Rate Constants for H-atom Transfer Reactions by the BEBO Method

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A detailed discussion of the calculation of rate constants for hydrogen atom transfer reactions based on the BEBO method is presented. Linear transition state models are used. A computer program using this method for determining rate constants is provided.

Key words: absolute chemical rate; BEBO activation energy; bond-order-bond-energy; chemical rate constants; computer program; hydrogen atom transfer.

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1. Introduction

The bond-energy-bond-order (BEBO) method is a procedure for calculating the activation energies of hydrogen transfer reactions from bond energies. When combined with absolute rate theory, it also yields values for the rate constants. It was formulated over 10 years ago by Johnston and Parr [1], and has since been applied with considerable success to the calculation of a large number of activation energies. Less frequently, it has been used to evaluate rate constants. Although the details of the BEBO method itself have

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1 Figures in brackets indicate literature references at the end of this paper.

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been published by Johnston [2], this aspect represents only a relatively small part of a rate constant
calculation. The purpose of this report is to give a detailed account, not only of the BEBO method and its
theoretical background, but also of the absolute rate theory portion of the calculation. In addition, instruc-
tions are provided for the use of a computer program which calculates rate constants based on the BEBO
method. The discussion is limited to linear transition state models.

2. Theory

2.1. Absolute Rate Theory & Transition State Model for BEBO Calculations

For a bimolecular reaction, \( A + B \rightarrow [AB]^* \rightarrow \text{products} \), absolute rate theory utilizes the concept of a
molecular complex made up of the two reactants. This complex is assumed to be in equilibrium with these
reactants. The resulting expression for the classical rate constant \( k_{cl} \) is

\[
k_{cl} = \frac{kT}{h} \frac{Q_{cl}^* e^{-\nu* kT}}{Q_d^* Q_e^*} \tag{1}
\]

where \( k \) is the Boltzmann constant, \( T \) is the absolute temperature, \( h \) is Planck's constant, \( Q_d^* \) and \( Q_e^* \) are the
classical partition functions per unit volume for reactants \( A \) and \( B \), \( Q_{cl}^* \) is the classical partition function per
unit volume for the complex, and \( \nu^* \) is the potential energy of the complex relative to that of the reactants.
The complex contains one unstable vibrational mode whose evolution brings about its dissociation into
product fragments. The partition function \( Q_{cl}^* \) is evaluated with this mode missing. A detailed derivation of
eq (1) which explains all its inherent assumptions has been given by Mahan [3]. Quantum mechanical correc-
tions to the partition functions at room temperature and above need be applied only to vibrational factors.
For a particular vibration of frequency \( \nu \), the quantum correction \( \Gamma_i \) is given by the expression

\[
\Gamma_i = \frac{u_i/2}{\sinh (u_i/2)} \quad \text{where} \quad u_i = \hbar \nu/kT \tag{2}
\]

We assume that all vibrational modes are independent so that the total quantum correction for a particular
species is simply the product of terms given by \( \text{eq (2)} \), one for each vibrational mode. There is also a quan-
tum correction to the unstable vibrational mode of the complex which we denote by \( \Gamma^* \). This results from
the effect of quantum mechanical tunneling through the potential barrier between reactants and products.
It will be considered in detail in section 2.5. Applying these quantum corrections to eq (1) yields the rate
expression

\[
k = \frac{kT}{h} \frac{Q_{cl}^* \{\pi \Gamma^*_i \} \Gamma^* e^{-\nu^* kT}}{Q_d^* \{\pi \Gamma^*_i \} Q_e^* \{\pi \Gamma^*_i \}} \tag{3}
\]

The general class of reactions we are considering has the form

\[
A-H + B^* \rightarrow A-H-B \rightarrow A^* + H-B \tag{4}
\]

Radical \( B^* \) abstracts a hydrogen atom attached to \( A \), the net result being the transfer of \( H \) from \( A \) to \( B \). For
this system, we take the most general transition state to be linear, having up to 5 mass points. Its structure
and the notation which we shall use are shown in figure 1a. There can be up to four internuclear distances,
\( R_a, R_b, R_c, \) and \( R_d \). The bonds associated with \( R_a \) and \( R_b \) will be assumed to be rigid. (The two vibrational
modes involving these bonds will have infinite frequencies and need not be formally included in the calcula-
tions.) Thus, there are only two vibrational stretching modes to be considered for this molecule, one of which
will be unstable. These modes arise from the stretching of the two central bonds \( b \) and \( c \) which are shown by
dotted lines to indicate their unstable character. Of the five possible masses, \( M_3 \) will normally be that of the
hydrogen atom; the other masses will be assigned values in the manner described below. The three angles
\( \Psi_2, \Psi_3, \) and \( \Psi_4 \) are defined by the bonds \( (a,b), (b,c), \) and \( (c,d) \) in the plane of the figure while the primed sym-
bols denote the corresponding angles in the plane perpendicular to the figure. Changes in these angles from
$180^\circ$ give rise to three doubly degenerate bending vibrations. To calculate the frequencies needed in eq (3), we require values for the two stretching force constants associated with bonds b and c, and three bending force constants arising from the three bond angles. As we shall see, these values can be generated by the BEBO process.

Within the framework of the transition state structure shown in figure 1a, it is possible to include all types of reactions implied by eq (4) by considering four cases; one having a 3 point transition state, two having 4 point states, and one having a 5 point state. These four cases are shown in figure 1b. In this figure, the subscript s appearing on the internuclear distances and force constants denote equilibrium values found in reactants or products. Because bonds a and d are assumed to be rigid, their bond distances will always be denoted by the single symbols $R_a$ and $R_d$, respectively. The bond distance between $M_2$ and $M_3$ goes from $R_b$ to $R_b$ in the transition state, while that between $M_3$ and $M_4$ goes from $R_0$ to $R_0$ in the transition state. In the transition state, the force constant $F_{bs}$ is modified and combined with that of the newly formed bond between $M_3$ and $M_4$ to produce two force constants $F_{b}$ and $F_{c}$. $F_b$ corresponds to the stable symmetric stretch and $F_c$ to the unstable asymmetric stretch. In cases IVa and V, the bending force constant $F_{b2}$ becomes $F_{b2}$ in the complex. The newly formed bond angle made by $M_2$, $M_3$, and $M_4$ leads to the force constant $F_{b2}$ in all cases. Finally, in cases IVb and V, we also have an additional bending force constant $F_{b4}$ which goes to $F_{b4}$ in the second product. The force constants associated with the out-of-plane bends are not shown since they are the same as the in-plane constants.

The way I have chosen to assign values to the mass points is somewhat arbitrary and is best explained by an example. Consider the reaction

$$\text{CH}_3\cdot\text{CH}_2\cdot\text{H} + \text{CH}_3 \rightarrow \text{CH}_3\cdot\text{CH}_2\cdot\text{H} = \text{CH}_3 \rightarrow \text{CH}_3\cdot\text{CH}_2\cdot + \text{H} = \text{CH}_3$$

Species A B C D

which is the abstraction of hydrogen from ethane by methyl radicals. The masses are assigned according to the following rules:

1) The mass of the transferred H is always assigned to $M_3$; therefore $M_3 = 1.008$ atomic mass units (a.m.u.).
2) The mass of the atom joined to the transferred H in reactant A is assigned to $M_2$; in this case $M_2 = 12.011$ a.m.u.
3) The masses of all the remaining atoms in A are added and assigned to $M_1$; thus in this example $M_1 = 17.051$ a.m.u.
4) The mass of the atom joined to the transferred H in the product D is assigned to $M_4$; here $M_4 = 12.011$ a.m.u.
5) The masses of all the remaining atoms in D are added and assigned to $M_5$; thus $M_5 = 3.024$ a.m.u. in this example.

Different models for the transition state, and different ways of arranging the masses in linear models have been explored in a limited way by Johnston [4] and by Sharp & Johnston [5]. They did find significant differences between various options. Presumably, complete vibrational analyses of the reactant and complex would yield more accurate rate constants than the linear models outlined above. Unfortunately, complete analyses are extremely complex even for fairly small molecules, and the ability to program the calculations
in a general manner would be lost by such an approach. Also, it is unlikely that all of the force constant values required would be available for a complete analysis. In view of the crudity of the rest of the calculation, it is unnecessary to strive for high accuracy in the vibrational analysis. Intuitively, one expects that the major features of these reactions are controlled by the nature of the atoms adjacent to the H atom being transferred, with the effects from the remainder of the molecule appearing in the bond energy values. If this is the case, then the linear models should at least be able to match trends within homologous series.

So far, we have seen in this section that evaluation of rate constants by the use of eq (3), based on the linear models shown in figure 1b, requires a knowledge of the potential energy $V^*$ of the complex, two stretching force constants, and from one to three bending force constants. The potential energy of all of these linear models could, if it were known, be shown on a 2-dimensional contour diagram like that shown in figure 2 where the independent variables are the bond distances $R_s$ and $R_c$. The required value of the potential energy $V$ is that at the saddle point position shown by the asterisk. For a region close to the saddle point,
FIGURE 2. Typical potential energy diagram for H atom exchange reaction. The position of the saddle point is shown by the asterisk. The direction $q$ is that in which the potential energy decreases most rapidly. The direction $\sigma$ is perpendicular to the $q$ direction.

it is customary to assume that the first derivatives of $V$ with respect to $R_b$ and $R_c$ are negligible, and that the potential energy can be approximated by a power series containing only quadratic terms. Thus, for small displacements from the saddle point, we have

$$2\delta V = F_{11}(\delta R_b)^2 + 2F_{12}(\delta R_b)(\delta R_c) + F_{22}(\delta R_c)^2$$

(5a)

where $F_{11} = \frac{\partial^2 V}{\partial R_b^2}$, $F_{12} = \frac{\partial^2 V}{\partial R_b \partial R_c}$, $F_{22} = \frac{\partial^2 V}{\partial R_c^2}$.

These derivatives are evaluated at the saddle point, and are, by definition, the stretching force constants of the complex. In matrix notation, this equation is

$$2\delta V = (\delta R)^t F_\sigma (\delta R)$$

(5b)

where $F_\sigma = \begin{bmatrix} F_{11} & F_{12} \\ F_{21} & F_{22} \end{bmatrix}$ and $\delta R = \begin{bmatrix} \delta R_b \\ \delta R_c \end{bmatrix}$.

This is the force constant matrix that will be used to calculate the vibrational stretching frequencies.

Starting at the saddle point, suppose we move in the direction in which $V$ decreases most rapidly; call this the $q$ direction, and let $\sigma$ denote the direction perpendicular to $q$. These directions define a rotated set of cartesian coordinates which we assume makes an angle $\alpha$ with the $R_b$ axis; (positive $\alpha$ is measured in the counter-clockwise direction). The transformation between the two sets of coordinates is given by the equation

$$R = \begin{bmatrix} R_b \\ R_c \end{bmatrix} = \begin{bmatrix} \cos \alpha & -\sin \alpha \\ \sin \alpha & \cos \alpha \end{bmatrix} \begin{bmatrix} q \\ \sigma \end{bmatrix} = U P$$

(6)

the matrix $U$ can now be used to express changes in $V$ at the saddle point in terms of changes in $q$ and $\sigma$ instead of $R_b$ and $R_c$. Thus, eq (5b) becomes

$$2\delta V = (\delta R)^t F_\sigma (\delta R) = (U\delta P)^t F_\sigma (U\delta P) = (\delta P)^t (U^t F_\sigma U)(\delta P)$$

(7)
The matrix $U^tF,U$ has the elements

$$(U^tF,U)_{11} = F_{11}\cos^2\alpha + 2F_{12}\cos\alpha\sin\alpha + F_{22}\sin^2\alpha$$

$$(U^tF,U)_{12} = (U^tF,U)_{21} = (F_{22} - F_{11})\cos\alpha\sin\alpha + F_{12}\cos^2\alpha - \sin^2\alpha$$

$$(U^tF,U)_{22} = F_{11}\sin^2\alpha - 2F_{12}\cos\alpha\sin\alpha + F_{22}\cos^2\alpha$$

As we shall see in the next section, the BEBO method provides values for the second derivatives of $V$ (i.e., the force constants) in the $\phi$ and $\alpha$ directions. This will allow us to evaluate the matrix $U^tF,U$. The stretching force constant matrix $F_r$, can then be obtained by inverting the transformation given by eq (6).

In this section I have presented a formula (eq (3)) for the rate constant and outlined the factors required to evaluate it. The details of the BEBO method will be given next. It will provide values for $V^{''}$ and all of the necessary force constants, both the stretching and the bending ones.

2. BEBO Method

The BEBO method is based on the concept of bond order. In the reactants the bond $b$ of figure 1a is said to have a bond order of unity, while in the products, its bond order is zero. The reverse of this situation holds for bond $c$. BEBO assumes that during the reaction, the total bond order of the two bonds is conserved; if $n$ is the order of bond $b$, and $m$ of bond $c$, then we have always $n + m = 1$. This is the basic assumption of the method. One bond is breaking at the same time that the other is forming. To apply this conservation condition it is necessary to relate the energies and lengths of bonds $b$ and $c$ to their bond orders $n$ and $m$.

For the relationship between order and length, Pauling [6] proposed the formula

$$R_a = R_s - \lambda \ln(n)$$

where $R_s$ is the length of the bond which is considered to be representative of a single bond between the two elements of interest. The parameter $\lambda$ is taken to have the same value for all element pairs. A plot of bond length versus the logarithm of the bond order is shown in figure 3 for certain element pairs. The data were

![Figure 3](image_url)

**Figure 3.** Plot of $R_a = R_s - \lambda \ln(n)$; bond distance versus the natural logarithm of the bond order for selected element pairs.
obtained from table 4–3 of Johnston’s book [2]. Values of λ obtained from least squares fits are given in the figure for the different bonds. The constancy of λ does not seem particularly striking. Pauling chose a value of 0.26 for λ; he writes, "This equation, which is based upon the study of interatomic distances for non-resonating and resonating covalent bonds in simple non-metallic substances of known structure, is found to agree reasonably well with those data for metallic crystals which are suited to a check on its validity, and its use permits a penetrating analysis of the structure of metals and intermetallic compounds to be made. There is some evidence that the constant . . . varies with the kind of atom and with the type of bond; but the evidence is not sufficiently extensive to lead to the determination of the nature of this variation." Certainly Pauling’s value doesn’t appear to have been based very heavily on the data in figure 3 since none of these λ values are close to 0.26. Although 0.26 can hardly be construed as universal, it has nevertheless been the value used for most BEBO calculations. There appears to be no reason why a different value shouldn’t be used if it gave better results.

Consider next the dependence of bond energy on bond order. Johnston [4] proposed the following relationship between the two quantities

\[ E_n = E_r n^p \]  

(10)

where \( E_r \) is the bond energy of a single bond and is analogous to \( R_r \) of eq (9). Note that this energy is the electronic dissociation energy of the bond in question; the zero point energy is not meant to be included in \( E_r \). Plots of \( \ln(E) \) versus \( \ln(n) \) are shown in figure 4 for the same bonds used in figure 3. The data are again from table 4–3 of Johnston [2]. We see that \( p \) depends on the kind of atoms in the bond. If more than one bond type occurs for a pair of atoms, then it is possible to extract values for \( p \) from plots like figure 4 provided we are not unduly bothered by a lack of linearity. When only a single bond type exists, then some other method must be devised. Actually, since we are interested in \( E_r \) and \( R_r \) for bond orders less than unity, even if multiple bonds were available for a plot like figure 4, some method of extrapolating to zero \( n \) would be necessary. Johnston [2], inspired by Badger’s rule for force constants, has devised a way. Let us first eliminate \( n \) between eqs (9) and (10); this yields

\[ \ln(E_r/E_n) = (p/\lambda)(R_r - R_n) \]  

(11)

This expression is analogous to Badger’s rule (see Herschbach & Laurie [7]), which is a universal empirical
relation having the form \( r = a_i - b_i \log(f) \), where \( r \) is the bond distance, \( f \) its force constant, \( a_i \) and \( b_i \) are constants, and \( i \) and \( j \) are the numbers of the rows in the periodic table in which the bound atoms are located. Johnston [8] found that plots of \( \log(f) \) versus \( r \) extrapolated very nicely to two-atom Lennard-Jones noble gas clusters. For clusters having Lennard-Jones parameters \( \sigma \) and \( \epsilon/\kappa \), the "bond" distance is \( 2^p \sigma \) and the "force" constant is \( 40.06(\epsilon/\kappa)^2 \). He then examined plots of \( \ln(E_a) \) versus \( R_n \) to see if a comparable extrapolation would be possible. The results are shown in figure 5. The data are mostly from Johnston [2], tables 4–3 and 4–1. Values of \( E_a \) and \( R_n \) for the He-Ne cluster were taken from Gilliom [9]. The energies for the bonds examined in figures 3 and 4 are supposed to extrapolate to the Ne-Ne cluster. The lines shown were drawn to connect the corresponding single bonds with this cluster. Points corresponding to multiple bonds fall more or less in the general direction of these lines. The assumption made in BEBO is that such an extrapolation adequately represents the bond energies for \( n < 1 \). Therefore, if we have a bond A-H, where A is some atom in the first row of the periodic table connected to an H atom, and \( R_n \) and \( E_a \) are its bond length and energy, then if this bond were perturbed in some fashion so that its bond length were greater than \( R_n \), then its bond energy would fall on the line drawn between the A-H and He-Ne points. Bonds involving atoms A from other rows of the periodic table will extrapolate to the appropriate rare gas-helium cluster. The slope of the line joining A-H to the cluster is, from eq (11), \(-p/\lambda\). Since the value of \( \lambda \) has been chosen, we have a way of calculating \( p \) for the A-H bond of interest. Formally, in this case,

\[
p = \frac{\lambda}{R_n - R_{Ne-Ne}} \ln(E_{Ne-Ne}/E_a)
\]

The parameter \( p \) thus depends on \( \lambda \), the bond energy and internuclear distance of A-H, and the interaction parameters for the appropriate rare gas cluster.

