EXPERIMENTAL PROCEDURES AND DATA COMPILATION TECHNIQUES FOR ULTRASONIC DOSIMETRIC ANALYSIS

BY

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THESIS

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Chapter 1

INTRODUCTION

In recent years the use of ultrasound in medical applications has increased dramatically. Projected sales of ultrasonic medical instrumentation should surpass one hundred million dollars by 1985 (Electronics, 1979). In an effort to bring new technology to the marketplace, the area of ultrasonic dosimetry has been neglected. Dosimetry is concerned with defining a quantitative relationship between some physical agent and the biological effect it produces. In the case of ultrasound a specific parameter such intensity, particle displacement, or acoustic pressure for example, would be related to the likelihood of the occurence of a biological alteration, usually located at the same irradiation site. This relationship is not meant to suggest that the clinical use of ultrasound is a hazard. There have been a number of studies though, which show that sufficient levels of ultrasound can produce irreversible biological damage. The increased clinical employment of ultrasonic instrumentation therefore, demands that information concerning the interaction of ultrasound energy with tissues be obtained.

To accomplish the objective of ultrasonic dosimetry, it is necessary to quantify accurately and precisely the ultrasonic source output parameters, determine in what manner the material effects the propagating energy, and relate these two items to a quantitative parameter determination at the site of interest. Therefore, the physical quantities which describe an interaction at some biological site must be defined and integrated into a concept that is applicable in ultrasonic radiation protection. The main objective of this thesis is to develop an ultrasonic dosimetric model from which the ultrasonic energy being absorbed or interacting with biological tissue can be determined.

To facilitate this model, the basic procedure will be to detect the distribution of the ultrasonic energy inside the specimen during irradiation. This will be done by first characterizing the complex ultrasonic field distribution patterns in the free field, and then inserting the specimen in the field to measure the <u>in vivo</u> field distribution within the specimen using the transient thermoelectric technique (Fry and Fry, 1954a and 1954b). Using this thermocouple technique to measure the tissue absorption coefficient, with a known spatial and temporal intensity and the exposure time, the amount of energy absorbed by the tissue may be found. This absorbed energy distribution with possible units of J/mm**3, can be related directly to the corresponding

biological effects. The parameter most widely used in bioeffect and biophysical studies is intensity. Given in mixed units of W/cm**2, intensity is primarily used because it is an easily measured parameter. It is not acceptable as a dosimetric quantity however, because it is an exposure condition, and not a measure of dose.

Some in vivo work has been done by other researchers. Most notably, the in utero work of (Hall and Robinson, 1974) and the $\underline{i}\underline{\hat{n}}$ $\underline{\hat{v}}\underline{\hat{v}}\underline{\hat{v}}\underline{\hat{o}}$ work of Bang (1972). A universal dosimetric response to ultrasonic exposure for different tissues suggested (Johnston and Dunn, 1976), but demonstrated in the case of suprathreshold lesions in brain tissue only. The experiment described in this thesis was not concerned with suprathreshold intensities as no biological damage assessment was undertaken. Rather, the philosophy work is to examine the interactions between the ultrasonic energy and the tissues, determine tissue characteristics such as the absorption coefficient, determine the energy absorbed within the specimen. Using the instrumentation and procedures described in the following chapters, a dosimetric model will be developed to find the absorption coefficient and absorbed energy of an irradiated tissue site. Improvements in both the experimental procedure and dosimetric model are proposed in Chapter 5.

CHAPTER 2

INSTRUMENTATION

This chapter details the instrumentation used in this experiment. Associated theory is presented where necessary. The instrumentation may be classified into three main categories 1) the ultrasonic source and associated driving hardware, 2) the thermocouple probe and recording device, and 3) the irradiation apparatus.

Section 2.1 The Ultrasonic Transducer

In the development of this experiment a number of constraints were placed on the acoustic field. First, the transducer must produce a field which can be directed at a particular location in the specimen. Also, for ease of analysis, it would be convenient to assume plane wave geometry at the irradiation site. To satisfy these criteria, a transducer which produces a focused field must be used. It was decided to use an existing focusing lens transducer. Other types of focusing transducers are discussed elsewhere (Fry and Dunn, 1962).

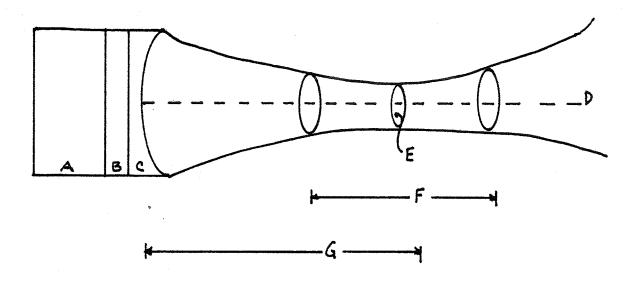
Lens transducers are often constructed in a sandwich arrangement consisting of a piezoelectric crystal, a spacing

material, and a plano concave lens. An arrangement of this type is shown in figure 1. In this transducer quartz is used for the oscillating plate. Quartz is often used for the crystal because of its high stability and high tensile strengths compared to ceramic crystals. Quartz has the disadvantage of requiring a relatively high electric field strength to obtain high power outputs due to the high input impedance of quartz. Thus, the power output of a quartz radiator is determined by the crystal and crystal holder.

The spacing material is used as a coupling medium between the crystal and the lens. Its thickness is usually one quarter of a wavelength to assure total transmission. A material this thin also has low sound attenuation. The spacer fluid in this case was Silicon 710 (Dow Corning Midland, Mich.), which was also used behind the transducer to aid in heat conduction.

The lens is a critcal component of the system as it must meet a number of criteria. The lens focuses the sound by the refraction of the wave as it passes through a media of differing sound velocity. However, when there is a refractive index n,

$$n = \frac{C_1}{C_2} \tag{1}$$



- A) QUARTZ OSCILLATOR
- B) SPACER MATERIAL
- C) LUCITE LENS
- D) FOCAL AXIS
- E) FOCAL AREA (DIAMETER = Dt)
- F) DEPTH OF FOCUS (Da)
- G) FOCAL LENGTH (F)

Figure 1. Schematic of transducer with focused field parameters (NEMA/AIUM Safety Standard, 1979)

where cl and c2 are the sound velocities in medium 1 and medium 2 respectively, there is an impedance discontinuity. This discontinuity may be designated by the ratio of the impedances given by;

$$m = P_1 C_1 / P_2 C_2$$
 (2)

where p_i C_i is the characteristic impedance of the lens and p_2 C_2 is the characteristic impedance of the propagating medium. This leads to a reflection coefficient R;

$$R = \left(\frac{\rho_2 C_2 - \rho_1 C_1}{\rho_2 C_2 + \rho_1 C_1}\right)^2 \tag{3}$$

The conditions for a good lens then must be n 1 and m=1 The material must also be machinable and have a low attenuation coefficient. The lens used in this transducer is Lucite. The characteristic parameters of this lens are $\rho = 1.18E03$ kg/m3, c=2.8E03 m/s, n=1.9, m=2.2 and R=0.14 (Hueter and Bolt,1955). This lens is mounted flush to the spacer/crystal face. This unit is contained in a stainless steel housing with a rear connector to accomodate a shielded high voltage cable.

The driving network used to supply the high voltage necessary to produce oscillation of the transducer crystal is shown in figure 2. This R.F system provides an output on any

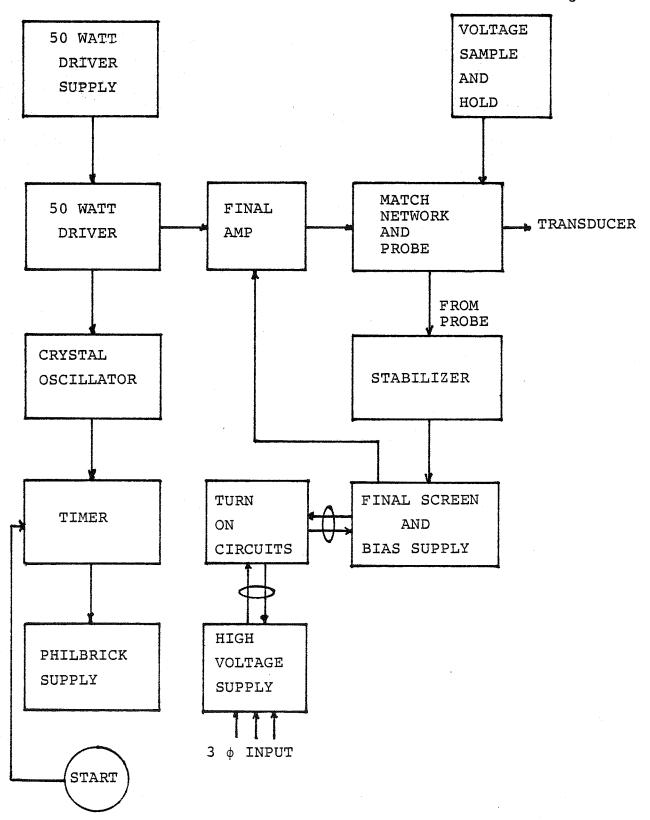


Figure 2. 2 kW driving system block diagram

of several frequencies at an R.F. power level up to 2 kW. It provides a closely controlled envelope (power level) from about 0.5 watts to over 2,000 watts by means of detected negative feedback. Frequencies are selected by means of a switch on the crystal oscillator and by changing several plug in coils. The R.F. frequency used in this experiment was 0.98 MHz which is the resonant frequency of the transducer. All interconnections for R.F. signals are made with matched 50 ohm coaxial cables.

Referring to figure 2, a start command (ground signal) is received by the timer chassis. This chassis generates square wave pulse with a time duration selected by a front panel switch. The pulse from the time chassis turns on buffer amplifier in the crystal oscillator chassis. oscillator is followed by a coarse and fine step attenuator which varies the output of the oscillator chassis from nearly 0 watts to 2 watts. The oscillator chassis drives a 50 watt amplifier, which in turn drives the 2 kW final amplifier. The output from the 2 kW final amplifier is connected to a matching network which transforms the 50 ohm output impedance of the final amplifier to the large input impedance of transducer. The rectified R.F. (now a changing d.c. level) connected to a high gain amplifier, the stabilizer. output of the stabilizer is connected to the 2 kW amplifier screen supply voltage. This causes an increase or decrease

in the R.F. envelope in the opposite sense of the original disturbance. This feedback is meant to hold the output of the system to within 0.1% of the desired level. The stabilizer can be connected to the screen supply of the 50 watt driver when the system is used in low level operation.

Included in the matching network is a capacitive voltage divider to produce the transducer driving voltage. The voltage divider ratio may be varied by a precision variable capacitor in the voltage divider. This network provides a means of obtaining variable acoustic intensities by changing the voltage applied to the transducer crystal. This applied voltage is given by the relation;

$$V = 2.86 \times 10^{2} + 4.98 * C_{5}$$
 (4)

where Cs is the variable capacitor dial setting. The resultant ultrasonic intensity produced by the transducer is described by;

$$I = \left(\frac{V}{V_{CAL}}\right)^2 * I_{CAL} \tag{5}$$

where Ical and Vcal are the calibrated intensity and voltage respectively. These two quantities are determined for the transducer in a process which involves the transient thermoelectric technique and the elastic sphere radiometer (Dunn et al, 1977). In a typical intensity calculation, the

voltage obtained with a capacitor setting of Cs=360 is V=2078.8 volts. The resulting intensity produced by this voltage using Ical=5.5 W/cm**2 and Vcal=685 V is I=50.8 W/cm**2. The intensities produced by the transducer in this experiment fall in the range of 10-60 W/cm**2.

It will be assumed for ease of analysis that the quartz plate may be modeled as a plane piston source. The field produced by such a source is mathematically described in a number of texts (Kinsler and Frey, 1962). This field possesses two distinct regions. In this analysis it is assumed that the transducer radius is much greater that than the wavelength. In the near field, known as the Fresnel zone, the axial intensity runs through a series of maxima of constant amplitude with intervening nulls. The last null occurs in the vicinity of $d=a**2/2*\lambda$, where d is the axial distance away from the source face, a is the radius of the piston, and λ is the wavelength. At d=a**2/ λ the intensity goes through its final maximum and as d>a**2/x, the intensity decreases inversely as the square of the distance. This region where $d>a**2/\lambda$ is called the far field or Fraunhofer zone. This complex field pattern has been modeled by a number of authors (Fry and Dunn, 1962).

When a plano concave lens is placed over this quartz source, the field structure is changed drastically. The

unfocused radiated sound for distances less than a**2/ may be thought of as being confined within a cylinder of radius a, whereas beyond this point there is at least approximate spherical divergence. On the other hand, the focused field intensity will converge at a designed focal length and then diverge. It is imperative for focusing that the focal length of the lens occurs inside the axial position of the zero order maxima of the unfocused field. This limits the choice of focal length for a particular crystal. In the case of the crystal used in the experimental transducer, the zero order maxima occurs at a**2/ which is 20 cm. As will be revealed shortly, the actual focal length for this transducer is well below this limit.

Having determined the intensity of the acoustic field, it is of great importance to determine its shape. This shape is determined by a number of lens characteristics. First, the focal length F is given by;

$$F = \frac{r}{1 - \frac{1}{n}} \tag{6}$$

where r is the lens radius of curvature, and n is the index of refraction for the lens. This formula is derived from the analogous formulas for optics. It must be noted that the sound velocity in the lens is greater than that of water, so the shape for an ultrasonic converging lens is opposite that

for an optical converging lens. Furthermore, the diameter to wavelength ratio is much smaller here than in optical lenses, thus the convergence in the acoustic case will be several orders of magnitude larger. Noting this, and assuming the lens is so thin that the axial length is negligible, equation 6 can be used to find the focal length.

To characterize the "working" part of the focused field, two other parameters are introduced. The schematic representation of these beam profile parameters are shown in figure 2. Dt is defined as the diameter of the focal area. This is the distance between the off axial points that are one half the intensity of the axial point. The relation to find this distance numerically is given by;

$$Dt = kt (F/D) * \lambda$$
 (7)

where the quantity kt is dimensionless and somewhat dependent upon the half aperture angle Ψ . This width Dt is taken in the neighborhood of the focal length and is not applicable further axially from the transducer. The other parameter is the axial length of the depth of focus, Da. This length is the distance between the two axial points which are at one half the maximal axial intensity. Da is given by;

$$Da = Ka * Dt$$
 (8)

where ka is dependent on the half aperture angle. approximations may be derived from the fundamental equations presented by O'Neil(1949) after making a few assumptions. First, the point under analysis is a large distance away from the transducer. Secondly, the radius of the transducer is larger than wavelength. Finally, the intensity distribution is gaussian, so specific values may be found on it. In the case of this particular transducer, it will shown that all these conditions are satisfied.

It is of great importance to quantify Da and Dt as given The lens was measured with a micrometer. From the geometry in figure 3, the radius of curvature was found to be 34.2 mm. From equation 6 the focal length was calculated to be 72.2mm. The half aperture angle " measured at the focal point is 13.6 degrees. To find Da and Dt, the k constants must be evaluated. For half aperture angle less than 50 degrees, Dt will be within 20% if kt=1. Therefore, for the transverse focal diameter is found to be 3.3 mm. value of ka is also dependent on the half aperture angle and in this case is about 13 (Fry and Dunn, 1962, figure 26). Using this value, the depth of focus should be about 40 mm. In the following chapter these field parameters determined experimentally, and the resultant values will be compared against those obtained above.

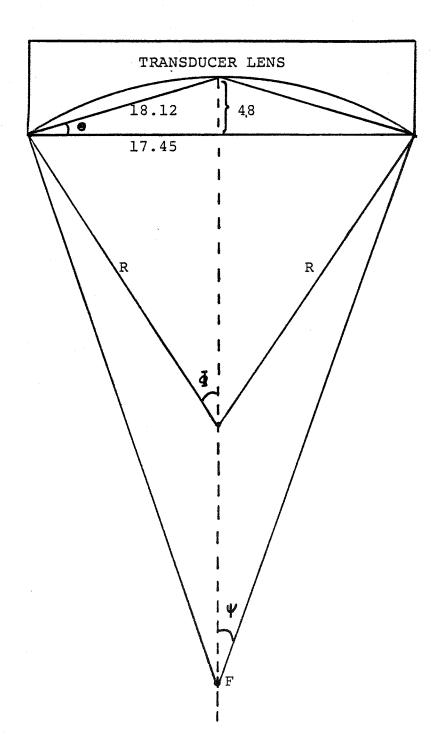


Figure 3. Lens geometry used for calculation of radius of curvature and focal length all lengths are in millimeters

Section 2.2 The Thermocouple Probe

The primary object of this study is to determine the absorption coefficient of an irradiated biological specimen and determine the absorbed energy. The ultrasonic probe which would be used to accomplish this must have a number of The probe should be small to mimimize qualities. insertion damage to the tissue and not interrupt ultrasonic beam propagation. Also, the probe should provide a means to obtain a value for absorption only. Furthermore, the probe should be mobile within the specimen to analysis at a number of different tissue sites. The only probe available which satisfies these criteria is the thermocouple (Fry and Fry, 1954a and 1954b). The probe consists of two wires of different metals welded or soldered together at one end to form a thermocouple junction. method involves placing the thermocouple junction in tissue of interest, and irradiating it with a ultrasonic pulse of known intensity, duration and carrier frequency. This pulse will interact with the tissue in such a manner as to increase the tissue temperature. When this occurs, emf of the thermocouple will change and a voltage difference will be measured across the thermocouple leads. This difference can be correlated to a time rate of change of tissue temperature. When the acoustic intensity at the

junction is known, the following relation can be used to find the absorption coefficient \mathbf{q} ,

where $\slash\hspace{-0.6em}P$ C is the product of the tissue density and heat capacity (J/°Ccm**3), I is the sound intensity (W/cm**2), and (dT/dt) is the initial time rate of temperature(°C/sec). Therefore, from directly measuring (dT/dt), we can obtain a value for the absorption coefficient for the tissue under test.

