Rotational Isomeric State Treatment of the Cystine Residue. Configuration Partition Function and Its Relationship to the Optical Activity Exhibited by the Disulfide Bond

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Abstract: The configuration partition function for two polypeptide chains cross-linked via a single cystinyl residue has been formulated. This objective is achieved using rotational isomeric state theory, in the form appropriate for branched molecules, to combine the configuration partition function for the cystinyl residue with appropriate representations for the configuration partition function of uncross-linked polypeptide chains. Rotational states, as well as preliminary estimates of the corresponding conformational energies, are obtained from consideration of semiempirical energy computations. The sequences -Cys-X- and -Cys-Pro- ($X \neq Pro$) lead to different results. The a priori probabilities computed for the two rotational states of the disulfide bond are found to be in excellent agreement with the sign, magnitude, and temperature dependence of the optical activity exhibited by the lowest energy electronic transition in simple derivatives of L-cystine.

Rotational isomeric state theory¹⁻⁴ has been successfully applied to homopolypeptides,⁵⁻¹⁵ oligopeptides,¹⁶ and both random¹⁷⁻¹⁹ and sequential²⁰ copolypeptides. It has also been applied to denatured proteins in which the disulfide bonds have been reduced so that covalent cross-links are absent.^{21,22} Recent extensions^{23,24} of rotational isomeric state theory to encompass the treatment of branched macromolecules now permit a study of disordered proteins containing an intact interchain disulfide bond. The first requirement is the formulation of the appropriate configuration partition function for the two cysteine residues which form the disulfide bond. The configuration partition function is presented in this paper. Subsequent communications will present the unperturbed dimensions computed for denatured proteins containing an interchain disulfide bond.

Geometry and Potential Functions. Figure 1 shows a representation of two polypeptide chains cross-linked via the disulfide bond formed by two cysteine residues. The C-S and S-S bond lengths are assigned the values 1.87 and 2.04 Å, respectively.²⁵ The C-C-S bond angle was 113°,²⁶ and the C-S-S bond angle was 107°.²⁷ All other bond lengths and bond angles were identical with those used by Brant et al.⁹ in their treatment of the L-alanyl residue. Virtual bonds⁵ are shown between successive α -carbon atoms. Each C^{α} , C', C^{β} , N, and S atom, as well as the intervening bonds and virtual bonds, is assigned a pre- and postsubscript according to the scheme described in ref 23. The presubscript denotes the branch to which that atom, bond, or virtual bond is assigned. The atoms in a given branch are numbered sequentially, as denoted by the postsubscript. The bond represented by bond vector $_{3}\mathbf{1}_{i}$ extends from atom *i*-1 to atom *i* in branch 3, while the virtual bond represented by $_{i}$ 1 $_{i}$ (j = 1, 2, 4, 5) extends from α -carbon atom 3*i*-3 to α -carbon 3*i* in branch *j*. The statistical weight matrix for bond (or virtual bond) i in branch j is denoted by $_{i}U_{i}$, and n_{i} is the number of virtual bonds in branch j.

Nonbonded energies were computed using the 6-12 functions described by Brant et al.⁹ The radius, polarizability, and effective number of electrons for the sulfur atom were taken from Scheraga.²⁶ The electrostatic interaction of the peptide groups was calculated in the monopole approximation as described by Brant et al.⁹ Torsional potentials for rotation about the N-C^{α} and C^{α}-C^{\prime} bonds were also taken from Brant et al.⁹ while the torsional potential for the C^{α}-C^{β} bond was that of Abe et al.²⁸ The torsional potential for the C^{β}-S bond was threefold, with minima at 0 and ±120° and a barrier height of 1.5 kcal/mol.²⁹ (We adopt here the convention that a trans placement corresponds to a dihedral angle of 0°.) For the S-S bond, the torsional potential is that used by Allinger et al.³⁰ Peptide bonds are maintained in the planar trans conformation.

Statistical Weight Matrices. The independence of the rotations about the N-C^{α} and C^{α}-C' bonds at adjacent α -carbon atoms⁵ permits immediate assignment of statistical weight matrices for virtual bonds which do not involve the α -carbon atoms in the cysteine residues. A 1 × 1 matrix whose sole element is unity is denoted by eq 1.

$$\mathbf{U}_{i} = {}_{2}\mathbf{U}_{j} = {}_{4}\mathbf{U}_{j} = {}_{5}\mathbf{U}_{j} = [1] \qquad i < n_{1}, j > 1 \qquad (1)$$

Atoms involved in the interactions pertinent to the assembly of ${}_1U_{n_1}$ are shown in Figure 2. The statistical weights for the rotational states about the ${}_{1}C^{\alpha}{}_{3n_{1}-3}C^{\beta}{}_{1}$ bond will depend on the values of φ and ψ . Consequently it is necessary for ${}_1U_{n_1}$ to be a row with the number of elements identical with the number of states for φ and ψ . Evaluation of the number of states, and the statistical weights, is based on the conformational energies for the fragment in Figure 2, evaluated at 10° intervals for φ and ψ . The result is shown in Figure 3. The convention adopted is that $\varphi, \psi = 0^{\circ}, 0^{\circ}$ for the fully extended chain. Figure 3 is, of course, identical with the conformational energy map obtained for the L-alanyl residue by Brant et al.9 The minimum energy is obtained when φ, ψ are 100°, 330°.⁹ This minimum and six local minima are denoted by the \times 's in Figure 3. They are listed in Table I in order of increasing energy.