We have now almost all of the information needed for the BEBO calculations. Consider a triatomic complex A-H-B; there are three interactions; two between H and the atoms A and B considered above, and the interaction between A and B themselves. If H is to form stable bonds its electron spin must be opposite each of the spins of A and B. Consequently, A and B will have parallel spins and must repel each other. Johnston uses one half the value of the Sato [10] triplet function to represent this interaction. He uses the modified function because it more closely approximates the calculated H-H triplet interaction. This function has the form
\[ V_r = E_{\alpha} E_\beta (1 + E) \]  
(13)

where \( E = \frac{1}{2} e^{-\beta R} \), \( \Delta R = R_r - R_a = R_s + R_c - R_u \). \( E_{\alpha} \) is the electronic dissociation energy, \( \Delta R \) the equilibrium internuclear distance, and \( \beta \) the Morse parameter (see Herzberg [11] p. 101) of the ground state of the diatomic molecule made up of \( A \) and \( B \). Values of these parameters for a number of such atom pairs are given in Table 11-1 of Johnston [2]. \( \Delta R \) is the difference between the actual distance \( R \) between \( A \) and \( B \) in the complex and the equilibrium distance \( R_e \) it would have in the diatomic molecule. It is worth pointing out that many people use \( E_{\alpha} \) as an adjustable parameter to fit the BEBO calculations to their experimental data. Other forms of the triplet function have been used and are discussed briefly in the Appendix. \( V_r \) can be expressed as a function of \( n \), the bond order of the \( b \) bond, through the conservation condition \( n + m = 1 \), and through eq (9) which gives the distances \( R_b \) and \( R_c \) in terms of \( n \) and \( m \).

We are now able to give the BEBO expression for the energy of the complex in terms of the bond order \( n \). The energy is assumed to be given by

\[ V(n) = E_{\alpha b} - E_{\alpha c} n^p - E_{\alpha c} m^q + V'(n) = E_{\alpha b}(1 - n^p) - E_{\alpha c}(1 - n)^q + V'(n) \]  
(14)

\( E_{\alpha b} \) and \( E_{\alpha c} \) are the single bond energies (electronic) for bonds \( b \) and \( c \), and the parameters \( p \) and \( q \) are calculated from eq (12) for \( b \) and \( c \), respectively. When \( n = 1 \), then \( m = 0 \), \( V = 0 \), and \( V' = 0 \), so that the energy is measured relative to the energy of the reactants. When \( n = 0 \), then \( m = 1 \), \( V = 0 \), and \( V = E_{\alpha b} - E_{\alpha c} \) which is the difference in the bond energies. BEBO assumes that the maximum value of \( V \) in the range \( 1 \geq n \geq 0 \) is the desired potential energy of the saddle point. This value \( V' \), is obtained by substituting into eq (14) that value of \( n \) which makes \( dV/dn = 0 \). In what follows, all quantities are considered to be evaluated at the saddle point.

Next, we must determine the stretching force constants in the \( \varphi \) and \( \sigma \) directions shown in figure 3. Equation (14) does not give the complete potential surface, but only that portion lying along the line of constant total bond order. BEBO assumes that at the saddle point, this path of constant bond order lies in the \( \varphi \) direction. This assumption will enable us to calculate the force constant \( F_{\varphi} = \frac{\partial^2 V}{\partial \varphi^2} \) from the second derivative of \( V \) with respect to \( n \), which we get by differentiating eq (14).

From eq (9), we can calculate the changes produced in \( R_b \) and \( R_c \) when \( n \) is changed. In vector notation these are

\[ \delta \mathbf{R} = \begin{bmatrix} \delta R_b \\ \delta R_c \end{bmatrix} = -\lambda \begin{bmatrix} 1/n \\ -1/m \end{bmatrix} \delta n; \ m = 1 - n. \]  
(15)

Because a change in \( n \) for constant total bond order is supposed to produce a move in the \( \varphi \) direction, the slope of a line in this direction can be gotten from eq (15). It is

\[ \frac{\delta R_b}{\delta R_c} = -\frac{n}{m} = \tan \alpha \]  
(16)

where \( \alpha \) is the angle which \( \varphi \) makes with the \( R_b \) axis as discussed earlier. From eq (15) we can show that

\[ \cos \alpha = m/\sqrt{(n^2 + m^2)}; \ \sin \alpha = -n/\sqrt{(n^2 + m^2)} \]  
(17)

The matrix \( \mathbf{U} \) defined in eq (6) can now be written in terms of \( n \) and \( m \).

\[ \mathbf{U} = \frac{1}{\sqrt{(n^2 + m^2)}} \begin{bmatrix} m & n \\ -n & m \end{bmatrix} \]  
(18)

By means of eq (6), \( \delta \mathbf{R} \) can be expressed in terms of \( \delta \mathbf{P} \); i.e., \( \delta \varphi \) and \( \delta \sigma \). Combining the differential form of eq (6) with eq (15), we get

\[ \mathbf{U} \delta \mathbf{P} = -\lambda \begin{bmatrix} 1/n \\ -1/m \end{bmatrix} \delta n \]  
(19)
Solving for $\delta P$ gives

$$
\delta P = \left[ \frac{\delta q}{\delta P} \right] = -\lambda U^{-1} \left[ \begin{array}{c} \frac{1}{n} \\ \frac{1}{m} \end{array} \right] \delta n = -\frac{\lambda}{\sqrt{(n^2 + m^2)}} \left[ \begin{array}{c} m - n \\ n - m \end{array} \right] \left[ \begin{array}{c} \frac{1}{n} \\ \frac{1}{m} \end{array} \right] \delta n
$$

As expected, $\sigma$ does not change when $n$ changes. From eq (20), we have for the derivative of $n$ with respect to $q$

$$
\frac{\delta n}{\delta q} = \frac{dn}{dQ} = -\frac{1}{\lambda} \frac{nm}{\sqrt{(n^2 + m^2)}}
$$

The second derivative of $V$ with respect to $q$ is obtained from the sequence

$$
\frac{dV}{dQ} = \frac{dn}{dn} \frac{dn}{dQ}
$$

$$
\frac{d^2V}{dQ^2} = \frac{d^2V}{dn^2} \left( \frac{dn}{dQ} \right)^2 + \frac{dV}{dn} \frac{d^2n}{dQ^2}
$$

Since $dV/dn = 0$ at the saddle point, we have

$$
F_s = \frac{d^2V}{dQ^2} = \frac{d^2V}{dn^2} \frac{n^2m^2}{\lambda^2(n^2 + m^2)}
$$

This gives one of the stretching force constants.

In the $\sigma$ direction, the stretching motion is assumed to be that of a normal molecule. Thus Badger's rule should be applicable. This says that the bond distance is proportional to the logarithm of the force constant, while eq (9) says that the bond distance is proportional to the logarithm of the bond order. Therefore, the force constant should be proportional to the bond order. We assume that

$$
F_s = F_{br}n, \quad F_c = F_{cm}
$$

where $F_{br}$ and $F_{cm}$ are single bond force constants. Consider the change in $V$ when $R_b$ and $R_c$ are changed by motion in the $\sigma$ direction. This is assumed to be given by

$$
2(\delta V)_\sigma = F_{br}n(\delta R_b)_\sigma^2 + F_{cm}(\delta R_c)_\sigma^2 + \frac{\partial^2V}{\partial R_c^2}(\delta R_c)_\sigma^2 = F_s(\delta \sigma)^2
$$

To evaluate $F_s$, we must express $(\delta R_b)_\sigma^2$, $(\delta R_c)_\sigma^2$, and $(\delta R_c)_\sigma^2$ in terms of $(\delta \sigma)^2$. From Eqs (6) and (18), we have

$$
\delta R = \left[ \begin{array}{c} \delta R_b \\ \delta R_c \end{array} \right] = \frac{1}{\sqrt{(n^2 + m^2)}} \left[ \begin{array}{c} m \\ n \\ -n \\ m \end{array} \right] \left[ \begin{array}{c} \delta_n \\ \delta_m \\ \delta_n \\ \delta_m \end{array} \right]
$$

For $\delta q = 0$,

$$
(\delta R_b)_\sigma = n\delta \sigma/\sqrt{(n^2 + m^2)}, \quad (\delta R_c)_\sigma = m\delta \sigma/\sqrt{(n^2 + m^2)}
$$

$$
(\delta R_c)_\sigma = (\delta R_b)_\sigma + (\delta R_c)_\sigma = (n + m)\delta \sigma/\sqrt{(n^2 + m^2)} = \delta \sigma/\sqrt{(n^2 + m^2)}.
$$

Therefore,

$$
2(\delta V)_\sigma = (F_{br}n^2 + F_{cm}n^2 + \frac{\partial^2V}{\partial R_c^2}) (\delta \sigma)^2(n^2 + m^2).
$$
Comparing this with eq (24) gives

$$F_a = \frac{F_b n^3 + F_c m^3 + \partial^2 V / \partial R_b^i}{n^2 + m^2}.$$  

(26)

The method assumes that if $V$ is expanded at the saddle point in terms of $\phi$ and $\sigma$ then there is no cross term; i.e., $\partial^2 V / \partial \phi \partial \sigma$ is assumed to be zero. Thus, we have

$$2 \delta V = (\delta \mathbf{P})^\dagger \begin{bmatrix} F_\phi & 0 \\ 0 & F_\sigma \end{bmatrix} (\delta \mathbf{P}).$$  

(27)

The use of eq (7) shows that

$$U^\dagger F_r U = \begin{bmatrix} F_\phi & 0 \\ 0 & F_\sigma \end{bmatrix}$$  

(28)

Inverting this equation gives

$$F_r = (U^\dagger)^{-1} \begin{bmatrix} F_\phi & 0 \\ 0 & F_\sigma \end{bmatrix} U^{-1} = U \begin{bmatrix} F_\phi & 0 \\ 0 & F_\sigma \end{bmatrix} U^\dagger$$  

(29)

where use has been made of the fact that $U^{-1} = U^\dagger$. Substituting eq (18) into (29) gives the desired stretching force constant matrix.

$$F_r = \frac{1}{n^2 + m^2} \begin{bmatrix} m & n \\ -n & m \end{bmatrix} \begin{bmatrix} F_\phi & 0 \\ 0 & F_\sigma \end{bmatrix} \begin{bmatrix} m - n \\ n - m \end{bmatrix}$$  

$$= \frac{1}{n^2 + m^2} \begin{bmatrix} F_\phi m^2 + F_\sigma n^2, -F_\phi mn + F_\sigma mn \\ -F_\phi mn + F_\sigma mn, F_\phi n^2 + F_\sigma m^2 \end{bmatrix} \begin{bmatrix} F_{11} & F_{12} \\ F_{21} & F_{22} \end{bmatrix}$$  

(30)

To complete the discussion of the BEBO method the bending force constants will now be evaluated. Consider first the one involving $M_3$ as the center mass. This will be $F_{s3}$ and appears in all of the transition states shown in figure 2. It is defined as the second partial derivative of $V$ with respect to the angle made by the bonds $b$ and $c$, with the bond lengths $R_b$ and $R_c$ held fixed. At equilibrium, this angle is $180^\circ$ for our transition state models. The geometry, when the angle is less than $180^\circ$ is shown in figure 6. To get $F_{s3}$, we differentiate $V$ twice,

$$\left( \frac{\partial V}{\partial \phi} \right)_{R_b, R_c} = \frac{\partial V}{\partial \phi} = \frac{\partial^2 V}{\partial R_b^i \partial \phi}$$  

$$\left( \frac{\partial^2 V}{\partial \phi^2} \right)_{R_b, R_c} = \frac{\partial^2 V}{\partial \phi^2} = \frac{\partial^2 V}{\partial R_b^i \partial \phi} \left( \frac{\partial R_b}{\partial \phi} \right)^2 + \frac{\partial V}{\partial R_c} \frac{\partial^2 R_c}{\partial \phi^2}$$  

(31)

![Figure 6. Definition of center bond angle.](image)

The derivatives of $V$, with respect to $R$, can be gotten from eq (13). The dependence of $R$ on $\phi$ can be determined from the following vector relationships,
\[ R_t = R_c - R_a \]
\[ R_t = R_c - R_b \]

\[ R_t^2 = R_t \cdot R_t = R_a^2 + R_b^2 - 2R_aR_b \cos \phi \]

\[ \frac{\partial R_t}{\partial \phi} = \frac{R_aR_b}{R_t} \sin \phi - 0 \quad \text{for } \phi = 180^\circ \]

\[ \frac{\partial^2 R_t}{\partial \phi^2} = \frac{R_aR_b \cos \phi}{R_t} - \frac{R_a^2 R_b^2}{R_t^2} \sin^2 \phi - \frac{R_aR_b}{R_t} \quad \text{for } \phi = 180^\circ \]

Thus,

\[ F_{\phi_3} = - \frac{\partial V_t}{\partial R_t} R_aR_b = - \frac{\partial V_t}{\partial R_t} \frac{R_aR_b}{R_a + R_b} \]

The other two bending force constants \( F_{\phi_2} \) and \( F_{\phi_4} \) are assumed to obey Badger's rule. We assume

\[ F_{\phi_2} = F_{\phi_2} \text{ and } F_{\phi_4} = F_{\phi_4} \text{ (34)} \]

This concludes the BEBO part of the calculation. It has provided us with the potential energy \( V^* \) of the saddle point, the stretching force constants \( F_{11}, F_{22}, \) and \( F_{12} \) and the bending force constants \( F_{\phi_2}, F_{\phi_3}, \) and \( F_{\phi_4}. \)

In the next section we shall use these force constants to carry out a frequency analysis for each of the transition state models shown in figure 2.

### 2.3. Vibrational Analysis

As we have seen in the force constant derivations, the potential energy \( V \) of the most general 5 mass point complex can be considered to depend on the variables \( R_a, R_b, R_c, R_d, \Psi_2, \Psi_3, \Psi_4, \Psi_5, \Psi_6, \) and \( \Psi_1. \) These are called the internal coordinates. Because our model is linear, \( V \) increases when any of the angles departs from 180°. Since we assume \( a \) and \( d \) to be rigid, \( R_a \) and \( R_d \) need not be included in the list of variables. For the time being, however, they will be included in the analysis. Let \( F \) be the complete force constant matrix for the complex. We have

\[ F = \begin{bmatrix}
F_{11} & F_{12} & \vdots \\
F_{21} & F_{22} & \vdots \\
\vdots & \vdots & \ddots \\
0 & \cdots & \cdots & 0
\end{bmatrix} \]

The two infinite force constants come from the use of rigid bonds for \( a \) and \( d. \) Let \( S \) be the (column) vector which denotes small changes in the saddle point values of the variables.

\[ S^\dagger = [\delta R_a, \delta R_b, \delta R_c, \delta R_d, \delta \Psi_2, \delta \Psi_3, \delta \Psi_4, \delta \Psi_5, \delta \Psi_6, \delta \Psi_1] \]

The potential energy is assumed to be given by

\[ V - V^* = \frac{1}{2} \dot{S}^\dagger F \dot{S} \]
Suppose there exists a matrix $G$, such that the kinetic energy in terms of the internal coordinates is

$$ T = \frac{1}{2} \dot{S}^T G^{-1} \dot{S} \quad (38) $$

Consider a new set of coordinates $Q$, the so-called normal coordinates, related to $S$ by the linear transformation

$$ S = LQ \quad (39) $$

such that

$$ V - V^* = \frac{1}{2} Q^T \Lambda Q = \Delta V \quad (40) $$

$$ T = \frac{1}{2} \dot{Q}^T \dot{E} \dot{Q} \quad (41) $$

where $\Lambda$ is a diagonal matrix having elements $\lambda_i$, and $E$ is the identity matrix. In this coordinate system there are no cross terms in $V$ and $T$.

Let $Q_i$ denote the $i$'th normal coordinate. The Lagrangian equations of motion for the system are

$$ \frac{d}{dt} \left( \frac{\partial L}{\partial \dot{Q}_i} \right) - \frac{\partial L}{\partial Q_i} = 0 \quad (42) $$

where $L = T - \Delta V = \frac{1}{2} \left[ \dot{Q}^T \dot{E} \dot{Q} - Q^T \Lambda Q \right] = \frac{1}{2} \left[ \sum \lambda_i Q_i^2 - \sum \lambda_i Q_i^2 \right]$,\n
$$ \frac{\partial L}{\partial \dot{Q}_i} = \dot{Q}_i \quad (43) $$

$$ \frac{\partial L}{\partial Q_i} = -\lambda_i Q_i \quad (44) $$

Therefore

$$ \ddot{Q}_i + \lambda_i Q_i = 0. \quad (46) $$

The solutions of this equation are

$$ Q_i = Q_i \cos(\lambda_i t + \epsilon_i). \quad (47) $$

Thus the $\lambda_i^2 = 2 \pi \nu_i$ are the frequencies of the vibrations of the $Q_i$ coordinates. These are called the normal mode vibrations.