The typical response of a thermocouple probe to a single acoustic pulse having a one second rectangular envelope is shown in figure 4. A relatively rapid rise occurs just after the initiation of the pulse followed by a slower "linear" rising phase for the remainder of the one second interval. After cessation of the pulse, a rapid fall in temperature occurs followed by a slower return of the temperature to value preceeding the acoustic pulse. The initial rapid increase in temperature results from the conversion of acoustic energy into heat by the viscous forces acting between the thermocouple wire and the imbedding medium. phase of the response approaches equilibrium rapidly. the magnitude of this viscous heating depends on the angle of incidence of the sound field with respect to the thermocouple

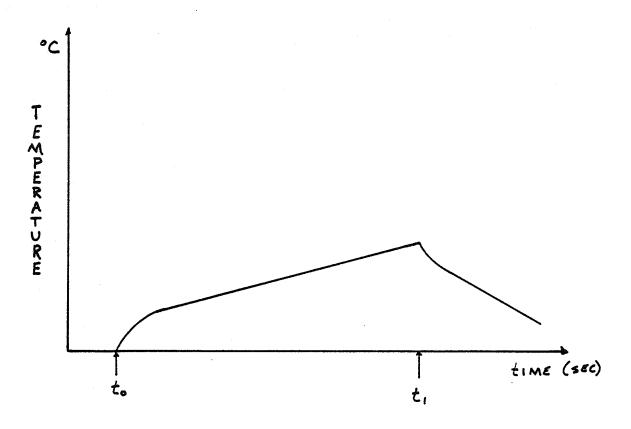


Figure 4. Typical thermocouple response (to is time at beginning of acoustic pulse. to time at end of acoustic pulse.

wire, it is assumed the wire is transverse to the direction of propagation. The second phase of the temperature sequence the "linear" part, is produced by absorption of sound in the body of the imbedding medium. The closeness of approach of this phase to linearity during irradiation is dependent on;

1) the acoustic amplitude; 2) the form of the variation of acoustic absorption coefficient with temperature; 3) the heat conductivity coefficients of the tissue and the wires; 4) the duration of the acoustic disturbance; 5) the acoustic field distribution.

thermocouple wire size and ultrasonic beam width play a major role in the accuracy of the determination of the absorption coefficient. Different combinations of thermocouple wire diameter and beam widths can produce widely varying thermocouple emf responses, which in turn yield erroneous absorption coefficient values (Goss et al, It was shown that the use of small diameter wires (13 µm) would minimize errors in the absorption coefficient determination. The use of small diameter wires corresponds to a mechanically weaker junction. As will be seen Chapter 3, the thermocouple will be subjected to mechanical The junction must be able to withstand this for the duration of the experiment. Previous work done in this lab suggested the use of 76 µm diameter wires compromise between small diameter and mechanical junction strength. The next section deals with the manufacturing process of a 76 jum diameter thermocouple.

<u>Section</u> 2.2.1 Thermocouple fabrication

As discussed previously, the thermocouple junctions to be used in this experiment must have a number of qualities. A butt welded 76 µm diameter chromel-constantan thermocouple was designed to satisfy these criteria. The design allows for smooth passage through the specimen, has a sensitivity of 60 µV/C (Omega Temperature Measurement Catalog,1980), can be subjected to experimental stresses without breaking, and uninterruption of the acoustic field may be assumed since the wire radius is much smaller than the wavelength. A magnified view of a 76 µm diameter thermocouple is shown in figure 5. Notice in this photo, the butt weld is well aligned. This junction has no protruding edge which might snag on tissue. The manufacture technique to assure this consistent alignment is as follows.

A square section of 76 µm mylar plastic was sandwiched between a glass slide and coverslip (figure 6). A fingertip held junction manipulator was also fashioned from 76 µm mylar. When the manipulator was inserted under the cover slip, a channel of variable width and 76 µm height was formed. The chromel (Hoskins Mfg Co, Detriot, Mich.) and

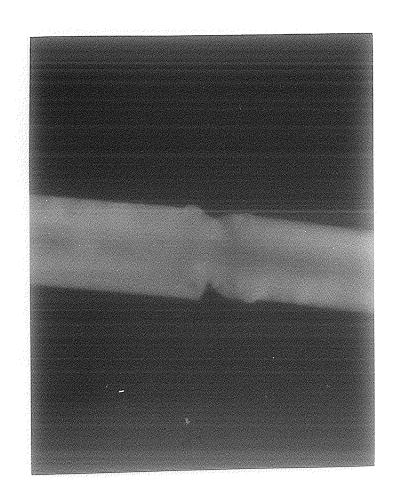


Figure 5. 76 µm thermocouple junction magnified 25x.

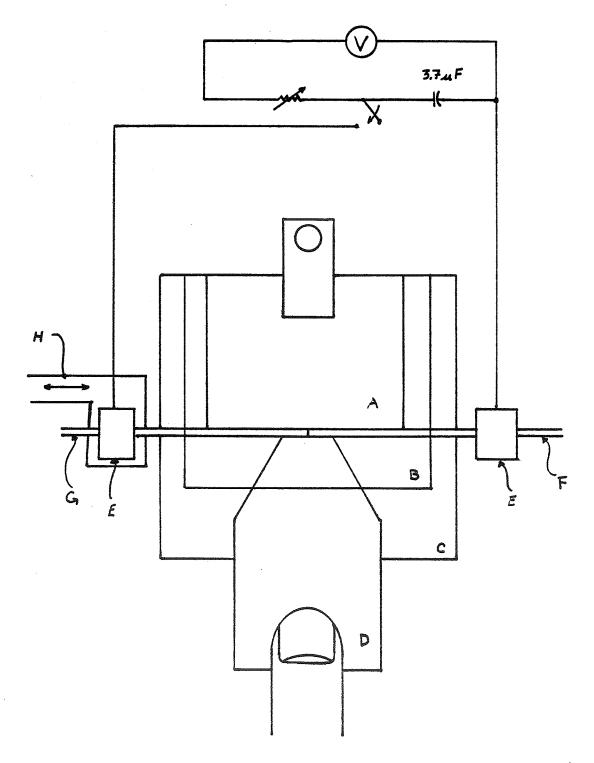


Figure 6. Schematic diagram of thermocouple manufacturing apparatus A) 76 µm mylar sheet, B) coverslip, C) glass slide, D) manipulator, E) clamping block, F) constantan wire, G) chromel wire, H) arm from worm gear

constantan (Cohen Wire Co., Mount Vernon, N.Y.) were into 15 cm lengths. The constantan wire was inserted under the coverslip from the left. When the tip to be welded was approximately at the midpoint of the channel, the wire clamped down next to the glass slide. This insured motional stability and a means of electrical connection. chromel wire was inserted from the right. It had already been clamped down onto an arm extending from a worm gear. This allowed the chromel wire to be moved in extremely increments to facilitate wire alignment. Thus one hand was used to control the manipulator while the other turned worm gear to move the chromel wire into position. After the chromel wire was butted against the constantan wire, the ends were aligned using the manipulator with the aid of a At this point, the microscope. junction resistance was measured to assure continuity. This capacitor was typically 9 to 10 ohms. To create the butt weld a 3.7 μ capacitor was discharged across the junction. This capacitor was in series with a variable resistor. This configuration allowed the capacitor to be charged to different voltages. A voltage of approximately 210 volts was first applied to weld the wires. If the junction appeared to have been formed properly, it was annealed by discharging the capacitor 20 times in 10 sec increments at a voltage of 120 volts. The junction resistance was once again measured and typically was about The thermocouple was removed from the alignment stage ohms.

and optically inspected from all sides using the microscope. Since the junction was difficult to locate with the unaided eye, a very small mark is made on both sides of the junction with blue machinists ink. To keep the ink bolus from catching on tissue, the bolus was smoothed with an alcohol saturated cotton swab. At this point, the thermocouple was ready for use.

Section 2.2.2 Thermocouple Response Acquisition

Two separate data acquisition systems were employed thermocouple response. monitor the The first system consisted of a Keithly microvoltmeter and an oscilloscope. The of the thermocouple was connected to output microvoltmeter. Using the microvoltmeter, the stability of response before heating could be examined and any d.c. bias nulled out. The signal was amplified by the Keithly with a gain of 10E05 and fed to the oscilloscope for visual inspection of the thermocouple response. This system was especially useful when beam plotting the transducer and permanent recordings of the response was not desired.

To obtain a permanent copy of the thermocouple response, a Hathaway oscillograph was used. The thermocouple signal was applied directly to the oscillograph which consisted of a sensitive galvanometer that deflects a light source that

was shining on light sensitive paper. The sensitivity of this galvanometer is such that an input thermocouple response of 60 uV (a 1°C change in temperature) will produce a 1.28 cm deflection on the oscillograph paper. This paper was darkroom processed after all the data had been recorded. The recordings produced look like the curve in figure 4. To obtain a value for the initial time rate of change of temperature (dT/dt), in °C/sec from this instrument, the following formula is used;

$$\frac{dT}{dt} = \left(\frac{dT}{dE}\right) * \left(\frac{di}{dx}\right) * \left(R_{TN} + R_{GAL}\right) * \left(SLOPE\right)$$
 (10)

where (dT/dE) is the thermocouple sensitivity (°C/uV), (di/dx) is the galvanometer sensitivity (0.545 uA/cm), Rgal is the resistance of the galvanometer(43.5 ohms), Rth is the resistance measured across the thermocouple leads in ohms, and Slope (cm/sec) is the slope of the thermocouple response curve at at 0.5 sec. All of the data discussed in Chapter 4 was obtained using the oscillograph.

<u>Sēction</u> 2.3 The <u>Irradiation</u> Apparatus

Having now detailed the physical characteristics and theory of the ultrasonic source and probe, attention is now turned to the exposure system. This system consists of a 36 liter Plexiglas tank placed on a 3 axis programmable milling machine base (Bridgeport). The specimen/thermocouple support

structure which was positioned opposite the transducer, was supported in the tank by a clamp extending from the back tank wall. The transducer was held in a fixed position relative to the tank and specimen structure by a clamp above the tank. A schematic of this exposure system is shown in figure 7.

The tank was filled with the ultrasonic propagation medium. Degassed distilled water was used in this experiment due to its extremely low acoustic attenuation, low cavitation probability, and limited ionization effects on thermocouple connections. In the bottom of the tank was a heating coil used to keep the water at a constant 37°C. At the end of the tank, behind the specimen structure was highly absorbing acoustic foam (SOAB) (B.F. Goodrich) to minimize standing wave effects. The position of the mill base was controlled by 3 separate Slo-Syn preset indexers, one for each axis of motion. Bidirectional increments of 0.001 in. to 9.999 in. could be selected using the indexer. The extent of motion recorded by feedback from 3 optical encoders which slide over 3 staionary transparent glass graticules. The digital information obtained from the encoders was displayed on separate X,Y, and Z Digi-Point digital readouts. The orientation of these coordinate axes, which will be of great importance in the following chapters, is also shown in fig 7.

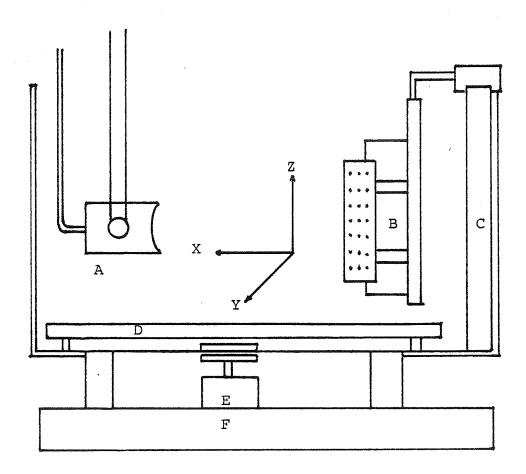


Figure 7. Schematic of exposure apparatus with coordinate system A) transducer
B) specimen structure, C) SOAB backing,
D) in-tank heater, E) magnetic stirrer,
F) 3 position milling base

A device which could uniformly move a thermocouple through the body was essential. Already existing was a mouse mount which secured the mouse for irradiation. decided to couple a suprastructure with this mount to facilitate thermocouple movement. There were a number of design aspects to be considered in the construction of suprastructure, such as: 1) the insertion and connection of the thermocouples require minimal thermocouple movement because of their fragility; 2) many thermocouple alignment positions to probe different tissues and individual mouse size variability; 3) and indicator of the amount of penetration of the thermocouple to correlate tissue probed with its thermocouple response curve; 4) bidirectional movement for replication of previous responses; 5) time and effort preparing the suprastructure should minimal; 6) ease of construction.

After several designs, consultations, and a few substitutions, the final design was arrived at. The complete system is shown is figure 8 and the detailed schematic of the suprastructure is shown in figure 9. The material selected for the building of the suprastructure was Plexiglas because of its availability, workability, and cost effectiveness. Stainless steel was used for the metal parts, due to its ability to resist the effects of saline solutions. The use

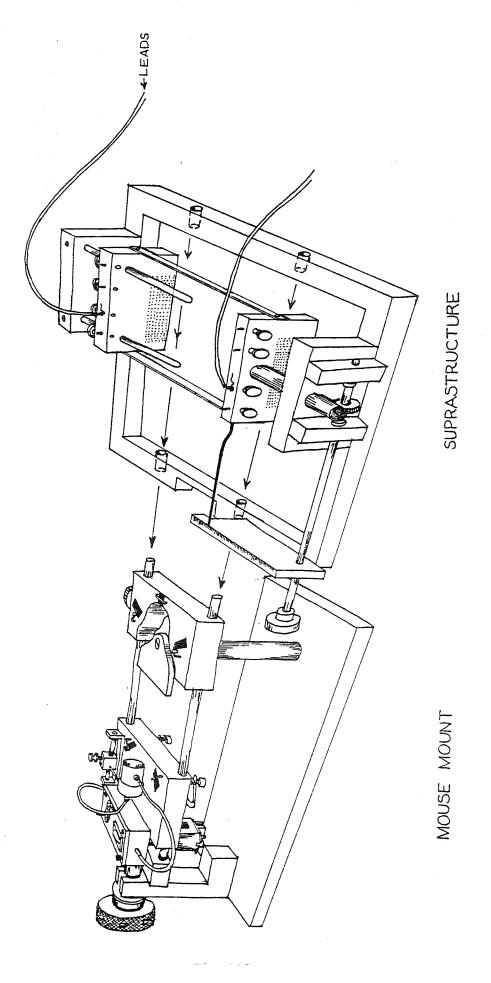


Figure 8. Complete specimen structure

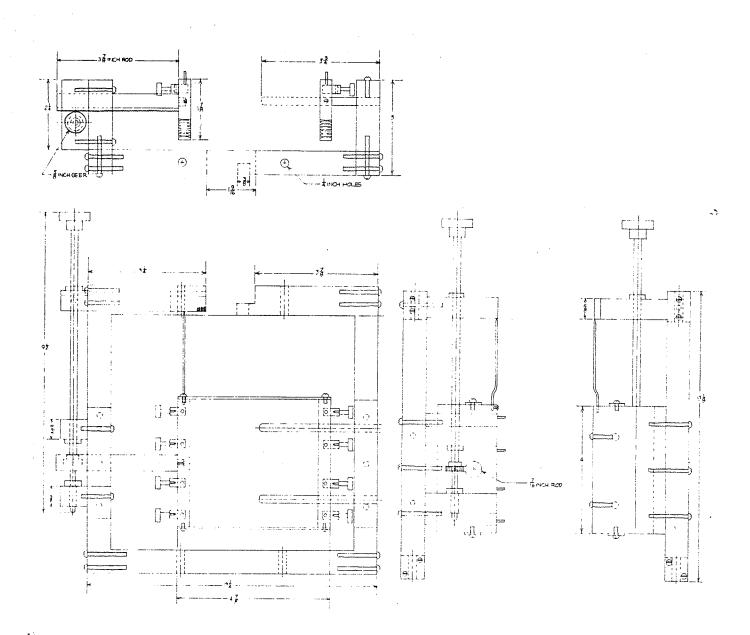


Figure 9. Detailed schematic of suprastructure

of the specimen/thermocouple structure in the experimental procedure is detailed in the following chapter.

CHAPTER 3

THE EXPERIMENTAL PROCEDURE

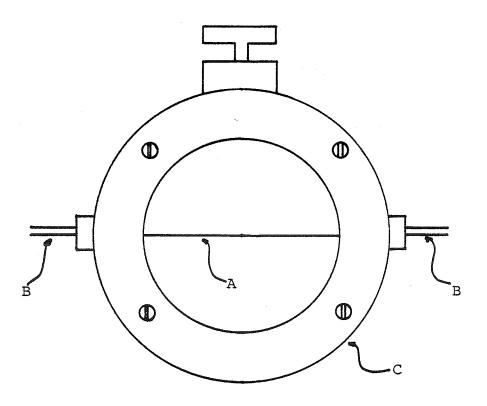
This chapter details the experimental procedures used to obtain the data necessary to compose an ultrasonic dosimetric model. This procedure was comprised of three distinct parts. First, the acoustic field was characterized without the specimen in place. Next, the specimen was prepared and irradiated. Finally, the tissue type of each irradiation site was determined through histological preparation of the specimen.

Section 3.1 Acoustic Field Characterization

The acoustic field produced by the transducer was considered to be a free field. A free field is characterized by plane progressive wave fronts of lateral extent where any standing wave effects are nonexistent. To construct a representation of this acoustic field, cross-sections of the field at various axial coordinates were obtained. These cross-sections are composed of point samples of some acoustic parameter taken at specific points in the plane. Since intensity was the field parameter to be used in the <u>fin</u> <u>vivo</u> absorption measurements (equation 9), it was logical to use intensity as the parameter to be sampled in the free field

characterization. To obtain these intensity samples, a transient thermoelectric probe was used for the precise calibration of the ultrasonic field. This technique has been used for many years in the Bioacoustics Research Lab and has been thoroughly described in the literature both for free field intensity determination (Dunn and Fry, 1957) and for $\underline{\text{in}}$ $\hat{f s}$ i $\hat{f t}$ ultrasonic absorption and intensity determination (Goss et al, 1979). For the free field detection of ultrasonic intensity, the probe consisted of a 0.5 mil lap soldered thermocouple imbedded in a fluid medium which was contained by two 76 um thick polyethylene acoustic windows and a steel housing. Two views of this probe are shown in figure 10. Silicon 710 (Dow Corning) was used as the embedding medium because its density and acoustic velocity are very similar to those of water and because its absorption coefficient is well known (Goss et al, 1977). Thus, the incident sound energy reflected was small and the temperature change could be related to intensity by using equation 9. This probe was phase insensitive device and no information about wave shapes could be determined using it.

