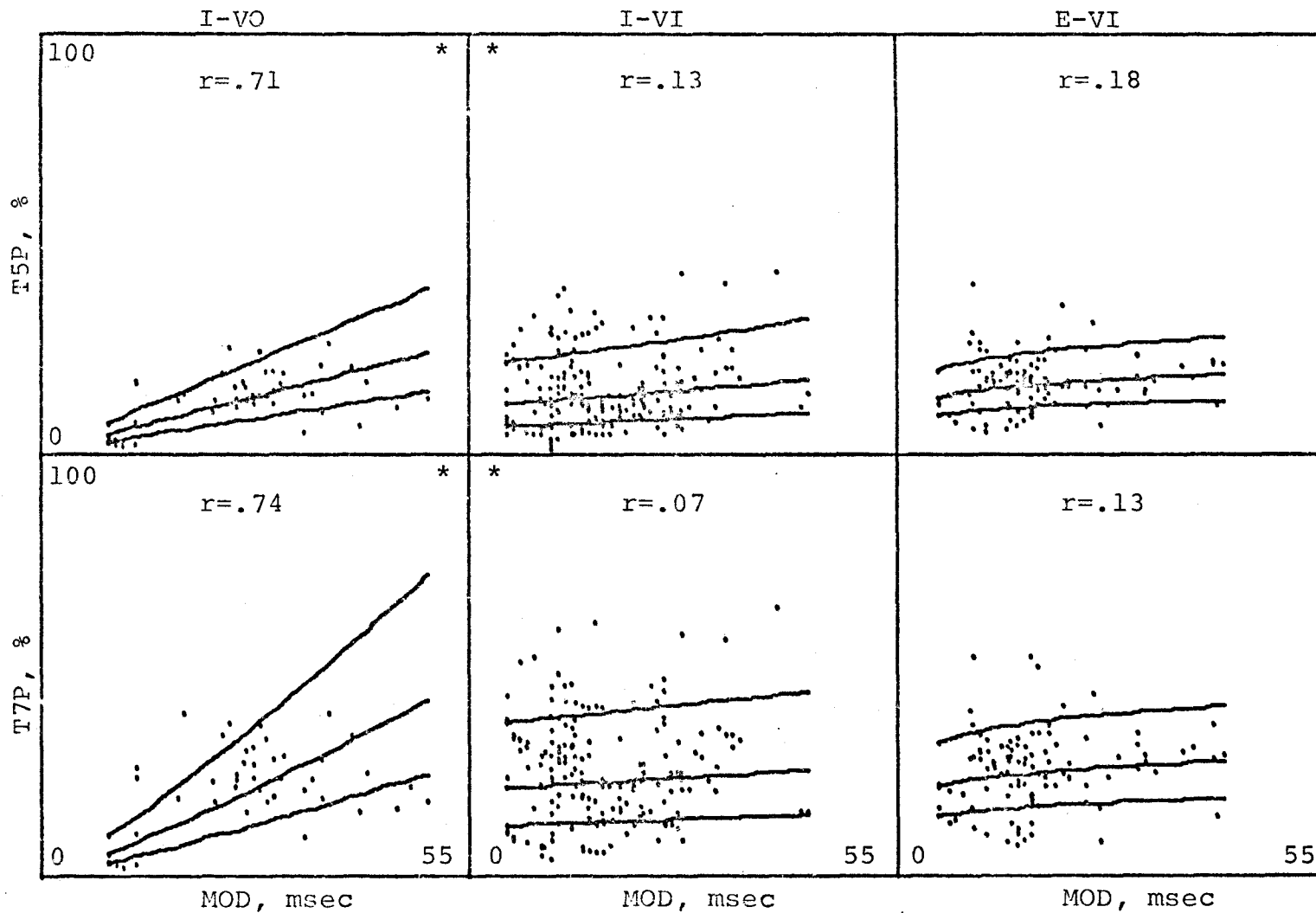


Figure 32. Regression of T5P and T7P on MOD for I-VO, I-VI and E-VI cells.



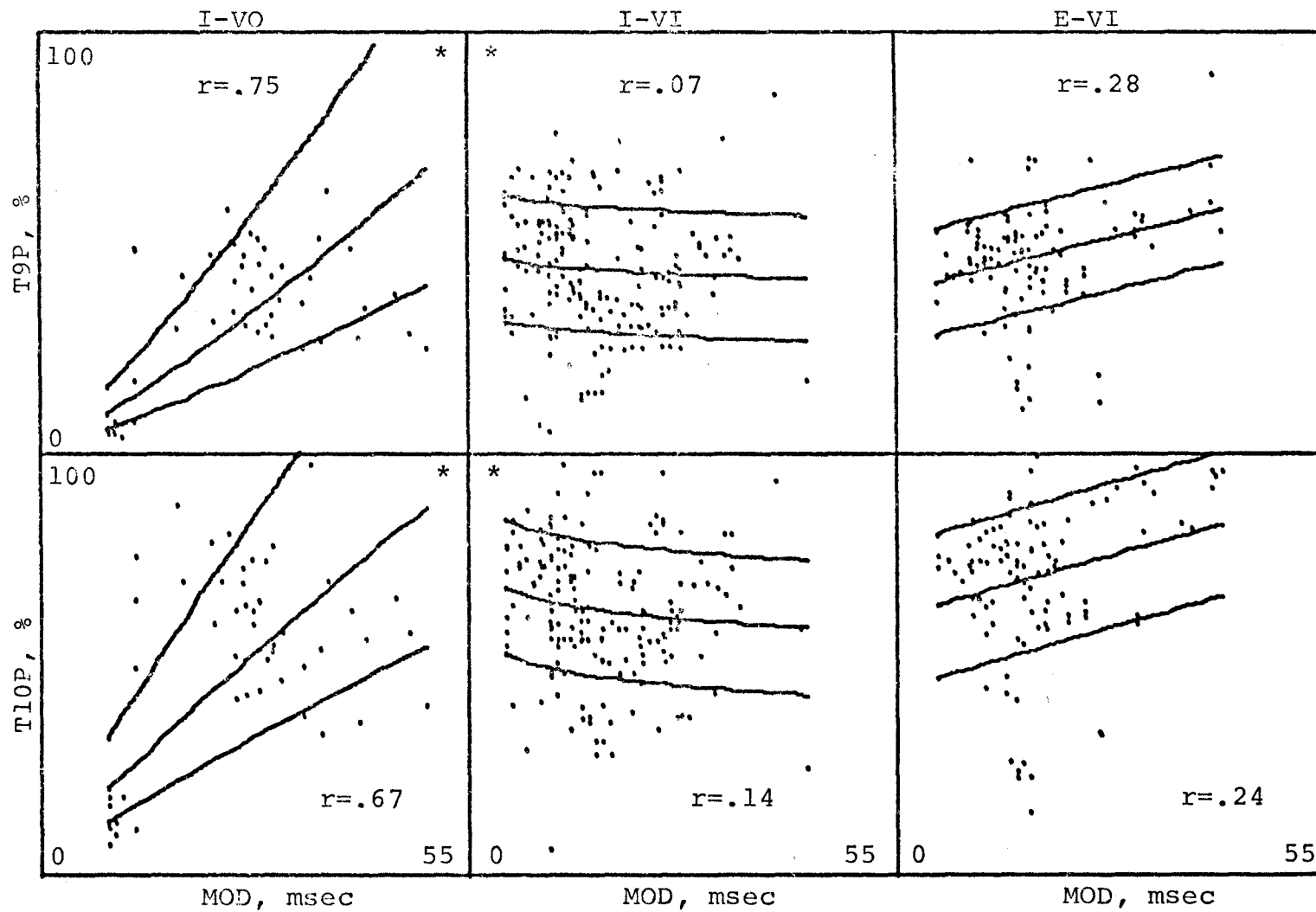


Figure 33. Regression of T9P and T10P on MOD for I-VO, I-VI and E-VI cells.

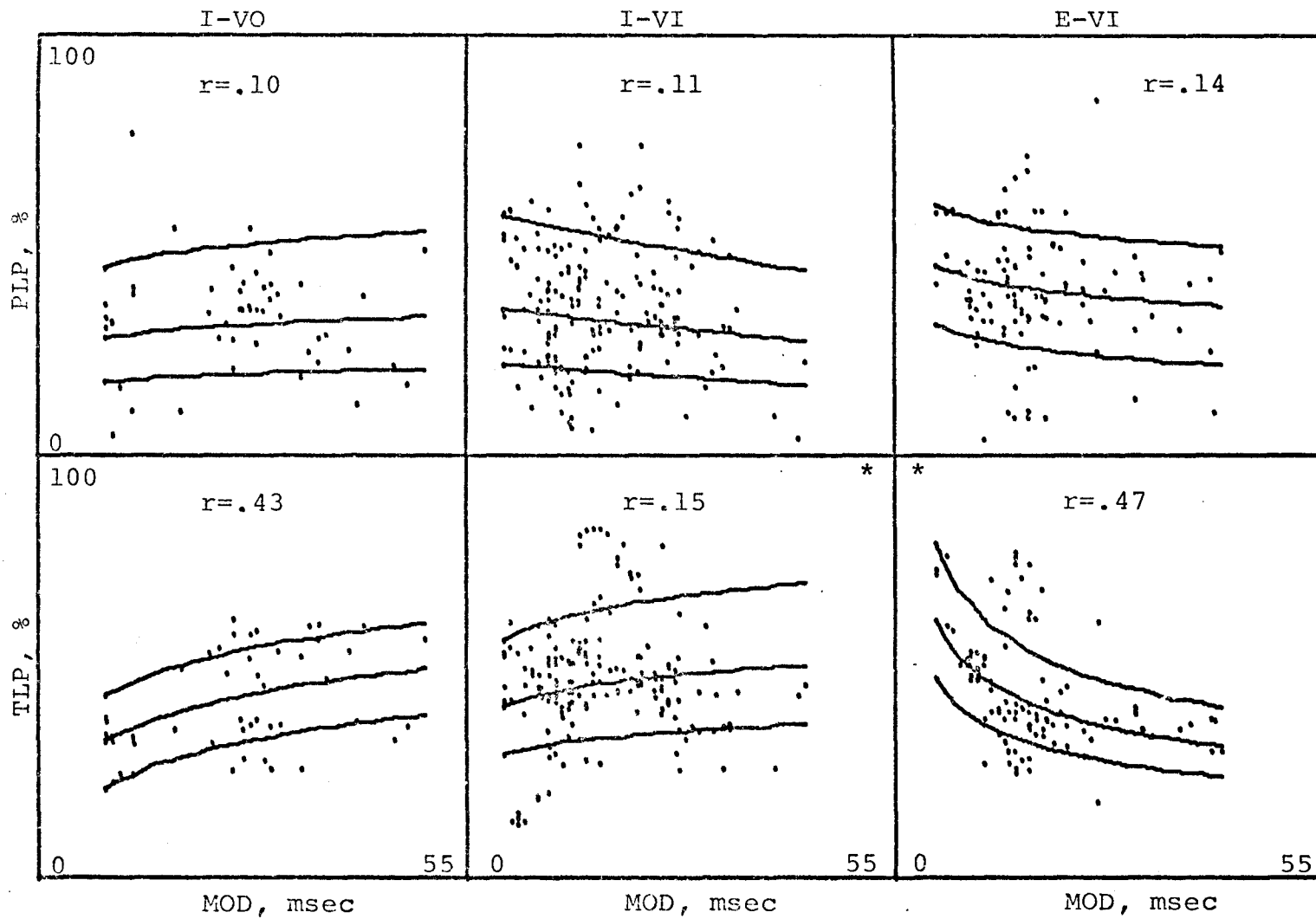


Figure 34. Regression of PLP and TLP on MOD for I-VO, I-VI and E-VI cells.

where applicable. Respiration rate (RR) is understood to be in breaths per minute.

The raw data points for each regression plot are placed in scattergram format. From Table III data, all I-VO plots contain 50 data points from 10 cells, all I-VI plots contain 175 data points from 51 cells and all E-VI plots contain 105 data points from 26 cells. In the regression analysis, each data point was considered as a new observation. No weighting factors were introduced to balance unequal data point contributions from various cells. The three lines drawn in each graph represent the mean regression line and error lines on either side of the mean for the best regression fit. The best fit was defined as that fit (linear, exponential, logarithmic or log-log) possessing the highest correlation coefficient for each set of data. The mean regression and error lines extend from low to high values of the independent variable which includes the entire rate or mode range observed for each case (I-VO, I-VI, E-VI) in this study. All plots are on a linear scale which renders curvilinear fits non-linear. Transformation of curvilinear regressions back to linear coordinates also distorts the symmetry of error lines on either side of the mean regression line. Still, the error lines include

68% of the raw data points or one standard deviation on either side of the mean regression line. The equations for all regression plots in Figures 11-34 are listed in the Appendix (Tables VIII-XIII). These equations can be used for accurate interpolation of dependent variables from independent variables. The type of fit (linear, exponential, logarithmic or log-log) used to correlate X and Y data for each regression plot are also listed in the Appendix (Table XIV).

Correlation coefficients or "r" values ranged from 0.04 to 0.99 and are given for each regression plot. Steep slopes and/or narrow error bands are associated with large "r" values and good fits. Flat slopes and/or wide error bands are associated with low "r" values and poor fits. The significant difference between "r" values of adjacent horizontal panels is designated by a number sign (#) for  $P < 0.05$  or an asterisk (\*) for  $P < 0.01$ . Blank panels indicate that the correlation coefficients between I-VO and I-VI or I-VI and E-VI plots are not significantly different at the 0.05 level.

In Figure 11, T50 is plotted as a function of RR. For I-VI and E-VI populations, T50 decreases as rates increase from 9.9 to 78.7 and 11.5 to 86.6 breaths per minute respectively. This indicates that at higher

breathing frequencies it takes less time on the average for inspiratory and expiratory trains to reach interspike intervals which are one half the difference between the first interval (FI) and minimum interval (MI). Comparison of I-VI and E-VI plots indicates that expiratory T50's are slightly longer in time than inspiratory T50's for equivalent respiratory rates. The expiratory regression has less scatter and a slightly greater slope as indicated by the significantly higher "r" value at the 0.05 level. For the I-VO population, respiration rate ranges from 7.9 to 30.9 breaths per minute. This decrease in rate response to drug stimulation or depression after vagotomy implicates the vagus in control of respiratory rate. For the T50 parameter, vagotomy significantly decreases the "r" value at the 0.05 level. That is, the correlation between T50 and RR is lost by cutting both vagus nerves (low slope, wide scatter). A similar analysis can be done by the reader for the remaining respiration rate regressions in Figures 11-22.

In Figure 23, T50 is plotted as a function of MOD. Mode ranges are approximately the same for I-VO, I-VI and E-VI populations and extend from 9.2 to 50.0, 5.2 to 43.6 and 5.2 to 42.0 milliseconds respectively. The plots show that T50 is directly related to the mode for

all populations. That is, as the mode decreases (increasing depth or tidal volume) it takes less time for respiratory trains to be activated. The inspiratory and expiratory populations are approximately the same and do not have significantly different "r" values at the 0.05 level. The relationship of T50 versus MOD is by far the best in the I-VO panel (highest "r" value) indicating that vagotomy significantly increases the correlation between T50 and MOD for inspiratory cells at the 0.01 level. Similar comparisons can be made by the reader for the remaining regressions in Figures 23-34.

Correlation coefficients ("r") are placed in rank order for the regression of I-VO, I-VI and E-VI cell discharge parameters on respiration rate and mode in Tables V and VI respectively. Correlations of inspiratory (I-VI) and expiratory (E-VI) parameters as a function of respiratory rate in Table V appear to parallel each other in rank order while vagotomy (I-VO) tends to invert the ordering for inspiratory (I-VI) cells. These same general relationships tend to hold for parameters plotted as a function of the mode in Table VI, but the I-VI - E-VI pairings and the I-VI - I-VO reciprocity are not as well defined. Comparison of respiration rate (Table V) and mode (Table VI) rankings for similar cell types reveals poor order

Table V. Rank ordering of correlation coefficients for the regression of 24 parameters on respiratory rate for I-VO, I-VI and E-VI cell types.

Rank Order	I-VO		I-VI		E-VI	
	Parameter	"r"	Parameter	"r"	Parameter	"r"
1	TL	.62	TL	.69	TL	.84
2	TLP	.59	T100	.68	T100	.79
3	T7P	.43	T90	.57	FI	.73
4	T5P	.39	T70	.55	T90	.72
5	T9P	.38	PL	.53	PL	.71
6	LI	.34	T50	.44	T70	.65
7	ST	.31	FI	.44	T50	.64
8	PL	.31	ST	.33	MEN	.62
9	SL	.30	MEN	.31	MED	.60
10	FI	.30	MED	.30	MOD	.59
11	N50	.27	MOD	.28	MI	.59
12	T10P	.26	MI	.27	LI	.44
13	PLP	.25	N100	.24	TLP	.43
14	N70	.24	T9P	.21	T7P	.38
15	MEN	.21	TLP	.21	N100	.38
16	N90	.18	LI	.20	T5P	.36
17	MED	.16	N70	.18	ST	.35
18	MI	.15	T5P	.16	PLP	.33
19	T70	.13	PLP	.15	T10P	.29
20	T100	.13	T10P	.12	N90	.24
21	MOD	.11	N90	.12	T9P	.17
22	N100	.10	T7P	.10	N70	.10
23	T50	.08	N50	.08	SL	.09
24	T90	.06	SL	.08	N50	.04

Table VI. Rank ordering of correlation coefficients for the regression of 24 parameters on mode for I-VO, I-VI and E-VI cell types.

Rank Order	I-VO		I-VI		E-VI	
	Parameter	"r"	Parameter	"r"	Parameter	"r"
1	MED	.99	MED	.99	MED	.99
2	MI	.99	MI	.99	MI	.99
3	MEN	.98	MEN	.97	MEN	.98
4	T50	.85	LI	.81	LI	.72
5	T70	.84	FI	.69	FI	.68
6	T90	.83	N100	.63	RR	.59
7	FI	.79	ST	.60	T90	.59
8	T100	.77	N90	.53	T50	.59
9	SL	.76	T50	.40	T70	.54
10	T9P	.75	N70	.35	T100	.51
11	T7P	.74	TL	.33	ST	.49
12	T5P	.71	T70	.33	TLP	.47
13	ST	.71	T90	.30	TL	.46
14	T10P	.67	RR	.28	N100	.35
15	N70	.56	N50	.28	N90	.30
16	N50	.53	T100	.26	T9P	.28
17	TL	.52	PL	.17	PL	.25
18	N90	.51	SL	.15	T10P	.24
19	LI	.48	TLP	.15	N70	.21
20	TLP	.43	T10P	.14	SL	.20
21	PL	.39	T5P	.13	T5P	.18
22	N100	.12	PLP	.11	N50	.17
23	RR	.11	T9P	.07	PLP	.14
24	PLP	.10	T7P	.07	T7P	.13



correlations which may even approach randomization. The reader can use Tables V and VI as references relating to those parameters which are correlated best with either respiration rate (RR) or mode (MOD) for each cell type studied (I-VO, I-VI, E-VI).

The numerous regression plots presented in Figures 11-34 are condensed in Figures 35-43 for reader convenience. In these latter plots, cells are still separated into I-VO, I-VI and E-VI types, but different parameters are superimposed in individual plots. Only the mean regression lines are plotted to avoid confusion from overlapping error lines and raw data points. In each figure, the same parameters are plotted both as a function of respiration rate (RR) and mode (MOD) for easy comparison. The respiration rate (RR) and mode (MOD) scalings are identical to those used in Figures 11-34 and the data ranges are the same. The dependent variable scalings are equivalent for both rate and mode plots for any one figure and correspond to values selected in Figures 11-34 with a few appropriate exceptions. Each axis is labeled with time or percent scales where applicable. Respiration rate (RR) is given in breaths per minute.

In Figure 35, T50, T70, T90 and T100 are plotted

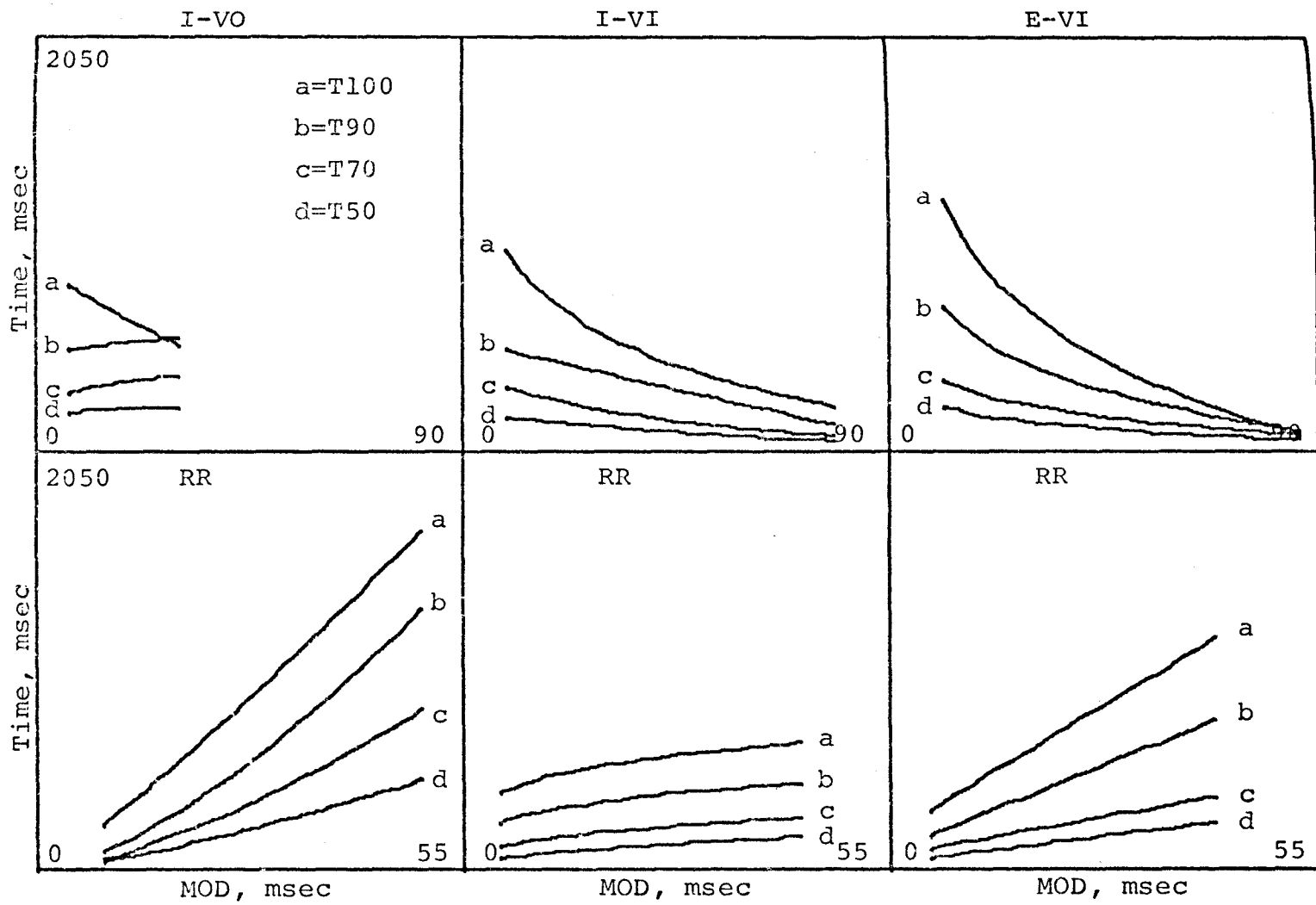


Figure 35. Regression of T100, T90, T70 and T50 on RR and MOD for I-VO, I-VI and E-VI cells.

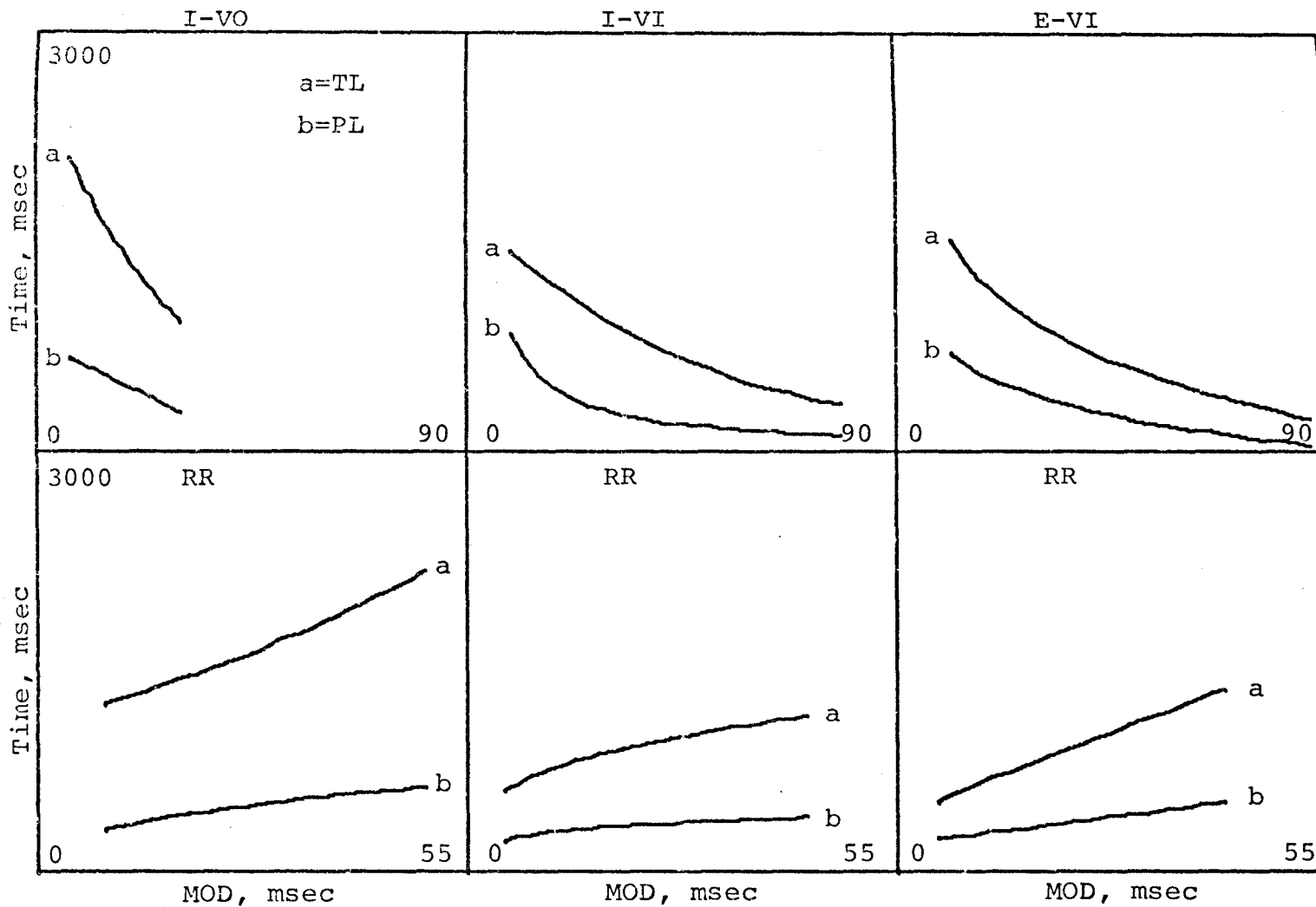


Figure 36. Regression of TL and PL on RR and MOD for I-VO, I-VI and E-VI cells.

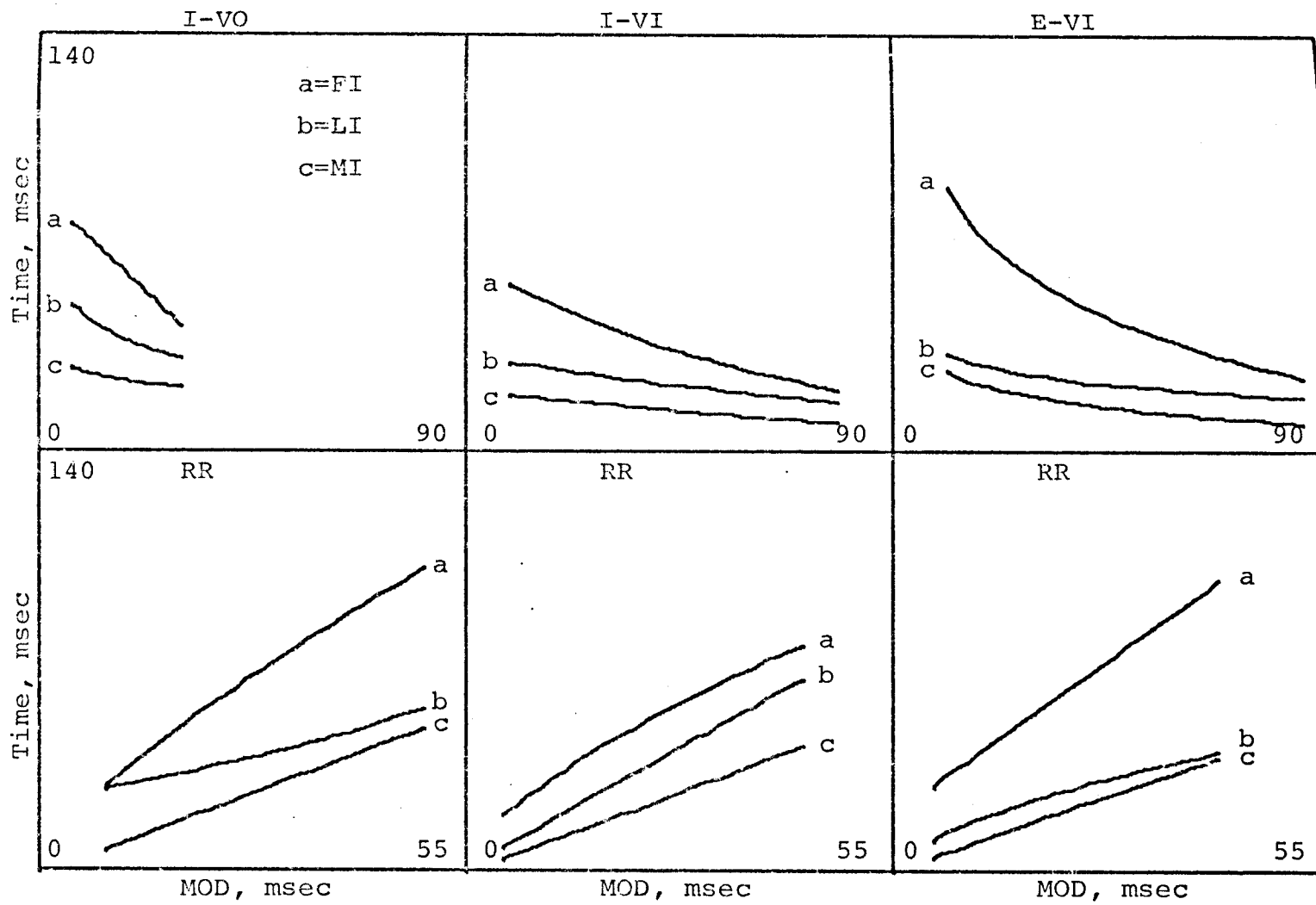


Figure 37. Regression of FI, LI and MI on RR and MOD for I-VO, I-VI and E-VI cells.

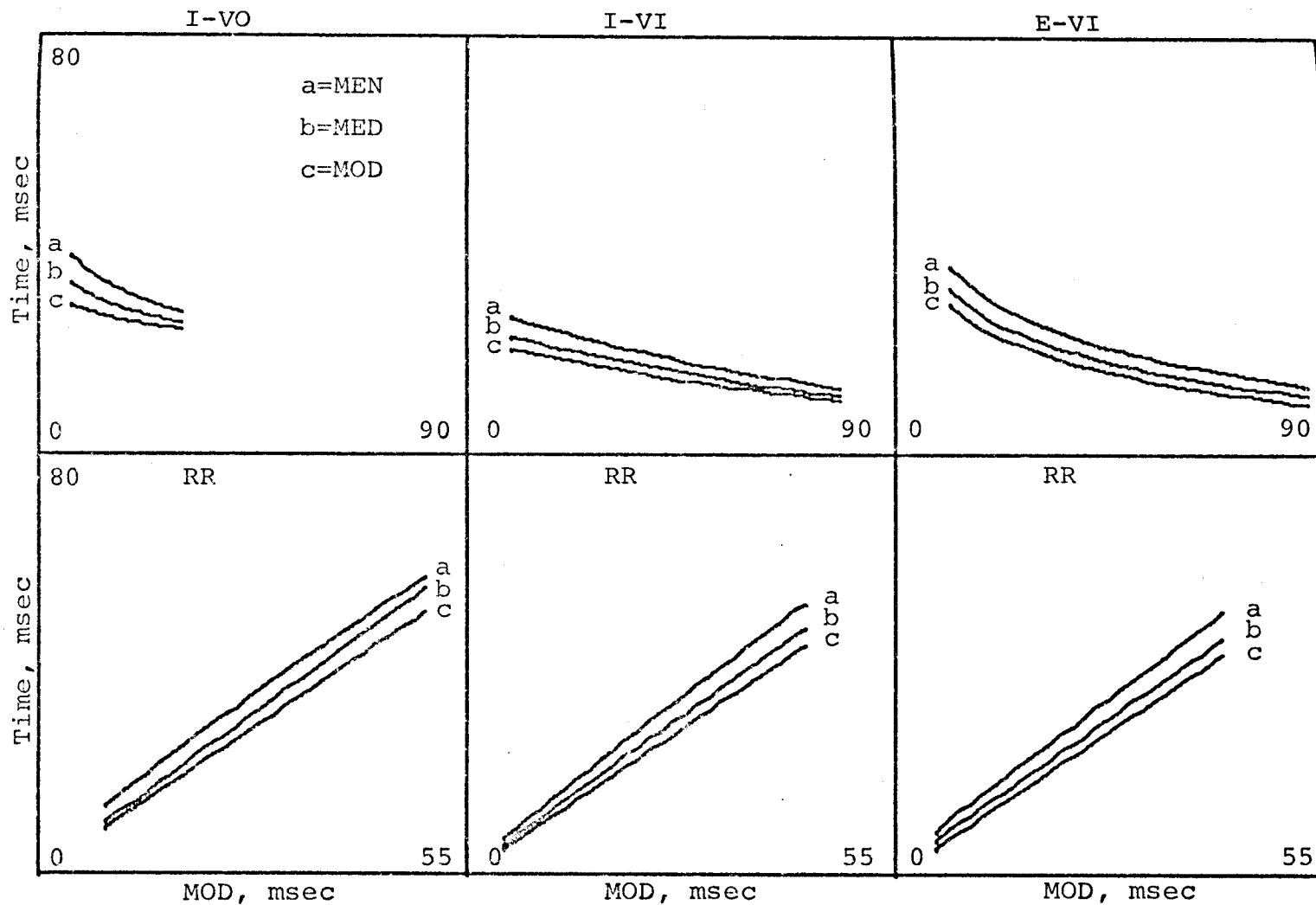


Figure 38. Regression of MEN, MED and MOD on RR and MOD for I-VO, I-VI and E-VI cells.

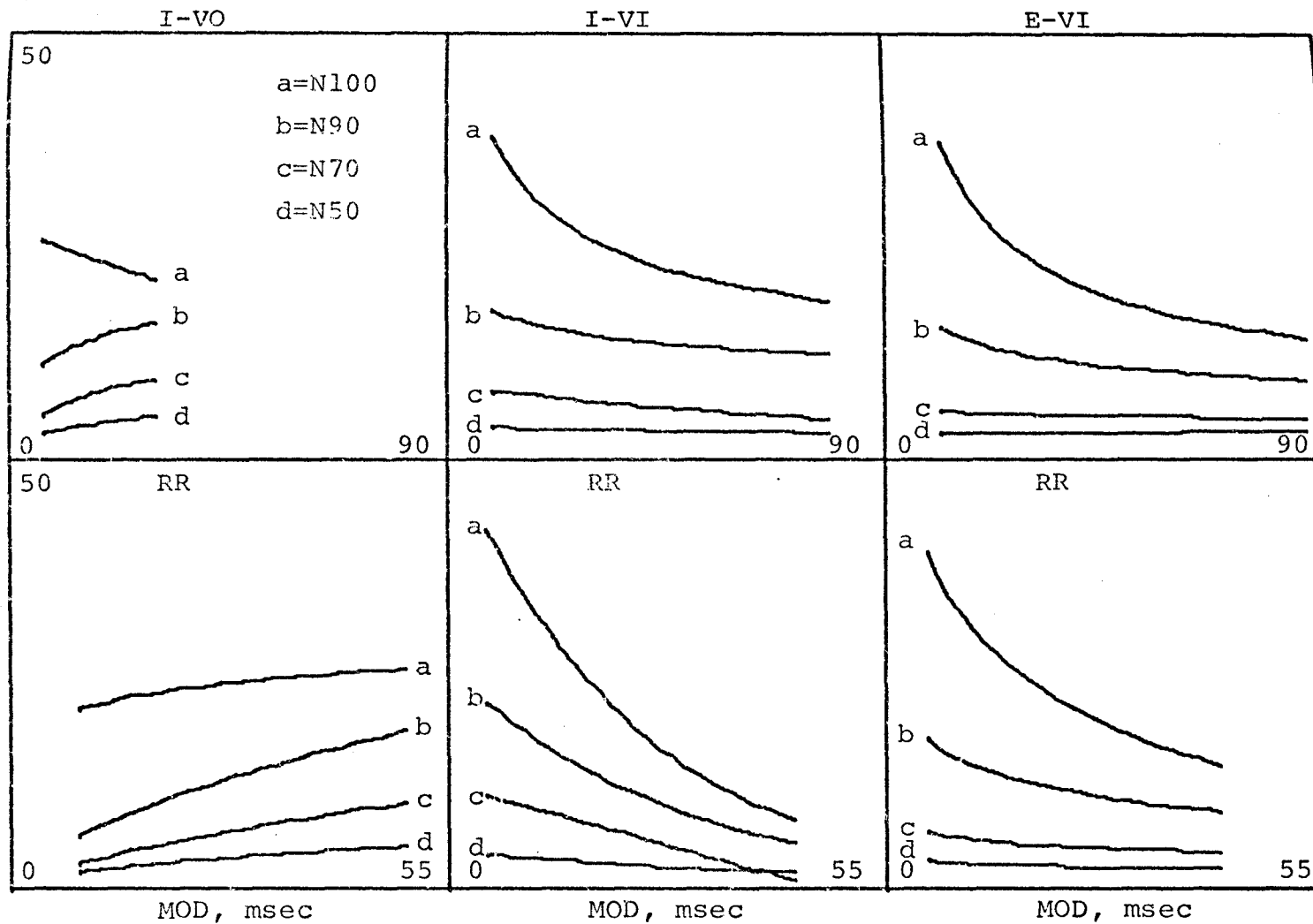


Figure 39. Regression of N100, N90, N70 and N50 on RR and MOD for I-VO, I-VI and E-VI cells.

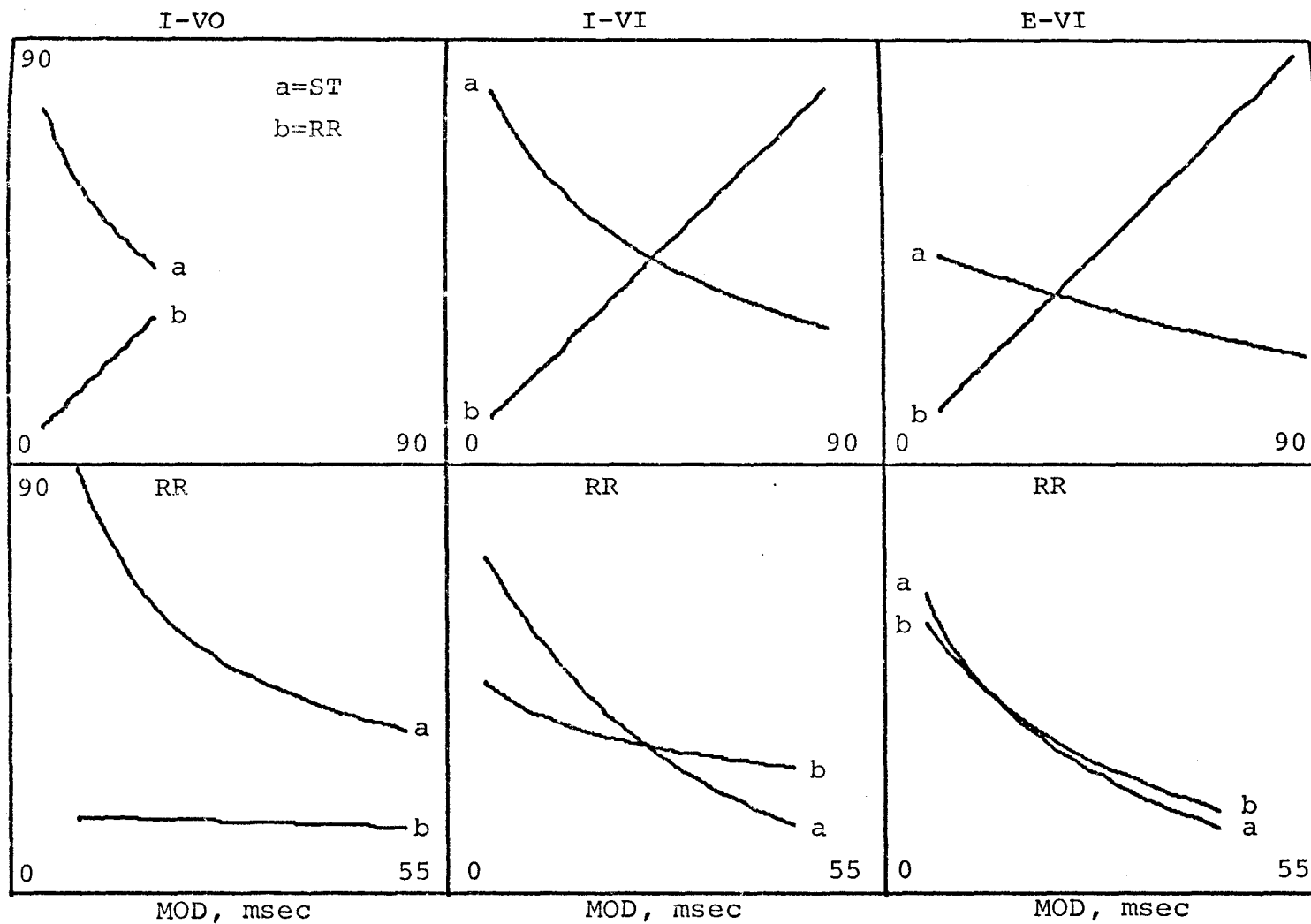


Figure 40. Regression of ST and RR on RR and MOD for I-VO, I-VI and E-VI cells.

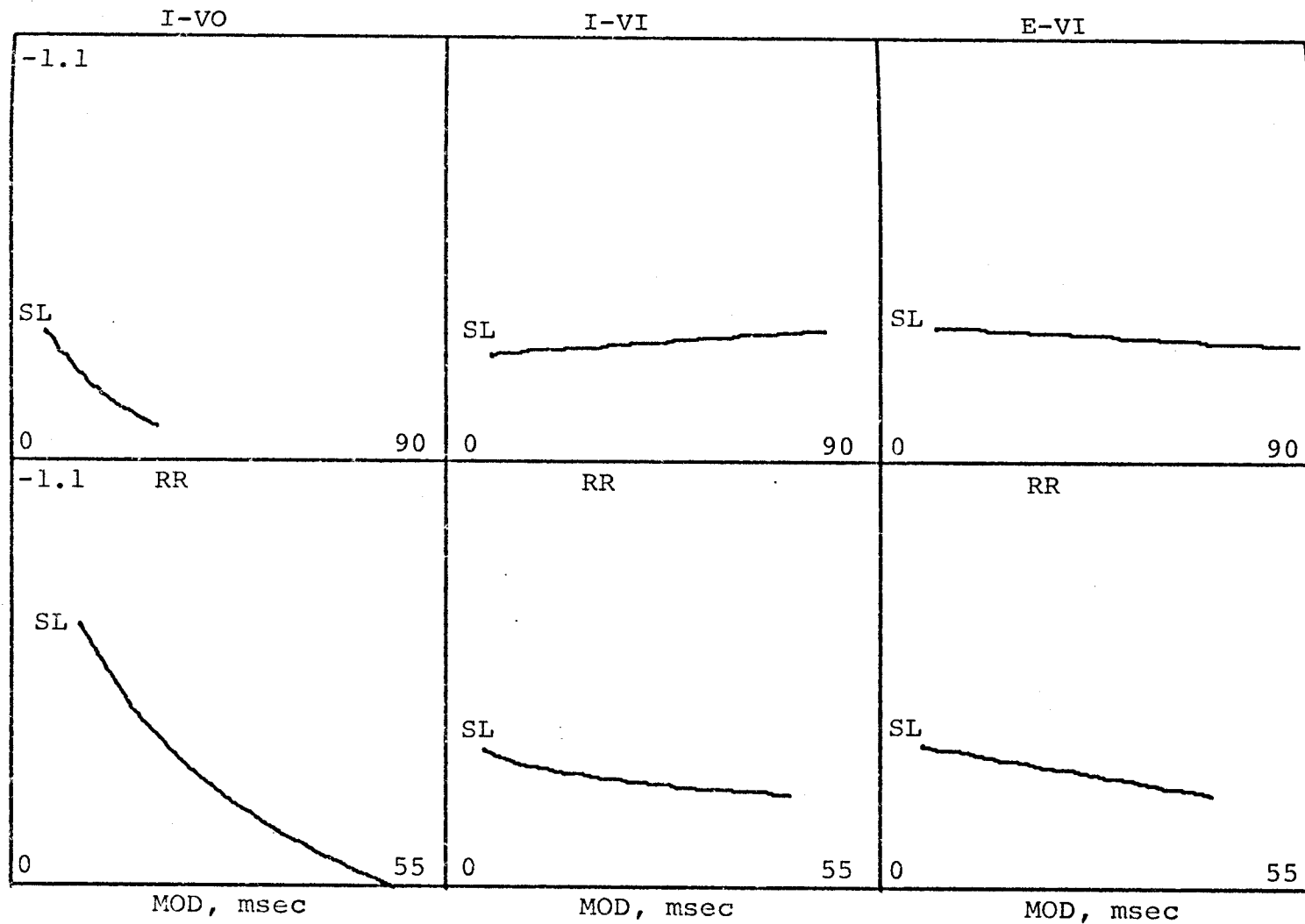


Figure 41. Regression of SL on RR and MOD for I-VO, I-VI and E-VI cells.



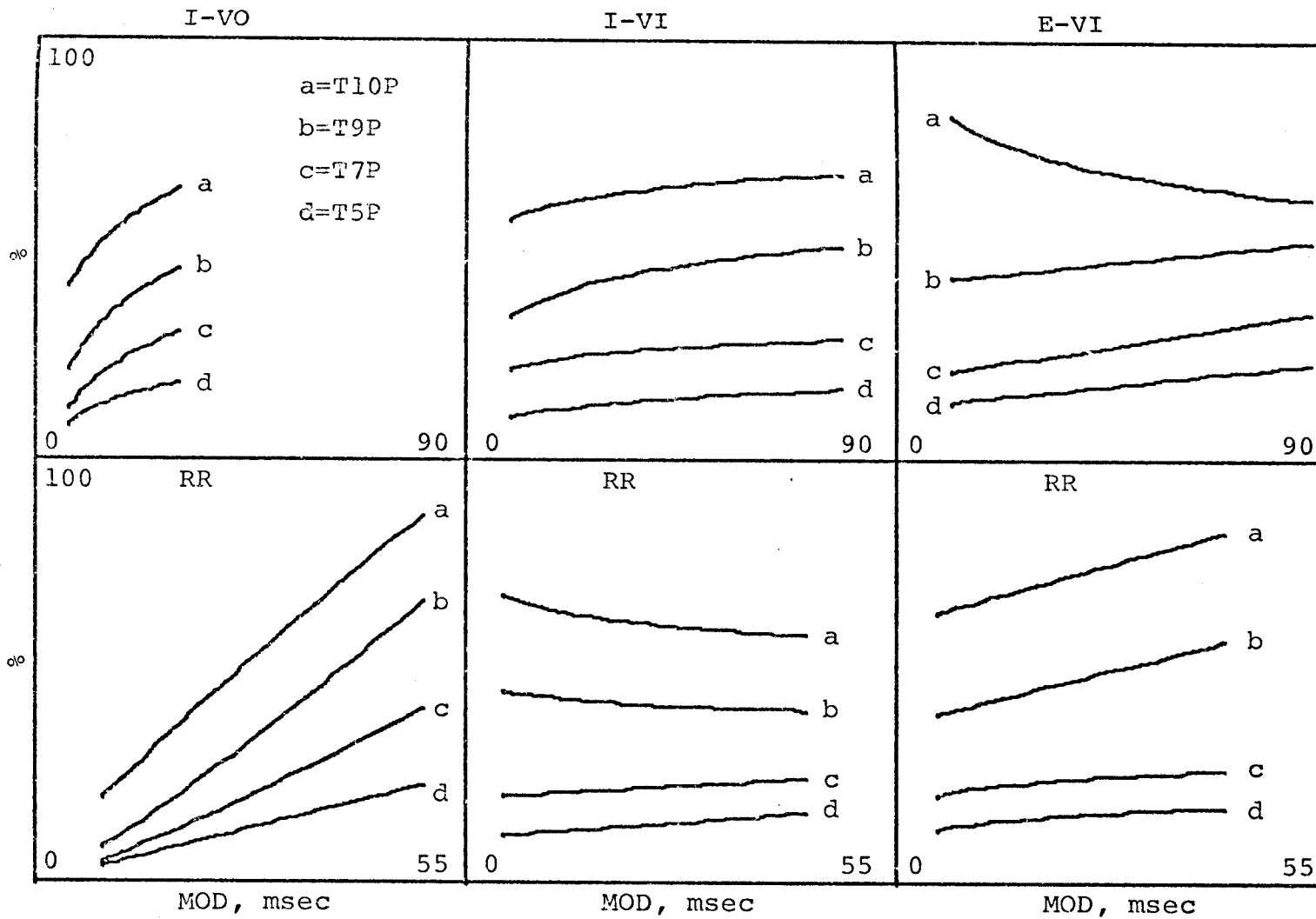


Figure 42. Regression of T10P, T9P, T7P and T5P on RR and MOD for I-VO, I-VI and E-VI cells.

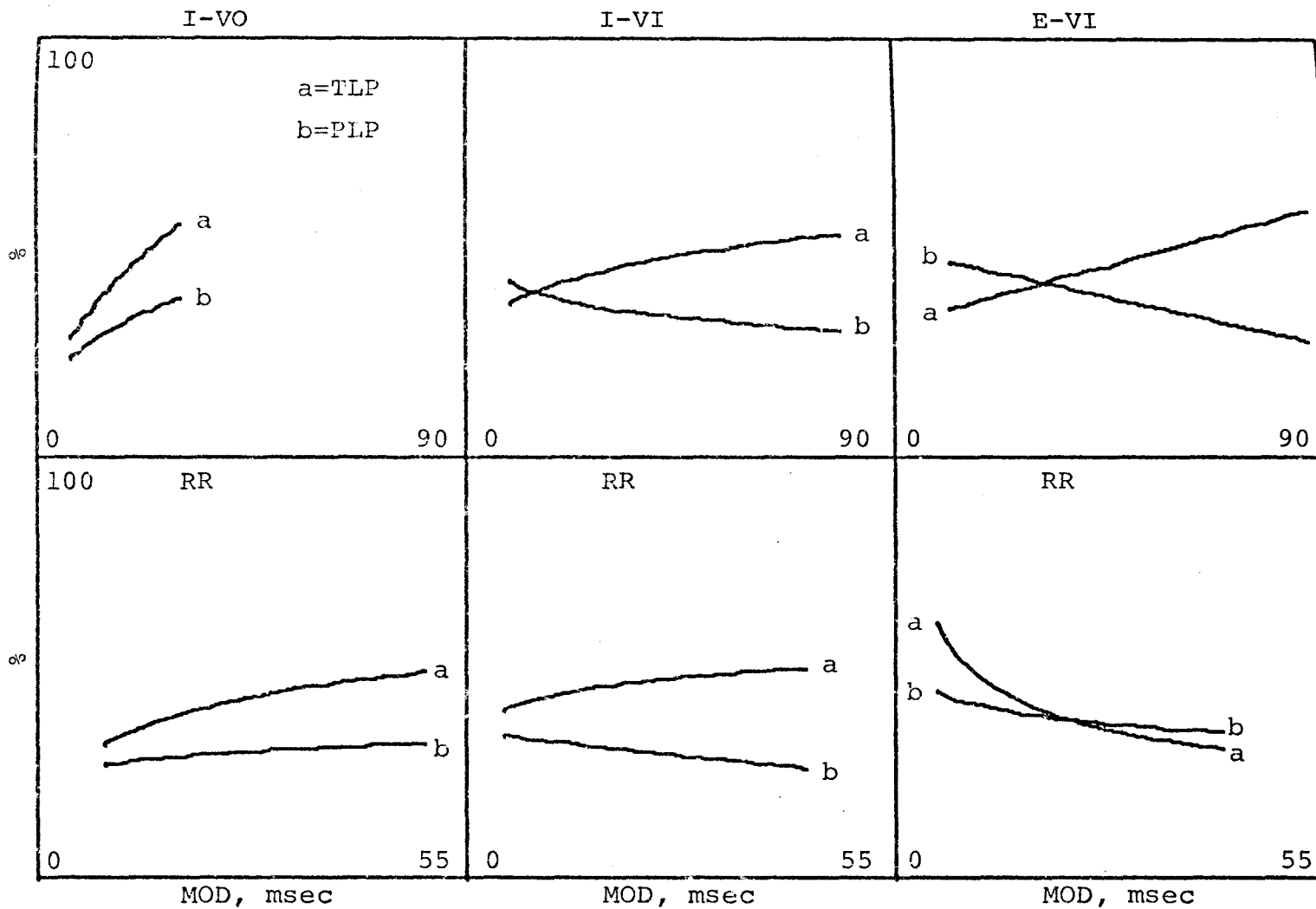


Figure 43. Regression of TLP and PLP on RR and MOD for I-VO, I-VI and E-VI cells.

as a function of both respiration rate (RR) and mode (MOD) for I-VO, I-VI and E-VI cells. In accordance with the definition of each parameter,  $T50 < T70 < T90 < T100$  for each of the six plots. Seven comparisons can be made between the six panels (e.g., I-VI to I-VO and I-VI to E-VI for RR and MOD plots and RR to MOD for I-VO, I-VI and E-VI plots). For example, the following observations can be made by comparing adjacent panels in the horizontal plane. Times to reach fractions of the way to the minimum interval (T50, T70, T90, T100) decrease as the respiration rate (RR) increases for both inspiratory and expiratory cells. Vagotomy abolishes this relationship for inspiratory cells. For inspiratory cells, T parameters are almost independent of changes in the mode (MOD) when the vagi are intact (I-VI), but become positively correlated when the vagal influence is removed (I-VO). For expiratory cells, T parameters follow changes in the mode (MOD).

Comparison of adjacent panels in the vertical plane reveals that T50, T70, T90 and T100 parameters for expiratory cells (E-VI) correlate with either respiration rate (RR) or mode (MOD), but parameter correlations for inspiratory cells (I-VI, I-VO) depend on the state of vagal innervation. That is, when the vagi are intact (I-VI),

T parameters correlate best with the respiration rate (RR). When the vagi are cut, the T parameters correlate best with the mode (MOD). This reversal is of importance to consider when comparing rate and depth outputs of the respiratory system. The reader can make similar comparisons between other cell and parameter types from the rest of the superimposed regression plots in Figures 36-43. Major findings from the population data will be elaborated in the Discussion.

#### C. Single Cell Data

From the 83 cells (Table III) contributing to the I-VO, I-VI and E-VI population data (Figures 11-43), 14 cells have been selected to illustrate single unit responses to induced or spontaneous changes in respiratory rate and/or depth. The reader should make careful comparisons between the single cell data presented and the population plots of Figures 35-43.

One of the important questions to consider in this study is how well do mode (MOD) interval times of single cells correlate with the depth of respiration. The theoretical reasoning diagrammed in Figure 10 predicts that for any one inspiratory cell, the mode (MOD) should be reciprocally related to the depth of respiration and independent of changes in spontaneous respiratory rate.

Examples from six inspiratory cells (I-VI) are presented to qualitatively establish the mode-depth relationship, point out quantitative cell-to-cell differences and suggest that mode interval times can fluctuate with the spontaneous respiratory rate in some circumstances. For each example, the depth of respiration is defined in relative terms as the average end inspiratory to end expiratory intrapleural pressure difference ( $\Delta$ IPP).

Three histograms and interspike interval modulation curves for inspiratory cell C32UI1R are superimposed in Figure 44A and B respectively. The curves are numbered consecutively, denoting a chronological data point acquisition sequence. Corresponding absolute and percent changes in the respiratory rate (RR), mode (MOD) and respiratory depth ( $\Delta$ IPP) measurements are listed for each data point in tabular format. In this series, there was a substantial spontaneous increase in the depth of respiration (73.1%) with only a small rate decrease (11.9%). The single unit response was one of activation. That is, the histogram interval distribution was shifted to the left (Figure 44A) resulting in a decreased mode time (34.5%) and the interspike interval modulation curve was shifted downward in a parallel fashion (Figure 44B). Rate and mode interaction, if any, cannot be ascertained

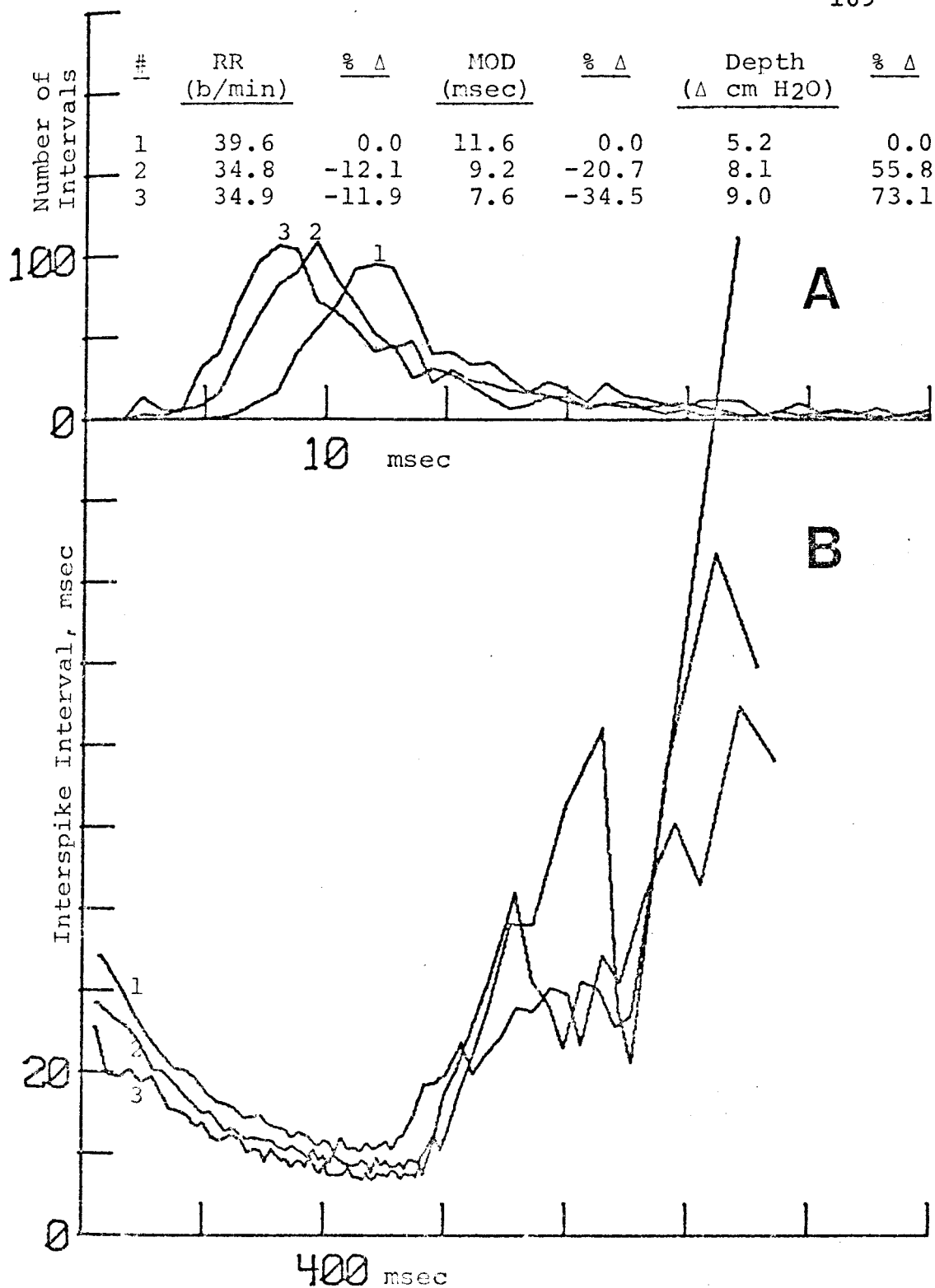


Figure 44. C32UIIR. Effect of spontaneous increase in respiratory depth on histogram (A) and interspike interval modulation curve (B). Data points numbered in time sequence. Vagi intact.

from these data. If one assumes that this cell possesses characteristics similar to the population data of Figure 16 where mode and rate are reciprocally related, the rate slowing observed in Figure 44 may limit the mode decrease during an increase in respiratory depth. From the data presented in Figure 44 it is reasonable to conclude that mode time and respiratory depth have a reciprocal relationship.

Figure 45 shows single cell data for cell C80UI1R obtained during a spontaneous decrease in respiratory depth ( $\Delta$ IPP). In this case, the depth decreases proportionately more than the rate increases (28.0% versus 13.8% respectively) suggesting that the mode increase (54.5%) is associated with the former. The cell displays a typical deactivation pattern in which the histogram interval distribution shifts to the right and the interspike interval modulation curve is shifted upward. In another cell, not pictured, C29UI7R, the increase in the mode from 10.8 to 13.2 milliseconds (22.2%) was associated with a depth change from 4.4 to 3.7 centimeters of water (-15.9%). The rate change was minimal from 30.7 to 29.7 breaths per minute (-3.3%). These single cell examples suggest that mode time and respiratory depth are reciprocally related.