Solving eq (39) for $Q$, and substituting into (40) and (41) yields

$$ V - V^* = \frac{1}{2} (L^{-1} S)^T \Lambda (L^{-1} S) = \frac{1}{2} S^T (L^{-1})^T \Lambda (L^{-1}) S \quad (48) $$

$$ T = \frac{1}{2} (L^{-1} \dot{S})^T \dot{E} (L^{-1} \dot{S}) = \frac{1}{2} \dot{S}^T (L^{-1})^T \dot{E} (L^{-1}) \dot{S} \quad (49) $$

Comparison with eqs (37) and (38) yields

$$ F = (L^{-1})^T \Lambda (L^{-1}) \quad (50) $$

$$ L^T FL = \Lambda \quad (51) $$

$$ G^{-1} = (L^{-1})^T \dot{E} (L^{-1}) \quad (52) $$

$$ L^T G^{-1} L = E \quad (53) $$
Next, solve eq (53) for \( L^\dagger = L^{-1}G \), substitute this into eq (51) and multiply by \( L \) on the left. This gives

\[
GFL = HL = LA
\]

as the set of equations which determine the transformation \( L \). Written out, eq (54) is

\[
\sum \delta_{ij} \lambda_L L_{jk} = 0
\]

This equation has solutions if the determinant

\[
|H - \lambda L| = 0
\]

This is the so-called secular equation which must be solved to get the \( \lambda_L \), the eigenvalues of \( H \) and thus the normal frequency values. Before doing this, it is first necessary to evaluate the matrix \( G \).

Equation (38) gives the kinetic energy in terms of the internal coordinates. As such, it does not include the kinetic energy of the center of mass or the rotational energy. We need to express the kinetic energy in terms of cartesian coordinates, transform the result to internal coordinates, and subtract out the center of mass and rotational energy. This will yield \( G^{-1} \). Let us begin by expressing the internal coordinates in terms of cartesian coordinates. Assume that the molecule lies along the x axis. A particular mass point \( M_i \) will have coordinates \((x_i, y_i, z_i)\) where \( y_i \) and \( z_i \) are small and describe the departures of the molecule from linearity during bending vibrations. Because \( y_i \) and \( z_i \) are small, the bond distances can be expressed as functions of the \( x_i \) only. Thus,

\[
\begin{align*}
R_a &= x_2 - x_1 \\
R_b &= x_3 - x_2 \\
R_c &= x_4 - x_3 \\
R_d &= x_5 - x_4 \\
R_e &= x_6 - x_5
\end{align*}
\]

Since there are 5 cartesian \( x \) coordinates we need one more coordinate for the internal system. This is taken to be the \( x \)-component of the center of mass of the molecule multiplied by the total mass, and is defined by the equation,

\[
Mx = \sum_{i=1}^5 M_i x_i
\]

where \( M = \sum_{i=1}^5 M_i \)

In matrix form these equations are

\[
\begin{bmatrix}
R_a \\
R_b \\
R_c \\
R_d \\
Mx
\end{bmatrix} =
\begin{bmatrix}
-1 & 1 & 0 & 0 & 0 \\
0 & -1 & 1 & 0 & 0 \\
0 & 0 & -1 & 1 & 0 \\
0 & 0 & 0 & -1 & 1 \\
M_1 & M_2 & M_3 & M_4 & M_5
\end{bmatrix}
\begin{bmatrix}
x_1 \\
x_2 \\
x_3 \\
x_4 \\
x_5
\end{bmatrix} = Mx
\]

Note that the vector \( R \) is basically that defined by eq (15). Here we have included \( R_a \) and \( R_d \).

We must next express the bond angles in terms of the cartesian coordinates. Consider \( \Psi_2 \), the angle formed by bonds \( a \) and \( b \). The geometry and notation for this angle are shown in figure 7. The two vectors along the bonds \( a \) and \( b \) are given by
The geometry of the bond angle $\Psi_2$ is related to the equation by

$$r_{21} + r_{23} = r_{21}r_{23}\cos\Psi_2.$$  

(71)

Substituting eq (70) into (71) gives

$$-R_aR_b + (y_1 - y_2)(y_3 - y_2) = \{[(R_a^2 + (y_1 - y_2)^2)(R_b^2 + (y_3 - y_2)^2)]^{1/2}\cos\Psi_2$$

(72)

Because the $y_i$ are small compared to $R_a$ and $R_b$, the radical can be expanded to give

$$-R_aR_b + (y_1 - y_2)(y_3 - y_2) = [R_aR_b + \frac{1}{2}(R_a/R_b)(y_1 - y_2)^2 + \frac{1}{2}(R_b/R_a)(y_3 - y_2)^2]$$

(73)

Let $\Psi_2 = 180^\circ + \delta\Psi_2$ where $\delta\Psi_2$ is small. Then

$$\cos\Psi_2 = -\cos(\delta\Psi_2) \approx -1 + \frac{1}{2}(\delta\Psi_2)^2$$

Substituting this into eq (73) and keeping terms through second order gives

$$(\delta\Psi_2)^2 = (1/R_a)(y_1 - y_2)^2 + (2/R_aR_b)(y_1 - y_2)(y_3 - y_2) + (1/R_b)(y_3 - y_2)^2$$

$$\delta\Psi_2 = -[(y_1 - y_2)/R_a + (y_3 - y_2)/R_b] = -y_1/R_a + (1/R_a + 1/R_b)y_2 - y_3/R_b$$

(74)

To see why the minus sign is needed, let $y_1 = y_3 = 0$; then for $y_2 > 0$, $\Psi_2 < 180^\circ$, so that $\delta\Psi_2$ must be $< 0$. There are analogous equations for the angles $\Psi_3$ and $\Psi_4$; there is also a set, identical in form, for the angles $\Psi'_i$ in the $x$-$z$ plane. These contain the $z_i$ rather than the $y_i$ coordinates. In these equations, the equilibrium values of $R_{a,\ldots,R_b}$ will be used.

The set of equations typified by eq (74) gives 3 equations in terms of the 5 $y_i$ coordinates; two more are needed. We have one defining the $y$ coordinate of the center of mass, like eq (58), and another defining a quantity $\eta_a$, which is given by the equation

$$\eta_a = \sum_{i=1}^{5} M_i x_i y_i$$

(75)

$\eta_a$ is related to the $z$ component of the angular momentum $m_z$ by the relation

$$m_z = \eta_a$$

(76)
The $x_i$ are the equilibrium $x$ values; these can be gotten relative to the center of mass component $x$, by inverting eq (60) and inserting equilibrium values for $R_a, \ldots, R_d$. In matrix form, these equations relating $y$, to the bond angles in the $x-y$ plane are,

$$\bar{\Psi} = \begin{bmatrix} \delta \psi_2 \\ \delta \psi_3 \\ \delta \psi_4 \\ M_y \end{bmatrix} = \begin{bmatrix} -q_a & q_a + q_b & -q_b & 0 & 0 \\ 0 & -q_b & q_b + q_c & -q_c & 0 \\ 0 & 0 & -q_c & q_c + q_d & -q_d \\ M_1x_1^2 & M_2x_2^2 & M_3x_3^2 & M_4x_4^2 & M_5x_5^2 \end{bmatrix} \begin{bmatrix} y_1 \\ y_2 \\ y_3 \\ y_4 \\ y_5 \end{bmatrix} = AY \quad (77)$$

where $q_a, \ldots, q_d$ are the reciprocals of the equilibrium values of $R_a, \ldots, R_d$. There is an analogous equation involving the $z$ coordinates.

Having obtained expressions (60) and (77) for the internal coordinates in terms of the cartesian coordinates, we can now invert these equations and insert them into the expression for the total kinetic energy which we shall call $T$. Therefore

$$\bar{T} = \frac{1}{2} \bar{X}^T \bar{D}_m \bar{X} + \frac{1}{2} \bar{Y}^T \bar{D}_m \bar{Y} + \frac{1}{2} \bar{Z}^T \bar{D}_m \bar{Z}$$

$$= \frac{1}{2} \hat{R}^T (M^{-1})^T \bar{D}_m (M^{-1}) \hat{R} + \frac{1}{2} \hat{\Psi}^T (A^{-1})^T \bar{D}_m (A^{-1}) \hat{\Psi} + z\text{-term}$$

$$= \frac{1}{2} \hat{R}^T \bar{G}_r^{-1} \hat{R} + \frac{1}{2} \hat{\Psi}^T \bar{G}_\varphi^{-1} \hat{\Psi} + z\text{-term}$$

$$= \frac{1}{2} T + \frac{1}{2} M(\dot{x}^2 + \dot{y}^2 + \dot{z}^2) + \frac{1}{2}(m_1^2 + m_2^2)/I$$

$$= \frac{1}{2} \hat{R}^T \bar{G}_r^{-1} \hat{R} + \frac{1}{2} \hat{\Psi}^T \bar{G}_\varphi^{-1} \hat{\Psi} + z\text{-term} + \frac{1}{2} M(\dot{x}^2 + \dot{y}^2 + \dot{z}^2) + \frac{1}{2}(m_1^2 + m_2^2)/I \quad (78)$$

where $I = \sum_{i=1}^{3} M_i x_i^2$ is the moment of inertia, and

$$D_m = \begin{bmatrix} M_1 & M_2 & 0 \\ 0 & M_3 & M_4 \\ 0 & 0 & M_5 \end{bmatrix} \quad (79)$$

We can satisfy eq (78) by writing $\bar{G}_r^{-1}$ and $\bar{G}_\varphi^{-1}$ in the partitioned forms

$$\bar{G}_r^{-1} = \begin{bmatrix} G_r^{-1} & 0 \\ 0 & M^{-1} \end{bmatrix} \quad (80)$$

$$\bar{G}_\varphi^{-1} = \begin{bmatrix} G_\varphi^{-1} & 0 \\ 0 & L^{-1} \end{bmatrix} = \bar{G}_\varphi^{-1} \quad (80)$$

We can get $G_r$ and $G_\varphi$ simply by inverting $\bar{G}_r^{-1}$ and $\bar{G}_\varphi^{-1}$.

This gives

$$\bar{G}_r = \begin{bmatrix} G_r & 0 \\ 0 & L \end{bmatrix} = MD_m^{-1}M^+ \quad (81)$$

$$\bar{G}_\varphi = \begin{bmatrix} G_\varphi & 0 \\ 0 & L \end{bmatrix} = AD_m^{-1}A^+$$
Since \( D_m \) is diagonal its inverse is easily evaluated and we therefore require only matrix multiplications to get \( G_r \) and \( G_{\phi} \).

The complete \( G \) matrix for the internal coordinates in partitioned form is

\[
G = \begin{bmatrix}
G_r & 0 & 0 \\
0 & G_{\phi} & 0 \\
0 & 0 & G_{\phi}^\prime
\end{bmatrix}
\]  

(82)

In partitioned form, the complete force constant matrix, eq (35), is

\[
F = \begin{bmatrix}
F_r & 0 & 0 \\
0 & F_{\phi} & 0 \\
0 & 0 & F_{\phi}^\prime
\end{bmatrix}
\]  

(83)

Note that \( F_r \) here is like eq (30), but contains the two infinite force constants corresponding to the rigid \( a \) and \( d \) bonds. The matrix \( H \) in partitioned form is

\[
H = \begin{bmatrix}
G_r F_r & 0 & 0 \\
0 & G_{\phi} F_{\phi} & 0 \\
0 & 0 & G_{\phi} F_{\phi}^\prime
\end{bmatrix}
= \begin{bmatrix}
H_r & 0 & 0 \\
0 & H_{\phi} & 0 \\
0 & 0 & H_{\phi}^\prime
\end{bmatrix}
\]  

(84)

Because \( H \) factors in this way, we can set up separate secular equations for the stretching and bending modes. Note that \( H \) is normally unsymmetric.

Before solving the secular equations, let us write down explicit expressions for \( G_r \) and \( G_{\phi} \). The direct evaluation of \( G_r \) from eq (81) yields

\[
\tilde{G}_r = \begin{bmatrix}
\mu_1 + \mu_2 & -\mu_2 & 0 & 0 & 0 \\
-\mu_2 & \mu_2 + \mu_3 & -\mu_3 & 0 & 0 \\
0 & -\mu_3 & \mu_3 + \mu_4 & -\mu_4 & 0 \\
0 & 0 & -\mu_4 & \mu_4 + \mu_5 & 0 \\
0 & 0 & 0 & 0 & M
\end{bmatrix}
\]  

(85)

where the \( \mu_i \) are the reciprocals of the masses \( M_i \). Comparison of this equation with eq (81) yields \( G_r \).

Because we are treating the \( a \) and \( d \) bonds as rigid, the stretching part of the problems is equivalent to a 3 mass point system where the first mass is \( M_1 + M_2 \) and the third is \( M_3 + M_4 \). The resulting \( 2 \times 2 \) matrix is the one actually used in the calculation. It is

\[
G_r(\text{rigid end bonds}) = \begin{bmatrix}
\frac{\mu_1 + \mu_2}{\mu_1} & \frac{\mu_2 + \mu_3}{\mu_2} & -\mu_3 \\
-\mu_3 & \frac{\mu_3 + \mu_4}{\mu_3} & \frac{\mu_4 + \mu_5}{\mu_4}
\end{bmatrix}
\]  

(86)

The stretching force constant matrix to be used with eq (86) is that \( F_r \) as given by eq (30).

The \( G_{\phi} \) matrix elements for this 5 point case are

\[
\begin{align*}
(G_{\phi})_{11} &= \varepsilon^2 \mu_1 + \varepsilon^2 \mu_3 + (\varepsilon_a + \varepsilon_b) \mu_2 \\
(G_{\phi})_{22} &= \varepsilon^2 \mu_2 + \varepsilon^2 \mu_4 + (\varepsilon_b + \varepsilon_c) \mu_3 \\
(G_{\phi})_{33} &= \varepsilon^2 \mu_3 + \varepsilon^2 \mu_5 + (\varepsilon_c + \varepsilon_d) \mu_4 \\
(G_{\phi})_{12} &= -\varepsilon_\phi [(\varepsilon_a + \varepsilon_b) \mu_2 + (\varepsilon_b + \varepsilon_c) \mu_3] \\
(G_{\phi})_{23} &= -\varepsilon_\phi [(\varepsilon_b + \varepsilon_c) \mu_3 + (\varepsilon_c + \varepsilon_d) \mu_4] \\
(G_{\phi})_{33} &= \varepsilon_\phi \varepsilon_\phi \mu_3 
\end{align*}
\]  

(87)
There are the expressions used in the calculation. Actually, they were not derived from eq (81) but were obtained from Wilson et. al. [12]. However, eq (81) was used for a numerical check of eq (87). To get the matrix elements for the two 4 mass point cases, simply delete from eq (87) those elements which contain either a missing $q$ or a missing $\mu$ or both. Do the same for the 3 point case, but delete also $(G_\phi)_{12}$; (there is only one element, $(G_\phi)_{22}$, in this case).

We are now ready to consider the secular equation. For the rate constant calculation only the $\lambda_k$ are required, so that a solution of eq (54) for the transformation matrix $L$ is not necessary. Nevertheless, $L$ is easily obtained and is convenient to have for the purpose of illustrating the actual vibrational motions of the complex. Thus we shall solve eq (54) as well as eq (56). According to eq (84), there are two secular equations to be solved ($H_\lambda$ and $H_\nu$, are equal). Because we are using rigid $a$ and $d$ bonds, the dimension of $H_\lambda$ is $2 \times 2$. The maximum dimension of $H_\nu$ is $3 \times 3$ and occurs for the 5 point model. Thus a solution of a $3 \times 3$ problem will suffice for our purpose and will also illustrate how an $n \times n$ problem is to be solved.

We begin by assuming that eq (56) has been solved. In the present work this was accomplished by expanding (56) and solving the resulting polynomial in $\nu$. In our case, the maximum degree was cubic, so that this part of the calculation was easily performed. As eq (47) shows, the desired frequencies are $\nu_k = \lambda_k^{-1/2} / 2\pi$. For the stretching modes of the complex one of the two frequencies will be imaginary because its $\lambda_k$ value will be negative. As mentioned earlier, this corresponds to the asymmetric stretch.

Consider now eq (55) for a general $3 \times 3 H$ matrix. Written out in full, it is

\[
(H_{11} - \lambda_k) L_{1k} + H_{12} L_{2k} + H_{13} L_{3k} = 0
\]

\[
H_{21} L_{1k} + (H_{22} - \lambda_k) L_{2k} + H_{23} L_{3k} = 0
\]

\[
H_{31} L_{1k} + H_{32} L_{2k} + (H_{33} - \lambda_k) L_{3k} = 0
\]

where $\lambda_k$ is one of the three values of $\lambda$ determined from the solution of the cubic (in this case) eq (56). Divide the first two of these equations by $L_{3k}$, and define the ratios $g_{1k} = L_{1k} / L_{3k}$. This yields two equations to be solved for the two unknowns $g_{1k}$ and $g_{2k}$.

\[
(H_{11} - \lambda_k) g_{1k} + H_{12} g_{2k} = -H_{13}
\]

\[
H_{21} g_{1k} + (H_{22} - \lambda_k) g_{2k} = -H_{23}
\]

We get two $g_{ik}$ values for each value of $\lambda_k$ substituted into eq (89), or six $g_{ik}$ values in all. Using these values, we can express $L$ in terms of the product of two matrices defined by

\[
L = \begin{bmatrix} g_{11} & g_{12} & g_{13} \\ g_{21} & g_{22} & g_{23} \\ 1 & 1 & 1 \end{bmatrix} \begin{bmatrix} L_{31} & 0 & 0 \\ 0 & L_{32} & 0 \\ 0 & 0 & L_{33} \end{bmatrix} = \Gamma \ell
\]

To determine the components of $\ell$, insert eq (90) into eq (51). We get

\[
\ell \Gamma \Gamma^\dagger \Gamma = \Lambda = \ell \ell^\dagger \Gamma \Gamma^\dagger \Gamma
\]

The final reordering is possible because $\ell$ and $\Lambda$ are diagonal and therefore $\Gamma \Gamma^\dagger \Gamma$ is diagonal. This equation is easily solved for the elements $\ell \ell^\dagger \ell$ to give

\[
(\ell \ell^\dagger)_{kk} = L_{3k} = \lambda_k (\Gamma \Gamma^\dagger \Gamma)_{kk}
\]

The other elements of $L$ are gotten from these values and the ratios $g_{ik}$ already determined.