The intensity at any point in the field can be calculated by using the temporal temperature response of the thermocouple probe, the known absorption coefficient and ρ C of Silicon 710, and the use of equation 9. If this absorption coefficient and ρ C are assumed always constant



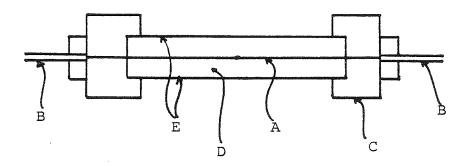


Figure 10. Schematic of calibrator
A) thermocouple wire, B) calibrator
leads, C) steel housing, D) imbedding
fluid, E) polyethylene acoustic windows

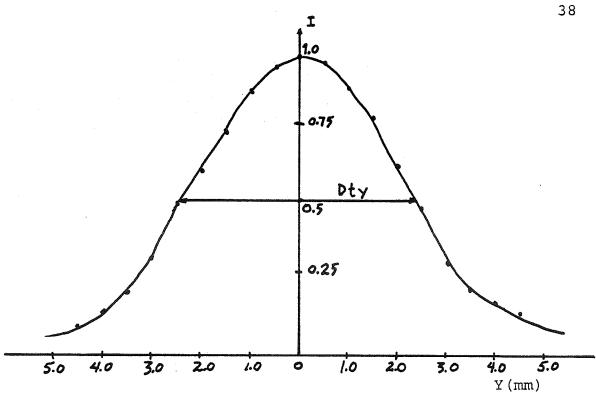
with respect to temperature and the time frame of this experiment, intensity and time rate of change of temperature are directly proportional. Furthermore, the magnitude of the net temperature change is also directly related to the time rate of change of temperature. An examination of the net temperature rise at the end of the acoustic disturbance will directly yield information about the magnitude of relative intensity. If the calibrator is moved on a line perpendicular to the focal axis, a relative intensity beam pattern may be plotted. The calibrator is then moved to a line that is also perpendicular to the focal axis, but orthogonal to the first line. Another beam plot is These two beam plots are combined to create a field cross-section. This process is repeated at succesive axial positions to obtain a three dimensional representation of the acoustic field.

The experimental set-up for the free field characterization was much the same as that in figure 7. place of the specimen structure, the calibrator was used. The tank was filled with 36 liters of degassed distilled water. syphoning process was used to reduce introduction of air into the solution. The tank heater was set for 37 °C. To aid in heating the solution, two other heaters were placed in the tank. A magnetic stirrer at bottom of the tank mixed the water. The transducer was positioned in the tank and its cable connector was attached to the driving network. The driving system was then switched on to allow proper warm-up. A 7.5 cm long pointer was placed on the front of the transducer housing. Since there is a 3 mm separation between the transducer face and the housing, the tip of this pointer is 7.8 cm away form the transducer This pointer tip indicates the region of maximal axial intensity (focal region) as determined by previous calibrations. After the water temperature had reached 37 C, the calibrator was placed in the tank parallel to transducer face so that the thermocouple wire was normal to the focal axis and parallel to the bottom of the tank. placement was done by visual inspection. The calibrator was moved foward until the thermocouple junction was as close possible to the transducer pointer tip. Care was taken to not puncture the acoustic window. This particular transducer/calibrator orientation was assigned the origin coordinate (0,0,0). Using the coordinate system of figure 7, the thermocouple was coincident with the Y axis, axis was the X axis, and the thermocouple junction was at the origin coordinate. All subsequent measurements were referenced to this point. In this calibration procedure probe was moved with respect to a stationary transducer. The calibrator was backed away from the transducer to allow the pointer to be removed. The calibrator was then retuned the origin.

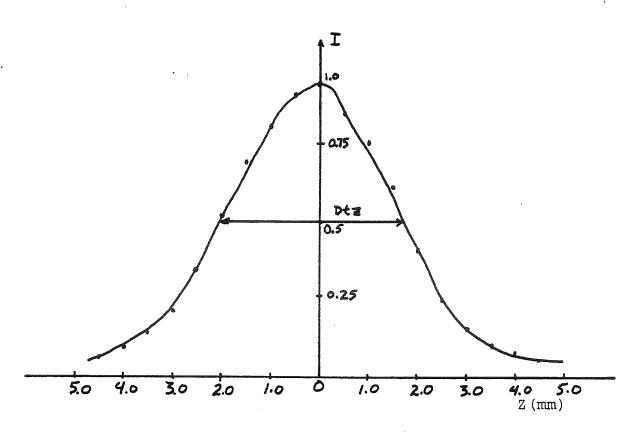
The variable capacitor was set so as to produce a maximal axial intensity of 2.44 W/cm**2. This intensity produced a suitable thermocouple output for beam plotting. The pulse length was chosen to be 0.3 seconds with a carrier frequency of 0.98 MHz. In this procedure, the thermocouple response was examined using the microvoltmeter/oscilloscope system.

The first field cross-section was obtained by moving the calibrator through the field in the YZ plane along the Y axis and then along the Z axis. The two relative intensity beam profiles obtained by this process are shown in figure 11. The actual shape of the focal region is slightly elliptical rather than circular. The transverse diameter in the Y direction is $5.0\,$ mm, while that in the Z direction is $0.0\,$ Dt= $0.0\,$ mm.

A series of field cross-sections were examined, using the same process as above, by incrementing X each time. The beam plots done in the diverging part of the field (-X) show a broadening of the intensity profile with a corresponding drop in the maximal axial intensity. The beam plots in the converging part of the field (+X) become very complicated and difficult to describe. These effects can be explained using the piston source theory discussed previously. The measured relative axial intensities are plotted in figure 12. It



A) Beam plot along Y axis



B) Beam plot along Z axis

Figure 11. Beam plots obtained along Y and Z axes

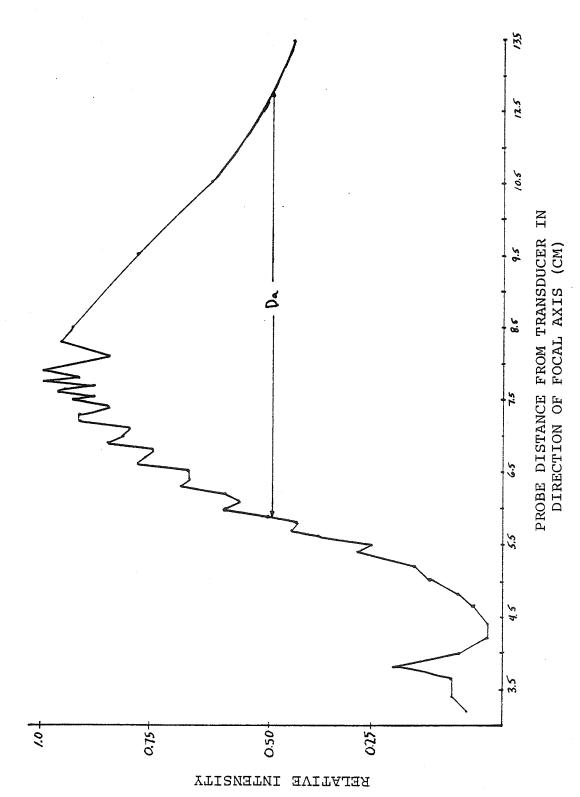


Figure 12. Relative intensity along focal axis

appears the true focal point of this transducer is indeed 7.8 cm from the face of the transducer.

The focal length determination from figure 12 was somewhat difficult due to the "ripple". The most likely cause of this was standing waves created by reflections from the SOAB tank backing. The true period of this standing wave was difficult to determine because the sampling increments used were integer multiples of $\frac{1}{3}$. The true standing wave pattern would be seen if sampling increments of one tenth the wavelength were used.

It was obvious that characterizing an acoustic field in this manner was a tedious process. Fortunately, frequent calibrations over the life of this transducer has shown the output to be quite stable with respect to time. Therefore this procedure does not have to be repeated often. The important field characteristics and dimensions have been determined. The interactions between this field and a specimen may now be investigated.

Section 3.2 Specimen Preparation and Irradiation

In this procedure the specimens to be irradiated were LAF1/J mice obtained from Jackson Labs in Bar Harbor, Maine. Each specimen was administered a diet consisting of only

water for the three days prior to the irradiation date. This starvation condition was necessary to minimize intestinal gas and fecal matter. If gas were to be present when the mouse was irradiated, the <u>in</u> <u>vivo</u> field would be drastically altered. The fecal matter is extremely heterogeneous, and thus it would be difficult to correlate the actual material at the irradiation site with the thermocouple response. Most of the body fat was also eliminated by this procedure. This is not desirable if body fat is to be a tissue of interest.

Prior to the experiment, the tissues to be irradiated were chosen. The thermocouple path through these tissues was limited by three constraints. They were: 1) the tissues to be irradiated had to spatially occur in a straight line; 2) the specimen structure allowed only a limited number of thermocouple wire orientations; 3) the irradiation procedure demanded that when the mouse was placed in the tank, the thermocouple wire should have been coincident with the Y axis (see Section 3.2.1). The second constraint can be alleviated through redesign of the specimen structure (Vaughn, 1980).

The experimental set-up is shown in figure 7. The preliminary experimental procedure was the same as in Section 3.1. As the water was coming up to temperature, the mouse was prepared. The starved mouse was placed in a glass chamber containing Metofane (Pittman-Moore Washington

Crossing, NJ) or chloroform. The mouse was removed when there was pedal response. The anesthetic dosage administered was lethal. This was a necessary part of the experimental procedure for a number of reasons. During irradiation the thermocouple response would be adversely effected by the respiration and other body movements of an anesthesized mouse. Secondly, the mouse orientation in the specimen structure in a number of irradiations would drown the mouse. Finally, the specimen would have to be sacrificed immediately after the irradiation in any event, to facilitate the histological preparation. Fortunately, it has been shown that in the three hours required to perform the irradiation procedure, tissue characteristics do not change drastically (Bamber et al, 1977). The results obtained from the specimen tissues should yield an accurate representation of fresh, if not living tissues.

After anesthetization, the mouse was prepared for placement in the specimen structure. First, all the fur was removed from the area between the fore limbs and the hind limbs on all sides of the body. Electric clippers were used to remove most of the fur, followed by an application of commercial cream depilatory to remove all remaining stubble. After 15 minutes, the depilatory was rinsed off and the mouse was dipped in a water/shampoo solution. This left a slick film on the mouse body which discourged bubble adherence.

To place the thermocouple wire in the tissues of interest, a 22 gauge 1.5 inch hypodermic needle was inserted into the mouse along the thermocouple path decided upon previously. The thermoucouple was then carefully threaded through the hypodermic needle, and was held while the needle and was removed from the body. This process left the thermocouple wire in the body on a path through the tissues of interest.

The mouse was then secured in the support structure. The orientation of the mouse depended on which side the acoustic energy was to be incident. In subjects numbered and 51 to 52, the mouse was placed head up, with the ventral side facing the transducer. In subjects 33 to 50, the mouse was placed head down, with the dorsal side facing the transducer. After the mouse was properly secured, thermocouple ends were threaded through the support holes. These two holes were chosen so that the thermocouple wire would slide through the mouse parallel to the bottom and back the tank. In otherwords, there should have been no deviation of the thermocouple path in the X or Z coordinates. The thermocouple was pulled through the mouse until junction appeared at the skin perforation at the mouse's right side. The blue ink marks were utilized to identify the exact location of the junction. The thermocouple wire ends were secured in the brass external lead connectors. This was

in such a manner as to tauten the wire but to leave the junction position undisturbed. The external thermocouple leads were attached to the brass connectors. The resistance across these leads was measured. This resistance value used for Rth in equation 10. The tiny perforations where the thermocouple entered and exited the skin were stained dark green with an astringent dye. The midway skin point between these two perforations was also stained. These markings were used as an aid in the histological preparation. The thermocouple junction was then moved from its present location through the mouse to the other skin perforation on the left side. The distance the thermocouple junction traveled from the right side skin perforation to the left side skin perforation was measured using a vernier/pointer configuration. This path length was recorded for use as a comparative standard in the irradiation procedure. thermocouple was then returned to the skin perforation on the right side for later alignment in the tank. Figure 13 shows a specimen prepared for irradiation.

Section 3.2.1 The Coordinate System and Alignment Procedure

The specimen/thermocouple structure was placed in the tank opposite the transducer. The transducer was brought up to the mouse so that the pointer tip was touching the thermocouple junction at the right side perforation. This

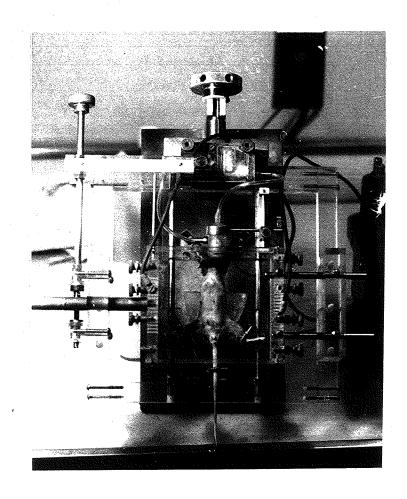


Figure 13. Prepared specimen secured for irradiation

specimen/transducer/thermocouple junction orientation was assigned the origin coordinate of (0,0,0) as seen in figure 14. Recall this tip indicated the location of the focal region. In this procedure the location of the focal region was used to report the position of the transducer. This was done in order to directly obtain the coordinates of the irradiation site within the specimen. Therefore, the relationship between the "in situ intensity" and thermocouple response could be examined in the same manner as Section 3.1.

the origin the thermocouple was coincident with the Y axis, the focal axis was the X axis and the thermocouple junction was at the intersection. The specimen stationary in space with respect to the coordinate system. The thermocouple junction moved only along the Y axis, and the junction positions were recorded in only one coordinate (Y). The focal region was moved in three dimensions and thus its coordinates were recorded as (X,Y,Z). Notice that both the thermocouple junction and focal region positions were recorded using the same coordinate system. Therefore, by simple inspection of the reported coordinate positions, the precise orientation between the thermocouple and focal region was known.

The length of the thermocouple path was measured again by moving the transducer to the left perforation. This new

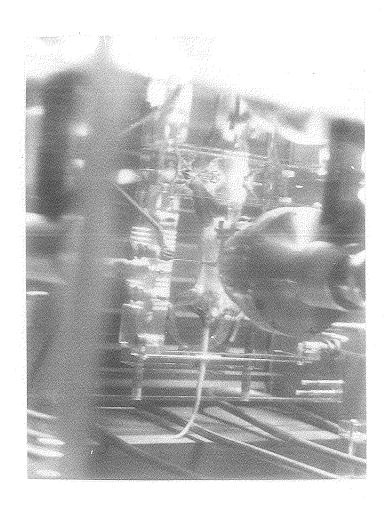


Figure 14. Specimen fixed in tank with transducer focal tip and thermocouple junction at position assigned (0,0,0)

transducer position should have read (0,+Y1,0) if the structure is properly aligned. The value of Yl should have equalled the path length previously measured outside the If the transducer were moved in the X or Z directions to place the focal tip on the left perforation, the specimen structure was realigned. Upon realignment the origin coordinate was reset. After any structural realignment was completed, the transducer tip was moved to the center marking. The transducer position then read (+X,+Y1/2,0). The X value was an indication of the thermocouple depth under the skin surface at this point. The transducer was backed away from the specimen and the pointer removed. transducer was then moved back so that the focal region was again at the origin.

To insure that the thermocouple junction was aligned properly in the focal region, beam plots like those described in Section 3.1 were done. The origin though, was not a satisfactory thermocouple location to do a beam plot, as the skin/water interface often yielded some unusual results. The thermocouple and transducer were both moved 4.0 mm in the +Y direction to a different irradiation site. In this position adequate <u>in</u> <u>vivo</u> beam plots were usually obtained. The criteria used for centering the junction in the focal region was attainment of the maximal relative intensity in both the Y and Z directions. It was intended that this location of

maximal intensity would coincide with the alignment obtained with the pointer centering process, although this was not always the case. Upon inspection of the two beam plots done in the Y and Z directions, it occasionally was found that the thermocouple was not in the locus of maximal intensity. The transducer position was readjusted so the maximal relative intensity in both the Y and Z directions was obtained. The thermocouple and transducer were moved back 4.0 mm in the -Y direction. If the transducer had to be realigned, the origin coordinate was reset.

A number of important assumptions were made in this alignment process and were used throughout the experiment. They were: 1) the movements of the thermocouple and transducer were always in the increments desired; 2) the scale for the thermocouple movement and transducer movement readout were accurate; 3) the thermocouple did not slip in the specimen structure, nor did the encoders on the mill base; 4) the ultrasonic beam was not refracted at the skin/water interface which would redirect the maximal intensity off the focal axis; 5) the focal region was always centered on the focal axis.

Section 3.2.2 Irradiation Algorithms

Once the thermocouple and transducer were properly

aligned, the irradiations were begun. There were a number of possible irradiation that could have been algorithms implemented. The transducer and thermocouple could be moved in a one-to-one correspondence in the Y direction. would keep the focal region over the thermocouple at irradiation site. Another approach would be to obtain beam plots at each thermocouple position by moving the transducer in its YZ plane. This algorithm might yield more information concerning beam permutations. The thermocouple might be incremented through a stationary field (a beam plot in the Y direction) , the transducer is then moved and the process is repeated. The algorithm used in this experiment was former. One-to-one correspondence yielded data which was easy to obtain and analyze. The second and third algorithms described above were used on a small number of specimens, but only to supplement the one-to-one data. It must be realized that the second and third algorithms are tedious time consuming processes, but will be easier to implement with a more automated precedure (Vaughn, 1980).

The implementation of the one-to-one algorithm was straightfoward. Beginning with the origin coordinate, the site was irradiated with a one second ultrasonic pulse of sufficient intensity to produce a legible, permanent copy of the thermocouple response output on an oscillograph. The thermocouple and transducer were simultaneously moved one

millimeter in the +Y direction. This new site was then irradiated using the same exposure conditions as the previous location. This process was continued until the junction exited the left side perforation. At each location the following data were recorded; 1) thermocouple position (Y), 2) focal region position (X,Y,Z); 3) pulse length; 4) value of the variable capacitor Cs; 5) observations of unusual thermocouple responses.

Occasionally, the number of thermocouple positions which were taken in one millimeter increments, yielded a distance which exceeded the thermocouple path length measured twice The source of this error stemmed thermocouple adhering to the tissues. Although the junction was designed for smooth passage, it still occasionally snagged on the tissues. This was especially true locations where the thermocouple entered or exited the body. The junction would snag on the skin and pull it along for 2-3 Obviously, this changed the tissue orientation from normal position, a situation which more than likely happened internally to varying degrees as well. To alleviate this problem, the junction was moved back and forth rapidly to dislodge the tissue and return it to its original location. This may have resulted in slight tissue damage, although it was felt such tissue damage did not effect the results.

After the last thermocouple response was recorded, the oscillograph paper was developed. If legible traces were obtained, the mouse was prepared for the histological sectioning process.

Section 3.3 Histological Preparation of the Specimen

To fully make use of the thermocouple response data obtained by the procedure in Section 3.2, the tissues comprising the thermocouple path must by identified. A method to precisely and accurately describe the path of the thermocouple was devised (Wyneken and O'Brien, 1980). This freeze-sectioning technique allows for both gross and semimicroscopic anatomical information to be obtained through thick sectioning.