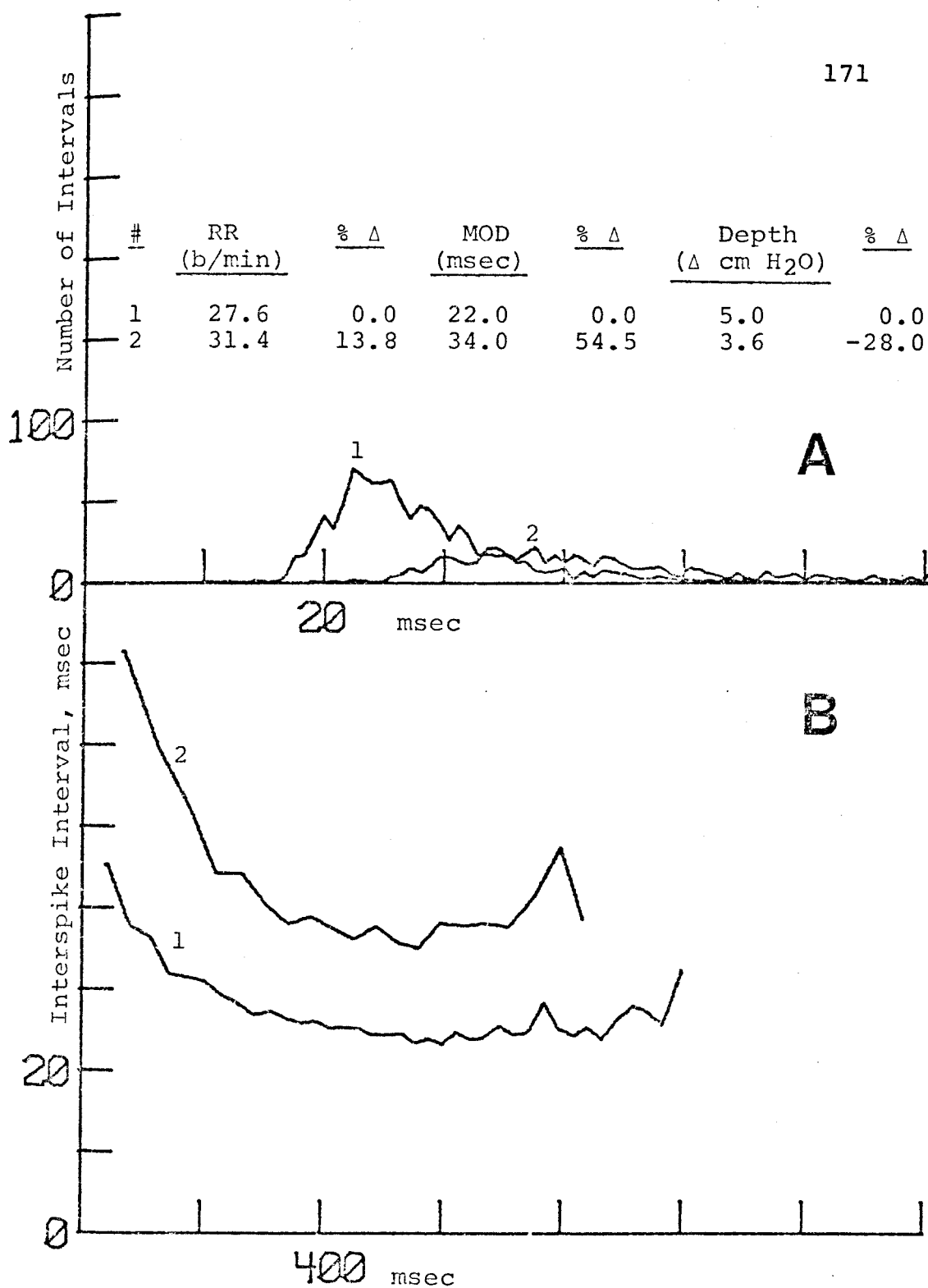


Figure 45. C80UI1R. Effect of spontaneous decrease in respiratory depth on histogram (A) and interspike interval modulation curve (B). Vagi intact.



Careful inspection of tabular data in Figures 44 and 45 shows that an absolute depth ( $\Delta$ IPP) of about 5.2 to 5.0 centimeters of water is paralleled by mode times of 11.6 and 22.0 milliseconds respectively. These differences in single cell firing frequencies at equivalent depths can be explained in two ways. First, the two cats may have had different (and unknown) compliances. Mode time may be directly related to compliance for any given volume or depth. Second, the difference may be attributed to characteristic variations among central inspiratory neurons. Both phenomena are probably operative in the respiratory control system, but the following experiment was addressed to the latter point.

In one cat breathing spontaneously, two inspiratory cells were recorded simultaneously with two separate microelectrodes inserted on opposite sides of the medulla. As shown in Figure 46A, the left cell (I2L) had a higher discharge frequency than the right cell (I3R). The histogram plots and frequency modulation curves of Figure 46B and C, respectively, show firing pattern differences between the fast (I2L) and slow (I3R) cells. Since both cells were recorded at the same time in the same animal, differences in mode time cannot be attributed to variations in mechanical or chemical

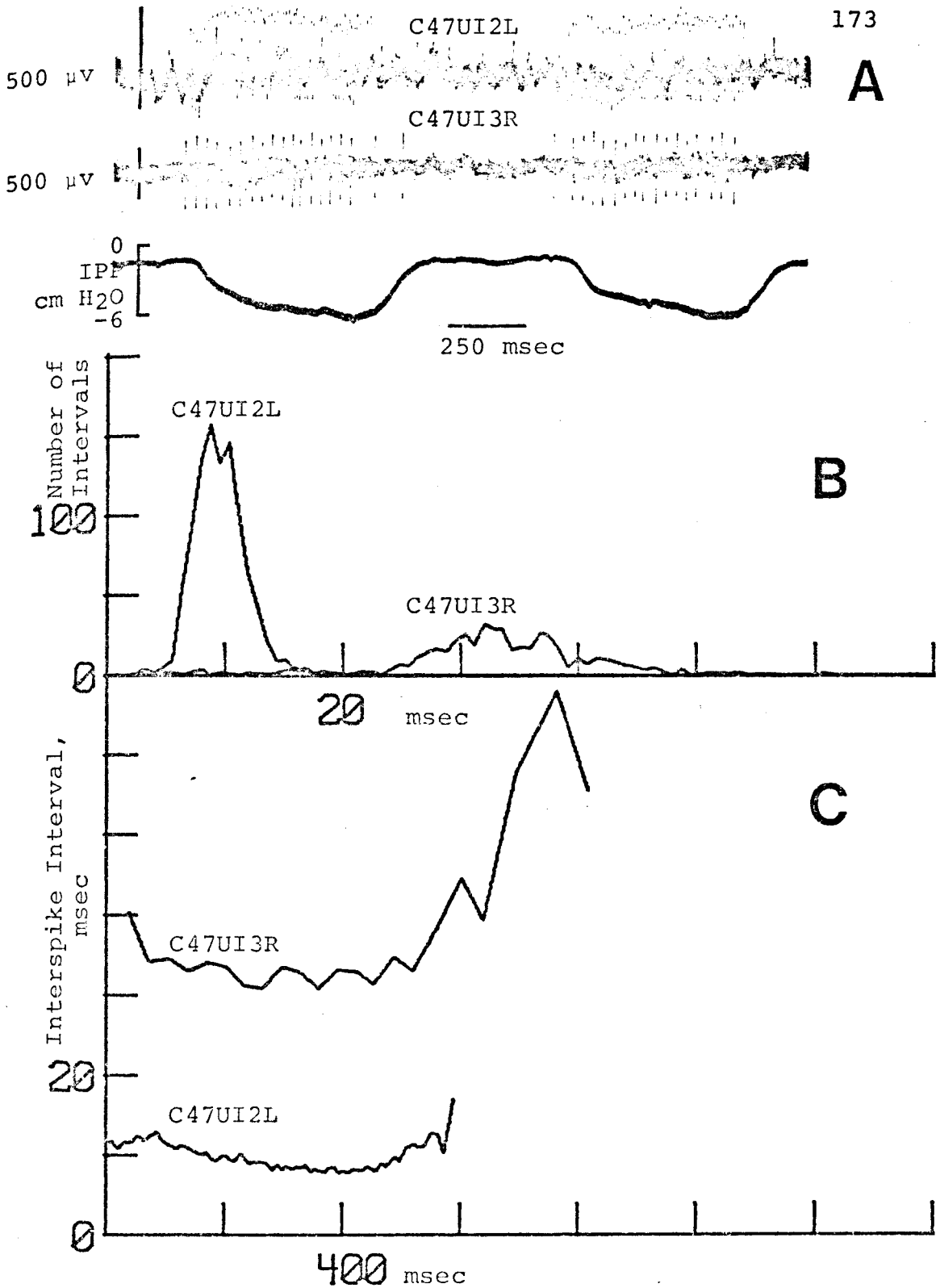


Figure 46. Two simultaneously recorded inspiratory cells from right and left medulla (A), their histograms (B) and interspike interval modulation curves (C). Vagi intact.

factors. Rather, it is suggested that inherent variability exists at the cellular level presumably due to differences in cell size, membrane characteristics and interneurone connections. This reasoning explains the scatter of mode (MOD) values plotted as a function of respiratory rate (RR) in Figure 16 and probably is a factor causing scatter of data points in all the regression plots (Figures 11-34). Because of these cell-to-cell differences, quantitation of parameter interactions (e.g., MOD versus  $1/\text{depth}$ ) cannot be done at the single cell level. That is, quantitation of one cell's characteristics represents a specific individual case. Quantitation of the whole population requires sufficient sampling of all representative cell types in the respiratory complex and statistical averaging of cell discharges in time and space. The latter is probably handled physiologically at several synaptic levels in the efferent pathway.

Although Figures 44 and 45 suggest that mode changes are associated with modifications in respiratory depth, Figure 47 presents evidence that this is not always the case. Here, a spontaneous increase in the respiratory rate occurs (40.8%) which is accompanied by an increase in the mode (MOD) interval time (31.6%). Since the maximal change in respiratory depth is at best

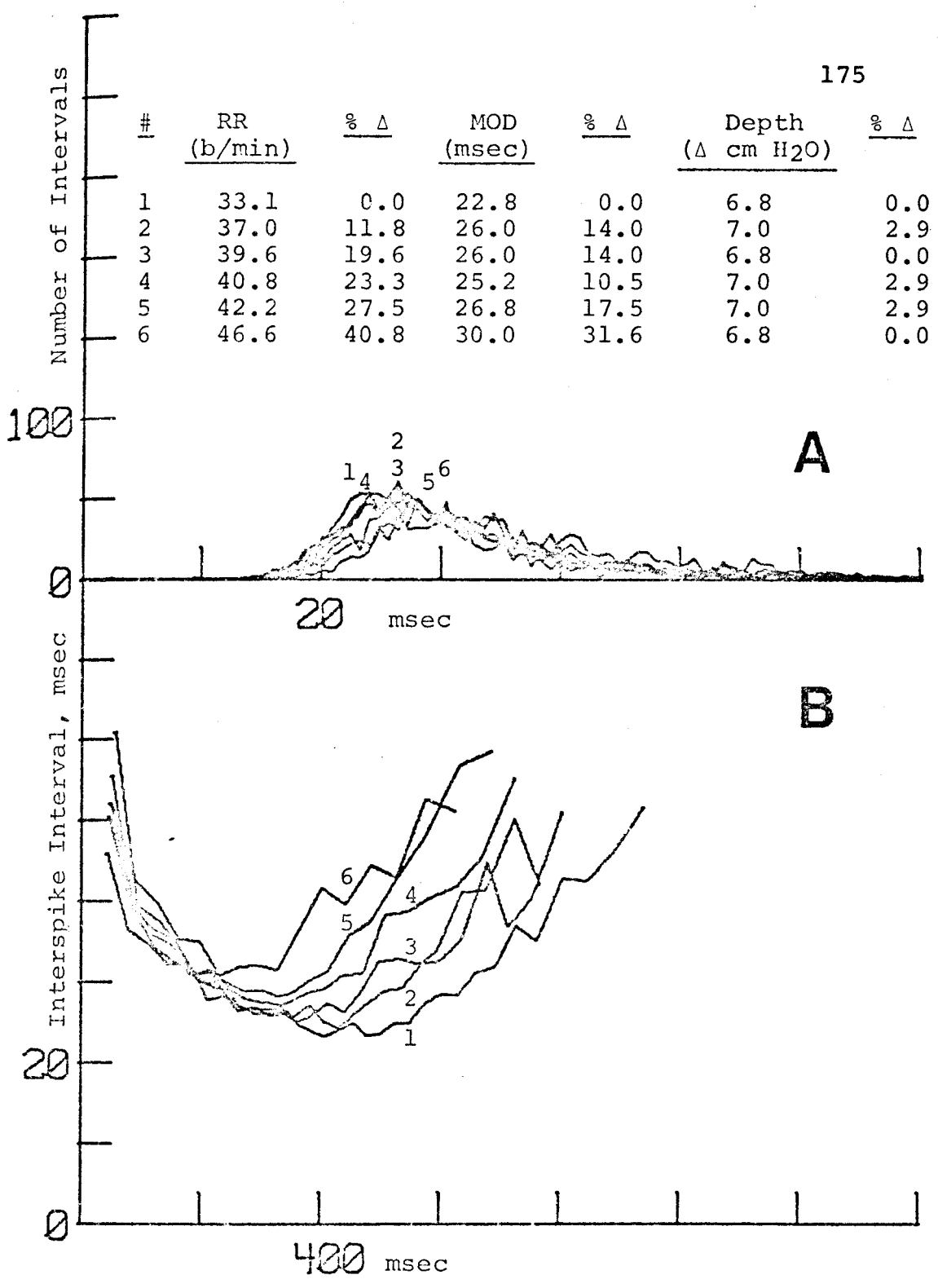


Figure 47. C28UI5. Effect of spontaneous increase in respiratory rate on histogram (A) and interspike interval modulation curve (B). Vagi intact.

only minimal (2.9%), the cellular depression must have been due to other factors. It is interesting to note that the histogram shift to the right (Figure 47A) is not a smooth transition as seen for the rate elevation. Also, the interspike interval modulation curve shift is not parallel (Figure 47B) as observed for other inspiratory cells (Figures 44-45). That is, the latter part of the train is affected to a larger extent than the initial phase of unit activity. These variations may explain the abnormal mode modification during spontaneous changes in breathing patterns involving rate, but not depth.

Seven cells have been selected to demonstrate the effects of certain drugs on discharge patterns. The pharmacologic agents (pentobarbital sodium, thiopental sodium, doxapram hydrochloride and morphine sulphate) were chosen because of their known effects on gross respiratory function. For example, Figures 48 and 49 illustrate the quantitative responses of cell C62UI3R to two injections of doxapram. Before discussing the physiological responses, however, the format of Figure 49 is described to facilitate reader comprehension of the data presentation. The data from the six remaining cells in this section have identical formats.

In Figure 49, the twenty five parameters examined

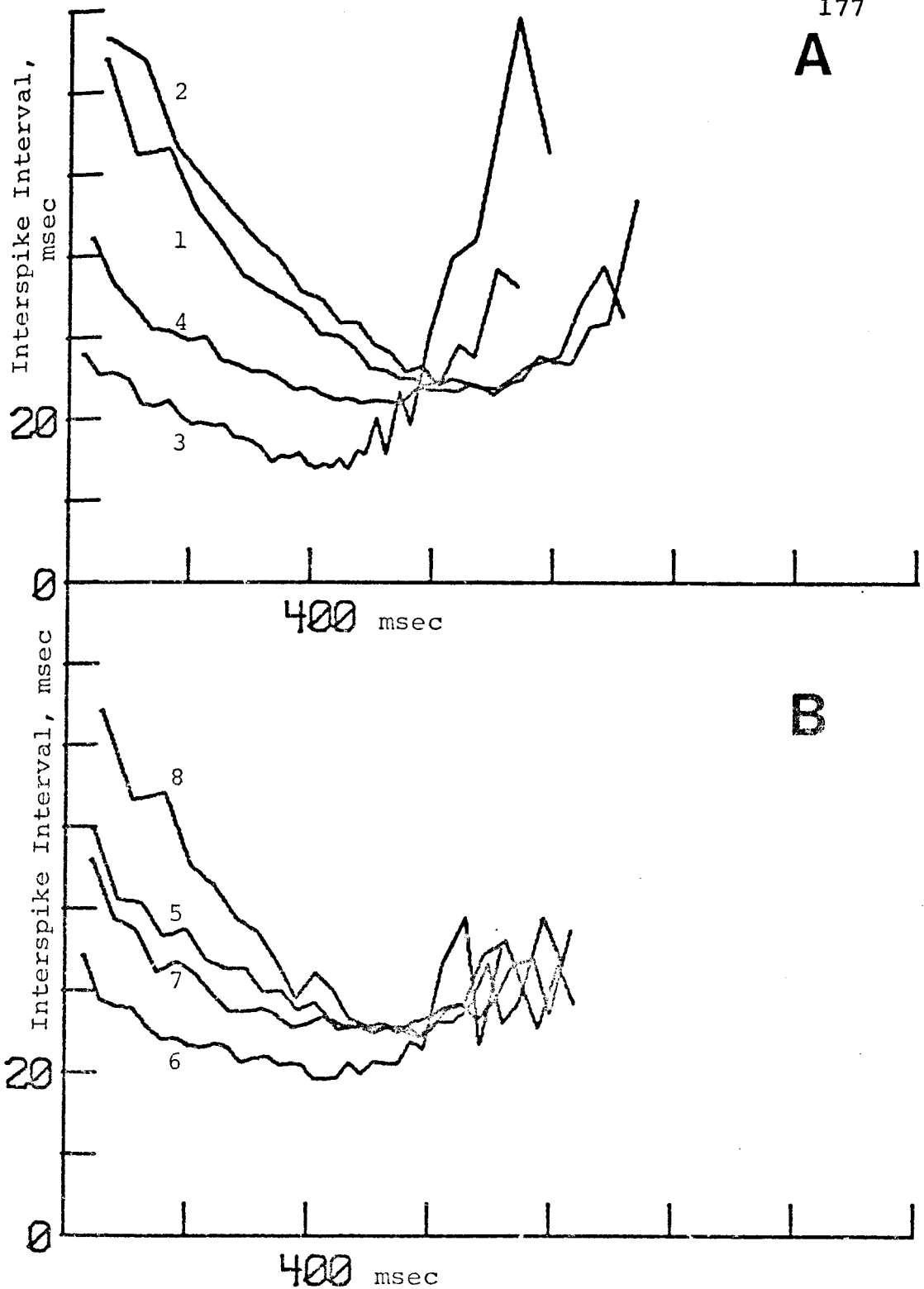


Figure 48. C62UI3R. Effect of doxapram on interspike interval modulation curve: A. 1.6 mg/Kg doxapram given between Curves 2 and 3; B. 1.6 mg/Kg doxapram given between Curves 5 and 6. Vagi intact.

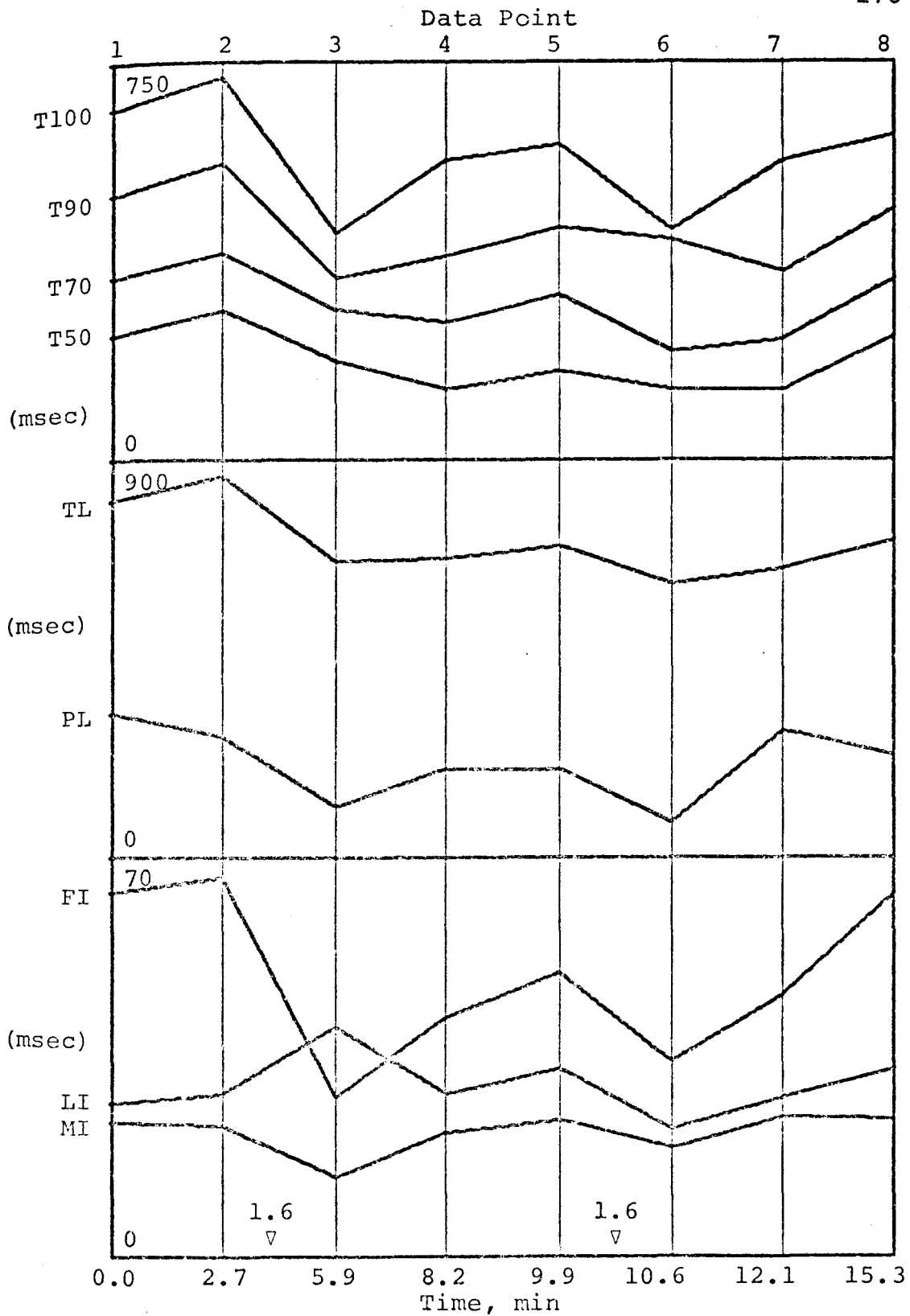


Figure 49A. C62UI3R. Effect of doxapram on parameters of inspiratory cell discharge. See text for details.

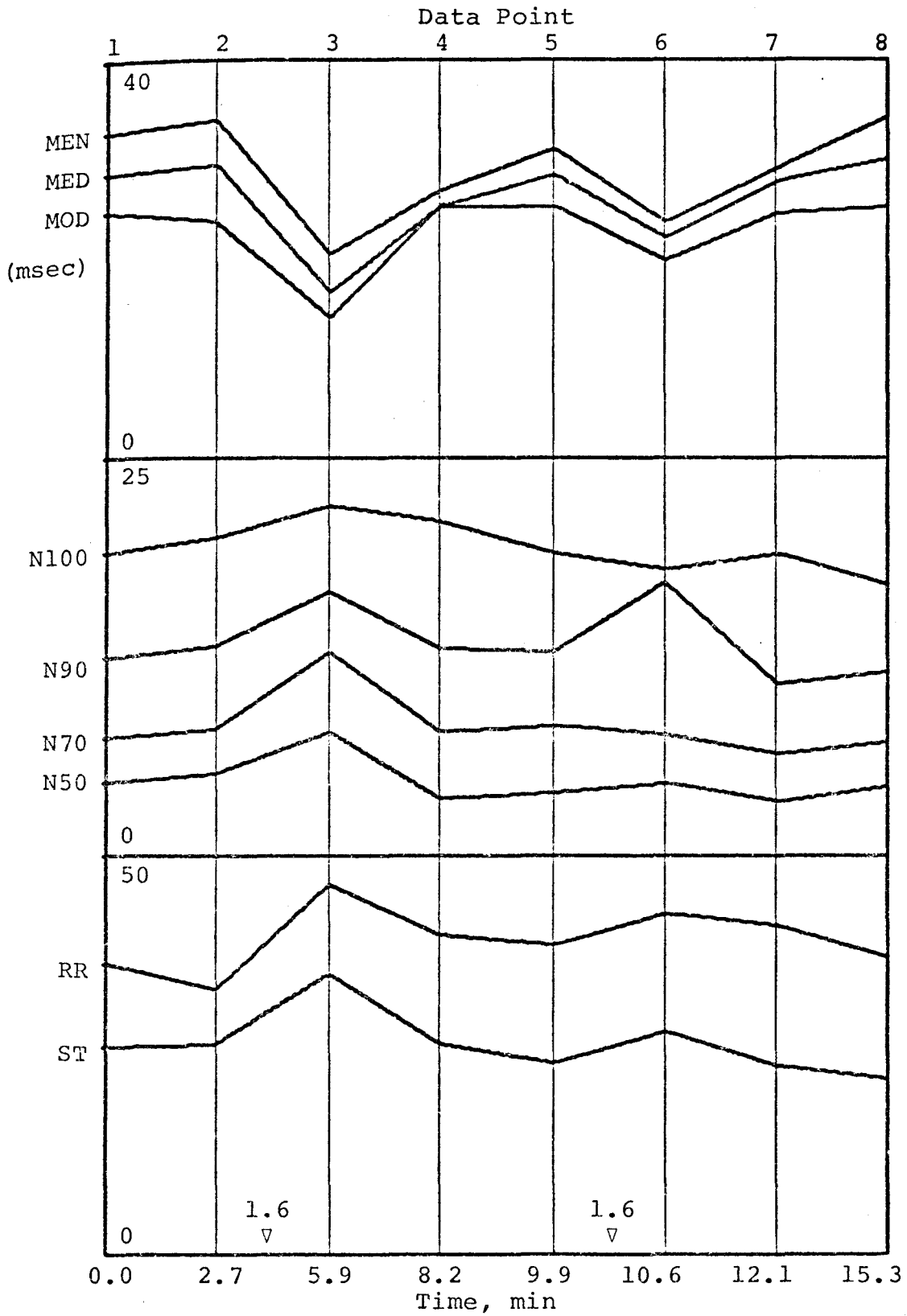


Figure 49B. C62UI3R. Effect of doxapram on parameters of inspiratory cell discharge. See text for details.



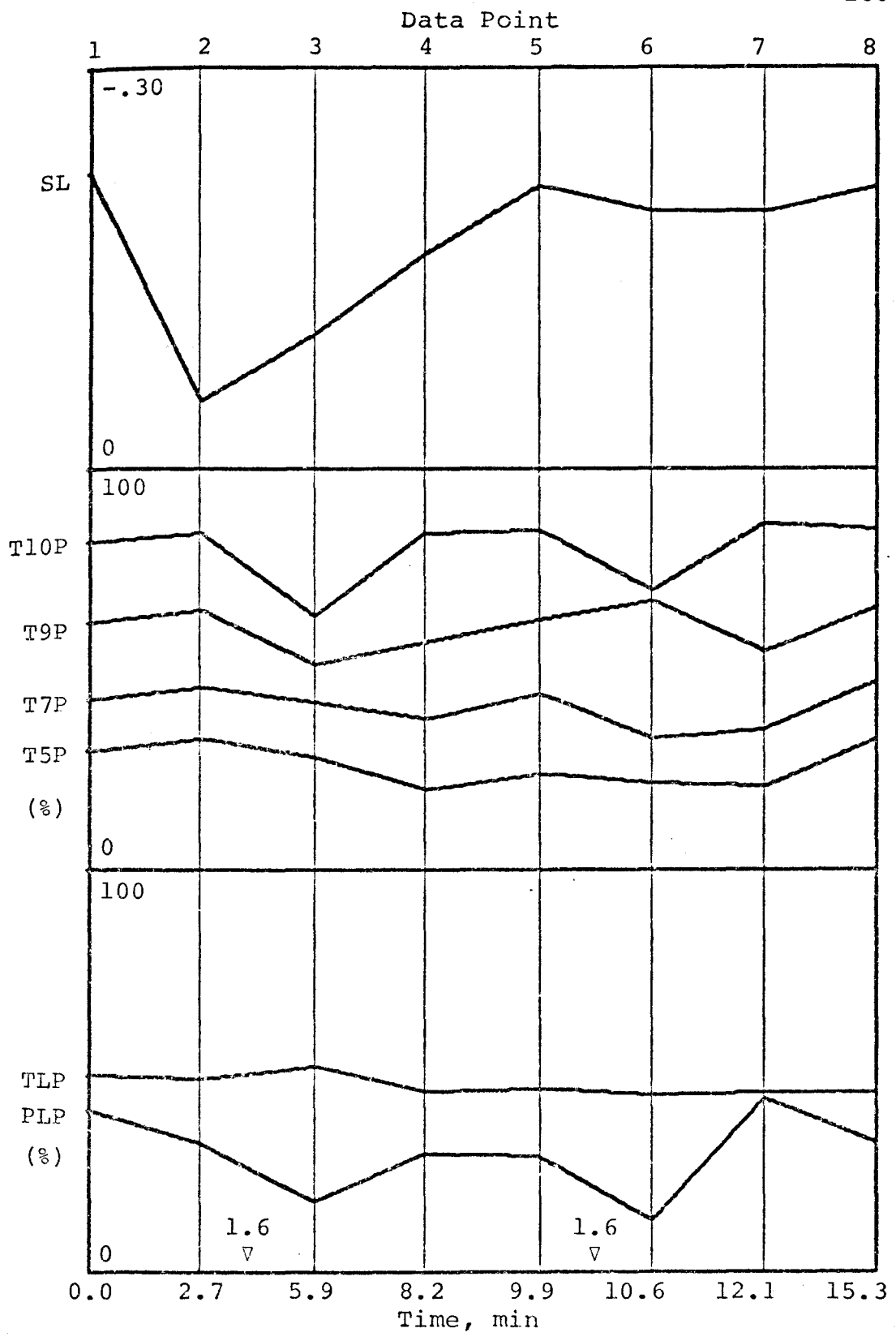


Figure 49C. C62UI3R. Effect of doxapram on parameters of inspiratory cell discharge. See text for details.

by regression analysis (Figures 11-43) are plotted as a function of time in nine panels on three pages. All parameters are identified according to the abbreviations listed in Table IV. By examination of mode and respiration rate trends in the upper and lower panels of Figure 49B, respectively, comparison of single cell characteristics with the mean population data is possible.

Each steady state measurement is marked off by a vertical line which is identified by a data point number at the top and a time measurement to the nearest tenth of a minute at the bottom. The uniformly spaced data points form a non-linear time function on the X-axis. Scaling of dependent variables is given for each group of parameters and appropriate dimensions are supplied. Respiration rate is in breaths per minute. Timings of doxapram injections are identified by inverted triangles with superscript numbers denoting drug dose in milligrams per kilogram (e.g.  $\nabla^{1.6}$  ).

Changes in interspike interval modulation curves for inspiratory cell C62UI3R following two separate injections of doxapram are shown in Figure 48A and B respectively. Superimposed Curves 1 and 2 in Figure 48A are pre-injection controls. Curve 3 represents the discharge pattern change shortly after injection of doxapram

(1.6 mg/kg) and Curve 4 shows partial recovery from drug activation. In Figure 48B, the series is continued using Curve 5 as the new control. The second doxapram injection (1.6 mg/kg) caused a pattern shift to Curve 6, but this activation level was not as high as in Curve 3, possibly due to tachyphylaxis. Curves 7 and 8 are recovery patterns.

Doxapram effects on the individual parameters for cell C62UI3R are shown in Figure 49. Doxapram accelerates the respiration rate (RR) and decreases the mode (MOD) interval time. Corresponding decreases in the first interval (FI), minimal interval (MI), times to reach portions of the way to the minimal interval (T50, T70, T90, T100), plateau level (PL), train length (TL), mean (MEN) and median (MED) times all parallel predicted parameter responses to increased rates and decreased modes from the mean population data (Figures 35-43). The last interval (LI) has an inconsistent biphasic response while the N50, N70, N90, N100 and ST parameters follow predicted changes in the mode but not the rate. The spikes per train response indicates that the mode time is decreasing faster than the train length. The slope (SL) has a very unpredictable pattern which must be associated with the poor correlation coefficients for

regressions of slope on rate (Figure 19, I-VI) and slope on mode (Figure 31, I-VI). Variable responses are seen for the normalized parameters (T5P, T7P, T9P, T10P, PLP) while the percent time the cell fires during the respiratory cycle (TLP) appears to be independent of rate and mode changes induced by doxapram.

Variations occur in the characteristics of single unit patterns in response to barbiturate administration. These are illustrated in the following two sets of data derived from two different cells. The vagus nerves were intact in each case which permits comparison of these single cell data with the mean I-VI population data in Figures 35-43. Data from the first inspiratory cell, C63UI4R, are presented in Figures 50 and 51. Identical doses of pentobarbital (1.5 mg/kg) were injected between each data point. With increasing accumulated dosage of pentobarbital it can be seen that respiration rate decreases from 44.7 to 24.0 breaths per minute (Figure 51B). The mode (MOD) time, however, only slightly increases from 14.0 to 13.8 milliseconds as illustrated in Figure 50A, where histogram distributions fall in overlapping populations. In contrast to the response of cell C62UI3R (Figure 49B) where both spikes per train (ST) and respiration rate (RR) showed parallel increases after doxapram

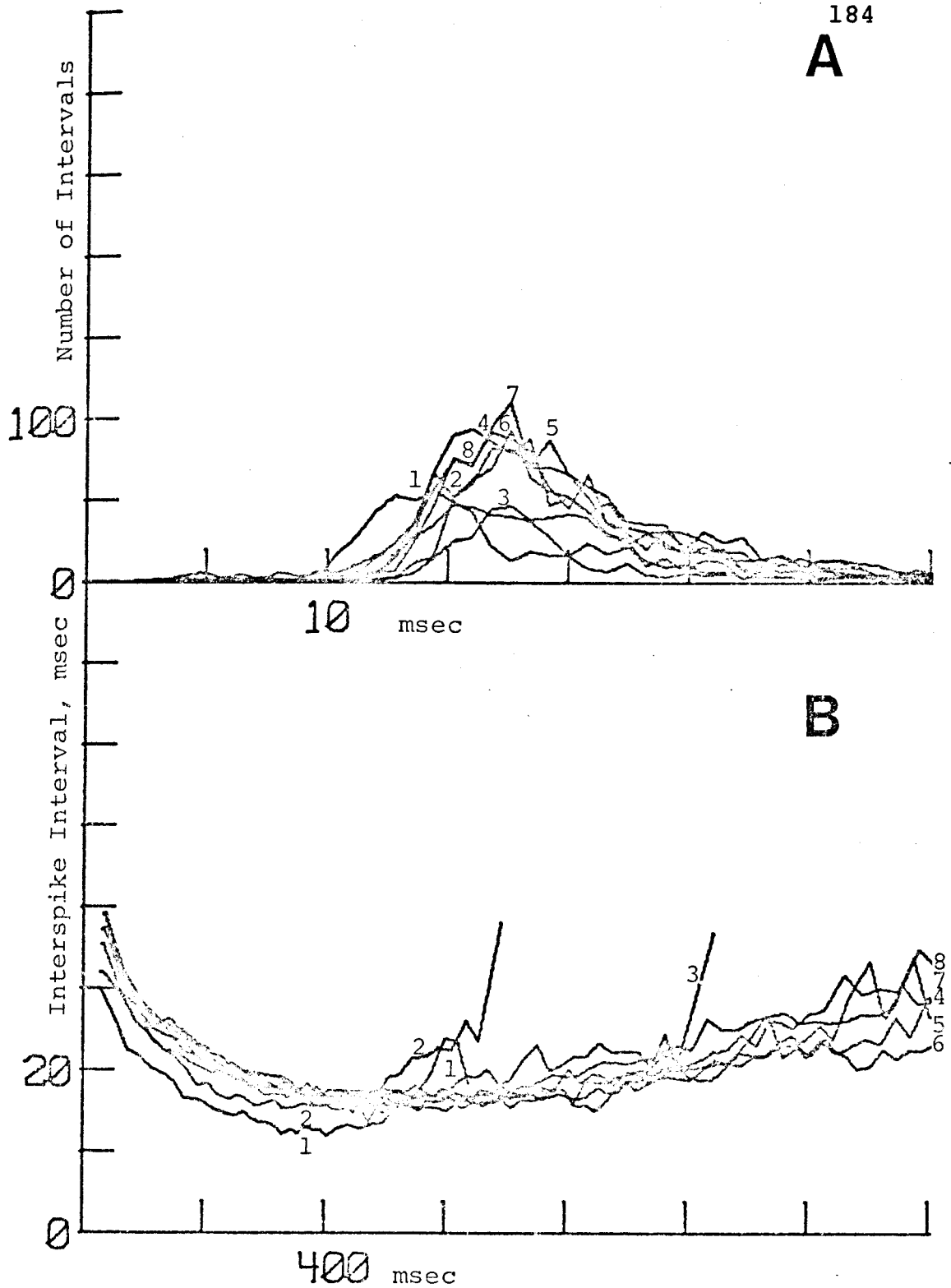


Figure 50. C63UI4R. Effect of cumulative doses of pentobarbital on histogram (A) and interspike interval modulation curve (B). Vagi intact.

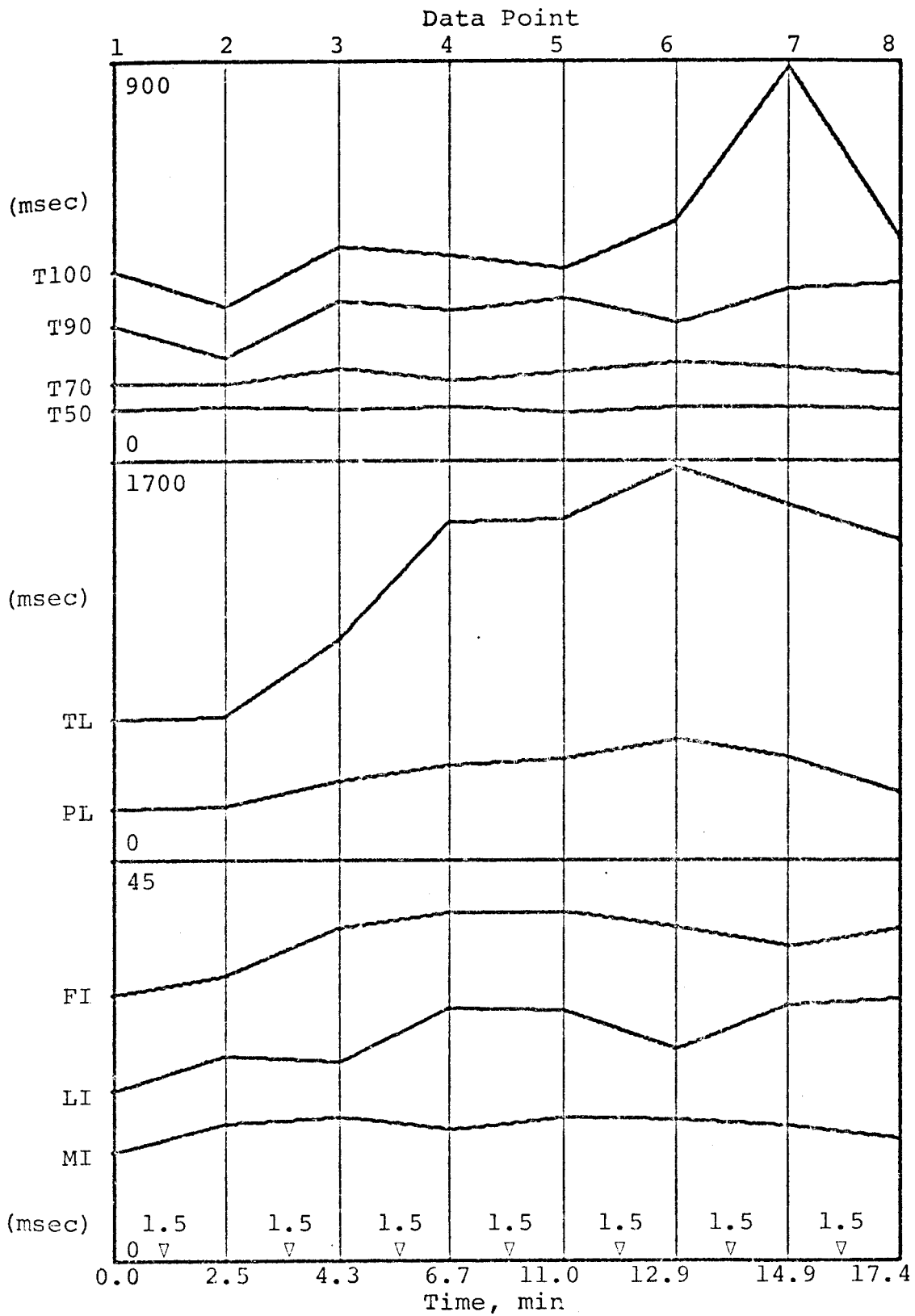


Figure 51A. C63UI4R. Effect of pentobarbital on parameters of inspiratory cell discharge. See text for details.

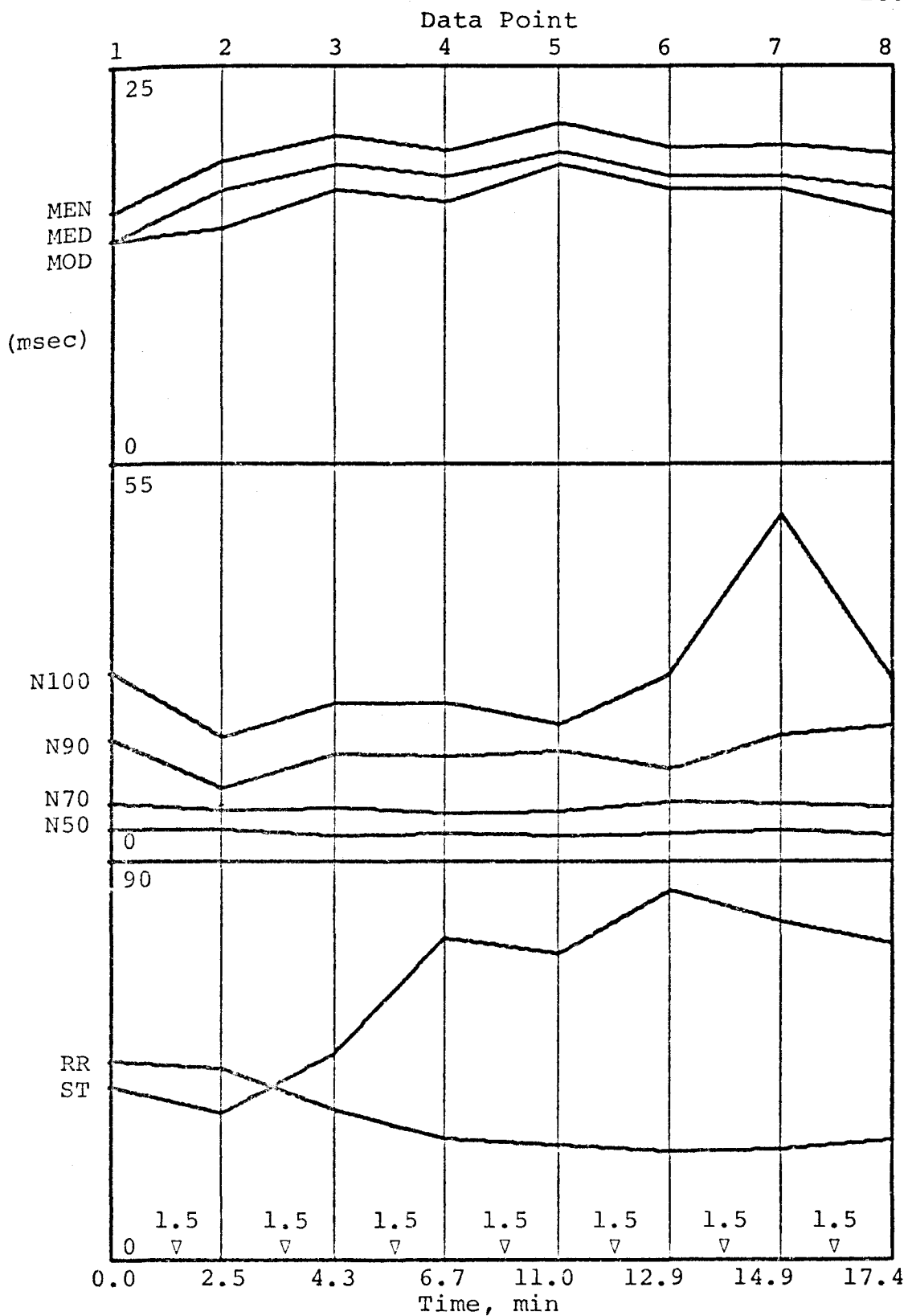


Figure 51B. C63UI4R. Effect of pentobarbital on parameters of inspiratory cell discharge. See text for details.

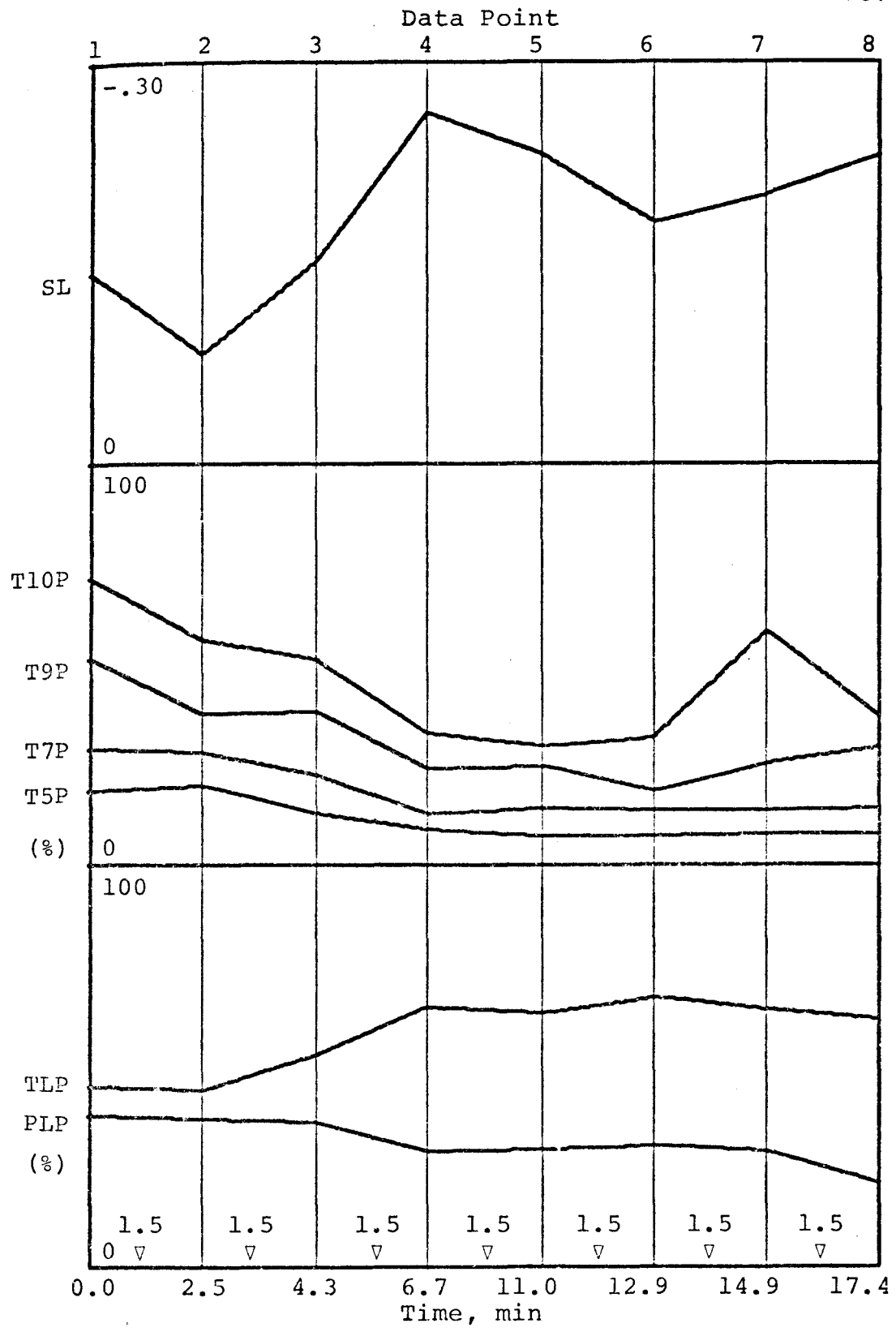


Figure 51C. C63UI4R. Effect of pentobarbital on parameters of inspiratory cell discharge. See text for details.



injections, ST and RR for cell C63UI4R change in opposite directions during barbiturate depression. This is due to the dramatic increase in train length (TL) following respiratory rate slowing with only a small decrease in the discharge frequency (Figure 50B). Parameters T50, T70, T90, T100 and PL show little change except for T100, which increases abruptly after the sixth injection of pentobarbital (Figure 51A). These observations, coupled with the longer train lengths, explain the decrease in the T5P, T7P, T9P, T10P and PLP measurements (Figure 51C). Vertical shifts of the interspike interval modulation curves are small as shown by the relatively small changes in first interval (FI), minimum interval (MI) and last interval (LI) during drug depression (Figure 51A). Similarly, N50, N70, N90 and N100 parameters show little change until high nembutal levels are reached (Figure 51B). In general, the slope (SL) tends to increase as a function of drug depression, but individual directional changes are not predictable. Finally, pentobarbital causes inspiratory train discharge to occupy more of the respiratory cycle time as indicated by the increase in TLP (Figure 51C). This is due to a greater increase in train length than in cycle time during the slowing of respiratory rate. Most of these pattern changes are consistent

with those predicted from the mean population data (Figures 35-43).

Data from the second inspiratory cell, C83UI1L, are presented in Figures 52 and 53. In this case, consecutive equal doses of pentobarbital (1.7 mg/kg) decreases both respiration rate from 71.8 to 29.9 breaths per minute and mode interval time from 21.2 to 14.8 milliseconds. The latter is illustrated in Figure 52A by a significant shift of the histogram interval distributions to the left. This mode response, which is quite different from that for cell C63UI4R (Figures 50 and 51), is indicative of unit activation (increased firing frequency). This presents an apparent paradox in which increased unit activity occurs during depression of respiratory rate. Data bearing specifically on this problem will be presented in the last part of Results. Similar to cell C63UI4R, the train length of cell C83UI1L was greatly increased by pentobarbital as shown in Figure 52B. The increasing train length and decreasing mode both contribute to an increase in spikes per train, and again the mode is inversely related to respiration rate. While the T50, T70, T90, T100 and PL parameters show predicted responses, an interesting inversion of first and last interval times is seen. That is, with accumulating doses

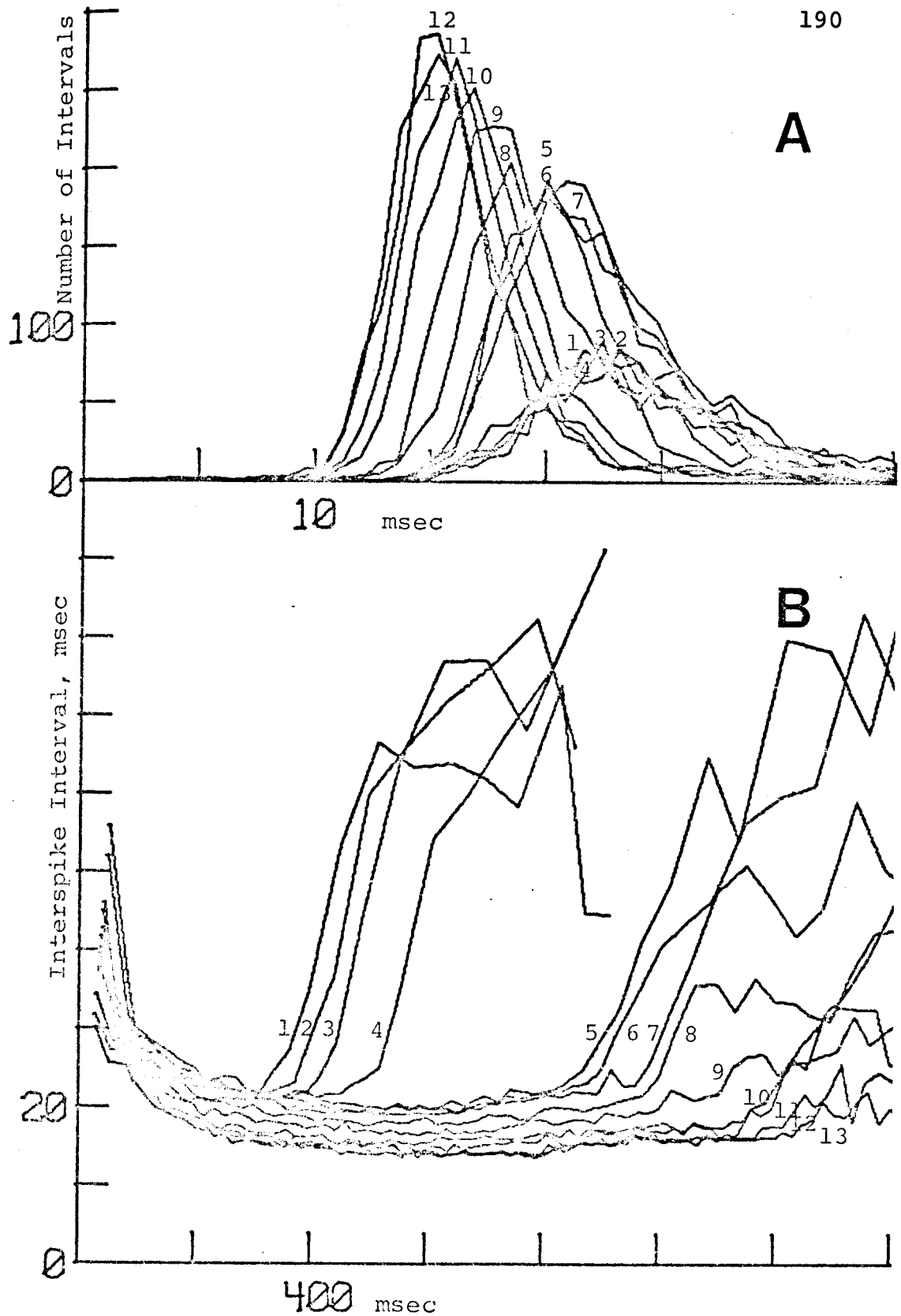


Figure 52. C83UI1L. Effect of cumulative doses of pentobarbital on histogram (A) and interspike interval modulation curve (B). Vagi intact.



Figure 53A. C83UILL. Effect of pentobarbital on parameters of inspiratory cell discharge. See text for details.

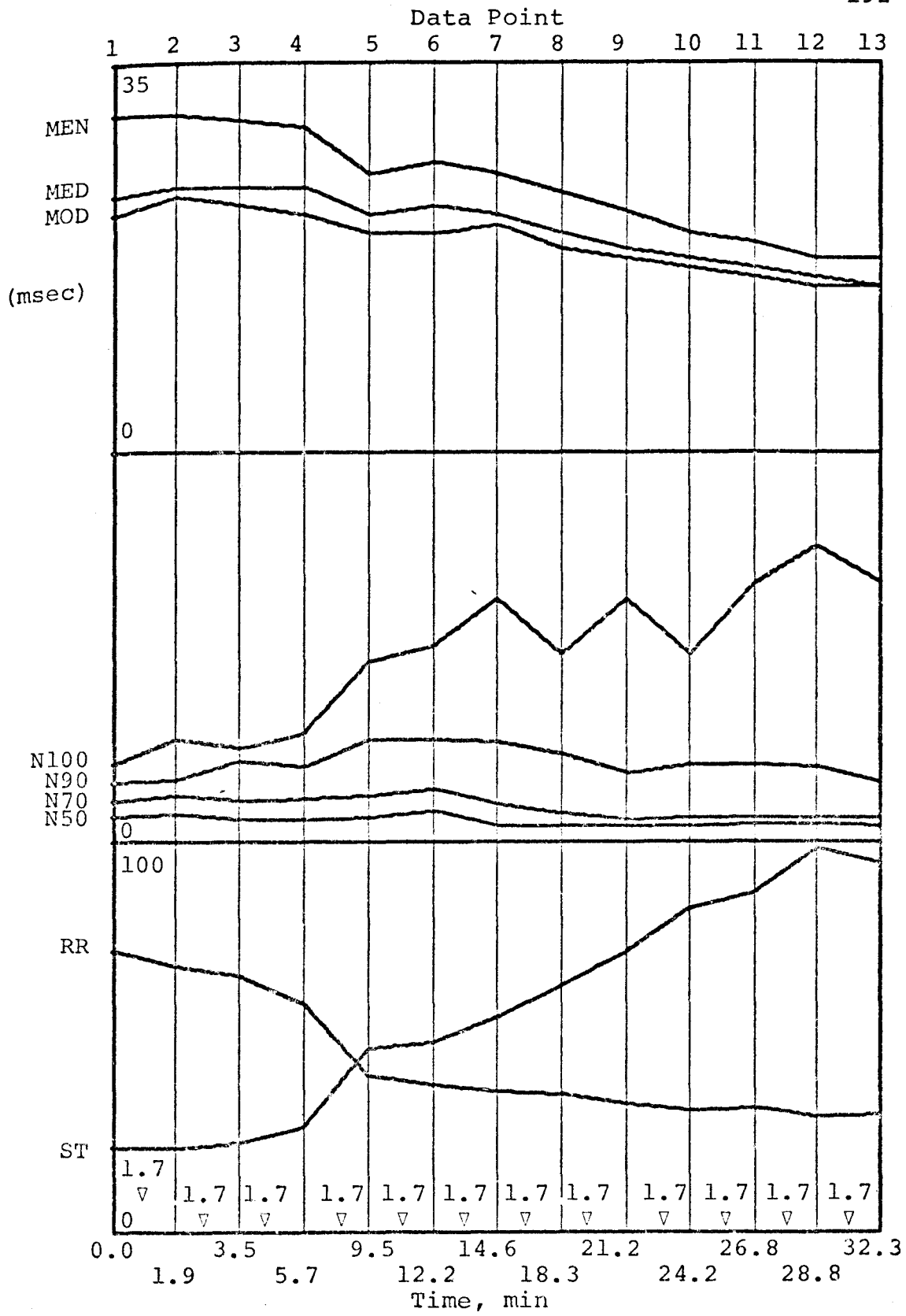


Figure 53B. C83UIII. Effect of pentobarbital on parameters of inspiratory cell discharge. See text for details.



Figure 53C. C83U11L. Effect of pentobarbital on parameters of inspiratory cell discharge. See text for details.

of pentobarbital, the first interval and the last interval change in opposite directions (Figures 52B and 53A). The slope again shows tendencies to increase at higher barbiturate doses. This is due mainly to first interval lengthening. Finally, the TLP remains independent of either rate or mode changes induced by pentobarbital depression. Single cell data and mean population data (Figures 35-43) are generally consistent for all parameters plotted as a function of rate, but not as a function of mode. The two significant exceptions are the regression of mode and last interval on respiratory rate.

The next inspiratory cell, C81UI5L, was selected to show single unit response to barbiturate depression in the absence of vagal afferent feedback. The data are shown in Figures 54 and 55 which can be correlated with the mean I-VO population data trends in Figures 35-43. Data point 1 represents a prevagotomy control. Data points 2 and 3 show parameter responses to section of the left (  $\nabla$  ) and right (  $\nabla$  ) vagus nerves respectively. Most parameter responses appear independent of unilateral vagotomy and become evident only after section of the second vagus nerve. The immediate effect of severing the left vagus on firing frequency is most likely related to the fact that this unit was located in the left medulla.

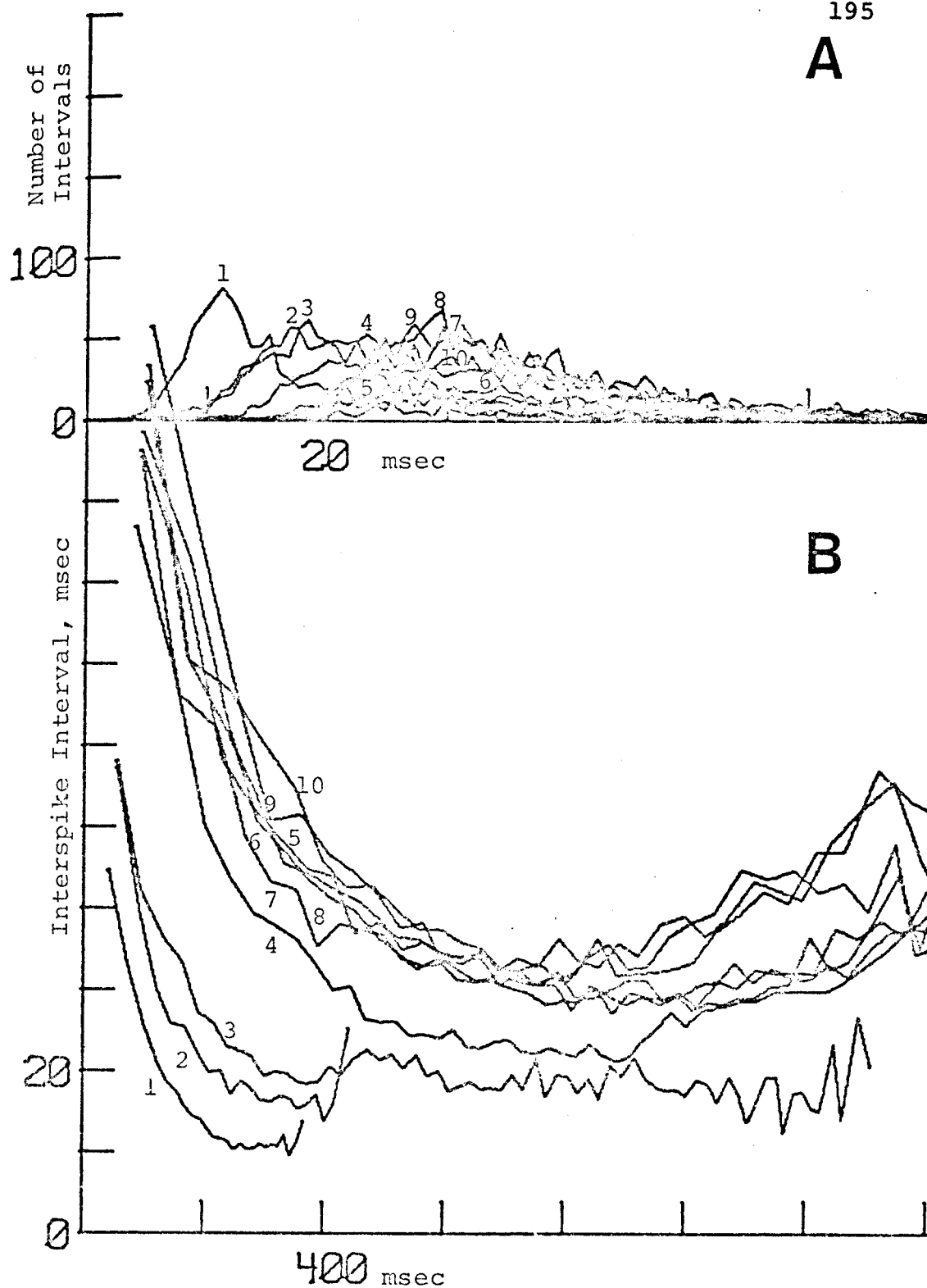


Figure 54. C81UI5L. Effect of cumulative doses of pentobarbital on histogram (A) and interspike interval modulation curve (B). Left vagus sectioned after #1; right vagus sectioned after #2.