The actual motions in the cartesian system can now be obtained by combining eq (39) with the inverse of eq (60) or eq (77). For the stretching motions we have
\[ X = M^{-1} \tilde{R} = M^{-1} \begin{bmatrix} L_i Q_i \\ 0 \end{bmatrix} \]  

where \( L_i \) arises from the secular equation containing \( H_i \). \( Q_i \) is the normal coordinate vector and the \( x \)-component of the center of mass has been set to zero. A similar equation results for the bending modes. This is

\[ Y = A^{-1} \tilde{\Psi} = A^{-1} \begin{bmatrix} L_i Q_i \\ 0 \\ 0 \end{bmatrix} \]  

where the \( z \)-component of the angular momentum and the \( y \)-component of the center of mass have been set to zero.

This completes the frequency analysis. In the next section we will consider the partition functions.

### 2.4. Partition Functions.

Herschbach et. al. [13] have shown how to express the classical partition function for polyatomic molecules in terms of local properties. We shall use their method because it allows for cancellations of considerable portions of the partition functions of the complex and reactants when their ratios are evaluated in the rate constant expression, eq (3). We begin the discussion with the classical partition function for a linear polyatomic molecule. This is (see Herzberg [14], pp. 502-509),

\[ q_{cl} = q_V (2\pi M k T / \hbar^2)^{3/2} \left( k T / (\hbar c B) \right)^{3N-5} \Pi_i \left( k T / (\hbar \omega_i B) \right) \]

where \( q_e \) is the electronic partition function, \( V \) is the volume, \( M \) is the total mass of the molecule, \( c \) is the velocity of light, \( \omega_i \) is the frequency of the \( i \)-th vibrational mode in cm\(^{-1} \) (\( \omega_i = \nu_i / c \)), \( N \) is the number of atoms in the molecule, \( B \) is the rotational constant; \( B = h/(8\pi^2 eI) \), where \( I \) is the moment of inertia of the molecule; \( \sigma \) is the symmetry number which is the number of indistinguishable positions into which the molecule can be turned by simple rigid rotations. For linear molecules \( \sigma = 1 \) or 2. Equation (95) neglects nuclear spins, anharmonicity, and non-rigidity of the molecule. Let us rewrite eq (95) in terms of \( I \) and \( u_i = h\nu_i/kT \). It becomes

\[ q_{cl} = q_e V 4\pi \sigma^{-1} (2\pi k T h^{-2})^{3/2} M^{3/2} I^{3N-5} \Pi_i u_i^{-1} \]

\( q_{cl} \) can also be written in the form

\[ q_{cl} = q_e \sigma^{-1} Z \prod_{a} \Lambda_a^3 \]

where

\[ Z = \int \cdots \int e^{-V/4\sigma^2} dx_1 \cdots dx_N 
\]

\[ \Lambda_a = h(2\pi M_a k T)^{-1/4} \]

\( Z \) is the so-called configuration integral, \( V \) is the potential energy, and \( x_1, y_1, z_1, \ldots, x_N, y_N, z_N \) are the cartesian coordinates of each of the \( N \) atoms. Eliminating \( q_{cl} \) between eqs (96) and (97) gives

\[ Z = V 4\pi (2\pi k T h^{-2})^{-1/4} M^{3N-5} I^{3N-5} \Pi_i M_a^{3/2} I u_i^{-1} \]

Consider now the matrix \( \mathbf{H} = GF \) defined by eq (84). A theorem of matrix algebra states that the determinant of \( \mathbf{H} \) equals the product of its eigenvalues (see Hohn [15], p. 283). There is also a theorem (Hohn, p. 65),
stating that the determinant of product of two matrices equals the products of the determinants of the matrices in the product. Consequently,

\[
|H|^{\frac{1}{N}} = |G|^{\frac{1}{N}} |F|^{\frac{1}{N}} = \prod_{i}^{3N-5} \lambda_{i}^{\frac{1}{N}} = (2\pi kT/h)^{3N-5} \prod_{i}^{3N-5} u_{i} \tag{100}
\]

Solving for the product over \( u_{i} \) gives

\[
\prod_{i}^{3N-5} u_{i}^{-1} = (2\pi kT/h)^{3N-5} |G|^{-\frac{1}{N}} |F|^{-\frac{1}{N}} \tag{101}
\]

Inserting eq (101) into (99) yields

\[
Z = V4\pi(2\pi kT)^{\frac{1}{2}(3N-5)} M^{3/2} I \prod_{i}^{N} M_{i}^{-3/2} |G|^{-\frac{1}{N}} |F|^{-\frac{1}{N}} \tag{102}
\]

This can be rearranged to give

\[
|F|^{\frac{1}{N}} (2\pi kT)^{-\frac{1}{2}(3N-5)} Z = V4\pi M^{3/2} I \prod_{i}^{N} M_{i}^{-3/2} |G|^{-\frac{1}{N}} = J_{N} \tag{103}
\]

The left side of eq (103) does not involve the masses, while the right side does not contain force constants. Therefore, the quantity denoted by \( J_{N} \) does not depend on either the force constants or the masses, but must depend only on geometrical parameters. Herschbach et. al. [13] have shown that for linear molecules

\[
J_{N} = V4\pi \prod_{i=1}^{N-1} R_{i+1,i} \tag{104}
\]

where \( R_{i+1,i} \) is the equilibrium distance between mass \( M_{i} \) and \( M_{i+1} \). For a general linear molecule, the classical partition function per unit volume can now be written

\[
q_{cl} = q_{cl}/V = V^{-1} q_{l}^{\sigma_{l}} J_{N}(2\pi kT)^{\frac{1}{2}(3N-5)} |F|^{-\frac{1}{N}} \prod_{i}^{N} \Lambda_{i}^{-3} \tag{105}
\]

This form of the partition function is suitable for the reactant molecules.

Let us now consider the partition function for the complex. Using eq (96), we have

\[
kT h^{-1} q^{*} = kT h^{-1} q_{l}^{\sigma_{l}} V4\pi(2\pi kT)^{-3N-5} M^{3/2} I \prod_{i}^{3N-6} u_{i}^{-1} \tag{106}
\]

Note that the product is over \( 3N-6 \); i.e., one less vibration than in a stable linear molecule. Consider next the quantity

\[
\left( \prod_{i}^{3N-6} u_{i}^{-1} \right) kT h^{-1} = (kT/h)^{3N-5} \prod_{i}^{3N-6} u_{i}^{-1} = (kT/h)^{3N-5} \prod_{i}^{3N-6} 2\pi \lambda_{i}^{\frac{1}{2}}
\]

\[
= (kT/h)^{3N-5} (2\pi)^{3N-6} \lambda^{\frac{1}{2}} \prod_{i}^{3N-6} \lambda_{i}^{\frac{1}{2}} \lambda_{i}^{-\frac{1}{2}}
\]

\[
= (2\pi kT/h)^{3N-5} \nu^{*} |F|^{-\frac{1}{N}} |G|^{-\frac{1}{N}} \tag{107}
\]

where \( \lambda^{*} \) is the negative eigenvalue and \( \nu^{*} \) is the associated imaginary frequency; eq (100) has been used. Using eq (107) in (106) gives

\[
kT h^{-1} Q^{*} = kT h^{-1} q^{*} V^{-1} = V^{-1} q_{l}^{\sigma_{l}} \left[ V4\pi M^{3/2} I \prod_{i}^{N} M_{i}^{-3/2} |G|^{-\frac{1}{N}} \nu^{*} |F|^{-\frac{1}{N}} (2\pi kT)^{3N-5/2} \right]
\]

\[
\cdot h^{-3N} \prod_{i}^{N} M_{i}^{-3/2}
\]

\[
= V^{-1} q_{l}^{\sigma_{l}} J_{N} \nu^{*} |F|^{-\frac{1}{N}} (2\pi kT)^{\frac{1}{2}(3N-5)} \prod_{i}^{N} \Lambda_{i}^{-3} \tag{108}
\]

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This equation is very similar to eq (105), the partition function of a stable molecule. Note that $|F|^\frac{1}{4}$ will be imaginary for the complex.

We can now write down the specific partition functions per unit volume for the four reaction cases shown in figure 1b.

Case III.

Species $A = M_2 - M_3$
Species $B = M_4$
Species $C = M_2 \ldots M_3 \ldots M_4$

$$Q_A = q_{c\alpha c\beta} 4\pi R_{ab}(2\pi kT)^{\frac{1}{2}} F_{a\beta}^{-1} (\Lambda_2 \Lambda_3)^{-3}$$
$$Q_B = q_{a\beta a\beta}$$
$$Q_C kT/h = q_{c\alpha c\alpha} 4\pi R_{ab} R_{bc}^{*} |F_r|^{-\frac{1}{2}} F_{a\beta}^{-1} (2\pi kT)^{2} (\Lambda_2 \Lambda_3 \Lambda_4)^{-3}$$

The matrix $F_r$ is the $2 \times 2$ one given by eq. (5b), and not the $4 \times 4$ used in eq (83).

Case IVa.

Species $A = M_2 - M_3$
Species $B = M_3$
Species $C = M_1 - M_2 \ldots M_3 \ldots M_4$

$$Q_A = q_{c\alpha c\beta} 4\pi R_{ab} R_{bc} (2\pi kT)^{\frac{1}{2}} F_{a\beta}^{-1} (F_{w} F_{w})^{-1} F_{w}^{-1} (\Lambda_1 \Lambda_2 \Lambda_3)^{-3}$$
$$Q_B = q_{a\beta a\beta}$$
$$Q_C kT/h = q_{c\alpha c\alpha} 4\pi R_{ab} R_{bc}^{*} |F_r|^{-\frac{1}{2}} F_{a\beta}^{-1} F_{a\beta} (2\pi kT)^{2} (\Lambda_1 \Lambda_2 \Lambda_3 \Lambda_4)^{-3}$$

Note that I have included $C_{w}$ in $Q_A$ and $Q_C$ even though it is supposed to be infinite; it will cancel out when the ratio $Q_C/Q_A$ is taken. Also note that the bending force constants appear with twice the power of the stretching force constants. This is because of the degeneracy.

Case IVb.

Species $A = M_3 - M_3$
Species $B = M_3$
Species $C = M_1 - M_2 \ldots M_3 \ldots M_4$

$$Q_A = q_{c\alpha c\beta} 4\pi R_{ab} R_{bc} (2\pi kT)^{\frac{1}{2}} F_{a\beta}^{-1} (F_{w} F_{w})^{-1} F_{w}^{-1} (\Lambda_1 \Lambda_2 \Lambda_3)^{-3}$$
$$Q_B = q_{a\beta a\beta}$$
$$Q_C kT/h = q_{c\alpha c\alpha} 4\pi R_{ab} R_{bc}^{*} |F_r|^{-\frac{1}{2}} F_{a\beta}^{-1} F_{a\beta} (2\pi kT)^{2} (\Lambda_1 \Lambda_2 \Lambda_3 \Lambda_4)^{-3}$$

Case V.

Species $A = M_3 - M_3$
Species $B = M_3$
Species $C = M_1 - M_2 \ldots M_3 \ldots M_4$

$$Q_A = q_{c\alpha c\beta} 4\pi R_{ab} R_{bc} (2\pi kT)^{\frac{1}{2}} F_{a\beta}^{-1} (F_{w} F_{w})^{-1} F_{w}^{-1} (\Lambda_1 \Lambda_2 \Lambda_3)^{-3}$$
$$Q_B = q_{a\beta a\beta}$$
$$Q_C kT/h = q_{c\alpha c\alpha} 4\pi R_{ab} R_{bc}^{*} |F_r|^{-\frac{1}{2}} F_{a\beta}^{-1} F_{a\beta} (2\pi kT)^{2} (\Lambda_1 \Lambda_2 \Lambda_3 \Lambda_4)^{-3}$$

We now have everything for eq (3) except the tunneling correction. This will be taken up in the next section.

### 2.5. Tunneling Correction

The one-dimensional Eckart potential function was used to approximate the barrier to quantum mechanical tunneling from reactants to products. Three parameters are required for its definition; these are shown in figure 8. Its functional form is
\[ V(x) = \frac{Ay}{1-y} - \frac{By}{(1-y)^2} \]  

where

\[ \gamma = -e^{2\pi \xi L} \]
\[ A = V_1 - V_2 \]
\[ B = (V_1^2 + V_2^2)^{1/2} \]
\[ L = 2\pi (-2/F)^{1/2} (V_1^{14} + V_2^{14})^{-1}, \text{ and} \]
\[ F = \frac{\partial^2 V}{\partial x^2} \]

evaluated at the maximum in the curve. \( F \) is a force constant. Using this potential function, Eckart [16] solved the wave equation and obtained the transmission coefficient for a particle with mass \( m \) approaching the barrier from the left with an energy \( E \). His result is

\[ K(E,V_1,V_2,F) = 1 - \frac{\cosh[2\pi(\alpha_1 - \alpha_2)] + \Delta}{\cosh[2\pi(\alpha_1 + \alpha_2)] + \Delta} \]  

where \( \Delta = \cosh[2\pi\delta] \) if \( \delta \) is real, and \( \Delta = \cos[2\pi|\delta|] \) if \( \delta \) is imaginary. The relationships of \( \alpha_1, \alpha_2, \) and \( \delta \) to the parameters of figure 8, are

\[ \alpha_1 = \frac{1}{2}[(E/C)^4 - 1] \]
\[ \alpha_2 = \frac{1}{2}[(E-A)/C]^4 \]
\[ \delta = \frac{1}{2}[(B-C)/C]^4 \]
\[ C = \frac{\hbar^2}{8mL^2} \]  

\[ \text{Figure 8. Eckart potential function.} \]

Given the transmission coefficient, Johnston [2], pp. 42 and 43, has derived the correction factor \( \Gamma^* \) which is the ratio of the quantum barrier crossing rate to the classical crossing rate. His result is

\[ \Gamma^* = e^{\gamma_1 \alpha T} \int_{E_o}^\infty K(E)e^{-E/kT} dE/kT \]  

where \( E_o = 0 \) when \( V_1 \leq V_2 \) and \( E_o = V_1 - V_2 \) when \( V_1 > V_2 \)
Let us rewrite this in a more symmetrical form. We define a new variable \( \epsilon = (E - V_i)/kT \), \( E = kTe + V_i \). Equation (112) becomes

\[
\Gamma^* = \int_{\epsilon_o}^{\infty} K(\epsilon) e^{-\epsilon} d\epsilon
\]  

(113)

where \( \epsilon_o = -V_1/kT \) when \( V_1 \leq V_2 \) and \( \epsilon_o = -V_2/kT \) when \( V_1 > V_2 \).

With this substitution, the parameters \( \alpha_1 \) and \( \alpha_2 \) become

\[
\alpha_i = \frac{1}{2}(kTe/C + V_i/C)^{1/2}, \quad i = 1 \& 2
\]  

(114)

From eq (110) we have

\[
K = K(\alpha_1, \alpha_2, \delta) = K(kTe/C, V_1/C, V_2/C, B/C)
\]

But

\[
B/C = V_1/C + 2[(V_1/C)(V_2/C)]^{1/2} + V_2/C
\]  

(115)

is a function of \( V_1/C \) and \( V_2/C \). Therefore

\[
K = K(\epsilon, p_1, p_2)
\]  

(116)

where

\[
p = kT/C, \quad p_1 = V_1/C \text{ and } p_2 = V_2/C.
\]

\( \Gamma^* \) thus depends on three parameters. Furthermore, it is invariant when \( p_1 \) and \( p_2 \) are interchanged; i.e., \( \Gamma^*(p_1, p_2, p) = \Gamma^*(p_2, p_1, p) \). To see this let \( p_1' = p_2 \) and \( p_2' = p_1 \). From eq (115) we see that

\[
(B/C)' = p_1' + 2(p_1'p_2')^{1/2} + p_2' = p_2 + 2(p_2p_1)^{1/2} + p_1 = B/C
\]

Thus, \( \delta' = \delta \). From eq (114) we have

\[
\alpha_1' = \frac{1}{2}(p\epsilon + p')^{1/2} = \frac{1}{2}(p\epsilon + p_1')^{1/2} = \alpha_1
\]

\[
\alpha_2' = \alpha_2
\]

Using these results in eq (110), we get

\[
K(\epsilon, p_1, p_2) = K(\epsilon, p_1, p_2) = K(\epsilon, p_1, p_2)
\]  

(117)

Suppose that \( p_1' > p_2' \); i.e., \( V_1' > V_2' \). Using eq (117), eq (113) becomes

\[
\Gamma^*(p_1, p_2) = \int_{\epsilon_o}^{\infty} K(\epsilon, p_1, p_2) e^{-\epsilon} d\epsilon = \int_{\epsilon_o}^{\infty} K(\epsilon, p_1, p_2) e^{-\epsilon} d\epsilon = \Gamma^*(p_1, p_2)
\]

The way Eq (113) was integrated to get \( \Gamma^* \) will be considered later when the computer program is discussed.