Following the irradiation procedure, the specimen still contained in the structure was submerged in an ice bath. In order to maintain temperatures suitable for rapid freezing at approximately -10 C, 4-5 liters of a saturated salt solution (290 gm/l) were mixed with 3600 grams of crushed ice. The specimen structure was maintained in the same orientation in the ice bath as it was during the irradiation process. The specimen could be removed from the ice bath without distorting the internal organ locations after 10 minutes. The mouse was removed from the experimental holder, taking

extra care not to disrupt the thermocouple wires, wrapped in plastic and placed in a $-10\,^{\circ}\text{C}$ freezer.

To obtain a section of the specimen, a specially constructed guillotine, cooled by a dry ice-90% alcohol bath, was used to cut rapidly through the animal along a dorsoventral plane (figure 15). Cuts were made on either side of the thermocouple wire yielding a cross-section piece approximately 8 mm thick. This section was packed in dry ice.

After cooling the freezing stage of a sliding arm microtome with a dry ice-90% alcohol bath, a paper collar was fitted around the 8 mm thick section on the stage and an ice base was built inside of the collar. This technique is similar to that of mounting a specimen on an object disk cryostat with an embedding medium which in this case is water, and the collar is required to contain the water prior to freezing. The cross-section piece was placed with its flatter surface down inside the collar. Care was taken note which surfaces were cranial or caudal. Small amounts of water were added around the base and the entire collar was then packed in crushed dry ice, freezing the 8 mm specimen to the base. This configuration is shown in figure Prior to sectioning the specimen, the paper collar removed.

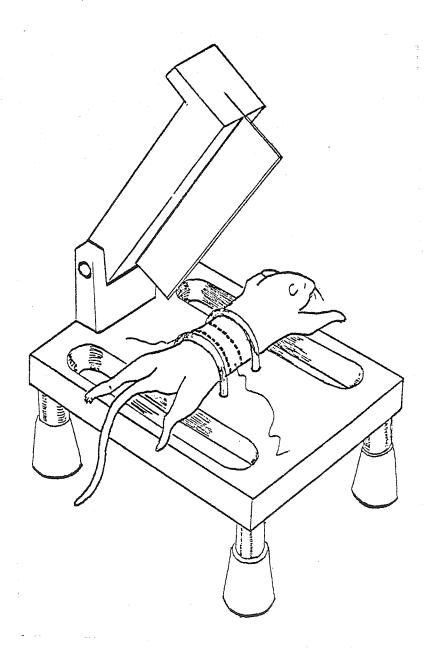


Figure 15. Specimen sectioning guillotine

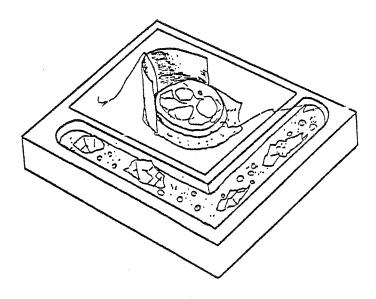


Figure 16. Specimen section freezing stage

The plane of section was determined using the entrance and exit skin perforations caused by the thermocouple wire and a third point determined by the position of the animal relative to the ultrasonic beam axis. These three spots had been stained dark green during the irradiation preparations to mark the plane of section.

Relatively thick serial sections (80 µm -120 µm) were made and mounted on gelatin coated slides. If these sections were to be saved, they were frozen or fixed using a treatment of absolute alcohol followed by flowing 0.75% celluloden solution in ether or alcohol, then 1 minute in 80% alcohol. Sections were removed until just before the level of the thermocouple was reached at which point the two ends of the thermocouple wire were attached to a six volt battery for approximately one minute to create a sear. The sear mark leaves a permanent visable record of the path of the thermocouple wire. The thermocouple wire was then removed and sectioning continued until the level of the sear was reached.

A scaled representation of the entire section at the level of the thermocouple was drawn with the aid of a Fowler Scale Magnifier equipped with a 10 mm grid reticule. All organs and tissues in this drawing were identified and labeled. The tissue type at each irradiation site was

recorded, as well as the distance between the site and the irradiated skin surface in the direction of the focal axis. Furthermore, each interposing tissue type between the skin surface and irradiation site was identified and dimensioned. These interposing tissues were recorded in order of their spatial occurence along the focal axis for each irradiation site. Thus, the acoustic propagation path to each irradiation site was characterized. This information combined with the dosimetric data obtained in Section 3.2 can yield the spatial distribution of ultrasonic energy within the specimen.

CHAPTER 4

RESULTS AND OBSERVATIONS

In this chapter the data obtained from the irradiation histological procedures described in Chapter 3 are collated and examined. The reasons for changes of the specimen orientation in the irradiation procedure over course of the experiment are explained. A model to calculate the acoustic intensity at each irradiation site within the specimen is proposed. Using this calculated "in situ intensity" value, the absorption coefficient of the tissue at each site is determined and entered in to a dosimetric file with pertinent irradiation and histological data. irradiation sites in this dosimetric file which have the same tissue type are grouped together. The absorption coefficient results for each tissue type are compared to literature values. Observations are made on this absorption coefficient data and improvements to the model are suggested.

<u>Section 4.1 Discussion of the Experimental Procedure</u>

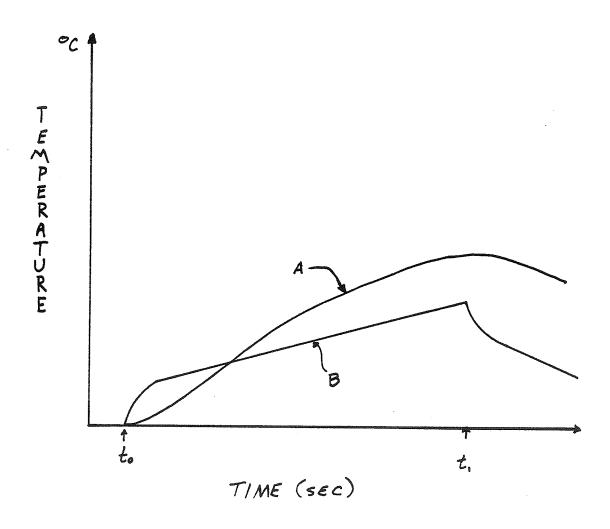
During this experiment, 53 specimens were irradiated. The results from a number of these specimens were not utilized in the analysis because: 1) the thermocouple responses were not legible on the oscillograph paper; 2)the

histological data were not obtained due to improper freezing or sectioning techniques; 3)a one-to-one correspondence did not exist between the number of irradiation sites exposed the irradiation procedure and the number of sites determined histologically. This last reason was discussed in Section 3.2 and concerned the snagging of the thermocouple junction in tissues. If the thermocouple did pull tissues out of their normal orientation, then more irradiations sites would be recorded than expected. If the tissues then pulled returned to their original orientation before the and specimen was frozen, then the histologist would find fewer irradiation sites than recorded in the irradiation procedure. Therefore, it would be subject to considerable error to correlate the thermocouple responses with the histological data.

valid specimen data were classified into two categories. These categories were partitioned by the orientation of the mouse in the support structure during irradiation. As it turned out, there was no need to analyze the data separately by this classification. Although it is convenient to discuss the experimental procedures in this manner. The first category consisted of the specimens (numbered 17 to 32, 52 and 53) which were irradiated head up, ventral side facing the transducer. Within this group two different thermocouple paths were utilized. In specimens 17

to 20, 25, 29 to 32, 52 and 53, the thermocouple wire was inserted just under the abdominal skin surface. In specimens 21 to 28 the thermocouple wire was inserted through the intestinal and gut area. It was found in these specimens that when the thermocouple junction was in gas or fecal matter an atypical thermocouple response was produced. In figure 17 a thermocouple response received from an area in specimen 25 known to contain gas and fecal matter is contrasted to a typical thermocouple response. Notice that it is nearly impossible to define a linear region and obtain a satisfactory value for dT/dt from trace A.

Since the contents of the intestinal tract markedly effects acoustic wave propagation, the irradiation procedure changed so as to avoid irradiating the intestinal tissues. Also, it was decided to probe a well characterized tissue. Kidney was chosen because of its accessability and known absorption and attenuation coefficients (Goss et al, 1979). To avoid having the sound beam pass through intestinal organs before striking the kidneys, the mice were irradiated from the back. To accomplish this using the existing specimen structure, the mouse was placed head down, dorsal side facing the transducer. Specimens 33 to 44, 47, 50, and 51 were irradiated in this fashion and comprise the second category.



Figure]7. Comparison of an intestinal thermocouple response (A) to a typical response (B)

Section 4.2 The Irradiation Files

Because of the large amount of data and to facilitate compilation and analysis, all irradiation histological data were stored in Interdata an 7/32 minicomputer(Perkin-Elmer Oceanside, NJ). The experimental irradiation data file (DOSEXPR.DTA) was arranged to accomodate data from very complex dosimetric experimental procedures. Although the file has provisions for accepting data from specimens probed with multiple thermocouple wires, only one thermocouple wire was used in this study. was ordered by specimen number beginning with specimen 17. The data entered in this file for each irradiation site were: 1) specimen number; 2) thermocouple probe number; 3) coordinates of the focal region (inches); 4) shot number; 5) thermocouple junction position as recorded with the vernier scale on the specimen structure (mm); 6) time rate of change of temperature measured from the oscillograph (cm/sec); 7) net temperature rise at the end of the acoustic disturbance obtained from the oscillograph (cm); 8) value of the variable capacitor Cs.

Some of these data were not entered in its most useful form for dosimetric analysis. A fortran program (DOSEXPR.FTN) was written which: 1) converted the focal region coordinates from inches to millimeters 2) normalized

the thermocouple position to a Y axis value with respect to the coordinate system (mm); 3) converted the dT/dt and net temperature rise values to °C/sec and °C respectively using equation 10; 4) calculated the acoustic intensity produced by the transducer using the Cs value and equations 4 and 5; 5) create a new file (DOSEXPC.DTA) to store this converted data and items 1, 2, and 4 from DOSEXPR.DTA. A final DOSEXPC.DTA file specimen entry is shown in Appendix A.

Section 4.3 The Histological Data File

The histological data file (DOSHIS.DTA) was set-up in the same form as the DOSEXPC.DTA file to simplify the combination of the two files. The information recorded irradiation site in a specimen were: 1) specimen number; 2)thermocouple 3)coordinate number; thermocouple junction position with respect to the specimen (mm); 4) thermocouple depth below the skin surface in the direction of the focal axis (mm); 5) tissue identification code; 6) interposing tissues. This interposing tissue entry consisted of a 4X2 array. Each interposing tissue and its depth were recorded in order of their occurence in relation irradiated skin surface. Occasionally, there were more than four interposing tissues between the irradiation site and the skin surface. In this case, 999 was placed in the specimen number column of the next line to denote a

continuation of the line before. The remaining interposing tissues and their thicknesses were entered on this line.

To facilitate tissue type data entry, a three letter was devised for each tissue encountered experiment. This coding system is listed in table 1. Provisions were made to describe different portions of an organ. In some cases, the first 2 letters of the code identified the organ and the third letter denoted a particular part of that organ. For example, in the case of kidney there were codes for the cortex (KIC), medulla (KIM), surface (KIS), and kidney in general (KIG). KIG was when particular part of the kidney could not be distinguished. Furthermore, the coding system was extended to cover the case of when the thermocouple junction was located in an interface between two different tissues. The letter of this interface code was the first letter of the tissue code through which the thermocouple junction had just passed. The second letter of the code was a J to denote tissue/tissue junction. The third letter of the interface code was the first letter of the tissue code through which the thermocouple junction will pass next. For example, if the thermocouple junction was located in the kidney/liver interface, the tissue code would be KJV. Using orientation of the scaled drawing (see Section 3.3) done from the histological specimen section, this code reads from

Table 1: TISSUE CODES FOR THE HISTOLOGY FILE DOSHIS.DTA

ADP-FAT BLO-BLOOD BVV-BLOOD VESSEL CNT-CONNECTIVE TISSUE CRU-FECAL MATERIAL DER-SKIN FAS~FASCIA GUG-STOMACH GENERAL GUL-STOMACH LUMEN GUS-STOMACH SURFACE KIC-KIDNEY CORTEX KIG-KIDNEY GENERAL KIM-KIDNEY MEDULLA KIS-KIDNEY SURFACE LCG-CAECUM GENERAL LCL-CAECUM LUMEN LCS-CAECUM SURFACE LIG-LARGE INTESTINE GENERAL LIL-LARGE INTESTINE LUMEN LIS-LARGE INTESTINE SURFACE MMA-ABDOMINAL MUSCLE/SUBCUTANEOUS MUSCLE MMB-BACK MUSCLE NSP-SPINAL COLUMN PLC-SPLEEN CORTEX PLG-SPLEEN GENERAL PLS-SPLEEN SURFACE SIG-SMALL INTESTINE GENERAL SIL-SMALL INTESTINE LUMEN SIS-SMALL INTESTINE SURFACE VEC-LIVER INTERIOR VEG-LIVER GENERAL VES-LIVER SURFACE WAT-WATER

FOR INTERFACES, USE J TRANSPOSED BETWEEN THE FIRST LETTER OF THE TISSUE CODES, GOING FROM LEFT TO RIGHT.

EXAMPLE; LIVER/KIDNEY INTERFACE IS VJK.

to right. A specimen entry from the DOSHIS.DTA file is presented in Appendix B.

<u>Section 4.3 The Dosimetric Model</u>

To determine the absorption coefficient of a tissue using a thermocouple probe, the acoustic intensity at that location must be known. In this experiment only the free field intensity is known. An iterative model was developed to calculate the "in situ intensity" knowing this free field intensity produced by the transducer and the interposing tissues.

The basis of this model is the theory of sound transmission through multiple layered media. This transmission theory is explained in a number of texts (Kinsler and Frey, 1962). In this case the layered media are the interposing tissues. To use the multiple layered media theory to model the interposing tissues a number of assumptions were made. They are: 1) all tissue boundries were normal to the direction of sound propagation; 2) all wave fronts were plane progressive; 3) there was no mode conversion; 4) the attenuation coefficient and characteristic acoustic impedance were spatially constant for each layer; 5) there were no standing wave effects; 6) all wave phenomena were of infinitesimal amplitude and linear. Once these

assumptions were made equations to calculate the "in situ intensity" could be formulated from the multiple tissue layer model.

As a plane progressive acoustic wave propagates through a medium which is lossy (tissue), the intensity decreases exponentially. The intensity at any point within the medium Ix, is given by;

$$I_{x} = I_{o} \exp(-2Ax) \tag{11}$$

where Io is the original incident intensity (W/cm**2), A is the attenuation coefficient of the medium (Np/cm), and X is the depth of acoustic penetration (cm). When this acoustic wave impinges upon a boundary with a contiguous second medium, a reflected wave is generated in the first medium, and a transmitted wave is generated in the second medium. The fractional energy transmitted into the second medium is given by the sound power transmission coefficient **C*;

$$dt = \frac{4Z_1 * Z_2}{(Z_1 + Z_2)^2}$$
 (12)

where Z1 and Z2 are the characteristic acoustic impedances of mediums 1 and 2 respectively. To find the intensity at an \underline{in} \underline{vivo} point P within the tissue of interest, where there are intervening layers of tissue between the point P and the skin

surface, an iterative process using equations 11 and 12 may be used. This process is expressed numerically as;

$$I_{5} = I_{0} \exp\left(-\frac{k}{2} 2 A_{n} \times n\right) * \prod_{n=1}^{K} dt_{n}$$
 (13)

where Is is the " $i\bar{n}$ $sit\bar{u}$ intensity" (W/cm**2), Io is the value of the free field incident intensity (W/cm**2), and K is the number of tissue boundries. In expressing equation 13 in this manner, it is assumed the degassed water is lossless.

To apply equation 13 to the data files, the attenuation coefficient and characteristic acoustic impedance of each interposing tissue must be known. A compilation of the literature values of these parameters for each interposing tissue is shown in Table 2. The values which are asterisked denote them to be approximated. These approximations were made from knowledge of the tissue constituents. Once the "in situ intensity" is calculated, the irradiated tissue absorption coefficient may be determined using equation 9. The value used for C in this equation is 3.77 J/ *Ccm**3 (Goss et al,1977).

Section 4.4 The Dosimetric File

To calculate the absorption coefficient at each irradiation site a fortran program (DOS.FTN) was written

TABLE 2: COMPILATION OF CHARACTERISTIC TISSUE PARAMETERS TO BE USED IN EQUATIONS 11 AND 13 (GOSS AND JOHNSTON, 1978)

TISSUE <u>CŌĐĒ</u>	TISSUE TYPE	VELOCITY (<u>m</u> / <u>s</u>)	DENSITY (<u>ā</u> / <u>ām</u>)	IMPEDANCE (<u>x10⁶ Râŷlŝ</u>)	ATTENUATION COEFFICIENT Np/cm
ADP	Fat	1467	0.95	1.39	0.07
BLO	Blood	1566	1.042	163	0.02
BVV	Blood Vessel	1590		1.82	0.17
CNT	Connective Tissue	1625	1.08	1.75	0.35
C RU	Fecal Material			1.40*	0.40
DER	Skin	1720	1.1	1.89	0.24
FAS	Fascia	1600	1.05	1.60*	0.18*
G U~	Stomach			1.60*	0.07*
KΙ	Kidney	1566	1.055	1.64	0.12
LC-	Caecum			1.60*	0.07*
LI~	Large Intestine			1.60	0.07*
MMA	Abdominal Muscle	1566	1.058	1.66	0.15
MMB	Back Muscle	1566	1.058	1.66	0.15
NSP	Spinal Bone Spinal Cord	3445 1615	1.82	7.8 1.66	1.6 0.25
PL≈	Spleen	1567	1.059	1.67	0.009
SIme	Small Intestine			1.60*	0.07*
VE	Liver	1578	1.055	1.66	0.10
WAT	Water	1500	1.00	1.5	0.0

which combined the DOSEXPC.DTA and DOSHIS.DTA data files. This program calculated the " $\underline{\hat{n}}$ $\underline{\hat{s}\hat{t}}\underline{\hat{u}}$ intensity" and absorption coefficient using equations 8, 12 and 13, and the experimental data for each irradiation site. A dosimetric file (DOS.DTA) was created which contained the data believed to be the most useful for the dosimetric analysis employed herein. The entries in this file were: 1) the specimen number; 2) thermocouple number; 3) thermocouple junction coordinates with respect to the specimen (mm); 4) tissue code; 5) thermocouple depth below the skin surface in the direction of the focal axis (mm); 6) time rate of change of temperature (${}^{\circ}$ C/sec); 7) " $\hat{\underline{n}}$ $\hat{\underline{s}}\hat{\underline{t}}\hat{\underline{u}}$ intensity ($\mathbb{W}/\mathbb{cm}**2$) 8) tissue absorption coefficient (Np/cm) at 1 MHz. A specimen entry from the DOS.DTA file is shown in Appendix C.