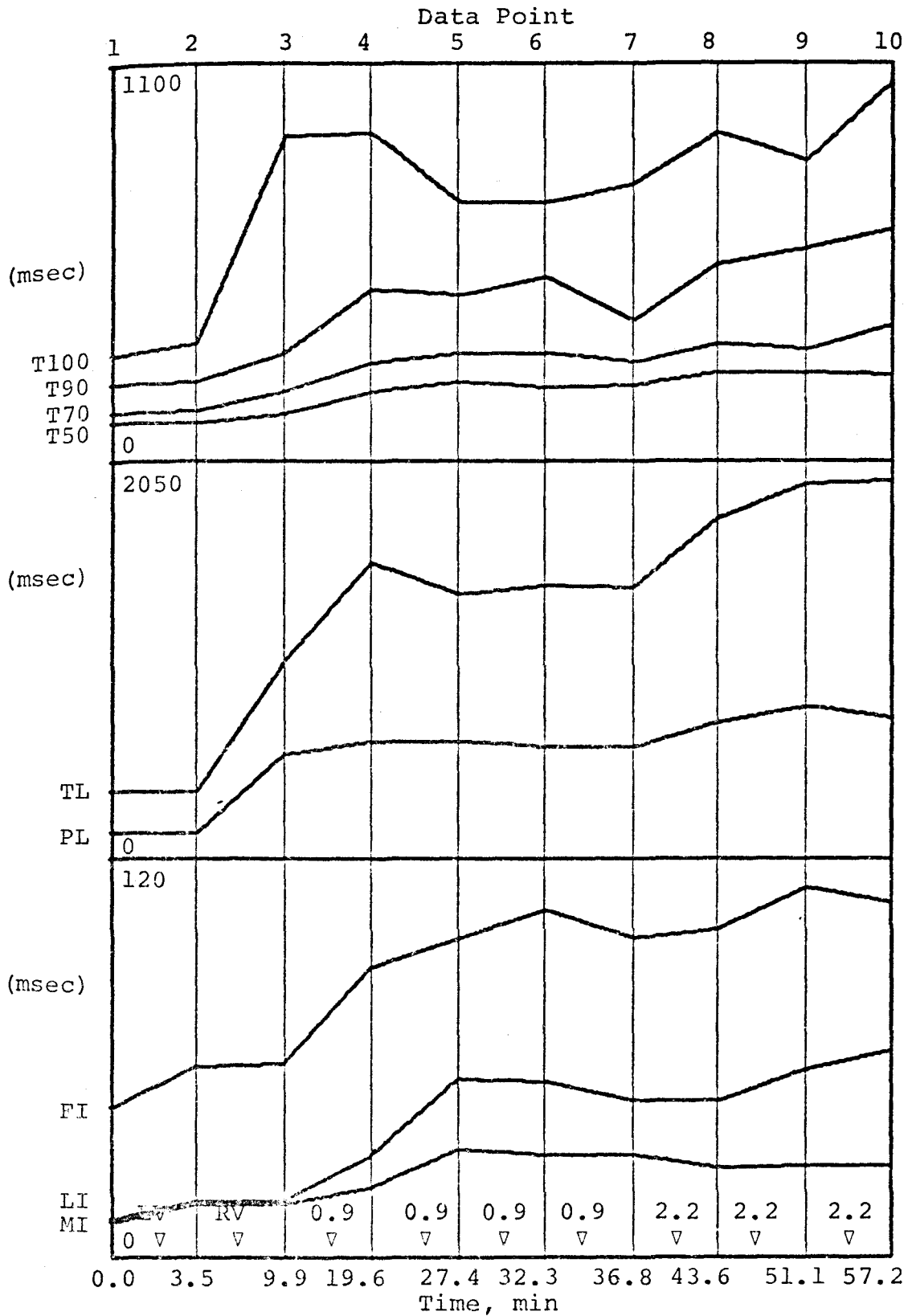


Figure 55A. C81UI5L. Effect of pentobarbital on parameters of inspiratory cell discharge. See text for details.

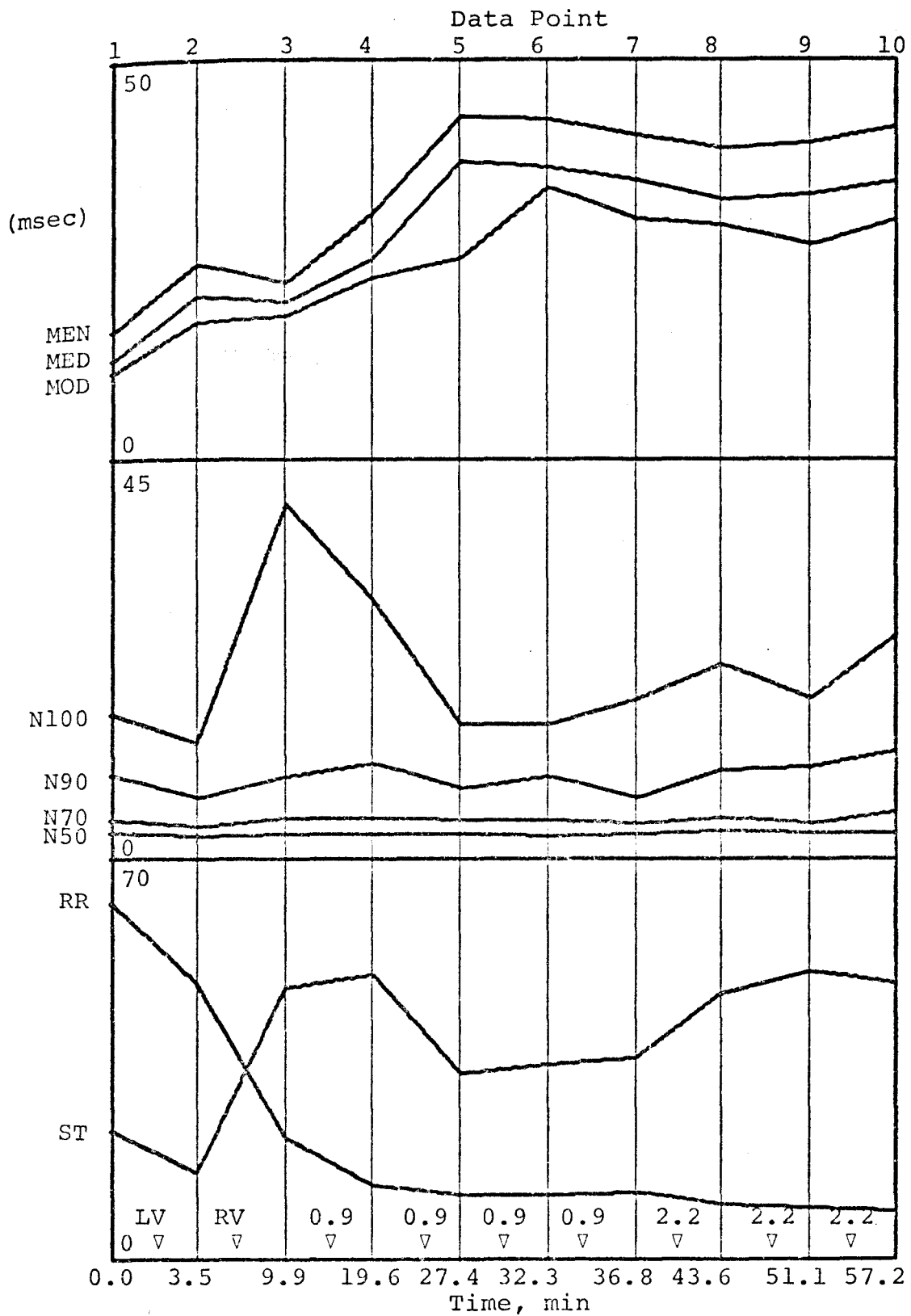


Figure 55B. C81UI5L. Effect of pentobarbital on parameters of inspiratory cell discharge. See text for details.

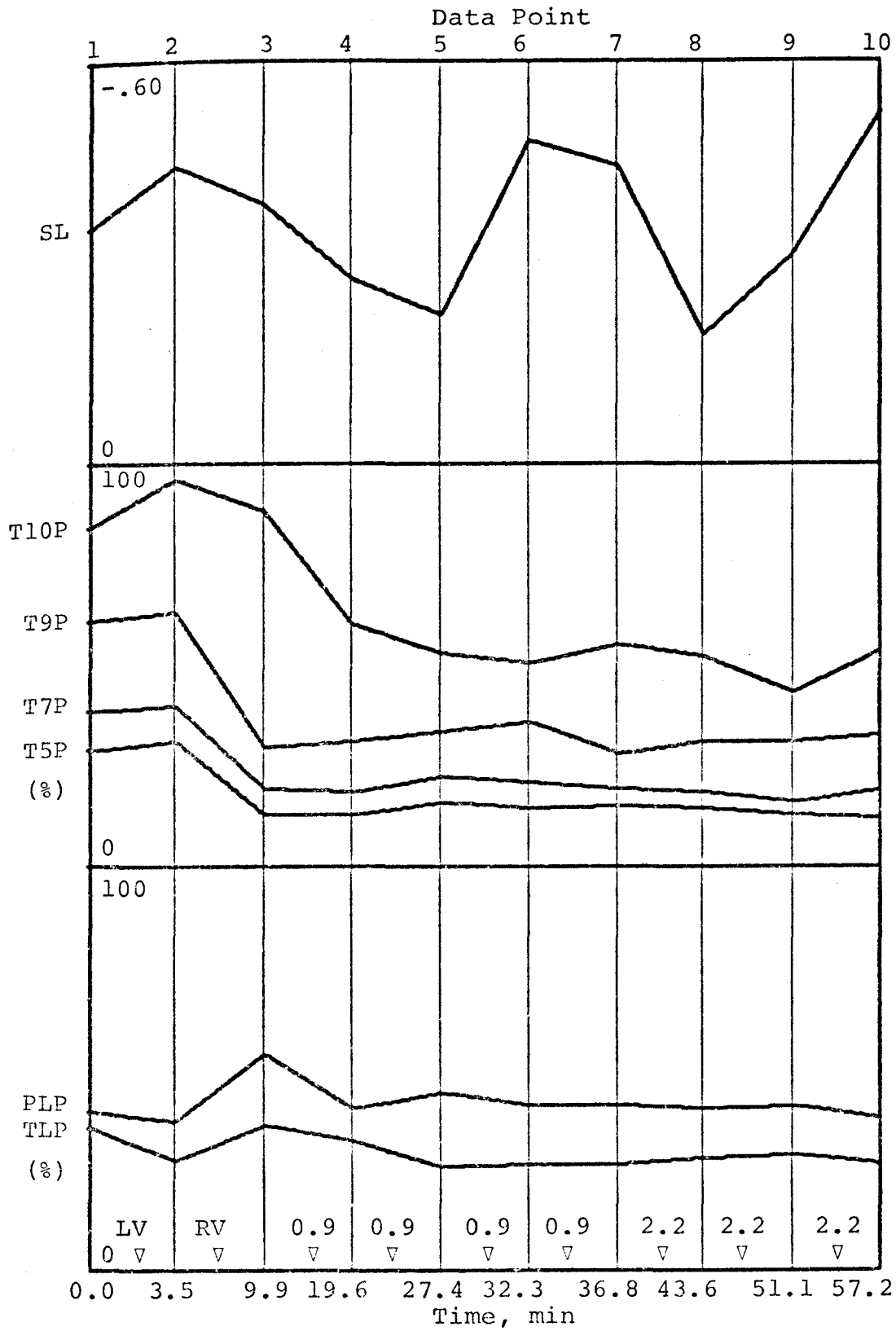


Figure 55C. C81UI5L. Effect of pentobarbital on parameters of inspiratory cell discharge. See text for details.

Starting with data point 3 as the vagotomized control, the parameters are plotted as a function of pentobarbital dose in Figure 55. Respiration rate decreases from 21.1 to 8.1 breaths per minute and mode time increases from 18.0 to 30.0 milliseconds. This is a typical depression response with the characteristic shift in the histogram to the right (Figure 54A) and an upward shift of the interspike interval modulation curve (Figure 54B). The train length increases, but due to the depression in firing frequency, the spikes per train remains relatively constant. The reciprocal relationship between ST and RR seems to be lost when the vagi are cut, at least for this cell. All other parameters follow predicted directional changes during rate and firing frequency depression when compared to the mean population data (Figures 35-43). It can be noted that the slope data are very unpredictable.

The responses of two expiratory cells during barbiturate accumulation are presented to show not only individual characteristics, but to reveal pattern similarities to inspiratory cells already discussed. The vagi are intact for each expiratory cell, which allows comparisons to be made with the mean E-VI population data in Figures 35-43. The data from the first expiratory

cell, C64UE4R, are plotted in Figures 56 and 57. Respiration rate decreases from 86.6 to 40.5 breaths per minute during pentobarbital depression while the mode shows little change except for a significant decrease from 17.2 to 12.4 milliseconds at data point 4. This single shift in the histogram can be seen in Figure 56A. As with inspiratory cells, the train length greatly increases (Figures 56B and 57A) which causes an increase in the spikes per train. An inverse relationship between ST and RR is thus established. A small increase in the plateau level is also observed. The T50, T70, T90 and T100 parameters tend to increase indicating that pentobarbital modifies the expiratory discharge pattern such that it takes more time to reach portions of the way to the minimum interval. These increases parallel train length increase since T5P, T7P, T9P and T10P measurements remain approximately constant for most of the drug range. The first, last and minimum intervals increase significantly, slightly and not at all as a function of accumulated drug respectively. The N parameters appear to be independent of drug modification except for N100 which increases after three pentobarbital injections. The slope response appears randomized as in other cells. Finally, TLP remains relatively constant indicating that this expiratory cell always fires

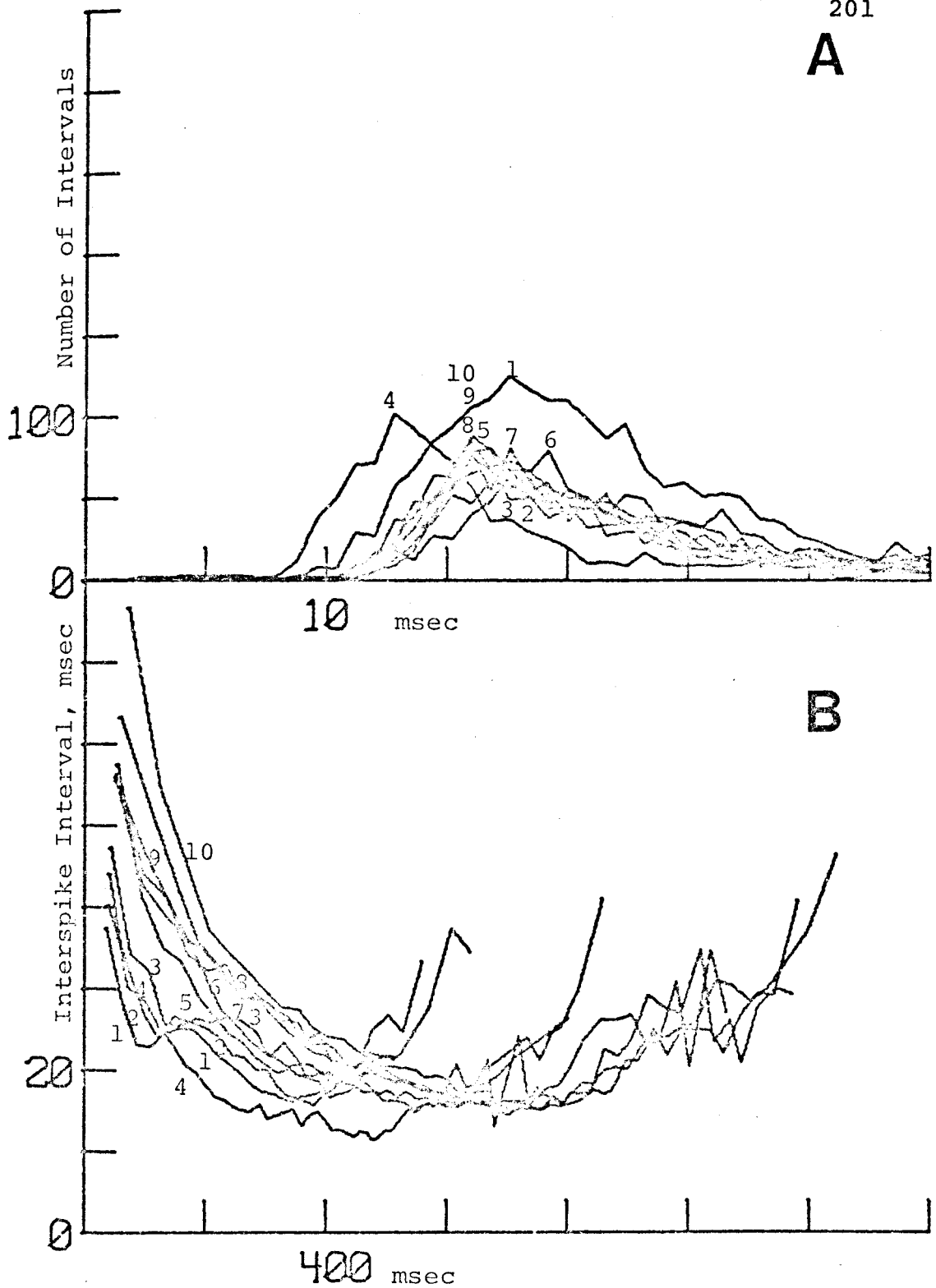


Figure 56. C64UE4R. Effect of cumulative doses of pentobarbital on histogram (A) and interspike interval modulation curve (B). Vagi intact.

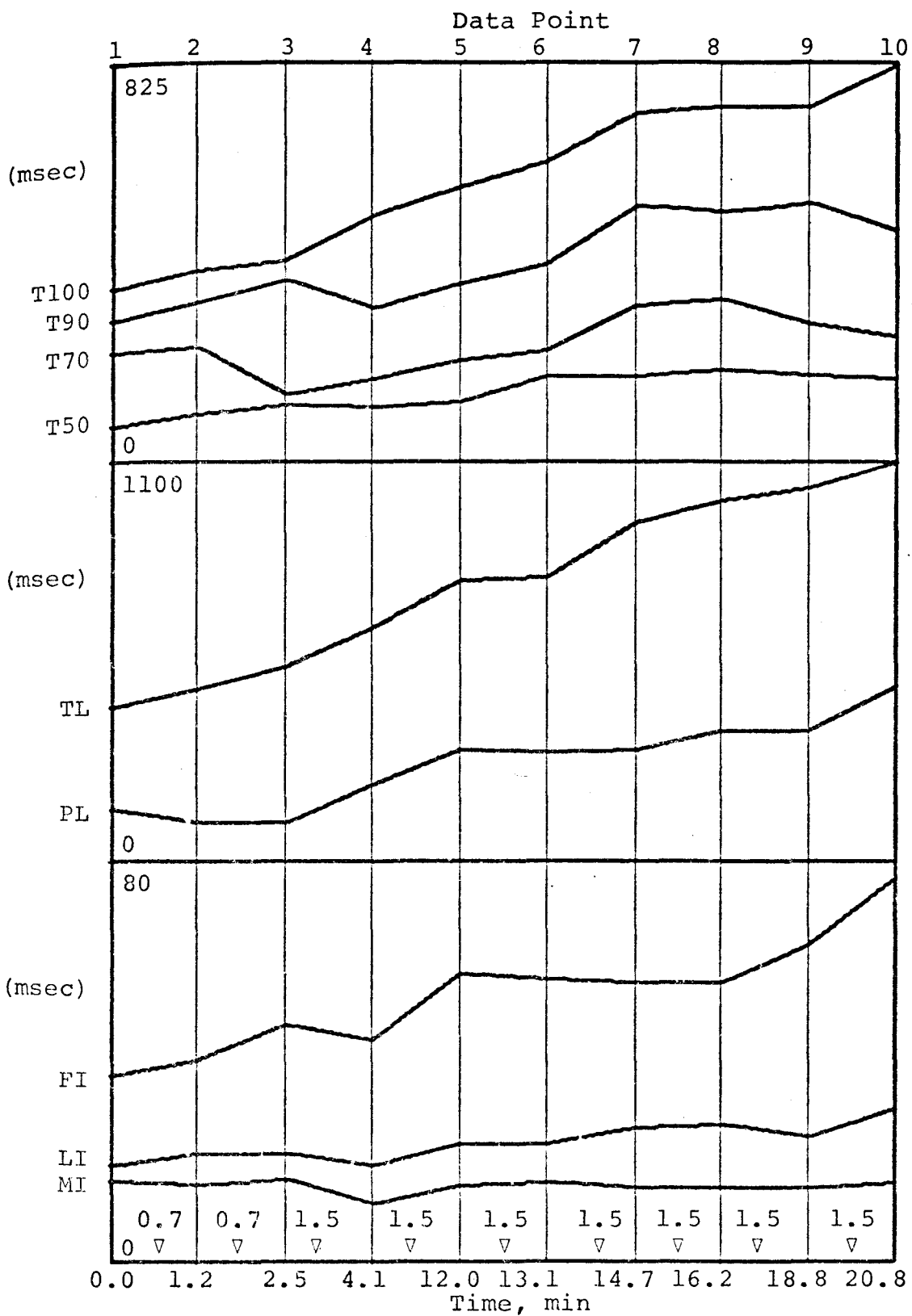


Figure 57A. C64UF4R. Effect of pentobarbital on parameters of expiratory cell discharge. See text for details.

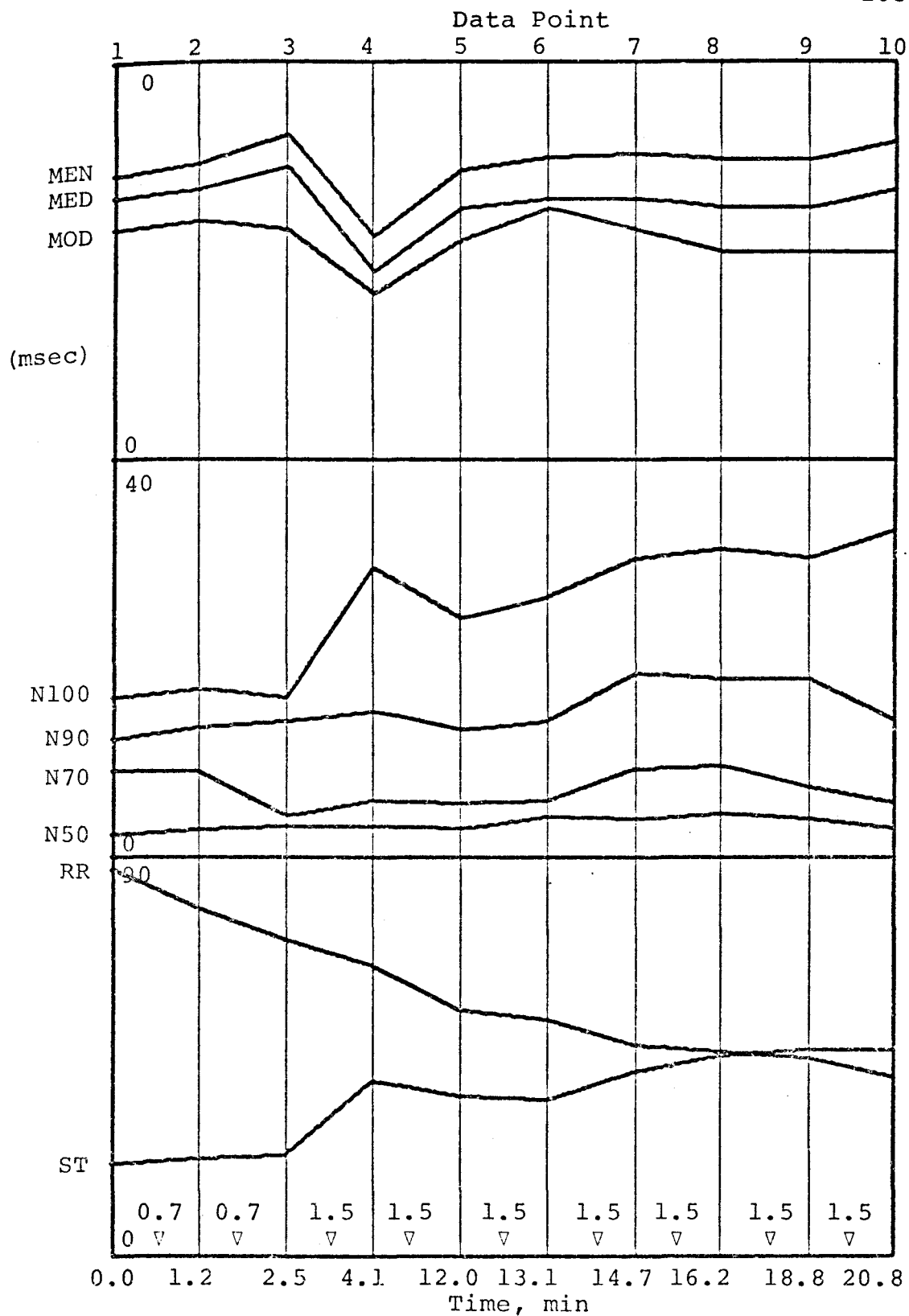


Figure 57B. C64UE4R. Effect of pentobarbital on parameters of expiratory cell discharge. See text for details.



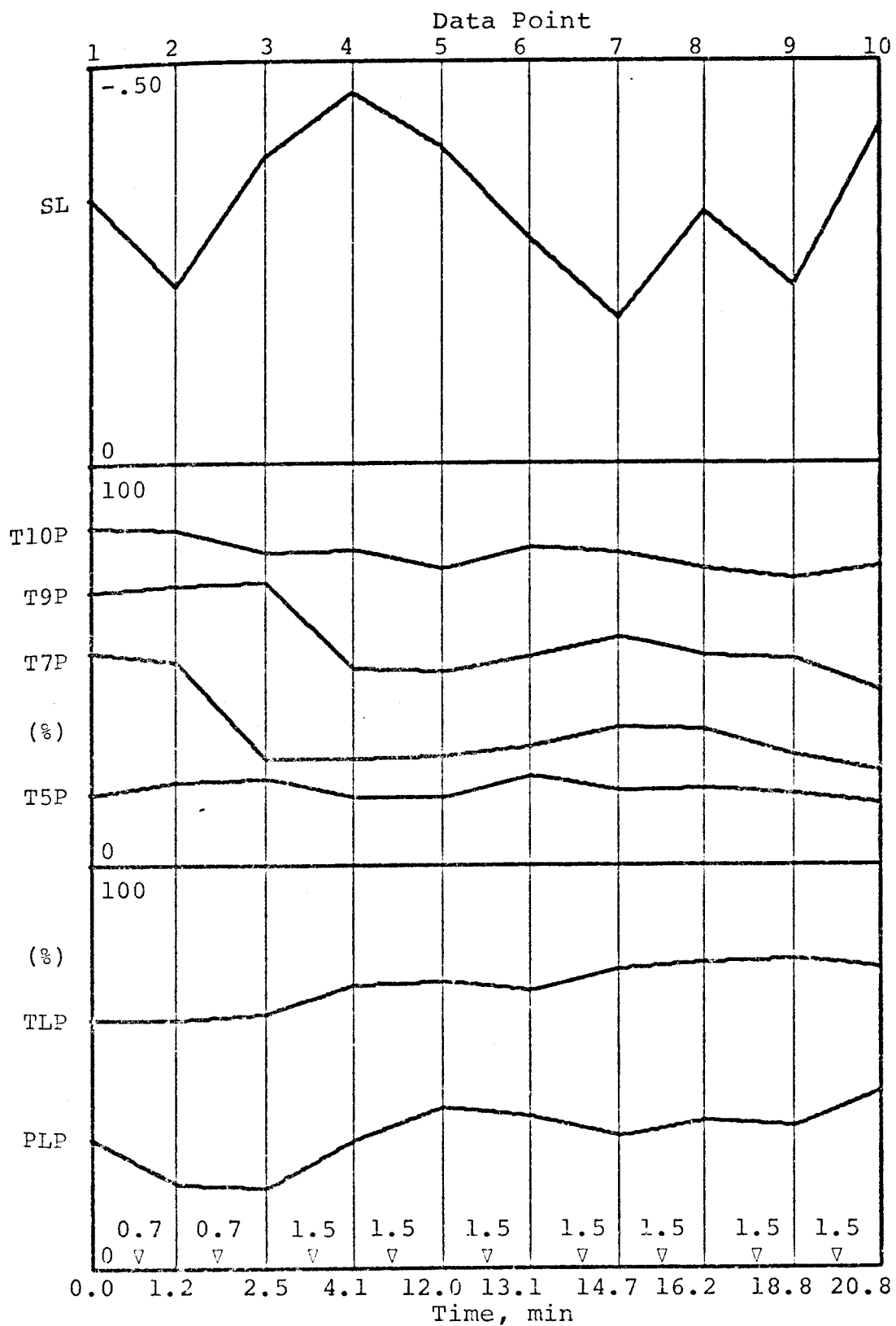


Figure 57C. C64UE4R. Effect of pentobarbital on parameters of expiratory cell discharge. See text for details.

for the same fraction of the respiratory cycle, regardless of the respiratory rate. All of these single cell observations on expiratory cell C64UE4R follow the mean E-VI population trends in Figures 35-43. It is significant that no outstanding differences between expiratory and inspiratory single cell responses to barbiturate depression can be observed.

Data from the second expiratory cell, C71UE2R, are presented in Figures 58 and 59. Successive barbiturate injections depressed the respiratory rate from 21.6 to 11.5 breaths per minute, but produced a triphasic shift in the mode. The mode first increased from 14.0 to 40.4 milliseconds, then decreased to 26.8 milliseconds and finally increased to 42.0 milliseconds. This can be seen as an inflection in the mode, median and mean curves between data points 9 and 12 in Figure 59B. Every other data point is plotted in Figure 58 to clarify histogram and interspike interval modulation curve shifts. This effect is best seen in Figure 58B. Increasing dosage of pentobarbital first causes an upward shift in the interspike interval modulation curve (data points 1, 3, 5, 7, 9), then a downward shift occurs (data point 11) followed by a final upward shift (data point 13). This pattern resembles the activation of

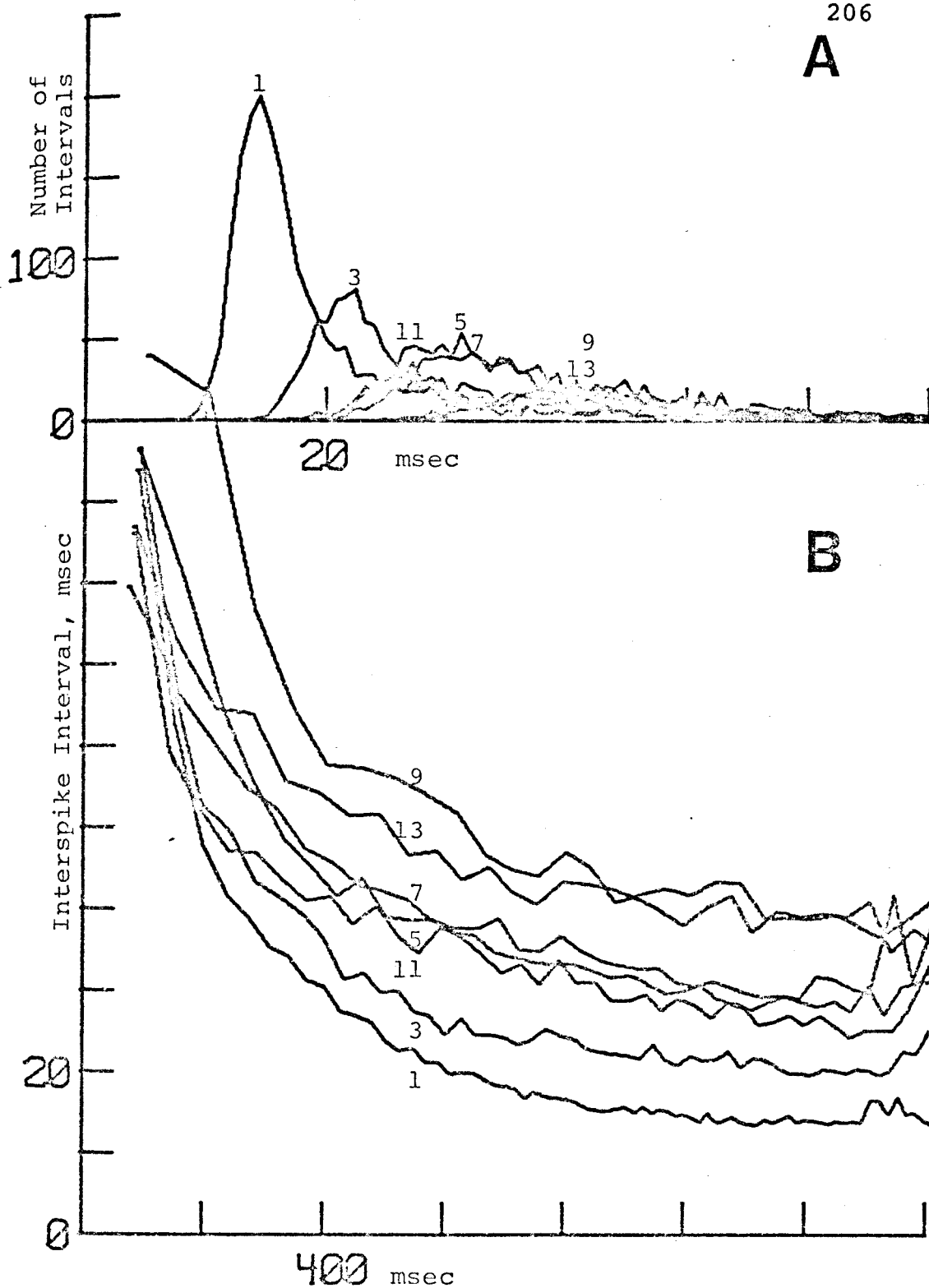


Figure 58. C71UE2R. Effect of cumulative doses of pentobarbital on histogram (A) and interspike interval modulation curve (B). Vagi intact.

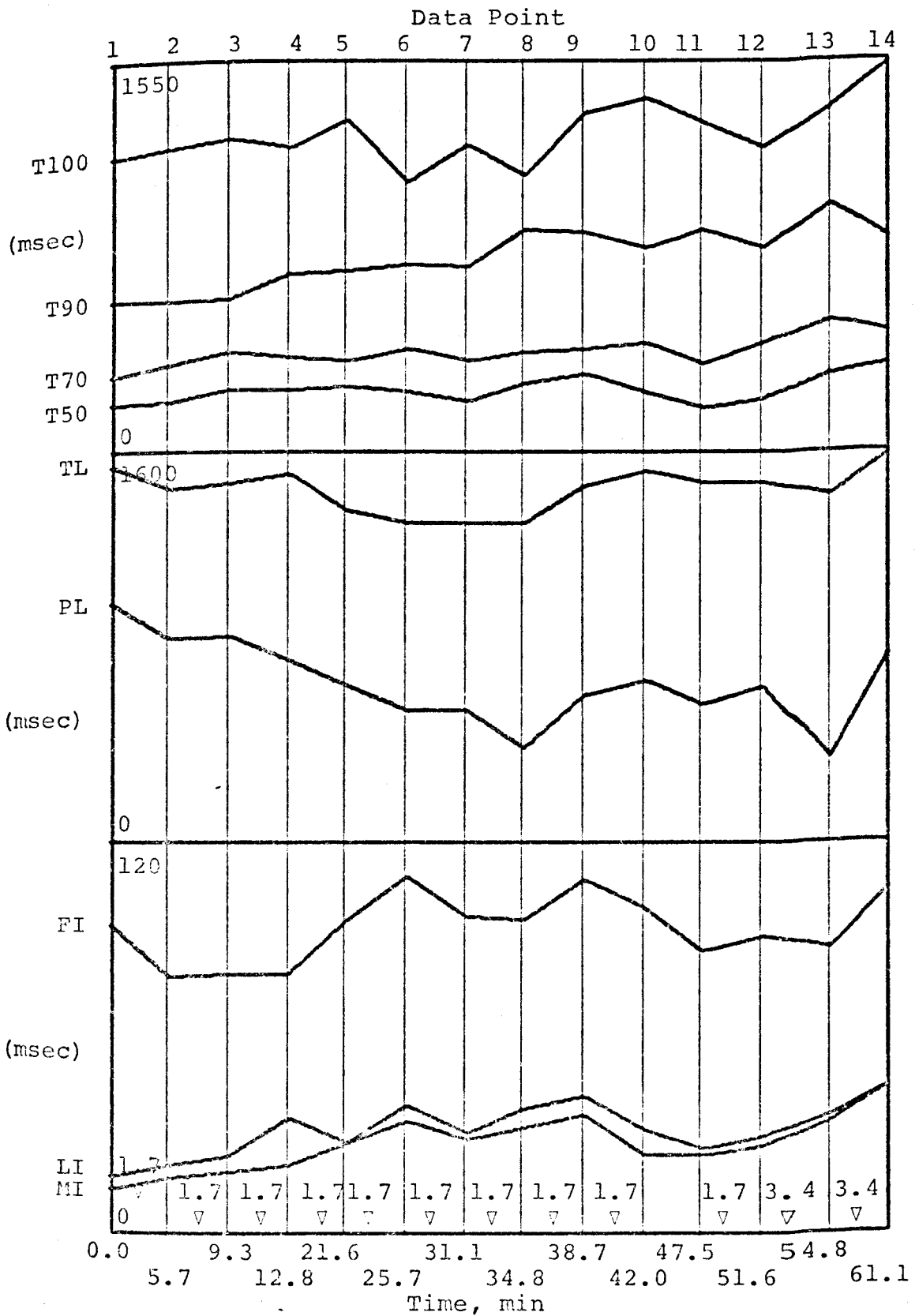


Figure 59A. C71UE2R. Effect of pentobarbital on parameters of expiratory cell discharge. See text for details.

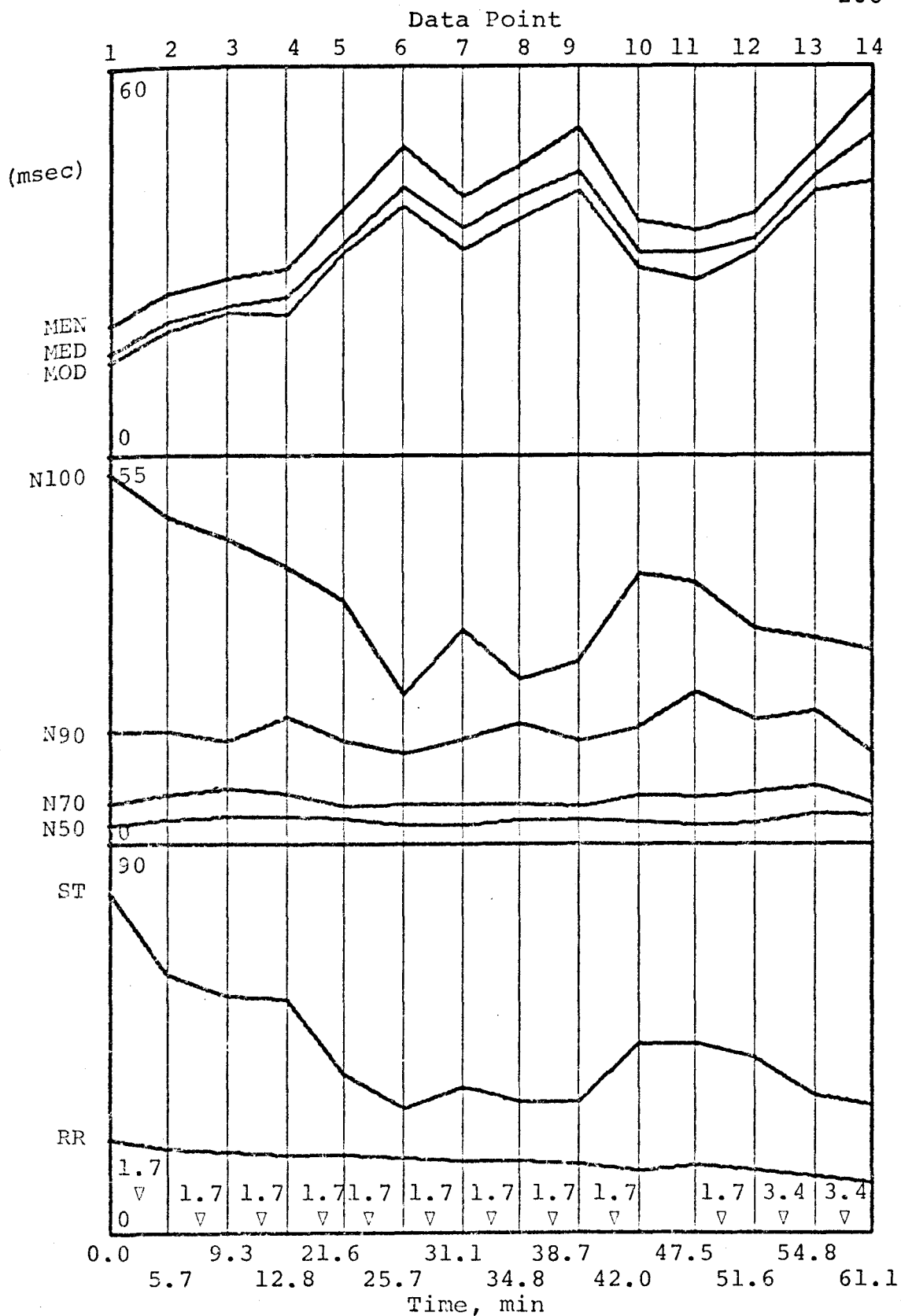


Figure 59B. C71UE2R. Effect of pentobarbital on parameters of expiratory cell discharge. See text for details.

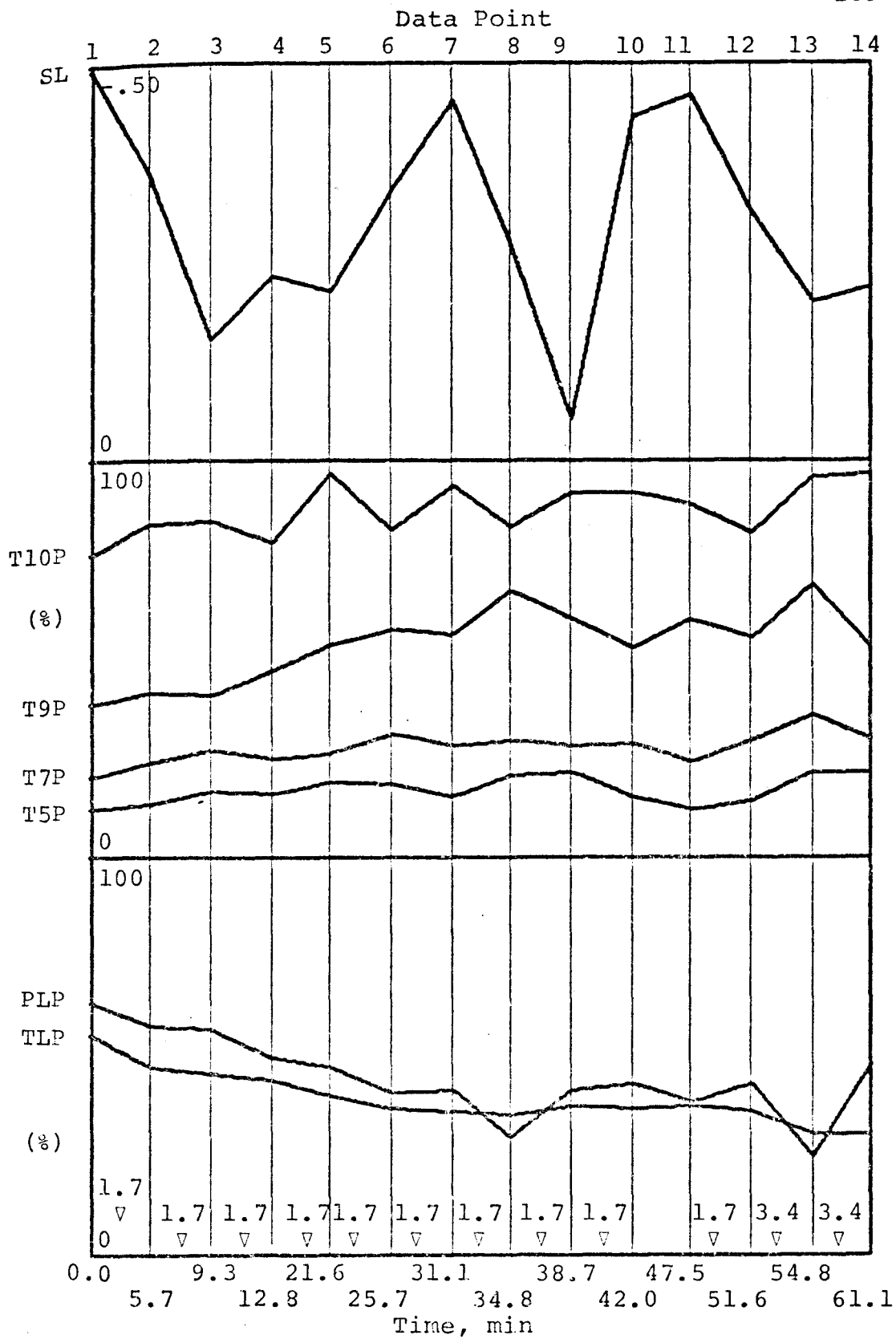


Figure 59C. C71UE2R. Effect of pentobarbital on parameters of expiratory cell discharge. See text for details.

inspiratory cell C83UI1L (Figures 50 and 51) during barbiturate administration and will be examined later.

Expiratory cell C71UE2R exhibits several atypical parameter responses during barbiturate depression. For example, pentobarbital has no effect on modifying the train length, a response which was observed consistently for other cells. For this reason, spikes per train mirror changes in the mode and the inverse relationship with respiratory rate is lost. Also, for the first time, the plateau level is seen to decrease as a function of barbiturate dose. Since the train length is not changing, PLP parallels the plateau level decrease and TLP decreases due to a lengthening of the respiratory cycle. The first, minimum and last intervals show variable fluctuations while N50, N70 and N90 parameters appear to be independent of pentobarbital effects. The N100 measurement, however, parallels the spikes per train and generally decreases at higher drug levels. A similar directional change for N100 is observed for inspiratory cell, C81UI5L (Figures 54 and 55) recorded when vagal afferents were removed. The slope pattern for this expiratory cell is very irregular and yields no useful information. Comparison of these parameter responses to the mean population data in Figures 35-43 reveals many

individual differences.

Another expiratory cell, C82UE3R, was examined during morphine administration as shown in Figures 60 and 61. Pattern shifts induced with morphine were considerably different from those seen in expiratory cells during barbiturate administration (Figures 56-59). After a small decrease in respiration rate from 29.4 to 24.0 breaths per minute, the rate was facilitated up to 53.8 with cumulative doses of morphine. This unexpected species-specific result may be attributed to sensitization of pulmonary receptors. A true depression of the discharge frequency is observed as shown by the shift to the right in the histogram distribution (Figure 60A) and the increased mean, median and mode times (Figure 61B). Figure 61 shows that most parameters are decreased by morphine. These effects are in direct opposition to those seen during pentobarbital depression of expiratory cells, but strict comparisons must be avoided since respiration rate alterations with each drug are going in reverse directions. These data indirectly demonstrate that morphine and pentobarbital probably act via different mechanisms. This is a significant observation worthy of further exploration.

Finally, the quantitative effects of vagotomy on



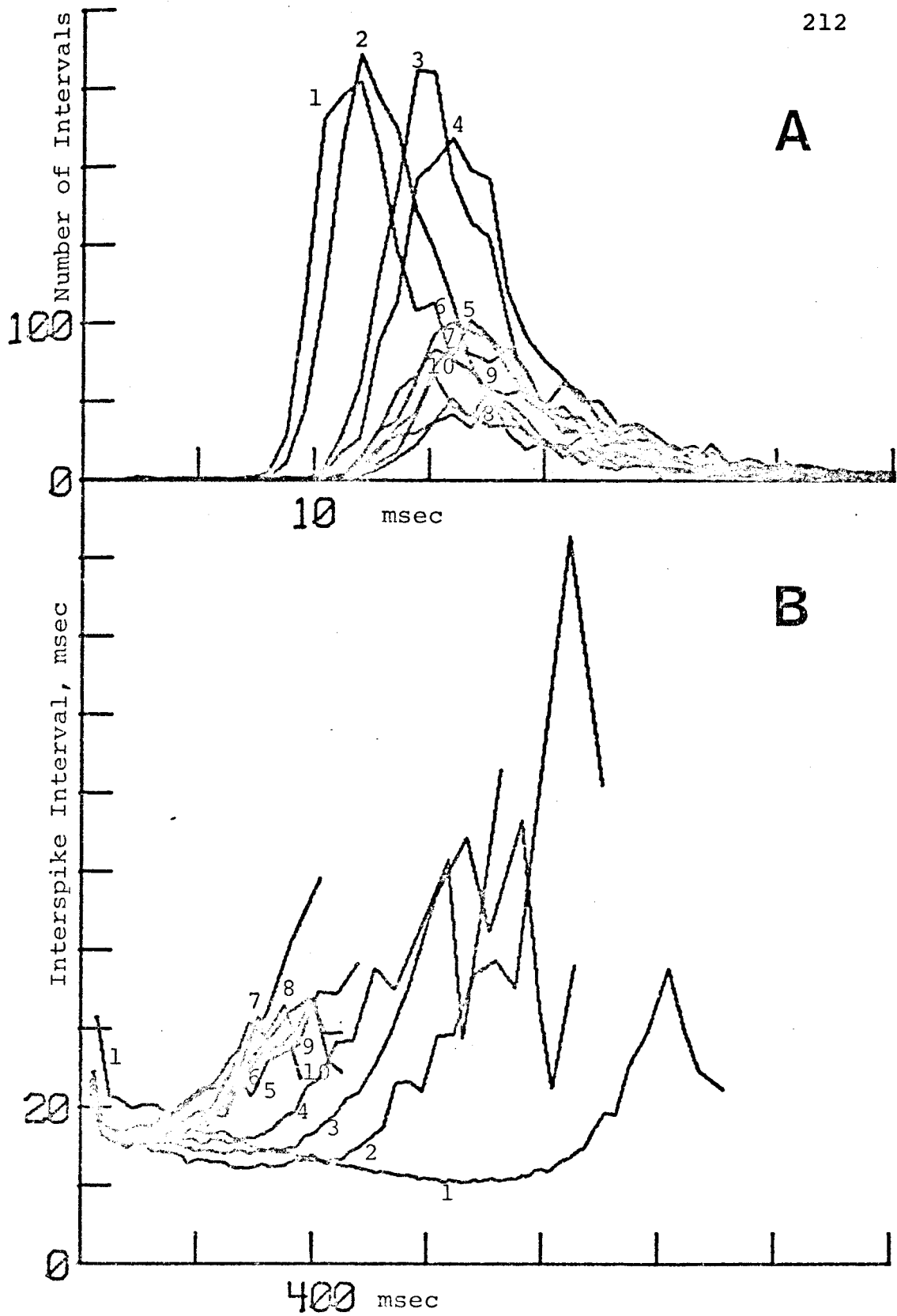


Figure 60. C82UE3R. Effect of cumulative doses of morphine sulphate on histogram (A) and interspike interval modulation curve (B). Vagi intact.

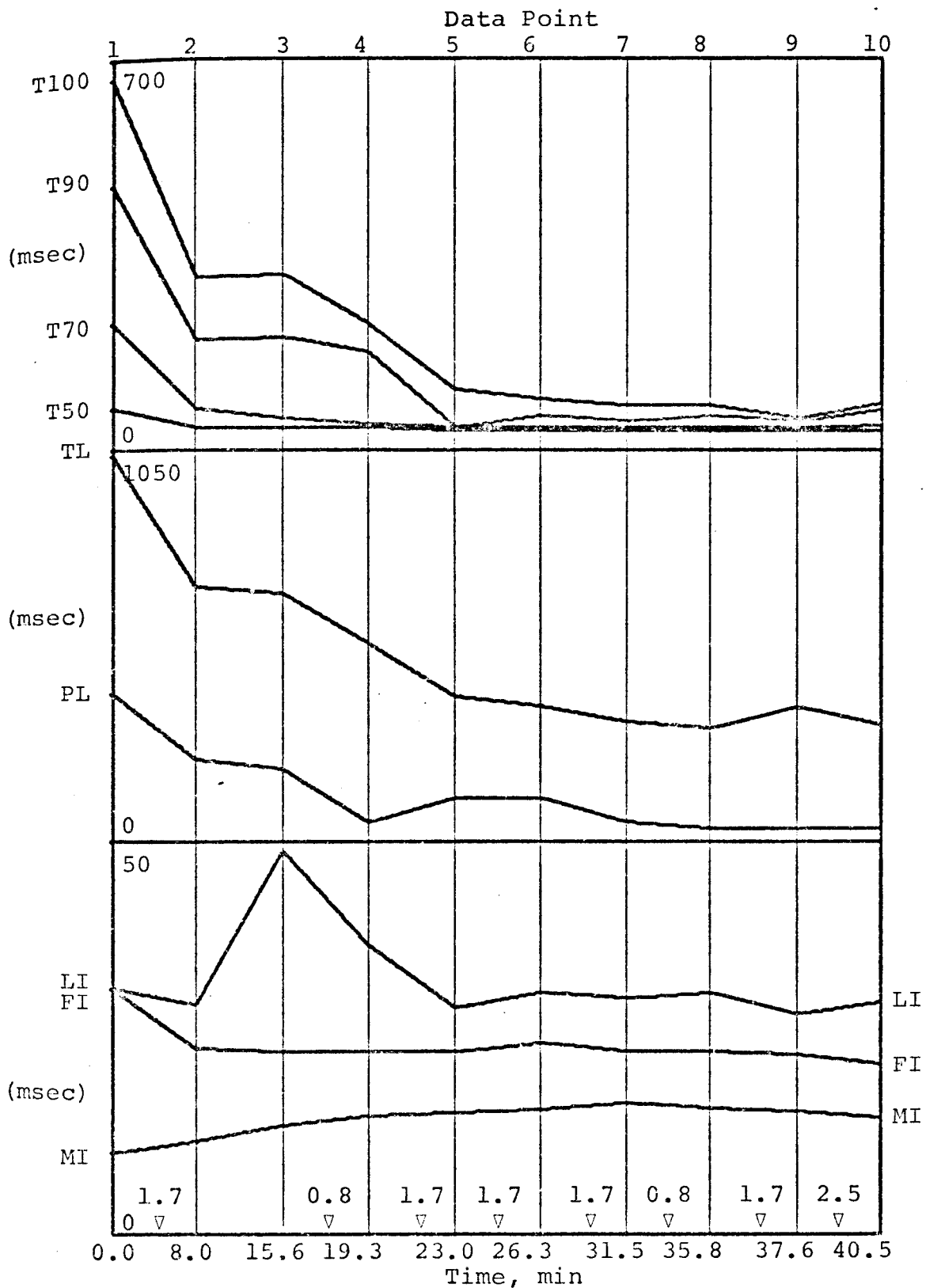


Figure 61A. C82UE3R. Effect of morphine on parameters of expiratory cell discharge. See text for details.

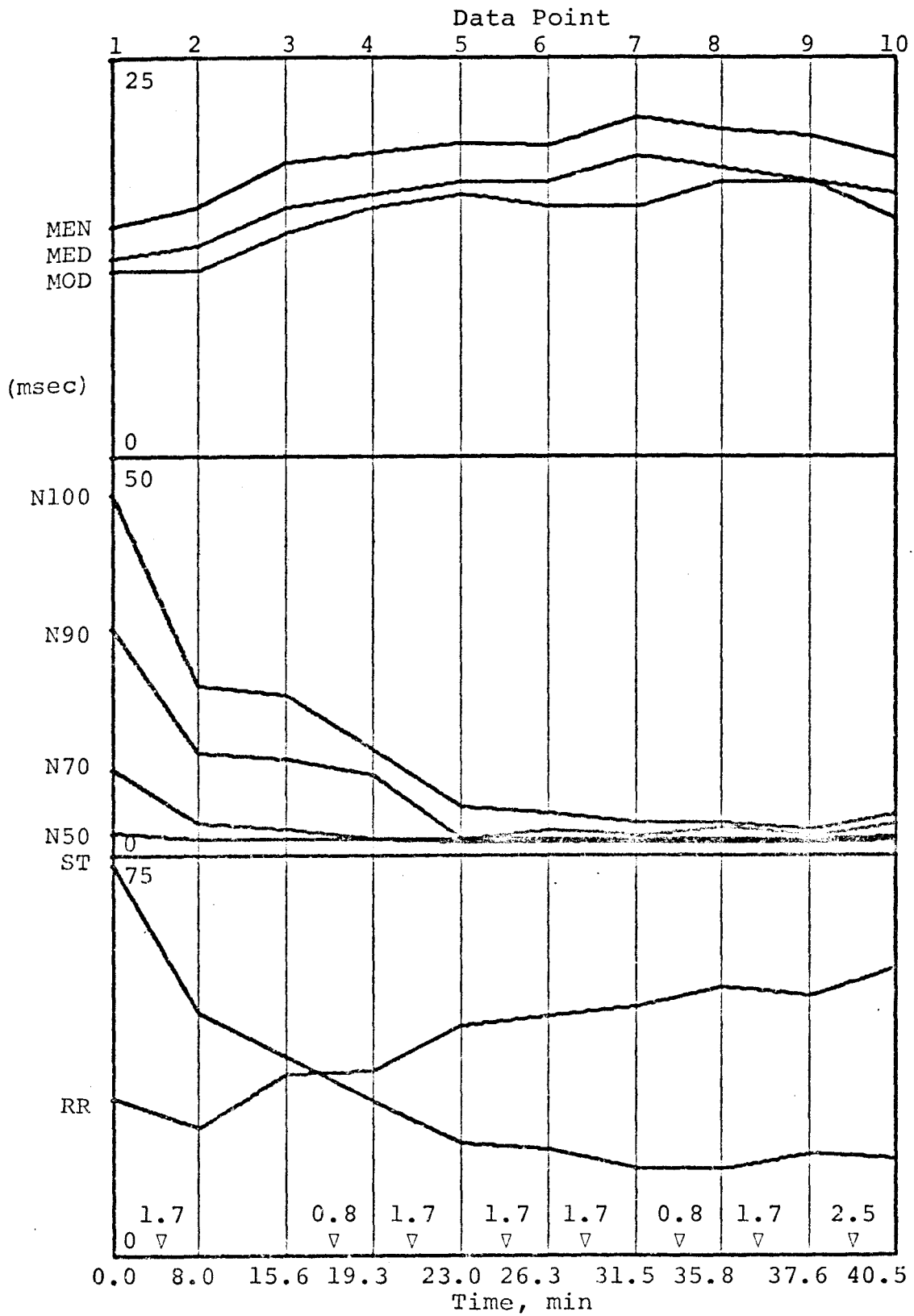


Figure 61B. C82UE3R. Effect of morphine on parameters of expiratory cell discharge. See text for details.

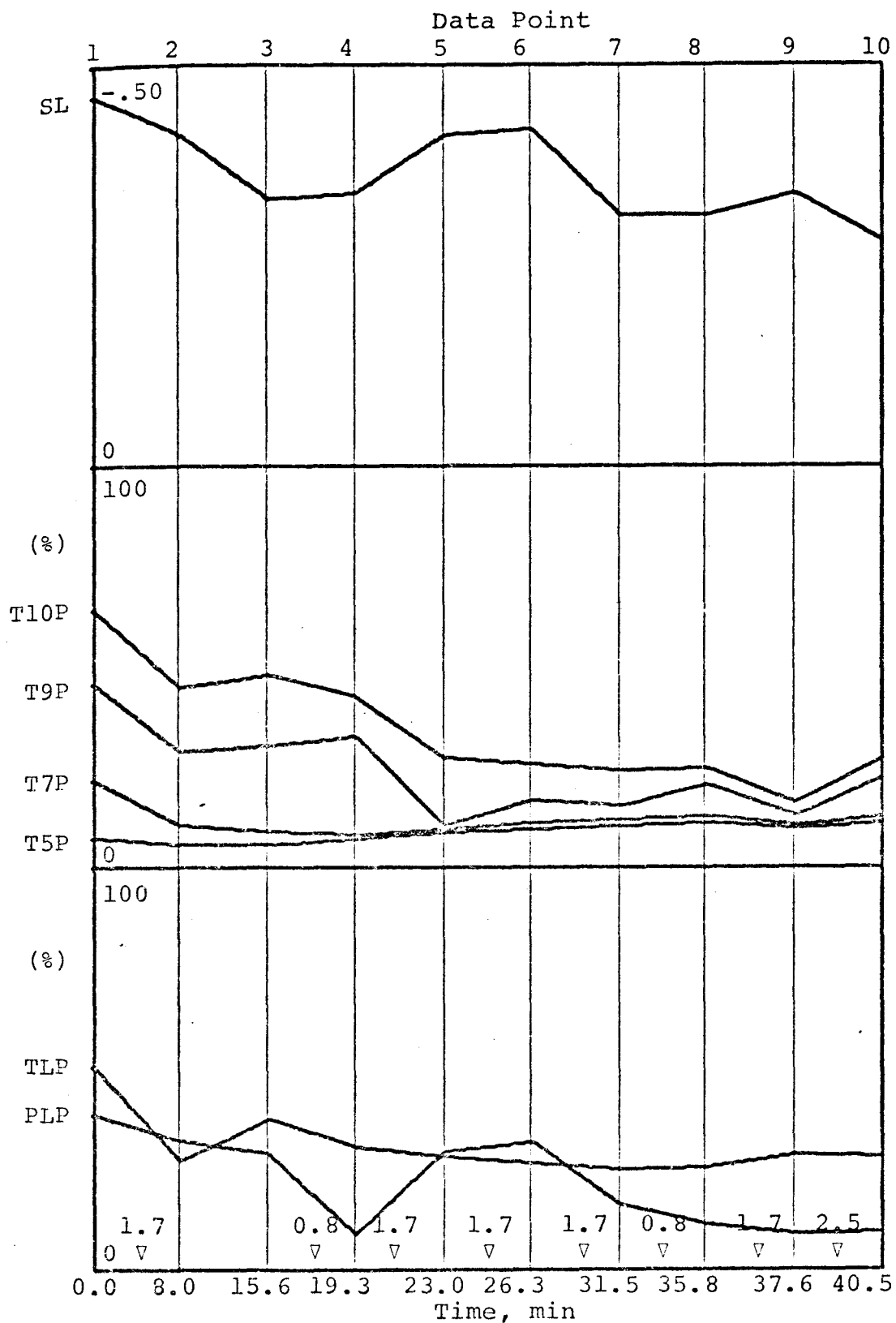


Figure 61C. C82UE3R. Effect of morphine on parameters of expiratory cell discharge. See text for details.

expiratory cell C68UE6R are shown in Figure 62. The first steady state measurement (Curve 1) is a prevagotomized control. One half a minute after cutting the left vagus nerve, a new steady state discharge was attained. The histogram was shifted to the right (Figure 62A) and the modulation curve shifted upward (Curve 2). After an additional 1.7 minutes, the curves assumed position 3. These shifts corresponded to an increase in the mode time from 10.8 to 19.6 to 30.8 milliseconds. This was associated with small respiratory rate decreases from 65.3 to 59.4 to 57.0 breaths per minute. The discharge pattern was analyzed up to 4.8 minutes after the initial unilateral vagotomy, but did not differ significantly from Curve 3. However, subsequent sectioning of the right vagus immediately terminated the expiratory discharge. Since the cell was anatomically located in the right medulla, it is postulated that the right vagal afferents were responsible for the activation of this expiratory unit. Removal of these afferents inhibited unit discharge. These vagotomy effects on an expiratory cell can be compared to the response of inspiratory cell C81UI5L in Figures 54 and 55 subjected to similar procedures.