In applying this correction, it is assumed that the \( x \) coordinate of Eckart's potential lies in the \( q \) direction discussed earlier. This is that direction at the saddle point in which the potential energy decreases most rapidly. It is also the direction of the path of constant total bond order. We therefore use the force constant \( F_\phi \) given by eq (22) for the second derivative of the Eckart potential at its maximum. The effective mass for tunneling, \( M^* \), is the proportionality factor between the kinetic energy and \( \frac{1}{2}q^2 \). We can calculate \( M^* \) in the following way: As far as tunneling is concerned, in the 4 and 5 mass point cases there are effectively 3
masses, since the end bonds are supposed to be rigid. Thus, there are only the two variables, \( R_b \) and \( R_e \), involved. (Bending modes are not considered.) The kinetic energy \( T_r \) for changes in these two bonds is given by

\[ T_r = \frac{1}{2} \dot{R} \cdot \mathbf{G} \cdot \dot{R} \]  

(118)

where \( \mathbf{R} \) is the 2-dimensional vector defined by eq (6), and \( \mathbf{G} \) is the \( 2 \times 2 \) matrix given by eq (86). The inverse of this matrix is easily calculated and found to be

\[ \mathbf{G}^{-1} = \begin{bmatrix} M_2(M_2 + M_4) & M_2M_4' \\ M_2M_4' & (M_2 + M_4)M_4' \end{bmatrix} M^{-1} \]  

(119)

where \( M_2' = M_1 + M_2 \) and \( M_4' = M_4 + M_5 \) in the 5 point case. The transformation between \( R_b, R_e \) and \( \varphi, \sigma \) is given by the matrix \( \mathbf{U} \) whose value, determined by the BEBD calculation, is given by eq (18). \( \mathbf{U} \) can be used to express \( T_r \) in terms of \( \varphi \) and \( \sigma \). Thus

\[ T_r = \frac{1}{2} \dot{R} \cdot \mathbf{G} \cdot \dot{R} = \frac{1}{2} \dot{\mathbf{P}} \cdot \mathbf{U} \cdot \mathbf{G}^{-1} \mathbf{U} \cdot \dot{\mathbf{P}} \]

The desired quantity \( M_1 \) is simply the matrix element \( (\mathbf{U} \cdot \mathbf{G}^{-1} \mathbf{U})_{\varphi \varphi} \). This is

\[ M_1 = \frac{M_2(M_2 + M_4)m^2 - 2M_2M_4'm + (M_2' + M_4)M_4'n^2}{(n^2 + m^2)M} \]  

(120)

where \( n \) and \( m \) are the bond orders from the BEBD calculation, and \( M \) is the total mass of the molecule.

The bases from which the tunneling parameters \( V_1 \) and \( V_2 \) are measured are taken to be the zero point energies of the reactants and products, respectively, and not the potential minimums as might be expected. The maximum of the potential, on the other hand, is placed at the potential minimum of the complex; i.e., at the saddle point. Johnston [2], pp. 190-196, gives reasons for this particular method of using the Eckart function for tunneling corrections.

We finally have everything needed for eq (3). In the next section explicit rate constant expressions will be given for the four reaction cases of figure 2.

### 2.6. Rate Constant Expressions

The rate constant expression eq (3) is not quite complete. It should be multiplied by the number of equivalent H atoms on the molecule being attacked. Let us call this factor the chemical multiplicity, \( \sigma_{ch} \). For example, there are 6 identical reaction paths for H abstraction of the 6 terminal H atoms on propane, and 2 paths for abstraction of the 2 central H atoms. Thus \( \sigma_{ch} = 6 \) in the first case, and 2 in the second. With this factor added, the rate constants for the four cases shown in figure 2 are

Case III. \( M_2 - M_3 + M_4 - M_2 \ldots M_3 \ldots M_4 \)

\[ k = SF \cdot (2\pi kT)^{3/2} (\Gamma \varphi)^2 \]

Case IVa. \( M_1 - M_2 - M_3 + M_4 - M_1 - M_2 \ldots M_3 \ldots M_4 \)

\[ k = SF_{\varphi\varphi}(F_{\varphi\varphi})^{-1} (2\pi kT)^{3/2} (\Gamma_2^2T_3^2) (\Gamma_2^2)^{-2} \]  

(121)

Case IVb. \( M_2 - M_3 + M_4 - M_2 - M_3 \ldots M_3 \ldots M_4 - M_3 \)

\[ k = SF_{\varphi\varphi}(4\pi)^{-1} \]  

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The calculated factors in S are:

1) \( R^* \) and \( R_e \); these are calculated from \( n \) and \( m \) through Pauling’s relation, eq (9).
2) \( v^* \) is the imaginary frequency obtained from the vibration analysis for the asymmetric stretch.
3) \( |F_r| \) is the determinant of the matrix given by eq (30). It is negative.
4) \( \Gamma_T^* \) is the quantum correction factor for the symmetric stretching frequency obtained from the vibrational analysis.
5) \( \Gamma^* \) is the tunneling correction factor obtained in section 2.5.
6) \( V^* \) is the saddle point potential energy given by the BEBO calculation.

Other calculated factors are:
1) \( F_{\psi_3} \) is the bending force constant given by eq (33).
2) \( F_{\psi_2} \) and \( F_{\psi_4} \) are the bending force constants given by eq (34).
3) The quantum correction factors \( \Gamma_T^*, \Gamma_T^*, \Gamma^* \) for the bending modes come from the frequency analysis via eq (2).

This concludes the theoretical part of this discussion. The next section contains a brief discussion of the computer program which was written to implement the rate constant calculations. This will be followed by instructions on how to use it.

### 3. Computer Implementation of BEBO

The computer program consists of a main section and six subroutines. It is written in an enhanced form of BASIC for use on a Hewlett-Packard 9845A computer.

#### 3.1. Description of the Main Program

The main program begins by reading the following data:

1) Runid$ This is a string variable having up to 79 alphanumeric characters to be used for the run identification.
2) Opt(M), \( M = 1,7 \)
   These are flags which provide a series of available options. These will be described in detail in the instruction section.
3) Ntemp
   This is the number of temperature values at which the rate constant is to be evaluated. A maximum of 16 values will be allowed.
4) Tmin, Tmax
   The minimum and maximum temperature values desired. The reciprocal temperature scale is divided into \( Ntemp - 1 \) equal intervals and the temperature evaluated from the reciprocal values. This gives a better distribution on an Arrhenius plot than if the temperature scale were divided into equal intervals.
5) M1, M2, M3, M4, M5
   These are the five mass point values determined according to the rules given in section 2.1.
6) Ras, Rbs, Rcs, Rds
   These are the equilibrium bond distances for single bonds.
7) Ebs, Ecs, P, Q
8) \( R_{ts}, E_{ts}, \beta \)
These are the bond distance, bond energy, and Morse parameter \( \beta \) for the triplet interaction.

9) \( F_{bs}, F_{cs}, F_{psis}, F_{psi4s} \)
These are the stretching force constants for single \( b \) and \( c \) bonds, and the bending force constants about the M2 and M4 masses.

10) \( S_a, S_b, S_c \)
These are the partition function symmetries \( \sigma_A, \sigma_B, \) and \( \sigma_C \).

11) \( \text{Schem} \)
This is the chemical multiplicity \( \sigma_{ch} \).

12) \( S_{ea}, S_{eb}, S_{ec} \)
These are the electronic degeneracies \( q_{eA}, q_{eB}, \) and \( q_{eC} \).

The program next prints out this input data to provide an easily read record and a check of the numbers.

After these preliminaries, the program then determines the saddle point position. This is done by an iterative procedure; \( n \) is initially set to 0.5; then the potential energy \( V \) is calculated according to eq (14) along with its first and second derivatives, \( V_n \) and \( V_{nn} \), with respect to \( n \). The subroutine Trpl is used to calculate the triplet part of \( V \). A new \( n \) is estimated by the Newton, Raphson method from the formula \( n' = n - V_n/V_{nn} \). The process is repeated using \( n' \) and continued until convergence is obtained. This yields a value of \( n \) which makes \( V \) zero; this will correspond to the desired maximum in \( V \). (I have not investigated the conditions for which a maximum is expected or if there could be more than one maximum.) Having obtained the value of \( n \) for the saddle point, the program calculates the stretching force constant matrix \( F_r \) given by eq (30), its determinant, and the saddle values of \( R_b \) and \( R_c \) from Pauling’s relation eq (9). It then evaluates the mass to be used for tunneling from a somewhat rearranged eq (120). Next, the \( 2 \times 2 \) matrix \( G_r \) is calculated from eq (86). This is then combined with \( F_r \) to form \( H_r \), and the stretching frequencies obtained by solving the resulting quadratic secular equation. The bending frequencies are next determined through the matrices \( F \) (eq (35)) and \( G \) (eq (87)). The sizes of these matrices will depend on the type of reaction. For the three mass point model there is only one element and thus a linear secular equation with one bending frequency. The two four point models require solving a quadratic secular equation for two frequencies. The five point model uses the subroutine Cubic to solve the cubic secular equation for three frequencies. The subroutine Normod then calculates the matrix for the normal coordinate transformation of the stretching modes.

At this point, the program prints out a number of properties of the complex. This will be discussed in detail in the instruction section.

The rate constants are then evaluated from eqs (121) at the different temperatures. The activation energy is gotten by numerically differentiating the logarithm of the rate constant by means of suitable finite difference formulas. Subroutine Fit is a least-squares routine which is used to fit Arrhenius equations through the calculated points. The program concludes with subroutine Pltk which draws an Arrhenius plot of the results.

3.2. Discussion of Subroutine Tun

The only subroutine worth discussing is Tun, the routine for evaluating the integral of eq (112) for the tunneling correction factor \( \Gamma^* \). Johnston and Heicklen [17] calculated this integral numerically by an unspecified method for a range of input parameter values. The three input parameters which they used were \( hv - kT \), where \( v^* = (-F/m)^q(2\pi), \) \( 2\pi V_1(hv^*), \) and \( 2\pi V_1(hv^*) \). Their results are in the form of a table. The method used in the present program is a modified 6-point Gaussian quadrature formula based on Legendre polynomials (see Abramowitz and Stegun [18]). This was used even though the nature of the integral suggests using a formula based on Laguerre polynomials. Neither of these formulas was satisfactory for the whole range of parameter values given by Johnston and Heicklen, so a modification of the first method was developed. It was based on the following ideas: When \( \epsilon \) gets large, the transmission approaches unity. The
idea is to use the Gaussian formula for that part of the integral where $K(e) < 1$. After $K(e)$ has gotten sufficiently close to unity, the remainder of the integral can be evaluated analytically; i.e., if $K(e) = 1$ for $e > e_s$, then
\[\int_{e_b}^{\infty} K(e) e^{-e} \, de \approx \int_{e_b}^{\infty} e^{-e} \, de = e^{-e_b}\]

The problem is to estimate $\epsilon_b$. Let us examine eqs (114) as $e \to \infty$. We get $\alpha_i \to \frac{1}{2} \xi^2$, where $\xi = kT \epsilon/C$. From eq (110), we have
\[K \to 1 - (1 + \Delta \chi \frac{1}{2} \exp(2\pi \xi^{1/2} + \Delta)^{-1} = K_a\]

We can set $K_a$ to some arbitrary value close to unity and solve this equation for $\epsilon$ and then $\xi$ and then $\epsilon_b$ which will be our cutoff point. The result is
\[\epsilon_b = C \left( (2\pi)^{-1} \ln \left[ 2(1 + \Delta)(1 - K_a) \right] \right)^2 \chi(kT)^{-1}\]

It turns out that this value is not entirely satisfactory and subtracting from this the average value of $V_1$ and $V_2$ works better. Also it can happen that $\epsilon_b$ as calculated from this formula can be very large when $K$ is close to unity. Thus, $\exp(-\epsilon_b)$ will be very small. There is no point in using a value for $\epsilon$ as the upper bound to the Gaussian formula if the integrand at this point is negligible because of the exponential factor. Thus $\epsilon_b$ was kept below a certain fixed value $\epsilon_{max}$. This yielded two parameters, $K_a$ and $\epsilon_{max}$ which were adjusted to minimize the squares of the differences between the results of this method and the results of Johnston and Heicklen. The differences averaged 1.3 percent with only two value differing by as much as 6 percent. Such accuracy should be quite adequate for the rate constant calculations.

4. INSTRUCTIONS FOR USING BEBO

4.1. Input

It will be assumed that the reader is familiar with the general operation and command system of the HP9845A. The program lines 5000 to 5240 contain a series of DATA statements which hold the input data. As an example, data for the ethane plus methyl radical reaction is contained in these statements. The general nature of the input has been discussed briefly in the last section; here this is considered in more detail.

1) Runid$ is a string variable containing identifying information; 79 characters can be used.
2) Opt(M), M = 1, 7 are flags for the following options:

Opt(0): This picks out the version of the triplet function $V_i$; these different forms of $V_i$ will be discussed in the Appendix.

Opt(1): As mentioned earlier, the activation energy $E_{act}$ at any temperature is obtained by numerically differentiating the logarithm of the rate constants. This is done in either of two ways. The more accurate method evaluates the rate constant three times at each temperature; at the particular point and slightly above and below the point. The derivative is then estimated from a 3 point finite difference formula. This is automatically the method used when only a single temperature point is requested. The second, less accurate, but faster method uses the rate constants calculated at $N_{temp}$ (see last section or below) points and uses a 5 point difference formula for the derivative. The more points requested and the narrower the temperature range, the more accurate is this method. The value of Opt(1) determines which of these methods will be used. Thus, when

Opt(1) = 1, 5 point difference formula used to get $E_{act}$ (fastest method).
Opt(1) = 2, 3 point difference formula used to get $E_{act}$ (most accurate method).
Opt(2): When
Opt(2)= 1, the natural logarithm of the rate constant is calculated.
Opt(2)= 2, the logarithm, base 10 of the rate constant is calculated.

Opt(3): When
Opt(3)= 1, the cathode ray tube is used for the printout. In this mode, execution of the program pauses before the Arrhenius plot is produced, and before the caption to the plot is generated. In each case execution can be resumed by pressing the "cont" key.
Opt(3)= 0, the internal printer is used for the output.

Opt(4): When
Opt(4)= 1, the rate constant is in cm$^3$/mole-s.
Opt(4)= 2, the rate constant is in cm$^3$/molecule-s.
Opt(4)= 3, the rate constant is in liters/mole-s.
Opt(4)= 4, the rate constant is in liters/molecule-s.

Opt(5): Not used.

Opt(6): When
Opt(6)= 0, the Eckart tunneling correction is not applied. It will automatically not be applied if the zero point energy of the reactants is greater than the potential energy $V^*$ of the saddle point.
Opt(6)= 1, the tunneling correction is applied.

Opt(7): When
Opt(7)= 3, the three parameter Arrhenius type equation, $AT_e^{-E_{act}+RT}$ is fit to the calculated rate constant values.
Opt(7)= 2, the standard two parameter Arrhenius equation $Ae^{-E_{act}+RT}$ is fit to the calculated rate constant values.
3) $N_{temp}$ is the number of temperature values (up to 16) at which the rate constant is to be evaluated. Use the absolute temperature scale.

4) $T_{min}$, $T_{max}$ are the minimum and maximum temperature values to be used. If $N_{temp}= 1$, then only one temperature value should be entered on this line.
5) $M_1, M_2, M_3, M_4, M_5$ are the five mass point values determined by the rules on page 5. For 3 point models set $M_1$ and $M_5$ to zero. The 4 point models will have either $M_1$ or $M_5$ equal to zero. Atomic mass units are to be used.

6) $R_{as}, R_{bs}, R_{cs}, R_{ds}$ are the single bond distances in Å. For 3 point models set $R_{as}$ and $R_{ds}$ to zero. For 4 point models, set either $R_{as}$ or $R_{ds}$ to zero.

7) $E_{bs}, E_{cs}, P, Q$: the first two parameters are the electronic energies for single bonds in kcal/mole. The quantity normally available is the bond dissociation energy $DH^*$ which is defined as the enthalpy change in the process in which one mole of the bond of interest is broken, with reactants and products being in their standard states as ideal gases at 1 atm and 25 °C. This is not the energy we want. The desired energy $E$ is shown in figure 9, which illustrates the energy relationships involved in the removal of an H atom from some group A. $Z_{A-H}$ and $Z_A$ are the zero point energies for the reactant and molecular product, and $H_{A-H}$, $H_A^*$, and $H_{A^*}$ are enthalpies of the species $A-H$, $A^*$, and $H^*$, respectively. In general, a particular enthalpy is the sum of the translational, rotational, vibrational, and $PV$ contributions. We have

$$H^* = H^*_{trans} + H^*_{rot} + H^*_{vib} + PV$$

By examining figure 9 it is easy to derive the relationship between $E$ and $DH^*$. It is

$$E = DH^* + (H_{A-H}^* - H_A^*) + (Z_{A-H} - Z_A) - H_{A^*}.$$  (122)
The second term is
\[ H^t_{A-H} - H^t_A = H^t_{A-H}^{(\text{trans})} - H^t_A^{(\text{trans})} + H^t_{A-H}^{(\text{rot})} - H^t_A^{(\text{rot})} + H^t_{A-H}^{(\text{vib})} \]

Assuming equipartition of energy, the translational and rotational enthalpies will be the same and the difference in vibrational enthalpies will normally be negligible. Thus, the second term in eq (122) can be neglected. The last term
\[ H^t_\text{rot} = E^\text{rot} + PV = 3RT/2 + RT, \]
where \(3RT/2\) is the translational energy of the H atom and RT is PV for an ideal gas. Thus, eq (122) becomes
\[ E = DH^0 = (Z_{A-H} - Z_A^{*}) - 5RT/2 \]

As an example, consider the process \(CH_3^-H+CH_3^+ + H^-\). To estimate the difference in zero point energies between \(CH_3^-H\) and \(CH_3^+\), I have assumed that one C-H stretch of 3100 cm\(^{-1}\) and two H-C-H bends of 1450 cm\(^{-1}\) have been lost in going from A-H to A\(^-\) and H\(^-\). This corresponds to a zero point energy difference of 8.575 kcal. For cases like this, the bond energy will be
\[ E_{cs} = DH^0 + 8.575 - 5RT_298/2 = DH^0 + 7.095 \text{ kcal} \]

The zero point energy difference for other types of bonds can probably be satisfactorily estimated in a similar manner. Having obtained \(E_{bs}\) and \(E_{cs}\) in this manner we can calculate \(P\) and \(Q\) from eq (12).