<u>Section</u> 4.5 <u>Data Analysis</u>

The data contained in the DOS.DTA file can be analyzed on two levels. On the tissue level, all irradiation sites of the same tissue type can be grouped together and their calculated absorption coefficients () compared. It was felt that the assumptions upon which the dosimetric model was based could more easily be substantiated by examining the sat each irradiation site. On the specimen level, the relationships between irradiation sites within a particular

specimen can be examined simultaneously. Useful information can be obtained by examining data on this level, especially in determining the energy absorbed by the specimen and the interaction of the acoustic field within the specimen. It is the intent of this section to examine the data on both levels and suggest improvements to the dosimetric model.

facilitate the examination of the s on the tissue level, the DOS.DTA file was partitioned by tissue type. The results of this partitioning process are shown in tabular form in Appendix D. Presented on the first page of appendix is the description of each of the columns in this table. Notice that not only are the tissues grouped with those of the same tissue code, but also with any pertinent tissue/tissue interfaces. For example, the skin group not only contains the irradiation sites which were skin (DER), but also contains the skin/muscle interfaces (MJD,DJM) skin/water interfaces (WJD, DJW). These interfaces were included to acount for histological irradiation site determination inaccuracies of ± 0.5 mm. It was possible that the thermocouple junction may have been in the tissue of interest rather than in the interface.

In Appendix D, two tissues types, kidney and muscle, have large data bases. These two tissue groups were more closely examined to assess the practicability of the

dosimetric model. The mean and standard deviation were calculated for the absorption coefficients of the two tissue groups. Recall that all the absorption coefficients calculated in this study were obtained using a 76 um thermocouple wire. When irradiated, this wire size will produce viscous heating effects which contribute to heating response throughout the duration of the acoustic disturbance. Therefore, the measured time rate of change of temperature will be higher than the rate of change without the wire in place. Goss et al (1977) examined the errors associated with the viscous heating contribution at 0.5 MHz and presented a plot of theoretical error vs. absorption coefficient. The 76 µm thermocouple line on this plot was used as a approximation of the error produced by the viscous heating at 1 MHz. Viscous heating is less dependent upon frequency than is the bulk heating due to absorption resulting in smaller errors due to this effect at higher frequencies. Thus, the error approximations taken from the Goss paper might be termed "worst case". Taking this viscous heating error into account reduces the mean absorption coefficient values by about 8%. The mean, standard deviation, "corrected" mean, and literature values for kidney and muscle are presented in Table 3. The kidney literature absorptin coefficient was measured using a 13 μm thermocouple junction (Goss et al, 1979). A literature value for the muscle absorption coefficient did not exist at the writing of

Literature 0.030 Value 0.045 z = = nuscle "Corrected" 0.0423 0.0520 0.0475 0.0355 0.0283 0.4342 Mean and kidney Standard Deviation 0.0274 0.0490 0.0213 0.0228 0.0174 0.4248 uo calculations 0.0453 0.0550 0.0510 0.0380 0.0315 0.4342 Mean absorption coefficient values. # of Irradiation Summary of Sites 67 73 27 54 17 1 1 Table 3: Tissue (under Code spine) KIG KIS KIC KIM 11111 HMB

this thesis. Goss et al (1979) reported the existance of a threefold difference between the attenuation and absorption coefficients for most soft tissues. Using this relationship and a value for the attenuation coefficient of 0.15 NP/cm (Table 2), the absorption coefficient for muscle is assumed to be 0.045 Np/cm. This is the literature value presented in Table 3.

The kidney data was inspected on the tissue level to detect flaws in the dosimetric model. In Table 3 it is shown that the three kidney "corrected" mean values are within 15% each other, while the standard deviation in each case is approximately 50%. Coupled with the fact that the irradiation sites represented by this kidney data located in many different orientations within the specimen (as noted by widely varying Y coordinate in Appendix D), the consistency of the "corrected" mean absorption coefficients suggest that the dosimetric model is substantiated in the case of kidney irradiation sites.

Attention is turned to the muscle absorption coefficient values. It is shown in Table 3 that the abdominal muscle and first back muscle entry "corrected" mean values are fairly close. The second back muscle entry is composed of back muscle irradiation sites which were under the spine. These sites are denoted by asterisks in Appendix D. The

"corrected" mean for this entry is an order of magnitude larger than the other back muscle "corrected" mean values. This absorption coefficient discrepancy arises from a dosimetric model inadequacy. To determine the nature of this inadequacy, the back muscle data were treated on the specimen level. In otherwords, relationships between the back muscle irradiation sites were inspected simultaneously to determine how the acoustic field interacted within the specimen.

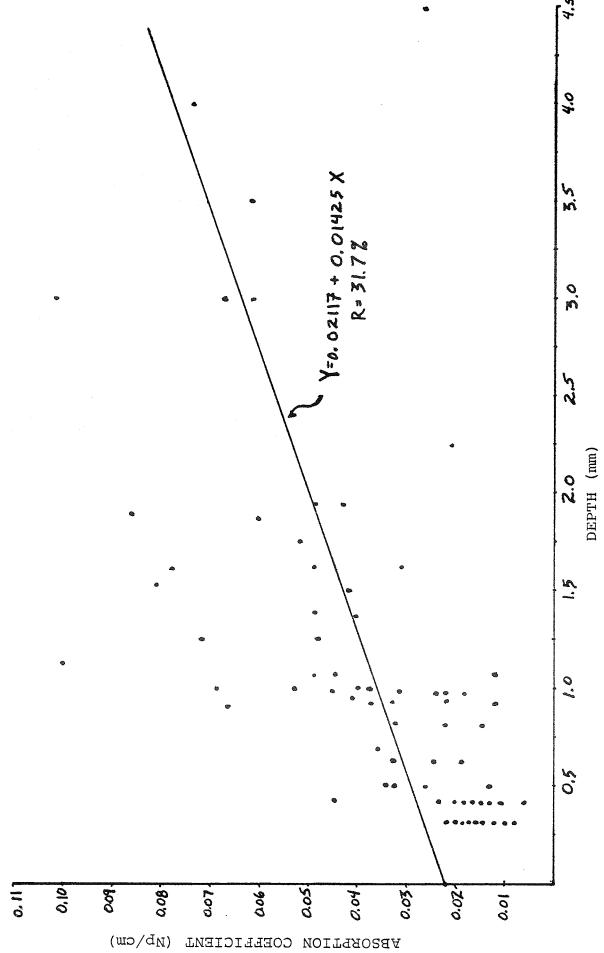
high calculated absorption coefficents for muscle under the spine are the result of a low calculated "in <u>sitū</u> intensity", as the dT/dt values obtained from these sites are in the same range as the dT/dt values obtained from back muscle sites not under the spine. This low "in situ intensity" is a result of faulty modeling of the interposing tissues. Recall from Section 4.2, that all the tissue boundries were modeled as plane interfaces of infinite lateral extent. Modeling the spinal column as a specular reflector results in an impedance discontinuity which allows only 25% of the incident acoustic intensity to transmitted. In actuality, the diameter of the spine is which is about twice the wavelength. The scattered and diffracted radiation has a distribution which is both complex and critically dependent upon the dimensions characteristic impedance of the spinal column. Furthermore, the spinal diameter is approximately the same size as the

minimum cross-sectional beam diameter (Dt). Thus, the spinal column will not interact with the entire acoustic wave as would an infinite interface. It can be concluded therefore, that the dosimetric model must be changed to produce a more suitable "in situ intensity" at irradiation sites which are under the spine.

Another consideration in the area of the spine is heat diffusion. During irradiation, the spinal column temperature will rise faster than the temperature of the surrounding tissues. This is due to the higher absorption coefficient of bone. Thus, a higher temperature rise will be measured in the tissues surrounding the spine than would be expected because of this added spinal heating source. No assumptions were made in the dosimetric model concerning this diffusion phenomena.

The abdominal muscle data were also inspected on the specimen level. Since the "corrected" mean value is fairly close to the literature value, one might infer that the dosimetric model is adequate for abdominal muscle irradiation sites. Upon closer inspection of the MMA data in Appendix D though, an interesting trend was discovered. It appeared that the magnitude of the calculated absorption coefficient was proportional to the depth of the thermocouple junction within the specimen in the direction of the focal axis, that

is , in the X direction. The muscle absorption coefficient though, should be independent of the tissue depth. In order to substantiate this trend, the calculated absorption coefficient was plotted against the thermocouple junction depth within the specimen (Figure 18). A first order least squares linear regression model was used to fit the points. The resulting regression equation was Y=0.02117+0.01425*X. Although this equation produces a which supports the trend, further analysis was performed to determine if there was significant regression and no lack of (Draper and Smith, 1966). The results of this analysis are shown in the Analysis of Variance Table (Table 4). two values of greatest interest in this table are R $% \left(1\right) =\left(1\right) +\left(1\right)$ The R value measures the proportion of the total variation about the mean Y explained by the regression. In this case R = 0.317. Thus, the regression equation obtained only 31.7% of the total variation. The F value denotes the significance of the lack of fit. From Table 4, F=7.0suggests that the lack of fit is significant. These two values indicate the first order regression equation obtained here is inadequate in describing the data points. A higher order regression model might be used to fit the data points. With the data presently available, the suspicion that the calculated absorption coefficient is proportional to depth can not be substantiated. Further investigation to



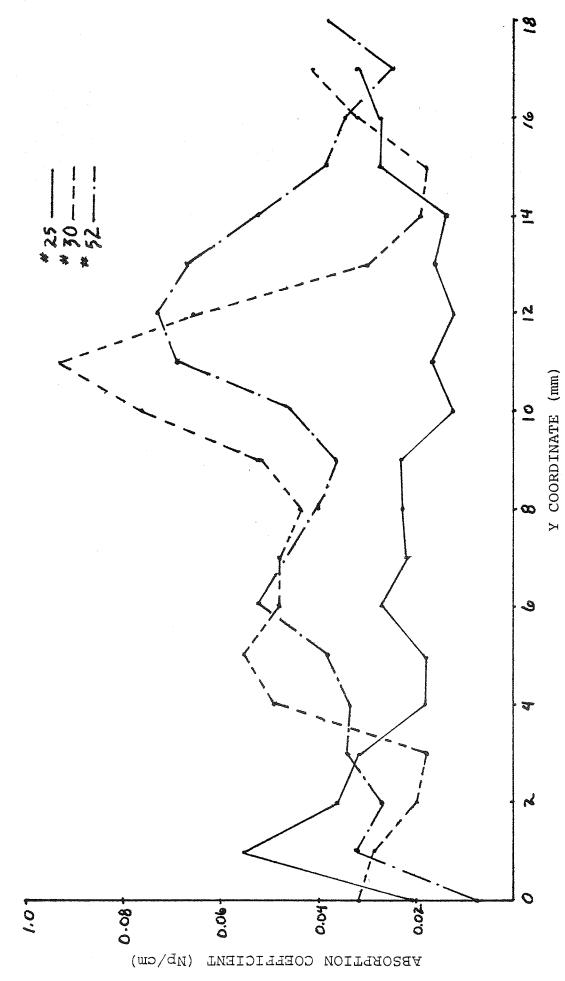
Absorption coefficient vs. tissue depth in direction of focal axis for abdominal ω muscle tissue Figure 18.

Table 4: Analysis of Variance Table

Sōūrĉē	Degrees of <u>Freêêdôm</u>	Sum of <u>Squares</u>	Mean <u>Square</u>	
Total	6 6	0.03624		R = 0.317
Regression	1	0.01149	0.01149	F = 30.24
Residual	65	0.02474	0.00038	
Lack of Fit	21	0.01901	0.00091	F = 7.0
Pure Error	4 4	0.00573	0.00013	

increase the data base size might produce data points which yield a significant regression equation with no lack of fit.

During the course of obtaining the irradiation data it was suspected that the angle of incidence of the acoustic beam with respect to the specimen affected the thermocouple This suspicion arose in a number of specimens which the thermocouple was inserted just under the skin. As the junction was pulled through the mouse higher dT/dt values were measured at in the central irradiation sites (normal acousic incidence) than at sites near the skin perforations (oblique incidence). In a few specimens though, higher dT/dt values were measured close to skin perforations. Since the irradiation sites near the perforations were the same tissue as those in the center of the mouse (skin/abdominal muscle interface), these dT/dt differences were not expected. Figure 19 is a comparison of specimens 25, 30 and Plotted is the calculated absorption coefficient thermocouple position along the Y axis. These specimens had thermocouple paths which were virtually identical in terms of the constituent tissues. Figure 19 shows that specimens 30 and 52 are examples of the first The central irradiation sites have a supposedly higher case. absorption coefficient than the identical tissues near skin perforations. Specimen 25 shows the opposite trend. can be seen in all three specimens that there is an increase



Aborption coefficient vs. thermocouple junction position on Y axis for abdominal muscle tissue in specimens numbered 25, 30, and 52Figure 19.

in the calculated absorption coefficient very close to the skin perforation. This is most likely due to the higher absorption coefficient of skin. It can not be concluded from this data whether or not acoustic incidence affects the determination of the absorption coefficient. This issue must be settled as new experimental procedures and data analysis algorithms would have to be developed if it were shown the assumption of normal acoustic incidence could not be substantiated at each irradiation site.

In summary, it has been shown that using computer files to manipulate the experimental data provides a successful means of achieving complex dosimetric analysis. The results of this analysis have shown the dosimetric model used to calculate the "iñ sitū intensity" to be incomplete. Allowances must be made for: 1) tissue boundaries which can not be modeled as infinite planes normal to the focal axis; 2) possible heat diffusion effects; 3) possible refraction of the acoustic beam at tissue boundaries. Combining these factors with the valid assumptions made in the first model, a second generation dosimetric model may be proposed.

CHAPTER 5

CONCLUSION

results discussed in the previous chapter show the approach used in the experiment to obtain dosimetric information is valid. Although the first generation dosimetric model was shown to have some inadequacies, a number of interesting interactions between the ultrasonic beam and the specimen tissues were characterized. Τt been shown that the transient thermoelectric technique was well suited for this application, and the associated irradiation instrumentation used to implement the data acquisition algorithm was adequate. The histological sectioning techniques developed here were not only valuable in this experiment, but may also be applied to other types of The data analysis performed in this study was primarily completed to substantiate the dosimetric model. Unfortunately, little progress was made in identifying dosimetric parameter. Before this parameter can be realized, number of steps must be taken to improve the experimental procedure, instrumentation, and data analysis techniques. The purpose of this chapter is to suggest possible experimental improvements, which upon implementation, will yield a complete dosimetric analysis.

In reviewing the experimental procedure, it becomes clear that the inherent flaw in this experiment is the uncertainty in the calculated "in situ intensity". intensity value was completely dependent upon the accuracy of the dosimetric model. If the interposing tissues were modeled incorrectly or beam refraction effects were present, an erroneous value for the tissue absorption coefficient would be obtained. Thus, an accurate method of obtaining the "iā <u>šifū</u> intensity" is imperative for experimental success. It would be desirable to measure the "in situ intensity" The probe used to measure this intensity must produce a response which is independent of the surrounding Such a probe can be developed using the transient thermoelectric technique. The thermocouple junction encased in a small bolus of polyethylene or epoxy resin to isolate the junction from the tissue. The absorption coefficients of these embedding materials can be determined. Using a short acoustic pulse (0.1 sec) to assure the heat from the surrounding tissue does not diffuse into the bolus, the thermocouple response used with equation 9 will yield the " $ilde{1}$ $ilde{n}$ $ilde{1}$ $ilde{n}$ $ilde{n}$ ilcoefficient at the irradiation site, the encased junction could be combined with a naked junction into a single thermocouple wire. These junctions would be separated by a distance of 3 cm to assure only one junction at a time is affected by the acoustic beam. To obtain both the ${\mathfrak C}$ and Is

at an irradiation site, the encased junction is first pulled into the site and irradiated with a short pulse. From the received thermocouple response the "in situ intesity" may be calculated. The naked thermocouple junction is then moved into the irradiation site, and irradiated with a one second pulse. This response, combined with the "in situ intensity" found moments earlier, will yield an accurate absorption coefficient value. This double junction probe will be especially useful in tissues which are difficult to model such as the intestinal organs and tissues surrounding the spine.

To successfully implement the procedures involved with this double junction probe, precise control over positions of the junctions must be obtained. The present specimen structure does not provide this control. Furthermore, this structure is not physically large enough to move both junctions (3 cm apart) through an entire specimen. A new structure which does not have these liabilities has been built (Vaughn, 1980). This structure has a number advantages over the specimen structure described in Section 3.2. They are: 1) precise computer controlled motor thermocouple junction positioning; 2) freedom of orientation with respect to the specimen structure greater thermocouple path choice; 3) multiple thermocouple wires may be used; 4) covered thermocouple lead connectors to

allow saline acoustic propagation solutions to be used; 5)coaxial thermocouple leads; 6)fewer Plexiglas obstructions around the specimen. The use of this specimen structure provides a means of changing the orientation of the mouse with respect to the focal axis as well as the ability to probe different tissue combinations.

To fully examine the interactions of the acoustic beam the tissues within the specimen, in vivo beam profiling should be done. Steps have already been taken to mechanize profiling procedure(Vaughn, 1980). The milling base has been placed under computer control. Sychronized movements of the thermocouple probe and milling base can be produced by programming the Interdata minicomputer. The output of a new data acquisition system is also fed to the Interdata. This system consists of a 2 stage amplifier which accepts the thermocouple response and an A/D converter. Thus, all thermocouple responses can be digitized, analyzed and stored for later use.

In this digitized form the thermocouple responses can be analyzed in a number of ways (Duback, 1980). The time rate of change of temperature can be investigated at any instant of the response. This investigation may yield information concerning heat diffusion effects. Detailed examination of the viscous rise may suggest a method for the

characterization of different tissues. It must be noted that the dT/dt value was a parameter not examined in Chapter 4. This value is a direct indication of the energy absorbed by the irradiated tissue. Therefore, the dT/dt values should be closely examined in further dosimetric analysis.

summary, it has been shown that the approach and experimental procedures discussed in this thesis provide useful information for dosimetric analysis. In the continuation of this study, the changes to the dosimetric model proposed in Chapter 4 should be incorporated into a second generation model. Using this new model, the new above, and the experimental instrumentation described procedures and data compilation techniques described Chapters 3 and 4, the information necessary to develop a total dosimetric concept will be obtained. Once this concept is complete, ultrasound may be used with greater confidence in medical applications.