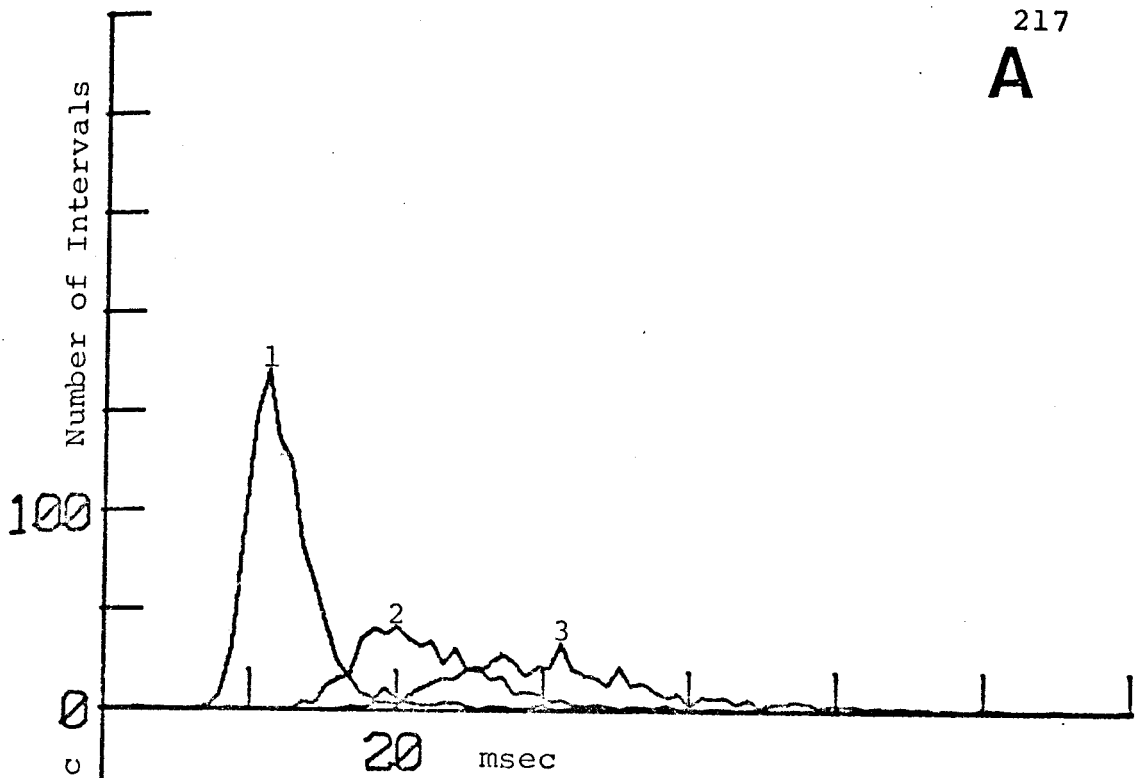
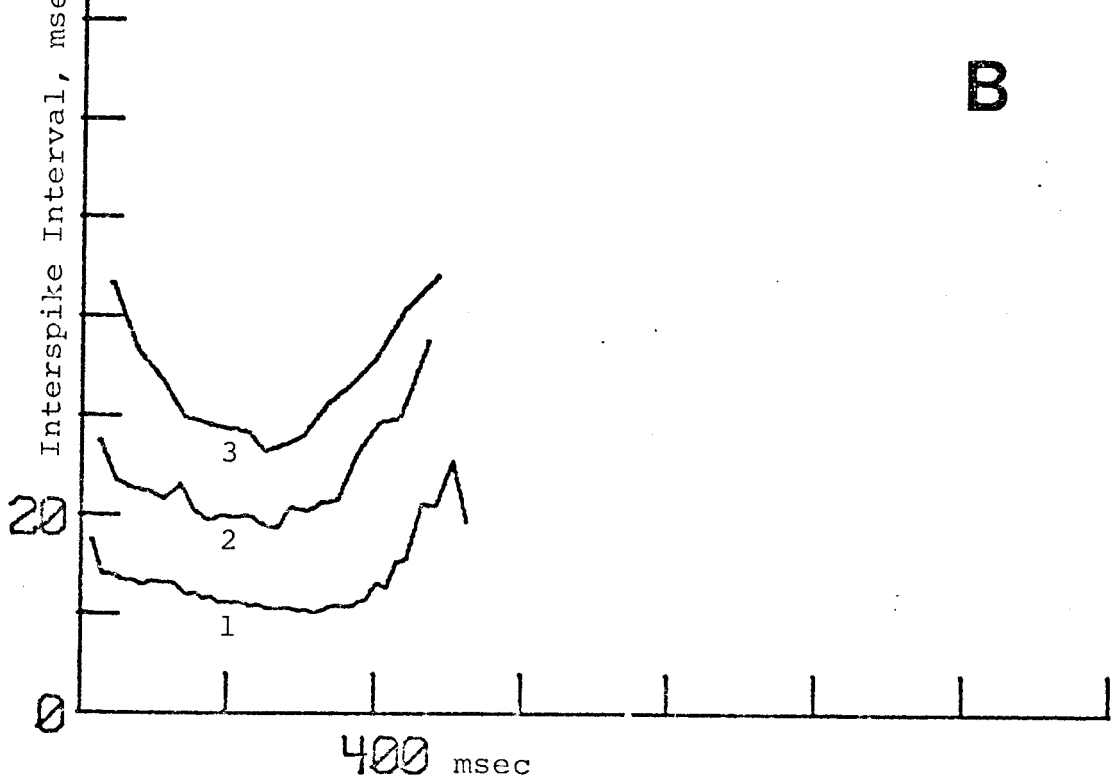
**A****B**

Figure 62. C68UE6R. Histogram (A) and interspike interval modulation curves (B) before (Curve 1) and after (Curves 2 and 3) unilateral left vagotomy.

D. Barbiturate Apneusis

During single cell depression studies, it was frequently observed that apneustic breathing patterns could be induced after summation of several serial injections of a barbiturate (pentobarbital or thiopental sodium). This well documented phenomenon (cf. Literature Review) was further investigated to facilitate interpretation of the mean population regressions (Figures 11-43) and single cell data (Figures 50-59) previously presented.

Classically, apneusis in the cat can be produced by surgical removal of the rostral pons (pneumotaxic area) coupled with bilateral vagotomy. This is illustrated in Figure 63. Previous to this record, the cerebellum of cat 12 was removed and the brainstem was transected at the level of the inferior colliculi. The cat continued to breathe rhythmically at about five breaths per minute and this breathing pattern was not altered significantly by unilateral vagotomy on the left side. Subsequent section of the right vagus nerve, however, initiated a sustained inspiratory effort as seen in the intrapleural pressure tracing (IPP). Clearly, pneumotaxic and vagal mechanisms play important roles in the maintenance of eupneic respiration. The literature also implicates the cerebellum as an additional source of inspiratory inhibition.

## C12 IC TRANSECTION

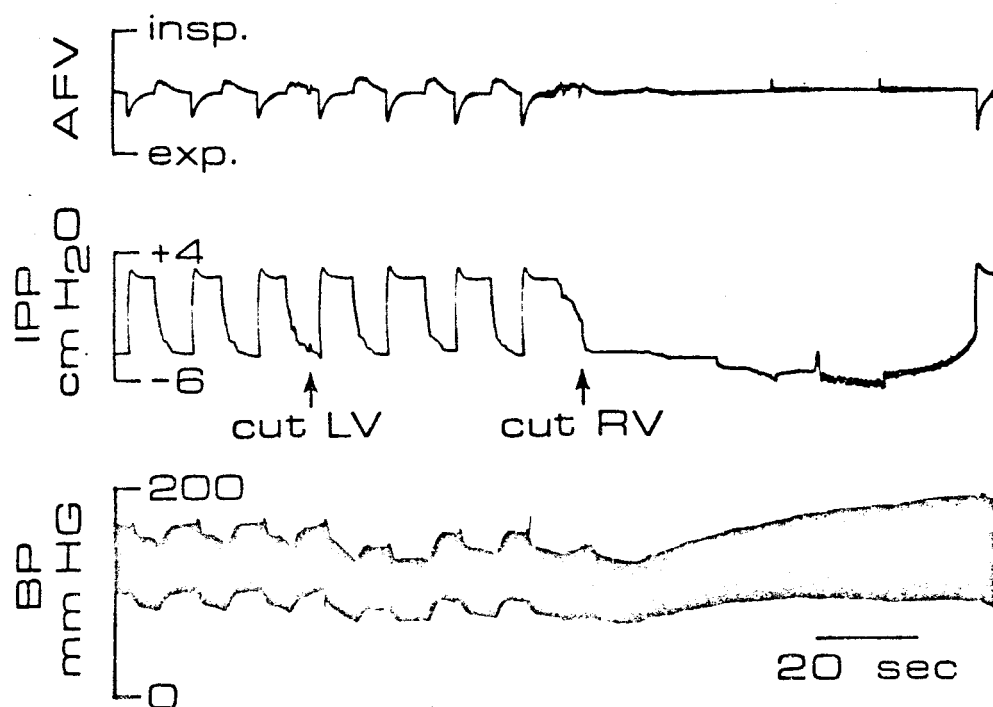


Figure 63. Genesis of classical apneusis in the cat. Prior to the beginning of record an inferior collicular transection was made. The left (LV) and right (RV) vagus nerves were sectioned at times indicated by arrows. Tracings top to bottom: air flow velocity (AFV), intrapleural pressure (IPP) and systemic blood pressure (BP).



Figure 64 shows a representative example of barbiturate induced apneusis in a cat with intact brainstem, cerebellum and vagi. At the arrow, 3.4 mg/kg pentobarbital was injected intravenously, raising the total dosage of barbiturate to 57.6 mg/kg, including the initial anesthetic dose of 30.0 mg/kg. After a short circulation delay, a prolonged inspiratory hold of more than 30 seconds was observed. A normal respiratory pattern then emerged, but the system was operating at a higher end-expiratory % CO<sub>2</sub> level. It is possible that this elevated CO<sub>2</sub> drive was responsible for the return to cyclicity. This example illustrates that pentobarbital can in some way block pneumotaxic, vagal and cerebellar inhibitions on inspiratory drive with the resultant genesis of apneusis. Similar results were found using thiopental.

Supportive evidence that the cerebellum is inhibitory to inspiration is given in Figure 65. The control respiratory pattern in panel A was obtained from cat 49 which had brainstem, cerebellum and vagi all intact. Between panels A and B the cerebellum was quickly removed by suction with very little blood loss. Immediately after cerebellectomy the eupneic respiration was transformed into one of apneusis with elevated end-expiratory % CO<sub>2</sub> values (panel B). Eight minutes later (panel C) the

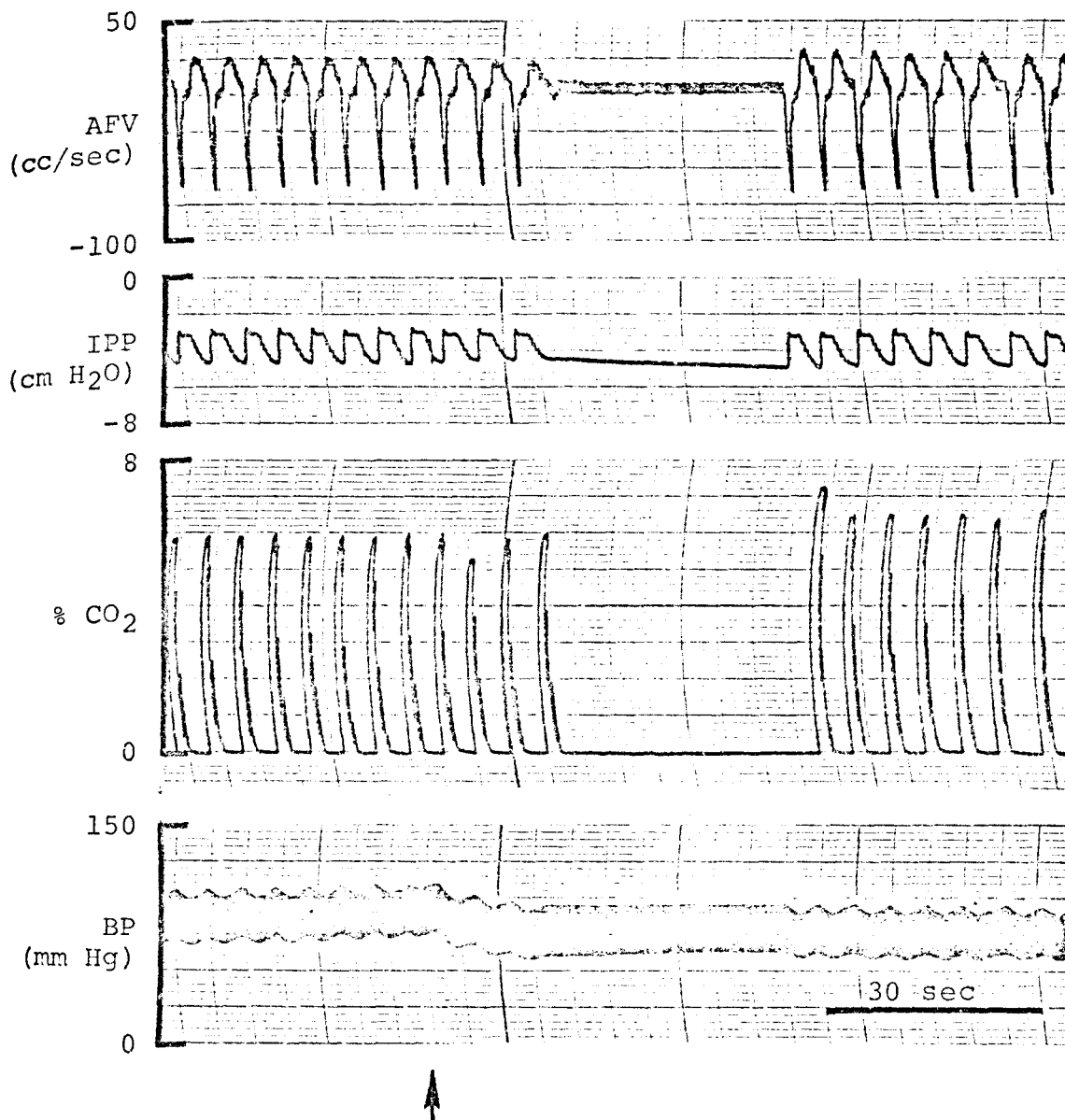


Figure 64. Pentobarbital apneusis. Cerebellum and vagi intact. Injection of 3.4 mg/Kg pentobarbital (arrow) induced a prolonged apneustic breath. Tracings as in Figure 63 with the addition of end-expiratory % CO<sub>2</sub> recording in third panel from top.

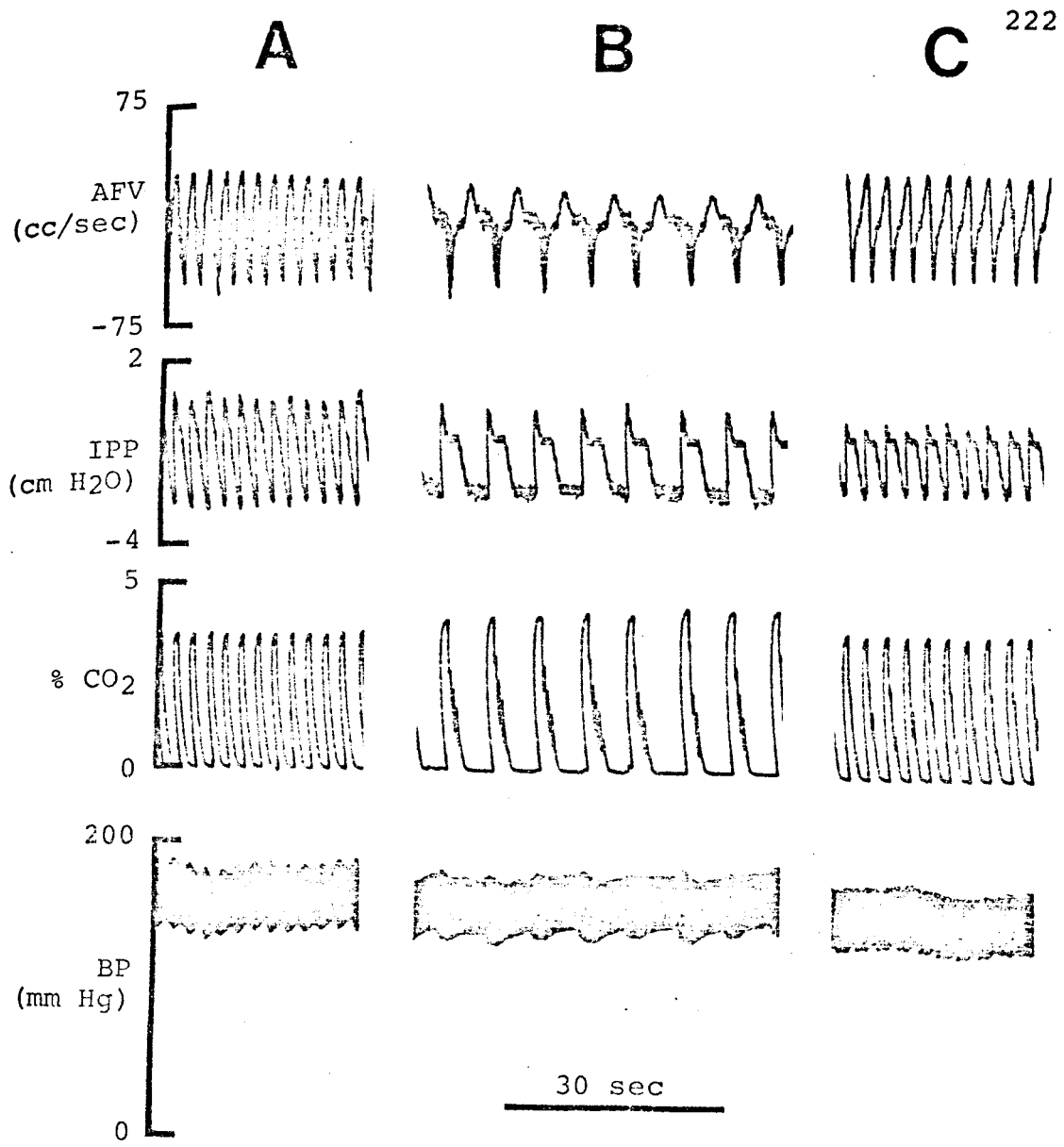


Figure 65. Apneustic pattern produced by acute cerebellectomy: A. control; B. three minutes later after cerebellectomy; C. recovery after eight minutes. Tracings as in Figure 64.

apneustic breaths were replaced by a normal respiratory pattern and the % CO<sub>2</sub> returned to control levels. Evidently the system is capable of readjustment, suggesting that cerebellar inhibition is of secondary importance (76).

Figure 66 summarizes apneustic threshold data for four different procedures. The apneustic threshold was defined as the amount of barbiturate required to induce the first apneustic breath under various experimental conditions. The total accumulated dose was expressed as the incremental amount over and above the initial anesthetic level of 30 mg/kg pentobarbital. For example, successive small increments of thiopental (A) and pentobarbital (B) were administered until the first apneustic breath was produced (cf. Figure 64). Approximately twice the dosage of pentobarbital was required to reach the apneustic threshold in cats with brainstem, cerebellum and vagi intact. The difference between pentobarbital and thiopental threshold doses was significant at the 0.04 level. This was probably due to the fast acting characteristics of thiopental, a short acting barbiturate that is rapidly absorbed.

In C, drug data are pooled for apneustic breaths induced by vagotomy at low pentobarbital levels and from

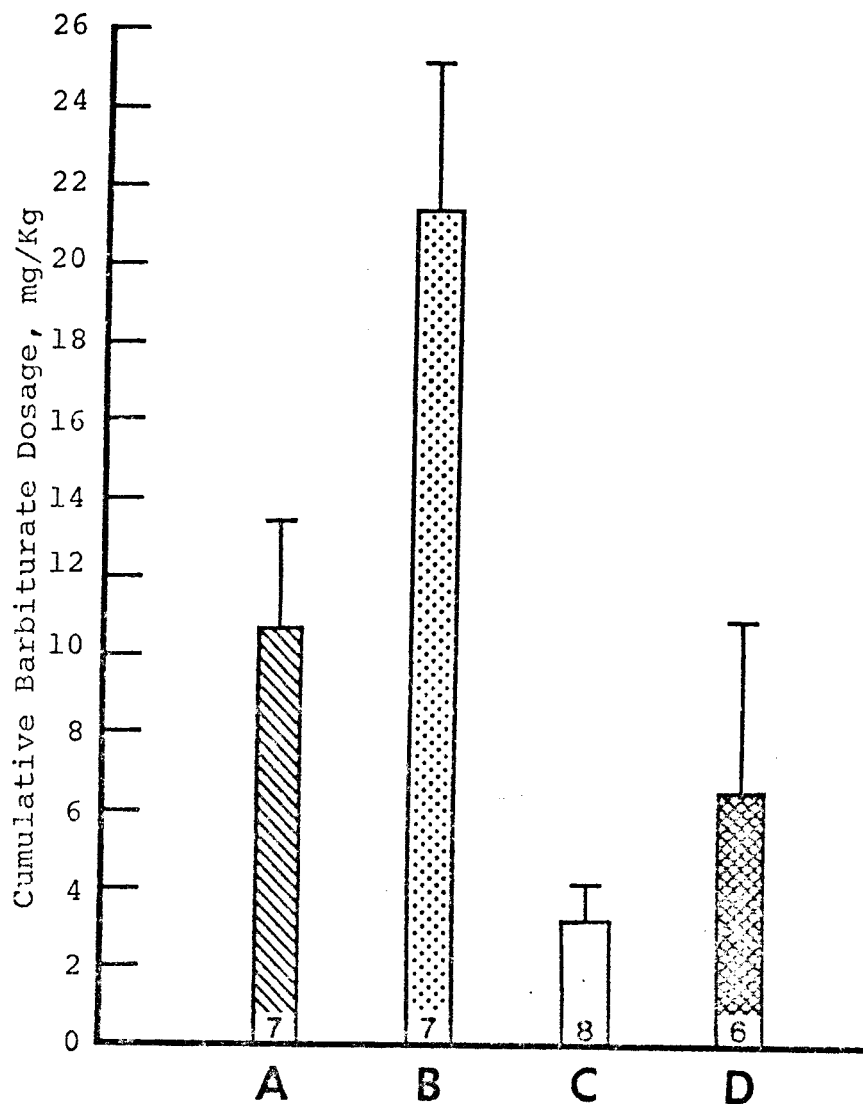


Figure 66. Mean and SEM of barbiturate dosage at apneustic threshold: A. thiopental; B. pentobarbital; C. pentobarbital after vagotomy only; D. pentobarbital after cerebellectomy only. Levels of significance: A vs. B,  $P < 0.04$ ; B vs. C,  $p < 0.0005$ ; C vs. D,  $p > 0.4$ ; B vs. D,  $p < 0.03$ . Ordinate expressed as cumulative dosage over and above initial anesthetic dose of 30 mg/Kg pentobarbital.

pentobarbital titration studies in vagotomized cats failing to show long inspiratory holds following vagal section. In the former group, the apneustic breath observed was not believed to be due to mechanical stimulation of the vagus, since this generally leads to expiratory apnea. Comparison of B and C shows that it was significantly easier ( $p < 0.0005$ ) to induce apneusis at lower pentobarbital levels when vagal inhibition was absent.

Finally, the cerebellum was removed in a group of cats with intact brainstem and vagi. The pentobarbital level to induce apneustic breaths (cf. Figure 65) in this group is shown in D. Comparison of B and D shows that cerebellectomy significantly decreased ( $p < 0.03$ ) the amount of pentobarbital required to induce sustained inspiratory holds in brainstem and vagi intact cats. Although there is no significant difference ( $p > 0.4$ ) between C and D, it cannot be concluded that vagal and cerebellar inhibitions are of equal strength because of the nature of these experiments. That is, apneusis often resulted immediately after vagotomy or cerebellectomy procedures indicating that the apneustic threshold was already exceeded by an unknown amount of barbiturate.

In conclusion, these results show that relatively large doses of barbiturate are required to block major

inhibitions on inspiration before apneusis is produced. However, partial reduction of inhibitory input (vagotomy or cerebellectomy) significantly decreases this barbiturate dose requirement. This implies that apneusis depends on the relative balance between excitatory and inhibitory inputs to the brainstem inspiratory cells.

Figure 67 shows the effect of vagotomy on cat 63 previously subjected to pentobarbital apneusis (17.8 mg/kg incremental dose) and cerebellectomy (22.2 mg/kg pentobarbital level). Thirty minutes after the latter procedure, the cat had recovered to the prevagotomy control pattern in Figure 67. At this point, section of the left vagus immediately prolonged the average inspiratory to expiratory time ratio and subsequent section of the right vagus further increased this ratio. At least two conclusions can be drawn from these results. First, the length of apneustic holds is directly related to the degree of blockade of inspiratory inhibitory systems. Second, a high pentobarbital level may block brainstem interconnections, since the control pattern was lost after unilateral vagotomy on the left side, a response which did not occur at lower pentobarbital levels (Figure 63).

Barbiturate induced apneusis was examined at the

C63

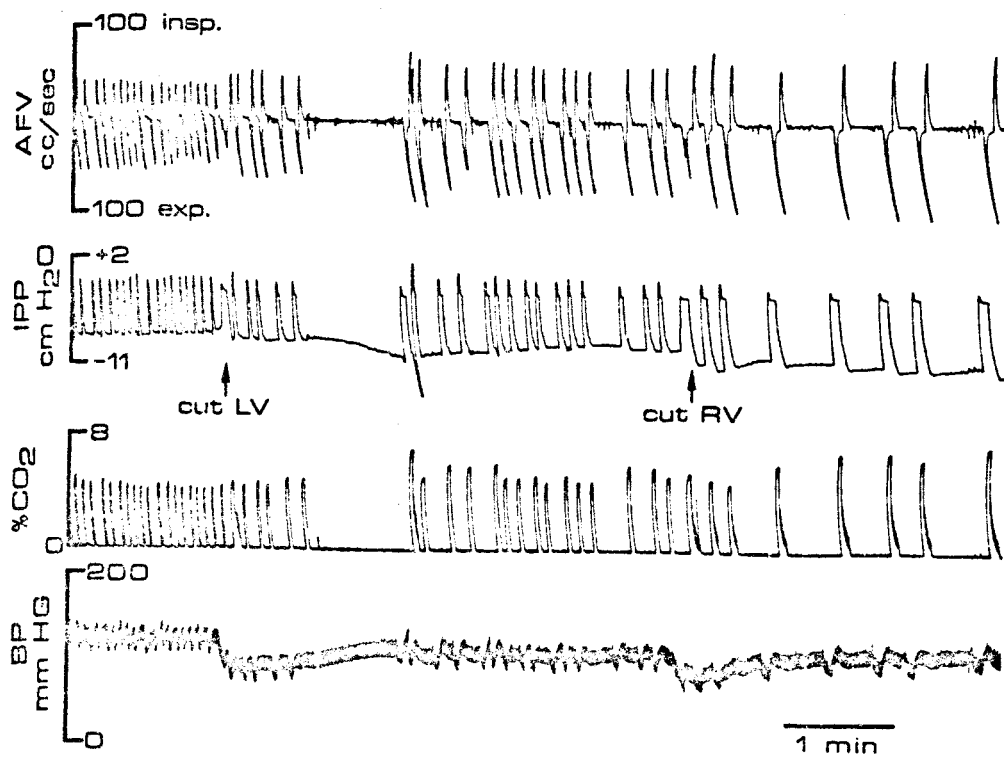


Figure 67. Effect of sequential left (LV) and right (RV) vagotomy on respiratory patterns. Cat previously subjected to cerebellectomy and high doses of pentobarbital. Tracings as in Figure 64.



single cell level. In Figure 68, an inspiratory cell was found to fire continuously during a sustained inspiratory hold induced by pentobarbital administration. An expiratory cell subjected to a similar procedure was inhibited throughout the apneustic breath as shown in Figure 69. These observations support the idea that inspiratory and expiratory cells are reciprocally inhibitory.

In another study, shifts in the histogram interspike interval distribution of a single inspiratory cell were followed after several thiopental injections as shown in Figure 70. In this preparation, which had intact brainstem, cerebellum and vagi, there was a consistent decrease in the respiration rate from 57 to 15 breaths per minute. However, an interesting triphasic shift in the histogram was observed as previously shown for another inspiratory cell during pentobarbital accumulation in Figures 52 and 53. For example, at low incremental thiopental doses (0-4 mg/kg) the histogram was first shifted to the right as the unit was deactivated (Figure 70, top panel). At medium thiopental levels (4-7 mg/kg), the unit became activated causing a shift in the histogram to the left (Figure 70, middle panel). Finally, at high thiopental levels (7-9 mg/kg), the inspiratory cell experienced a second depression

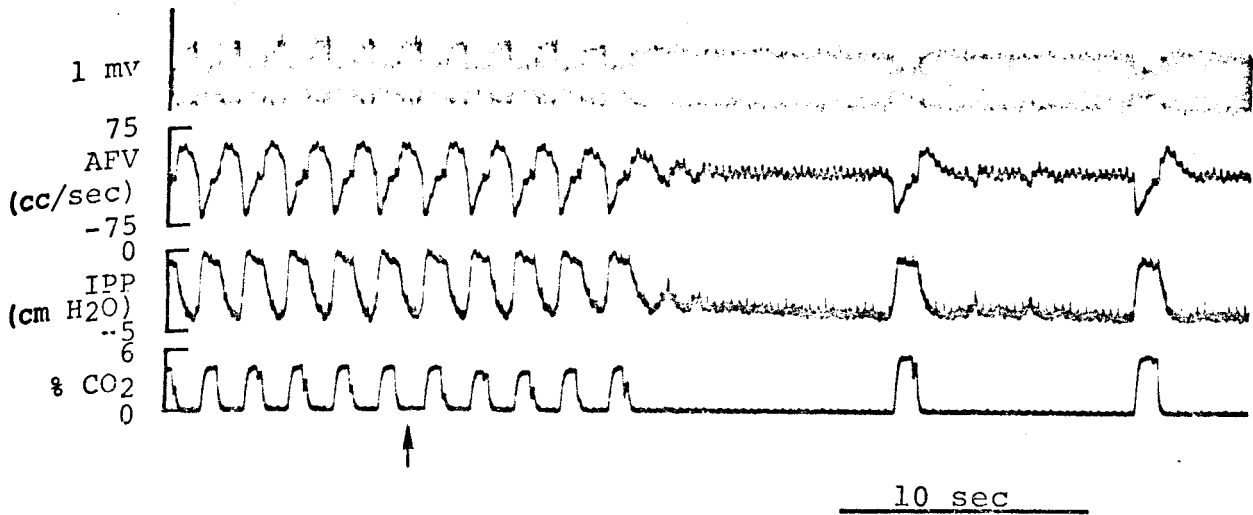


Figure 68. C50UI5R. Single inspiratory cell recording during induction of apneusis after administration of 1.0 mg/Kg pentobarbital (arrow). Tracings top to bottom: unit activity, air flow, intrapleural pressure and end-expiratory % CO<sub>2</sub>.

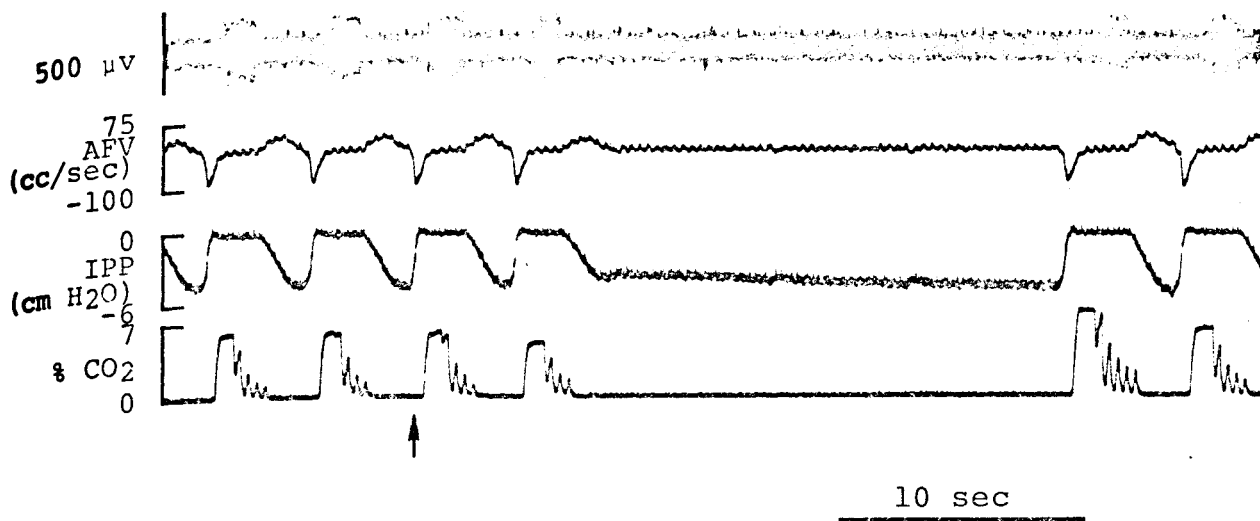


Figure 69. C64UE5R. Single expiratory cell recording during induction of apneusis after administration of 3.0 mg/Kg pentobarbital (arrow). Tracings as in Figure 68.

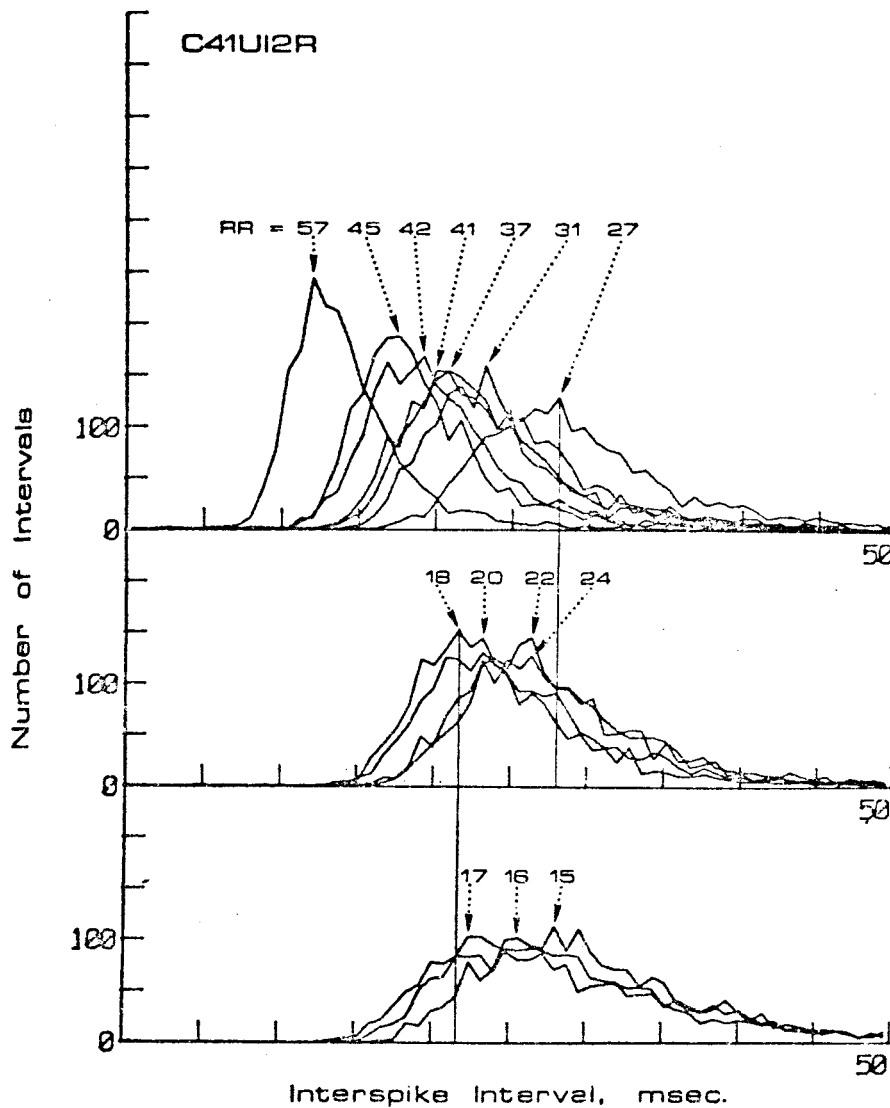


Figure 70. Changes in single inspiratory cell histogram during progressive depression of respiration rate (RR) produced by serial thiopental injections. Corresponding respiration rates and histogram modes are indicated by arrows.

trend and the histogram shifted back to the right (Figure 70, bottom panel). Apneusis was never produced.

Explanation of this response most likely involves differential thiopental depression of inhibitory versus excitatory inputs to the medullary inspiratory mechanism. That is, at medium thiopental levels (4-7 mg/kg), a significant fraction of total inhibition could be sufficiently depressed by the barbiturate, thereby allowing inspiratory unit activation. This would result from pneumotoxic, cerebellar or vagal blockade, or partial depression of each by thiopental.

The mode, median and mean interval times from each histogram distribution in Figure 70 are plotted as a function of thiopental dose and respiration rate in Figure 71A and B respectively. The dose is expressed as the incremental level over the initial anesthetic dose of 30 mg/kg pentobarbital. Similar data from expiratory cell C47UE10R are included for comparison. The triphasic shift in interval times for inspiratory cell C41UI2R are clearly seen in each plot. Qualitatively speaking, a mirror image triphasic shift can also be observed for expiratory cell C47UE10R. This was previously seen for expiratory cell C71UE2R (Figures 58 and 59). For example, when the inspiratory unit was depressed, the expiratory

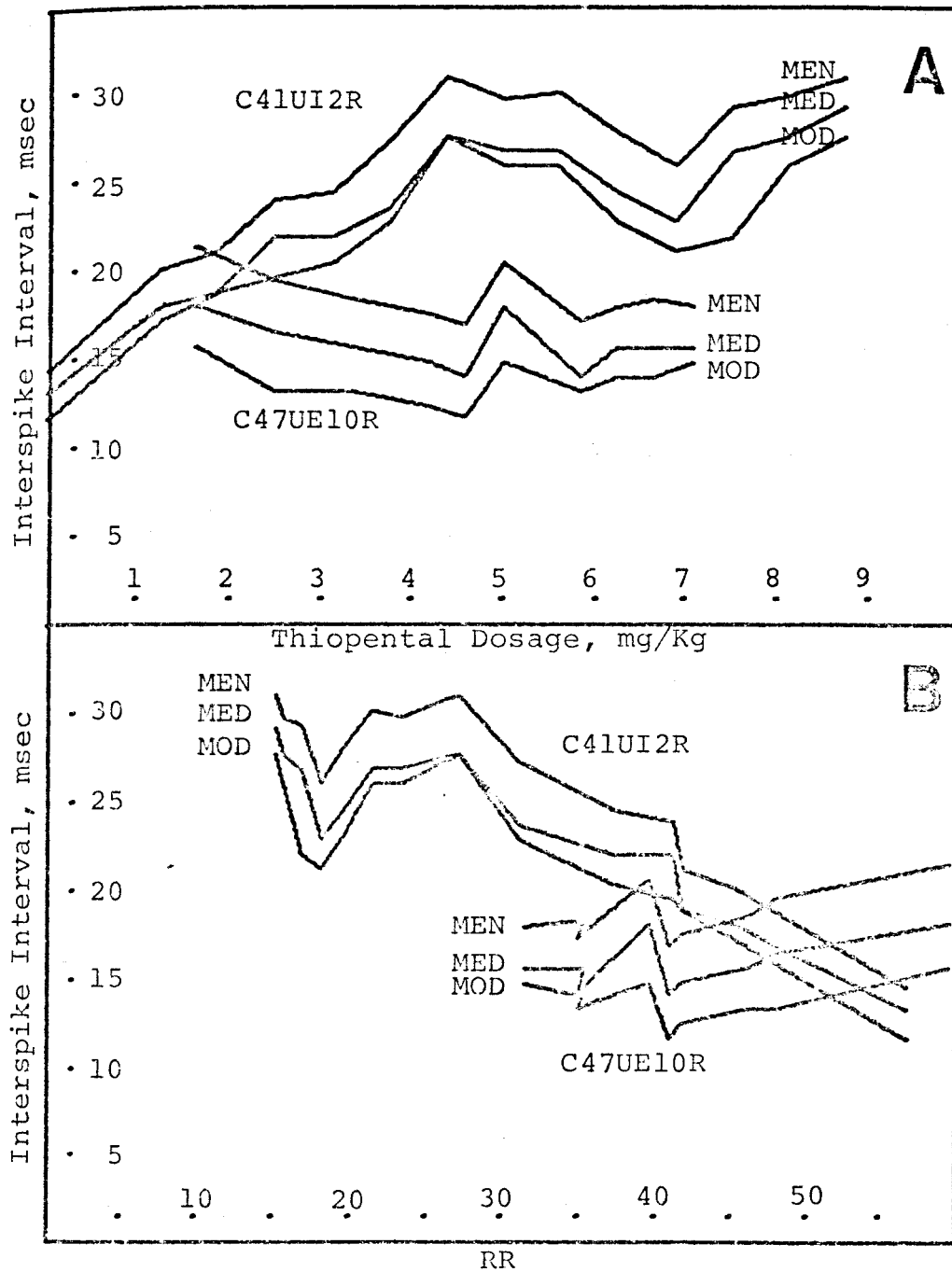


Figure 71. Histogram data for cell C41UI2R (Figure 70) replotted to show changes in mean (MEN), median (MED) and mode (MOD) as a function of thiopental dosage (A) and respiration rate (B). Expiratory cell C47UE10R plotted for comparison.

unit was activated and vice versa. This implies a negative reciprocal relationship between firing frequencies of inspiratory and expiratory cells.

An attempt was made to determine if pentobarbital could differentially block pneumotoxic regions before medullary areas. The cerebellum was removed from cat 65 and an electrode was inserted into the classical pneumotoxic region of the rostral pons. The vagi were intact. Figure 72 shows the frequency discharge response of three constantly firing (non-respiratory) cells plotted as a function of time before and after pentobarbital administration. Cell K1, the first cell found, was completely inhibited by a 2.7 mg/kg injection of pentobarbital which brought the total sum dose over the anesthetic level to 4.0 mg/kg. Another cell, K2 was soon found which had a discharge frequency approximately half that of cell K1. After a 2.7 mg/kg pentobarbital injection, the discharge frequency was sharply inhibited, but rebounded back to the control level in less than a minute. This unit could not be completely inhibited by two additional pentobarbital injections (3.3 and 2.7 mg/kg). Finally, a third constantly firing cell (K3) was located in the upper pons. The firing frequency of this unit was very low presumably due to the high pentobarbital

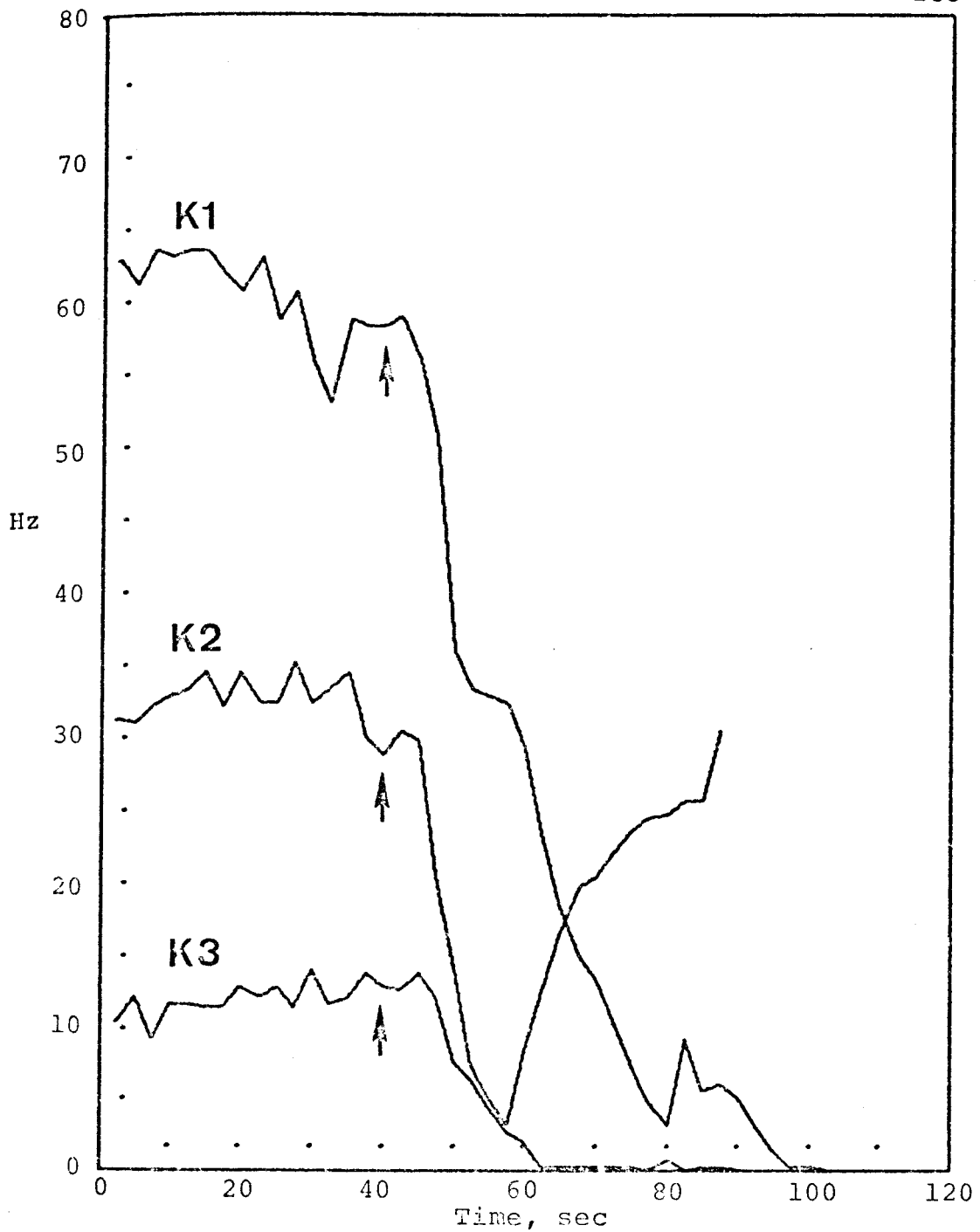


Figure 72. Effect of pentobarbital (arrows) on discharge frequency of three non-respiratory pontine cells. See text for details.



level (12.7 mg/kg). When 3.3 mg/kg pentobarbital was given, this cell was completely inhibited like cell K1 and a tendency toward apneusis was observed. Subsequent searching for non-specific units in the rostral pons was not successful and it was concluded that this region of the brainstem was deeply depressed by the total sum level of pentobarbital present (16.0 mg/kg). Replacement of the same recording electrode in the medullary architecture, however, immediately yielded unit activity. Soon an inspiratory cell was located which correlated with the oscillating peripheral parameters. This demonstrated two points. First, the failure to find single cell discharges in the pons could not be attributed to microelectrode failure. Second, the medulla was not as highly depressed as the rostral pons at an incremental pentobarbital level of 16.0 mg/kg. This information suggests that the inflection of the mode curve for inspiratory cell C41UI2R in Figure 71 may be due to depression of pneumotaxic circuits by pentobarbital.

In final studies, miscellaneous procedures were performed to induce apneusis in the spontaneously breathing cat. Figure 73 illustrates the effect of bilateral carotid occlusion (BCO) on modification of the breathing pattern at different pentobarbital levels. In this

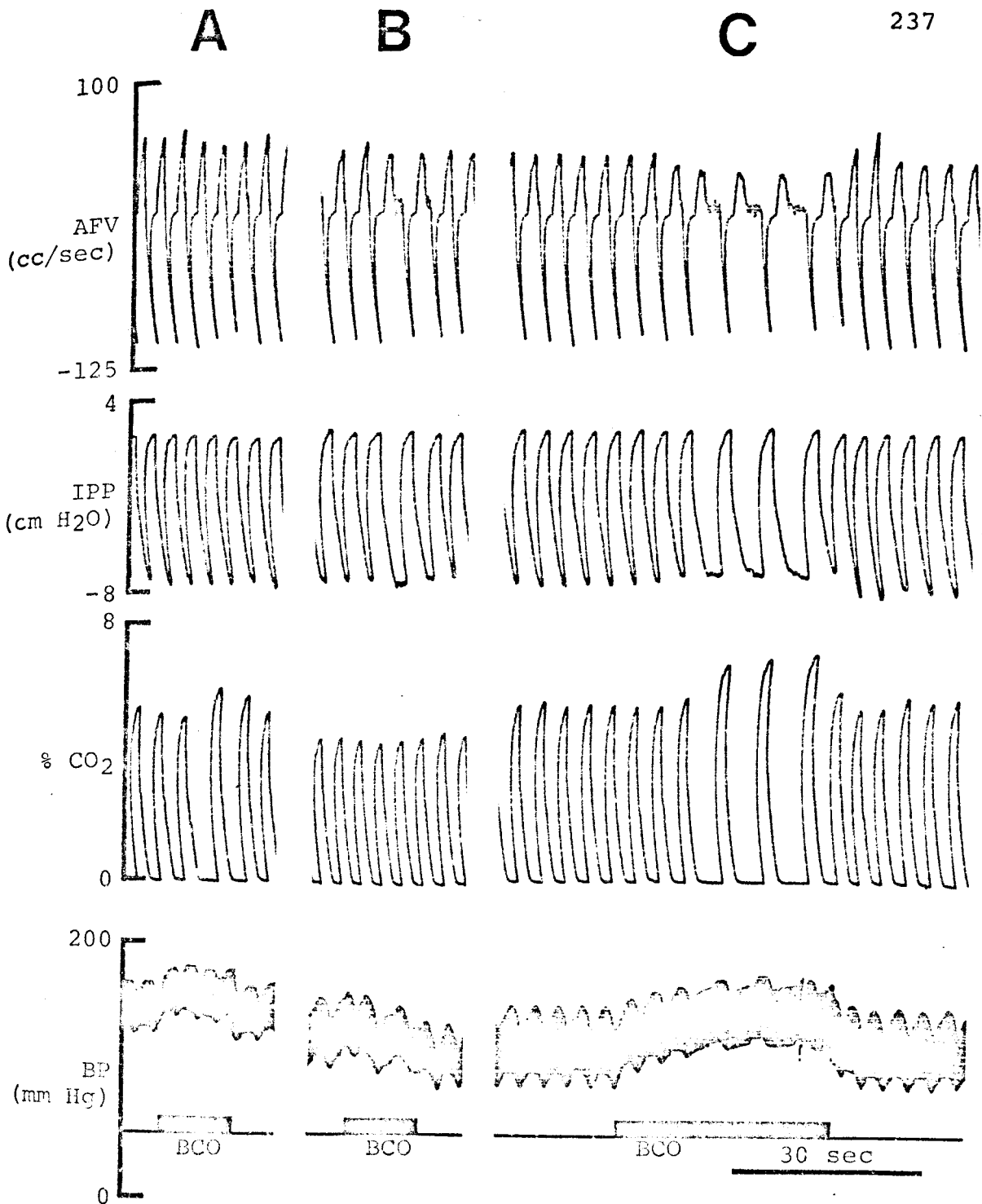


Figure 73. Changes in respiratory patterns during bilateral carotid occlusion (BCO) in a vagotomized cat with increasing pentobarbital dosage: A. after 31.0 mg/Kg; B. after 32.8 mg/Kg; C. after 34.5 mg/Kg. Tracings as in Figure 64.

preparation, vagi were cut but the cerebellum was still intact. As the total barbiturate level was elevated from 31.0 to 32.8 to 34.5 mg/kg in panels A, B and C, respectively, there was an increased tendency toward apneusis during each BCO procedure. This demonstrates that a change in inhibitory input to respiratory centers over carotid sinus afferents (baroreceptor or chemoreceptor) can induce apneusis provided the barbiturate depression is at an appropriate level.

Similar results were found in another cat in which carotid sinus pressure was decreased via intravenous administration of histamine as shown in Figure 74. The vagi were cut and the cerebellum was intact as in Figure 73. The accumulated total dose of thiopental over the initial anesthetic dose of pentobarbital was 34.4 mg/kg. The effective dose was lower, since thiopental injections were given over a five hour period. After histamine injection there was a precipitous fall in systemic blood pressure from about 150 to 85 millimeters mercury due to peripheral vasomotor dilatation. A corresponding modification in the breathing pattern resulted with prolongation of the inspiratory duration resembling apneusis. As the blood pressure returned to a pre-injection control level over the following five

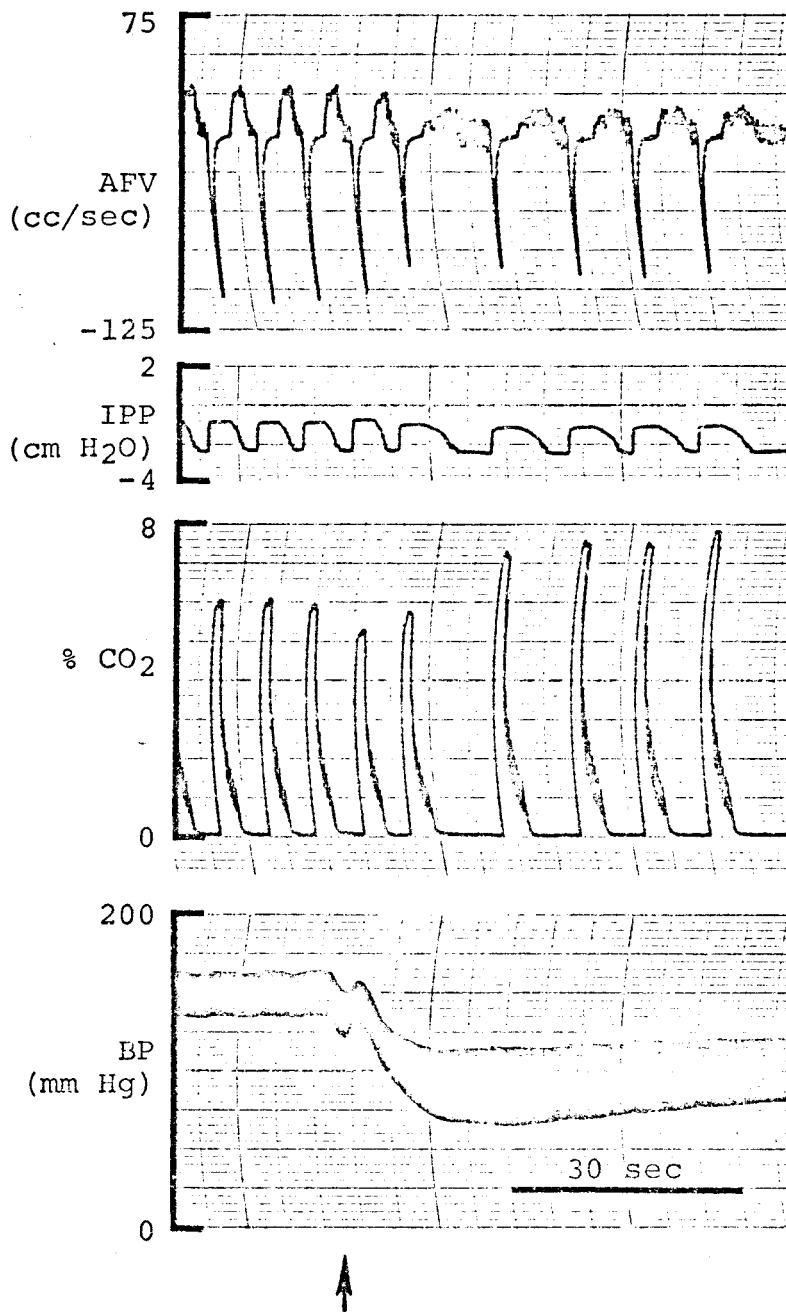


Figure 74. Changes in respiratory pattern produced by 17.0  $\mu$ gm histamine phosphate (arrow) in a vagotomized cat. Tracings as in Figure 64.

minutes, the respiratory pattern lost its apneustic form. These results substantiate the conclusions made for Figure 73; namely, a decrease in the inhibitions to the inspiratory mechanism can lead to apneusis when other inhibitory systems are partially blocked by barbiturate depression.

A final attempt was made to induce apneusis by facilitating excitatory inputs to the inspiratory complex. In Figure 75, doxapram, a respiratory stimulant, was given intravenously to a cat with severed vagi and intact cerebellum. The pentobarbital level was at 4.4 mg/kg over the initial anesthetic dose. Following a transient dip in the blood pressure, two prolonged apneustic breaths were produced. This response is consistent with the concept that inspiratory duration is a "simple" algebraic summation function of all excitatory and inhibitory inputs impinging on inspiratory cells in the medulla at any given point in time.

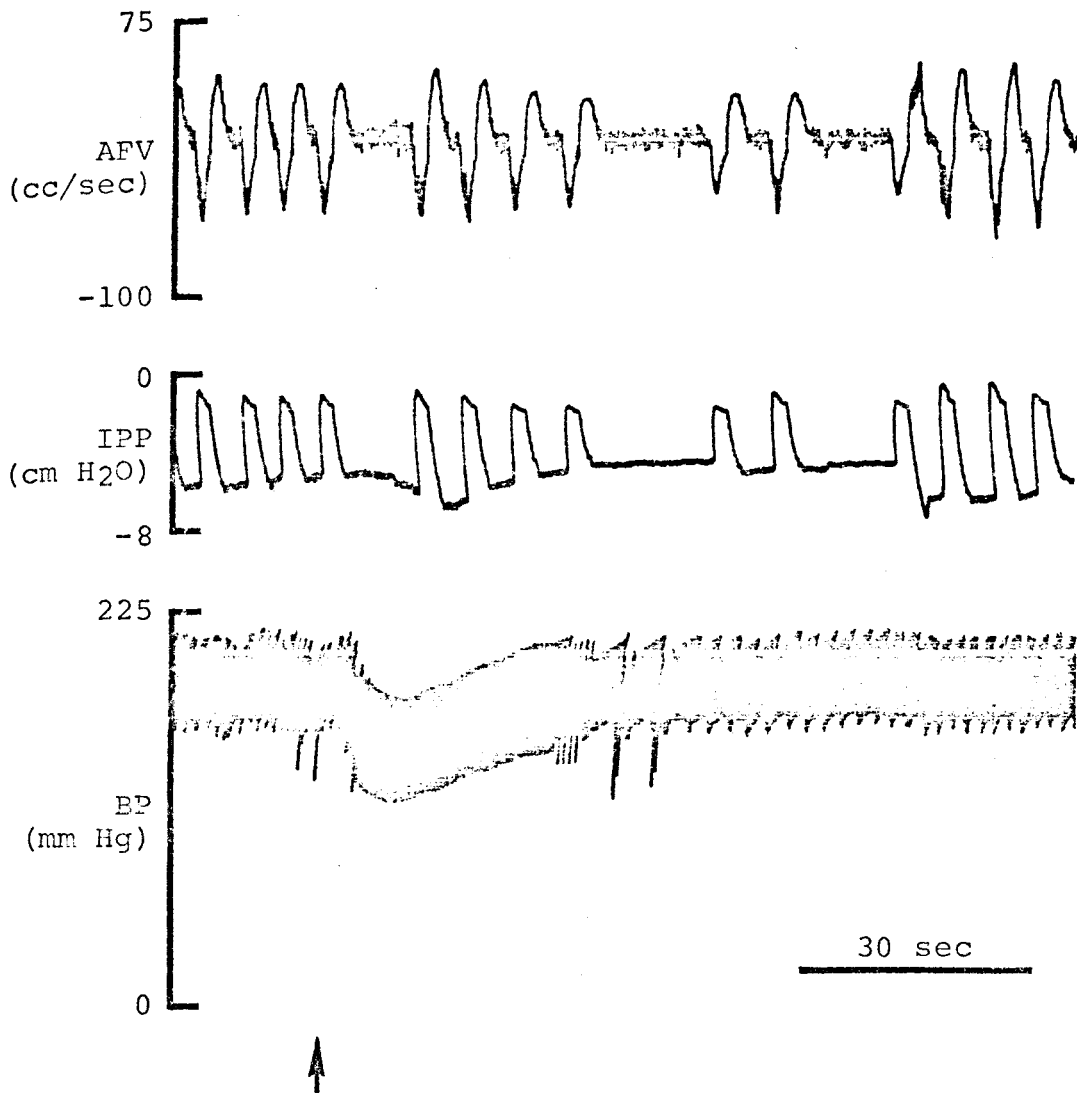


Figure 75. Changes in respiratory pattern produced by 4.4 mg/Kg doxapram (arrow) in a vagotomized cat. Tracings as in Figure 63.

## CHAPTER VI

### DISCUSSION

#### A. Population Modeling

In assessing the experimental data presented in this dissertation, two crucial questions arise: first, how are the limits of a cell population determined, and second, how many single cell elements need to be combined to represent physiological response characteristics of the defined population?