8) \(Rts,Ets,Beta\) are the triplet interaction parameters in \(\text{A kcal}\) and \(\text{A}^*\), respectively. I have been using the values given in Johnston [1966], table 11-1.
9) \(F_{bs}, F_{cs}, F_{psi2s}, F_{psi4s}\) are the single bond force constants. The first two are the stretching constants in dynes/cm; the second two are bending force constants in dyne-cm. In the 3 mass point case, both the bending force constants are set to zero. For 4 point models, only one of the bending force constants will have a value of zero.

10) \(Sa, Sb, Sc\) are the partition function symmetries for \(A\cdot H, B\cdot\), and \(A\cdot H\cdot B\), respectively.

11) Schem is the chemical multiplicity.

12) \(Sea, Seb, Sec\) are the electronic degeneracies for \(A\cdot H, B',\) and \(A\cdot H\cdot B\). \(Sea\) will normally have the value one. Since \(B\cdot\) and \(A\cdot H\cdot B\) each have an unpaired electron, \(Seb\) and \(Sec\) will normally have the value two.

4.2. Output

BEBO first prints out the input data. It then the following properties of the complex:

1) The potential energy of activation \(V^*\) in kcal/mole.
2) The bond orders \(n\) and \(m\) of the \(b\) and \(c\) bonds.
3) The bond distances \(R_b\) and \(R_c\) in \(\text{Å}\).
4) The force constant in the \(a\) direction in dynes/cm and the angle \(\theta\) makes with the \(R_b\) axis on a contour plot like figure 3.
5) The force constant in the \(\sigma\) in dynes/cm, and the angle to the \(R_b\) axis.
6) The force constant in the unstable normal mode direction in dynes/cm, and the angle to the \(R_b\) axis.
7) The force constant in the stable normal mode direction in dynes/cm, and the angle to the \(R_b\) axis. Note that the normal mode directions are usually not orthogonal.
8) The stretching force constant matrix \(F_r\) in dynes/cm.
9) The equations for transforming back and forth between the normal mode and valence bond coordinates.
10) The bending force constants in dyne-cm.
11) The two stretching frequencies in \(\text{cm}^{-1}\).
12) The one to three bending frequencies in \(\text{cm}^{-1}\).
13) The zero point energy of the complex in kcal/mole.
14) The zero point energy of the reactants in kcal/mole.
15) The zero point energy of the products in kcal/mole.
16) The Eckart potential function parameters \(V_1\) and \(V_2\) in kcal/mole.
17) The reduced mass for tunneling \(M_r = M_{rho}\).
18) The second two of Johnston and Heicklen’s tunneling parameters (see section 2.5).

The program then prints out the rate constants as a function of temperature. Also given at each temperature is the logarithm of the rate constant, the logarithm of the Arrhenius preexponential factor, the activation energy, the tunneling correction factor, and the first of Johnston and Heicklen’s tunneling parameters. Since the tunneling algorithm has not been checked outside the parameter ranges used by Johnston and Heicklen, their parameters values are listed to make sure that they are within the proper ranges. The limits are \(\Delta 1\) and \(\Delta 2 = 0\) to 20, and \(U^* = 0\) to 16.

Finally, there are listed the differences between the calculated values of the logarithm of the rate constant and the values predicted by the least squares fitted Arrhenius equation. This fitted curve is shown by the dotted line on the Arrhenius plot. The fitted Arrhenius parameters are given in the caption of the plot. On the next two pages there is a sample output for the ethane and methyl radical reaction.

5. References

6. APPENDIX: Various Triplet Functions

The subroutine Trpl is able to provide three different triplet functions which are selected according to the value of flag Opt(0). They are as follows:

Opt(0) = 0: This is the modified Sato triplet function with a small portion neglected. Instead of Eq. (13), $V_r = E_r$ is used. This simpler formula seems to have been used in the days of mechanical desk calculators. This option is useful when attempts are being made to reproduce the results of earlier workers.

Opt(0) = 1: Eq. (13) is used.

Opt(0) = 2: Arthur et al. [19] have developed a triplet energy formula by fitting a function to the H-H triplet potential energy values given by Hirschfelder and Linnett [20]. Their formula is

$$V_r = 5.873E_r e^{-1.74\beta(R_s+R_t)(\beta(R_s+R_t))^{1.525}}$$

They claim better results in certain cases when this function is used.
### BEBO Calculations

**Function Identification:**
CH3CH2-H + CH3 = CH3CH2 + H-CH3

**Options Used in Calculations:**
- Modified Sato triplet function.
- Five-point difference formulas used to get activation energy.
- Base 10 logarithm of the rate constant.
- Rate constant units in liters/mole-sec.

**Masses, in Atomic Mass Units:**
- M1 = 17.0510
- M2 = 12.0110
- M3 = 1.0080
- M4 = 12.0110
- M5 = 3.0240

**Single Bond Distances, in Angstroms:**
- R1 = 1.52600, R2 = 1.09000, R3 = 1.09000
- Single Bond Energies of Center Bonds, in kcal; also p & q Parameters.
- E1 = 95.100, E2 = 111.100, p = 1.085, q = 1.093

**Single Bond Energy, Distance, & Morse Parameter for Triplet Interaction.**
- E = 84.400, R = 1.54000, Beta = 1.4250

**Electronic Degeneracy for Species A, B, & C:**
- SA = 1, SB = 1, SC = 1

**Chemical Multiplicity:**
- Scheme: 6

**Pauling's Bond-Order Parameter:**
- Lambda = 0.2300

**Properties of Complex**
- Potential Energy of Activation: V = 14.596 kcal
- Bond Order Parameters: N = 0.5804, M = 0.4196

**Bond Distances for Center Bonds:**
- Rb = 1.24235, Rc = 1.33313 Angstroms

**Force Const. in Rho Direction:**
- Frho = -8.25341E+04 dynes/cm, Angle = -54.13 deg

**Force Const. in Sigma Direction:**
- Fsigma = 2.90227E+05 dynes/cm, Angle = 35.87 deg

**Force Const. in 0 or n.m. direction:**
- Frho = -7.58801E+04 dynes/cm, Angle = -46.42 deg

**F Matrix for Stretches in dynes/cm:**

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<tr>
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<th>1.82224E+05</th>
<th>1.77020E+05</th>
<th>1.77020E+05</th>
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<td>1.94388E+04</td>
<td>1.94388E+04</td>
<td>1.94388E+04</td>
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</tbody>
</table>

**Normal Coordinate Transformations:**
- Cs = 3.2115 Rb + 3.0558 Rc
- Rs = 0.1525 Rb + 0.8279 Rc

**F Matrix Elements for Bends; (these equal the bending force constants):**
- Fpsi1 = 5.30935E-12, Fpsi2 = 7.54513E-13, Fpsi3 = 2.29551E-12 dynes/cm

**Stretching Frequencies:**
- 910.401, 522.47 wave nos.

**Bending Frequencies:**
- 1132.37, 148.99, 440.72 wave nos.

**Zero Point Energy of Complex:**
- 5.689 kcal

**Potential Energy of Reactants:**
- 7.996 kcal

**Potential Energy of Products:**
- 7.478 kcal

**Energy Basis for Eclipsed Tunneling:**
- V1 = 6.690, V2 = 13.117 kcal

**Reduced Mass for Tunneling:**
- Mrho = 0.9314

**Johnston's Tunneling parameters:**
- A1 = 0.812, A2 = 17.279
<table>
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<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>1000/T</td>
<td>T</td>
<td>K</td>
</tr>
<tr>
<td>0.50</td>
<td>2000</td>
<td>1.796E+09</td>
</tr>
<tr>
<td>0.72</td>
<td>1286</td>
<td>1.030E+08</td>
</tr>
<tr>
<td>1.06</td>
<td>947</td>
<td>5.106E+06</td>
</tr>
<tr>
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<td>750</td>
<td>1.036E+06</td>
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<tr>
<td>1.61</td>
<td>621</td>
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<td>529</td>
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<td>462</td>
<td>3.577E+03</td>
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<tr>
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<td>409</td>
<td>6.501E+02</td>
</tr>
<tr>
<td>2.72</td>
<td>367</td>
<td>1.264E+02</td>
</tr>
<tr>
<td>3.00</td>
<td>333</td>
<td>2.599E+01</td>
</tr>
</tbody>
</table>

Figure 1. Plot of logarithm of rate constant as a function of the reciprocal of the temperature.

A least squares fit of the calculated rate constant to the Arrhenius expression, \( k = A \times (T^n) \times \exp(-E_{\text{Arr}}/RT) \), yields the following values for the parameters:

\[
\log(A) = -2.362E+00, \quad n = 3.821, \quad E_{\text{Arr}} = 8.374
\]
This is a listing of the computer program for calculating the rate constants of hydrogen atom transfer reactions according to the bond-energy-bond-order method (customarily referred to as BEBO).

It is written in an enhanced form of the BASIC language for use on a Hewlett-Packard 9845A computer.

```plaintext
10 OVERLAP
20 READ H,K,Lam,Pmass,Econv,Precl,Imax,C,Null
30 DATA 6.6234E-27,1.38033E-16,1.65972E-24,6.94612E-14,1E-9,15,2.99776E10,0
40 Tpfac=2*PI*K*1.4399581E13
50 DIM Runid$(79)
60 DIM Opt(7),Temp(1:16),Tpktkcal(1:16),Ktkcal(1:16),Kqpr(1:16),Eactpr(1:16)
70 DIM Lkqpr(1:16),Lap(1:16),Ustr(1:16),Error(1:16),Par(1:3)
80 COM Delta,Lam,Nbebo,Arb,Rcs,Rts,Betab,Betac,Beta,Es,Ecs,Ets
90 READ Runid$ !Run identification.
110 READ Ntemp !Number of temperature values.
120 IF Ntemp<16 THEN L62
130 PRINT "Error-03"
140 BEEP
150 STOP
160 L62: IF Ntemp<5 THEN Opt(1)=2
170 REDIM Temp(1:Ntemp),Tpktkcal(1:Ntemp),Ktkcal(1:Ntemp),Kqpr(1:Ntemp)
180 REDIM Eactpr(1:Ntemp),Lkqpr(1:Ntemp),Lap(1:Ntemp),Ustr(1:16),Error(1:Ntemp)
190 IF Ntemp=1 THEN READ Tmax !A single temperature entered.
200 IF Ntemp=1 THEN Tmin=1
210 IF Ntemp=1 THEN Opt(1)=2
220 IF Ntemp>1 THEN READ Tmin,Tmax !The minimum and maximum temperatures.
230 Rtmn=1000/Tmax
240 Rtnax=1000/Tmin
250 IF Ntemp>1 THEN Delrt=(Rtnax-Rtmn)/(Ntemp-1)
260 FOR L=1 TO Ntemp
270 Temp(L)=1000/(Rtmn+(L-1)*Delrt)
280 Tpktkcal(L)=Tpfac*Temp(L)
290 Ktkcal(L)=Tpktkcal(L)/(2*PI)
300 NEXT L
310 Delrt=Delrt/1000 !Atomic mass units.
320 READ M1,M2,M3,M4,M5
330 IF M1>0 THEN Mu1=1/M1
340 IF M5>0 THEN Mu5=1/M5
350 READ Ras,Arb,Rcs,Rds !Angstroms.
360 IF Ras>0 THEN Rhoa=1/Ras
370 IF Rds>0 THEN Rhod=1/Rds
380 READ Ebs,Ecs,P,Q !Energies in kcal/mole.
390 READ Rs,Et,Beta
400 READ Fbs,Fcs,Fpsi2s,Fpsi4s !Stretching constants in dynes/cm.
410 READ Sa,Sb,Sc !Bending constants in dynes-cm.
420 READ Schem !Partition function symmetries.
430 READ Schem !Chemical multiplicity.
440 READ Sa,Seb,Sec !Electronic degeneracy.
460 PRINTER IS 16
```
L72: IF Opt(3)=0 THEN PRINTER IS 0
480 PRINT "****************************************************************************** " BEEDO Calculations"
490 PRINT "Run Identification:"
500 PRINT "****************************************************************************** "
510 PRINT "Options Used in Calculations:"
520 PRINT "****************************************************************************** "
530 IF Opt(0)=0 THEN PRINT "Simplified, modified Sato triplet function."
540 IF Opt(0)=1 THEN PRINT "Modified Sato triplet function."
550 IF Opt(0)=2 THEN PRINT "Two-parameter Arthur et.al., triplet function."
560 IF Opt(0)=3 THEN PRINT "Three-parameter Arthur et.al., style triplet function."
570 L64: IF Opt(1)=1 THEN PRINT "Five-point difference formulas used to get activation energy."
580 IF Opt(1)=2 THEN PRINT "Three-point difference formulas used to get the activation energy."
590 IF Opt(2)=1 THEN PRINT "Natural logarithm of the rate constant."
600 IF Opt(2)=2 THEN PRINT "Base 10 logarithm of the rate constant."
610 IF Opt(4)=1 THEN PRINT "Rate constant units in cc/mole-s."
620 IF Opt(4)=2 THEN PRINT "Rate constant units in cc/molecule-s."
630 IF Opt(4)=3 THEN PRINT "Rate constant units in liters/mole-s."
640 IF Opt(4)=4 THEN PRINT "Rate constant units in liters/molecule-s."
650 IF Opt(6)=0 THEN PRINT "Tunneling correction not applied."
660 PRINT "****************************************************************************** "
670 PRINT "Masses, in Atomic Mass Units."
680 PRINT "****************************************************************************** "
690 PRINT USING Format31;M1,M2,M3,M4,M5
700 Format31: IMAGE " M1=",3D.4D"," M2=",3D.4D"," M3=",3D.4D"," M4=",3D.4D"," M5=",3D.4D"
710 PRINT "Single Bond Distances, in Angstroms."
720 PRINT USING Format32;Ras,Rbs,Rcs,Rds
730 Format32: IMAGE " Ras="Z.5D"," Rbs="Z.5D"," Rcs="Z.5D"," Rds="Z.5D"
740 PRINT "Single Bond Energies of Center Bonds, in kcal; also p & q Parameters."
750 PRINT USING Format33;Ebs,Ecs,p,q
760 Format33: IMAGE " Ebs="Z.3D"," Ecs="Z.3D"," p="Z.3D"," q="Z.3D"
770 PRINT "Single Bond Energy, Distance, & Morse Parameter for Triplet Interaction."
780 PRINT USING Format34;Ets,Hts,Beta
790 Format34: IMAGE " Ets="Z.5D"," Hts="Z.5D"," Beta="Z.4D"
800 PRINT "Single Bond Stretching Force Constants in dynes/cm for Center Bonds."
810 PRINT USING Format35;Fbs,Fcs
820 Format35: IMAGE " Fbs="Z.5DE"," Fcs="Z.5DE"
830 PRINT "Single Bond Bending Force Constants in dyne-cm for Outer Masses."
840 PRINT USING Format36;Fpsi2s,Fpsi4s
850 Format36: IMAGE " Fpsi2s="Z.5DE"," Fpsi4s="Z.5DE"
860 PRINT "Partition Function Symmetry Numbers for Species A, B, & C."
870 PRINT USING Format37;Sa,Sb,Sc
880 Format37: IMAGE " Sa="Z"," Sb="Z"," Sc="Z"
890 PRINT "Chemical Multiplicity."
900 PRINT USING Format38;Scheme
910 Format38: IMAGE " Scheme="Z"
920 PRINT "Electronic Degeneracy for Species A, B, & C."
930 PRINT USING Format39;Sea,Seb,Sec
PRINT "Pauling's Bond-Order Parameter."
PRINT USING Format41; Lam
PRINT "Sym=Schem*Sec*Sa*Sb/(Sea*Seb*Sc)"
Fbs=1E-16*Fbs/Econv  !Force constants are converted to kcal energy units.
Fcs=1E-16*Fcs/Econv
Fpsi2s=Fpsi2s/Econv
Fpsi4s=Fpsi4s/Econv
I ************~******************************************************************
N=.5 Icount=0
L11: IF N<0 THEN N=.5*Nold
IF N>1 THEN N=.5*(Nold+1)
Nold=N
CALL TrplfRbs,Rcs,Rts,Ets,Beta,Lam,N,Vt,Vtn,Vtnn,Vtr,Vtrr,OptOJJ
M=1-N
V=Ebs*(1-N*P)-Ecs*M*Q+Vt
Vn=-Ebs*P*N*(P-1)+Ecs*Q*M*(Q-1)+Vtn
Vnn=-Ebs*P*(P-1)*N*(Q-2)-Ecs*Q*(Q-1)*M*(Q-2)+Vtnn
N=Nold-Vn/Vnn
IF ABS[(N-Nold)/Nold]<Prec1 THEN L10
Icount=Icount+1
IF Icount>Imax THEN L12
GOTO L11
L12: PRINT "Error-01"
STOP
L10: N=Nbebo=Nold  *************************************************************************
Next, the stretching part of the force constant matrix is calculated.
Nsq=N*M*N
Msq=M^2
N2m2=Nsq+Msq
Nm=N*M
Frho=Vnn*Nm^-2/(N2m2*Lam^2)
Fsigma=(Fbs*N^3+Fcs*M^3+Vtrr)/N2m2
DIM Fr[1:2,1:2]  
Fr[1,1]=Frho*Msq+Fsigma*Nsq/N2m2 
Fr[1,2]=Fr[2,1]=Frho*Msq+Fsigma*Nsq/N2m2 
Fr[2,2]=Frho*Msq+Fsigma*Msq/N2m2
Rb=Rbs-Lam*LOG(N) 
Rc=Rcs-Lam*LOG(M)
Dfr=DET(Fr)
Dfr=SGN(Dfr)*SQR(ABS(Dfr))
Jfac=(Rb*Rc/Rbs)^2
Rhob=1/Rb
Rhoc=1/Rc
Cc=-Nbebo/[1-Nbebo]
1440 \text{Ma} = \text{M1} + \text{M2}
1450 \text{Mx} = \text{M3}
1460 \text{Mb} = \text{M4} + \text{M5}
1461 \text{Maxb} = \text{Ma} + \text{Mx} + \text{Mb}
1470 \text{Mrho} = \frac{\text{Ma} \cdot \text{Mb} \cdot (1 + \text{Cc}^2) + \text{Mb} \cdot \text{Mx} \cdot \text{Cc}^2 + \text{Ma} \cdot \text{Mx}}{\text{Maxb} \cdot (1 + \text{Cc}^2)}
1471 \text{DIM} \ 	ext{Azr}[1:3,1:2], \text{Alq}[1:3,1:2]
1472 \text{Azr}[1,1] = \frac{-\text{Mb} + \text{Mx}}{\text{Maxb}}
1473 \text{Azr}[1,2] = \text{Azr}[2,2] = \frac{-\text{Mb}}{\text{Maxb}}
1474 \text{Azr}[2,1] = \text{Azr}[3,1] = \frac{\text{Ma}}{\text{Maxb}}
1475 \text{Azr}[3,2] = \frac{\text{Ma} + \text{Mx}}{\text{Maxb}}
1480 \text{Mrho} = 1