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APPENDIX A

EXAMPLE ENTRY FROM IRRADIATION FILE DOSEXPC.DTA

DESCRIPTION OF COLUMNS IS FOUND IN SECTION 4.2

INCIDENT INTENSITY																							ର ପ୍ର	38. 2	
DEFLECTION	1.51	1.56																					2.90		
Tara	0.73	0.79	1.22	0.97	0.53	0.53	0.46	0.66	0.61	0.47	0.53	0.64	0.46	0.38	0.45	0.96	0.79	1.57	1.60	1.06	0.56	1.36	1.59	0.46	
TC POSITION	75	7.6	77	78	79	80	81	85	63	40	92	88	87	88	68	06	9.1	92	63	94	95	9.6	67	86	
SHOT#		Ci	n	4	IJ	<i>ব</i>	7	۵	4	10	11	2	13	14	15	16	17	18	15	50	70	eg G		5.4	
7	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
×	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	O :	O :	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
>-	O :	0.	€. O	ල ල	4.0	ુ: ○	6.0	7.0	9.0	٥. ٥.	10.0	11.0	12.0	13.0	14.0	15.0	15.0	17.0	13.0	19.0	20.0	21.0	22.0	23.0	
#31	 i	0	a	0	0	0	٥	0	0	0	0	0	0	0	0	0	0	C	0	٥	Ç	0	C	0	
MCNSE#	38	٥	a	Q	¢	0	٥	Ģ	٥	0	۵	٥.	C	૦ ,	a	\$	0	٥	0	٥	٥	0	\$	Q	

APPENDIX B

EXAMPLE ENTRY FROM HISTOLOGY FILE DOSHIS.DTA

DESCRIPTION OF COLUMNS IS FOUND IN SECTION 4.3

#3SAOM	#31	>	><	7	DEPTH T	SSUE CODE	INTE	ERPOS	SING	!	4088					
36		0.0	0.0	0.0	0.0	OC.W		0.0					_	C	c	
Ç	٥	1. O	0.0	0.0	0.2	DER	DER	O. 23		<u>.</u>	0	0.0		Ċ	0	
¢	٥	2.0	0.0	0.0	D. C.	nro	DER							0	0	
٥	0	3.0	0.0	0.0	m m	MUM	DER							Ċ		
٥	0	4.0	0.0	0.0	4. 5	KIC	DER							Ó		
Ç	0	5.0	0.0	0.0	5.2	KIC	DER						_	O		
Ç	0	6.0	0.0	0.0	6. 2	K16	DER							Ó		
٥	٥	7.0	0.0	0.0	7.0	KIM	DER		MMA		ADP 5		KIG		CT.	
¢	0	O	0.0	0.0	7.8	KIC	DER									
¢	٥	9.0	0.0	0.0		KIC	DER									
666	٥	٠ . ٥	0.0	0.0	9. 1	KIC	K1C									
C	٥	10.0	0.0	0.0	8.0	BLO	DER									
Ç	٥	11.0	0.0	0.0	7.2	BLO	DER									
٥	0	12.0	0.0	0.0	7.0	CNC	DER		MMB) MFIB			
666	0	12.0	0.0	0.0	7.0	CNT	CNJ				_					
٥	0	13.0	0.0	0.0	7.0	FAS	DER		MMB		4 NSP					
666	0	13.0	0.0	0.0	7.0	FAS	FAS				<u> </u>					
٥	0	14.0	0.0	0.0	6.9	FAS	DER				dSM C					
666	0	14.0	0.0	0.0	6.9	·FAS	ADP		FAS		J7					
٥	0	15.0	0.0	0.0	6.8	FAS	DER				4 ADP		2 181.0			
666	0	15.0	0.0	0.0	6.8	F-AS	FAS				\Box					
Ç	0	16.0	0.0	0.0	ь. Б	FAS	DER		MME		_					
٥	0	17.0	0.0	0.0	6.0	FAS	DER				5 ADP		F.AS			
٥	0	18.0	0.0	0.0	O	PLG	DER				-					
666	0	18.0	0.0	0.0	5.0	PL 6	PL 6				\sim				c	
٥	0	19.0	0.0	0.0	ත ස	PL.0	DER		MMA		1 ADP		3 14	,,,		
\$	0	20.0	0.0	0.0	1.7	MMA	DER		MMV				<u>^</u>	0	٥.	
٥	٥	21.0	O .C	0.0	1.2	MMM	DER		MMA		m		_	0	<u>۔</u>	
0	0	22.0	0.0	0.0	Ç.	NER.	DEX				\cap		_	0	<u>۔</u>	
ಎ	O	23.0	0.0	0.0	0.0	MCG				<u>.</u>	\sim		_	0	C	

APPENDIX C

EXAMPLE ENTRY FROM DOSIMETRIC FILE DOS.DTA

DESCRIPTION OF COLUMNS IS FOUND IN SECTION 4.4

ALPHA (NP/CM)	0.038	0.040	0.066	0.056	0.030	0.030	0.027	0.040	0.03B	0.034	0.033	0.040	0.552	1.272	0.507	0.049	0.049	0,075	0.078	070.0	0.030	0.072	0.000	0.023
I (W/CM2)	38. 2	37.3	34.9	32. 9	33. 4	32. 9	32. 1	31.2	30.7	26. 1	30. 5	30.5	1.6	9.0	1.7	26.3	30. 2	30.8	30.7	33. 1	05. B	35. 9	37.3	36. 2
DTDT(DEG/SEC)													0.46											
DEPTH(MM)													7.0							Э. В				0.0
TISSUE CODE	W.J.D.	DER	BLO	M.M.	KIC	KIC	KIG	KIM	KIC	KIC	BLC	BLO	CNI	FAS	FAS	FAS	FAS	FAS	PL¢	974	MMA	PIFIA	DER	MC Q
Ν	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
×	0.0	0.0	0.0	0.0	0.0	0.0	O :O	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
>-	0.0	0.1	2.0	3.0	4.0	5.0	6.0	7.0	9.0	9.0	10.0	11.0	12.0	13.0	14.0	15.0	15.0	17.0	18.0	19.0	20.0	21.0	22.0	23.0
TC#	1 i	0	c	٥	0	0	0	0	0	0	0	٥	0	0	0	0	0	0	0	0	0	O	0	0
*COUSE#	38	36	36	-9 -9 -0	38	36	36	36	36	36	36	36	36	36	36	38	36	38	36	36	36	38	36	36

APPENDIX D

THIS TABLE IS A LISTING OF DOS.DTA DIVIDED BY TISSUE TYPE

COLUMN 1- SPECIMEN NUMBER

COLUMN 2- THERMOCOUPLE NUMBER

COLUMN 3-THERMOCOUPLE JUNCTION COORDINATES WITH RESPECT TO THE SPECIMEN (mm).

COLUMN 4- TISSUE CODE

COLUMN 5- THERMOCOUPLE DEPTH BELOW THE SKIN SURFACE IN THE DIRECTION OF THE FOCAL AXIS (mm)

COLUMN 6- TIME RATE OF CHANGE OF TEMPERATURE (C/SEC)

COLUMN 7- "IN SITU INTENSITY" (W/CM**2)

COLUMN 8- TISSUE ABSORPTION COEFFICIENT (NP/CM)

7 8 10 9 30 31 9 25 25 33 33 36 40 40 41 40 40 40 40 40 40 40 40 40 40 40 40 40	18 (18 (18 (18 (18 (18 (18 (18 (18 (18 (13.0 7.0 9.0 14.0 3.0 4.0 15.0 23.0 24.0 1.0 22.0 1.0 22.0 1.0 22.0 1.0 22.0 1.0 22.0 1.0 22.0 1.0 22.0 1.0 22.0 1.0 22.0 1.0 22.0 1.0 23.0 24.0 1.0 25.0 26.0 10.0			ADPPPPPPPAAAAAAAAAAAAAAAAAAAAAAAAAAAAA	0.3 0.3 0.3 0.3 0.3 1.1 0.6 1.2 1.0 0.8 1.1 1.7 1.7 1.7 1.7 1.7 1.7 1.7	0.74 0.78 0.32 0.93 1.15 0.37 0.45 1.38 0.95 1.08 1.08 1.08 1.09	49.7 49.7 49.7 49.8 48.9 36.3 18.5 148.1 49.6 18.8 11.7 17.9 17.9 17.9 17.9 18.8 18.6 18.7 17.9 17.9 17.9 18.8 18.6 18.7 18.6 18.7 17.9 17.9 18.8 18.6 18.6 18.7 18.6	0.030 0.028 0.030 0.013 0.036 0.044 0.019 0.046 0.045 0.054 0.030 0.036 0.106 0.049 0.131 0.022 0.402 0.087 0.063 0.079 0.045 0.054 0.055 0.054 0.055 0.
168 189 205 213 214 277 376 378	34 0 36 0 36 0 36 0 39 0 43 0	2.0 0 10.0 0 11.0 0 9.0 0 11.0	0.0 0.0 0.0 0.0 0.0	0.0 0.0 0.0 0.0 0.0	BLO BLO BLO BLO BLO BLO BLO	9.7 7.5 2.0 8.0 7.2 8.8 8.8	0.22 0.41 1.22 0.53 0.64 1.06 0.56	24.6 30.7 34.9 30.5 30.5 40.5 1.9	0.017 0.025 0.066 0.033 0.040 0.049 0.563
379	44 0		0.0	0.0	BLO	8.2		28.9	0.059

```
400
         44
            0 11.0
                      0.0
                           0.0 BLO
                                     8.5 0.30 21.9 0.026
401
         44
            0 12.0
                      0.0
                           0.0 BLO
                                     8.3 0.20 29.3 0.013
303
            0 11.0
         40
                      0.0
                           0.0 BVV
                                     7.3 0.40
                                                 2.1 0.357
377
         43
            0 12.0
                      0.0
                           0.0 BVV
                                     9.0 0.24
                                                 3.3 0.137
402
            0 13.0
         44
                      0.0
                           0.0 BVV
                                     8.2 0.21
                                                 0.4 0.921
403
         44
            0 14.0
                      0.0
                           0.0 BVV
                                     7.9
                                          0.31
                                                 0.4 1.360
427
         47
            0 12.0
                      0.0
                           0.0 BVV
                                     6.3 0.27
                                                 0.3 1.620
472
         51
            0 12.0
                      0.0
                           0.0 BVV
                                     5.3 0.44
                                                 5.2 0.161
235
         37
            0
                9.0
                      0.0
                           0.0 BJV
                                     7.0 0.46 19.4 0.045
236
         37
            0 10.0
                     0.0
                           0.0 BJV
                                     7.3 0.27
                                               23.3 0.022
191
         34
            0 11.0
                           0.0 MJB
                     0.0
                                     8.2 1.20
                                                 7.5 0.301
193
         34 0 13.0
                     0.0
                           0.0 MJB
                                     8.1 0.52 29.3 0.034
  2
         17 0
                2.0
                      0.0
                           0.0 CNT
                                     1.5 0.96 48.3 0.038
  3
         17 0
                4.0
                     0.0
                           0.0 CNT
                                     1.2 0.92 48.8 0.036
  5
         17
            0
                8.0
                      0.0
                           0.0 CNT
                                     0.6 1.37 50.1 0.052
169
         33
            0 13.0
                     0.0
                           0.0 CNT
                                     9.8 0.17 25.2 0.013
170
         33
            0 14.0
                                     9.7 0.18 20.8 0.016
                      0.0
                           0.0 CNT
215
         36
            0 12.0
                     0.0
                           0.0 CNT
                                     7.0 0.46
                                                1.6 0.552
237
         37
            0 11.0
                                     7.8 0.23 18.4 0.024
                     0.0
                           0.0 CNT
238
         37
            0 12.0
                      0.0
                           0.0 CNT
                                     8.0 0.21 19.3 0.021
         39 0 10.0
278
                     0.0
                           0.0 CNT
                                     9.2
                                          0.67
                                                 1.5 0.862
354
         42
            0 12.0
                     0.0
                           0.0 CNT
                                     7.6
                                          1.83
                                                 1.5 2.349
         47 0 11.0
426
                     0.0
                           0.0 CNT
                                     6.1 0.85
                                                 5.9 0.270
355
         42 0 13.0
                     0.0
                           0.0 CJM
                                     7.8 2.45
                                                 5.5 0.839
 37
         23
            0
                5.0
                     0.0
                           0.0 CRU
                                     3.7 0.94 45.4 0.039
 38
         23
            Ω
                7.0
                     0.0
                           0.0 CRU
                                     4.1 3.25 11.5 0.535
 39
                8.0
         23
            0
                     0.0
                           0.0 CRU
                                     4.1 1.44 44.2 0.062
         24 1
 44
                9.0
                     0.0
                           0.0 CRU
                                     1.5 1.52 46.7 0.061
 45
         24
            0 10.0
                     0.0
                           0.0 CRU
                                     1.4 1.74 47.1 0.070
 46
         24
            0
              11.0
                     0.0
                           0.0 CRU
                                     1.4 2.73 46.2 0.112
 89
         28
            0
                5.0
                     0.0
                           0.0 CRU
                                     0.6 0.69 36.1 0.036
 92
         28
            0
                8.0
                     0.0
                           0.0 CRU
                                     0.7 0.54 35.8 0.028
 93
         28
            0
               9.0
                     0.0
                           0.0 CRU
                                     0.8
                                         0.42 18.4 0.043
         28
 94
            0 10.0
                     0.0
                           0.0 CRU
                                     0.9 0.44 18.3 0.046
172
         33 0 16.0
                     0.0
                           0.0 CRU
                                     9.0
                                         0.17 26.9 0.012
173
         33
            0 17.0
                     0.0
                           0.0 CRU
                                     8.8 0.00 15.2 0.000
174
         33
            0 18.0
                     0.0
                           0.0 CRU
                                     8.5 0.12 15.7 0.014
175
         33 0 19.0
                     0.0
                           0.0 CRU
                                     8.0 0.21 16.3 0.024
176
         33 0 20.0
                     0.0
                           0.0 CRU
                                     7.5 0.19 17.0 0.021
 25
         20 0 13.0
                     0.0
                           0.0 DER
                                     0.2 0.34 50.9 0.013
 51
         24 0
              17.0
                     0.0
                           0.0 DER
                                     0.1 0.32 51.2 0.012
 54
         25 0
               1.0
                     0.0
                           0.0 DER
                                     0.5 1.45 50.2 0.055
```

```
69
         25
            0 15.0
                      0.0
                            0.0
                                DER
                                       0.5 0.69 50.2 0.026
 85
         28
            0
                1.0
                      0.0
                            0.0 DER
                                       0.1 0.49 37.5
                                                       0.025
 86
         28
            0
                2.0
                      0.0
                            0.0 DER
                                       0.2
                                           0.42 37.3 0.021
115
         29
            0
               16.0
                            0.0 DER
                                      0.2 0.72 50.9 0.027
                      0.0
151
         32
            0
                1.0
                      0.0
                            0.0 DER
                                       0.2
                                           0.08 19.3 0.008
157
         33
             0
                1.0
                      0.0
                            0.0 DER
                                      1.8
                                           0.82 27.3 0.057
204
         36
             0
                1.0
                      0.0
                            0.0 DER
                                       0.2
                                           0.79
                                                 37.3 0.040
225
         36
            0
               22.0
                      0.0
                            0.0 DER
                                       0.2 1.59 37.3 0.080
250
         38
             0
                1.0
                      0.0
                            0.0 DER
                                      0.2 0.80 19.3 0.078
267
         38
             0
               21.0
                      0.0
                            0.0 DER
                                       0.2
                                           1.32
                                                   2.7 0.912
269
         39
             0
                1.0
                            0.0 DER
                      0.0
                                       0.2
                                           0.53 50.9 0.020
293
         40
                1.0
            0
                      0.0
                            0.0 DER
                                      1.8 0.59 19.4 0.058
340
         41
            0
               21.0
                      0.0
                            0.0 DER
                                       0.2 1.34 19.3 0.131
388
         43
             0
               23.0
                      0.0
                            0.0
                                DER
                                       0.6
                                           1.17 36.6 0.060
389
         44
            1
                0.0
                      0.0
                            0.0 DER
                                       0.0
                                           0.12 38.2 0.006
414
         44
            0
               25.0
                      0.0
                            0.0 DER
                                       0.0
                                           0.26
                                                  7.4 0.066
415
         47
            1
                0.0
                      0.0
                            0.0 DER
                                      0.0 1.52 19.8 0.145
436
         50
            0
                1.0
                      0.0
                            0.0 DER
                                      2.4 0.52 34.0 0.029
 12
         19
            1
                1.0
                      0.0
                            0.0
                                DJA
                                      0.3 0.53 50.7 0.020
 13
         19
             0
                3.0
                      0.0
                            0.0 DJA
                                       0.2 0.54 50.9 0.020
 19
         20
            ï
                1.0
                      0.0
                            0.0 DJM
                                       0.2
                                           0.39 50.9 0.014
 20
         20
             0
                3.0
                      0.0
                            0.0 DJM
                                       0.2 0.49 50.9 0.018
                5.0
7.0
 21
         20
             0
                      0.0
                            0.0
                                DJM
                                       0.3
                                           0.59
                                                 50.7
                                                       0.022
 22
         20
             0
                      0.0
                            0.0
                                DJM
                                      0.5
                                           0.58 49.8 0.022
 24
         20
            0
               11.0
                      0.0
                            0.0 DJM
                                      0.2 0.45 50.9
                                                       0.017
 55
         25
            0
                2.0
                      0.0
                            0.0
                                DJM
                                       0.3 0.96 50.5 0.036
 56
         25
             0
                3.0
                      0.0
                            0.0
                                       0.2
                                DJM
                                           0.86 50.9 0.032
 57
         25
                4.0
             0
                                DJM
                                      0.3
                                           0.50 50.7 0.019
                      0.0
                            0.0
 5.8
         25
             0
                5.0
                      0.0
                            0.0 DJM
                                      0.3 0.47 50.7 0.018
 59
         25
             0
                6.0
                      0.0
                            0.0
                                DJM
                                       0.3
                                           0.72 50.7 0.027
 60
         25
            0
                7.0
                      0.0
                            0.0
                                DJM
                                       0.2
                                           0.59 50.9
                                                       0.022
 6ĩ
         25
            0
                8.0
                      0.0
                            0.0 DJM
                                      0.3
                                           0.62 50.7 0.023
 62
         25
            0
                9.0
                      0.0
                            0.0
                                DJM
                                      0.3
                                           0.63 50.7 0.023
 63
         25
              10.0
            0
                      0.0
                            0.0
                                DJM
                                      0.4
                                           0.34 50.3 0.013
 64
         25
            0 11.0
                      0.0
                            0.0
                                DJM
                                      0.3 0.46 50.7 0.017
 65
         25
            0
              12.0
                      0.0
                            0.0
                                DJM
                                      0.2 0.33 50.9 0.012
 66
         25
            0
               13.0
                      0.0
                            0.0 DJM
                                      0.4 0.42 50.4 0.016
 67
         25
            0 14.0
                      0.0
                            0.0 DJM
                                      0.5
                                           0.38 50.2 0.014
 68
         25
            0
              15.0
                      0.0
                            0.0 DJM
                                      0.6
                                           0.68 48.5
                                                       0.027
 99
         28
            0 15.0
                      0.0
                            0.0 DJM
                                      0.6 0.30 36.7 0.015
114
         29
            0
               15.0
                      0.0
                            0.0
                                DJM
                                      0.6
                                           0.32 50.0 0.012
118
         30
            0
                1.0
                      0.0
                            0.0
                                DJM
                                      0.8
                                          0.76 49.5
                                                       0.029
119
         30
            0
                2.0
                      0.0
                            0.0
                                DJM
                                      0.7
                                           0.53 49.7
                                                       0.020
120
         30
            0
                3.0
                      0.0
                            0.0
                                DJM
                                      0.6
                                           0.48 49.9
                                                       0.018
121
         30
            0
                4.0
                      0.0
                            0.0 DJM
                                      0.6 1.30 49.9 0.049
122
         30
            0
                5.0
                      0.0
                            0.0 DJM
                                      0.5 1.46 50.2 0.055
123
         30
            0
                6.0
                      0.0
                            0.0 DJM
                                      0.5 1.28 50.2 0.048
124
         30
            0
                7.0
                      0.0
                            0.0 DJM
                                      0.6 1.27 49.9 0.048
         3 0
3 0
125
                      0:0
                            8:8
                                DJM
DJM
                                      8:7
                                           1:35
                                                 43:3
                                                       8:844
127
         30
            0 10.0
                      0.0
                            0.0 DJM
                                      0.6 2.02 49.9 0.076
```

```
128
         30
            0 11.0
                      0.0
                           0.0 DJM
                                     0.7 2.44 49.7 0.093
131
         30
                                     0.7
            0 14.0
                      0.0
                           0.0 DJM
                                          0.50 49.7
                                                      0.019
132
                     0.0
         30 0 15.0
                           0.0 DJM
                                     0.6 0.47 50.0 0.018
133
         30
            0 16.0
                      0.0
                           0.0 DJM
                                     0.4 0.85 50.4 0.032
135
         31
            1
                1.0
                      0.0
                           0.0 DJM
                                     0.2 0.43 50.9
                                                     0.016
148
              15.0
         31
            0
                      0.0
                           0.0 DJM
                                      0.6 0.45 50.0 0.017
227
         37
            1
                1.0
                           0.0 DJM
                      0.0
                                     1.2 0.91 26.3 0.065
103
         28
            0 19.0
                           0.0 MJD
                      0.0
                                      0.2 0.46 37.3 0.023
149
         31
            0 16.0
                      0.0
                           0.0 MJD
                                      0.1 0.37 51.2 0.014
317
         40
            0 25.0
                      0.0
                           0.0 MJD
                                      0.3 0.95 50.7 0.035
 11
         18
            0 15.0
                      0.0
                           0.0 DJW
                                     0.0 0.38 52.1 0.014
 52
         24
            0 18.0
                      0.0
                           0.0 DJW
                                     0.0 0.72 52.1 0.026
 70
         25
            0 17.0
                      0.0
                           0.0 DJW
                                     0.0 0.83 52.1 0.032
104
         28
            0 20.0
                      0.0
                           0.0 DJW
                                     0.0 0.22 38.2 0.011
116
         29
            0 17.0
                      0.0
                           0.0 DJW
                                     0.0 0.18 52.1 0.007
134
         30
            0 17.0
                      0.0
                           0.0 DJW
                                     0.0 1.12 52.1 0.041
179
         33
            0 23.0
                      0.0
                           0.0 DJW
                                     0.0 0.39 30.2 0.024
            0 22.0
202
         34
                      0.0
                           0.0 DJW
                                     0.0 0.69 38.2 0.034
226
         36 0 23.0
                      0.0
                           0.0 DJW
                                     0.0 0.46 38.2 0.023
248
         37 0 22.0
                           0.0 DJW
                      0.0
                                      0.0 0.25 28.3 0.017
318
         40 0 26.0
                      0.0
                           0.0 DJW
                                      0.0 0.70 52.1 0.025
341
         41 0 22.0
                      0.0
                           0.0 DJW
                                     0.0 1.11 19.8 0.106
459
         50
              24.0
            0
                      0.0
                           0.0 DJW
                                      0.0 0.45 38.2 0.022
460
         51
            1
                0.0
                      0.0
                           0.0 DJW
                                     0.0 0.69 19.8 0.066
483
         52
            1
                0.0
                      0.0
                           0.0 DJW
                                     0.0 0.14 28.3 0.009
502
         53
            1
                0.0
                           0.0 DJW
                      0.0
                                     0.0 0.35 19.8 0.033
  Ĭ
         17
            ì
                0.0
                      0.0
                           0.0 WJD
                                      0.0 0.08 52.1 0.003
 53
         25
            ĩ
                0.0
                      0.0
                           0.0 WJD
                                      0.0 0.58 52.1 0.021
 71
         26
            1
                0.0
                      0.0
                           0.0 WJD
                                      0.0 0.72 52.1 0.026
 84
         28
            1
                0.0
                      0.0
                           0.0 WJD
                                      0.0 0.75 38.2 0.037
117
         30
            1
                0.0
                      0.0
                           0.0 WJD
                                      0.0 0.85 52.1 0.031
150
         32
                0.0
            1
                      0.0
                           0.0 WJD
                                      0.0 0.08 19.8 0.008
156
         33
            1
                0.0
                                      0.0 0.71 30.2 0.044
                      0.0
                           0.0 WJD
180
         34
            1
                0.0
                      0.0
                           0.0 WJD
                                      0.0 1.29 38.2 0.064
203
         36
            1
                0.0
                      0.0
                           0.0 WJD
                                      0.0 0.73 38.2 0.036
249
         38
            1
                0.0
                      0.0
                           0.0 WJD
                                      0.0 1.11 19.8 0.106
268
         39
            ĭ
                0.0
                      0.0
                           0.0 WJD
                                      0.0 0.39 52.1 0.014
292
         40
                           0.0 WJD
            ĭ
                0.0
                      0.0
                                      0.0 0.47 21.4 0.042
319
         41
            1
                0.0
                      0.0
                           0.0 WJD
                                      0.0
                                          1.14 19.8 0.109
342
         42
                           0.0 WJD
            1
                0.0
                      0.0
                                      0.0 0.84 38.2 0.042
365
         43
            1
                0.0
                      0.0
                           0.0 WJD
                                      0.0 0.73 12.9 0.107
435
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                0.0
                     0.0
                           0.0 WJD
                                      0.0 0.20 38.2 0.010
216
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                      0.0
                           0.0 FAS
                                     7.0 0.38
                                                 0.6 1.272
217
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                      0.0
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                                      6.9
                                          0.45
                                                 1.7 0.507
218
         36
           0 15.0
                      0.0
                           0.0 FAS
                                      6.8
                                          0.96 26.3 0.069
219
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                      0.0
                           0.0 FAS
                                     6.5
                                          0.79
                                               30.2 0.049
220
         36
           0 17.0
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                                     6.0 1.57 30.8 0.096
229
         37
            0
                3.0
                           0.0 FAS
                      0.0
                                     4.3 0.98 21.6 0.086
```