Concerning the first question, the assumption must be made that "singular" biological populations possess inherent variability. Population characteristics may follow normal or skewed distributions, but in either case, variations are present. Recognition of this fact is important, since it may prevent establishment of such rigid criteria for a population (minimization of variability) that only a few elements are eligible. This, by necessity, requires the postulation of a number of other so-called populations. Cohen (40, 41, 42, 43, 45) and others (14, 15, 128, 169, 173), in describing no less than eight different

respiratory phasing patterns, may have failed to recognize inherent biological variability. If so, the elaborate theories on rate and depth control proposed by different individuals (15, 42) are highly questionable. Under these conditions, the experimental analysis no longer deals with real biological entities, but with artifactual systems whose parameters are mere functions of the experimental techniques employed.

A second, but less common error, is the tendency to be over-inclusive when defining the limits of a population. In this case, cells rightfully belonging to separate populations are included within the same category. Subsequent averaging of data results in the masking of real and important physiological trends. Obviously, a balance between over and under selectivity should be sought when attempting to assign single cell observations into appropriate groups. As a guideline, it is suggested that population limits be set such that standard deviations of individual elements approach predictable magnitudes as determined from statistical theory.

Assuming that real physiological populations can be defined, the second question posed above seeks realistic means whereby population characteristics can be accurately assessed. Since it is unlikely that all components



in a population can be sampled, functional description of many systems, notably CNS subsystems, must rely on approximation techniques (31, 41, 51, 101, 102, 127, 129). Two powerful techniques stand out in importance. First, Knight (101, 102) has stressed the importance of averaging repetitive observations from a single cell when describing activity in a population of neurons. In his own words (101), "While at present it is not feasible to record individually from a uniform population of neurons, it is possible to do something equivalent: to record from a single member over repeated stimuli." That is, processes spacially distributed over many components, but occurring at the same point in time, can be approximated by temporal summation of a single component's output at one point in space. Second, Nelson (127) has emphasized the importance of grouping characteristics of individual cells for accurate population description. He writes, "Ideally, the electrode records from but a single element in the system; with data from a large number of such elements it is possible to 'resynthesize' the total population and draw conclusions about the location of units having similar temporal patterns of firing." These two mutually interactive techniques can be coupled as follows in population description studies:

<u>Population</u>	<u>Single Cell Discharge Pattern</u>	<u># Cells</u>	<u># Cycles</u>
Homogeneous	Smooth	1	1
Homogeneous	Rough	1	N
Heterogeneous	Smooth	N	1
Heterogeneous	Rough	N	N

The number of cells and number of cycles averaged for each cell are listed as either 1 or N (arbitrary large number) in specific combination pairs depending on the type of population and type of single cell discharge pattern. For example, if all the components of a population are identical (homogeneous) and if each discharge is not subject to change fluctuations (smooth), population characteristics can be identified by examination of a single burst discharge from one cell in the population. If all the cells are identical (homogeneous), but are subject to change fluctuations (rough), then it is necessary to average many output cycles from one cell before description of population characteristics is possible. In situations where the population contains a variety of cellular individuality (heterogeneous), it is necessary to sample from a large representative fraction of the population for accurate description. Considering the argument of inherent biological variability presented above, the

first three cases possessing population homogeneity and/or single cell regularity must be rejected as untenable. For the respiratory system, this is confirmed by the rough interspike interval modulation curves for single burst discharges (Figure 9A) and the presence of error lines (standard deviation) in all regression plots (Figures 11-34). This leaves the fourth case which demands that many cycles from single cells and repeated observations on different cells be averaged for population assessment. Description errors are, of course, minimized by maximizing the N values.

In this study, four respiratory neuron populations were defined which subdivided inspiratory and expiratory cell types into vagi-intact and vagotomized groups (I-VO, I-VI, E-VO, E-VI). The limits of each population were inclusive enough to allow for inherent variability, but exclusive enough to avoid masking of important physiological data (determined via range of error lines in Figures 11-34). Data analysis of single cell discharges included generation of common histogram interval distributions (66, 67, 68, 106, 139, 165 and Figure 6) and the new interspike interval modulation curve (Figure 9). The latter quantitatively defined the average interval ordering for a single respiratory cell discharging under

steady state conditions. This technique proved superior to those of Cohen (40) and Bertrand and Hugelin (24) who used modified histogram techniques. From the histogram (Figure 6) and interspike interval modulation curve (Figure 9B), 34 parameters were defined (Table IV) which successfully quantitated all phases of spike activity (initial, middle, terminal). This analysis compares favorably with other work published in the literature in which only a few parameter measurements were made in any one study. For example, the following parameters form an almost complete list of measurements made on single respiratory discharges (listed in decreasing frequency of occurrence - author references not exhaustive): ST (31, 40, 41, 51, 63, 91, 97, 127, 129); TL (31, 41, 51, 63, 91, 97, 127, 129); mean discharge frequency or  $1/MEN$  (31, 40, 51, 69, 72, 91, 97); RR (31, 127, 129); CT (127, 129); peak discharge frequency or  $1/MI$  (40, 97); T100 (97); TLP (129).

Discharge characteristics of the I-VO, I-VI, and E-VI populations were determined by regression analysis of 24 parameter groups on respiration rate and mode (Figures 11-43). The E-VO population data were not reported since only three cells contributed to this group (cf. Table III). Before discussing the parameter trends observed, it is important to consider the degree to which

separate populations have been sampled.

Waldron (174) has estimated that at any one time there are at least 2000 respiratory (inspiratory and expiratory) neurons in the total respiratory complex of the rostral medulla. Assuming this to be a reasonable estimate and assuming an inspiratory to expiratory cell ratio of 2:1 (155), there must be at least 1333 inspiratory and 667 expiratory cells involved in respiratory function (cf. Table III). In this study, only 10 I-VO, 51 I-VI and 23 E-VI cells were analyzed (Table III) which corresponds to 0.8%, 3.8% and 3.4% of each respective population. However, assuming that each burst discharge in time represents one cell in space (101), corrections can be made for the repetitive observations on single cells during curve smoothing procedures. For example, 1159 I-VO, 6079 I-VI and 3776 E-VI trains (Table III) or simulated cell samples contributed 86.9%, 456.0% and 566.1%, respectively, to each population. Within the constraints of these assumptions, the mathematical analysis used in this study has provided an excellent estimate of the inspiratory (I-VO, I-VI) and expiratory (E-VI) populations. Data interpretation should be weighed accordingly.

Parameter regressions plotted in Figures 11-43 represent specific population trends during barbiturate

and doxapram induced modifications in the spontaneous respiration rate and mode interval time. If it can be assumed that mode and tidal volume are reciprocally related as the rationale in Results proposes, specific characteristics in respiratory cell discharge patterns can be correlated with the two major outputs of the respiratory system namely, respiration rate and depth. Both rate and mode curves (Figures 11-43) can be viewed as barbiturate depression curves since pentobarbital or thiopental administration leads to decreased rates and tidal volumes (82, 131). Relatively speaking, the contribution from doxapram-facilitated data points is minimal.

Two series of figures are presented to summarize I-V0, I-VI and E-VI population characteristics. In the first series, Figures 76-80, regression correlation coefficients ("r" values) are plotted in bar graph format for each parameter as a function of either respiration rate or mode. All populations are represented. The "r" value reveals the degree of population homogeneity toward any one parameter. For example, as "r" values range from 0.00 to 1.00, there is a population trend or shift from complete heterogeneity to complete homogeneity. By definition, all parameters fall with these limits. For specific respiration rate or mode plots, different populations

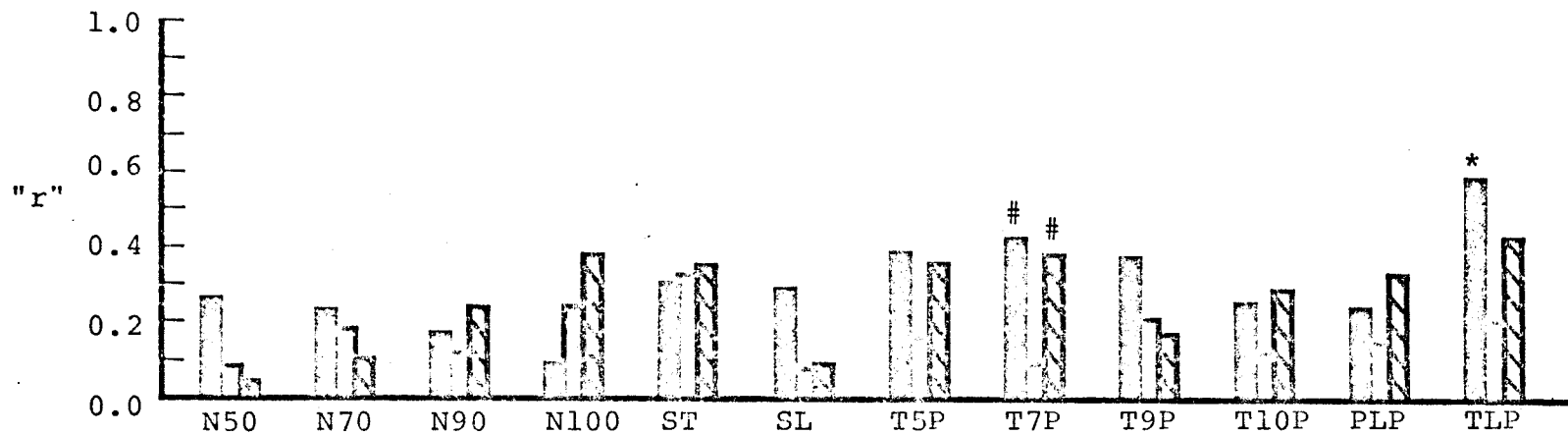
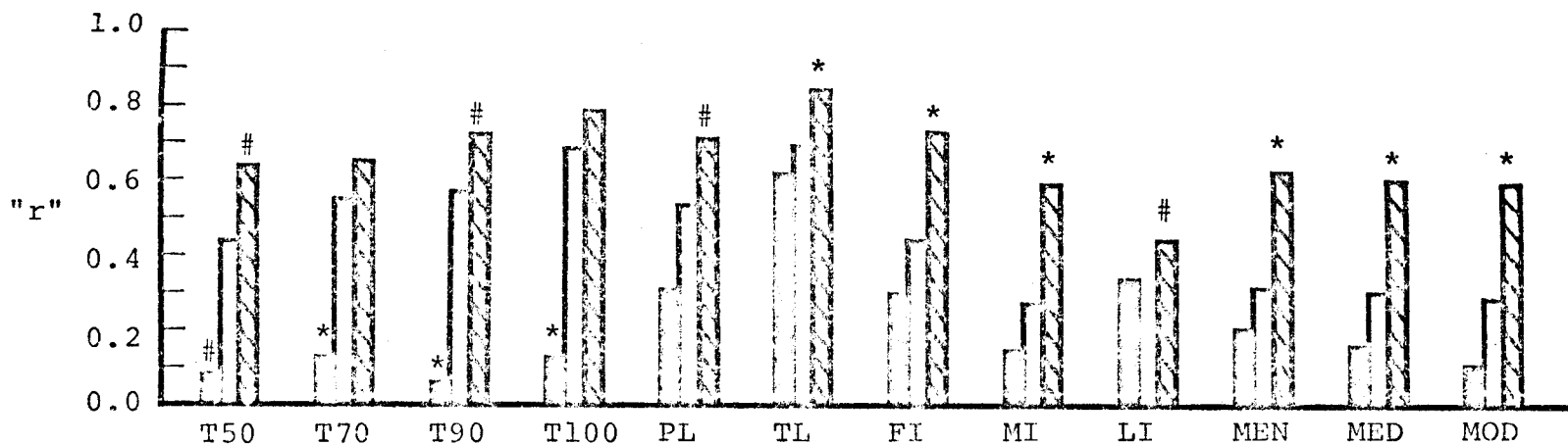


Figure 76. Comparison of correlation coefficients for 24 parameters on respiration rate for I-VO (solid), I-VI (open) and E-VI (striped) cells: # = P<0.05; \* = P<0.01.

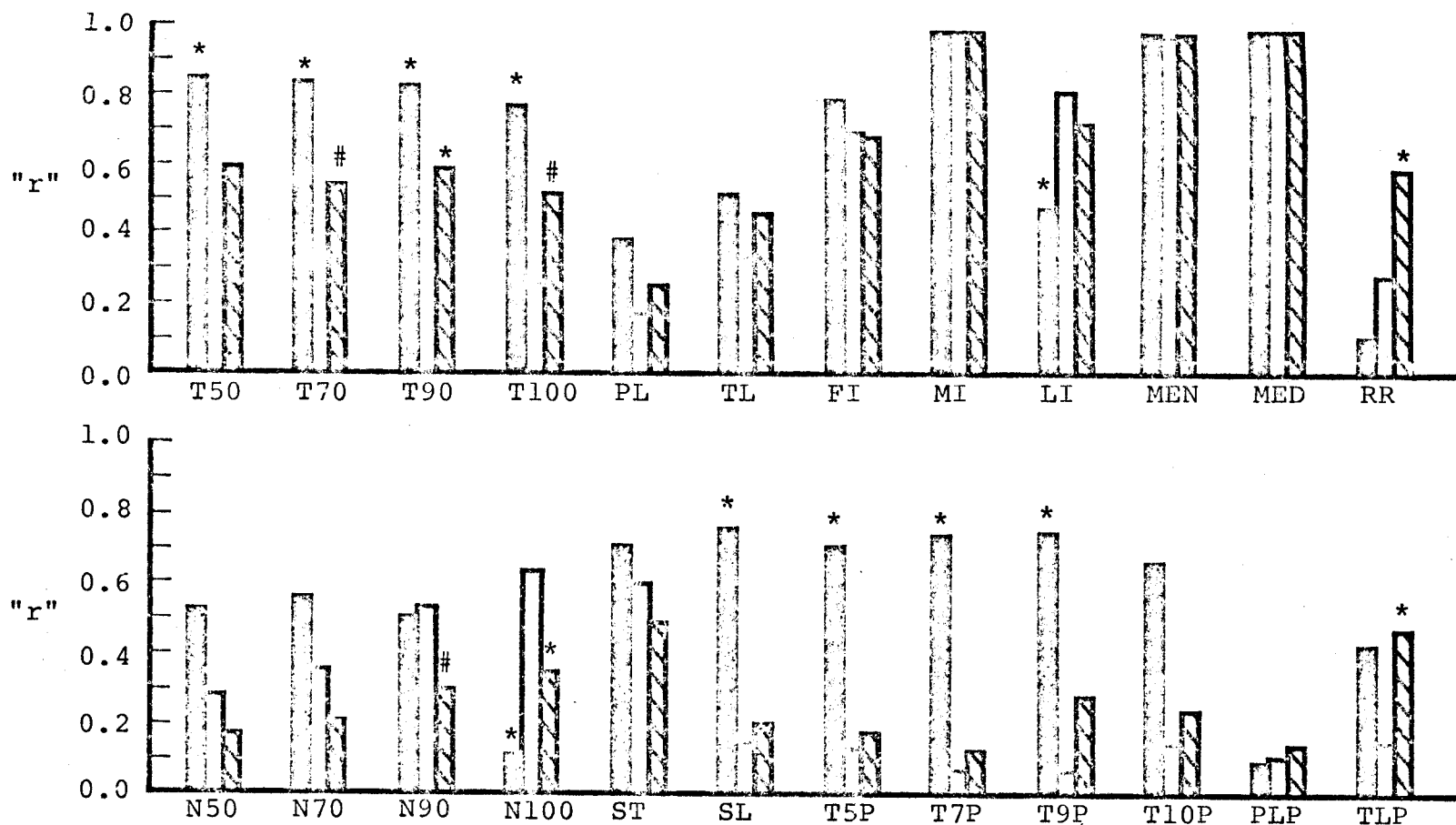


Figure 77. Comparison of correlation coefficients of 24 parameters on mode for I-VO (solid), I-VI (open) and E-VI (striped) cells: # = P < 0.05; \* = P < 0.01.



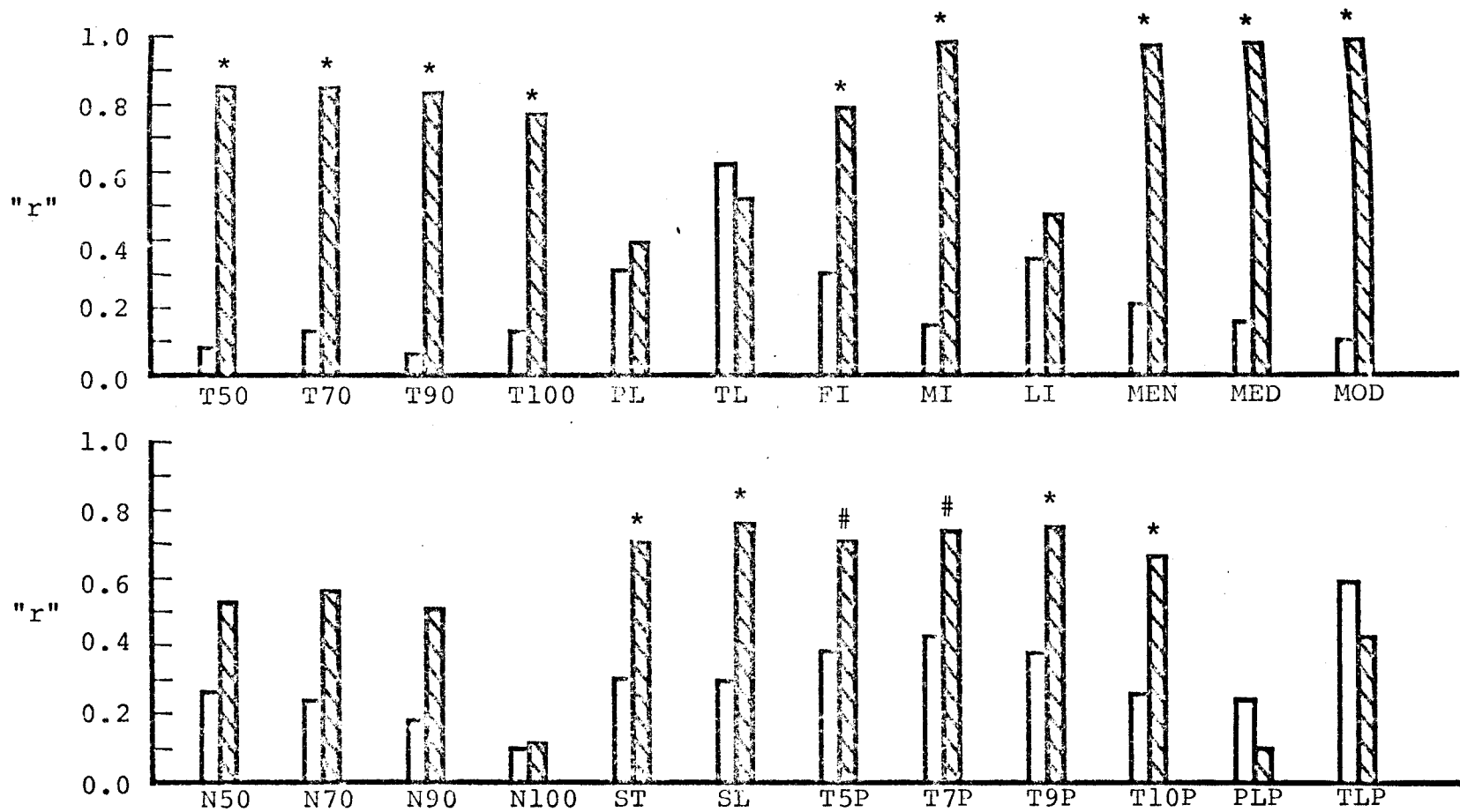


Figure 78. Comparison of correlation coefficients for 24 parameters on respiration rate (open) and mode (striped) for I-VO cells: # = P<0.05; \* = P<0.01.

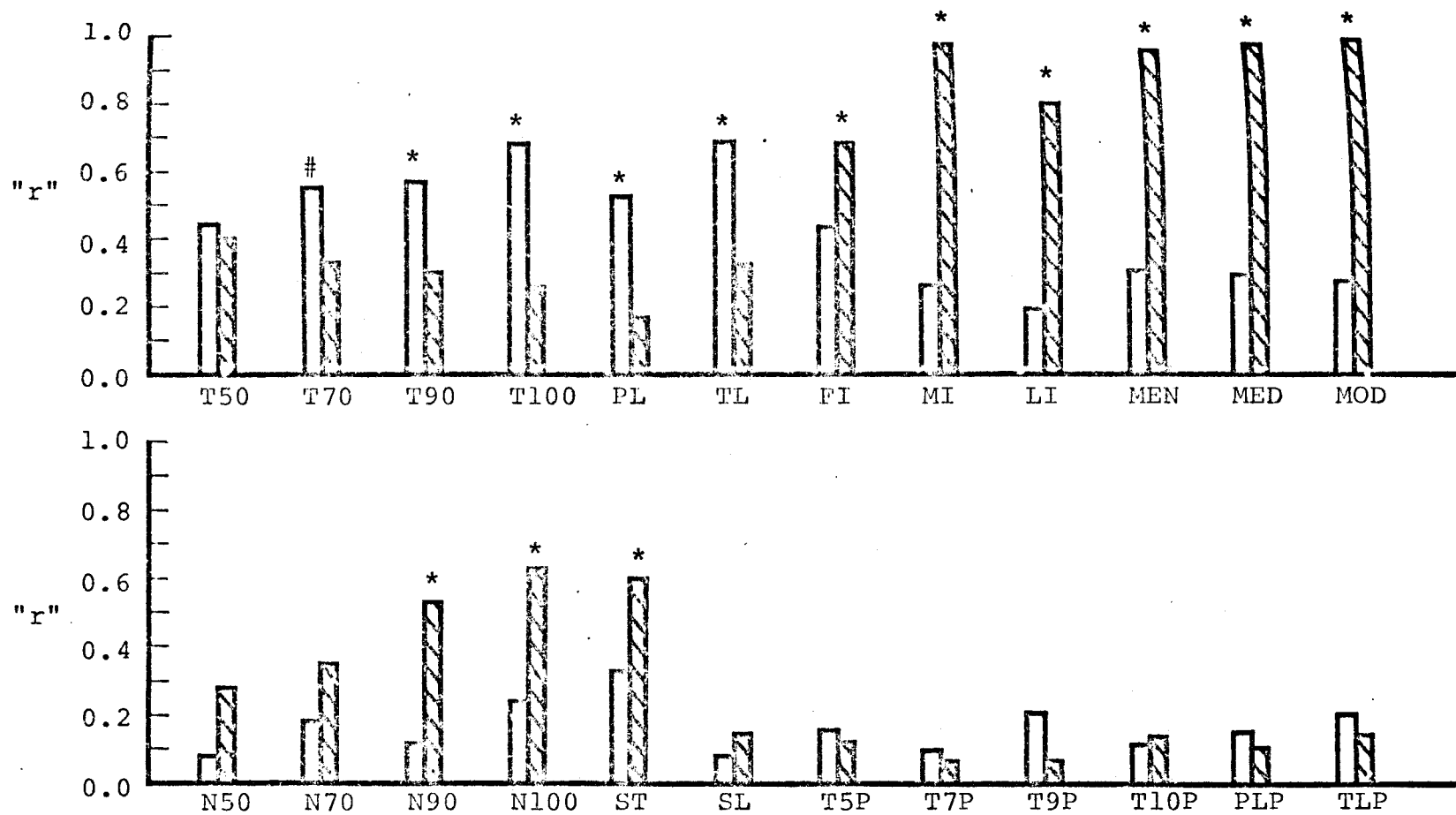


Figure 79. Comparison of correlation coefficients for 24 parameters on respiration rate (open) and mode (striped) for I-VI cells: # = P < 0.05; \* = P < 0.01.

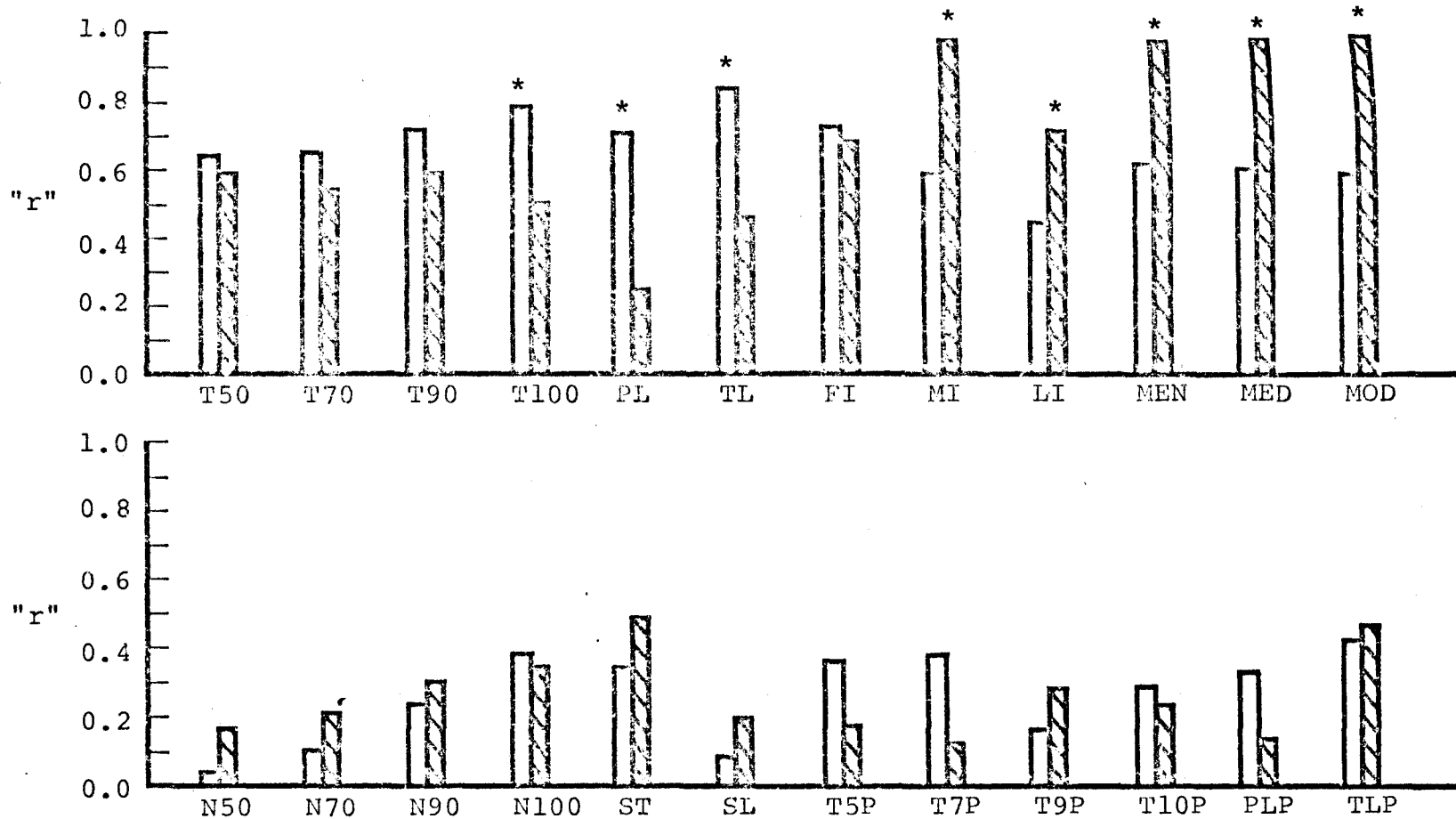


Figure 80. Comparison of correlation coefficients for 24 parameters on respiration rate (open) and mode (striped) for E-VI cells: \* = P < 0.01.

exhibit different "r" values. Also, any one parameter usually correlates better with respiration rate than mode and vice versa. In such cases, the significant difference between "r" values is indicated by a number sign (#) for  $P < 0.05$  and an asterisk (\*) for  $P < 0.01$ . Lack of a symbol designates that "r" values are not significantly different at the 0.05 level.

Figure 76 shows that for the first 12 parameters (T50 - MOD) expiratory cells have a higher degree of correlation with respiration rate than do inspiratory cells. This indicates that for these parameters, the expiratory population possesses less individual cell-to-cell differences and is relatively more homogeneous than either inspiratory population. Comparison of I-VO and I-VI data for the first 12 parameters shows that vagotomy usually decreases the correlation coefficients of inspiratory cell characteristics on rate. This evidence supports the concept that the control of respiratory rate operates through vagal afferent feedback (55, 57, 73, 129, 161, 162, 171). None of the latter 12 parameters (N50 - TLP) for any cell type in Figure 76 display high "r" values, illustrating that these measurements are relatively independent of rate changes.

Figure 77 shows that I-VO parameters usually correlate

best with mode, indicating that vagotomy significantly entrains many inspiratory discharge characteristics with alterations in respiratory depth. The expiratory and I-VI orderings are fairly consistent with rate data (Figure 76), with E-VI cells showing higher correlations on mode than I-VI cells. Also, the last 12 parameters for these two populations correlate very poorly with mode suggesting that, with the exception of I-VO parameters on mode, these parameters are poor indicators of either rate or depth changes.

In Figure 78, I-VO parameter "r" values are presented for both rate and mode regressions. Similar plots are found in Figure 79 for I-VI cells. When the vagi are intact (Figure 79), parameters derived from the summation of intervals (T50 - TL) correlate best with respiration rate, while individual interspike intervals (FI - MOD) correlate best with the mode. Parameters N50 - ST show an increased tendency to correlate with the mode, but the latter parameters (SL - TLP) fail to correlate with either rate or mode for I-VI cells. Sectioning of the vagi (Figure 78) significantly alters the correlation pairing for inspiratory T50 - PL parameters from rate to mode. The significant TL correlation with rate when the vagi are intact is reduced to no significant difference

from the mode regression after cutting both vagi. The FI - MOD parameters retain their strong mode correlations after vagal section. Finally, vagotomy tends to increase all mode correlations for the last 12 parameters (N50 - TLP) in the inspiratory population. These results show that, with the exception of parameters T50 - TL when the vagi are intact, most parameters correlate best with fluctuations in the mode irrespective of the presence or absence of vagal afferents. In general, this indicates that vertical ordering of the interspike interval modulation curve is more sensitive to depth changes, while horizontal interval ordering ( $\Sigma$  intervals) varies with respiration rate changes. The latter reverts to mode correlations after vagal section.

Figure 80 presents similar data for E-VI cells. Expiratory cells tend to show the same qualitative "r" value pattern as I-VI cells for most parameters (compare with Figure 79). This includes a respiratory rate coupling for horizontal time measurements (T50 - TL) and mode coupling for vertical time measurements (LI - MOD) with the exception of FI. As with I-VI cells, the last 12 E-VI parameters (N50 - TLP) show poor correlations with either rate or mode.

It has been indicated previously that correlation

coefficients or "r" values reveal the degree of homogeneity or degree of fit. This concept of correlation also involves the type of fit since population data were represented by the best of four different regression equations; namely, linear, exponential, logarithmic or log-log. The best type of equation fits for all cell type parameters are listed in the Appendix (Table XIV) for both respiration rate and mode regressions. Taking all 24 parameters at a time for each cell type reveals that 16 I-VO, 11 I-VI, and 13 E-VI parameters are fit best with logarithmic, exponential and logarithmic equations, respectively, as a function of respiration rate. For mode regressions, 19 I-VO, 11 I-VI and 12 E-VI parameters fit best with logarithmic and linear equations respectively. The interpretations of these data are highly speculative but they are suggestive of different processes in operation for different cell populations as a function of either rate or mode.

The second series of summarization data for population studies are presented in Figures 81-83. For each cell type, I-VO, I-VI and E-VI, a family of theoretical interspike interval modulation curves were reconstructed by interpolation of parameter values from regression equations in the Appendix (Tables VIII-XIII) at five

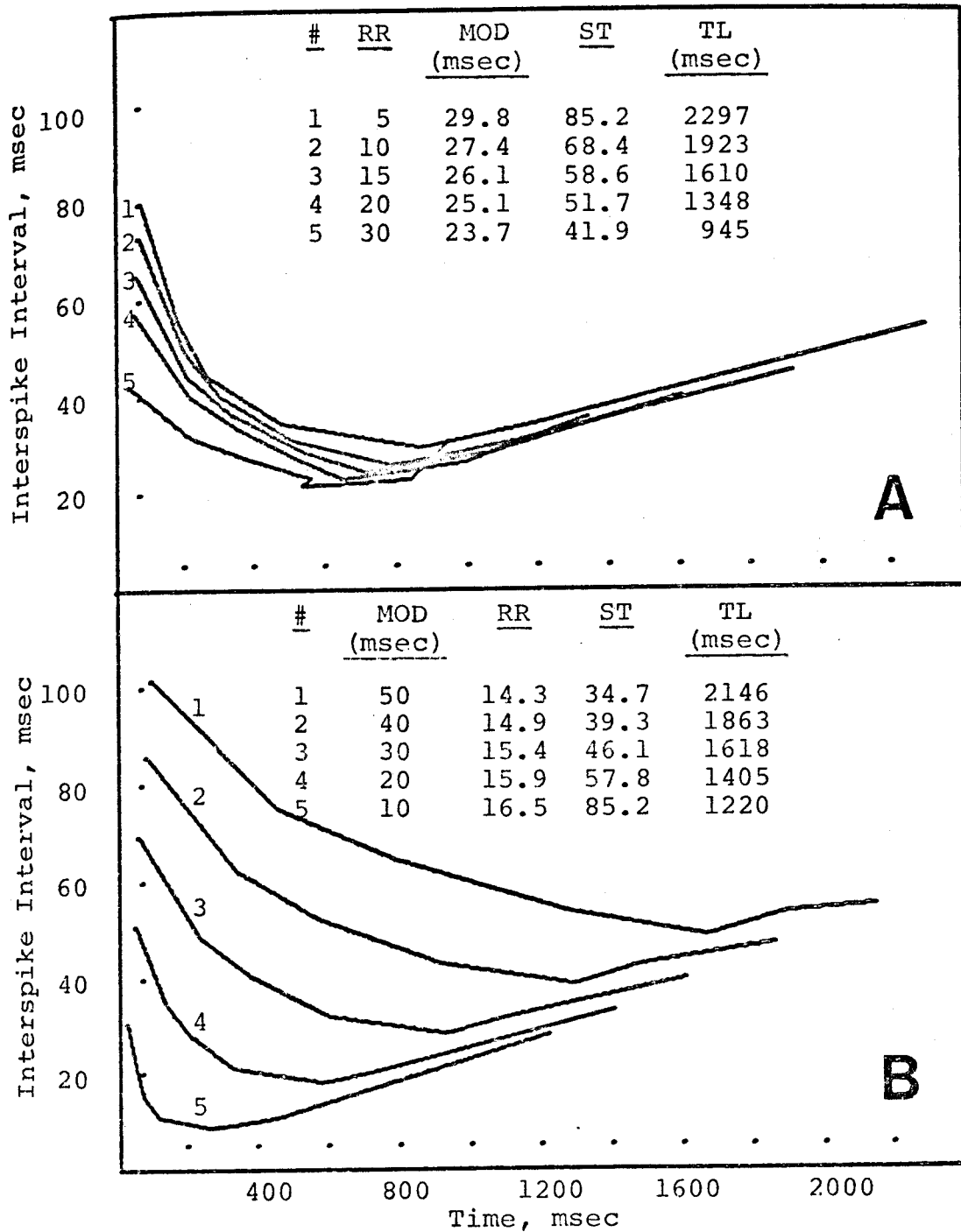


Figure 81. Theoretical reconstruction of interspike interval modulation curves for I-VO cells. Pattern shifts are shown for increasing respiration rate (A) and for decreasing mode (B). Tabular data is given for selected parameters.



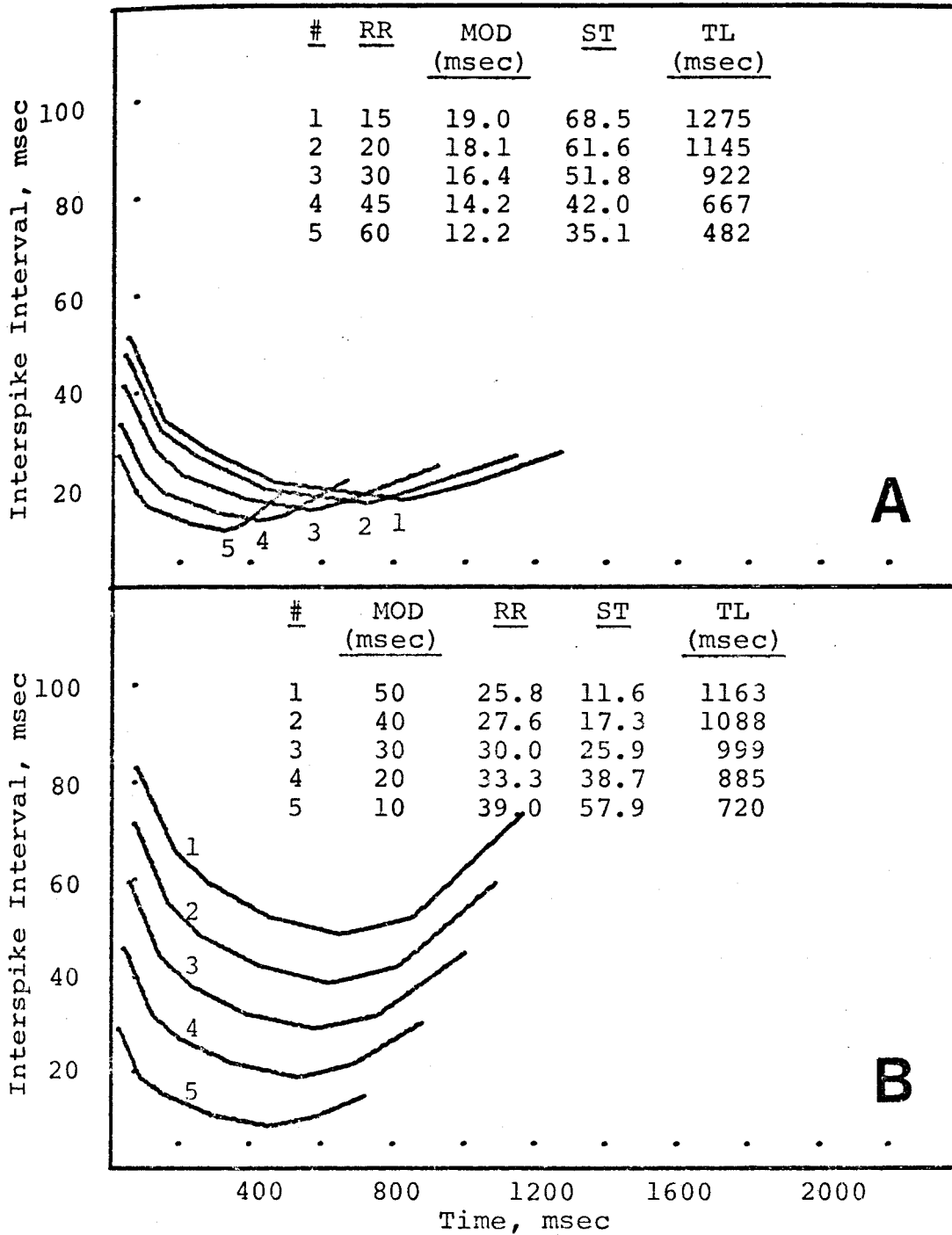


Figure 82. Theoretical reconstruction of interspike interval modulation curves for I-VI cells. Pattern shifts are shown for increasing respiration rate (A) and for decreasing mode (B). Tabular data is given for selected parameters.

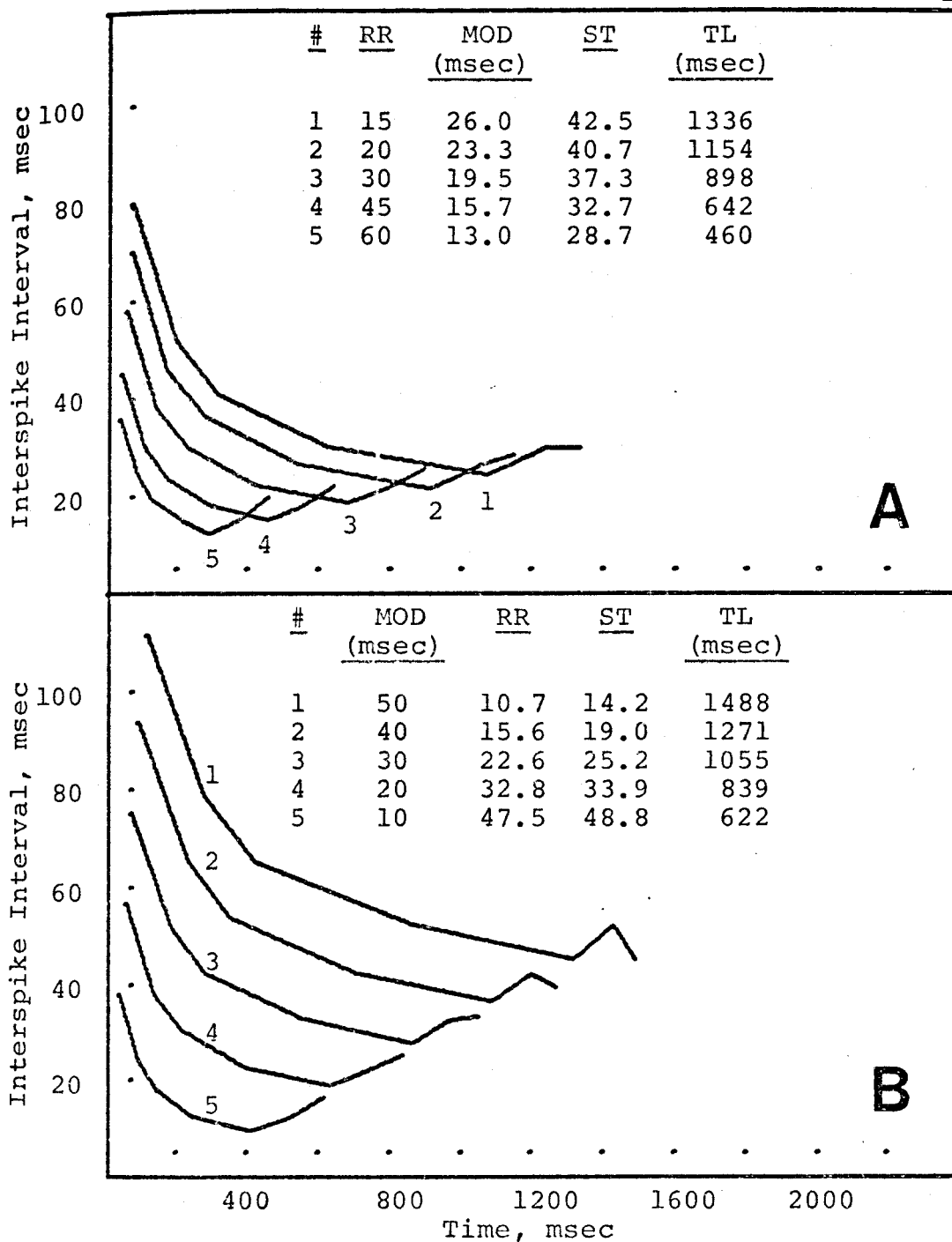


Figure 83. Theoretical reconstruction of interspike interval modulation curves for E-VI cells. Pattern shifts are shown for increasing respiration rate (A) and for decreasing mode (B). Tabular data is given for selected parameters.

different rates and five different modes. Each theoretical curve consists of seven points with the following coordinates:

<u>#</u>	<u>X</u>	<u>Y</u>
1	FI	FI
2	T50	$0.5(FI - MI) + MI$
3	T70	$0.3(FI - MI) + MI$
4	T90	$0.1(FI - MI) + MI$
5	T100	MI
6	T90 + PL	$0.1(FI - MI) + MI$
7	TL	LI

Corresponding respiration rate, mode, spikes per train and train length parameters are identified for each curve in tabular format. Consistently, the first parameter is the independent variable and the remaining three are dependent variables. For respiration rate plots,  $MOD \propto TL/(ST-1) \sim k$ . For mode plots,  $TL \propto MOD(ST-1) \sim k$ . Rate and mode curves are plotted in the same figures for each cell population. Panel A represents the projected population response to increasing rates. Panel B predicts the population response to decreasing modes (increasing depths). All plots in Figures 81-83 have identical scalings for easy comparison.

Theoretical interspike interval modulation curves

from I-VO population data are plotted in Figure 81. Panel A shows that the family of curves form a tight distribution group, differing significantly only in train length and possibly first interval as the respiration rate is increased. Only a small decrease in mode is observed due to a simultaneous proportional decrease in train length and spikes per train. Panel B shows significant pattern changes in both horizontal (X-axis) and vertical (Y-axis) interval ordering during decrease in mode time (increase in depth). This is due to the high correlation of many I-VO parameters on mode (Figure 78). As the mode decreases, a minimal increase in respiration rate occurs, presumably due to the simultaneous decrease in train length ( $TLP \sim k$ ). The latter results from a decrease in mode which is faster than the increase in spikes per train.

Data from the I-VI population are plotted in Figure 82. The curves in panel A show a downward shift as the respiratory rate is increased. Disproportionate decreases in spikes per train and train length produce moderate mode interval shortening. Panel B shows theoretical parallel shifts of the I-VI curves as a function of decreasing mode. The reduction in train length (and, therefore, the increase in respiration rate) follows predicted mode and spikes per train combinations. Comparison of Figures 82

and 81 reveals interesting inspiratory cell pattern differences when the vagi are intact and severed.

The last set of theoretical interspike interval modulation curves are plotted in Figure 83 for E-VI cells. In panel A, an increase in respiration rate is accompanied by a decrease in mode due to a faster decrease in train length than spikes per train. As the mode decreases in panel B, the modulation curve shifts downward in a characteristic activation pattern. Unlike I-VI curves in Figure 82B, E-VI curve shifts are not parallel since a simultaneous decrease in train length occurs. This results from a decreasing mode - spikes per train product as the respiratory depth is increased. Presumably this explains the simultaneous increase in respiration rate as a function of mode. Comparison of E-VI and I-VI patterns in Figures 83 and 82, respectively, demonstrates qualitative similarities between corresponding curves. Quantitatively, however, expiratory cells consistently have lower discharge frequencies than inspiratory cells at equivalent respiratory rates (panel A) as shown by the higher modulation curves for E-VI cells. In mode plots (panel B), expiratory first interval times are significantly longer than those for inspiratory cells.

In summary, the reconstructed discharge patterns

for I-VO, I-VI and E-VI cell groups in Figures 81-83 constitute specific population characteristics corresponding to medullary outputs of respiration rate (panel A) and respiratory depth (1/MOD, panel B). A major conclusion from these data is that respiratory cell discharge patterns have a two-fold dynamic response. In one sense, individual curves represent dynamic modulation of the interspike interval during the train progression as indicated by the "U" shaped quality of all curves. This property of discharge has been recognized by many investigators (51, 69, 72, 74, 91) who have described frequency modulation curves for single respiratory cells. In a second sense, however, respiratory discharge patterns are dynamically locked to the respiration rate and mode outputs. Only a few workers have realized this for rate (31, 127, 129) and depth changes (129). From the data presented in this dissertation, it is suggested that the interpretation of single respiratory cell discharge characteristics must incorporate respiration rate and depth modifications. This suggests that single cell data acquired under differing rate and depth conditions should not be averaged together as Nesland et al. (129) failed to recognize.

## B. Single Cell Elements

As previously discussed, all regression plots (Figures 11-34) characteristically possessed standard deviations (error lines) from the mean regression line, indicative of inherent biological variation within each defined population of cells. Selected single cell responses to induced or spontaneous modifications in the breathing pattern were presented in Results to illustrate this cell to cell variability in otherwise homogeneous neuron populations. Figure 46, for example, presents evidence that two cells recorded at the same time can exhibit wide variation in discharge frequency. To summarize single cell differences from and similarities to population characteristics, interspike interval modulation curves for individual units (Figures 44-62) and theoretical units (Figures 81-83) can be compared for corresponding cell types.

Deviations from I-VO population predictions (Figure 81) are seen for inspiratory cell C81UI5L in Figure 54B. After severing both vagi (data points 1 and 2) data points 3-10 represent interspike interval modulation curve response to cumulative doses of pentobarbital. A simultaneous decrease in respiration rate and increase in mode interval time resulted in a predicted

increase in train length, although no overall change in spikes per train was observed. Induced modifications in the latter parameter were masked by significant depression of both rate and depth components by the barbiturate (82, 131).

Many single cell comparisons can be made with the theoretical I-VI population data (Figure 82). Inspiratory pattern responses for two cells, C32UI1R (Figure 44B) and C80UI1R (Figure 45B), show parallel shifts during spontaneous depth increase and decrease, respectively. Since respiration rate modification in each case was at best only minimal, both modulation curve sets match predicted I-VI population patterns as a function of mode (Figure 82B). During a spontaneous increase in respiratory rate with minimal depth change, inspiratory cell C28UI5 (Figure 47B) showed a simultaneous increase in mode interval time. This rendered the single unit pattern response atypical compared to I-VI population characteristics as a function of rate (Figure 82A). For another inspiratory cell, C62UI3R (Figure 48), doxapram-induced increases in respiratory rate revealed typical discharge pattern responses. That is, the interspike interval shifted downward as the mode interval time decreased with a simultaneous decrease in train length and



spikes per train. This response paralleled I-VI population predictions during rate increase (Figure 82A). Finally, inspiratory cells C63UI4R (Figure 50B) and C83UI1L (Figure 52B) were followed during pentobarbital administration. The response pattern of cell C63UI4R during respiration rate decrease closely followed population trends (Figure 82A) with increases in mode, spikes per train and train length. The response pattern of cell C83UI1L, was atypical. In this case, respiration rate was decreased and train length was lengthened, but a significant decrease in the mode interval time caused an exaggerated increase in spikes per train. The modulation curve shift (Figure 52B) appeared to possess characteristics of theoretical I-VI population curves as a function of both rate and mode.

A few single expiratory cell examples can be compared with E-VI population predictions in Figure 83. Cell C64UE4R (Figure 56B) and cell C71UE2R (Figure 58B) were subjected to cumulative doses of pentobarbital. In the former case, only a slight decrease in the mode was observed as the rate was significantly decreased. There was a corresponding increase in spikes per train and train length. With the exception of a peculiar inflection in the interspike interval modulation curve

for the first three data points (Figure 56B), this pattern response resembled theoretical E-VI population trends during rate decrease (Figure 83A). In general, cell C71UE2R pattern shifts paralleled E-VI population constructions as a function of mode (Figure 83B). During pentobarbital accumulation, there was only a slight respiration rate decrease while the mode was significantly increased. The train length was not lengthened as predicted, but spikes per train decreased nevertheless. In Figure 60B, expiratory cell C82UE3R was depressed by cumulative doses of morphine sulphate. With the exception of the control pattern at data point 1, successive injections of morphine produced atypical interspike interval modulation curve shifts which did not correlate with E-VI population projections related to either respiration rate or mode changes (Figure 83). Morphine caused a slight increase in respiration rate which was inconsistent with Ngai's (132) observations. Also, the mode was increased while the spikes per train decreased, producing a significant shortening in the train length. Examination of Figure 60B reveals that morphine, in this case, exerts its greatest effect on the latter portion of the interspike interval modulation curve. Finally, expiratory cell C68UE6R (Figure 62B) was analyzed before (Curve 1)

and after (Curves 2 and 3) unilateral left vagotomy. The mode interval time and spikes per train changed in opposite directions so that train length, and therefore respiration rate, remained relatively constant. Disregarding the latter two characteristics, the single cell pattern shifts resembled E-VI population data as a function of mode (Figure 83B).

From the comparison of inspiratory (I-VI) and expiratory (E-VI) single cell data and population responses, it is difficult to find major differences between these two neuron groups. Both cell types possess indistinguishable variability during barbiturate depression, and interspike interval modulation curves (Figures 82 and 83) reveal distinct similarities as a function of either respiration rate or mode. Interpretation of these data is consistent with the hypothesis that inspiratory and expiratory neurons are functionally similar. It is possible that these cell types belong to a similar reticular formation cell class, differing only in intercellular connections.

Gross characterizations of respiratory cell discharge patterns include assessment of spike frequency, burst duration and phasing with the respiratory cycle. Variations in the first two parameters throughout the

respiratory complex can be explained on the basis of threshold differences among cells (91) and/or differing combinations of neuronal connections (36). The pairing of cell inputs (excitatory and inhibitory) with threshold characteristics of the membrane defines the level of excitability for any given respiratory neuron. This excitability level for individual respiratory cells and grouped neuron populations fluctuates in phase with the respiratory cycle due to oscillating inputs (156) and variations in threshold values (18, 154).

The data presented in this dissertation indicate that expiratory cells have lower discharge frequencies than inspiratory cells at equivalent respiratory rates and modes (compare Figures 82 and 83) suggesting that the former have less excitatory inputs and/or higher firing thresholds than the latter. Also, the observation that expiratory cell parameters correlate better with rate and mode (cf. Figures 76 and 77) indicates that the E-VI population is relatively more homogeneous than the I-VI cell group. This decreased variability in the expiratory population may be indicative of fewer modulatory inputs to these cells, or of the dominance of one or more inputs with relatively lower variability.

Finally, both inspiratory and expiratory cells

tend to fire for a constant fraction of the respiratory cycle ( $TLP \sim k$ ) irrespective of respiration rate or depth changes (Figures 49-61). A wide variety of percentages are found for I-VO, I-VI and E-VI cell groups (Figures 22, 34 and 43) indicating that all cells in any defined population do not have identical starting and stopping times. Further study into this phenomenon, preferably at the intracellular level, should provide important data on single cell inputs and threshold characteristics.

### C. Respiratory Complex Organization

The most consistent definition of apneusis refers to the situation where inspiratory mechanisms are locked in a hold position for an extended period of time. This respiratory pattern can be induced either by appropriate neural transections (112, 113, 114, 116, 117, 136, 167, 176 and Figure 63) or high barbiturate levels (31, 82, 131, 151 and Figure 64). Although both techniques result in prolonged inspiratory times, significant differences occur in the depth of inspiration. Neural transections produce near maximal inspiratory efforts while barbiturate administration produces a prolonged, but shallower inspiratory hold. Although this difference cannot be seen by comparing Figures 63 and 64, this result is presumably due to partial depression of the

inspiratory mechanism by barbiturate.

The production and abolition of apneusis has been useful in modeling the organization of brainstem respiratory mechanisms. For example, localized lesions in the rostral pons in vagotomized preparations have resulted in apneusis (22, 93, 134, 147, 151, 152, 170, 172). Investigators have suggested that this pneumotaxic area is inhibitory to inspiration. Transection between the medullary-pontine border in apneustically breathing animals converts this pattern into one of gasping (26, 87, 112, 133, 136, 167, 170). Coupled with the former data, this suggests that the lower pons contains a constantly discharging cell population which is excitatory to rhythmically active medullary inspiratory neurons. From this work and similar studies, a general concept of respiratory organization developed which suggested that medullary inspiratory neurons are inhibited by vagal, pneumotaxic, cerebellar and expiratory cell connections and are facilitated by apneustic drives.

The fact that barbiturates can induce apneusis suggests that respiratory inhibitory mechanisms are differentially more susceptible to pharmacologic blockade than facilitatory systems. If this were not the case, barbiturate administration should result in apnea or

expiratory holds as seen with morphine (132). Figure 72 implies that rostral pontine cells are depressed before medullary cells at equivalent barbiturate levels in the same cat. This finding is consistent with the observation that barbiturates act at high levels of the CNS and then work their way down the neuraxis as the dose is increased. Anatomically speaking, then, the pneumotaxic region may be depressed before the apneustic or medullary areas with cumulative barbiturate administration.

When the vagi are intact, it takes a higher dose of barbiturate to induce apneusis than when the vagi are severed (Figure 66). Assuming that pneumotaxic circuits are the first to be blocked as discussed above, this drug level difference suggests that either vagal influences or inspiratory mechanisms are mediated at sub-pneumotaxic stations (lower pons) or inhibitory synapses are more susceptible to barbiturate blockade than facilitatory synapses. In either case, high barbiturate levels appear to disrupt brainstem connections so that unilateral vagal influence cannot cross over to the opposite brainstem side (compare Figures 63 and 67).

The notion that apneusis results from a preponderant influence of facilitatory versus inhibitory drives on inspiration was introduced in Results. This concept

is actually a combination of Gray's (77) multiple factor theory ( $\Sigma$  chemical factors) and Gesell's et al. (71) reflexogenic components ( $\Sigma$  neural factors). For example, removal of inhibition via cerebellectomy (76, 83 and Figure 65), vagotomy (Figure 67) or decreasing carotid sinus pressure (BCO, Figure 73; histamine, Figure 74) all produced apneustic breaths. Similarly, increased facilitatory drive via doxapram (Figure 75) produced identical pattern changes. These results could only be observed when the barbiturate level was significantly higher than the initial anesthetic dosage (30 mg/kg pentobarbital). At lower levels, cerebellectomy, showed little, and vagotomy showed marked rate and depth changes, while decreased sinus pressure (86) and doxapram produced definite rate acceleration and depth increase. From these data it is suggested that pneumotoxic inhibitors must be blocked by barbiturate before apneusis can be induced by various experimental procedures. Krieger, Christensen, Sapru and Wang (105) have recently postulated the presence of another inhibitory afferent pathway in the cervical cord, which modulates pontine apneustic mechanisms. If this is a real entity, it represents another inhibitory input to the inspiratory system.

Results derived from single cell studies support



the conclusion that barbiturates differentially block inhibitory before facilitatory inputs. Figure 68 shows that inspiratory cells fire continuously during apneusis indicating that they are not depressed to an appreciable extent by barbiturate. Expiratory cells, on the other hand, are completely inhibited during the apneustic breath (Figure 69) due to direct barbiturate depression or more probably to reciprocal inhibition from constantly active inspiratory cells. Robson et al. (148) found similar results, but also described constant expiratory activity in some circumstances.

An interesting exception was observed for inspiratory cell C54UI13L as shown in Figure 84. A prolonged apneustic breath was produced one and one-half minutes after pentobarbital injection (new incremental dose level at 19.6 mg/kg). Unlike the inspiratory cell response in Figure 68, cell C54UI13L failed to fire continuously during the inspiratory hold. Rather, the unit turned off at a train length that was not significantly different from control. The cell recording was not lost at this time since the cell fired during succeeding breaths. This atypical unit discharge could arise from an inspiratory cell on the fringe of the I-VO population that was blocked by barbiturate or from a starter neuron

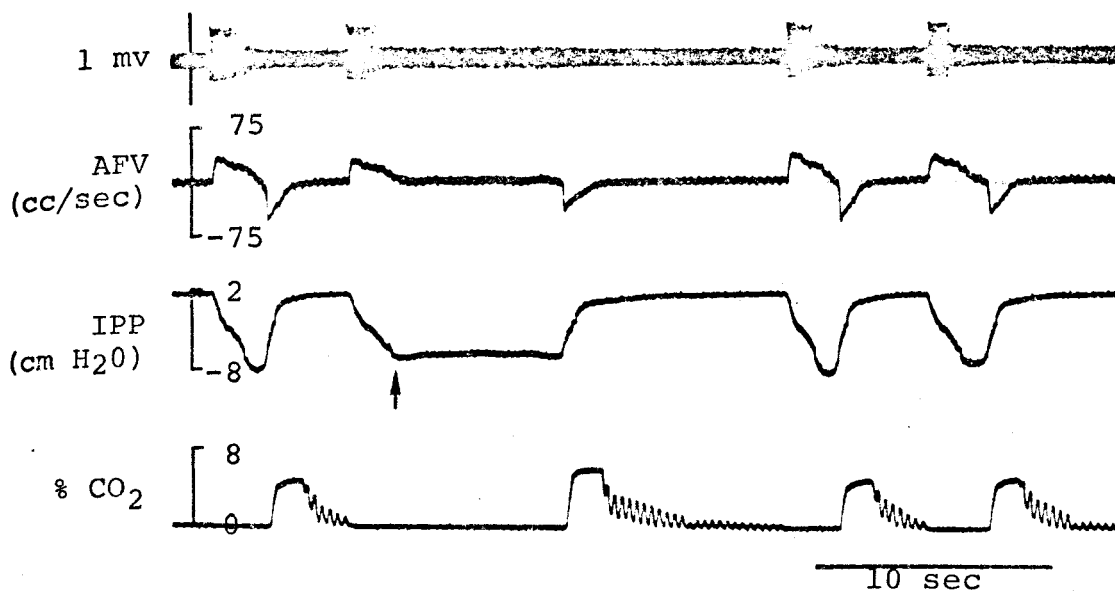


Figure 84. C54UI13L. Single inspiratory cell recording during induction of apneustic breath (arrow) 90 seconds after administration of 1.8 mg/Kg pentobarbital. Note failure of unit to maintain discharge for duration of the inspiratory hold as in Figure 68. Vagi sectioned.

or pacemaker cell responsible for activating the inspiratory complex. The latter speculation is an exciting possibility which merits further investigation.

The triphasic shift in the histogram distribution in Figures 70 and 71 shows that an inspiratory cell with vagi intact can, in fact, be activated during barbiturate accumulation. Since the respiration rate was consistently depressed, the coupling between rate and unit discharge frequency was destroyed. In normal cases, rate and cell discharge frequency change in the same direction (31). Therefore, this uncoupling at medium barbiturate levels represents a significant change in brainstem functional activity during drug depression. More specifically, by comparison of Figures 66 and 71, the inflection in the inspiratory curves in the latter figure can be attributed to pneumotaxic blockade. That is, the inflection occurred at a low barbiturate level (4-7 mg/kg thiopental) not sufficient to block vagal inhibitory feedback (Figure 66). Also, apneusis was not seen at this drug level.

Following apneustic breaths there is usually an elevated end-expiratory % CO<sub>2</sub> (Figures 64, 65, 67, 68, 69, 73, 74). It was suggested that this central chemodrive was of sufficient magnitude to periodically

interrupt inspiratory holds. This view is inconsistent with Stella's (168) data, which showed prolonged apneustic breaths induced by either CO<sub>2</sub> administration or re-breathing. On the other hand, Ngai (130) showed that 10% CO<sub>2</sub> accelerated the apneustic cycling, a response independent of peripheral chemoreceptor drives. Åström (9) has published excellent work on the combined action of CO<sub>2</sub> excess and O<sub>2</sub> deficiency in the regulation of breathing. The work of Mitchell and Herbert (123) should also be considered. These workers showed that CO<sub>2</sub> does have a direct effect on central inspiratory neurons in terms of rhythm generation (increased frequency discharge) although CO<sub>2</sub> does not change the resting membrane potential.

#### D. Information Transfer

The data presented in this dissertation have established certain quantitative relationships between respiratory cell discharge characteristics and medullary outputs of respiration rate and depth (1/MOD). It is suggested that if this study were repeated at the phrenic motoneuron or vagal afferent level, valuable information on gross properties of intersystem information transfer could be evaluated.