1500 \text{Next, the Gr matrix is calculated. This will hold for 3,4, & 5 atom models with rigid bonds on the ends.}
1510 \text{First, the type of model being used is determined. (Note, all are linear)}
1520 \text{DIM} \text{Gr}[1:2,1:2]
1530 \text{Model} = 5
1540 \text{IF} \ (\text{M1} = 0) \ \text{OR} \ (\text{M5} = 0) \ \text{THEN} \ \text{Model} = 4
1550 \text{IF} \ (\text{M1} = 0) \ \text{AND} \ (\text{M5} = 0) \ \text{THEN} \ \text{Model} = 3
1570 \text{Model} = \text{Model} - 2
1580 \text{Mu} 2 = \frac{1}{\text{M2}}
1590 \text{Mu} 3 = \frac{1}{\text{M3}}
1600 \text{Mu} 4 = \frac{1}{\text{M4}}
1610 \text{Gr}(1,1) = \text{Mu} 2 / (1 + \text{M1} / \text{M2}) + \text{Mu} 3
1620 \text{Gr}(1,2) = \text{Gr}(2,1) = -\text{Mu} 3
1630 \text{Gr}(2,2) = \text{Mu} 3 + \text{Mu} 4 / (1 + \text{M5} / \text{M4})
1640 \text{Stretching frequencies evaluated.}
1650 \text{DIM} \text{Hr}[1:2,1:2]
1660 \text{MAT} \text{Hr} = \text{Gr} \cdot \text{Fr}
1670 \text{Bh} = \text{Hr}[1,1] + \text{Hr}[2,2]
1680 \text{Ch} = \text{Hr}[1,1] \cdot \text{Hr}[2,2] - \text{Hr}[1,2] \cdot \text{Hr}[2,1]
1690 \text{Dh} = \text{SQR}(\text{Bh}^2 - 4 \cdot \text{Ch})
1700 \text{DIM} \text{Evs}[1:2], \text{Fq} = \text{Fr}[1:2], \text{Ls}[1:2]
1710 \text{Evs}[1] = 0.5 \cdot (\text{Bh} + \text{Dh})
1720 \text{Evs}[2] = 0.5 \cdot (\text{Bh} - \text{Dh})
1730 \text{FOR} \text{I} = 1 \ \text{TO} \ 2
1740 \text{Fq}[\text{I}] = \text{SGN}(\text{Evs}[\text{I}]) \cdot 682.427 \cdot \text{SQR}(\text{ABS}(\text{Evs}[\text{I}])) / (2 \cdot \text{PI}) \ \text{In} \ 1/\text{cm.}
1750 \text{NEXT} \text{I}
1760 \text{Bending frequencies now calculated.}
1770 \text{Fps} 13 = -\text{Rb} \cdot \text{Ac} \cdot \text{Vtr} / (\text{Rb} + \text{Ac})
1780 \text{DIM} \text{G}[1:3,1:3], \text{F}[1:3,1:3], \text{H}[1:3,1:3], \text{Evb}[1:3], \text{Frqb}[1:3], \text{Lb}[1:3]
1790 \text{REDIM} \text{G}[1:3,1:3], \text{F}[1:3,1:3], \text{H}[1:3,1:3], \text{Evb}[1:3], \text{Frqb}[1:3], \text{Lb}[1:3]
1800 \text{REDIM} \text{Ev} = [1:Model[2.1,1:Model[2], \text{H}[1:Model[2,1,1:Model[2]]
1810 \text{REDIM} \text{Ev} = [1:Model[2], \text{Frqb}[1:Model[2], \text{Lb}[1:Model[2]]
1820 \text{ON} \text{Mode} = 2 \ \text{GOTO} \ \text{La} 3, \text{La} 4, \text{La} 5
1830 \text{I} \ \text{Three atom model.}
1840 \text{La} 3: \text{G}[1,1] = \text{Mu} 2 \cdot \text{Rho}^2 + \text{Mu} 4 \cdot \text{Rho}^2 + \text{Mu} 3 \cdot (\text{Rho} + \text{Rho})^2
1850 \text{F}[1,1] = \text{Fps} 13
1860 \text{MAT} \text{H} = \text{G} \cdot \text{F}
Evb(1) = H(1, 1) 
GOTO L20

I ..............................

I Four atom models.
La4: IF M1 = 0 THEN La4sub
G(1, 1) = Mu1*Rhoa^2 + Mu3*Rhob^2 + Mu2*(Rhoa + Rhob)^2
G(2, 1) = G(2, 1) = Rhoc*(Mu2 + (Rhoa + Rhob)*Mu3)
G(2, 2) = Mu3*Rhob^2 + Mu4*Rhoc^2 + Mu2*(Rhoa + Rhob)^2
G(1, 1) = Fpsi2 = Fpsi2s*N
G(1, 2) = F(2, 1) = 0
G(2, 2) = Fpsi3
MAT H = G*F
GOTO La4end

La4sub: G(1, 1) = Mu2*Rhob^2 + Mu4*Rhoc^2 + Mu3*(Rhoa + Rhob)^2
G(2, 1) = G(2, 1) = Rhoc*(Mu2 + (Rhoa + Rhob)*Mu3)
G(2, 2) = Mu3*Rhob^2 + Mu4*Rhoc^2 + Mu2*(Rhoa + Rhob)^2
G(1, 1) = Fpsi2 = Fpsi2s*N
G(1, 2) = F(2, 1) = 0
G(2, 2) = Fpsi3
MAT H = G*F
GOTO La4end

Bending frequencies.
L20: FOR I = 1 TO Model2
Freq(I) = 8B^2 * 427 * SQRT(Evb(I)) / (2*PI)
2380 NEXT I
2390 PRINT "Properties of Complex."
2400 PRINT USING Format1; V
2410 Format1: IMAGE "Potential Energy of Activation: V=" DD.DD, kcal"
2420 PRINT USING Format2; N, M
2430 Format2: IMAGE "Bond Order Parameters: N=" ZZ.DD, " M=" ZZ.DD
2440 PRINT USING Format3; Rb, Rc
2450 Format3: IMAGE "Bond Distances for Center Bonds: Rb=" ZZ.DD," Rc=" ZZ.DD, " Angstroms"
2460 PRINT USING Format4; Frho*Econv*1E16, ATN(Cc)*180/PI
2470 Format4: IMAGE "Potential Energy of Activation: V=" DD.DD," kcal"
2480 PRINT USING Format5; Fsigma*Econv*1E16, ATN(-1/Cc)*180/PI
2490 Format5: IMAGE "Bond Order Parameters: N=" ZZ.DD, " M=" ZZ.DD
2500 DIM Lqi(1:2,1:2), Lq(1:2,1:2)
2510 CALL NormodlHr(*J, Evs(*J, Fr(*J, Lqi(*J))
2520 MAT Lq=INV(Lqi)
2530 DEF FNGg(Zr)=Fr(1,1)*COS(Zr)+2*Fr(1,2)*SIN(Zr)*COS(Zr)+Fr(2,2)*SIN(Zr)^2
2540 Z=ATN(Z)
2550 PRINT USING Format4a; FNGg(Zr)*Econv*1E16, Zr*180/PI
2560 Format4a: IMAGE "Potential Energy of Activation: V=" DD.DD," kcal"
2570 MAT Alq=Azr*Lq
2580 PRINT Alq"
2590 DIM Rcomplex(1:2), Zeq(1:3)
2595 MAT Zeq=Azr*Rcomplex
2600 PRINT Zeq"
2610 PRINT "F Matrix for Stretches in dyne/cm"
2620 PRINT USING Format5; Fr(1,1)*Econv*1E16, Fr(1,2)*Econv*1E16
2630 PRINT USING Format6; Fr(2,1)*Econv*1E16, Fr(2,2)*Econv*1E16
2640 PRINT "Normal Coordinate Transformations"
2650 PRINT USING Format45; Lqi(1,1), Lqi(1,2), Lq(1,1), Lq(1,2)
2660 Format45: IMAGE " Qs=" ZZ.DD, Rb=" ZZ.DD," Rc=" ZZ.DD
2670 PRINT Using Format46; Lqi(2,1), Lqi(2,2), Lq(2,1), Lq(2,2)
2680 Format46: IMAGE " Qr=" ZZ.DD, Rb=" ZZ.DD," Rc=" ZZ.DD
2690 PRINT "F Matrix Elements for Bends; [these equal the bending force constants]"
2695 PRINT USING Format6; Fpsi2*Econv, Fpsi3*Econv, Fpsi4*Econv
2700 Format6: IMAGE " Fpsi2=" ZZ.DD," Fpsi3=" ZZ.DD," Fpsi4=" ZZ.DD, " dyne-cm"
2705 PRINT Using Format7; Frqs(2), Frqs(1)
2710 Format7: IMAGE "Stretching Frequencies: " 4D.DD," i, " 4D.DD, " wave nos."
2700 IF Model=3 THEN PRINT USING Format8; Frqbf(1), Null, Null
2710 IF Model=4 THEN PRINT USING Format8; Frqbf(1), Frqbf(2), Null
2720 IF Model=5 THEN PRINT USING Format8; Frqbf(1), Frqbf(2), Frqbf(3)
2740 Zcmplx=Frqbf(1)
2750 FOR I=1 TO Model2
2760 Zcmpl=
2770 NEXT I
2780 Zcmpl=.5*Zcmpl*2.8585E-3
2790 PRINT USING Format9; Zcmpl
2800 Format9: IMAGE "Zero Point Energy of Complex=",3D.3D. kcal"
2810 I The rate constants are now evaluated at the different temperatures.
2820 Unit=1
2830 ON Opt(4) GOTO Lunit1,Lunit2,Lunit3,Lunit4
2840 Lunit2: Unit=6.02E23
2850 GOTO Lunit1
2860 Lunit3: Unit=1000
2870 GOTO Lunit1
2880 Lunit4: Unit=6.02E26
2890 GOTO Lunit1
2900 Lunit1: I Continue
2910 DEF FN0(F)=Z*EXP(-.5*Z)/(1-EXP(-Z))
2920 DIM Lograte(-1 :11
2930 NRt=0
2940 IF Opt(1)=1 THEN L61
2950 Oert=1.987E-3*.05/V
2960 Npt=1
2970 L61: Evbs=Fbs*{(Mu2+Mu3)
2980 Frqbs=682.427*SQR(Evbs)/(2*PI)
2990 Zreact=Frqbs/2
3000 Zprod=Frqbs/2
3010 IF Model=3 THEN L40
3020 Frqcs=Fcs*{(Mu3+Mu4)
3030 Frqcs=682.427*SQR(Evcs)/(2*PI)
3040 Zreact=Zreact+Frqcs/2
3050 IF Opt(6)=0 THEN Tunnel=0
3060 L40: IF Opt(6)=0 THEN L85
3070 Zreact=Zreact*2.8585E-3
Zprod = Zprod * 2.8585E-3
PRINT USING Format10; Zreact
PRINT USING Format11; Zprod
V1 = V - Zreact
Tunnel = 1
IF V1 > 0 THEN LB4
Tunnel = 0
PRINT "Eckart tunneling factor not applied because zero point energy of reactants is" 
GOTO LB5
LB4: V2 = V - Ebs + Ecs - Zprod
PRINT USING Format12; V1, V2
PRINT USING Format13; Mrho
PRINT "Johnston's tunneling factors calculated. Nu = SQR[-Ev(2)*Econv*1E16/(Mrho*Pmass)]/2*PI)
A1 = V1*Econv*2*PI/[H*Nu)
A2 = V2*Econv*2*PI/[H*Nu)
Tunnel = 0
PRINT "Eckart tunneling factor not calculated because A1 or A2 is greater then 20.",
GOTO LB5
L300: PRINT USING Format16; A1, A2
PRINT "Johnston's tunneling parameters: A1 = " DZ.3D", A2 = " DZ.3D
PRINT "Calculating rate const. at " DD.3D", "th temperature value out of DD.3D values.
Rtemp = 1/Temp[L]
FOR Nt = -Npt TO Npt
Rt = Rtemp + Nt*Delrt
T = 1/Rt
Tpk*tcalsub = Tpfac*T
Ktckalsub = Tpk*tcalsub/(2*PI)
Kfac = EXP[-V/Ktckalsub]*Sym*Jfac*C*Frqs(2)*SQR[Fbs]/(Dfr*Unit)
Tf = 1.439/T
ON Model2 GOTO Lb3, Lb4, Lb5
Lb3: Kcl = Kfac*Tpk*tcalsub*1.5/Fpsi3
G3 = FNG(Tf*Frqs[1])*FNG(Tf*Frq[1])^2/FNG(Tf*Frq[2])
Kg = G3*Kcl
GOTO L30
Lb4: IF M1 = 0 THEN Lb4sub
Kcl = Kfac*Tpk*tcalsub*1.5*Fpsi2s/(Fpsi2s*Fpsi3)
G4a = FNG(Tf*Frqs[1])*(FNG(Tf*Frq[1])*FNG(Tf*Frq[2]))^2
G4a = G4a / (FNG(Tf*Frqbs) * FNG(Tf*Frqpsi^2s)^2)
Kg = G4a * KcL
GOTO L30
L4bsub: Kg = Kg / (FNG(Tf*Frqbs) * FNG(Tf*Frqpsi^2s)^2) * Kg
GOTO L30
G4b = FNG(Tf*Frqbs) * (FNG(Tf*Frqpsi^2s) * FNG(Tf*Frqbs))^2
Kg = G4b * KcL
GOTO L30
L5: Kg = Kg / (FNG(Tf*Frqbs) * FNG(Tf*Frqpsi^2s)^2) * Kg
GOTO L30
Gam = 1
IF Tunnel = 0 THEN L87
CALL TunL(V1*Econv, V2*Econv, Frho*Econv*1E16, Mrho*Pmass, T, Gam)
KcL = 6.025E23 * Kcl
Kq = 6.025E23 * 1E-24 * KcL
IF Nt = 0 THEN Kqtab = Kg
IF Nt = 0 THEN Gamtab = Gam
Lograte(Nt) = LOG(Kq)
GOTO L59
IF Npt = 0 THEN L59
Eactpr(L) = -1.987E-3 * (-Lograte(-1) + Lograte(1)) / (2*Delrt)
IF Opt(3) = 1 THEN Lapr(L) = LOG(Kqtab * EXP(Eactpr(L) / Ktkcal[L]))
IF Opt(3) = 2 THEN Lapr(L) = LGT(Kqtab * EXP(Eactpr(L) / Ktkcal[L]))
L59: Kqpr(L) = Kgtab
IF Opt(3) = 1 THEN Kqpr(L) = Lograte(0)
IF Opt(3) = 2 THEN Kqpr(L) = LGT(Kqtab)
Ustr[L] = Nu*H / (K*Temp[L])
Gampr(L) = Gamtab
NEXT L
IF Opt(3) = 0 THEN PRINTER IS 0
IF Npt = 1 THEN L60
DIM DD, D1, D1(1:5), D1(1:5)
D0(1) = -2.06333333333
D0(2) = 4
D0(3) = -3
D0(4) = -1.33333333333
D0(5) = -0.25
D0(6) = 1.5
D0(7) = 0.5
D0(8) = 8.33333333333E-2
D0(9) = 8.33333333333E-2
D0(10) = 8.33333333333E-2
D0(11) = 8.33333333333E-2
D0(12) = -6.666666666667
D0(13) = 0
D0(14) = 0
D0(15) = 0
D1 = 0
D1(1) = 0
D1(2) = 0
D1(3) = 0
D1(4) = 0
D1(5) = 0
D1(6) = 0
D1(7) = 0
D1(8) = 0
D1(9) = 0
D1(10) = 0
D1(11) = 0
D1(12) = 0
D1(13) = 0
D1(14) = 0
D1(15) = 0
MAT Eactpr = ZER
FOR J=1 TO 5
Eactpr(1)=Eactpr(1)+0(J)*Lkqpr(J)
Eactpr(2)=Eactpr(2)+0(J)*Lkqpr(J)
Eactpr(Ntemp-1)=Eactpr(Ntemp-1)-D(J)*Lkqpr(Ntemp-J+1)
Eactpr(Ntemp)=Eactpr(Ntemp)-D(J)*Lkqpr(Ntemp-J+1)
FOR Na=3 TO Ntemp-2
Eactpr(Na)=Eactpr(Na)+D(J)*Lkqpr(Na+J-3)
NEXT Na
END