305 450 451 35 72 304	50 50 23 26	0 13.0 0 15.0 0 16.0 1 1.0 0 1.0 0 12.0	0.0 0.0 0.0 0.0 0.0	0.0 0.0 0.0	FAS FAS MJF MJF MJF	6.7 6.4 6.1 1.3 1.0 6.8	0.46 0.45 0.84 0.56 0.76 0.39	40.8 30.1 30.3 49.0 50.0 2.6	0.021 0.028 0.052 0.022 0.029 0.284
279 280		0 11.0 0 12.0	0.0	0.0	GUG GUG		1.88		0.349
182 207 208 211 212 239 240 243 244 272 275 285 296 299 300 307 311 321 325 333 344 345 353 369 370 371 372	36 36 36 37 37 37 37 39 40 40 40 40 41 41 41 41 41 42 42 42 43 43 43 43 43	0 2.0 0 4.0 0 5.0 0 8.0 0 9.0 0 13.0 0 14.0 0 17.0 0 14.0 0 14.0 0 15.0 0 15.0 0 14.0 0 15.0 0 15.0 0 18.0 0 15.0 0 18.0 0 19.0 0 14.0 0 15.0 0 18.0 0 19.0 0 19.0 0 110.0 0 10.0 0		0.0 0.0 0.0 0.0 0.0 0.0	KIC	2.1 2.8 4.0 7.5 4.0 6.0 6.5 7.5	0.53 0.61 0.47 0.20 0.34 0.46 0.57 2.14 0.82 0.31 0.26 0.55 0.37 0.07 0.47 0.067 0.44 0.45 0.39 1.39 0.72 0.46 0.46 0.72 0.46 0.47 0.46 0.57 0.57 0.67 0.47 0.46 0.46 0.47	35.7 33.8 33.1 29.8 10.2 9.8 24.2 23.9	0.041 0.050 0.076 0.089 0.031 0.008
373 375 391 406 407	44 (0 8.0 0 10.0 0 2.0 0 17.0 0 18.0	0.0 0.0 0.0 0.0	0.0	KIC KIC KIC KIC	8.5 9.2 2.8 17.0 6.1	0.50 0.91 0.51	21.5 4.3 34.4 30.1 31.0	0.000 0.219 0.050 0.032 0.043

408 418 423 431 441 442 443 444 445 446 453 464 469 475	44 47 47 50 50 50 50 51 51 51	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	19.0 3.0 8.0 6.0 7.0 8.0 9.0 10.0 11.0 18.0 3.0 4.0 9.0 15.0		0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0	KIC KIC KIC KIC KIC KIC KIC KIC	5.5 4.4 6.3 16.0 7.1 7.3 7.6 8.0 8.1 8.3 5.7 4.2 4.8 5.2	0.62 4.72 0.50 0.39 0.40 0.58 0.54 0.62 0.46 0.84 0.92 0.15	27.4 17.1 16.2 16.5 25.6 29.5 29.0 28.8 29.1 30.9 34.7 31.9	0.072 0.069 0.551 0.057 0.029 0.026 0.035 0.026 0.040 0.028 0.046 0.054
479	51	0	19.0	0.0	0.0	KIC	5.8 5.0	0.63 1.40	31.6 29.7	0.038
183	34	0	3.0	0.0	0.0	KIM	5.0	0.46	32.4	0.027
184	34	0	4.0	0.0	0.0	KIM	5.5	0.60	32.0	0.035
185	34	0	5.0	0.0	0.0	KIM	5.8	0.61	31.8	0.036
186	34	0	6.0	0.0	0.0	KIM	5.6	0.46	32.0	0.027
210 241	36 37	0	7.0 15.0	0.0 0.0	0.0	KIM	7.0	0.66	31.2	0.040
242	37	0	16.0	0.0	0.0	KIM KIM	7.5 6.8	0.34	22.5	0.029
273	39	Õ	5.0	0.0	0.0	KIM	6.0	1.67	42.2	0.028
274	39	0	6.0	0.0	0.0	KIM	7.0	1.39	42.9	0.061
283	39	0	15.0	0.0	0.0	KIM	6.8	0.38	42.3	0.017
284	39	0	16.0	0.0	0.0	KIM	6.1	3.50	43.4	0.152
285	39		17.0	0.0	0.0	KIM	5.8	4.45	43.9	0.192
286	39		18.0	0.0	0.0	KIM	5.4	2.43	44.1	0.104
287	39		19.0	0.0	0.0	KIM	5.0	2.60	45.4	0.108
288	39	0	20.0	0.0	0.0	KIM	4.3	1.22	44.7	0.052
297	40	0	5.0	0.0	0.0	KIM	4.7	0.17	43.5	0.007
298 308	40 40	0	6.0 16.0	0.0	0.0	KIM	6.7	0.44	37.6	0.022
309	40	0	17.0	0.0	0.0	KIM KIM	6.2 6.1	0.51	42.6 43.3	0.023
310	40		18.0	0.0	0.0	KIM	6.0	0.00	44.2	0.011
323	41	0	4.0	0.0	0.0	KIM	3.6	0.33	17.5	0.036
324	41	0	5.0	0.0	0.0	KIM	4.2	0.19	33.0	0.011
335	41	0	16.0	0.0	0.0	KIM	4.0		34.2	0.039
336	41		17.0	0.0	0.0	KIM	3.3	0.62	33.7	0.035
374	43	0	9.0	0.0	0.0	KIM	9.0	0.16	28.1	0.011
419	47	0	4.0	0.0	0.0	KIM	5.0	0.62	16.9	0.069
420	47	0	5.0	0.0	0.0	KIM	5.5	0.95	16.8	0.107
421	47	0	6.0	0.0	0.0	KIM	6.2	1.27	16.5	0.146
422 161	47	0	7.0 5.0	0.0	0.0	KIM	6.6	2.90	16.2	0.339
162	33 33	0	6.0	0.0	0.0	KIG KIG	6.9 7.8	0.71	23.0 23.1	0.058
163	33	0	7.0	0.0	0.0	KIG	8.7		22.4	0.084
164	33	Ö	8.0	0.0	0.0	KIG	9.0	0.39	21.7	0.047
165	33	0	9.0	0.0	0.0	KIG	9.3		22.2	0.043
166	33	0	10.0	0.0	0.0	KIG	9.9		22.2	0.057