In a control systems approach, it would be ideal to derive transfer functions (ratio of output to input)

relating discharge pattern differences to synaptic processes intervening between successive stations of the respiratory arc. Due to numerous complexities inherent in the system, including proprioceptive modulation of rate and depth outputs at the spinal cord level (7, 54, 159, 160) and rate and depth interactions (37), the application of rigid mathematical criteria in transfer function derivation is quite difficult. Nevertheless, by comparison of idealized interspike interval modulation curves from various respiratory populations (expiratory, inspiratory, internuncial, phrenic, vagal afferent) it may be possible to initiate quantitative description of system couplings.

Gesell et al. (72) studied frequency modulation patterns of respiratory cells and found similar discharge patterns along major arcs of the respiratory system. For two of these components, interspike interval modulation curves for a vagal afferent and phrenic motoneuron were reconstructed in Figure 85A and B from the data of Adrian (2) and Pitts (142), respectively. These curves, generated from single train discharges, have a relatively smooth "U" shaped form which is characteristic of inspiratory and expiratory cells only after averaging of many successive bursts (cf. Figure 9). The regularity

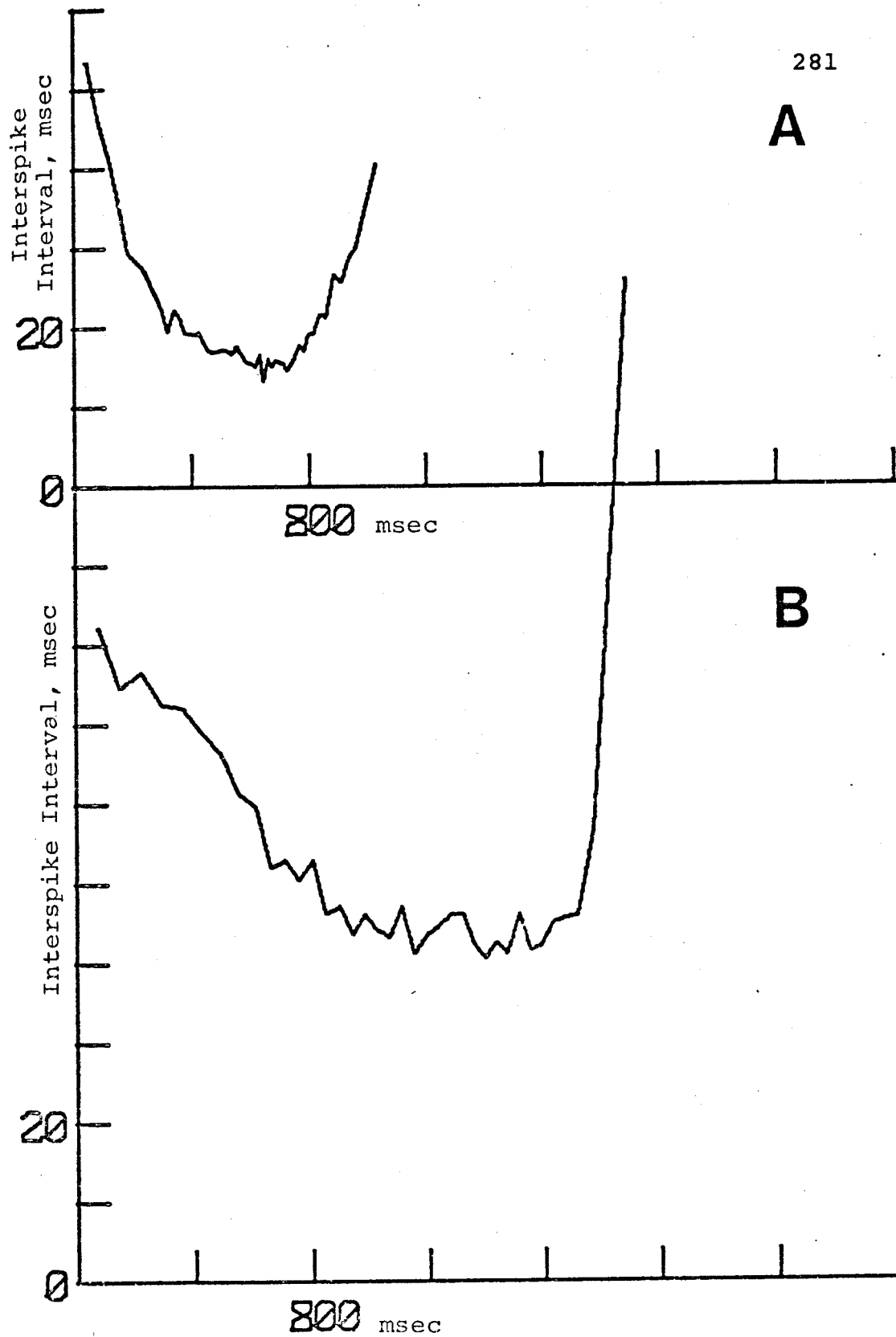


Figure 85. Interspike interval modulation curves for single train of vagal afferent (A) and phrenic motoneuron (B).

in vagal afferent interval ordering for a single burst (Figure 85A) derives from the highly predictable relationships between lung volume and vagal afferent discharge frequency (2). The regular phrenic pattern (Figure 85B) for a single burst indicates that this motoneuron is acting as a physiological integrator. Spatial summation of several respiratory neurons and/or spinal neurons arriving at phrenic cell bodies results in transformation of the variable medullary output into a smoothed pattern before proceeding to the respiratory musculature (75, 143). Comparison of Figure 85B (RR  $\sim$  15) with Figure 82A (Curve 1) shows that average medullary outputs also undergo frequency depression as they are converted into phrenic outputs. The phrenic motoneuron may act as a frequency converter to provide suitable impulse frequency to muscles of respiration for tidal volume control (144).

Dirken and Woldring (51) suggested that expiratory cells were activated by vagal afferents. If this is so, the inflation reflex which inhibits inspiration may be mediated through expiratory cells which have inhibitory inputs to inspiratory neurons. The data in Table III shows that only a few expiratory cells could be located when the vagus nerves were sectioned. Moreover,

Figure 62 shows that an expiratory discharge could be inhibited by vagotomy, suggesting that some kind of functional connection exists between vagal afferents and expiratory neurons. Assuming this to be true, the greater homogeneity of expiratory than inspiratory cells may be due to the dominance of highly regular input signals carried over vagal afferents (Figure 85A).

If one studies the respiratory system in terms of respiratory rate and depth outputs, it should be recognized that the transfer function for rate is equal to unity between serially-linked components. This is true since all components of the respiratory arc oscillate at the same frequency. The transfer function for depth, however, remains undefined since it is coded and recoded at each level of the respiratory system.



## CHAPTER VII

### CONCLUSION

1. The experiments presented in this dissertation were designed to investigate three aspects of central respiratory control utilizing computer techniques: (a) population characteristics of inspiratory and expiratory cells in the medulla; (b) characteristics of single elements in the above populations; (c) genesis of apneustic breathing by barbiturate administration and other experimental procedures.

2. The cells examined in these experiments comprised three neuronal populations: inspiratory, vagus nerves sectioned; inspiratory, vagus nerves intact; and expiratory, vagus nerves intact.

3. A new mathematical technique was devised to describe average interspike interval ordering of respiratory trains (interspike interval modulation curves) by temporal summation of consecutive trains discharging under steady state conditions. This technique provides

a capability for detailed analysis of the respiratory train, not possible by histogram analysis.

4. Twenty-five parameters, most of which have not been reported in the literature, were derived from the modulation curves. These parameters describe specific characteristics of the initial, middle and terminal phases of a respiratory train.

5. All parameters for similar populations were correlated with two functional respiratory outputs, rate and depth ( $\approx 1/\text{mode interval time}$ ) using regression analyses. The implications of similarities and dissimilarities in the response of inspiratory and expiratory cells were discussed.

6. Theoretical modulation curves were reconstructed from regression equations for each population. These curves predict characteristic alterations in respiratory cell discharge patterns as a function of rate and mode.

7. Single cell parameter responses were analyzed during experimental manipulations designed to modify respiratory rate and depth. Variations in discharge patterns, observed among many single units, were consistent with the expected variation within population distributions.

8. Various experimental procedures were found to

elicit apneustic breathing in cats under pentobarbital anesthesia. Additional doses of depressant drugs appeared to induce apneusis by differential inhibition of negative feedback loops in the brainstem. At appropriate barbiturate levels, decreasing carotid sinus pressure or doxapram administration elicited apneustic breathing, presumably by activation of facilitatory input. In other cases, vagotomy and cerebellectomy also produced apneusis.

9. The data in this study were discussed in the context of possible interneuronal connections within the brainstem respiratory complex.

## CHAPTER VIII

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CHAPTER IX

APPENDIX

Table VII. Core memory assignments for program DATA.

<u>LINC</u> <u>Segments</u>	<u>PMODE</u> <u>Fields</u>	<u>Absolute</u> <u>Location</u>	<u>Contents</u>
IF* 0	0	0000 - 0777 1000 - 1777	Decwriter Output X - Y Plotter Output
IF 1	0	2000 - 2777 } 3000 - 3777 }	X - Y Plotter Subroutine
IF 2	0	4000 - 4777 } 5000 - 5777 }	Mainline Program and Tape Storage Output
IF 3	0	6000 - 6777 } 7000 - 7777 }	CRT Display HISTO, ISIM, AXIS and TITLE
DF** 4	1	10000 - 10777 11000 - 11777	$\Sigma$ Interval Time (LO) Array, Bits 0-11 (LOOP B); ISIM Array (LOOP C) First Interval (FI) Array
DF 5	1	12000 - 12777 13000 - 13777	Divisor N (DIV) Array $\Sigma$ Interval Time (HI) Array; Bits 12-23 (LOOP B)
DF 6	1	14000 - 14777 15000 - 15777	Intervals per Train (ST-1) Array Respiration Rate (RR) Array
DF 7	1	16000 - 16777 17000 - 17777	HISTO Array Analog Input Array (LOOP A); ISIM Array (LOOP B)

\*Instruction Field

\*\*Data Field

```

0000          *20
0001          /PROGRAM ERASE
0002          /
0003          /ZERO STORED IN EACH WORD OF
0004          /1000 OCTAL BLOCK LINC MAGNETIC TAPE
0005          /
0006          0020 0067 ST,      SET I 7
0007          0021 3777          3777
0010          0022 0011          CLR
0011          0023 1067          STA I 7
0012          0024 0207          XSK 7
0013          0025 6023          JMP .-2
0014          0026 0000          HLT
0015          0027 1020          LDA I
0016          0030 4000          4000
0017          0031 4033          STC W0
0020          0032 0736 A,      WRI I U
0021          0033 0000 W0,    0
0022          0034 1020          LDA I
0023          0035 0001          1
0024          0036 1140          ADM
0025          0037 0033          W0
0026          0040 1460          SAE I
0027          0041 4777          4777
0030          0042 6032          JMP A
0031          0043 0712          RDE U
0032          0044 4000          4000
0033          0045 0000          HLT
0034          0046 0000          HLT
0035          0047 6020          JMP ST

```

## NO ERRORS

```

A      4032
ST     4020
W0     4033

```



```

0000          *20
0001          /PROGRAM DATA
0002          /
0003          /TIME SCALED 4X SLOWER
0004          /
0005          /LOOP A- DISPLAY CHANNEL 12 INPUT
0006          /LOOP B- HISTO- ISIM CONSTRUCTION
0007          /LOOP C- OUTPUT SELECTION
0010          /
0011          /ANALOG INPUTS
0012          /
0013          /12 A+B- SINGLE UNIT RECORDING
0014          /04 B - RATE OF RISE TRIGGER LEVEL
0015          /00 B - RESET DELAY 0-4095 MSEC
0016          /05 A - POSITION CHANNEL 12 DISPLAY
0017          /01 C - SHIFT HISTO DISPLAY U TO D
0020          /01 C" - SHIFT ISIM DISPLAY R TO L
0021          /
0022          /PROGRAM CONTROL
0023          /
0024          /LEFT SWITCHES - N FOR HISTOGRAM
0025          /RIGHT SWITCHES- DATA POINT NUMBER
0026          /
0027          /SENSE SWITCHES
0030          /
0031          /0 A - CHANNEL 12 DISPLAY FREEZE
0032          /0 B - MANUAL PULSE REJECTION
0033          /0 C - HISTO- ISIM DISPLAY SELECTION
0034          /0 C"- EXIT FROM ANY OUTPUT PROCEDURE
0035          /1 B - CONTINUOUS HISTOGRAM
0036          /1 C - START TELETYPE OUTPUT
0037          /2 C - START XY PLOT
0040          /3 C - START TAPE STORAGE
0041          /4 C - PROGRAM RESTART
0042          /5 A - EXIT CHANNEL 12 DISPLAY LOOP
0043          /5 B - EXIT HISTO- ISIM CONSTRUCT LOOP
0044          /5 B"- ENTER OUTPUT LOOP
0045          /5 C - COMPLETE TELY, XY OR TAPE OUTPUT
0046          /
0047          0020  0002          PDP
0050          PHODE
0051          4021  5046          TLS          /CLOCK PRESETS
0052          4022  6135          CLSA
0053          4023  7300          CLA CLL
0054          4024  1235          TAD KCLLR
0055          4025  6132          CLLR
0056          4026  7200          CLA
0057          4027  1236          TAD KCLAB
0060          4030  6133          CLAB
0061          4031  7200          CLA
0062          4032  1237          TAD KCLEN
0063          4033  6134          CLEN
0064          4034  5240          JMP GO
0065          4035  2100  KCLLR, 2100          /100K HZ MODE 1
0066          4036  7730  KCLAB,  -50          /PRESET -40
0067          4037  0300  KCLEN,  300          /ENABLE
0070          /
0071          /LOOP A
0072          /
0073          4040  6141  GO,          LINC
0074          LMCDE
0075          0041  0647          LDF 7

```

0076	0042	0073	CO,	SET I 13	/SAMPLE AND
0077	0043	2777		2777	/DISPLAY CHAN 12
0100	0044	0500	D,	IOB	
0101	0045	6131		6131	/CLOCK OVERFLOW
0102	0046	6044		JMP -2	/EVERY 0.1 MSEC
0103	0047	0500		IOB	
0104	0050	6135		6135	/CLSA
0105	0051	0112		SAM 12	
0106	0052	0341		SCR 1	
0107	0053	1073		STA I 13	
0110	0054	0213		XSK 13	
0111	0055	6044		JMP D	
0112	0056	0073	DS,	SET I 13	/START DISPLAY
0113	0057	2777		2777	
0114	0060	0445		SNS 5	
0115	0061	0456		SKP	
0116	0062	6102		JMP ST	/EXIT TO LOOP B
0117	0063	7727		JMP TITL	/TO TITLE
0120	0064	7727		JMP TITL	/ROUTINE
0121	0065	0105	SD,	SAM 5	
0122	0066	0341		SCR 1	
0123	0067	4972		STC +3	
0124	0070	1033		LDA I 13	
0125	0071	1120		ADA I	
0126	0072	0000		0	
0127	0073	0153		DIS 13	
0130	0074	0213		XSK 13	
0131	0075	6065		JMP SD	
0132	0076	0440		SNS 0	
0133	0077	0456		SKP	
0134	0100	6056		JMP DS	/FREEZE DISPLAY
0135	0101	6042		JMP CO	/RESAMPLE
0136			/		
0137	0102	0063	ST,	SET I 3	/CLEAR MEMORY
0140	0103	3777		3777	
0141	0104	0011		CLR	
0142	0105	0644		LDF 4	
0143	0106	1063		STA I 3	
0144	0107	0545		LDF 5	
0145	0110	1043		STA 3	
0146	0111	0546		LDF 6	
0147	0112	1043		STA 3	
0150	0113	0647		LDF 7	
0151	0114	1043		STA 3	
0152	0115	0203		XSK 3	
0153	0116	6105		JMP -11	
0154			/		
0155	0117	5760		STC R1	
0156	0120	5761		STC R2	
0157	0121	5762		STC R3	
0160	0122	5763		STC R4	
0161	0123	5766		STC M1	
0162	0124	5767		STC M2	
0163	0125	5770		STC M3	
0164	0126	5757		STC L510	
0165			/		
0166	0127	0517		LSW	/SELECT N FOR
0167	0130	1120		ADA I	/HISTO
0170	0131	0001		1	
0171	0132	4512		STC HN	
0172	0133	5755		STC ZRO	
0173			/		
0174	0134	0064		SET I 4	/FI

0175	0135	2777	2777	
0176	0136	0070	SET I 10	/RESP RATE
0177	0137	3000	3000	
0200	0140	6145	JMP .+5	
0201			/	
0202			/LOOP B	
0203			/	
0204	0141	0011	N, CLR	/REMOVE RESET
0205	0142	0647	LDF 7	/DELAY TIME
0206	0143	1052	STA 12	
0207	0144	0643	LDF 3	
0210			/	
0211	0145	0063	SET I 3	/LOW
0212	0146	3777	3777	
0213	0147	0065	SET I 5	/DIVISOR
0214	0150	3777	3777	
0215	0151	0066	SET I 6	/HIGH
0216	0152	3000	3000	
0217	0153	0072	SET I 12	/INTERVAL TIMES
0220	0154	3000	3000	
0221			/	
0222	0155	0223	NA, XSK I 3	/LOW
0223	0156	0226	XSK I 6	/HIGH
0224	0157	0647	LDF 7	
0225	0160	1032	LDA I 12	/INTERVAL TIMES
0226	0161	0643	LDF 3	
0227	0162	0450	AZE	
0230	0163	0456	SKP	
0231	0164	6231	JMP NN	/EXIT
0232	0165	5752	STC CONT	
0233			/	
0234	0166	0500	NC, IOB	/CLOCK OVERFLOW
0235	0167	6131	6131	/EVERY 0.1 MSEC
0236	0170	6211	JMP NJ	
0237	0171	0500	IOB	
0240	0172	6135	6135	/CLSA
0241			/	
0242	0173	1020	LDA I	
0243	0174	0001	1	
0244	0175	1140	ADM	
0245	0176	1755	ZRO	
0246	0177	1460	SRE I	
0247	0200	0144	144	/100
0250	0201	6211	JMP NJ	
0251			/	
0252	0202	1020	LDA I	/INCREMENT RATE
0253	0203	0001	1	/ARRAY EVERY
0254	0204	0546	LDF 6	/0.01 SEC
0255	0205	1150	ADM 10	
0256	0206	0643	LDF 3	
0257	0207	0011	CLR	
0260	0210	5755	STC ZRO	
0261			/	
0262	0211	1020	NJ, LDA I	/TRANSFER INT
0263	0212	0001	1	/TIMES
0264	0213	0644	LDF 4	
0265	0214	1203	LAM 3	
0266	0215	0643	LDF 3	
0267	0216	5747	STC GAR	
0270	0217	0645	LDF 5	
0271	0220	1206	LAM 6	
0272	0221	0643	LDF 3	
0273	0222	1020	LDA I	

0274	0223	7776	-1	
0275	0224	1140	ADM	
0276	0225	1752	CONT	
0277	0226	0450	AZE	
0300	0227	6166	JMP NC	/CONT TRANSFER
0301	0230	6155	JMP NA	/READ NEW INT
0302			/	
0303	0231	0072	NN,	SET I 12
0304	0232	3000		3000
0305	0233	0011	CLR	/CLEAR INTERVAL
0306	0234	0647	LDF 7	/TIMES ARRAY
0307	0235	1072	STA I 12	
0310	0236	0643	LDF 3	
0311	0237	0212	XSK 12	
0312	0240	6234	JMP .-4	
0313	0241	0072	SET I 12	
0314	0242	3000	3000	
0315			/	
0316	0243	1020	LDA I	
0317	0244	0377	377	
0320	0245	5742	STC 52	
0321	0246	5757	STC L510	
0322	0247	0603	LIF 3	
0323	0250	6040	JMP DISR	/DISPLAY R
0324			/	
0325	0251	0500	C,	IOB
0326	0252	6131	6131	/CLOCK OVERFLOW
0327	0253	6251	JMP .-2	/EVERY 0.1 HSEC
0330	0254	0500	IOB	
0331	0255	6135	6135	/CLSA
0332			/	
0333	0256	1020	LDA I	
0334	0257	0001	1	
0335	0260	1140	ADM	
0336	0261	1755	ZRO	
0337	0262	1460	SAE I	
0340	0263	0144	144	/100
0341	0264	6274	JMP .+10	
0342			/	
0343	0265	1020	LDA I	/INCREMENT RATE
0344	0266	0001	1	/ARRAY EVERY
0345	0267	0646	LDF 6	/0.01 SEC
0346	0270	1150	ADM 10	
0347	0271	0643	LDF 3	
0350	0272	0011	CLR	
0351	0273	5755	STC ZRO	
0352			/	
0353	0274	0440	SNS 0	/MANUAL PULSE
0354	0275	0456	SKP	/REJECTION
0355	0276	6251	JMP C	
0356			/	
0357	0277	6552	JMP TRIG	
0360	0300	0451	APD	
0361	0301	6251	JMP C	/NO TRIGGER
0362			/	
0363	0302	0603	LIF 3	/TRIGGER
0364	0303	6016	JMP DISS	/DISPLAY S
0365			/	
0366	0304	1000	QT,	LDA
0367	0305	0512	HN	
0370	0306	4314	STC .+6	
0371	0307	1000	LDA	/EXIT TO DISPLAY
0372	0310	1770	M3	/IF REACH HISTO

0373	0311	1120	ADA I	/N AND TRAIN
0374	0312	0001	1	/FINISHED
0375	0313	1460	SAE I	
0376	0314	0000	0	
0377	0315	0456	SKP	
0400	0316	6572	JMP RR	
0401				
0402	0317	0445	SNS 5	/MANUAL EXIT
0403	0320	0456	SKP	/TO DISPLAY
0404	0321	6572	JMP RR	
0405	0322	0230	XSK I 10	
0406	0323	0456	SKP	/EXIT IF MORE
0407	0324	6572	JMP RR	/THAN 510 TRAINS
0410				
0411	0325	1020	LDA I	
0412	0326	0001	1	
0413	0327	1140	ADM	/INCREMENT NUM
0414	0330	1763	R4	/TRAINS
0415				
0416	0331	1020	LDA I	/SET FI MARKER
0417	0332	0001	1	
0420	0333	5743	STC FI	
0421				
0422	0334	0225	XSK I 5	/DIVISOR
0423	0335	0232	XSK I 12	/INTERVAL TIMES
0424	0336	6342	JMP .+4	
0425	0337	1020	LDA I	/TRAIN LENGTH
0426	0340	0007	7	/>510 INTERVALS
0427	0341	5757	STC L510	/SET MARKER
0430				
0431	0342	1020	LDA I	
0432	0343	0377	377	
0433	0344	5742	STC S2	
0424	0345	5751	STC STOR	/OVERFLOW STORE
0435				
0436	0346	0500	IOB	/CLOCK OVERFLOW
0437	0347	6131	6131	/EVERY 0.1 MSEC
0440	0350	6346	JMP .-2	
0441	0351	0500	IOB	
0442	0352	6135	6135	/CLSA
0443				
0444	0353	1020	LDA I	/INCREMENT TIME
0445	0354	0001	1	/COUNTER
0446	0355	1140	ADM	
0447	0356	1751	STOR	
0450				
0451	0357	0100	SAM 0	/RESET DELAY
0452	0360	0451	APD	
0453	0361	0011	CLR	
0454	0362	0243	ROL 3	
0455	0363	1100	ADA	
0456	0364	1751	STOR	
0457	0365	1460	SAE I	/4095 0.1 MSEC
0460	0366	7777	7777	/MAX DELAY
0461	0367	0456	SKP	
0462	0370	6141	JMP N	/EXIT
0463				
0464	0371	1000	LDA	/BYPASS INTERVAL
0465	0372	1757	L510	/INCREMENT IF
0466	0373	1460	SAE I	/TRAIN >510 INT
0467	0374	0007	7	
0470	0375	0456	SKP	
0471	0376	6410	JMP .+12	

0472			LDA I	/INCREMENT INT
0473	0377	1020	1	/TIME
0474	0400	0001	LDF 7	
0475	0401	0647	ADM 12	
0476	0402	1152	LDF 3	
0477	0403	0643		
0500			LDA I	
0501	0404	1020	1	
0502	0405	0001	ADM	
0503	0406	1140	ZRO	
0504	0407	1755	SAE I	
0505	0410	1460	144	/100
0506	0411	0144	JMP +10	
0507	0412	6422		
0510			LDA I	/INCREMENT RATE
0511	0413	1020	1	/ARRAY EVERY
0512	0414	0001	LDF 6	/0.01 SEC
0513	0415	0646	ADM 10	
0514	0416	1150	LDF 3	
0515	0417	0643	CLR	
0516	0420	0011	STC ZRO	
0517	0421	5755		
0520			SNS 0	/MANUAL PULSE
0521	0422	0440	SKP	/REJECTION
0522	0423	0456	JMP N	
0523	0424	6141		
0524			JMP TRIG	
0525	0425	6552	APD	
0526	0426	0451	JMP A	/NO TRIGGER
0527	0427	6346		
0530			LDA	
0531	0430	1000	STOR	
0532	0431	1751	ADA I	/REJECT TRIGGER
0533	0432	1120	-24	/IF <2 MSEC
0534	0433	7753	APD	/SINCE LAST
0535	0434	0451	JMP A	/TRIGGER
0536	0435	6346		
0537			LIF 3	/TRIGGER
0540	0436	0603	JMP DIST	/DISPLAY T
0541	0437	6027		
0542			LDA	/FI?
0543	0440	1000	FJ	
0544	0441	1743	SAE I	
0545	0442	1460	1	
0546	0443	0001	JMP +10	/NO
0547	0444	6454	LDA	/YES, SAVE FI
0550	0445	1000	STOR	
0551	0446	1751	LDF 4	
0552	0447	0644	STA 1 4	
0553	0450	1064	LDF 3	
0554	0451	0643	CLR	
0555	0452	0011	STC FI	
0556	0453	5743		
0557			LDA	
0560	0454	1000	L510	
0561	0455	1757	SAE I	/BYPASS ISIM
0562	0456	1460	7	/CALCULATIONS
0563	0457	0007	SKP	/IF TRAIN
0564	0460	0456	JMP 0	/LENGTH>510 INT
0565	0461	6474		
0566			LDA I	/INCREMENT
0567	0462	1020	1	/DIVISOR
0570	0463	0001		

0571	0464	0645	LDF 5	
0572	0465	1145	ADM 5	
0573	0466	0643	LDF 3	
0574				
0575	0467	1020	LDA I	/INCREMENT N
0576	0470	0001	1	/FOR ISIM
0577	0471	1140	ADM	
0600	0472	1766	M1	
0601	0473	6476	JMP .+3	
0602				
0603	0474	0441	SNS 1	/CONTINUE TO
0604	0475	6342	JMP E	/FILL HISTO
0605	0476	1000	LDA	/BYPASS HISTO
0606	0477	1751	STOR	/CONSTRUCTION
0607	0500	1120	ADA I	/IF INTERVAL
0610	0501	6003	-1774	/TIME >102 MSEC
0611	0502	0451	APC	
0612	0503	0456	SKP	
0613	0504	6544	JMP MS	
0614				
0615	0505	1000	LDA	
0616	0506	1770	M3	
0617	0507	1120	ADA I	
0620	0510	0001	1	
0621	0511	1460	SAE I	
0622	0512	0000	0	
0623	0513	6522	JMP .+7	
0624	0514	1000	LDA	
0625	0515	1757	L510	
0626	0516	1460	SAE I	
0627	0517	0007	7	
0630	0520	6334	JMP B	
0631	0521	6141	JMP N	
0632	0522	1020	LDA I	/INCREMENT N
0633	0523	0001	1	/FOR HISTO
0634	0524	1140	ADM	
0635	0525	1770	M3	
0636				
0637	0526	1000	LDA	/CONSTRUCT HISTO
0640	0527	1751	STOR	/0.4 MSEC BINS
0641	0530	0341	SCR 1	
0642	0531	1560	BCL I	
0643	0532	0001	1	
0644	0533	1120	ADA I	
0645	0534	2000	2000	
0646	0535	4542	STC .+5	
0647	0536	1020	LDA I	
0650	0537	0001	1	
0651	0540	0647	LDF 7	
0652	0541	1140	ADM	
0653	0542	0000	0	
0654	0543	0643	LDF 3	
0655				
0656	0544	1000	LDA	
0657	0545	1757	L510	
0660	0546	1460	SAE I	
0661	0547	0007	7	
0662	0550	6334	JMP B	/RETURN & BUMP
0663	0551	6342	JMP E	/RETURN- NO BUMP
0664				
0665	0552	1000	LDA	/TRIGGER
0666	0553	1742	S2	/ROUTINE
0667	0554	0017	COM	

0670	0555	5741	STC	51	
0671	0556	0104	SAM	4	
0672	0557	0341	SCR	1	
0673	0560	0017	COM		
0674	0561	5746	STC	TRG	
0675	0562	0112	SAM	12	
0676	0563	0341	SCR	1	
0677	0564	5742	STC	52	
0700	0565	1000	LDA		
0701	0566	1741	S1		
0702	0567	3742	ADD	52	
0703	0570	3746	ADD	TRG	
0704	0571	6000	JMP	0	/RETURN
0705					
0706	0572	0063	RR,	SET I 3	/LOW
0707	0573	3777		3777	
0710	0574	0065		SET I 5	/DIVISOR
0711	0575	3777		3777	
0712	0576	0066		SET I 6	/HIGH
0713	0577	3000		3000	
0714					
0715	0600	0645	LDF	5	
0716	0601	1025	LDA	I 5	
0717	0602	0643	LDF	3	
0720	0603	0450	AZE		
0721	0604	0456	SKP		/EXIT TO DISPLAY
0722	0605	7525	JMP	KR	/IF DIV=0
0723	0606	0065	SET	I 5	
0724	0607	3777		3777	
0725					
0726	0610	0645	RR,	LDF 5	/CALCULATE MEAN
0727	0611	1025		LDA I 5	/INTERVAL TIMES
0730	0612	0643		LDF 3	
0731	0613	0450		AZE	
0732	0614	0456		SKP	
0733	0615	6634		JMP RP	/EXIT
0734	0616	5015		STC KD	
0735	0617	0644		LDF 4	
0736	0620	1023		LDA I 3	/LOW
0737	0621	0643		LDF 3	
0740	0622	5016		STC KL	
0741	0623	0645		LDF 5	
0742	0624	1026		LDA I 6	/HIGH
0743	0625	0643		LDF 3	
0744	0626	5017		STC KH	
0745	0627	7000		JMP DIV	
0746	0630	0644		LDF 4	
0747	0631	1043		STA 3	/RESTORE AVG
0750	0632	0643		LDF 3	
0751	0633	6610		JMP RK	
0752					
0753	0634	0070	RP,	SET I 10	/CALCULATE
0754	0635	3000		3000	/INDIVIDUAL
0755	0636	0646	PR,	LDF 6	/RESP RATES
0756	0637	1030		LDA I 10	
0757	0640	0643		LDF 3	
0760	0641	0450		AZE	
0761	0642	0456		SKP	
0762	0643	6660		JMP RA	/EXIT
0763	0644	5015		STC KD	
0764	0645	1020		LDA I	
0765	0646	5140		5140	/60000 0.01
0766	0647	5016		STC KL	/SEC/10 MIN



0767	0650	1020		LDA I	
0770	0651	0016		16	
0771	0652	5017		STC KH	
0772	0653	7000		JMP DIV	
0773	0654	0646		LDF 6	
0774	0655	1050		STA 10	
0775	0656	0643		LDF 3	
0776	0657	6636		JMP PR	
0777					
1000	0660	0070	RA,	SET I 10	/CALCULATE MEAN
1001	0661	3000		3000	/RESP RATE
1002	0662	0646	RA,	LDF 6	
1003	0663	1030		LDA I 10	
1004	0664	0643		LDF 3	
1005	0665	0450		AZE	
1006	0666	0456		SKP	
1007	0667	6707		JMP DV	
1010	0670	5751		STC STOR	
1011	0671	1020	RU,	LDA I	
1012	0672	0001		1	
1013	0673	1200		LAM	
1014	0674	1762		R3	
1015	0675	5747		STC GAR	
1016	0676	1200		LAM	
1017	0677	1761		R2	
1020	0700	1020		LDA I	
1021	0701	7776		-1	
1022	0702	1140		ADM	
1023	0703	1751		STOR	
1024	0704	0450		AZE	
1025	0705	6671		JMP RU	
1026	0706	6662		JMP RN	
1027	0707	1000	DV,	LDA	
1030	0710	1763		R4	
1031	0711	5015		STC KD	
1032	0712	1000		LDA	
1033	0713	1762		R3	
1034	0714	5016		STC KL	
1035	0715	1000		LDA	
1036	0716	1761		R2	
1037	0717	5017		STC KH	
1040	0720	7000		JMP DIV	
1041	0721	1040		STA	
1042	0722	1760		R1	/MEAN RESP RATE
1043					
1044	0723	0070		SET I 10	/FIND HIGH AND
1045	0724	3000		3000	/LOW RESP RATE
1046	0725	0646		LDF 6	
1047	0726	1030		LDA I 10	
1050	0727	0643		LDF 3	
1051	0730	1040		STA	
1052	0731	1762		R3	/LOW RR
1053	0732	1040		STA	
1054	0733	1761		R2	/HIGH RR
1055	0734	0646	TR,	LDF 6	
1056	0735	1030		LDA I 10	
1057	0736	0643		LDF 3	
1060	0737	0450		AZE	
1061	0740	0456		SKP	
1062	0741	7070		JMP KX	/EXIT
1063	0742	0017		COM	
1064	0743	1100		ADM	
1065	0744	1761		R2	/HIGH RR

1066	0745	0451		APD	
1067	0746	0456		SKP	
1070	0747	0756		JMP .+7	
1071	0750	0646		LDF 6	
1072	0751	1010		LDA 10	
1073	0752	0643		LDF 3	
1074	0753	1010		STA	
1075	0754	1761		R2	/HIGH RR
1076	0755	0734		JMP TR	
1077	0756	0646		LDF 6	
1100	0757	1010		LDA 10	
1101	0760	0643		LDF 3	
1102	0761	0017		COH	
1103	0762	1100		ADA	
1104	0763	1762		R3	/LOW RR
1105	0764	0451		APD	
1106	0765	0734		JMP TR	
1107	0766	0646		LDF 6	
1110	0767	1010		LDA 10	
1111	0770	0643		LDF 3	
1112	0771	1010		STA	
1113	0772	1762		R3	/LOW RR
1114	0773	0734		JMP TR	/EXIT
1115					
1116				*1000	
1117	1000	0002	DIY,	PDP	/DIVIDE
1120				PMODE	/ROUTINE
1121	5001	7300		CLA CLL	
1122	5002	1215		TAD KD	
1123	5003	3210		DCR .+5	
1124	5004	1216		TAD KL	
1125	5005	7421		MOL	
1126	5006	1217		TAD KH	
1127	5007	7407		DVI	
1130	5010	0000		0	
1131	5011	7200		CLA	
1132	5012	7501		MGR	
1133	5013	6141		LINC	
1134				LMODE	
1135	1014	0000		JMP 0	/RETURN
1136	1015	0000	ED,	0	
1137	1016	0000	KL,	0	
1140	1017	0000	KH,	0	
1141					
1142	1020	0064	KX,	SET I 4	/CORRECT FI
1143	1021	2777		2777	/ARRAY
1144	1022	0644	KX2,	LDF 4	
1145	1023	1024		LDA I 4	
1146	1024	0643		LDF 3	
1147	1025	0450		AZE	
1150	1026	0456		SKP	
1151	1027	7036		JMP AB	
1152	1030	1020		LDA I	
1153	1031	0001		1	
1154	1032	0644		LDF 4	
1155	1033	1144		ADM 4	
1156	1034	0643		LDF 3	
1157	1035	7022		JMP KX2	
1160					
1161	1036	0063	AB,	SET I 3	/CORRECT FI
1162	1037	2000		2000	
1163	1040	1020		LDA I	
1164	1041	0001		1	

1165	1042	0644		LDF 4	
1166	1043	1143		ADM 3	
1167	1044	0643		LDF 3	
1170					
1171	1045	0065		SET I 5	/CALCULATE ST
1172	1046	2000		2000	
1173	1047	0070		SET I 10	
1174	1050	3765		3765	
1175	1051	0645		LDF 5	
1176	1052	1005		LDA 5	/NUM TRAINS
1177	1053	0643		LDF 3	
1200	1054	5015		STC KD	
1201	1055	5016		STC KL	
1202	1056	5017		STC KH	
1203	1057	1000	571,	LDA	
1204	1060	1706		M1	/FMN
1205	1061	5752		STC CONT	
1206	1062	1020	572,	LDA I	
1207	1063	0001		1	
1210	1064	1200		LAM	
1211	1065	1016		KL	
1212	1066	5747		STC GAR	
1213	1067	1200		LAM	
1214	1070	1017		KH	
1215	1071	1020		LDA I	
1216	1072	7776		-1	
1217	1073	1140		ADM	
1220	1074	1752		CONT	
1221	1075	0450		AZE	
1222	1076	7062		JMP ST2	
1223	1077	0230		XSK I 10	
1224	1100	7057		JMP ST1	
1225	1101	7000		JMP DIV	
1226	1102	1120		ADA I	
1227	1103	0912		12	
1230	1104	5764		STC R5	/ST
1231					
1232	1105	0063		SET I 3	/FIND MI
1233	1106	2000		2000	
1234	1107	0644		LDF 4	
1235	1110	1003		LDA 3	
1236	1111	0643		LDF 3	
1237	1112	1040		STA	
1240	1113	1765		R6	/MI
1241	1114	0644	MI1,	LDF 4	
1242	1115	1023		LDA I 3	
1243	1116	0643		LDF 3	
1244	1117	0450		AZE	
1245	1120	0456		SKP	
1246	1121	7134		JMP AT	/EXIT
1247	1122	0017		COM	
1250	1123	1100		ADA	
1251	1124	1765		R6	/MI
1252	1125	0451		RPO	
1253	1126	7114		JMP MI1	
1254	1127	0644		LDF 4	
1255	1130	1003		LDA 3	
1256	1131	0643		LDF 3	
1257	1132	5765		STC R6	/MI
1260	1133	7114		JMP MI1	
1261					
1262	1134	0011	AT,	CLR	/CALCULATE
1263	1135	5751		STC STOR	/NUMBER OF

1264	1136	0065	SET I 5	/INTERVALS PER
1265	1137	2000	2000	/TRAIN
1266	1140	0067	SET I 7	
1267	1141	3777	3777	
1270	1142	1020	LDA I	
1271	1143	0001	1	
1272	1144	1140	ADM	
1273	1145	1751	STOR	
1274	1146	0645	LDF 5	
1275	1147	1005	LDA 5	
1276	1150	0643	LDF 3	
1277	1151	0450	AZE	
1300	1152	0456	SKP	
1301	1153	7175	JMP KQ	/EXIT
1302	1154	0017	COM	
1303	1155	0645	LDF 5	
1304	1156	1125	ADA I 5	
1305	1157	0643	LDF 3	
1306	1160	0450	AZE	
1307	1161	0456	SKP	
1310	1162	7142	JMP KQ	
1311	1163	0017	COM	
1312	1164	0646	LDF 6	
1313	1165	1067	STA I 7	/NUM TRAINS
1314	1166	0643	LDF 3	
1315	1167	1000	LDA	
1316	1170	1751	STOR	
1317	1171	0646	LDF 6	
1320	1172	1067	STA I 7	/NUM INTERVALS
1321	1173	0643	LDF 3	
1322	1174	7142	JMP KQ	
1323				
1324	1175	0071	SET I 11	/SUM 0.4 MSEC
1325	1176	2001	2001	/BINS TO FORM
1326	1177	0072	SET I 12	/0.8 MSEC BINS
1327	1200	3777	3777	
1330	1201	0073	SET I 13	
1331	1202	1577	1577	
1332	1203	0647	LDF 7	
1333	1204	1031	LDA I 11	
1334	1205	0231	XSK I 11	
1335	1206	0016	NOP	
1336	1207	1131	ADA I 11	
1337	1210	1072	STA I 12	
1340	1211	0643	LDF 3	
1341	1212	0231	XSK I 11	
1342	1213	0016	NOP	
1343	1214	0233	XSK I 13	
1344	1215	7203	JMP .-12	
1345				
1346	1216	0063	SET I 3	/FIND MEAN
1347	1217	3777	3777	
1350	1220	0067	SET I 7	
1351	1221	3777	3777	
1352	1222	0011	CLR	
1353	1223	5016	STC KL	
1354	1224	5017	STC KH	
1355	1225	0646	LDF 6	
1356	1226	1027	LDA I 7	
1357	1227	0643	LDF 3	
1360	1230	5753	STC NOM	
1361	1231	0646	LDF 6	
1362	1232	1027	LDA I 7	

1363	1233	0643	LDF 3	
1364	1234	0450	AZE	
1365	1235	0456	SKP	
1366	1236	7313	JMP MN4	/EXIT
1367	1237	5754	STC LEN	
1370	1240	0644	LDF 4	
1371	1241	1023	LDA I 3	
1372	1242	0643	LDF 3	
1373	1243	5752	STC CONT	
1374	1244	1020	LDA I	
1375	1245	0001	1	
1376	1246	1200	LAM	
1377	1247	1016	KL	
1400	1250	5747	STC GAR	
1401	1251	1200	LAM	
1402	1252	1017	KH	
1403	1253	1460	SAE I	
1404	1254	7777	7777	
1405	1255	7261	JMP .+4	
1406	1256	0011	CLR	/CLR MEAN IF
1407	1257	5773	STC T2	/SUM > 24 BITS
1410	1260	7320	JMP M0	/EXIT
1411	1261	1020	LDA I	
1412	1262	7776	-1	
1413	1263	1140	ADM	
1414	1264	1752	CONT	
1415	1265	0450	AZE	
1416	1266	7244	JMP MN3	
1417	1267	1020	LDA I	
1420	1270	7776	-1	
1421	1271	1140	ADM	
1422	1272	1754	LEN	
1423	1273	0450	AZE	
1424	1274	7240	JMP MN2	
1425	1275	0063	SET I 3	
1426	1276	3777	3777	
1427	1277	1020	LDA I	
1430	1300	7776	-1	
1431	1301	1140	ADM	
1432	1302	1753	NOM	
1433	1303	0450	AZE	
1434	1304	0456	SKP	
1435	1305	7225	JMP MN1	
1436	1306	0646	LDF 6	
1437	1307	1007	LDA 7	
1440	1310	0643	LDF 3	
1441	1311	5754	STC LEN	
1442	1312	7240	JMP MN2	
1443	1313	1000	LDA	
1444	1314	1766	M1	/FHN
1445	1315	5015	STC KD	
1446	1316	7000	JMP DIV	
1447	1317	5773	STC T2	/MEAN
1450			/	
1451	1320	0071	MO, SET I 11	/FIND NODE
1452	1321	3777	3777	
1453	1322	0073	SET I 13	
1454	1323	1677	1677	
1455	1324	0647	LDF 7	
1456	1325	1031	LDA I 11	
1457	1326	0643	LDF 3	
1460	1327	5772	STC T1	
1461	1330	1020	LDA I	

1462	1331	0004		4	
1463	1332	5751		STC STOR	
1464	1333	0233	KP,	XSK I 13	
1465	1334	0456		SKP	
1466	1335	7363		JMP KS	/EXIT
1467	1336	1020		LDA I	
1470	1337	0010		10	/INCREMENT BY
1471	1340	1140		ADM	/0.8 MSEC
1472	1341	1751		STOR	
1473	1342	1000		LDA	
1474	1343	1772		T1	/MON
1475	1344	0017		COM	
1476	1345	0647		LDF 7	
1477	1346	1131		ADR I 11	
1500	1347	0643		LDF 3	
1501	1350	0451		APO	
1502	1351	7333		JMP KP	
1503	1352	0647		LDF 7	
1504	1353	1011		LDA 11	
1505	1354	0643		LDF 3	
1506	1355	5772		STC T1	/MON
1507	1356	1000		LDA	
1510	1357	1751		STOR	
1511	1360	1040		STA	
1512	1361	1767		M2	/MOD
1513	1352	7333		JMP KP	
1514			/		
1515	1363	0071	KS,	SET I 11	/FIND MEDIAN
1516	1364	3777		3777	
1517	1365	1000		LDA	
1520	1366	1770		M3	/HSN
1521	1367	0341		SCR 1	
1522	1370	1560		BCL I	
1523	1371	4000		4000	
1524	1372	5745		STC MD	
1525	1373	1020		LDA I	
1526	1374	0004		4	
1527	1375	1040		STA	
1530	1376	1771		M4	/MED
1531	1377	0647	KA,	LDF 7	
1532	1400	1031		LDA I 11	
1533	1401	0643		LDF 3	
1534	1402	0017		COM	
1535	1403	1140		ADM	
1536	1404	1745		MD	
1537	1405	0451		APO	
1540	1406	7414		JHP KE	/EXIT
1541	1407	1020		LDA I	
1542	1410	0010		10	
1543	1411	1140		ADM	
1544	1412	1771		M4	/MED
1545	1413	7377		JHP KA	
1546			/		
1547	1414	0065	KE,	SET I 5	/STORE CONSTANTS
1550	1415	2000		2000	/AT END OF TL
1551	1416	0067		SET I 7	/QUARTER 4
1552	1417	2340		2340	
1553	1420	0063		SET I 3	
1554	1421	2000		2000	
1555	1422	1000		LDA	
1556	1423	1760		R1	/MEAN RR
1557	1424	0647		LDF 7	
1560	1425	1047		STA 7	/2340-RR

1561	1426	0645	LDF 5	
1562	1427	1005	LDA 5	/NT
1563	1430	0647	LDF 7	
1564	1431	1067	STA I 7	/2341-NT
1565	1432	0643	LDF 3	
1566	1433	1000	LDA	
1567	1434	1764	R5	/ST
1570	1435	0647	LDF 7	
1571	1436	1067	STA I 7	/2342-ST
1572	1437	0011	CLR	
1573	1440	1067	STA I 7	/2343-SD
1574	1441	1067	STA I 7	/2344-SE
1575	1442	0644	LDF 4	
1576	1443	1003	LDA 3	/FI
1577	1444	0647	LDF 7	
1600	1445	1067	STA I 7	/2345-FI
1601	1446	0011	CLR	
1602	1447	1067	STA I 7	/2346-SD
1603	1450	1067	STA I 7	/2347-SE
1604	1451	0643	LDF 3	
1605	1452	1000	LDA	
1606	1453	1765	R6	/MI
1607	1454	0647	LDF 7	
1610	1455	1067	STA I 7	/2350-MI
1611	1456	0643	LDF 3	
1612	1457	1000	LDA	
1613	1460	1766	M1	/ISIM N
1614	1461	0647	LDF 7	
1615	1462	1067	STA I 7	/2351-FMN
1616	1463	0643	LDF 3	
1617	1464	1000	LDA	
1620	1465	1770	M3	/HISTO N
1621	1466	0647	LDF 7	
1622	1467	1067	STA I 7	/2352-HSN
1623	1470	0643	LDF 3	
1624	1471	1000	LDA	
1625	1472	1773	T2	/MEAN
1626	1473	0647	LDF 7	
1627	1474	1067	STA I 7	/2353-MEN
1630	1475	0643	LDF 3	
1631	1476	1000	LDA	
1632	1477	1771	M4	/MEDIAN
1633	1500	0647	LDF 7	
1634	1501	1067	STA I 7	/2354-MED
1635	1502	0643	LDF 3	
1636	1503	1000	LDA	
1637	1504	1767	M2	/HISTO MODE
1640	1505	0647	LDF 7	
1641	1506	1067	STA I 7	/2355-MOD
1642	1507	0643	LDF 3	
1643	1510	1000	LDA	
1644	1511	1772	T1	/MON
1645	1512	0647	LDF 7	
1646	1513	1067	STA I 7	/2356-MON
1647	1514	0011	CLR	
1650	1515	1067	STA I 7	
1651	1516	0207	XSK 7	
1652	1517	7515	JMP -2	
1653	1520	0643	LDF 3	
1654				
1655	1521	0445	SNS 5	/HLT UNTIL
1656	1522	7525	JMP KR	/RESET SNS 5
1657	1523	0603	LIF 3	

1660	1524	6104		JMP DISS	
1661			/		
1662	1525	7734	KR,	JMP DSPY	/TO DISPLAY
1663	1526	7734		JMP DSPY	/ROUTINE
1664	1527	0603		LIF 3	
1665	1530	6366		JMP AXIS	/TO AXIS DISPLAY
1666			/		
1667	1531	7727	KJ,	JMP TITL	/TO TITLE
1670	1532	7727		JMP TITL	/ROUTINE
1671	1533	0441		SNS 1	
1672	1534	7547		JMP KW	
1673	1535	0440		SNS 0	
1674	1536	0456		SKP	
1675	1537	7547		JMP KW	/EXIT
1676	1540	0603		LIF 3	
1677	1541	6051		JMP DIS1	
1700	1542	0445	BB1,	SNS 5	
1701	1543	7535		JMP .-6	
1702	1544	0002		PDP	/TO TELETYPE
1703				PMODE	/OUTPUT
1704	5545	4422		JMS I T00	
1705	5546	6141		LINC	
1706				LMODE	
1707			/		
1710	1547	0442	KW,	SNS 2	
1711	1550	7562		JMP KF	
1712	1551	0440		SNS 0	
1713	1552	0456		SKP	
1714	1553	7562		JMP KF	/EXIT
1715	1554	0603		LIF 3	
1716	1555	6062		JMP DIS2	
1717	1556	0445	BB2,	SNS 5	
1720	1557	7551		JMP .-6	
1721	1560	0002		PDP	/TO XY PLOT
1722				PMODE	
1723	5561	4405		JMS I XY	
1724				LMODE	
1725			/		
1726	1562	0443	KF,	SNS 3	/TAPE STORAGE
1727	1563	7724		JMP KU	
1730	1564	0440		SNS 0	
1731	1565	0456		SKP	
1732	1566	7724		JMP KU	/EXIT
1733	1567	0603		LIF 3	
1734	1570	6073		JMP DIS3	
1735	1571	0445	BB3,	SNS 5	
1736	1572	7564		JMP .-6	/DELAY
1737			/		
1740	1573	1020		LDA I	/CHECK FOR
1741	1574	6000		6000	/OVER WRITING
1742	1575	5622		STC W00	
1743			/		
1744	1576	0516		RSH	
1745	1577	1560		BCL I	
1746	1600	4000		4000	
1747	1601	1120		ADA I	
1750	1602	7632		-145	/RETURN IF
1751	1603	0451		AP0	/DATA POINT
1752	1604	0456		SKP	/NUMBER > 101
1753	1605	7571		JMP BB3	
1754			/		
1755	1606	0516		RSH	/READ DATA POINT
1756	1607	1040		STA	/NUMBER AND MULT



Address	Hex	Hex	Label	Comment
1757	1610	1613	.+3	
1760	1611	0242	ROL 2	
1761	1612	1120	ADA 1	
1762	1613	0000	0	
1763	1614	1040	STA	
1764	1615	1750	BLK	
1765	1616	1140	ADM	
1766	1617	1622	WT0	
1767	1620	0647	LDF 7	
1770	1621	0712	RDE U	
1771	1622	0000	0	
1772	1623	0072	SET I 12	
1773	1624	3000	3000	
1774	1625	1012	LDA 12	/FI OF ARRAY
1775	1626	0643	LDF 3	
1776	1627	0450	AZE	
1777	1630	0456	SKP	
2000	1631	7641	JMP SV	/BLOCKS EMPTY
2001				
2002	1632	0440	SNS 0	/BLOCKS FILLED
2003	1633	0456	SKP	
2004	1634	7724	JMP KU	/EXIT
2005	1635	0603	LIF 3	
2006	1636	6115	JMP DISF	
2007	1637	0445	SNS 5	
2010	1640	7632	JMP .-6	
2011				
2012	1641	1020	LDA I	/STORE DATA
2013	1642	4000	4000	
2014	1643	5652	STC W0	
2015	1644	1000	LDA	
2016	1645	1750	BLK	
2017	1646	1140	ADM	
2020	1647	1652	W0	
2021	1650	0644	LDF 4	
2022	1651	0734	WRC I U	/FIRST HALF FM
2023	1652	0000	0	
2024	1653	0643	LDF 3	
2025	1654	1020	LDA I	
2026	1655	1001	1001	
2027	1656	1140	ADM	
2030	1657	1652	W0	
2031	1660	5653	STC W1	
2032	1661	0644	LDF 4	
2033	1662	0734	WRC I U	/LAST HALF FM
2034	1663	0000	0	
2035	1664	0643	LDF 3	
2036				
2037	1665	1020	LDA I	
2040	1666	1001	1001	
2041	1667	1140	ADM	
2042	1670	1663	W1	
2043	1671	5674	STC W2	
2044	1672	0644	LDF 4	
2045	1673	0734	WRC I U	/FI
2046	1674	0000	0	
2047	1675	0643	LDF 3	
2050				
2051	1676	1020	LDA I	
2052	1677	4003	4003	
2053	1700	5707	STC W3	
2054	1701	1000	LDA	
2055	1702	1750	BLK	

2056	1703	1140		ADM	
2057	1704	1707		W3	
2060	1705	0646		LDF 6	
2061	1706	0734		WRC I U	/TL
2062	1707	0000	W3,	0	
2063	1710	0643		LDF 3	
2064			/		
2065	1711	1020		LDA I	
2066	1712	4004		4004	
2067	1713	5722		STC W4	
2070	1714	1000		LDA	
2071	1715	1750		BLK	
2072	1716	1140		ADM	
2073	1717	1722		W4	
2074	1720	0647		LDF 7	
2075	1721	0714		WRC U	/HISTO AND
2076	1722	0000	W4,	0	/CONSTANTS
2077	1723	0643		LDF 3	
2100			/		
2101	1724	0444	KU,	SNS 4	
2102	1725	7525		JMP KR	/DISPLAY AGAIN
2103	1726	6102		JMP ST	/RESTART
2104			/		
2105	1727	1000	TITL,	LDA	/SAVE RETURN
2106	1730	0000		0	/ADDRESS
2107	1731	5756		STC RETN	
2110	1732	0603		LIF 3	
2111	1733	6443		JMP LET	
2112			/		
2113	1734	1000	DSPV,	LDA	/SAVE RETURN
2114	1735	0000		0	/ADDRESS
2115	1736	5756		STC RETN	
2116	1737	0603		LIF 3	
2117	1740	6126		JMP DISP	
2120			/		
2121	1741	0000	S1,	0	
2122	1742	0000	S2,	0	
2123	1743	0000	FI,	0	
2124	1744	0000	RC,	0	
2125	1745	0000	MD,	0	
2126	1746	0000	TRG,	0	
2127	1747	0000	GAR,	0	
2130	1750	0000	BLK,	0	
2131	1751	0000	STOR,	0	
2132	1752	0000	CONT,	0	
2133	1753	0000	NOM,	0	
2134	1754	0000	LEN,	0	
2135	1755	0000	ZRO,	0	
2136	1756	0000	RETN,	0	
2137	1757	0000	LS10,	0	
2140	1760	0000	R1,	0	
2141	1761	0000	R2,	0	
2142	1762	0000	R3,	0	
2143	1763	0000	R4,	0	
2144	1764	0000	R5,	0	
2145	1765	0000	R6,	0	
2146	1766	0000	H1,	0	
2147	1767	0000	H2,	0	
2150	1770	0000	H3,	0	
2151	1771	0000	H4,	0	
2152	1772	0000	T1,	0	
2153	1773	0000	T2,	0	
2154			/		

2155				PMODE	
2156				*6014	/IF 3
2157				LMODE	
2160	0014	1000		1000	/PRESETS FOR
2161	0015	1000		1000	/TITLE ROUTINE
2162					
2163	0016	0011	DIS5,	CLR	/DISPLAY S
2164	0017	0061		SET I 1	
2165	0020	0364		364	
2166	0021	0062		SET I 2	
2167	0022	0646		646	
2170	0023	1742		DSC 2	
2171	0024	1762		DSC I 2	
2172	0025	0602		LIF 2	
2173	0026	6304		JMP QT	
2174					
2175	0027	0011	DIST,	CLR	/DISPLAY T
2176	0030	0061		SET I 1	
2177	0031	0364		364	
2200	0032	0062		SET I 2	
2201	0033	0650		650	
2202	0034	1742		DSC 2	
2203	0035	1762		DSC I 2	
2204	0036	0602		LIF 2	
2205	0037	6440		JMP TB	
2206					
2207	0040	0011	DISR,	CLR	/DISPLAY R
2210	0041	0061		SET I 1	
2211	0042	0364		364	
2212	0043	0062		SET I 2	
2213	0044	0644		644	
2214	0045	1742		DSC 2	
2215	0046	1762		DSC I 2	
2216	0047	0602		LIF 2	
2217	0050	6251		JMP C	
2220					
2221	0051	0011	DIS1,	CLR	/DISPLAY W
2222	0052	0061		SET I 1	
2223	0053	0364		364	
2224	0054	0062		SET I 2	
2225	0055	0656		656	
2226	0056	1742		DSC 2	
2227	0057	1762		DSC I 2	
2230	0060	0602		LIF 2	
2231	0061	7542		JMP BB1	
2232					
2233	0062	0011	DIS2,	CLR	/DISPLAY P
2234	0063	0061		SET I 1	
2235	0064	0364		364	
2236	0065	0062		SET I 2	
2237	0066	0640		640	
2240	0067	1742		DSC 2	
2241	0070	1762		DSC I 2	
2242	0071	0602		LIF 2	
2243	0072	7556		JMP BB2	
2244					
2245	0073	0011	DIS3,	CLR	/DISPLAY A
2246	0074	0061		SET I 1	
2247	0075	0364		364	
2250	0076	0062		SET I 2	
2251	0077	0602		602	
2252	0100	1742		DSC 2	
2253	0101	1762		DSC I 2	

2254	0102	0602	LIF 2	
2255	0103	7571	JMP BB3	
2256			/	
2257	0104	0011	DISS, CLR	/DISPLAY 5
2260	0105	0061	SET 1 1	
2261	0106	0364	364	
2262	0107	0062	SET 1 2	
2263	0110	0752	752	
2264	0111	1742	DSC 2	
2265	0112	1762	DSC 1 2	
2266	0113	0602	LIF 2	
2267	0114	7521	JMP BB5	
2270			/	
2271	0115	0011	DISF, CLR	/DISPLAY F
2272	0116	0061	SET 1 1	
2273	0117	0364	364	
2274	0120	0062	SET 1 2	
2275	0121	0614	614	
2276	0122	1742	DSC 2	
2277	0123	1762	DSC 1 2	
2300	0124	0602	LIF 2	
2301	0125	7637	JMP BB4	
2302			/	
2303	0126	1020	DISP, LDA 1	
2304	0127	7400	-377	
2305	0130	0061	SET 1 1	
2306	0131	0000	0	
2307	0132	0062	SET 1 2	
2310	0133	0566	566	
2311	0134	1742	DSC 2	/DISPLAY 6
2312	0135	1762	DSC 1 2	/X AXIS
2313			/	
2314	0136	0440	SNS 0	/SELECT DATA
2315	0137	6265	JMP MK	/DISPLAY
2316			/	
2317	0140	1020	LDA 1	
2320	0141	7540	-237	
2321	0142	0061	SET 1 1	
2322	0143	0000	0	
2323	0144	0062	SET 1 2	
2324	0145	0572	572	
2325	0146	1742	DSC 2	/DISPLAY 20
2326	0147	1762	DSC 1 2	/Y AXIS
2327	0150	1762	DSC 1 2	
2330	0151	1762	DSC 1 2	
2331			**	
2332	0152	1020	LDA 1	
2333	0153	7400	-377	
2334	0154	0061	SET 1 1	
2335	0155	0040	40	
2336	0156	0062	SET 1 2	
2337	0157	0572	572	
2340	0160	1742	DSC 2	/DISPLAY 200
2341	0161	1762	DSC 1 2	/X AXIS
2342	0162	1762	DSC 1 2	
2343	0163	1762	DSC 1 2	
2344	0164	1762	DSC 1 2	
2345	0165	1762	DSC 1 2	
2346			/	
2347	0166	0101	SAN 1	/R TO L ISIM
2350	0167	0451	AF0	/DATA DISPLAY
2351	0170	0011	CLR	/SHIFT
2352	0171	1120	ADR 1	

2353	0172	2000		2000	
2354	0173	4175		STC .42	
2355	0174	0065		SET I 5	
2356	0175	0000		0	
2357	0176	6242		JMP SCLE	/SCALE PLOT
2360	0177	4201		STC .42	
2361	0200	0071		SET I 9	
2362	0201	0000		0	
2363			/		
2364	0202	0644	RE,	LDF 4	
2365	0203	1005		LDA 5	
2366	0204	0643		LDF 3	
2367	0205	1120		ADA I	
2370	0206	0005		5	
2371	0207	0341		SCR 1	/SCALE INT TIMES
2372	0210	1040		STA	
2373	0211	0220		SCL	
2374	0212	1120		ADA I	
2375	0213	6776		-1001	/BYPASS DISPLAY
2376	0214	0451		APD	/IF >101.4 MSEC
2377	0215	0456		SKP	
2400	0216	6224		JMP .46	
2401	0217	1020		LDA I	/DISPLAY
2402	0220	0000	SCL,	0	/INTERVALS
2403	0221	1120		ADA I	
2404	0222	7400		-377	
2405	0223	0151		DIS 9	
2406	0224	0644		LDF 4	
2407	0225	1025		LDA I 5	/EXIT ISIM
2410	0226	0643		LDF 3	
2411	0227	0450		AZE	/DISPLAY IF
2412	0230	0456		SKP	/INTERVAL = 0
2413	0231	6364		JMP DL	
2414	0232	6242		JMP SCLE	/SCALE PLOT
2415	0233	1140		ADM	
2416	0234	0011		9	/HORIZONTAL
2417	0235	1120		ADA I	/INCREMENT
2420	0236	7000		-777	
2421	0237	0451		APD	
2422	0240	6202		JMP RE	
2423	0241	6364		JMP DL	
2424			/		
2425	0242	1000	SCLE,	LDA	/SCALING ROUTINE
2426	0243	0000		0	/SAVE RETURN ADD
2427	0244	4264		STC BACK	
2430	0245	0644		LDF 4	
2431	0246	1005		LDA 5	
2432	0247	0643		LDF 3	
2433	0250	1120		ADA I	
2434	0251	0005		5	
2435	0252	0343		SCR 3	
2436			/		
2437	0253	0002		PDF	
2440				PHODE	
2441	6254	7421		NOL	
2442	6255	7200		CLA	
2443	6256	7407		DVI	/DIV 5
2444	6257	0005		5	
2445	6260	7200		CLA	
2446	6261	7501		NON	
2447	6262	6341		LINC	
2450				LHODE	
2451	0263	6264		JMP BACK	