FOR J=1 TO 5
Eactpr(1)=Eactpr(1)-1.987E-3/Delrt
Eactpr(2)=Eactpr(2)-1.987E-3*LOG(10)/Delrt
IF Opt(2)=1 THEN MAT Eactpr=Eactpr * (-1.987E-3/De L rt J
IF Opt(2)=2 THEN MAT Eactpr=Eactpr*[-1.987E-3*LOG[101/0eLrt]
NEXT J

NEXT L

FOR L=1 TO Ntemp
Afac=Kqpr(L)*EXP(Eactpr(L)/Ktkca(L))
IF Opt(2)=1 THEN Lapr(L)=LOG(Afac)
IF Opt(2)=2 THEN Lapr(L)=GT(Afac)
NEXT L

IF Opt(3)=0 AND (Ntemp>1) THEN PRINT PAGE

PRINT "Temp., Rate Const., A factor, Act. Energy, Eckart U* Factor, Deviation from Fit"
FOR Opt(2)=1 THEN PRINT "1000/T T K Ln(K) Ln(A) Eact Gam U*
Fit"
IF Opt(2)=2 THEN PRINT "1000/T T K Log(K) Log(A) Eact Gam U*
Fit"
NEXT L

PRINT USING Format14;1000/Temp(L),Temp(L),Kqpr(L),Lkqpr(L),Lapr(L),Eactpr(L),Gampr(L),Us
tr(L),Error(L)

EXIT GRAPHICS

PRINT "Figure 1. Plot of logarithm of rate constant as a function of the reciprocal
of the temperature."
PRINT "A least squares fit of the calculated rate constant to the Arrhenius expression,"
PRINT "k=A*EXP[-Earr/RT], yields the following values for the parameter values:")
PRINT "k=A*[T'n]*EXP(-Earr/RT), yields the following values for the parameter values:"
5000 I Run Identification
5010 DATA CH3CH2-H + CH3 = CH3CH2 + H-CH3
5020 I Options:
5021 I 0,1,2,3,4,5,6,7
5030 DATA 1,1,2,1,3,0,1,3
5040 I Number of temperature values.
5050 DATA 10
5060 I Minimum and maximum temperatures, (or a single temperature if Ntemp=1).
5070 DATA 333.3333333,2000
5080 I Masses: M1, M2, M3, M4, M5
5090 DATA 17.051,12.011,1.008,12.011,3.024
5100 I Single bond distances: Ra, Rb, Rc, Rd
5110 DATA 1.526,1.09,1.09,1.09
5120 I Bond energies: Ebs and Ecs; BEBO parameters: p and q
5130 DATA 105.1,111.1,1.0852,1.0925
5140 I Triplet single bond distance, bond energy, and Morse parameter:
5150 I Rts, Ets, and Beta
5160 DATA 1.54,84.4,1.425
5170 I Stretching constants Fbs, Fcs and bending constants Fpsi2s, Fpsi4s
5180 DATA 4.79E5,4.79E5,9.14837E-12,5.4653E-12
5190 I Partition function symmetries for species A, B, and C
5200 DATA 1,1,1
5210 I Chemical multiplicity
5220 DATA 6
5230 I Electronic degeneracies for species A, B, and C
5240 DATA 1,2,2
SUB Trpl(Rbs, Rcs, Rts, Ets, Beta, Lam, N, Vt, Vtn, Vtnn, Vtr, Vtrr, Version)

<table>
<thead>
<tr>
<th>Line</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>5400</td>
<td>SUB Trpl(Rbs, Rcs, Rts, Ets, Beta, Lam, N, Vt, Vtn, Vtnn, Vtr, Vtrr, Version)</td>
</tr>
<tr>
<td>5402</td>
<td>Version=0; Simplified, modified Sato function.</td>
</tr>
<tr>
<td>5404</td>
<td>Version=1; Modified Sato function.</td>
</tr>
<tr>
<td>5406</td>
<td>Version=2; Arthur et al., function (2 parameter form).</td>
</tr>
<tr>
<td>5410</td>
<td>Delrs=Rbs+Rcs-Rts</td>
</tr>
<tr>
<td>5412</td>
<td>IF Version=2 THEN Delrs=Delrs+Rts</td>
</tr>
<tr>
<td>5414</td>
<td>M=1-N</td>
</tr>
<tr>
<td>5416</td>
<td>R=Delrs-Lam<em>LOG(N</em>N*M)</td>
</tr>
<tr>
<td>5418</td>
<td>Rn=Lam*[1/M-1/N]</td>
</tr>
<tr>
<td>5420</td>
<td>Rnn=Lam*[1/N-2+1/M-2]</td>
</tr>
<tr>
<td>5422</td>
<td>IF (Version=1) OR (Version=0) THEN L10</td>
</tr>
<tr>
<td>5426</td>
<td>P1=5.873</td>
</tr>
<tr>
<td>5428</td>
<td>P2=1.747</td>
</tr>
<tr>
<td>5430</td>
<td>P3=1.525</td>
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<tr>
<td>5440</td>
<td>D=Ets<em>P1</em>Beta*P3</td>
</tr>
<tr>
<td>5442</td>
<td>E=EXP(-P2<em>Beta</em>R)</td>
</tr>
<tr>
<td>5444</td>
<td>Er=-P2<em>Beta</em>E</td>
</tr>
<tr>
<td>5446</td>
<td>En=Er*Rn</td>
</tr>
<tr>
<td>5448</td>
<td>Enn=-P2<em>Beta</em>[Rnn<em>E+Rn</em>En]</td>
</tr>
<tr>
<td>5450</td>
<td>Err=-P2<em>Beta</em>Er</td>
</tr>
<tr>
<td>5452</td>
<td>Rp3=R*P3</td>
</tr>
<tr>
<td>5454</td>
<td>Vt=D*P3</td>
</tr>
<tr>
<td>5456</td>
<td>Vtn=D<em>Rp3</em>{En+E<em>P3</em>Rn/R}</td>
</tr>
<tr>
<td>5458</td>
<td>Vtnn=D<em>Rp3</em>{En+P3*(2<em>En+E</em>(P3-1)*Rn/R)<em>Rn+E</em>Rnn}/R</td>
</tr>
<tr>
<td>5460</td>
<td>Vtr=D<em>Rp3</em>{Er+P3*E/R}</td>
</tr>
<tr>
<td>5462</td>
<td>Vtrr=D<em>Rp3</em>{Err+2<em>P3</em>E/R+P3*(P3-1)*E/R^2}</td>
</tr>
<tr>
<td>5464</td>
<td>SUBEXIT</td>
</tr>
<tr>
<td>5466</td>
<td>L10: E=.5<em>EXP(-Beta</em>R)</td>
</tr>
<tr>
<td>5468</td>
<td>Er=-Beta*E</td>
</tr>
<tr>
<td>5470</td>
<td>En=Er*Rn</td>
</tr>
<tr>
<td>5472</td>
<td>Enn=-Beta*[Rnn<em>E+Rn</em>En]</td>
</tr>
<tr>
<td>5474</td>
<td>Err=-Beta*Er</td>
</tr>
<tr>
<td>5476</td>
<td>IF Version=0 THEN L15</td>
</tr>
<tr>
<td>5478</td>
<td>Vt=Ets<em>E</em>[1+E]</td>
</tr>
<tr>
<td>5480</td>
<td>Efac=1.2+E</td>
</tr>
<tr>
<td>5482</td>
<td>Vtn=Ets<em>En</em>Efac</td>
</tr>
<tr>
<td>5484</td>
<td>Vtr=Ets<em>Er</em>Efac</td>
</tr>
<tr>
<td>5486</td>
<td>Vtnn=Ets<em>En</em>Efac</td>
</tr>
<tr>
<td>5488</td>
<td>Vtrr=Ets<em>Er</em>Efac</td>
</tr>
<tr>
<td>5500</td>
<td>Vtrr=Ets*Err</td>
</tr>
<tr>
<td>5502</td>
<td>SUBEND</td>
</tr>
</tbody>
</table>
5600 SUB Cubic(Cf(*),Rt(*))
5602 R=-Cf(0)
5604 Q=-Cf(1)
5606 P=-Cf(2)
5608 A=(3*Q-P^2)/3
5610 B=(2*P^3-9*P*Q+27*R)/27
5612 Fc1=A^3/27
5614 Fc=B^2/4+Fc1
5616 IF Fc<0 THEN L89
5618 PRINT "Error-005"
5620 STOP
5622 L89: Psi=-B/(2*SQR(-Fc1))
5624 DEG
5626 Psi=ACS(Psi)/3
5628 Fc=2*SQR(-A/3)
5630 Rt(1)=Fc*COS(Psi)-P/3
5632 Rt(2)=Fc*COS(Psi+120)-P/3
5634 Rt(3)=Fc*COS(Psi+240)-P/3
5636 RAD
5638 SUBEND
5700 SUB Normod(H(*),Ev(*),F(*),Li(*))
5710 DIM H(1:2,1:2),Ev(1:2,1:2),F(1:2,1:2),G(1:2,1:2),L(1:2,1:2)
5720 FOR I=1 TO 2
5730 G(1,I)=-H(1,2)/(H(1,1)-Ev(I))
5740 G(2,I)=1
5750 NEXT I
5760 MAT Fg=F*G
5770 MAT Gt=TRN(G)
5780 MAT Gfg=Gt*Fg
5790 FOR I=1 TO 2
5800 L(2,I)=SQR(Ev(I)/Gfg(I,I))
5810 L(1,I)=G(1,I)*L(2,I)
5820 NEXT I
5830 MAT Li=INV(L)
5840 SUBEND
6000 SUB Tun1(V1, V2, F, M, T, Gam) ! 6 point Gaussian Legendre
6005 OPTION BASE 1
6010 DIM X(6), W(6)
6015 DATA .238619186083, .661209386466, .932469514203
6020 FOR N = 4 TO 6
6025 READ X(N)
6030 X(7 - N) = -X(N)
6035 NEXT N
6040 DATA .467913934573, .360761573048, .171324492379
6045 FOR N = 4 TO 6
6050 READ W(N)
6055 W(7 - N) = W(N)
6060 NEXT N
6065 H = 6.6234E-27
6070 K = 1.38033E-16
6075 Kt = K * T
6080 A = V1 - V2
6085 V1h = SQRT(V1)
6090 V2h = SQRT(V2)
6095 B = [V1h + V2h] * 2
6100 Pi2 = 2 * PI
6105 L = Pi2 * SQRT(-2/F) / [1/V1h + 1/V2h]
6110 C = H * 2 / ([B * M * L * 2]
6115 DEF FNCosh(Z) = .5 * (EXP(Z) + EXP(-Z))
6120 Delta = (B - C) / C
6125 IF Delta < 0 THEN L10
6130 Dfac = FNCosh(PI * SQRT(Delta))
6135 GOTO L11
6140 L10: Dfac = COS(PI * SQRT(-Delta))
6145 L11: IF V2 > V1 THEN EO = -V1 / Kt
6150 IF V1 > V2 THEN EO = -V2 / Kt
6155 Vav = 5 * (V1 + V2)
6160 Eb1 = (C * LOG([2 * (1 + Dfac)] / .014) / Pi2) * 2 - Vav) / Kt
6165 Eb2 = 3.2
6170 Eb = MIN(Eb1, Eb2)
6175 Em = .5 * (Eb - EO)
6180 Ep = .5 * (Eb + EO)
6185 Gam = 0
6190 FOR N = 1 TO 6
6195 E = Em * X(N) + Ep
6200 Kt = Kt * E
6205 Alph1 = PI * SQRT([Kt + V1] / C)
6210 Alph2 = PI * SQRT([Kt + V2] / C)
6215 Facp = FNCosh(Alph1 + Alph2)
6220 Facm = FNCosh(Alph1 - Alph2)
6225 Ke = (Facp - Facm) / (Facp + Dfac)
6230 Y = Ke * EXP(-E)
6235 Gam = Gam + W(N) * Y
6240 NEXT N
6245 L50: Gamfac = EXP(-Eb)
6250 Gam = Em * Gam + Gamfac
6255 SUBEND
SUB Pltk(Opt(*), Nm, T(*), Lk(*), E(*))

DEF

PLOTTER IS 13, "GRAPHICS"

GRAPHICS

LOCATE 19, 119, 10, 95

DIM Xtics(1:11), Ytics(1:11)

DATA .01, .02, .025, .05, .1, .2, .25, .5, 1, 2, 2.5

MAT READ Xtics

Xmin = 1000 / T(1)

Xmax = 1000 / T(Nm)

Xspan = Xmax - Xmin

FOR I = 1 TO 11

Itab = I

IF INT(Xspan / Xtics(I)) <= 6 THEN L30

NEXT I

L30: Xtic = Xtics(Itab)

Xstart = INT(Xmin / Xtic) * Xtic

Xstop = INT(Xmax / Xtic) * Xtic

IF (Xstop > Xmax) AND (Xstop - Xmax > 1E-2 * Xtic) THEN Xstop = (INT(Xmax / Xtic) + 1) * Xtic

Ymin = Ymax = Lk(1)

FOR N = 2 TO Nm

IF Ymin > Lk(N) THEN Ymin = Lk(N)

IF Ymax < Lk(N) THEN Ymax = Lk(N)

NEXT N

Yspan = Ymax - Ymin

MAT Ytics = Xtics

FOR I = 1 TO 11

If INT(Yspan / Ytics(I)) <= 6 THEN L40

NEXT I

L40: Ytic = Ytics(Itab)

Ystart = INT(Ymin / Ytic) * Ytic

Ystop = (INT(Ymax / Ytic) + 1) * Ytic

SCALE Xstart, Xstop, Ystart, Ystop

LINE TYPE 3

GRID Xtic, Ytic, Xstart, Ystart

LINE TYPE 1

AXES Xtic, Ytic, Xstart, Ystart, 2, 2, 6

FRAME

CSIZE 3

LOG 8

FOR Ypos = Ystart TO Ystop STEP Ytic

MOVE Xstart, Ypos

LABEL USING "MDT.DDX"; Ypos

NEXT Ypos

LOG 6

FOR Xpos = Xstart + Xtic TO Xstop STEP Xtic

MOVE Xpos, Ystart - Ytic / 10

LABEL USING "Z.DD"; Xpos

NEXT Xpos
6598  LDIR  90
6600  CSIZE  4.5
6602  LORG  1
6604  ON Opt(2)  GOTO  L1, L2
6606  L1:  ON Opt(4)  GOTO  La1, La2, La3, La4
6608  La1:  MOVE  Xstart-Xspan*.15, Ystart+Yspan*.27
6610  LABEL  USING "K"; "Ln(k) cc/mole-s"
6612  GOTO  Lout
6614  La2:  MOVE  Xstart-Xspan*.15, Ystart+Yspan*.18
6616  LABEL  USING "K"; "Ln(k) cc/molecule-s"
6618  GOTO  Lout
6620  La3:  MOVE  Xstart-Xspan*.15, Ystart+Yspan*.18
6622  LABEL  USING "K"; "Ln(k) liters/mole-s"
6624  GOTO  Lout
6626  La4:  MOVE  Xstart-Xspan*.15, Ystart+Yspan*.12
6628  LABEL  USING "K"; "Ln(k) liters/molecule-s"
6630  GOTO  Lout
6632  L2:  ON Opt(4)  GOTO  Lb1, Lb2, Lb3, Lb4
6634  Lb1:  MOVE  Xstart-Xspan*.15, Ystart+Yspan*.25
6636  LABEL  USING "K"; "Log(k) cc/mole-s"
6638  GOTO  Lout
6640  Lb2:  MOVE  Xstart-Xspan*.15, Ystart+Yspan*.16
6642  LABEL  USING "K"; "Log(k) cc/molecule-s"
6644  GOTO  Lout
6646  Lb3:  MOVE  Xstart-Xspan*.15, Ystart+Yspan*.16
6648  LABEL  USING "K"; "Log(k) liters/mole-s"
6650  GOTO  Lout
6652  Lb4:  MOVE  Xstart-Xspan*.15, Ystart+Yspan*.08
6654  LABEL  USING "K"; "Log(k) liters/molecule-s"
6656  Lout:  LDIR  0
6658  MOVE  Xstart+.43*Xspan, Ystart-Yspan*.12
6660  LABEL  USING "K"; "1000/T"
6662  MOVE  T(1), Lk(1)
6664  FOR  N=1 TO  Nm
6666  PLOT  1000/T(N), Lk(N)
6668  NEXT  N
6669  LINE  TYPE  3
6670  MOVE  1000/T(1), Lk(1)-E(1)
6672  FOR  N=1 TO  Nm
6674  PLOT  1000/T(N), Lk(N)-E(N)
6676  NEXT  N
6678  RAD
6680  SUBEND
6800  SUB _Fit(Opt2,N,T(*),Xob(*),E(*),P(*),E2,Opt7)
6801  OPTION BASE 1
6810  DIM A(N,3),At(3,N),Ata(3,3),Atai(3,3),Atx(3),X(N)
6820  FOR I=1 TO N
6830    A(I,1)=1
6831    A(I,2)=0
6840  IF Opt2=2 THEN L10
6850  IF Opt7=3 THEN A(I,2)=LOG(T(I))
6870  A(I,3)=-1/(1.987*T(I))
6880  GOTÔ L11
6890  L10: IF Opt7=3 THEN A(I,2)=LG(T(I))
6900  A(I,3)=-.43429/(1.987*T(I))
6910  L11: NEXT I
6920  MAT At=TRN(A)
6930  MAT Ata=At*A
6940  MAT Atai=INV(Ata)
6950  MAT Atx=At*Xob
6960  MAT P=Atai*Xtx
6970  MAT X=A^P
6980  MAT E=Xob-X
6990  E2=DOT(E,E)
7000  END