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167
         33 0 11.0
                      0.0
                            0.0 KIG
                                      9.4 0.24 24.4 0.019
187
                            0.0 KIG
         34
            0
                7.0
                      0.0
                                      6.3 0.26 31.7 0.015
209
            0
                6.0
         36
                      0.0
                            0.0 KIG
                                      6.2 0.46
                                                32.1
                                                      0.027
322
         41
            0
                3.0
                      0.0
                            0.0 KIG
                                      2.9
                                           0.63 17.9
                                                      0.067
346
         42
            0
                4.0
                                      5.2 0.86 30.5 0.053
                      0.0
                            0.0 KIG
347
         42
                                           0.68 27.6 0.046
            0
                5.0
                      0.0
                            0.0 KIG
                                      6.1
348
         42
            0
                6.0
                            0.0 KIG
                      0.0
                                      6.7
                                           0.64 26.4 0.046
349
         42
            0
                7.0
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                            0.0 KIG
                                      7.1
                                           0.44 25.9
                                                      0.032
350
         42
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                8.0
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                            0.0 KIG
                                      7.3
                                           0.26 24.7 0.020
351
         42
            0
                9.0
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                            0.0 KIG
                                      7.3
                                           0.62 29.6 0.040
352
         42
            0
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                            0.0 KIG
                                      7.4
                                           0.84
                                                30.2 0.053
392
         44
            0
                3.0
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                                      3.9 1.01 33.4 0.057
393
         44
            0
                4.0
                      0.0
                            0.0 KIG
                                      5.4 1.08 32.7 0.062
394
            0
                            0.0 KIG
         44
                5.0
                      0.0
                                      6.6 1.27 31.5 0.076
395
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            0
                6.0
                      0.0
                            0.0 KIG
                                      7.4 1.27 31.3 0.077
396
         44
                7.0
            0
                      0.0
                            0.0 KIG
                                      8.1 1.04 26.9 0.073
397
         44
            0
                8.0
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                                      8.1
                                          0.96 29.3 0.062
398
         44
            0
                9.0
                      0.0
                            0.0 KIG
                                      8.6 0.73 29.2 0.047
432
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            0 17.0
                      0.0
                            0.0 KIG
                                      5.6 0.71 16.9 0.079
433
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            0 18.0
                      0.0
                            0.0 KIG
                                      5.0
                                           0.72 16.9 0.080
434
         47
            0
               19.0
                      0.0
                            0.0 KIG
                                      4.3 0.43 17.1 0.048
465
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            0
                5.0
                      0.0
                            0.0 KIG
                                      5.1
                                           0.79 33.2 0.045
466
         51
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                                      5.0
                                           0.71 33.2 0.040
467
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            0
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                                      5.0 0.40 32.9 0.023
468
         51
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                8.0
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                            0.0 KIG
                                      5.2
                                          0.30 32.3 0.018
476
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            0 16.0
                      0.0
                            0.0 KIG
                                      5.7
                                           0.78 31.9 0.046
477
         51 0 17.0
                      0.0
                            0.0 KIG
                                      5.4
                                          0.74 33.0 0.042
478
         51 0 18.0
                      0.0
                            0.0 KIG
                                      5.2 1.01 31.0 0.061
160
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                4.0
                      0.0
                            0.0 KIS
                                      6.3
                                          0.88 23.3 0.071
188
         34
            0
                8.0
                      0.0
                            0.0 KIS
                                          0.31 31.1 0.019
                                      6.7
195
         34
            0 15.0
                      0.0
                                          0.78
                            0.0 KIS
                                      7.0
                                                30.2 0.049
196
         34
            0 16.0
                      0.0
                            0.0 KIS
                                      6.5
                                          0.59 31.2 0.036
197
              17.0
         34
            0
                      0.0
                            0.0 KIS
                                      6.2 0.58
                                                31.4 0.035
232
         37
            0
                6.0
                            0.0 KIS
                      0.0
                                      6.2
                                          0.98 22.7 0.082
233
         37
            0
                7.0
                      0.0
                            0.0 KIS
                                      6.5
                                           0.90 22.1 0.077
234
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                8.0
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                            0.0 KIS
                                      6.7
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271
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276
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                                          1.43 34.0 0.079
281
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                            0.0 KIS
                                      8.0
                                          0.58 40.6 0.027
         40
301
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                      0.0
                            0.0 KIS
                                      7.4
                                           0.51 40.9 0.024
                7.0
326
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                                          0.74
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                                      5.1
                                                  2.3 0.597
368
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                3.0
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                            0.0 KIS
                                           0.49 10.8 0.086
                                      3.3
380
         43
            0 15.0
                      0.0
                            0.0
                                KIS
                                      7.5
                                           0.89
                                                29.5 0.057
381
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         43
                      0.0
                            0.0
                                KIS
                                      6.9
                                           0.53 29.9
                                                      0.034
            0 17.0
382
         43
                      0.0
                            0.0 KIS
                                      6.6
                                          0.00 23.5 0.000
383
         43 0 18.0
                      0.0
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                                      5.3
                                           0.00 25.3 0.000
384
         43 0 19.0
                      0.0
                            0.0 KIS
                                           0.36 28.9 0.024
                                      3.9
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385
                      0.0
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                                      3.6
                                          1.24 33.7 0.069
386
         43
            0 21.0
                      0.0
                            0.0 KIS
                                      3.2
                                          1.06 33.6 0.060
405
         44
            0 16.0
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                            0.0 KIS
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440
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                                    5.8 0.70 30.7 0.043
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454
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                                    4.9 0.39
                                               28.8 0.026
456
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                                    4.2 0.69 31.0 0.042
457
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                                    3.0 0.77 34.1 0.043
474
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                          0.0 KIS
                                    5.4 0.42 31.7 0.025
181
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            0
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                          0.0 MJK
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356
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                          0.0 MJK
                                    7.5 0.41 14.7 0.053
357
         42 0 15.0
                     0.0
                          0.0 MJK
                                    6.8 0.86 30.4 0.053
358
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                     0.0
                          0.0 MJK
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         42 0 17.0
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                          0.0 MJK
360
         42 0 18.0
                          0.0 MJK
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361
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                     0.0
                           0.0 MJK
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 48
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 49
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                     0.0
 90
         28
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                                    0.7 1.12 36.7 0.058
 91
         28 0
               7.0
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                          0.0 LIS
                                    0.7 0.87 36.8 0.045
 17
         19
            0 11.0
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                          0.0 LCS
                                    0.4 0.88 49.2 0.034
 32
         22 0 11.0
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 33
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171
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                          0.0 LIG
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  4
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        19
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                                    0.8 0.86 49.8 0.033
 34
         22 0 15.0
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 50
           0 16.0
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100
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                                    1.1 0.22 36.2 0.011
101
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102
         28
           0 18.0
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                                    0.3 0.19 37.1 0.010
105
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            ï
               6.0
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                                    1.6 1.99 48.5 0.078
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106
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107
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108
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                                    1.7 1.30 48.4 0.051
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110
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                                     1.5 1.08 48.7 0.042
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                           0.0 MMA
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112
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            0 13.0
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                           0.0 MMA
                                     0.9 0.60 49.6 0.023
113
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                           0.0 MMA
                                     0.5 0.35 50.2 0.013
129
         30
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                           0.0 MMA
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137
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                           0.0 MMA
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142
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145
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                                     0.3 0.33 50.5 0.012
146
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                                     0.3 0.26 50.5 0.010
147
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                           0.0 MMA
                                     0.3 0.41 50.5 0.015
152
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                2.0
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                           0.0 MMA
                                     0.3 0.16 19.2 0.016
153
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                3.0
                     0.0
                           0.0 MMA
                                     0.4 0.23 19.2 0.023
154
         32
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               4.0
                     0.0
                           0.0 MMA
                                    0.4 0.18 19.2 0.018
155
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               5.0
                     0.0
                           0.0 MMA
                                     0.5 0.35 19.1 0.035
158
         33 0
                2.0
                     0.0
                           0.0 MMA
                                     4.0 1.00 26.0 0.073
177
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                     0.0
                           0.0 MMA
                                         0.38 25.7 0.028
                                     4.5
178
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              22.0
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                                    1.3 0.73 28.3 0.049
223
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                                    1.7 0.56 35.3 0.030
224
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                     0.0
                           0.0 MMA
                                    1.2 1.36 35.9 0.072
228
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            0
               2.0
                     0.0
                                         0.80 24.9 0.061
                          0.0 MMA
                                     3.5
247
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            0 21.0
                          0.0 MMA
                     0.0
                                    0.9 0.61 27.0 0.043
           0 19.0
266
         38
                     0.0
                          0.0 MMA
                                    1.2 2.87 49.2 0.110
270
            0
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               2.0
                     0.0
                           0.0 MMA
                                    3.0 1.49 45.9 0.061
291
         39
            0 24.0
                     0.0
                           0.0 MMA
                                     1.8 2.22 48.3 0.087
294
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            0
               2.0
                     0.0
                           0.0 MMA
                                     3.2 0.65 19.1 0.064
314
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            0
              22.0
                     0.0
                           0.0 MMA
                                     2.3 0.52 47.5 0.021
387
         43 0
              22.0
                     0.0
                           0.0 MMA
                                    1.6 1.49 34.8 0.081
437
         50 0
                2.0
                     0.0
                           0.0 MMA
                                    3.0 1.81 33.7 0.101
458
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              23.0
                     0.0
                           0.0 MMA
                                    1.9 0.80 34.8 0.043
         52 0
484
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                     0.0
                          0.0 MMA
                                    0.8 0.46 26.4 0.033
485
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               2.0
                     0.0
                          0.0 MMA
                                     0.9 0.40 26.3 0.029
436
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               3.0
                     0.0
                          0.0 MMA
                                     0.9 0.46 26.3 0.033
487
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               4.0
                     0.0
                          0.0 MMA
                                     0.9 0.45 26.3 0.032
         52 0
488
               5.0
                          0.0 MMA
                     0.0
                                    1.0 0.53 26.2 0.038
489
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               6.0
                     0.0
                          0.0 MMA
                                    1.0 0.74 26.4 0.053
490
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               7.0
                     0.0
                          0.0 MMA
                                    1.1 0.66 26.2 0.048
491
         52 0
               8.0
                     0.0
                                     1.3 0.55 26.0 0.040
                          0.0 MMA
492
         52
            0
               9.0
                     0.0
                          0.0 MMA
                                    1.1 0.52 26.2 0.037
493
         52 0 10.0
                     0.0
                          0.0 MMA
                                     1.0 0.62 26.1 0.045
494
           0 11.0
         52
                     0.0
                          0.0 MMA
                                    1.0 0.95 26.1 0.069
498
         52 0 15.0
                     0.0
                          0.0 MMA
                                     0.8 0.51 26.4 0.037
499
        52 0 16.0
                     0.0
                          0.0 MMA
                                    0.5 0.48 26.6 0.034
500
        52 0 17.0
                     0.0
                          0.0 MMA
                                    0.5 0.36 26.6 0.026
```

14 194 367 19 20 21 22 24 55	34 (43 (20 1) 20 (20 (20 (25 (25 (25 (25 (25 (25 (25 (25 (25 (25	2.0 1.0 3.0 5.0 7.0 11.0 2.0 3.0	0.0 0.0 0.0 0.0 0.0 0.0 0.0	0.0 0.0 0.0 0.0 0.0 0.0 0.0	MJS MJS MJS MJO MLD MDJM DJM DJM DJM DJM DJM DJM	1.0 7.9 3.0 0.2 0.2 0.3 0.5 0.2	0.62 0.64 0.39 0.49 0.59 0.58 0.45 0.96	49.5 29.5 11.1 50.9 50.9 50.7 49.8 50.9 50.5 50.9	0.029 0.040 0.109 0.014 0.018 0.022 0.022 0.017 0.036 0.032
57 58 59 60 61	25 (25 (25 (25 (25 (5.0 6.0 7.0	0.0 0.0 0.0	0.0	DJM DJM DJM	0.3 0.3 0.3	0.50 0.47 0.72 0.59	50.7 50.7 50.7	0.019 0.018 0.027 0.022
62 63 64 65	25 (25 (25 (25 (9.0 10.0 11.0	0.0 0.0 0.0 0.0	0.0 0.0 0.0 0.0	DJM DJM DJM DJM	0.3 0.3 0.4 0.3	0.62 0.63 0.34 0.46	50.7 50.7 50.3 50.7 50.9	0.023 0.023 0.013 0.017 0.012
66 67 68 99	25 0 25 0 25 0 28 0	13.0 14.0 15.0 15.0	0.0 0.0 0.0	0.0 0.0 0.0	DJM DJM DJM DJM	0.4 0.5 0.6 0.6	0.42 0.38 0.68 0.30	50.4 50.2 48.5 36.7	0.012 0.016 0.014 0.027 0.015
114 118 119 120	29 0 30 0 30 0	1.0 2.0 3.0	0.0	0.0 0.0 0.0	DJM DJM DJM	0.6 0.8 0.7 0.6	0.32 0.76 0.53 0.48	50.0 49.5 49.7 49.9	0.012 0.029 0.020 0.018
121 122 123 124 125	30 0 30 0 30 0 30 0	5.0 6.0 7.0	0.0 0.0 0.0 0.0	0.0 0.0 0.0 0.0	DJM DJM DJM DJM	0.6 0.5 0.5 0.6	1.30 1.46 1.28 1.27	49.9 50.2 50.2 49.9	0.049 0.055 0.048 0.048
126 127 128 131	30 0 30 0 30 0	9.0 10.0 11.0	0.0	0.0	DJM DJM DJM DJM	0.6 0.6 0.7	1.37 2.02 2.44 0.50	49.7 49.9 49.9 49.7	0.044 0.052 0.076 0.093 0.019
132 133 135 148	30 0 30 0 31 1 31 0	15.0 16.0 1.0 15.0	0.0 0.0 0.0	0.0	MLD MLD MLC	0.6 0.4 0.2 0.6	0.47 0.85 0.43 0.45	50.0 50.4 50.9	0.018 0.032 0.016 0.017
227 35 72 304	40 0	1.0 1.0 12.0	0.0 0.0 0.0	0.0 0.0 0.0	MJF MJF	1.0	0.56 0.76 0.39	50.0	0.065 0.022 0.029 0.284
43 103 149 317 159		19.0 16.0 25.0	0.0	0.0	MJD DUM DUM	0.2 0.1 0.3	0.95	37.3 51.2 50.7	0.005 0.023 0.014 0.035
190 192 *	33 0 34 0 34 0			0.0	MMB	8.3		25.0 29.2 6.7	0.066 0.034 0.133

25567890 2567890 2627893 3328978** 444890 44773 1367890 12538 33333333333333333333333333333333333	42 0 42 0	7.0 9.0 10.0 11.0 12.0 13.0 14.0 10.0 11.0 12.0 13.0 14.0 12.0 13.0 14.0 11.0 13.0 14.0 12.0	0.0			8.2 8.1 1.0 7.9 3.0 1.4 1.9 3.0 2.35 6.8 4.7 4.4 4.1 2.8 7.8 2.2	0.30 0.47 0.47 0.47 0.42 0.42 0.42 0.42 0.42 0.42 0.54	44.8 9.3 45.0 9.3 45.0 45.0 45.0 62.4 4.4 9.2 2.4 3.3 2.9 3.0 3.0 3.0 3.0 3.0 3.0 3.0 3.0	0.013 0.0043 0.093 0.093 0.057 0.045 0.025 0.045 0.045 0.025 0.045 0.055 0
199 200 221 222 245 265	34 0 36 0 36 0 37 0	19.0 20.0 18.0 19.0 19.0	0.0 0.0 0.0 0.0 0.0	0.0 0.0 0.0 0.0	PLG PLG PLG PLG PLG PLG		1.06 0.52	33.3 33.7 30.7 33.1 25.4 48.8	0.098 0.060 0.039

```
339
        41 0 20.0
                     0.0
                          0.0 PLG
                                    1.2 1.47 36.4 0.076
343
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                          0.0 PLG
              1.0
                     0.0
                                    2.0 0.72 34.5 0.039
480
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                     0.0
                          0.0 PLG
                                    4.3 1.46 32.9 0.084
481
         51
            0 21.0
                     0.0
                          0.0 PLG
                                    2.8 0.55 17.8 0.058
201
         34
            0 21.0
                     0.0
                          0.0 PJM
                                    1.4 1.02 35.0 0.055
246
         37 0 20.0
                     0.0
                          0.0 PJM
                                    1.9 0.47 26.2 0.034
15
        19 0
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                     0.0
                          0.0 SIG
                                    0.7 0.70 50.2 0.026
 28
         21 0
               6.0
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                          0.0 SIG
                                    3.0 0.59 48.3 0.023
 42
              14.0
         23
            0
                     0.0
                          0.0 SIG
                                    3.9 0.62 48.1 0.024
 77
         26
            0
                                     3.2 1.32 48.3 0.052
               6.0
                     0.0
                          0.0 SIG
 78
         26
           0
               7.0
                     0.0
                          0.0 SIG
                                    3.8 1.96 47.8 0.077
 95
           0 11.0
         28
                     0.0
                          0.0 SIG
                                    1.0 0.60 19.0 0.060
 96
         28
           0 12.0
                     0.0
                          0.0 SIG
                                    1.0 0.36 19.0 0.036
 97
         28 0 13.0
                     0.0
                          0.0 SIG
                                    1.1 0.13 19.0 0.013
                          0.0 SIG
251
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               2.0
                     0.0
                                    0.8 0.42 18.9 0.042
               3.0
438
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                     0.0
                          0.0 SIG
                                    3.3 1.06 17.8 0.113
 36
         23 0
               3.0
                     0.0
                          0.0 SIS
                                    3.2 0.81 48.3 0.032
 40
         23 0 10.0
                     0.0
                          0.0 SIS
                                    4.2 0.96 43.4 0.042
 75
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                     0.0
                          0.0 SIS
                                    2.4 0.10 48.8 0.004
 76
         26
           0
               5.0
                     0.0
                          0.0 SIS
                                    2.8 0.41 48.7 0.016
 79
         26 0
               8.0
                     0.0
                          0.0 SIS
                                    3.9 0.66 47.7 0.026
 80
         26 0
               9.0
                     0.0
                          0.0 SIS
                                    3.8 0.91 47.9 0.036
312
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                          0.0 SIS
                                    5.4 0.00 39.2 0.000
313
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                                    5.0 0.36 40.5 0.017
                     0.0
                          0.0 SIS
412
        44 0 23.0
                     0.0
                          0.0 SIS
                                    2.8 1.85
                                                6.7 0.519
413
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                     0.0
                                    1.8 0.53
                          0.0 SIS
                                                6.8 0.147
424
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                     0.0
                          0.0 SIS
                                    6.0 3.08
                                                6.0 0.963
425
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                     0.0
                          0.0 SIS
                                    6.0 1.04
                                                6.1 0.324
                                    4.9 0.84 33.3 0.048
439
         50 0
               4.0
                     0.0
                          0.0 SIS
 73
         26
           0
               2.0
                     0.0
                          0.0 SIL
                                    1.0 0.14 49.9 0.005
 74
         26
           0
               3.0
                     0.0
                          0.0 SIL
                                    1.8 0.15 49.1 0.006
81
         26 0 10.0
                     0.0
                          0.0 SIL
                                    3.7 0.68 48.0 0.027
         26 0 11.0
82
                     0.0
                          0.0 SIL
                                    3.7 1.17 47.9 0.046
83
         26 0 12.0
                     0.0
                          0.0 SIL
                                    3.6 1.28 48.0 0.050
14
        19
            0
               5.0
                     0.0
                                    1.0 0.76 49.5 0.029
                          0.0 MJS
194
         34 0 14.0
                     0.0
                                    7.9 0.62 29.5 0.040
                          0.0 MJS
367
         43 0
               2.0
                     0.0
                          0.0 MJS
                                    3.0 0.64 11.1 0.109
409
        44 0 20.0
                     0.0
                          0.0 KJS
                                    5.0 0.75 28.5 0.050
410
        44 0 21.0
                     0.0
                          0.0 KJS
                                    4.3 1.18 33.4 0.067
                     0.0
411
        44 0 22.0
                          0.0 KJS
                                    3.3 1.82 17.7 0.195
230
         37 0
               4.0
                     0.0
                          0.0 VES
                                    5.7 1.17 24.1 0.092
231
         37 0
               5.0
                     0.0
                                    6.0 1.17 19.5 0.113
                          0.0 VEG
```

264	38	0	16.0	0.0	0.0	WAT	3.7	0.77	47.8	0.030
289	39	0	22.0	0.0	0.0	WAT	3.2	2.32	48.3	0.091
1	17	1	0.0	0.0	0.0	WJD	0.0	0.08	52.1	0.003
53	25	1	0.0	0.0	0.0	WJD	0.0	0.58	52.1	0.021
71	26	1	0.0	0.0	0.0	WJD	0.0	0.72	52.1	0.026
84	28	1	0.0	0.0	0.0	WJD	0.0	0.75	38.2	0.037
117	. 30	1	0.0	0.0	0.0	WJD	0.0	0.85	52.1	0.031
150	32	1	0.0	0.0	0.0	WJD	0.0	0.08	19.8	0.008
156	. 33	1	0.0	0.0	0.0	WJD	0.0	0.71	30.2	0.044
180	34	1	0.0	0.0	0.0	WJD	0.0	1.29	38.2	0.064
203	36	1	0.0	0.0	0.0	WJD	0.0	0.73	38.2	0.036
249	38	1	0.0	0.0	0.0	WJD	0.0	1.11	19.8	0.106
268	39	1	0.0	0.0	0.0	WJD	0.0	0.39	52.1	0.014
292	40	1	0.0	0.0	0.0	WJD	0.0	0.47	21.4	0.042
319	41	1	0.0	0.0	0.0	WJD	0.0	1.14	19.8	0.109
342	42	1	0.0	0.0	0.0	WJD	0.0	0.84	38.2	0.042
365	43	1	0.0	0.0	0.0	WJD	0.0	0.73	12.9	0.107
435	50	1	0.0	0.0	0.0	WJD	0.0	0.20	38.2	0.010
11	18	0	15.0	0.0	0.0	DJW	0.0	0.38	52.1	0.014
5 2	24	0	18.0	0.0	0.0	DJW	0.0	0.72	52.1	0.026
70	25	0	17.0	0.0	0.0	DJW	0.0	0.88	52.1	0.032
104	28	0	20.0	0.0	0.0	DJW	0.0	0.22	38.2	0.011
116	29	0	17.0	0.0	0.0	DJW	0.0	0.18	52.1	0.007
134	30	0	17.0	0.0	0.0	DJW	0.0	1.12	52.1	0.041
179	33	0	23.0	0.0	0.0	DJW	0.0	0.39	30.2	0.024
202	34	0	22.0	0.0	0.0	DJW	0.0	0.69	38.2	0.034
226	36	0	23.0	0.0	0.0	DJW	0.0	0.46	38.2	0.023
248	37	0	22.0	0.0	0.0	DJW	0.0	0.25	28.3	0.017
318	4 0	0	26.0	0.0	0.0	DJW	0.0	0.70	52.1	0.025
341	41	0	22.0	0.0	0.0	DJW	0.0	1.11	19.8	0.106
459	5 0	0	24.0	0.0	0.0	DJW	0.0	0.45	38.2	0.022
460	51	1	0.0	0.0	0.0	DJW	0.0	0.69	19.8	0.066
483	5 2	1	0.0	0.0	0.0	DJM	0.0	0.14	28.3	0.009
502	5 3	1	0.0	0.0	0.0	DJW	0.0	0.35	19.8	0.033
4 3	23	0	16.0	0.0	0.0	WJM	1.0	0.12	49.4	0.005