2452			/		
2453	0264	0000	BACK,	0	
2454			/		
2455	0265	0071	HK,	SET I 11	/HISTO START ADD
2456	0266	3777		3777	
2457			/		
2460	0267	1020		LDA I	
2461	0270	7700		-77	
2462	0271	0061		SET I 1	
2463	0272	0000		0	
2464	0273	0062		SET I 2	
2465	0274	0572		572	
2466	0275	1742		DSC 2	/DISPLAY 200
2467	0276	1762		DSC I 2	/Y AXIS
2470	0277	1762		DSC I 2	
2471	0300	1762		DSC I 2	
2472	0301	1762		DSC I 2	
2473	0302	1762		DSC I 2	
2474			/		
2475	0303	1020		LDA I	
2476	0304	7400		-377	
2477	0305	0061		SET I 1	
2500	0306	0717		717	
2501	0307	0062		SET I 2	
2502	0310	0564		564	
2503	0311	1742		DSC 2	
2504	0312	1762		DSC I 2	
2505	0313	1762		DSC I 2	/DISPLAY 100
2506	0314	1762		DSC I 2	/X AXIS
2507	0315	1762		DSC I 2	
2510	0316	1762		DSC I 2	
2511			/		
2512	0317	0073	LB,	SET I 13	/DISPLAY HISTO
2513	0320	0000		0	/0.8 MSEC BINS
2514	0321	0101		SAM 1	/U TO D HISTO
2515	0322	0451		RPO	/DISPLAY SHIFT
2516	0323	0011		CLR	
2517	0324	0017		COM	
2520	0325	4346		STC XC	
2521	0326	0647	LK,	LDF 7	
2522	0327	1011		LDA 11	
2523	0330	0643		LDF 3	
2524	0331	0450		RZE	
2525	0332	0456		SKP	
2526	0333	6352		JHP LY	
2527	0334	1040		STA	
2530	0335	0344		DT	
2531	0336	1120		ADA I	
2532	0337	6003		-1774	
2533	0340	0451		RPO	
2534	0341	0456		SKP	
2535	0342	6352		JHP LY	
2536	0343	1020		LDA I	
2537	0344	0000	DT,	0	
2540	0345	1120		ADA I	
2541	0346	0000	XC,	0	
2542	0347	1120		ADA I	
2543	0350	7400		-377	
2544	0351	0153		DIS 13	
2545	0352	0231	LY,	XSK I 11	
2546	0353	0916		NOP	
2547	0354	1020		LDA I	
2550	0355	0004		4	

2551	0356	1140		ADM	
2552	0357	0013		13	
2553	0360	1120		ADA I	
2554	0361	7001		-776	
2555	0362	0451		RPO	
2556	0363	6326		JMP LK	
2557	0364	0602	DL,	LIF 2	
2560	0365	7756		JMP RETN	/EXIT
2561			/		
2562	0366	0063	AXIS,	SET I 3	/HORIZONTAL AXIS
2563	0367	0000		0	/DISPLAY
2564	0370	0070		SET I 10	
2565	0371	6777		-1000	
2566	0372	1020		LDA I	
2567	0373	7400		-377	
2570	0374	0073	AK,	SET I 13	
2571	0375	7715		-62	
2572	0376	0143	BR,	DIS 3	
2573	0377	1120		ADA I	
2574	0400	0001		1	
2575	0401	1460		SAE I	
2576	0402	7412		-365	
2577	0403	6376		JMP BR	
2600	0404	1020		LDA I	
2601	0405	7400		-377	
2602	0406	0163		DIS I 3	
2603	0407	0230		XSK I 10	
2604	0410	0456		SKP	
2605	0411	6415		JMP BM	/EXIT
2606	0412	0233		XSK I 13	
2607	0413	6406		JMP -5	
2610	0414	6374		JMP AK	
2611			/		
2612	0415	1020	BM,	LDA I	/VERTICAL AXIS
2613	0416	7400		-377	/DISPLAY
2614	0417	0063	RZ,	SET I 3	
2615	0420	7000		7000	
2616	0421	0070		SET I 10	
2617	0422	7715		-62	
2620	0423	0073		SET I 13	
2621	0424	7767		-10	
2622	0425	0143	RT,	DIS 3	
2623	0426	1120		ADA I	
2624	0427	0001		1	
2625	0430	1460		SAE I	
2626	0431	0400		400	
2627	0432	6435		JMP AM	
2630	0433	0602		LIF 2	
2631	0434	7531		JMP KJ	/EXIT
2632	0435	0230	AM,	XSK I 10	
2633	0436	6425		JMP RT	
2634	0437	0163		DIS I 3	
2635	0440	0233		XSK I 13	
2636	0441	6437		JMP -2	
2637	0442	6417		JMP RZ	
2640			/		
2641	0443	1020	LET,	LDA I	/TITLE ROUTINE
2642	0444	0200		200	/IN IF 1
2643	0445	0004		ESF	
2644	0446	0500	LISN,	10B	
2645	0447	6031		6031	/KSF
2646	0450	6467		JMP TITLE	
2647	0451	0500		10B	

2650	0452	6036	6036	/KRB
2651	0453	0500	IOB	
2652	0454	6046	6046	/TLS
2653	0455	1560	BCL I	
2654	0456	7700	7700	
2655	0457	0241	ROL 1	
2656	0460	1120	ADA I	
2657	0461	0600	600	
2660	0462	1075	STA I 15	
2661	0463	1460	SRE I	
2662	0464	0706	706	
2663	0465	6446	JMP LISN	
2664	0466	6515	JMP EXIT	
2665	0467	0074	TITLE, SET I 14	
2666	0470	1000	1000	
2667	0471	0061	SET I 1	
2670	0472	0000	0	
2671	0473	1034	INTITL, LDA I 14	/BUMP
2672	0474	1000	LDA	
2673	0475	0015	15	
2674	0476	0017	COM	
2675	0477	1100	ADA	
2676	0500	0014	14	
2677	0501	0451	APD	
2700	0502	6504	JMP .+2	
2701	0503	6523	JMP END	
2702	0504	1014	LDA 14	
2703	0505	4507	STC .+2	
2704	0506	0076	SET I 16	
2705	0507	0000	0	
2706	0510	1020	LDA I	
2707	0511	0370	370	
2710	0512	1756	DSC 16	
2711	0513	1776	DSC I 16	
2712	0514	6473	JMP INTITL	
2713	0515	0075	EXIT, SET I 15	
2714	0516	1000	1000	
2715	0517	0500	IOB	
2716	0520	6042	6042	/TCF
2717	0521	0602	LIF 2	
2720	0522	7756	JMP RETN	
2721	0523	0074	END, SET I 14	
2722	0524	1000	1000	
2723	0525	0500	IOB	
2724	0526	6042	6042	/TCF
2725	0527	0602	LIF 2	
2726	0530	7756	JMP RETN	
2727				
2730			*564	
2731	0564	2101	2101	/1
2732	0565	0177	0177	
2733	0566	4136	4136	/0
2734	0567	3641	3641	
2735	0570	4136	4136	/0
2736	0571	3641	3641	
2737	0572	4523	4523	/2
2740	0573	2151	2151	
2741	0574	4136	4136	/0
2742	0575	3641	3641	
2743	0576	4136	4136	/0
2744	0577	3641	3641	
2745				
2746			*602	/IF 3



2747	0602	4477	4477	
2750	0603	7744	7744	/A
2751	0604	5177	5177	
2752	0605	2651	2651	/B
2753	0606	4136	4136	
2754	0607	2241	2241	/C
2755	0610	4177	4177	
2756	0611	3641	3641	/D
2757	0612	4577	4577	
2760	0613	4145	4145	/E
2761	0614	4477	4477	
2762	0615	4044	4044	/F
2763	0616	4136	4136	
2764	0617	2645	2645	/G
2765	0620	1077	1077	
2766	0621	7710	7710	/H
2767	0622	7741	7741	
2770	0623	0041	0041	/I
2771	0624	4142	4142	
2772	0625	4076	4076	/J
2773	0626	1077	1077	
2774	0627	4324	4324	/K
2775	0630	0177	0177	
2776	0631	0301	0301	/L
2777	0632	3077	3077	
3000	0633	7730	7730	/M
3001	0634	3077	3077	
3002	0635	7706	7706	/N
3003	0636	4177	4177	
3004	0637	7741	7741	/O
3005	0640	4477	4477	
3006	0641	3044	3044	/P
3007	0642	4276	4276	
3010	0643	0376	0376	/Q
3011	0644	4477	4477	
3012	0645	3146	3146	/R
3013	0646	5121	5121	
3014	0647	4651	4651	/S
3015	0650	4040	4040	
3016	0651	4077	4077	/T
3017	0652	0177	0177	
3020	0653	7701	7701	/U
3021	0654	0176	0176	
3022	0655	7402	7402	/V
3023	0656	0677	0677	
3024	0657	7701	7701	/W
3025	0660	1463	1463	
3026	0661	6314	6314	/X
3027	0662	0770	0770	
3030	0663	7007	7007	/Y
3031	0664	4543	4543	
3032	0665	6151	6151	/Z
3033	0666	4177	4177	
3034	0667	0000	0000	/[
3035	0670	2040	2040	
3036	0671	0410	0410	/\
3037	0672	0000	0000	
3040	0673	7741	7741	/]
3041	0674	2000	2000	
3042	0675	2077	2077	/^
3043	0676	3410	3410	
3044	0677	1010	1010	/BACK ARROW
3045	0700	0000	0000	

3046	0701	0000	0000	/SPACE
3047	0702	7500	7500	
3050	0703	0000	6000	/!
3051	0704	6000	6000	
3052	0705	6000	6000	/"
3053	0706	3614	3614	
3054	0707	1436	1436	/NUMBER SIGN
3055	0710	7721	7721	
3056	0711	4677	4677	/\$
3057	0712	1446	1446	
3060	0713	6130	6130	/%
3061	0714	5166	5166	
3062	0715	0523	0523	/&
3063	0716	5000	5000	
3064	0717	0060	0060	/APOSTROPHY
3065	0720	4136	4136	
3066	0721	0000	0000	/(
3067	0722	0000	0000	
3070	0723	3641	3641	/)
3071	0724	2050	2050	
3072	0725	0050	0050	/*
3073	0726	0404	0404	
3074	0727	0437	0437	/+
3075	0730	0605	0605	
3076	0731	0000	0000	/,
3077	0732	0404	0404	
3100	0733	0404	0404	/-
3101	0734	0001	0001	
3102	0735	0000	0000	/.
3103	0736	0601	0601	
3104	0737	4030	4030	//
3105	0740	4136	4136	
3106	0741	3641	3641	/0
3107	0742	2101	2101	
3110	0743	0177	0177	/1
3111	0744	4523	4523	
3112	0745	2151	2151	/2
3113	0746	4122	4122	
3114	0747	2651	2651	/3
3115	0750	2414	2414	
3116	0751	0477	0477	/4
3117	0752	5172	5172	
3120	0753	0651	0651	/5
3121	0754	1506	1506	
3122	0755	4225	4225	/6
3123	0756	4443	4443	
3124	0757	6050	6050	/7
3125	0750	5126	5126	
3126	0761	2651	2651	/8
3127	0762	5120	5120	
3130	0763	3651	3651	/9
3131	0764	2200	2200	
3132	0765	0000	0000	/:
3133	0766	2601	2601	
3134	0767	0000	0000	/;
3135	0770	2410	2410	
3136	0771	0042	0042	/<
3137	0772	1212	1212	
3140	0773	1212	1212	/=
3141	0774	4200	4200	
3142	0775	1024	1024	/>
3143	0776	2000	2000	
3144	0777	3045	3045	/?

3145					
3146				PMODE	
3147				PLOT=JMS I 2	
3150				WRITE=JMS I 3	
3151				MARK=JMS I 4	
3152				NUM=JMS I 20	
3153				DUMP=JMS I 21	
3154				DOWN=CLA	
3155				UP=CLA IAC	
3156				INIT=CLA CMA	
3157				*2	/8 MODE PAGE 0
3160	0002	3400		PLOTX	
3161	0003	3313		TYPLOT	
3162	0004	3276		PLT1	
3163	0005	1020	XY,	XYPLT	
3154	0006	0000	CT,	0	
3165				*10	
3166	0010	0000	PXTABLE,	0	/AUTO INDEXES
3167	0011	0000	PYTABLE,	0	/10 AND 11
3170				*20	
3171	0020	3200		DECPLT	
3172	0021	2200		PLTSTG	
3173				*1000	
3174	1000	7777	KFM,	7777	/FMODM
3175	1001	5777	KHS,	5777	/HISTO
3176	1002	1006	PTEXT1,	TEXT1	
3177	1003	1010	PTEXT2,	TEXT2	
3200	1004	1012	PTEXT3,	TEXT3	
3201	1005	1015	PTEXT4,	TEXT4	
3202	1006	6000			
3202			TEXT1,	TEXT \$0\$	
3203	1007	0000		0	
3204	1010	6260			
3204			TEXT2,	TEXT \$20\$	
3205	1011	0000		0	
3206	1012	6160			
3206	1013	6000			
3206			TEXT3,	TEXT \$100\$	
3207	1014	0000		0	
3210	1015	6460			
3210	1016	6000			
3210			TEXT4,	TEXT \$400\$	
3211	1017	0000		0	
3212	1020	0000	XYPLT,	0	/START XY-PLOT
3213	1021	7300		CLA CLL	/ROUTINE
3214	1022	1200		TAD KFM	
3215	1023	3010		DCA PXTABLE	
3216	1024	1201		TAD KHS	/RESTORE TABLE
3217	1025	3011		DCA PYTABLE	/START ADDRESSES
3220	1026	7240		INIT	
3221	1027	4402		PLOT	
3222	1030	7201		UP	
3223	1031	4402		PLOT	
3224	1032	0074		74	/0.6"
3225	1033	0106		106	/0.7"
3226	1034	7200		CLA	
3227	1035	1202		TAD PTEXT1	/0
3230	1036	4403		WRITE	
3231	1037	0003		0003	
3232					
3233	1040	7201		UP	
3234	1041	4402		PLOT	
3235	1042	0055		55	/0.45"

3236	1043	0252	252	/1.7"
3237	1044	7200	CLA	
3240	1045	1203	TAD PTEXT2	/20
3241	1046	4403	WRITE	
3242	1047	0003	0003	
3243				
3244	1050	7201	UP	
3245	1051	4402	PLOT	
3246	1052	0074	74	/0.6"
3247	1053	1072	1072	/5.7"
3250	1054	7200	CLA	
3251	1055	1202	TAD PTEXT1	/0
3252	1056	4403	WRITE	
3253	1057	0003	0003	
3254				
3255	1060	7201	UP	
3256	1061	4402	PLOT	
3257	1062	0036	36	/0.3"
3260	1063	1236	1236	/6.7"
3261	1064	7200	CLA	
3262	1065	1204	TAD PTEXT3	/100
3263	1066	4403	WRITE	
3264	1067	0003	0003	
3265				
3266	1070	7201	UP	
3267	1071	4402	PLOT	
3270	1072	0330	330	/2.2"
3271	1073	0062	62	/0.5"
3272	1074	7200	CLA	
3273	1075	1205	TAD PTEXT4	/400
3274	1076	4403	WRITE	
3275	1077	0003	0003	
3276				
3277	1100	7201	UP	/1518 0.0
3300	1101	4402	PLOT	
3301	1102	0120	120	
3302	1103	0120	120	
3303	1104	7240	INIT	
3304	1105	4402	PLOT	
3305				
3306	1106	7200	DOWN	/Y-AXIS
3307	1107	4402	PLOT	
3310	1110	0000	0	
3311	1111	1750	1750	
3312				
3313	1112	7200	CLA	
3314	1113	1353	TAD KINC2	
3315	1114	3335	DCR INC2	
3316	1115	1351	TAD KCT	
3317	1116	3006	DCR CT	
3320				
3321	1117	7200	BACK1, CLA	
3322	1120	1355	TAD INC2	
3323	1121	3325	DCR INC1	
3324	1122	7200	DOWN	
3325	1123	4402	PLOT	
3326	1124	0024	24	
3327	1125	0000	INC1, 0	
3330	1126	7200	CLA	
3331	1127	1352	TAD M5	
3332	1130	1335	TAD INC2	
3333	1131	3335	DCR INC2	
3334	1132	7201	UP	

3335	1133	4402		PLOT	
3336	1134	0000		0	
3337	1135	0000	INC2,	0	
3340	1136	2006		ISZ CT	
3341	1137	5317		JMP BACK1	
3342			/		
3343	1140	7201		UP	/151M X-AXIS
3344	1141	4402		PLOT	
3345	1142	0000		0	
3346	1143	0000		0	
3347	1144	7200		DOWN	
3350	1145	4402		PLOT	
3351	1146	1356		1356	
3352	1147	0000		0	
3353	1150	5377		JMP PAGE1	
3354			/		
3355	1151	7754	KCT,	-24	
3356	1152	7716	M5,	-62	
3357	1153	1750	KINC2,	1750	
3360			/		
3361				*1177	
3362	1177	7200	PAGE1,	CLA	
3363	1200	1342		TAD KINC4	
3364	1201	3224		DCA INC4	
3365	1202	1340		TAD CTK	
3366	1203	3006		DCA CT	
3367			/		
3370	1204	7200	BACK2,	CLA	
3371	1205	1224		TAD INC4	
3372	1206	3211		DCA INC3	
3373	1207	7200		DOWN	
3374	1210	4402		PLOT	
3375	1211	0000	INC3,	0	
3376	1212	0024		24	
3377	1213	2006		ISZ CT	
3400	1214	7410		SKP	
3401	1215	5227		JMP AXS2	
3402	1216	7200		CLA	
3403	1217	1341		TAD M75	
3404	1220	1224		TAD INC4	
3405	1221	3224		DCA INC4	
3406	1222	7201		UP	
3407	1223	4402		PLOT	
3410	1224	0000	INC4,	0	
3411	1225	0000		0	
3412	1226	5204		JMP BACK2	
3413			/		
3414	1227	7201	AXS2,	UP	/HISTO X-AXIS
3415	1230	4402		PLOT	
3416	1231	0000		0	
3417	1232	0764		764	
3420	1233	7200		DOWN	
3421	1234	4402		PLOT	
3422	1235	1356		1356	
3423	1238	0764		764	
3424			/		
3425	1237	7201		UP	
3426	1240	4402		PLOT	
3427	1241	1306		1306	/7.1"
3430	1242	0726		726	/4.7"
3431	1243	7200		CLA	
3432	1244	1261		TAD PTEXT5	/100
3433	1245	4403		WRITE	

3434	1246	0003		0003
3435			/	
3436	1247	7201		UP
3437	1250	4402		PLOT
3440	1251	1356		1356
3441	1252	0764		764
3442			/	
3443	1253	7200		CLA
3444	1254	1342		TAD KINC4
3445	1255	3305		DCA INC6
3446	1256	1340		TAD CTK
3447	1257	3006		DCA CT
3450	1260	5265		JMP BACK3
3451			/	
3452	1261	1262	PTEXT5,	TEXT5
3453	1262	6160		
3453	1263	6000		
3453			TEXT5,	TEXT \$100\$
3454	1264	0000		0
3455			/	
3456	1265	7200	BACK3,	CLA
3457	1266	1305		TAD INC6
3460	1267	3272		DCA INC5
3461	1270	7200		DOWN
3462	1271	4402		PLOT
3463	1272	0000	INC5,	0
3464	1273	1010		1010
3465	1274	2006		ISZ CT
3466	1275	7410		SKP
3467	1276	5310		JMP HISP
3470	1277	7200		CLA
3471	1300	1341		TAD M75
3472	1301	1305		TAD INC6
3473	1302	3305		DCA INC6
3474	1303	7201		UP
3475	1304	4402		PLOT
3476	1305	0000	INC6,	0
3477	1306	0764		764
3500	1307	5265		JMP BACK3
3501			/	
3502	1310	7201	HISP,	UP
3503	1311	4402		PLOT
3504	1312	0000		0
3505	1313	0764		764
3506	1314	7300		CLA CLL
3507	1315	3333		DCA FL1
3510	1316	1344		TAD M175
3511	1317	3006		DCA CT
3512	1320	7200	BAK1,	CLA
3513	1321	6211		CPF 10
3514	1322	1411		TAD I PYTABLE
3515	1323	6201		CPF 0
3516	1324	1345		TAD K764
3517	1325	3334		DCA FL2
3520	1326	1343		TAD PG
3521	1327	1333		TAD FL1
3522	1330	3333		DCA FL1
3523	1331	7200		DOWN
3524	1332	4402		PLOT
3525	1333	0000	FL1,	0
3526	1334	0000	FL2,	0
3527	1335	2006		ISZ CT
3530	1336	5320		JMP BAK1

/PLOT HISTO

/125

/HISTO START ADD

3531	1337	5377		JMP PAGE2	
3532			/		
3533	1340	7766	CTK,	-12	
3534	1341	7665	M75,	-113	
3535	1342	1356	KINC4,	1356	
3536	1343	0006	PG,	6	
3537	1344	7603	M175,	-175	/-125
3540	1345	0764	K764,	764	/5"
3541			/		
3542				*1377	
3543	1377	7201	PAGE2,	UP	/PLOT ISIM
3544	1400	7300		CLA CLL	
3545	1401	3276		DCA FL3	
3546	1402	1316		TAD N02	
3547	1403	3314		DCA KTC	
3550	1404	1315		TAD N00	
3551	1405	3006		DCA CT	
3552	1406	7300	BAK2,	CLA CLL	
3553	1407	3274		DCA FL5	
3554	1410	6211		CDF 10	
3555	1411	1410		TAD I PXTABLE	
3556	1412	6201		CDF 0	
3557	1413	7450		SNA	/EXIT IF TRAIN
3560	1414	5301		JMP EX	/DONE
3561	1415	1312		TAD K5	
3562	1416	3277		DCA FL4	
3563			/		
3564	1417	2314		ISZ KTC	
3565	1420	5236		JMP .+16	
3566	1421	7300		CLA CLL	
3567	1422	1277		TAD FL4	
3570	1423	7421		NOL	
3571	1424	7200		CLA	
3572	1425	7407		DVI	/SCALE X AXIS
3573	1426	0032		32	
3574	1427	7200		CLA	
3575	1430	7501		NOR	
3576	1431	1276		TAD FL3	
3577	1432	3276		DCA FL3	
3600	1433	1316		TAD N02	
3601	1434	3314		DCA KTC	
3602	1435	5250		JMP .+13	
3603			/		
3604	1436	7300		CLA CLL	
3605	1437	1277		TAD FL4	
3606	1440	7421		NOL	
3607	1441	7200		CLA	
3610	1442	7407		DVI	
3611	1443	0033		33	
3612	1444	7200		CLA	
3613	1445	7501		NOR	
3614	1446	1276		TAD FL3	
3615	1447	3276		DCA FL3	
3616			/		
3617	1450	7300		CLA CLL	/SCALE Y AXIS
3620	1451	1277		TAD FL4	
3621	1452	1317		TAD N4	
3622	1453	7110		CLL RAR	
3623	1454	3277		DCA FL4	
3624			/		
3625	1455	7300		CLA CLL	
3626	1456	1276		TAD FL3	
3627	1457	1310		TAD M135G	

3630	1460	7500		SMA	/EXIT IF TRAIN
3631	1461	5301		JMP EX	/LENGTH >2 SEC
3632			/		
3633	1462	7300		CLA CLL	
3634	1463	2006		ISZ CT	
3635	1464	5270		JMP .+4	
3636	1465	7300		CLA CLL	
3637	1466	1311		TAD K1	/UP
3640	1467	3274		DCA FL5	
3641			/		
3642	1470	7300		CLA CLL	
3643	1471	1274		TAD FL5	
3644	1472	1313		TAD K7200	/DOWN
3645	1473	3274		DCA FL5	
3646	1474	0000	PL5,	0	/PEN POSITION
3647	1475	4402		PLOT	
3650	1476	0000	PL3,	0	
3651	1477	0000	PL4,	0	
3652	1500	5206		JMP BAK2	
3653			/		
3654	1501	7201	EX,	UP	/REPOSITION PEN
3655	1502	4402		PLOT	/FOR NEXT PLOT
3656	1503	1402		1402	
3657	1504	0000		0	
3660			/		
3661	1505	6141		LINC	
3662				LHODE	
3663	1506	0602		LIF 2	
3664	1507	7562		JMP KF	/EXIT
3665			/		
3666	1510	6421	M1356,	-1356	
3667	1511	0001	K1,	1	
3670	1512	0005	K5,	5	
3671	1513	7200	K7200,	7200	
3672	1514	0000	K10,	0	
3673	1515	7777	N00,	7777	
3674	1516	7775	N02,	7775	
3675	1517	7774	N4,	7774	
3676			/		
3677				PMODE	
3700				*2000	/PLOT SUBROUTINE
3701				PAGE	/IN IF 1
3702	2000	0000	MOVE,	0	
3703	2001	1276		TAD MNN5	
3704	2002	3301		DCA MNN500	
3705	2003	3271		DCA OLDRX	
3706	2004	1277	11H1,	TAD MNN2	
3707	2005	3300		DCA MNN200	
3710	2006	1672		TAD I L	
3711	2007	3273		DCA M	
3712	2010	1273	11H2,	TAD M	
3713	2011	0275		AND LLL7	
3714	2012	3265		DCA RABY	
3715	2013	1273		TAD M	
3716	2014	7012		RTR	
3717	2015	7010		RGR	
3720	2016	0275		AND LLL7	
3721	2017	1276		TAD MNN5	
3722	2020	7500		SMA	
3723	2021	4703		JMS I RPNUPD	
3724	2022	1274		TAD LLL5	
3725	2023	3264		DCA RABX	
3726	2024	1264		TAD RABX	



3727	2025	3271	DCA OLDRX
3730	2026	4313	JMS SCLPLT
3731	2027	4304	JMS PLOTIT
3732	2030	7200	CLA
3733	2031	1273	TAD M
3734	2032	7012	RTR
3735	2033	7012	RTR
3736	2034	7012	RTR
3737	2035	3273	DCA M
3740	2036	2300	ISZ MMM2G0
3741	2037	5210	JMP TIM2
3742	2040	2301	ISZ MMM5G0
3743	2041	7410	SKP
3744	2042	5245	JMP SPC
3745	2043	2272	ISZ L
3746	2044	5204	JMP TIM1
3747	2045	4703	SPC, JMS I RPNUPD
3750	2046	7200	CLA
3751	2047	1270	TAD LINRUN
3752	2050	7650	SNA CLA
3753	2051	5256	JMP .+5
3754	2052	1275	TAD LLL7
3755	2053	3265	DCA RAWY
3756	2054	3264	DCA RAWX
3757	2055	5261	JMP .+4
3760	2056	1274	TAD LLL5
3761	2057	3264	DCA RAWX
3762	2060	3265	DCA RAWY
3763	2061	4313	JMS SCLPLT
3764	2062	4304	JMS PLOTIT
3765	2063	5600	JMP I MOVE
3766			/
3767	2064	0000	RAWX, 0
3770	2065	0000	RAWY, 0
3771	2066	0000	SCLMLT, 0
3772	2067	0000	TRANSQ, 0
3773	2070	0000	LINRUN, 0
3774	2071	0000	OLDRX, 0
3775	2072	0000	L, 0
3776	2073	0000	M, 0
3777	2074	0005	LLL5, 5
4000	2075	0007	LLL7, 7
4001	2076	7773	MMM5, -5
4002	2077	7776	MMM2, -2
4003	2100	0000	MMM2G0, 0
4004	2101	0000	MMM5G0, 0
4005	2102	0000	CNTR, 0
4006	2103	2264	RPNUPD, PNPUDN
4007			/
4010	2104	0000	PLOTIT, 0
4011	2105	7201	PNSTAT, CLA IAC
4012	2106	4712	JMS I PLUT
4013	2107	0000	XHVT, 0
4014	2110	0000	YHVT, 0
4015	2111	5704	JMP I PLOTIT
4016			/
4017	2112	3400	PLUT, PLOTX
4020	2113	0000	SCLPLT, 0
4021	2114	1267	TAD TRANSQ
4022	2115	7450	SNA
4023	2116	5354	JMP STRAIT
4024	2117	1277	TAD MMM2
4025	2120	7450	SNA

4026	2121	5334		JMP ROTRIT
4027	2122	7510		SPA
4030	2123	5343		JMP INVRT
4031			/	
4032	2124	7200	ROTLFT,	CLA
4033	2125	1264		TAD RAWX
4034	2126	3310		DCA TRY
4035	2127	1265		TAD RAWY
4036	2130	7041		CIA
4037	2131	1275		TAD LLL7
4040	2132	3307		DCA TRX
4041	2133	5360		JMP SCALE
4042			/	
4043	2134	1264	ROTRIT,	TAD RAWX
4044	2135	7041		CIA
4045	2136	1274		TAD LLL5
4046	2137	3310		DCA TRY
4047	2140	1255		TAD RAWY
4050	2141	3307		DCA TRX
4051	2142	5360		JMP SCALE
4052			/	
4053	2143	1264	INVRT,	TAD RAWX
4054	2144	7041		CIA
4055	2145	1274		TAD LLL5
4056	2146	3307		DCA TRX
4057	2147	1265		TAD RAWY
4060	2150	7041		CIA
4061	2151	1275		TAD LLL7
4062	2152	3310		DCA TRY
4063	2153	5360		JMP SCALE
4064			/	
4065	2154	1264	STRAIT,	TAD RAWX
4066	2155	3307		DCA TRX
4067	2156	1265		TAD RAWY
4070	2157	3310		DCA TRY
4071			/	
4072	2160	1266	SCALE,	TAD SCLMLT
4073	2161	3302		DCA CNTR
4074	2162	1307		TAD TRX
4075	2163	2302		ISZ CNTR
4076	2164	5362		JMP -2
4077	2165	3307		DCA XMYT
4100	2166	1266		TAD SCLMLT
4101	2167	3302		DCA CNTR
4102	2170	1310		TAD TRY
4103	2171	2302		ISZ CNTR
4104	2172	5370		JMP -2
4105	2173	3310		DCA YMYT
4106	2174	5713		JMP I SCLPLT
4107				TRX=XMYT
4110				TRY=YMYT
4111			/	
4112				PAGE
4113	2200	0000	PLTSTG,	0
4114	2201	3317		DCA STRADD
4115	2202	1731		TAD I OLDX
4116	2203	3327		DCA COLDX
4117	2204	1732		TAD I OLDY
4120	2205	3330		DCA COLDY
4121	2206	1324	RDPACK,	TAD MM2
4122	2207	3320		DCA MM2GO
4123	2210	1717		TAD I STRADD
4124	2211	3321		DCA SAV

4125	2212	1321	RDPK2,	TAD SAV
4126	2213	7012		RTR
4127	2214	7012		RTR
4130	2215	7012		RTR
4131	2216	0325	RDPK3,	AND LL77
4132	2217	7450		SNA
4133	2220	5256		JMP TYPFIN
4134	2221	3322	BASAD,	DCA CODE
4135	2222	1326		TAD MM5
4136	2223	3323		DCA CNTR2
4137	2224	1322		TAD CODE
4140	2225	2323		ISZ CNTR2
4141	2226	5224		JMP .-2
4142	2227	1336		TAD BASE
4143	2230	3733		DCA I RL
4144	2231	4304	UPDATE,	JMS SCL00
4145	2232	3731		DCA I OLDX
4146	2233	1734		TAD I RYMYT
4147	2234	3732		DCA I OLDY
4150	2235	4735		JMS I RMOVE
4151	2236	4304		JMS SCL00
4152	2237	7041		CIA
4153	2240	1731		TAD I OLDX
4154	2241	1327		TAD COLDX
4155	2242	3327		DCA COLDX
4156	2243	1734		TAD I RYMYT
4157	2244	7041		CIA
4160	2245	1732		TAD I OLDY
4161	2246	1330		TAD COLDY
4162	2247	3330		DCA COLDY
4163	2250	1321	RDPK4,	TAD SAV
4164	2251	2320		ISZ MM2G0
4165	2252	5216		JMP RDPK3
4166	2253	7200		CLA
4167	2254	2317		ISZ STRADD
4170	2255	5206		JMP RDPACK
4171	2256	7200	TYPFIN,	CLA
4172	2257	1327		TAD COLDX
4173	2260	3731		DCA I OLDX
4174	2261	1330		TAD COLDY
4175	2262	3732		DCA I OLDY
4176	2263	5600		JMP I PLTSTG
4177			/	
4200	2264	0000	PNUPDN,	0
4201	2265	1324		TAD MM2
4202	2266	7640		SZA CLA
4203	2267	5273		JMP .+4
4204	2270	1300		TAD DN
4205	2271	3702		DCA I RPNSTA
4206	2272	5275		JMP .+3
4207	2273	1301		TAD PUP
4210	2274	3702		DCA I RPNSTA
4211	2275	1703		TAD I ROLDRX
4212	2276	1326		TAD MM5
4213	2277	5664		JMP I PNUPDN
4214	2300	7200	DN,	CLA
4215	2301	7201	PUP,	CLA IAC
4216	2302	2105	RPNSTA,	PNSTAT
4217	2303	2071	ROLDRX,	OLDRX
4220			/	
4221	2304	0000	SCL00,	0
4222	2305	7200		CLA
4223	2306	3713		DCA I RRAWX

4224	2307	3714	DCA I RRAWY
4225	2310	4715	JMS I RSCLPL
4226	2311	1716	TAD I RXMYT
4227	2312	5704	JMP I SCL00
4230	2313	2064	RRAWX, RAWX
4231	2314	2065	RRAWY, RAWY
4232	2315	2113	RSCLPL, SCLPLT
4233	2316	2107	RXMYT, XMYT

4234			/VARIABLES FOR PLTSTG
4235	2317	0000	STRADD, 0
4236	2320	0000	MM260, 0
4237	2321	0000	SAV, 0
4240	2322	0000	CODE, 0
4241	2323	0000	CNTR2, 0
4242	2324	7776	MM2, -2
4243	2325	0077	LL77, 77
4244	2326	7773	MMS, -5
4245	2327	0000	COLDX, 0
4246	2330	0000	COLDY, 0
4247	2331	3562	OLDX, PLOTX+162
4250	2332	3563	OLDY, PLOTX+163
4251	2333	2072	RL, L
4252	2334	2110	RMYT, YMYT
4253	2335	2000	RMOVE, MOVE
4254	2336	2332	BASE, -4

4255 /

4256 /MOVEMENT CODES FOR EACH CHARACTER

4257 /

4260	2337	1370	1370;
4260	2340	1333	1333;
4260	2341	4026	4026;
4260	2342	4040	4040;
4260	2343	4040	4040 /A
4261	2344	4676	4676;
4261	2345	3343	3343;
4261	2346	3303	3303;
4261	2347	4042	4042;
4261	2350	4000	4000 /B
4262	2351	4660	4660;
4262	2352	1676	1676;
4262	2353	0105	0105;
4262	2354	4010	4010;
4262	2355	4040	4040 /C
4263	2356	3676	3676;
4263	2357	4145	4145;
4263	2360	0030	0030;
4263	2361	0000	0000;
4263	2362	0000	0000 /D
4264	2363	4676	4676;
4264	2364	0306	0306;
4264	2365	0333	0333;
4264	2366	4000	4000;
4264	2367	4040	4040 /E
4265	2370	4676	4676;
4265	2371	0306	0306;
4265	2372	0333	0333;
4265	2373	0000	0000;
4265	2374	0000	0000 /F
4266	2375	4660	4660;
4266	2376	1676	1676;
4266	2377	0105	0105;
4266	2400	4010	4010;
4266	2401	2242	2242 /G

4267	2402	0376	0376;	
4267	2403	4643	4643;	
4267	2404	4040	4040;	
4267	2405	4040	4040;	
4267	2406	4040	4040	/H
4270	2407	7010	7010;	
4270	2410	2030	2030;	
4270	2411	1626	1626;	
4270	2412	3636	3636;	
4270	2413	3636	3636	/I
4271	2414	0072	0072;	
4271	2415	3630	3630;	
4271	2416	4626	4626;	
4271	2417	4646	4646;	
4271	2420	4646	4646	/J
4272	2421	0276	0276;	
4272	2422	4013	4013;	
4272	2423	4613	4613;	
4272	2424	4646	4646;	
4272	2425	4646	4646	/K
4273	2426	0076	0076;	
4273	2427	4040	4040;	
4273	2430	4040	4040;	
4273	2431	4040	4040;	
4273	2432	4040	4040	/L
4274	2433	2376	2376;	
4274	2434	4046	4046;	
4274	2435	4040	4040;	
4274	2436	4040	4040;	
4274	2437	4040	4040	/M
4275	2440	4076	4076;	
4275	2441	4646	4646;	
4275	2442	4646	4646;	
4275	2443	4646	4646;	
4275	2444	4646	4646	/N
4276	2445	7561	7561;	
4276	2446	3616	3616;	
4276	2447	4145	4145;	
4276	2450	1030	1030;	
4276	2451	0501	0501	/O
4277	2452	4676	4676;	
4277	2453	0343	0343;	
4277	2454	4643	4643;	
4277	2455	0006	0006;	
4277	2456	0000	0000	/P
4300	2457	1645	1645;	
4300	2460	4146	4146;	
4300	2461	0030	0030;	
4300	2462	2262	2262;	
4300	2463	4072	4072	/Q
4301	2464	4676	4676;	
4301	2465	0343	0343;	
4301	2466	4013	4013;	
4301	2467	4040	4040;	
4301	2470	4040	4040	/R
4302	2471	3070	3070;	
4302	2472	4241	4241;	
4302	2473	1333	1333;	
4302	2474	0504	0504;	
4302	2475	4616	4616	/S
4303	2476	7620	7620;	
4303	2477	0646	0646;	
4303	2500	0606	0606;	

4303	2501	0606	0606;	
4303	2502	0506	0606	/T
4304	2503	7601	7601;	
4304	2504	1001	1001;	
4304	2505	4130	4130;	
4304	2506	4646	4646;	
4304	2507	4646	4646	/U
4305	2510	7606	7606;	
4305	2511	4620	4620;	
4305	2512	4646	4646;	
4305	2513	4646	4646;	
4305	2514	4646	4646	/V
4306	2515	7606	7606;	
4306	2516	2310	2310;	
4306	2517	4630	4630;	
4306	2520	4646	4646;	
4306	2521	4646	4646	/W
4307	2522	4670	4670;	
4307	2523	0666	0666;	
4307	2524	4076	4076;	
4307	2525	4040	4040;	
4307	2526	4040	4040	/X
4310	2527	7320	7320;	
4310	2530	2306	2306;	
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4313	2552	2545	2545	/A
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4334	2674	6333	6333;
4334	2675	3333	3333;
4334	2676	3333	3333;

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/NUMBER SIGN

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4351	3000	1424	1424 /:
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4356	3030	2123	2123;
4356	3031	7060	7060 /?
4357			/

4360			PAGE
4361	3200	0000	DECPLT, 0
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4363	3202	5207	JMP PLUS
4364	3203	7041	CIA
4365	3204	3261	DCA VALUE
4366	3205	1272	TAD L55
4367	3206	5211	JMP .+3
4370	3207	3261	PLUS, DCA VALUE
4371	3210	1274	TAD L40
4372	3211	4276	JMS FLT1
4373	3212	3270	DCA SKPF
4374	3213	1253	TAD M7
4375	3214	3271	DCA CNT4
4376	3215	3262	DCA DIGIT
4377	3216	1253	TAD CNTR2A
4400	3217	3263	DCA CNTR2B
4401	3220	1252	TAD ADDR2A
4402	3221	3226	DCA ARROW
4403	3222	7410	SKP
4404	3223	3261	DCA VALUE
4405	3224	7100	CLL
4406	3225	1261	TAD VALUE
4407	3226	1254	ARROW, TAD TENFNR
4410	3227	7430	SZL
4411	3230	2262	ISZ DIGIT
4412	3231	7430	SZL
4413	3232	5223	JMP ARROW-3
4414	3233	7200	CLA
4415	3234	1262	TAD DIGIT
4416	3235	2271	ISZ CNT4
4417	3236	7410	SKP
4420	3237	5242	JMP .+3
4421	3240	7450	SNA

4422	3241	5264		JMP CK
4423	3242	2270		ISZ SKPF
4424	3243	1275		TAD K60
4425	3244	4276	OUT,	JMS PLT1
4426	3245	3262		DCA DIGIT
4427	3246	2226		ISZ ARROW
4430	3247	2263		ISZ CNTRZB
4431	3250	5225		JMP ARROW-1
4432	3251	5600		JMP I DECPLT
4433	3252	1254	ADDRZA,	TAD TENPWR
4434	3253	7774	CNTRZA,	-4
4435	3254	6030	TENPWR,	-1750
4436	3255	7634		-0144
4437	3256	7766		-0012
4440	3257	7777		-0001
4441	3260	7777		-0001
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4443	3262	0000	DIGIT,	0
4444	3263	0000	CNTRZB,	0
4445	3264	1270	CK,	TAD SKPF
4446	3265	7650		SNA CLA
4447	3266	1273		TAD M20
4450	3267	5243		JMP OUT-1
4451	3270	0000	SKPF,	0
4452	3271	0000	CNT4,	0
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4454	3272	0055	L55,	55
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4460	3276	0000	PLT1,	0
4461	3277	7006		RTL
4462	3300	7006		RTL
4463	3301	7006		RTL
4464	3302	0312		AND L7700
4465	3303	3310		DCA STG
4466	3304	1311		TAD STG+1
4467	3305	4743		JMS I RPLTST
4470	3306	7300		CLA CLL
4471	3307	5676		JMP I PLT1
4472	3310	0000	STG,	0
4473	3311	3310		STG
4474	3312	7700	L7700,	7700
4475	3313	0000	TYPLOT,	0
4476	3314	3346		DCA 5
4477	3315	1713		TAD I TYPLOT
4500	3316	0344		AND L77
4501	3317	7041		CIA
4502	3320	3740		DCA I RSCLML
4503	3321	1713		TAD I TYPLOT
4504	3322	7012		RTR
4505	3323	7012		RTR
4506	3324	7012		RTR
4507	3325	0345		AND L7
4510	3326	3741		DCA I RTRANS
4511	3327	1713		TAD I TYPLOT
4512	3330	7006		RTL
4513	3331	7006		RTL
4514	3332	0345		AND L7
4515	3333	3742		DCA I RLINRU
4516	3334	2313		ISZ TYPLOT
4517	3335	1346		TAD 5
4520	3336	4743		JMS I RPLTST

4521	3337	5713		JMP I TYFLOT
4522			/	
4523	3340	2066	RSCLML,	SCLMLT
4524	3341	2067	RTRANS,	TRANSQ
4525	3342	2070	RLINKU,	LINRUN
4526	3343	2200	RPLTST,	PLTSTG
4527	3344	0077	L77,	77
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4541				PLPL=6521
4542				PLUD=6522
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4545	3400	0000	PLOTX,	0
4546	3401	7510		SPA
4547	3402	5220		JMP PLOTA
4550	3403	1361		TAD PLOTX
4551	3404	7112		CLL RTR
4552	3405	7710		SPA CLA
4553	3406	5227		JMP PLOT1
4554	3407	7620		SNL CLA
4555	3410	5214		JMP .+4
4556	3411	3361		DCA PLOTX
4557	3412	6504		PLPU
4560	3413	5216		JMP .+3
4561	3414	2361		ISZ PLOTX
4562	3415	6524		PLPD
4563	3416	4370		JMS PLOTX
4564	3417	5227		JMP PLOT1
4565	3420	7200	PLOTA,	CLA
4566	3421	6504		PLPU
4567	3422	3361		DCA PLOTX
4570	3423	3362		DCA PLOTX
4571	3424	3363		DCA PLOTX
4572	3425	4370		JMS PLOTX
4573	3426	5600		JMP I PLOTX
4574	3427	1362	PLOT1,	TAD PLOTX
4575	3430	7141		CIA CLL
4576	3431	1600		TAD I PLOTX
4577	3432	7420		SNL
4600	3433	7041		CIA
4601	3434	3364		DCA PLOTX
4602	3435	7004		RAL
4603	3436	3367		DCA PLOTX
4604	3437	1600		TAD I PLOTX
4605	3440	3362		DCA PLOTX
4606	3441	2200		ISZ PLOTX
4607	3442	1363		TAD PLOTX
4610	3443	7141		CIA CLL
4611	3444	1600		TAD I PLOTX
4612	3445	7420		SNL
4613	3446	7041		CIA
4614	3447	3365		DCA PLOTX
4615	3450	1367		TAD PLOTX
4616	3451	7004		RAL
4617	3452	3367		DCA PLOTX

4620	3453	1600	TAD I PLOTX
4621	3454	3363	DCA PLOTNY
4622	3455	2200	ISZ PLOTX
4623	3456	1364	TAD PLOTDX
4624	3457	7141	CIA CLL
4625	3460	1365	TAD PLOTDY
4626	3461	7620	SNL CLA
4627	3462	5275	JMP PLOT2
4630	3463	1364	TAD PLOTDX
4631	3464	3366	DCA PLOTNA
4632	3465	1365	TAD PLOTDY
4633	3466	3364	DCA PLOTDX
4634	3467	1366	TAD PLOTNA
4635	3470	3365	DCA PLOTDY
4636	3471	7001	JAC
4637	3472	0367	AND PLOTNY
4640	3473	1342	TAD PLOT11
4641	3474	5300	JMP .14
4642	3475	1367	TAD PLOTNY
4643	3476	7110	CLL RAR
4644	3477	1345	TAD PLOT12
4645	3500	3366	DCA PLOTNA
4646	3501	1766	TAD I PLOTNA
4647	3502	3340	DCA PLOT4
4650	3503	1365	TAD PLOTDY
4651	3504	7640	SZA CLA
4652	3505	5310	JMP .13
4653	3506	1766	TAD I PLOTNA
4654	3507	5314	JMP .15
4655	3510	1367	TAD PLOTNY
4656	3511	1350	TAD PLOT13
4657	3512	3367	DCA PLOTNY
4660	3513	1767	TAD I PLOTNY
4661	3514	3334	DCA PLOTDB
4662	3515	1365	TAD PLOTDY
4663	3516	7041	CIA
4664	3517	3365	DCA PLOTDY
4665	3520	1364	TAD PLOTDX
4666	3521	7040	CNA
4667	3522	3367	DCA PLOTNY
4670	3523	1364	TAD PLOTDX
4671	3524	7110	CLL RAR
4672	3525	2367	ISZ PLOTNY
4673	3526	7410	SKP
4674	3527	5600	JMP I PLOTX
4675	3530	7100	CLL
4676	3531	1365	TAD PLOTDY
4677	3532	7430	SZL
4700	3533	5340	JMP PLOT4
4701	3534	0000	FLOTDB, 0
4702	3535	1364	TAD PLOTDX
4703	3536	4370	JNS PLOTHT
4704	3537	5325	JMP PLOT3
4705	3540	0000	FLOT4, 0
4706	3541	5336	JMP .-3
4707	3542	3543	FLOT11, .11
4710	3543	6511	FLPR
4711	3544	6521	FLPL
4712	3545	3546	FLOT12, .11
4713	3546	6512	FLDU
4714	3547	6514	FLDD
4715	3550	3551	FLOT13, .11
4716	3551	6513	FLDU!FLPR

4717	3552	6523	FLUD!PLPL
4720	3553	6515	FLDD!PLPR
4721	3554	4355	JMS +1
4722	3555	0000	0
4723	3556	6514	FLDD
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4725	3560	5755	JMP 1 --3
4726	3561	0000	PLOTIN, 0
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4730	3563	0000	PLOTIN, 0
4731	3564	0000	PLOTIX, 0
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4736	3571	6501	PLSF
4737	3572	5371	JMP --1
4740	3573	6502	PLCF
4741	3574	5770	JMP 1 PLOTIN
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4747			PMODE
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4753	0201	7300	CLA CLL
4754	0202	4162	JMS CRLF
4755	0203	4162	JMS CRLF
4756	0204	1062	TAD K322 /R
4757	0205	4154	JMS TYPE
4760	0206	1066	TAD K240 /SPACE
4761	0207	4154	JMS TYPE
4762	0210	1062	TAD K322 /R
4763	0211	4154	JMS TYPE
4764	0212	1050	TAD K301 /R
4765	0213	4154	JMS TYPE
4766	0214	1064	TAD K324 /T
4767	0215	4154	JMS TYPE
4770	0216	1052	TAD K305 /E
4771	0217	4154	JMS TYPE
4772	0220	1066	TAD K240 /SPACE
4773	0221	4154	JMS TYPE
4774	0222	1066	TAD K240 /SPACE
4775	0223	4154	JMS TYPE
4776	0224	1057	TAD K315 /M
4777	0225	4154	JMS TYPE
5000	0226	1066	TAD K240 /SPACE
5001	0227	4154	JMS TYPE
5002	0230	1426	TAD I PR1 /MEAN RESP RATE
5003	0231	4114	JMS TYP
5004	0232	1054	TAD K310 /H
5005	0233	4154	JMS TYPE
5006	0234	1066	TAD K240 /SPACE
5007	0235	4154	JMS TYPE
5010	0236	1427	TAD I PR2 /HIGH
5011	0237	4114	JMS TYP
5012	0240	1056	TAD K314 /L
5013	0241	4154	JMS TYPE
5014	0242	1066	TAD K240 /SPACE
5015	0243	4154	JMS TYPE

5016	0244	1430	TAD I PR3	/LOW
5017	0245	4114	JMS TYP	
5020	0246	1060	TAD K316	/N
5021	0247	4154	JMS TYPE	
5022	0250	1066	TAD K240	/SPACE
5023	0251	4154	JMS TYPE	
5024	0252	1431	TAD I PR4	/NUMBER TRAINS
5025	0253	4114	JMS TYP	
5026	0254	6211	CDF 10	
5027	0255	1045	TAD K5000	/INDIVID RATES
5030	0256	4423	JMS I T01	
5031	0257	6201	CDF 0	
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5033	0261	4154	JMS TYPE	
5034	0262	1062	TAD K322	/R
5035	0263	4154	JMS TYPE	
5036	0264	1050	TAD K301	/A
5037	0265	4154	JMS TYPE	
5040	0266	1055	TAD K311	/I
5041	0267	4154	JMS TYPE	
5042	0270	1060	TAD K316	/N
5043	0271	4154	JMS TYPE	
5044	0272	1063	TAD K323	/S
5045	0273	4154	JMS TYPE	
5046	0274	1066	TAD K240	/SPACE
5047	0275	4154	JMS TYPE	
5050	0276	1066	TAD K240	/SPACE
5051	0277	4154	JMS TYPE	
5052	0300	1055	TAD K311	/I
5053	0301	4154	JMS TYPE	
5054	0302	1060	TAD K316	/N
5055	0303	4154	JMS TYPE	
5056	0304	1064	TAD K324	/T
5057	0305	4154	JMS TYPE	
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5061	0307	4154	JMS TYPE	
5062	0310	1066	TAD K240	/SPACE
5063	0311	4154	JMS TYPE	
5064	0312	1063	TAD K323	/S
5065	0313	4154	JMS TYPE	
5066	0314	1064	TAD K324	/T
5067	0315	4154	JMS TYPE	
5070	0316	1066	TAD K240	/SPACE
5071	0317	4154	JMS TYPE	
5072	0320	1432	TAD I PR5	/ST
5073	0321	4114	JMS TYP	
5074	0322	1060	TAD K316	/N
5075	0323	4154	JMS TYPE	
5076	0324	1066	TAD K240	/SPACE
5077	0325	4154	JMS TYPE	
5100	0326	6211	CDF 10	
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5102	0330	4114	JMS TYP	
5103	0331	1044	TAD K3777	/TRAIN LENGTH
5104	0332	4425	JMS I T03	
5105	0333	6201	CDF 0	
5106	0334	1053	TAD K306	/F
5107	0335	4154	JMS TYPE	
5110	0336	1055	TAD K311	/I
5111	0337	4154	JMS TYPE	
5112	0340	1062	TAD K322	/R
5113	0341	4154	JMS TYPE	
5114	0342	1063	TAD K323	/S

5115	0343	4154	JMS TYPE	
5116	0344	1064	TAD K324	/T
5117	0345	4154	JMS TYPE	
5120	0346	1066	TAD K240	/SPACE
5121	0347	4154	JMS TYPE	
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5123	0351	4154	JMS TYPE	
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5127	0355	4154	JMS TYPE	
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5132	0360	4423	JMS I T01	
5133	0361	6201	CDF 0	
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5145	0373	4154	JMS TYPE	
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5151	0377	4154	JMS TYPE	
5152	0400	1065	TAD K326	/V
5153	0401	4154	JMS TYPE	
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5155	0403	4154	JMS TYPE	
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5175	0423	4154	JMS TYPE	
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5177	0425	4114	JMS TYP	
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5231	0457	4154	JMS TYPE	
5232	0460	1066	TAD K240	/SPACE
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5241	0467	4154	JMS TYPE	
5242	0470	1051	TAD K304	/D
5243	0471	4154	JMS TYPE	
5244	0472	1066	TAD K240	/SPACE
5245	0473	4154	JMS TYPE	
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5247	0475	4114	JMS TYP	
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5255	0503	4154	JMS TYPE	
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5262	0510	1057	TAD K315	/M
5263	0511	4154	JMS TYPE	
5264	0512	1052	TAD K305	/E
5265	0513	4154	JMS TYPE	
5266	0514	1051	TAD K304	/D
5267	0515	4154	JMS TYPE	
5270	0516	1066	TAD K240	/SPACE
5271	0517	4154	JMS TYPE	
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5273	0521	4114	JMS TYP	
5274	0522	1060	TAD K316	/N
5275	0523	4154	JMS TYPE	
5276	0524	1066	TAD K240	/SPACE
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5301	0527	4114	JMS TYP	
5302	0530	6211	CDF 10	
5303	0531	1046	TAD K5777	/HISTO TABLE
5304	0532	4424	JMS I T02	
5305	0533	6201	CDF 0	
5306	0534	4162	JMS CRLF	/SPACE PAPER
5307	0535	4162	JMS CRLF	
5310	0536	4162	JMS CRLF	
5311	0537	4162	JMS CRLF	
5312	0540	4162	JMS CRLF	



5313	0541	4162	JMS CRLF	
5314	0542	4162	JMS CRLF	
5315	0543	4162	JMS CRLF	
5316	0544	4162	JMS CRLF	
5317	0545	4162	JMS CRLF	
5320	0546	4162	JMS CRLF	
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5324	0600	0000	OUT1, 0	/TELETYPE OUT
5325	0601	3012	DCA 12	
5326	0602	6046	TLS	/ROUTINE
5327	0603	7300	CLA CLL	
5330	0604	4162	JMS CRLF	
5331	0605	4162	JMS CRLF	
5332	0606	7200	CLA	
5333	0607	1072	TAD M12	
5334	0610	3073	DCA DON	
5335	0611	7200	AX, CLA	
5336	0612	1412	TAD I 12	
5337	0613	7440	SZA	
5340	0614	5220	JMP +4	
5341	0615	4162	JMS CRLF	
5342	0616	4162	JMS CRLF	
5343	0617	5600	JMP I OUT1	
5344	0620	4114	JMS TYP	
5345	0621	2073	ISZ DON	
5346	0622	5227	JMP DB	
5347	0623	4162	JMS CRLF	
5350	0624	7200	CLA	
5351	0625	1072	TAD M12	
5352	0626	3073	DCA DON	
5353	0627	4101	DB, JMS KEY	
5354	0630	7440	SZA	
5355	0631	5211	JMP AX	
5356	0632	1070	TAD K260	
5357	0633	3074	DCA CP	
5360	0634	4162	JMS CRLF	
5361	0635	4162	JMS CRLF	
5362	0636	5600	JMP I OUT1	
5363				
5364	0637	0000	OUT2, 0	
5365	0640	3012	DCA 12	
5366	0641	6046	TLS	
5367	0642	7300	CLA CLL	
5370	0643	4162	JMS CRLF	
5371	0644	4162	JMS CRLF	
5372	0645	7200	CLA	
5373	0646	1072	TAD M12	
5374	0647	3073	DCA DON	
5375	0650	1076	TAD N175	
5376	0651	3077	DCA CNT	
5377	0652	7200	AX2, CLA	
5400	0653	1412	TAD I 12	
5401	0654	4114	JMS TYP	
5402	0655	2077	ISZ CNT	
5403	0656	5260	JMP +2	
5404	0657	5637	JMP I OUT2	
5405	0660	2073	ISZ DON	
5406	0661	5266	JMP DB2	
5407	0662	4162	JMS CRLF	
5410	0663	7200	CLA	
5411	0664	1072	TAD M12	

5412	0665	3073		DCA DON
5413	0666	4101	DB2,	JMS KEY
5414	0667	7440		SZA
5415	0670	5252		JMP AX2
5416	0671	1070		TAD K260
5417	0672	3074		DCA CP
5420	0673	5637		JMP I OUT2
5421			/	
5422	0674	0000	OUT3,	0
5423	0675	3012		DCA 12
5424	0676	6046		TLS
5425	0677	7300		CLA CLL
5426	0790	4162		JMS CRLF
5427	0701	4162		JMS CRLF
5430	0702	7200		CLA
5431	0703	1071		TAD N2
5432	0704	3073		DCA DON
5433	0705	7200	AX3,	CLA
5434	0706	1412		TAD I 12
5435	0707	7440		SZA
5436	0710	5313		JMP .+3
5437	0711	4162		JMS CRLF
5440	0712	5674		JMP I OUT3
5441	0713	4114		JMS TYP
5442	0714	1066		TAD K240
5443	0715	4154		JMS TYPE
5444	0716	1066		TAD K240
5445	0717	4154		JMS TYPE
5446	0720	2073		ISZ DON
5447	0721	5305		JMP AX3
5450	0722	4162		JMS CRLF
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5452	0724	1071		TAD N2
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5454	0726	4101		JMS KEY
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5464				*22
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5466	0023	0600	T01,	OUT1
5467	0024	0637	T02,	OUT2
5470	0025	0674	T03,	OUT3
5471	0026	5760	PR1,	R1
5472	0027	5761	PR2,	R2
5473	0030	5762	PR3,	R3
5474	0031	5763	PR4,	R4
5475	0032	5764	PR5,	R5
5476	0033	5765	PR6,	R6
5477	0034	5766	PH1,	M1
5500	0035	5767	PH2,	M2
5501	0036	5770	PH3,	M3
5502	0037	5771	PH4,	M4
5503	0040	5772	PT1,	T1
5504	0041	5773	PT2,	T2
5505	0042	0777	K777,	777
5506	0043	2000	K2000,	2000
5507	0044	3777	K3777,	3777
5510	0045	5000	K5000,	5000

/SPACE

/SPACE

/8 MODE PAGE 0

5511	0046	5777	K5777,	5777	
5512	0047	7777	K7777,	7777	
5513	0050	0301	K301,	301	/A
5514	0051	0304	K304,	304	/D
5515	0052	0305	K305,	305	/E
5516	0053	0306	K306,	306	/F
5517	0054	0310	K310,	310	/H
5520	0055	0311	K311,	311	/I
5521	0056	0314	K314,	314	/L
5522	0057	0315	K315,	315	/M
5523	0060	0316	K316,	316	/N
5524	0061	0317	K317,	317	/O
5525	0062	0322	K322,	322	/R
5526	0063	0323	K323,	323	/S
5527	0064	0324	K324,	324	/T
5530	0065	0326	K326,	326	/V
5531	0066	0240	K240,	240	/SPACE
5532	0067	0257	K257,	257	//
5533	0070	0260	K260,	260	/0
5534	0071	7776	N2,	-2	
5535	0072	7766	M12,	-12	
5536	0073	0000	00N,	0	
5537	0074	0000	CP,	0	
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5541	0076	7603	N175,	-175	
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5544			/		
5545	0101	0000	KEY,	0	/CHECK FOR HIT
5546	0102	6031		KSF	/S (STOP) KEY
5547	0103	5106		JMP +3	
5550	0104	6036		KRB	
5551	0105	3074		DCA CP	
5552	0106	6032		KCC	
5553	0107	6046		TLS	
5554	0110	7200		CLA	
5555	0111	1074		TAD CP	
5556	0112	1075		TAD M323	
5557	0113	5501		JMP I KEY	
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5561	0114	0000	TYP,	0	/OCTAL TO
5562	0115	7421		MQL	/DECIMAL
5563	0116	7407		DVI	/CONVERSION
5564	0117	1750		1750	/ROUTINE
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5566	0121	7501		MQA	
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5575	0130	3100		DCA REM	
5576	0131	7501		MQA	
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5615	0150	4154		JMS TYPE	
5616	0151	1066		TAD K240	
5617	0152	4154		JMS TYPE	
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5623	0155	6041		TSF	
5624	0156	5155		JMP -1	
5625	0157	6046		TL5	
5626	0160	7200		CLA	
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5630					
5631	0162	0000	/	CRLF, 0	/CARRIAGE RETURN
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5635	0166	4154		JMS TYPE	
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5640	0171	0212	LF,	212	

## NO ERRORS

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AB	5036
ADDRZA	3252
AK	-6374
AM	-6435
ARROW	3226
AT	5134
AX	0611
AXIS	6366
AX52	1227
AX2	0652
AX3	0705
B	4334
BACK	6264
BACK1	1117
BACK2	1204
BACK3	1265
BAK1	1320
BAK2	1406
BASAD	2221
BASE	2336
BB1	5542
BB2	5556
BB3	5571
BB4	5637
BB5	5521
BLK	5750
BM	6415
BR	6376
C	4251
CK	3264
CNT	0077
CNTR	2102
CNTRZA	3253
CNTRZB	3263

CNTR2	2323
CNT4	3271
CO	4042
CODE	2322
COLDX	2327
COLDY	2330
CONT	5752
CP	0074
CR	0170
CRLF	0162
CT	0006
CTK	1340
D	4044
DB	0627
DB2	0666
DECPLT	3200
DIGIT	3262
DISF	6115
DISP	6126
DISR	6040
DISS	6016
DIST	6027
DIS1	6051
DIS2	6062
DIS3	6073
DISS	6104
DIV	5000
DL	6364
DN	2300
DN	0073
DDWN	7200
DS	4056
DSPY	5734
DT	6344
DUMP	4421
DV	4707
DVI	7407
E	4342
END	6523
EX	1501
EXIT	6515
FI	5743
GAR	5747
GD	4040
HISP	1310
HN	4512
INC1	1125
INC2	1135
INC3	1211
INC4	1224
INC5	1272
INC6	1305
INIT	7240
INTITL	6473
INVRT	2143
KA	5377
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KCLEN	4037
KCLLR	4035
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KEY	0101

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KO	5175
KP	5333
KQ	5142
KR	5525
KS	5363
KTC	1514
KU	5724
KW	5547
KX	5020
KX2	5022
K1	1511
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K240	0066
K257	0067
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K323	0063
K324	0064
K326	0065
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K5000	0045
K5777	0046
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K7200	1513
K764	1345
K777	0042
K7777	0047
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LF	0171
LINRUN	2070
LISN	6446
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LLL5	2074
LLL7	2075
LL77	2325
LY	6352
L40	3274
L510	5757
L55	3272
L7	3345
L77	3344

L7700	3312
M	2073
MARK	4404
MD	5745
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MM260	2180
MM5	2076
MM560	2101
MN2	2324
MN260	2320
MN5	2326
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MN2	5240
MN3	5244
MN4	5313
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N00	1515
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N4	1517
OLDEX	2071
OLDX	2331
OLDY	2332
OUT	3244
OUT0	0177
OUT1	0600
OUT2	0637
OUT3	0674
PAGE1	1177
PAGE2	1377
PLCF	6502
PLDD	6514
PLDU	6512
PLOT	4402
PLOT0	3420
PLOT0B	3534
PLOT0X	3564

PL01DY 3565  
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PL01T2 3545  
PL01T3 3550  
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PL01X 3400  
PL01 3427  
PL012 3475  
PL013 3525  
PL014 3540  
PLFD 6524  
PLFL 6521  
PLFR 6511  
PLFU 6504  
PLSF 6501  
PLTSTG 2200  
PLT1 3276  
PLUD 6522  
PLUS 3207  
PLUT 2112  
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PL2 1334  
PL3 1476  
PL4 1477  
PL5 1474  
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PM2 0035  
PM3 0036  
PM4 0037  
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PR5 0032  
PR6 0033  
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RANY 2065  
RC 5744  
RDPACK 2206  
RDPK2 2212



RDPK3	2216
RDPK4	2258
RE	6202
REN	0100
RETN	5756
RK	4610
RL	2333
RLINRU	3342
RHOVE	2335
RN	4662
ROLDX	2303
ROILET	2124
ROTRIT	2134
RP	4634
RPLTST	3343
RPNSTA	2302
RPNUPD	2103
RR	4572
RRANK	2313
RRANY	2314
RSCLHL	3340
RSCLPL	2315
RT	6425
RTRANS	3341
RU	4671
RXMVT	2316
RYMVT	2334
RZ	6417
R1	5760
R2	5761
R3	5762
R4	5763
R5	5764
R6	5765
S	3346
SAV	2321
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SFC	2045
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STOR	5751
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ST2	5062
SV	5641
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S2	5742
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TENWR	3254
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TEXT3	1012
TEXT4	1015
TEXT5	1262
TIM1	2084

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T02	0024
T03	0025
TR	4734
TRANSQ	2067
TRG	5746
TRIG	4552
TRX	2107
TRY	2110
TYP	0114
TYPE	0154
TYFFIN	2256
TYPLOT	3313
T1	5772
T2	5773
UF	7201
UPDATE	2231
V	4371
VALUE	3261
WRITE	4403
W10	5622
W0	5652
W1	5663
W2	5674
W3	5707
W4	5722
XC	6346
XHVT	2107
XY	0005
XYPLT	1020
YHVT	2110
ZRD	5755

\*L L, \$CALC1, 0

\*N

C FOCAL-12

01. 10 L 0, F1, U, #0, 1

01. 20 R !!, "FDP", PF, "LDP", PL

01. 30 F DP=PF, PL; D 2

01. 40 T !!; L C, F1; Q

02. 10 S  $J = \langle DP * 5 + 4 \rangle * 256 + 221$ ; S  $NT = F1 \langle J + 4 \rangle$ ; S  $ST = F1 \langle J + 5 \rangle / 10$

02. 20 S  $MI = F1 \langle J + 11 \rangle$ ; S  $L = MI + \langle F1 \langle J + 8 \rangle - MI \rangle / 10$ ; S  $D = ST - FITR \langle ST \rangle$

02. 30 D 3

02. 40 S  $J = DP * 5 * 256$ ; S  $SL = \langle F1 \langle J + 1 \rangle - F1 \langle J \rangle \rangle / F1 \langle J + 1 \rangle$

02. 50 D 4

02. 60 D 5

03. 10 S  $K = 0$ ; S  $C1 = 0$ ; S  $C2 = 0$

03. 20 S  $J = \langle DP * 5 + 3 \rangle * 256$ ; S  $C3 = F1 \langle J + K \rangle$

03. 30 S  $C1 = C1 + C3 * F1 \langle J + K + 1 \rangle$ ; S  $C2 = C2 + C3 * F1 \langle J + K + 1 \rangle ^ 2$

03. 40 S  $K = K + 2$ ; S  $C3 = F1 \langle J + K \rangle$ ; I  $\langle F1 \langle J + K \rangle \rangle 3. 5, 3. 5, 3. 3$

03. 50 S  $DF = FSQT \langle \langle C2 - C1 ^ 2 \rangle / NT \rangle / \langle NT - 1 \rangle$

03. 55 S  $DS = DF$

03. 60 D 3. 1; S  $J = \langle DP * 5 + 2 \rangle * 256$ ; S  $K = -1$ ; S  $C3 = 1$

03. 70 D 3. 3

03. 80 S  $K = K + 1$ ; I  $\langle F1 \langle J + K \rangle \rangle 3. 9, 3. 9, 3. 7$

03. 90 D 3. 5

04. 10 D 3. 1; S  $C3 = 0$

04. 20 I  $\langle K - ST + 2 \rangle 4. 3, 4. 3, 4. 7$

04. 30 S  $C1 = C1 + F1 \langle J + K \rangle$

04. 40 I  $\langle F1 \langle J + K \rangle - L \rangle 4. 5, 4. 6, 4. 6$

04. 50 S  $C2 = C2 + F1 \langle J + K \rangle$

04. 60 S  $K = K + 1$ ; G 4. 2

04. 70 S  $C = F1 \langle J + K \rangle * D$ ; S  $C1 = C1 + C$

04. 80 S  $LI = D * \langle F1 \langle J + K \rangle - F1 \langle J + K - 1 \rangle \rangle + F1 \langle J + K - 1 \rangle$

05. 10 S  $J = \langle DP * 5 + 4 \rangle * 256 + 221$

05. 20 S  $F1 \langle J + 1 \rangle = C1 / 10 + . 5$ ; S  $F1 \langle J + 2 \rangle = 60000 / F1 \langle J + 3 \rangle + . 5$

05. 30 S  $F1 \langle J + 6 \rangle = 05 * 10 + . 5$ ; S  $F1 \langle J + 7 \rangle = 100 * DS / FSQT \langle NT \rangle + . 5$

05. 40 S  $F1 \langle J + 9 \rangle = DF + . 5$ ; S  $F1 \langle J + 10 \rangle = 10 * DF / FSQT \langle NT \rangle + . 5$

05. 50 S  $F1 \langle J + 18 \rangle = C2 / 10 + . 5$ ; S  $F1 \langle J + 19 \rangle = LI + . 5$ ; S  $F1 \langle J + 20 \rangle = -SL + 100 + . 5$

05. 60 S  $F1 \langle J + 33 \rangle = C2 * 1000 / C1 + . 5$ ; S  $F1 \langle J + 34 \rangle = C1 * 10 / F1 \langle J + 2 \rangle + . 5$

\*

\*L L, #CALC2, 0

\*W

C FOCAL-12

01. 10 L 0, F1, U, #0, 1

01. 20 A !!, "FDP", PF, "LDP", PL

01. 30 F DP=PF, PL; D 2

01. 40 T !!; L C, F1; Q

02. 10 S  $J=(DP*5+4)*256+221$

02. 20 S  $MI=F1(J+11)$ ; S  $D=F1(J+8)-MI$

02. 30 S  $K5=.5*D+MI$ ; S  $K7=.3*D+MI$ ; S  $K9=.1*D+MI$

02. 40 D 4

02. 50 D 6

04. 10 S  $K=0$ ; S  $T=0$ ; S  $J=DP*5*256$ ; S  $L=K5$ ; D 5

04. 20 S  $T5=D$ ; S  $N5=N$ ; S  $L=K7$ ; D 5

04. 30 S  $T7=D$ ; S  $N7=N$ ; S  $L=K9$ ; D 5

04. 40 S  $T9=D$ ; S  $N9=N$ ; S  $L=MI$ ; D 5

04. 50 S  $T1=D$ ; S  $N1=N$

05. 10 I  $(F1(J+K)-L)5.3, 5.3, 5.2$

05. 20 S  $T=T+F1(J+K)$ ; S  $K=K+1$ ; G 5.1

05. 30 S  $M=(F1(J+K)-F1(J+K-1))/F1(J+K)$

05. 40 S  $D=T+(L-F1(J+K-1))/M$

05. 50 S  $N=K-(F1(J+K-1)-L)/(F1(J+K)-F1(J+K-1))$

06. 10 S  $J=(DP*5+4)*256+221$

06. 20 S  $F1(J+21)=T5/10+.5$ ; S  $F1(J+22)=T7/10+.5$

06. 30 S  $F1(J+23)=T9/10+.5$ ; S  $F1(J+24)=T1/10+.5$

06. 40 S  $F1(J+25)=N5*10+.5$ ; S  $F1(J+26)=N7*10+.5$

06. 50 S  $F1(J+27)=N9*10+.5$ ; S  $F1(J+28)=N1*10+.5$

06. 60 S  $D=F1(J+1)/100$ ; S  $F1(J+29)=T5/D+.5$

06. 70 S  $F1(J+30)=T7/D+.5$ ; S  $F1(J+31)=T9/D+.5$

06. 80 S  $F1(J+32)=T1/D+.5$

\*

\*L L, \$READ, 0

\*W

C FOCAL-12

01. 10 L 0, F1, U, #0, 1

01. 20 A !, "TAPE", T, "FDP", PF, "LDP", PL

01. 30 F DP=PF, PL; D 2

01. 40 D 2. 05; T %8. 04; L C, F1; 0

02. 05 S J=(DP\*5+4)\*256+221; F I=1, 31; T " ", !

02. 10 T %3, "DATA POINT", DP+(T-1)\*100

02. 15 T !, %11, "TL", F1(J+1), !, "CT", F1(J+2)\*10

02. 20 T !, %10. 01, "RR", F1(J+3)/10, !, %11, "NT", F1(J+4)

02. 25 T !, %10. 01, "ST", F1(J+5)/10, !, "SD", F1(J+6)/10

02. 30 T !, %11. 02, "SE", F1(J+7)/100, !, %10. 01, "FI", F1(J+8)/10

02. 35 T !, "SD", F1(J+9)/10, !, %11. 02, "SE", F1(J+10)/100

02. 40 T !, %10. 01, "MI", F1(J+11)/10, !, %10, "FMN", F1(J+12)

02. 45 T !, "H5N", F1(J+13), !, %9. 01, "MEN", F1(J+14)/10

02. 50 T !, "MED", F1(J+15)/10, !, "MOD", F1(J+16)/10

02. 55 T !, %10, "MON", F1(J+17), !, %11, "PL", F1(J+18)

02. 60 T !, %10. 01, "LI", F1(J+19)/10, !, %4. 02, "SL", -F1(J+20)/100

02. 65 T !, %10, "T50", F1(J+21), !, "T70", F1(J+22)

02. 70 T !, "T90", F1(J+23), !, %9. 0, "T100", F1(J+24)

02. 75 T !, %9. 01, "N50", F1(J+25)/10, !, "N70", F1(J+26)/10

02. 80 T !, "N90", F1(J+27)/10, !, %8. 01, "N100", F1(J+28)/10

02. 85 T !, %9. 01, "T5P", F1(J+29)/10, !, "T7P", F1(J+30)/10

02. 90 T !, "T9P", F1(J+31)/10, !, %8. 01, "T10P", F1(J+32)/10

02. 95 T !, %9. 01, "PLP", F1(J+33)/10, !, "TLP", F1(J+34)/10

\*

\*L L, \$SUM1, 0

\*W

C FOCAL-12

01. 10 L 0, F1, U, #0, 1  
 01. 20 R "TAPE", T, "FDP", PF, "LDP", PL  
 01. 30 T "!!," DP RR NT ST SD"  
 01. 40 T " SE FI SD SE MI"  
 01. 50 F DP=PF, PL; D 2  
 01. 60 T "!!," %8. 04; L C, F1; Q

02. 10 S J=(DP\*5+4)\*256+221  
 02. 20 T !, %3, DP+(T-1)\*100  
 02. 30 T %6. 01, F1(J+3)/10, %7, F1(J+4), %6. 01, F1(J+5)/10  
 02. 40 T F1(J+6)/10, %6. 02, F1(J+7)/100, %6. 01, F1(J+8)/10  
 02. 50 T F1(J+9)/10, %6. 02, F1(J+10)/100, %6. 01, F1(J+11)/10  
 \*

\*L L, \$SUM2, 0

\*W

C FOCAL-12

01. 10 L 0, F1, U, #0, 1  
 01. 20 R "TAPE", T, "FDP", PF, "LDP", PL  
 01. 30 T "!!," DP FMN HSN MEN MED"  
 01. 40 T " MOD MON FL LI"  
 01. 50 F DP=PF, PL; D 2  
 01. 60 T "!!," %8. 04; L C, F1; Q

02. 10 S J=(DP\*5+4)\*256+221  
 02. 20 T !, %3, DP+(T-1)\*100  
 02. 30 T %7, F1(J+12), F1(J+13), %6. 01, F1(J+14)/10  
 02. 40 T F1(J+15)/10, F1(J+16)/10, %7, F1(J+17)  
 02. 50 T F1(J+18), %6. 01, F1(J+19)/10  
 \*

\*L L, \$SUM3, 0

\*N

C FOCAL-12

01. 10 L O, F1, U, #0, 1

01. 20 A "TAPE", T, "FDP", PF, "LDP", PL

01. 30 T "!!," DP SL T50 T70 T90

01. 40 T " T100 N50 N70 N90 N100"

01. 50 F DP=PF, PL; D 2

01. 60 T "!!," #8. 04; L C, F1; Q

02. 10 S  $J = (DP * 5 + 4) * 256 + 221$

02. 20 T !, #3, DP+(T-1)\*100, " "

02. 30 T #3. 02, -F1(J+20)/100, #7, F1(J+21), F1(J+22)

02. 40 T F1(J+23), F1(J+24), #6. 01, F1(J+25)/10

02. 50 T F1(J+26)/10, F1(J+27)/10, F1(J+28)/10

\*

\*L L, \$SUM4, 0

\*N

C FOCAL-12

01. 10 L O, F1, U, #0, 1

01. 20 A "TAPE", T, "FDP", PF, "LDP", PL

01. 30 T "!!," DP TL CT T5P T7P"

01. 40 T " T9P T10P PLP TLP"

01. 50 F DP=PF, PL; D 2

01. 60 T "!!," #8. 04; L C, F1; Q

02. 10 S  $J = (DP * 5 + 4) * 256 + 221$

02. 20 T !, #3, DP+(T-1)\*100

02. 30 T #7, F1(J+1), F1(J+2)\*10, #6. 01, F1(J+29)/10, F1(J+30)/10

02. 40 T F1(J+31)/10, F1(J+32)/10, F1(J+33)/10, F1(J+34)/10

\*

\*L L, \$REORD, 0

\*H

C FOCAL-12

01. 10 L 0, F0, U, #0, 0

01. 20 A !!, "TAPE", T, "FDP", PF, "LDP", PL; S T=(T-1)\*100

01. 30 D 2

01. 40 F R=1, 34; S L=R\*1000; F DP=PF, PL; D 3

01. 50 T !!; L C, F0; Q

02. 10 S A(1)=2; S A(2)=21; S A(3)=22; S A(4)=23

02. 20 S A(5)=24; S A(6)=18; S A(7)=1; S A(8)=3

02. 30 S A(9)=5; S A(10)=8; S A(11)=11; S A(12)=19

02. 40 S A(13)=14; S A(14)=15; S A(15)=16; S A(16)=25

02. 50 S A(17)=26; S A(18)=27; S A(19)=28; S A(20)=29

02. 60 S A(21)=30; S A(22)=31; S A(23)=32; S A(24)=33

02. 70 S A(25)=34; S A(26)=20; S A(27)=7; S A(28)=10

02. 80 S A(29)=6; S A(30)=9; S A(31)=4; S A(32)=12

02. 90 S A(33)=13; S A(34)=17

03. 10 L 0, F1, U, #0, 1

03. 20 S J=(DP\*5+4)\*256+221

03. 30 S F1(L+DP+T)=F0(J+A(R))

03. 40 L C, F1

\*



\*L L, \$IVD, 0

\*M

C FOCAL-12

01. 10 L 0, F0, U, #0, 0

01. 20 F J=1, 34; S N=0; S L=J\*1000; F K=1, 50; S N=N+1; D 2; D 3

01. 30 T !!!; L C, F0; R

02. 10 I <N-2>2. 2, 2. 15, 2. 2

02. 15 S N=N+41; R

02. 20 I <N-50>2. 3, 2. 25, 2. 3

02. 25 S N=N+32; R

02. 30 I <N-85>2. 4, 2. 35, 2. 4

02. 35 S N=N+10; R

02. 40 I <N-112>2. 5, 2. 45, 2. 5

02. 45 S N=N+6; R

02. 50 I <N-120>2. 6, 2. 55, 2. 6

02. 55 S N=N+11; R

02. 60 I <N-140>2. 7, 2. 65, 2. 7

02. 65 S N=N+37; R

02. 70 I <N-179>2. 8, 2. 75, 2. 8

02. 75 S N=N+11; R

02. 80 I <N-191>2. 9, 2. 85, 2. 9

02. 85 S N=N+7

02. 90 R

03. 10 L 0, F1, U, #0, 1

03. 20 S F1(L+K)=F0(L+N)

03. 30 L C, F1

\*

\*L L, \$IVI, 0

\*N

C FOCAL-12

01.10 L 0, F0, U, #0, 0

01.20 F J=1, 34; S N=0; S L=J\*1000; F K=1, 175; S N=N+1; D 2; D 3

01.30 T !!!; L C, F0; 0

02.05 I (N-1)2.15, 2.1, 2.15

02.10 S N=N+1; R

02.15 I (N-43)2.25, 2.2, 2.25

02.20 S N=N+7; R

02.25 I (N-82)2.35, 2.3, 2.35

02.30 S N=N+3; R

02.35 I (N-95)2.45, 2.4, 2.45

02.40 S N=N+17; R

02.45 I (N-118)2.55, 2.5, 2.55

02.50 S N=N+2; R

02.55 I (N-131)2.65, 2.6, 2.65

02.60 S N=N+9; R

02.65 I (N-177)2.75, 2.7, 2.75

02.70 S N=N+2; R

02.75 I (N-190)2.85, 2.8, 2.85

02.80 S N=N+1; R

02.85 I (N-198)2.95, 2.9, 2.95

02.90 S N=N+8

02.95 R

03.10 L 0, F1, U, #0, 1

03.20 S F1(L+K)=F0(L+N)

03.30 L C, F1

\*

\*L L,\$EVO,0

\*W

C FOCAL-12

01.10 L 0,F0,U,#0,0

01.20 F J=1,34;S N=0;S L=J\*1000;F K=1,3;S N=N+1;D 2;D 3

01.30 T !!!;L C,F0;Q

02.10 I (N-1)2.3,2.2,2.3

02.20 S N=N+19;R

02.30 I (N-21)2.5,2.4,2.5

02.40 S N=N+45;R

02.50 I (N-67)2.7,2.6,2.7

02.60 S N=N+15

02.70 R

03.10 L 0,F1,U,#0,1

03.20 S F1(L+K)=F0(L+N)

03.30 L C,F1

\*

\*L L,\$EVI,0

\*W

C FOCAL-12

01.10 L 0,F0,U,#0,0

01.20 F J=1,34;S N=0;S L=J\*1000;F K=1,105;S N=N+1;D 2;D 3

01.30 T !!!;L C,F0;Q

02.10 I (N-20)2.3,2.2,2.3

02.20 S N=N+1;R

02.30 I (N-66)2.5,2.4,2.5

02.40 S N=N+1;R

02.50 I (N-82)2.7,2.6,2.7

02.60 S N=N+1

02.70 R

03.10 L 0,F1,U,#0,1

03.20 S F1(L+K)=F0(L+N)

03.30 L C,F1

\*

```
*L L, $ROWM, 0
```

```
*N
```

```
C FOCAL-12
```

```
01. 10 L 0, F1, U, #0, 1
```

```
01. 20 A !!, "FC", CF, "LC", CL
```

```
01. 30 F U=CF, CL; D 2; D 3; D 4
```

```
01. 40 T ! ! ! ; L C, F1; R
```

```
02. 10 I <U-1> 15, 2, 15, 2, 2
```

```
02. 15 S D=. 1; R
```

```
02. 20 I <U-7> 2, 25, 2, 25, 2, 3
```

```
02. 25 S D=1; R
```

```
02. 30 I <U-25> 2, 35, 2, 35, 2, 4
```

```
02. 35 S D=10; R
```

```
02. 40 I <U-28> 2, 45, 2, 45, 2, 5
```

```
02. 45 S D=100; R
```

```
02. 50 I <U-30> 2, 55, 2, 55, 2, 6
```

```
02. 55 S D=10; R
```

```
02. 60 S D=1
```

```
03. 10 S N=0; S C1=0; S C2=0; S H=0
```

```
03. 20 S K=U*1000+1; S L=F1(K)/D
```

```
03. 30 S X=F1(K+N)/D
```

```
03. 35 I <X> 3, 4, 3, 7, 3, 4
```

```
03. 40 I <X-L> 3, 45, 3, 5, 3, 5
```

```
03. 45 S L=X
```

```
03. 50 I <H-X> 3, 55, 3, 6, 3, 6
```

```
03. 55 S H=X
```

```
03. 60 S N=N+1; S C1=C1+X; S C2=C2+X^2; G 3, 3
```

```
03. 70 S M=C1/N; S SD=FSQRT((C2-C1^2/N)/(N-1))
```

```
04. 10 T ! ! ! !, %3"ROW", U, ! !, %8. 04, "MEAN ", M, !, "%SD ", SD
```

```
04. 20 T !, "SE ", SD/FSQRT(N), !, "RANGE", H-L
```

```
04. 30 T !, "LOW ", L, !, "HIGH ", H
```

```
*
```

\*L L, \$RLIN, 0

\*W

C FOCAL-12

01.10 L D, F1, U, #0, 1  
 01.20 R !!, "FC", CF, "LC", CL  
 01.30 F U=CF, CL; D 2  
 01.40 D S; T %8.04; L C, F1; 0

02.10 D 5  
 02.20 F V=1.13; D 3  
 02.30 D 5  
 02.40 F V=14.26; D 3

03.10 S A=U; D 6  
 03.20 S XD=D  
 03.30 S A=Y; D 6  
 03.40 S YD=D  
 03.50 D 4

04.10 S N=9; S C1=0; S C2=0; S C3=0; S C4=0; S C5=0  
 04.15 S XL=U\*1000+1; S YL=Y\*1000+1  
 04.20 S X=F1(XL+N)/XD; S Y=F1(YL+N)/YD  
 04.25 I (X)4.3, 4.45, 4.3  
 04.30 S N=N+1  
 04.35 S C1=C1+X; S C2=C2+X^2; S C3=C3+Y  
 04.40 S C4=C4+Y^2; S C5=C5+X+Y; G 4.2  
 04.45 S DX=C2-C1^2/N; S DY=C4-C3^2/N  
 04.50 S NU=C5-C1\*C3/N; S R=NU/FSQT(DX\*DY)  
 04.55 S M=NU/DX; S B=(C3-C1\*M)/N  
 04.60 S S=DY-DX\*M^2; S E=FSQT(S/(N-2))  
 04.65 T ! ! !, %2.0, "X", U, " Y", Y, %9.04  
 04.70 T " B", B, " M", M\*1000, " E", E, " R", R,  
 04.75 T !, " N", N, " D", DX, " S", S, " V", R^2

05.10 F I=1.14; T " ", !  
 05.20 T "LINEAR FIT- Y=B+X\*M/1000+E"

06.10 I (R-1)6.2, 6.2, 6.3  
 06.20 S D=.1; R  
 06.30 I (R-7)6.4, 6.4, 6.5  
 06.40 S D=1; R  
 06.50 I (R-25)6.6, 6.6, 6.7  
 06.60 S D=10; R  
 06.70 S D=100

\*

\*L L, \$REXP, 0

\*N

C FOCAL-12

01. 10 L 0, F1, U, #0, 1

01. 20 R !!, "FC", CF, "LC", CL

01. 30 F U=CF, CL; D 2

01. 40 D 5; T %8. 04; L C, F1; 0

02. 10 D 5

02. 20 F V=1, 13; D 3

02. 30 D 5

02. 40 F V=14, 26; D 3

03. 10 S R=U; D 6

03. 20 S XD=D

03. 30 S R=Y; D 6

03. 40 S YD=D

03. 50 D 4

04. 10 S N=0; S C1=0; S C2=0; S C3=0; S C4=0; S C5=0

04. 15 S XL=U\*1000+1; S YL=Y\*1000+1

04. 20 S X=F1(XL+N)/XD; S Y=F1(YL+N)/YD

04. 25 I (X)4, 3, 4, 45, 4, 3

04. 30 S Y=FLOG(Y); S N=N+1

04. 35 S C1=C1+X; S C2=C2+X^2; S C3=C3+Y

04. 40 S C4=C4+Y^2; S C5=C5+X\*Y; G 4, 2

04. 45 S DX=C2-C1^2/N; S DY=C4-C3^2/N

04. 50 S NU=C5-C1\*C3/N; S R=NU/FSQT(DX\*DY)

04. 55 S M=NU/DX; S B=(C3-C1\*M)/N

04. 60 S S=DY-DX\*M^2; S E=FSQT(S/(N-2))

04. 65 T ! ! !, %2, 0, "X", U, " Y", V, %9, 04

04. 70 T " B", B, " M", M\*1000, " E", E, " R", R

04. 75 T !, " N", N, " D", DX, " S", S, " V", R^2

05. 10 F I=1, 14; T " ", !

05. 20 T "EXPONENTIAL FIT- Y=EXP(B+X\*M/1000+E)"

06. 10 I (A-1)6, 2, 6, 2, 6, 3

06. 20 S D=1; R

06. 30 I (A-7)6, 4, 6, 4, 6, 5

06. 40 S D=1; R

06. 50 I (A-25)6, 6, 6, 6, 6, 7

06. 60 S D=10; R

06. 70 S D=100

\*

\*L L, \$RLOG, 0

\*W

C FOCAL-12

01. 10 L D, F1, U, #0, 1  
 01. 20 A !!, "FC", CF, "LC", CL  
 01. 30 F U=CF, CL; D 2  
 01. 40 D 5; T %8. 04; L C, F1; Q

02. 10 D 5  
 02. 20 F V=1, 13; D 3  
 02. 30 D 5  
 02. 40 F V=14, 26; D 3

03. 10 S A=U; D 6  
 03. 20 S XD=D  
 03. 30 S A=V; D 6  
 03. 40 S YD=D  
 03. 50 D 4

04. 10 S N=0; S C1=0; S C2=0; S C3=0; S C4=0; S C5=0  
 04. 15 S XL=U\*1000+1; S YL=Y\*1000+1  
 04. 20 S X=F1(XL+N)/XD; S Y=F1(YL+N)/YD  
 04. 25 I (X)4. 3, 4. 45, 4. 3  
 04. 30 S X=FLOG(X); S N=N+1  
 04. 35 S C1=C1+X; S C2=C2+X^2; S C3=C3+Y  
 04. 40 S C4=C4+Y^2; S C5=C5+X\*Y; G 4. 2  
 04. 45 S DX=C2-C1^2/N; S DY=C4-C3^2/N  
 04. 50 S NU=C5-C1\*C3/N; S R=NU/FSQT(DX\*DY)  
 04. 55 S M=NU/DX; S B=(C3-C1\*M)/N  
 04. 60 S S=DY-DX\*M^2; S E=FSQT(S/(N-2))  
 04. 65 T ! !, %2. 0, "X", U, " Y", V, %9. 04  
 04. 70 T " B", B, " M", M, " E", E, " R", R  
 04. 75 T !, " N", N, " D", DX, " S", S, " V", R^2

05. 10 F I=1, 14; T " ", !  
 05. 20 T "LOGARITHMIC FIT- Y=B+LN(X)\*M+E"

06. 10 I (A-1)6. 2, 6. 2, 6. 3  
 06. 20 S D=. 1; R  
 06. 30 I (A-7)6. 4, 6. 4, 6. 5  
 06. 40 S D=1; R  
 06. 50 I (A-25)6. 6, 6. 6, 6. 7  
 06. 60 S D=10; R  
 06. 70 S D=100

\*

\*L L,\$RLLG,0

\*N

C FOCAL-12

01.10 L 0, F1, U, #0, 1  
 01.20 A !!, "FC", CF, "LC", CL  
 01.30 F U=CF, CL; D 2  
 01.40 D 5; T %8.04; L C, F1; 0

02.10 D 5  
 02.20 F V=1, 13; D 3  
 02.30 D 5  
 02.40 F V=14, 26; D 3

03.10 S A=U; D 6  
 03.20 S XD=D  
 03.30 S A=V; D 6  
 03.40 S YD=D  
 03.50 D 4

04.10 S N=0; S C1=0; S C2=0; S C3=0; S C4=0; S C5=0  
 04.15 S XL=U\*1000+1; S YL=Y\*1000+1  
 04.20 S X=F1(XL+N)/XD; S Y=F1(YL+N)/YD  
 04.25 I (X)4.3; 4.45, 4.3  
 04.30 S X=FLOG(X); S Y=FLOG(Y); S N=N+1  
 04.35 S C1=C1+X; S C2=C2+X^2; S C3=C3+Y  
 04.40 S C4=C4+Y^2; S C5=C5+X\*Y; G 4.2  
 04.45 S DX=C2-C1^2/N; S DY=C4-C3^2/N  
 04.50 S NU=C5-C1\*C3/N; S R=NU/FSQT(DX\*DY)  
 04.55 S M=NU/DX; S B=(C3-C1\*M)/N  
 04.60 S S=DY-DX\*M^2; S E=FSQT(S/(N-2))  
 04.65 T !!, %2.0, "X", U, " Y", V, %9.04  
 04.70 T " B", B, " M", M\*1000, " E", E, " R", R  
 04.75 T !, " N", N, " D", DX, " S", S, " Y", R^2

05.10 F I=1, 14; T " ", !  
 05.20 T "LOG-LOG FIT- Y=EXP(B+LN(X)\*M/1000+E)"

06.10 I (A-1)6.2, 6.2, 6.3  
 06.20 S D=.1; R  
 06.30 I (A-7)6.4, 6.4, 6.5  
 06.40 S D=1; R  
 06.50 I (A-25)6.6, 6.6, 6.7  
 06.60 S D=10; R  
 06.70 S D=100

\*



\*L L, \$REGSD, 0

\*N

C FDCAL-12

01. 10 A !, "N1", N1, !, "N2", N2  
 01. 20 A !, "D1", D1, !, "D2", D2  
 01. 30 A !, "S1", S1, !, "S2", S2  
 01. 40 A !, "M1", M1, !, "M2", M2  
 01. 50 S DF=N1+N2-4; S 5=(S1+S2)/DF  
 01. 60 S DI=F5QT(S/(D1+D2))  
 01. 70 S T=FRBS((M1-M2)/DI)

02. 10 S C=. 63661977  
 02. 20 S D=F5QT(DF)  
 02. 30 S TH=FATN(T/D); S X=DF/(DF+T^2)  
 02. 40 I (DF-1)2. 5, 2. 5, 3. 1  
 02. 50 S R=C\*TH; G 4. 5

03. 10 S TS=FITR(DF/2); S TS=2\*TS  
 03. 20 I (DF-TS)3. 3, 3. 3, 3. 4  
 03. 30 S IX=1; G 3. 5  
 03. 40 S IX=2  
 03. 50 S A=1; S Y=1; S Z=IX; S W=Z+1  
 03. 60 I (DF-W)4. 1, 4. 1, 3. 7  
 03. 70 S Y=X\*Y\*Z/W  
 03. 80 S A=A+Y; S Z=Z+2; S W=W+2; G 3. 6

04. 10 I (2-IX)4. 2, 4. 2, 4. 4  
 04. 20 S N=F5QT(X-X^2)  
 04. 30 S A=C\*(TH+A\*N); G 4. 5  
 04. 40 S N=F5QT(1-X); S A=A\*N  
 04. 50 S P=1-A

05. 10 T !, "DF=", DF, !, "T= ", T, !, "P= ", P, !!!

\*

\*L L, \$RSIGD, 0

\*N

C FOCAL-12

01.10 A !, "N1", N1, "N2", N2, "R1", R1, "R2", R2  
 01.20 S  $Z1 = \langle \text{FLOG}(1+R1) - \text{FLOG}(1-R1) \rangle / 2$   
 01.30 S  $Z2 = \langle \text{FLOG}(1+R2) - \text{FLOG}(1-R2) \rangle / 2$   
 01.40 S  $S = 1 / \langle N1 - 3 \rangle + 1 / \langle N2 - 3 \rangle$   
 01.50 S  $C = \text{FABS}(\langle \langle Z1 - Z2 \rangle / \text{FSQT}(S) \rangle)$ ; T !, "C=", C  
 01.60 D 2  
 01.70 G 1.1

02.10 I  $\langle C - 1.959964 \rangle 2.14, 2.17, 2.2$   
 02.14 T " N5 P=.05"; R  
 02.17 T " S P=.05"; R  
 02.20 I  $\langle C - 2.053749 \rangle 2.24, 2.27, 2.3$   
 02.24 T " S P<.05"; R  
 02.27 T " S P=.04"; R  
 02.30 I  $\langle C - 2.170090 \rangle 2.34, 2.37, 2.4$   
 02.34 T " S P<.04"; R  
 02.37 T " S P=.03"; R  
 02.40 I  $\langle C - 2.326348 \rangle 2.44, 2.47, 2.5$   
 02.44 T " S P<.03"; R  
 02.47 T " S P=.02"; R  
 02.50 I  $\langle C - 2.575829 \rangle 2.54, 2.57, 2.6$   
 02.54 T " S P<.02"; R  
 02.57 T " S P=.01"; R  
 02.60 T " S P<.01"; R  
 \*

\*L L, \$PVALR, 0

\*N

C FOCAL-12

01.10 A !, "N", N, !, "R", R  
 01.20 S DF=N-2; S D=FSQT(DF)  
 01.30 S T=FSQT(1-R^2); S T=R\*D/T  
  
 02.10 S C=.63661977  
 02.20 S D=FSQT(DF)  
 02.30 S TH=FATN(T/D); S X=DF/(DF+T^2)  
 02.40 I (DF-1)2.5, 2.5, 3.1  
 02.50 S A=C\*TH; G 4.5  
  
 03.10 S TS=FITR(DF/2); S TS=2\*TS  
 03.20 I (DF-TS)3.3, 3.3, 3.4  
 03.30 S IX=1; G 3.5  
 03.40 S IX=2  
 03.50 S A=1; S Y=1; S Z=IX; S W=Z+1  
 03.60 I (DF-W)4.1, 4.1, 3.7  
 03.70 S Y=X\*Y\*Z/W  
 03.80 S A=A+Y; S Z=Z+2; S W=W+2; G 3.6  
  
 04.10 I (2-IX)4.2, 4.2, 4.4  
 04.20 S N=FSQT(X-X^2)  
 04.30 S A=C\*(TH+A\*N); G 4.5  
 04.40 S N=FSQT(1-X); S A=A\*N  
 04.50 S P=1-A  
  
 05.10 T !, "DF=", DF, !, "T= ", T  
 05.20 T !, %10.10, "P= ", P, %8.04  
 05.30 G 1.1  
 \*

\*L L, \$STTEST, 0

\*M

C FOCAL-12

01.10 S C1=0; S C2=0; S R=0  
 01.20 A !!, "NX", N  
 01.30 F J=1, N; A !, V; S C1=C1+V; S C2=C2+V^2  
 01.40 S MX=C1/N; S K=(C2-C1^2/N)/(N-1); S DX=FSQTK(K)  
 01.50 S EX=DX/FSQTK(N); S DF=N  
 01.60 D 1.1; A !!, "NY", N; D 1.3  
 01.70 S MY=C1/N; S K=(C2-C1^2/N)/(N-1); S DY=FSQTK(K)  
 01.80 S EY=DY/FSQTK(N); S DF=DF+N-2  
 01.85 A !!, "PAIRED?", Y; I (Y-25)1.95, 1.9, 1.95  
 01.90 A !, "R", R; S DF=DF/2  
 01.95 S T=FABS((MX-MY)/FSQTK(EX^2+EY^2-2\*R\*EX\*EY))

02.10 S C=.63661977  
 02.20 S D=FSQTK(DF)  
 02.30 S TH=FATN(T/D); S X=DF/(DF+T^2)  
 02.40 I (DF-1)2.5, 2.5, 3.1  
 02.50 S A=C\*TH; G 4.5

03.10 S TS=FITR(DF/2); S T5=2\*TS  
 03.20 I (DF-T5)3.3, 3.3, 3.4  
 03.30 S IX=1; G 3.5  
 03.40 S IX=2  
 03.50 S A=1; S Y=1; S Z=IX; S W=Z+1  
 03.60 I (DF-W)4.1, 4.1, 3.7  
 03.70 S Y=X\*Y\*Z/W  
 03.80 S A=A+Y; S Z=Z+2; S W=W+2; G 3.6

04.10 I (2-IX)4.2, 4.2, 4.4  
 04.20 S N=FSQTK(X-X^2)  
 04.30 S A=C\*(TH+A\*N); G 4.5  
 04.40 S N=FSQTK(1-X); S A=A\*N  
 04.50 S P=1-A

05.10 T !!, "MX=", MX, !, "SD=", DX, !, "SE=", EX  
 05.20 T !!, "MY=", MY, !, "SD=", DY, !, "SE=", EY  
 05.30 T !!, "DF=", DF, !, "T= ", T, !, "P= ", P, !!!; Q  
 \*

\*L L,\$TIMFOL,0

\*N

C FOCRL-12

01.10 S M=0;S D=0;S S=0;T %8.02

01.20 T !,"TL MIN DIFF"

01.30 A !!,L;D 5;S Z=T;T M,D

01.40 A !!,L;D 5;S M=T-Z;S D=M-S

01.45 I <L>1.5,1.6,1.5

01.50 T M,D;S S=M;G 1.4

01.60 T !!!,%8.04;Q

05.10 I <L-50>5.15,5.15,5.2

05.15 S T=<L-8>/10.5;R

05.20 I <L-130>5.25,5.25,5.3

05.25 S T=<L-10>/10;R

05.30 I <L-206>5.35,5.35,5.4

05.35 S T=<L-16>/9.5;R

05.40 I <L-278>5.45,5.45,5.5

05.45 S T=<L-26>/9;R

05.50 I <L-380>5.55,5.55,5.6

05.55 S T=<L-40>/8.5;R

05.60 I <L-476>5.65,5.65,5.7

05.65 S T=<L-60>/8;R

05.70 I <L-596>5.75,5.75,5.8

05.75 S T=<L-86>/7.5;R

05.80 I <L-708>5.85,5.85,5.95

05.85 S T=<L-120>/7;R

05.95 S T=<L-162>/6.5;R

\*

\*L L,%PLOT,0

\*M

C FOCAL-12

01.10 L O,F0,I,#0,0

01.15 L O,F1,I,#0,1

01.20 O C

01.25 A !,"PEN DOWN",G;S P=0;S C1=0;S C2=0;S T=0

01.30 S A=FDIS(0,0);S A=FDIS(0,.68)

01.35 S A=FDIS(1.02,.68);S A=FDIS(1.02,0)

01.40 S A=FDIS(0,0);O P

01.45 O C

01.50 S A=FDIS(0,.34);S A=FDIS(1.02,.34);O P

01.55 O C

01.60 S A=FDIS(.34,0);S A=FDIS(.34,.68);O P

01.65 O C

01.70 S A=FDIS(.68,0);S A=FDIS(.68,.68);O P

01.75 O C

01.80 A "PEN UP",G

02.10 S P=P+1;S C1=FABS(C1-.34)

02.20 I (P-3)2.4,2.3,2.4

02.30 S C2=.34;S T=1

02.40 I (P-5)2.6,2.5,2.6

02.50 S C2=.68;S T=0

02.60 I (P-6)3.1,3.1,2.7

02.70 O R

02.80 T !!!;S G=F0(0);L C,F0;L C,F1;Q

03.10 T !!!,%2,"PLOT",P,%8.04

03.20 A !,"X ROW",XR,!,"X SCALE",XS

03.30 S U=XR\*1000;S D=XR;D 5;S XS=XS\*D

03.40 A !,"Y ROW",YR,!,"Y SCALE",YS

03.50 S V=YR\*1000;S D=YR;D 5;S YS=YS\*D

04.10 S K=0

04.15 S K=K+1

04.20 I (T)4.25,4.25,4.3

04.25 S X=F0(U+K);S Y=F0(V+K);G 4.35

04.30 S X=F1(U+K);S Y=F1(V+K)

04.35 I (X)4.4,4.65,4.45

04.40 S X=X+4096

04.45 I (Y)4.5,4.65,4.55

04.50 S Y=Y+4096

04.55 S A=FDIS(C2+X\*.34/XS,C1+Y\*.34/YS)

04.60 G 4.15

04.65 O P

04.70 O C

04.75 G 2.1

05.10 I (D-1)5.2,5.2,5.3

05.20 S D=.1;R

05.30 I (D-7)5.4,5.4,5.5

05.40 S D=1;R

05.50 I (D-25)5.6,5.6,5.7

05.60 S D=10;R

05.70 S D=100

\*

\*L L%REGP1.0

\*W

C FOCAL-12

```

01.10 A !!!, "PEN DOWN", G
01.20 S P=0; S C1=0; S C2=0
01.30 S P=P+1; S C1=FABS(C1-.34)
01.40 I (P-3)1.6, 1.5, 1.6
01.50 S C2=.34
01.60 I (P-5)1.8, 1.7, 1.8
01.70 S C2=.68
01.80 I (P-6)2.1, 2.1, 1.9
01.90 D R
01.95 G 1.2

02.10 D C
02.15 T !!!, Z2.0, "PLOT", P, Z8.04
02.20 A !, "REG FIT", R, !, "B", B, !, "M", M, !, "E", E
02.25 A !, "X LOW", XL, !, "X HIGH", XH; S B=B-2*E
02.30 A !, "X SCALE", XS, !, "Y SCALE", YS; S S=(XH-XL)/25
02.35 I (R-1)2.4, 2.4, 2.5
02.40 F I=1, 3; D 3
02.45 G 1.3
02.50 I (R-2)2.55, 2.55, 2.65
02.55 F I=1, 3; D 4
02.60 G 1.3
02.65 I (R-3)2.7, 2.7, 2.8
02.70 F I=1, 3; D 5
02.75 G 1.3
02.80 I (R-4)2.85, 2.85, 2.15
02.85 F I=1, 3; D 6
02.90 G 1.3

03.10 S B=B+E
03.20 D C
03.30 F J=XL, S, XH; S Y=(B+M*J)*.34/Y5; D 7
03.40 D P

04.10 S B=B+E
04.20 D C
04.30 F J=XL, S, XH; S Y=FEXP(B+M*J)*.34/Y5; D 7
04.40 D P

05.10 S B=B+E
05.20 D C
05.30 F J=XL, S, XH; S Y=(B+M*FLOG(J))* .34/Y5; D 7
05.40 D P

06.10 S B=B+E
06.20 D C
06.30 F J=XL, S, XH; S Y=FEXP(B+M*FLOG(J))* .34/Y5; D 7
06.40 D P

07.10 I (Y)7.4, 7.2, 7.2
07.20 I (Y-.34)7.3, 7.3, 7.4
07.30 S A=FDIS(C2+J*.34/XS, C1+Y)
07.40 R
*
```

\*L L, %REGP2, 0

\*N

C FOCAL-12

```

01.10 A "PEN DOWN",G,!!!;D C
01.15 S P=0;S C1=0;S C2=0
01.20 S A=FDIS<0,0>;S A=FDIS<0,.68>
01.25 S A=FDIS<1.02,.68>;S A=FDIS<1.02,0>
01.30 S A=FDIS<0,0>;D P
01.35 D C
01.40 S A=FDIS<0,.34>;S A=FDIS<1.02,.34>;D P
01.45 D C
01.50 S A=FDIS<.34,0>;S A=FDIS<.34,.68>;D P
01.55 D C
01.60 S A=FDIS<.68,0>;S A=FDIS<.68,.68>;D P
01.65 D C

02.10 S P=P+1;S C1=FABS<C1-.34>
02.15 I <P-3>2.25,2.2,2.25
02.20 S C2=.34
02.25 I <P-5>2.35,2.3,2.35
02.30 S C2=.68
02.35 I <P-6>2.5,2.5,2.4
02.40 D R
02.45 G 1.15
02.50 A !!!,"N",N
02.55 F K=1,N;D 3
02.60 D C
02.65 G 2.1

03.10 D C
03.15 T !!,%2,"PLOT",P,%8.04
03.20 A !,"REG FIT",R,!,"B",B,!,"M",M
03.25 A !,"X LOW",XL,!,"X HIGH",XH
03.30 A !,"X SCALE",XS,!,"Y SCALE",YS;S S=(XH-XL)/25
03.35 I <R-1>3.4,3.4,3.5
03.40 D C
03.42 F J=XL,S,XH;S Y=(B+M*J)*.34/YS;D 4
03.44 D P
03.46 R
03.50 I <R-2>3.55,3.55,3.65
03.55 D C
03.57 F J=XL,S,XH;S Y=FEXP<B+M*J>*.34/YS;D 4
03.59 D P
03.61 R
03.65 I <R-3>3.7,3.7,3.8
03.70 D C
03.72 F J=XL,S,XH;S Y=(B+M*FLOG<J>)*.34/YS;D 4
03.74 D P
03.76 R
03.80 I <R-4>3.85,3.85,3.1
03.85 D C
03.87 F J=XL,S,XH;S Y=FEXP<B+M*FLOG<J>)*.34/YS;D 4
03.89 D P

04.10 I <Y>4.4,4.2,4.2
04.20 I <Y-.34>4.3,4.3,4.4
04.30 S A=FDIS<C2+J*.34/XS,C1+Y>
04.40 R
*
```



\*L L,ZLINEP,0

\*W

C FOCAL-12

01.10 L 0,F1,I,#0,1

01.20 A !!, "PEN DOWN", G

01.30 D 7

01.40 A !!, "FDP", PF, !, "LDP", PL

02.10 A !!, "T100", D; S M=.34

02.20 F J=2,5; D 5

02.30 A !!, "TL", D; S M=.68

02.40 F J=6,7; D 5

02.50 A !!, "F1/LI", D; S D=D\*10; S M=1.02

02.60 F J=10,12; D 5

02.70 O R

02.80 D 7

03.10 A !!, "MEN", D; S D=D\*10; S M=.34

03.20 F J=13,15; D 5

03.30 A !!, "N100", D; S D=D\*10; S M=.68

03.40 F J=16,19; D 5

03.50 A !!, "RR/ST", D; S D=D\*10; S M=1.02

03.60 F J=8,9; D 5

03.70 O R

03.80 D 7

04.10 A !!, "SL", D; S D=-D\*100; S M=.34

04.20 S J=26; D 5

04.30 S D=1000; S M=.68

04.40 F J=20,23; D 5

04.50 S M=1.02

04.60 F J=24,25; D 5

04.70 O R

04.80 G 1.3

05.10 S L=J\*1000

05.20 F K=PF, PL; S V=F1(L+K); D 6

05.30 O P

05.40 O C

06.10 I (V)6.2,6.3,6.3

06.20 S V=V+4096

06.30 S A=FDIS(M-V\*.34/D,.68\*(K-PF)/(PL-PF))

07.10 O C

07.20 S A=FDIS(0,0); S A=FDIS(0,.68)

07.30 S A=FDIS(1.02,.68); S A=FDIS(1.02,0)

07.40 S A=FDIS(0,0); O P

07.50 O C

07.60 S A=FDIS(.34,0); S A=FDIS(.34,.68); O P

07.70 O C

07.80 S A=FDIS(.68,0); S A=FDIS(.68,.68); O P

07.90 O C

\*

Table VIII. Regression equations of 24 parameters on respiration rate for I-VO cells.

1. T50 = 147.2220 + 21.1092 · ln(RR) ± 98.7780
2. T70 = 156.0950 + 63.3316 · ln(RR) ± 165.1760
3. T90 = 405.9660 + 44.2711 · ln(RR) ± 253.2590
4. T100 = EXP[6.8533 - 0.0193(RR) ± 0.7480]
5. PL = 792.7180 - 16.8713(RR) ± 273.1040
6. TL = EXP[7.9172 - 0.0355(RR) ± 0.2342]
7. FI = 88.0191 - 1.5262(RR) ± 25.5480
8. MI = 36.6944 - 4.5819 · ln(RR) ± 10.8402
9. LI = 75.3811 - 13.1186 · ln(RR) ± 12.7498
10. MEN = 52.9135 - 7.5962 · ln(RR) ± 12.4183
11. MED = 43.3729 - 5.4107 · ln(RR) ± 11.9047
12. MOD = 35.1646 - 3.3601 · ln(RR) ± 10.9786
13. N50 = -0.0234 + 1.4496 · ln(RR) ± 1.7945
14. N70 = -1.4086 + 3.1496 · ln(RR) ± 4.4809
15. N90 = 3.9172 + 3.4890 · ln(RR) ± 6.5986
16. N100 = EXP[3.3038 - 0.0088(RR) ± 0.4769]
17. ST = 124.0940 - 24.1712 · ln(RR) ± 25.8741
18. SL = EXP[-0.6432 - 0.0583(RR) ± 0.9638]
19. T5P = -6.4697 + 7.1477 · ln(RR) ± 6.0545
20. T7P = -14.6178 + 12.9528 · ln(RR) ± 9.6405
21. T9P = 14.2368 + 17.3266 · ln(RR) ± 14.9481
22. T10P = 6.1723 + 16.8245 · ln(RR) ± 21.9979
23. PLP = EXP[2.4491 + 0.3447 · ln(RR) ± 0.4723]
24. TLP = EXP[2.3434 + 0.4872 · ln(RR) ± 0.2374]

Table IX. Regression equations of 24 parameters on respiration rate for I-VI cells.

1. T50 = EXP[5.2791 - 0.0168(RR) ± 0.4879]
2. T70 = EXP[5.9359 - 0.0207(RR) ± 0.4527]
3. T90 = 544.7000 - 5.3169(RR) ± 110.3810
4. T100 = 1822.4400 - 367.3070 · ln(RR) ± 151.5740
5. PL = EXP[8.9515 - 0.9727 · ln(RR) ± 0.5897]
6. TL = EXP[7.4751 - 0.0216(RR) ± 0.3241]
7. FI = EXP[4.1577 - 0.0144(RR) ± 0.4247]
8. MI = EXP[3.0236 - 0.0097(RR) ± 0.4929]
9. LI = EXP[3.4636 - 0.0084(RR) ± 0.5742]
10. MEN = EXP[3.3497 - 0.0103(RR) ± 0.4487]
11. MED = EXP[3.2064 - 0.0101(RR) ± 0.4596]
12. MOD = EXP[3.0901 - 0.0097(RR) ± 0.4704]
13. N50 = EXP[1.3283 - 0.0029(RR) ± 0.5420]
14. N70 = EXP[2.1464 - 0.0079(RR) ± 0.6085]
15. N90 = EXP[3.2366 - 0.1684 · ln(RR) ± 0.5423]
16. N100 = EXP[4.4251 - 0.3482 · ln(RR) ± 0.5332]
17. ST = 133.7710 - 24.1028 · ln(RR) ± 26.2316
18. SL = 0.2676 + 0.0009(RR) ± 0.1664
19. T5P = EXP[1.6797 + 0.2574 · ln(RR) ± 0.6053]
20. T7P = 13.8721 + 3.2756 · ln(RR) ± 12.3565
21. T9P = 14.4688 + 8.2056 · ln(RR) ± 14.5038
22. T10P = 44.8520 + 5.2108 · ln(RR) ± 15.8760
23. PLP = 53.8008 - 5.4440 · ln(RR) ± 13.9808
24. TLP = EXP[3.1710 + 0.1823 · ln(RR) ± 0.3230]

Table X. Regression equations of 24 parameters on respiration rate for E-VI cells.

1. T50 = 414.4040 - 80.5027 · ln(RR) ± 49.1951
2. T70 = 675.8920 - 133.5940 · ln(RR) ± 78.0391
3. T90 = 1438.1000 - 299.7580 · ln(RR) ± 147.7610
4. T100 = 2617.3300 - 569.2290 · ln(RR) ± 219.7980
5. PL = 1495.4300 - 325.7690 · ln(RR) ± 162.7690
6. TL = 3048.0900 - 632.1220 · ln(RR) ± 203.9220
7. FI = 164.2190 - 31.3311 · ln(RR) ± 15.0115
8. MI = 47.4782 - 8.5725 · ln(RR) ± 5.9890
9. LI = 50.0437 - 7.2535 · ln(RR) ± 7.5328
10. MEN = 63.2751 - 11.2802 · ln(RR) ± 7.1998
11. MED = 55.7401 - 10.0034 · ln(RR) ± 6.7460
12. MOD = 51.5672 - 9.4249 · ln(RR) ± 6.4786
13. N50 = 2.8954 + 0.0026(RR) ± 1.0860
14. N70 = EXP[1.9225 - 0.0856 · ln(RR) ± 0.4301]
15. N90 = EXP[3.3632 - 0.2598 · ln(RR) ± 0.5371]
16. N100 = EXP[4.8054 - 0.4883 · ln(RR) ± 0.5994]
17. ST = EXP[3.8809 - 0.0087(RR) ± 0.4402]
18. SL = 0.3539 - 0.0007(RR) ± 0.1330
19. T5P = 12.1021 + 0.1206(RR) ± 5.9030
20. T7P = 18.5201 + 0.1848(RR) ± 8.2920
21. T9P = 41.1972 + 0.1200(RR) ± 12.9830
22. T10P = 105.7470 - 9.9173 · ln(RR) ± 16.8018
23. PLP = 49.2392 - 0.2522(RR) ± 13.3991
24. TLP = 31.3642 + 0.3127(RR) ± 12.2442

Table XI. Regression equations of 24 parameters on mode for I=VO cells.

1. T50 = EXP[1.0320 + 1.2984 · ln(MOD) ± 0.4121]
2. T70 = EXP[0.8938 + 1.4780 · ln(MOD) ± 0.4866]
3. T90 = EXP[1.3322 + 1.4884 · ln(MOD) ± 0.5060]
4. T100 = EXP[2.8979 + 1.1556 · ln(MOD) ± 0.4785]
5. PL = EXP[4.8709 + 0.3974 · ln(MOD) ± 0.4734]
6. TL = EXP[6.9655 + 0.0141(MOD) ± 0.2555]
7. FI = EXP[1.6716 + 0.7526 · ln(MOD) ± 0.2953]
8. MI = EXP[-0.2865 + 1.0628 · ln(MOD) ± 0.0819]
9. LI = EXP[3.1848 + 0.0164(MOD) ± 0.3305]
10. MEN = EXP[0.6723 + 0.8603 · ln(MOD) ± 0.0963]
11. MED = EXP[0.1401 + 0.9860 · ln(MOD) ± 0.0701]
12. RR = EXP[2.8361 - 0.0035(MOD) ± 0.3519]
13. N50 = EXP[-0.2682 + 0.4897 · ln(MOD) ± 0.3919]
14. N70 = EXP[-0.3260 + 0.6783 · ln(MOD) ± 0.5045]
15. N90 = EXP[0.5364 + 0.6099 · ln(MOD) ± 0.5160]
16. N100 = EXP[2.7940 + 0.1159 · ln(MOD) ± 0.4755]
17. ST = EXP[5.7306 - 0.5586 · ln(MOD) ± 0.2817]
18. SL = 1.5777 - 0.4054 · ln(MOD) ± 0.1762
19. T5P = EXP[-0.7570 + 1.0007 · ln(MOD) ± 0.4944]
20. T7P = EXP[-0.9009 + 1.1821 · ln(MOD) ± 0.5395]
21. T9P = EXP[-0.4554 + 1.1903 · ln(MOD) ± 0.5328]
22. T10P = EXP[1.1015 + 0.8601 · ln(MOD) ± 0.4767]
23. PLP = EXP[3.0794 + 0.1005 · ln(MOD) ± 0.4852]
24. TLP = 9.3865 + 10.1569 · ln(MOD) ± 10.7925

Table XII. Regression equations of 24 parameters on mode for I-VI cells.

1. T50 = EXP[3.4807 + 0.4385 · ln(MOD) ± 0.4992]
2. T70 = EXP[4.2152 + 0.3596 · ln(MOD) ± 0.5112]
3. T90 = EXP[5.0715 + 0.2638 · ln(MOD) ± 0.4108]
4. T100 = EXP[5.5763 + 0.2304 · ln(MOD) ± 0.4115]
5. PL = 118.0350 + 74.1879 · ln(MOD) ± 214.8230
6. TL = EXP[5.8916 + 0.2984 · ln(MOD) ± 0.4226]
7. FI = EXP[1.8325 + 0.6608 · ln(MOD) ± 0.3431]
8. MI = EXP[-0.1469 + 1.0296 · ln(MOD) ± 0.0848]
9. LI = EXP[0.5052 + 0.9691 · ln(MOD) ± 0.3438]
10. MEN = EXP[0.4260 + 0.9319 · ln(MOD) ± 0.1187]
11. MED = EXP[0.1874 + 0.9697 · ln(MOD) ± 0.0754]
12. RR = 57.9943 - 8.2390 · ln(MOD) ± 13.6773
13. N50 = EXP[1.5490 - 0.0186(MOD) ± 0.5220]
14. N70 = 12.3624 - 0.2541(MOD) ± 5.6031
15. N90 = EXP[3.2714 - 0.0358(MOD) ± 0.4619]
16. N100 = EXP[3.9485 - 0.0425(MOD) ± 0.4265]
17. ST = EXP[4.4601 - 0.0402(MOD) ± 0.4317]
18. SL = 0.4408 - 0.0518 · ln(MOD) ± 0.1649
19. T5P = EXP[2.4029 + 0.0101(MOD) ± 0.6075]
20. T7P = EXP[3.0127 + 0.0048(MOD) ± 0.5512]
21. T9P = 49.1428 - 2.1869 · ln(MOD) ± 14.7922
22. T10P = 75.3259 - 4.4685 · ln(MOD) ± 15.8466
23. PLF = EXP[3.5634 - 0.0067(MOD) ± 0.4934]
24. TLP = EXP[3.5274 + 0.1022 · ln(MOD) ± 0.3264]

Table XIII. Regression equations of 24 parameters on mode for E-VI cells.

1. T50 = 43.8866 + 4.6491(MOD) ± 51.7531
2. T70 = 75.7932 + 6.8930(MOD) ± 87.0515
3. T90 = 92.4733 + 15.4179(MOD) ± 171.5270
4. T100 = 178.3820 + 22.8032(MOD) ± 312.2210
5. PL = 205.4960 + 7.1587(MOD) ± 224.2730
6. TL = 406.0200 + 21.6303(MOD) ± 336.7980
7. FI = 19.3363 + 1.8472(MOD) ± 15.9847
8. MI = EXP[0.0741 + 0.9541 · ln(MOD) ± 0.0576]
9. LI = EXP[1.4074 + 0.6132 · ln(MOD) ± 0.2543]
10. MEN = 2.9307 + 1.1204(MOD) ± 1.7804
11. MED = 1.4173 + 1.0386(MOD) ± 1.1502
12. RR = EXP[4.2340 - 0.0372(MOD) ± 0.4073]
13. N50 = 4.1876 - 0.4250 · ln(MOD) ± 1.0718
14. N70 = 8.5037 - 1.0773 · ln(MOD) ± 2.2008
15. N90 = 24.4363 - 4.1020 · ln(MOD) ± 5.6749
16. N100 = 58.5247 - 11.7915 · ln(MOD) ± 13.4960
17. ST = 98.2588 - 21.4823 · ln(MOD) ± 16.2235
18. SL = 0.3880 - 0.0034(MOD) ± 0.1308
19. T5P = EXP[2.2873 + 0.1666 · ln(MOD) ± 0.3945]
20. T7P = EXP[2.8490 + 0.1194 · ln(MOD) ± 0.3965]
21. T9P = 37.6028 + 0.4654(MOD) ± 12.6321
22. T10P = 60.9340 + 0.5264(MOD) ± 17.0150
23. PLP = 52.2560 - 4.6586 · ln(MOD) ± 14.0615
24. TLP = EXP[4.6441 - 0.32.46 · ln(MOD) ± 0.2601]

Table XIV. Best fitting regression equations for 24 parameters on respiration rate and mode for I-VO, I-VI and E-VI cells: linear (LIN); exponential (EXP); logarithmic (LOG); log-log (LLG).

	<u>Parameter</u>	<u>I-VO</u> <u>(RR)</u>	<u>I-VI</u> <u>(RR)</u>	<u>E-VI</u> <u>(RR)</u>	<u>I-VO</u> <u>(MOD)</u>	<u>I-VI</u> <u>(MOD)</u>	<u>E-VI</u> <u>(MOD)</u>
1	T50	LOG	EXP	LOG	LLG	LLG	LIN
2	T70	LOG	EXP	LOG	LLG	LLG	LIN
3	T90	LOG	LIN	LOG	LLG	LLG	LIN
4	T100	EXP	LOG	LOG	LLG	LLG	LIN
5	PL	LIN	LLG	LOG	LLG	LOG	LIN
6	TL	EXP	EXP	LOG	EXP	LLG	LIN
7	FI	LIN	EXP	LOG	LLG	LLG	LIN
8	MI	LOG	EXP	LOG	LLG	LLG	LLG
9	LI	LOG	EXP	LOG	EXP	LLG	LLG
10	MEN	LOG	EXP	LOG	LLG	LLG	LIN
11	MED	LOG	EXP	LOG	LLG	LLG	LIN
12	MOD (RR)	LOG	EXP	LOG	(EXP)	(LOG)	(EXP)
13	N50	LOG	EXP	LIN	LLG	EXP	LOG
14	N70	LOG	EXP	LLG	LLG	LIN	LOG
15	N90	LOG	LLG	LLG	LLG	EXP	LOG
16	N100	EXP	LLG	LLG	LLG	EXP	LOG
17	ST	LOG	LOG	EXP	LLG	EXP	LOG
18	SL	EXP	LIN	LIN	LOG	LOG	LIN
19	T5P	LOG	LLG	LIN	LLG	EXP	LLG
20	T7P	LOG	LOG	LIN	LLG	EXP	LLG
21	T9P	LOG	LOG	LIN	LLG	LOG	LIN
22	T10P	LOG	LOG	LOG	LLG	LOG	LIN
23	PLP	LLG	LOG	LIN	LLG	EXP	LOG
24	TLP	LLG	LLG	LIN	LOG	LLG	LLG



APPROVAL SHEET

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The final copies have been examined by the director of the dissertation and the signature which appears below verifies the fact that any necessary changes have been incorporated and that the dissertation is now given final approval by the Committee with reference to content and form.

The dissertation is therefore accepted in partial fulfillment of the requirements for the degree of Doctor of Philosophy.

August 9, 1973  
Date

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