A preliminary evaluation was made of the feasibility of using seven states for rotation about the last virtual bond in branch 1. Statistical weights for the states were assigned as shown in eq 2.

$$\frac{\sum_{\varphi' \psi'} \exp[-E(\varphi', \psi')/RT]}{\sum_{\varphi' \psi} \exp[-E(\varphi, \psi)/RT]}$$
(2)

The summations are taken at 10° intervals for φ and ψ , ignoring locations where the energy exceeds the minimum by more than 5 kcal/mol. The range for the double sum in the numerator is over that section of the conformational energy map assigned to the state in question, while the range for the double sum in the denominator extends over the entire conformational energy map. The boundaries for the states are provided by the 5 kcal/mol contour line and the dashed lines in Figure 3. The dashed lines approximate the location of the



Figure 1. Representation of two polypeptide chains cross-linked by the disulfide bond formed by two cysteine residues.



Figure 2. Atoms considered in the formulations of ${}_{1}\mathbf{U}_{n_{1}}$.

energy barriers between the minima. Average values of φ and ψ were determined for each state, using eq 3 and 4. The summations extend

$$\langle \varphi \rangle = \frac{\sum_{\varphi' \psi'} \varphi' \exp[-E(\varphi', \psi')/RT]}{\sum_{\varphi' \psi'} \exp[-E(\varphi', \psi')/RT]}$$
(3)

$$\langle \psi \rangle = \frac{\sum_{\varphi' \ \psi'} \varphi' \exp[-E(\varphi', \psi')/RT]}{\sum_{\varphi' \ \psi'} \exp[-E(\varphi', \psi')/RT]}$$
(4)

over the region of the conformational energy map assigned to the state in question. Results are summarized in Table I for the case where RT = 0.6 kcal/mol.

The final two states listed in Table I were eliminated from further consideration because of their small combined weight. An additional simplification is provided by combination of the fourth and fifth entries to yield a state with $\langle \varphi \rangle$, $\langle \psi \rangle = 86^{\circ}$, 126° and a weight of 0.043. This procedure yields the four states denoted by the squares in Figure 3. The statistical weight matrix ${}_{1}U_{n_{1}}$ is shown in eq 5.



Figure 3. Conformational energy map obtained for the atoms in Figure 2. Contours are drawn at 1, 2, and 5 kcal/mol above the energy minimum, which occurs at φ , $\psi = 100^{\circ}$, 330°. Minima are denoted by X's. The squares represent the rotational states used (see text).

 Table I. Minima in the L-Alanyl Residue Conformational Energy Map

Energy, kcal/ mol	arphi,deg	ψ, deg	$\langle \varphi angle,$ deg	$\langle \psi \rangle$. deg	Statistical wt
0.00	100	330	94	328	0.415
0.21	100	280	92	275	0.173
0.27	30	330	26	312	0.363
0.91	110	130	107	128	0.033
1.33	10	120	15	119	0.010
2.34	230	240	236	248	0.004
2.38	250	350	247	348	0.002

Table II. States Used for φ , ψ

Designation	$\langle \varphi \rangle$, deg	$\langle \psi \rangle$. deg	Statistical wt	1 <i>P</i> ₁₁ <i>n</i> ₁
β_1	94	328	w1 (0.418)	0.426
β_2	26	312	$w_2(0.365)$	0.380
β_3	92	275	$w_3(0.174)$	0.154
α_1	86	126	w4 (0.043)	0.040

$$\beta_1 \quad \beta_2 \quad \beta_3 \quad \alpha_1 \\ \mathbf{U}_{n_1} = [w_1 \quad w_2 \quad w_3 \quad w_4] \tag{5}$$

Table II provides a summary in which the w_i have been adjusted so that their sum is unity.

The dimensions of ${}_{1}U_{n_{1}}$ and ${}_{2}U_{2}$ require that ${}_{2}U_{1}$ be a column consisting of four elements. Independence of events at neighboring α -carbon atoms requires that these four elements be identical.

$$_{2}\mathbf{U}_{1} = \operatorname{col}(1, 1, 1, 1)$$
 (6)

The dihedral angle, χ_1 , about the ${}_1C^{\alpha}{}_{3n_1}-C^{\beta}{}_1$ bond specifies the position of ${}_3S_2$. This angle will be considered to be 0° for a trans placement of ${}_1N_{3n_1-1-1}C^{\alpha}{}_{3n_1-3}C^{\beta}{}_{1-3}S_2$. Interactions contributing to the statistical weights represented in ${}_3U_1$ are the torsional potential about the ${}_1C^{\alpha}{}_{3n_1-3}C^{\beta}{}_1$ bond and the



Figure 4. Atoms considered in the formulation of ${}_{3}U_{1}$.



Figure 5. Conformational energy as a function of χ_1 for the four indicated combinations of φ, ψ .

nonbonded interactions between ${}_{3}S_{2}$ and all other atoms in Figure 4 except ${}_{1}C^{\alpha}{}_{3n_{1}}$ and ${}_{3}C^{\beta}{}_{1}$. The resulting energies are shown in Figure 5 for the four states assigned by φ, ψ . This energy is essentially identical for all four combinations of φ , ψ when $\chi_1 = 0^\circ$. When $\chi_1 = 120^\circ$ and φ is 26°, the ${}_{3}S_2$ atom is 2.96 Å from the oxygen atom bonded to ${}_{1}C'_{3n_{1}-2}$. However, the energy of this interaction is less than 0.1 kcal/mol and can be alleviated by a slight increase in φ , so it will be ignored. The $_{3}S_{2}$ atom interacts with the oxygen atom bonded to $_{2}C'_{1}$ when $\chi_1 = 240^\circ$ and $\psi = 275^\circ$. The estimated energy is about 0.6 kcal/mol, and the statistical weight is s_1 . This interaction is relieved at larger values of ψ and is practically nonexistent when ψ is in the β_1 or β_2 states. A repulsive interaction occurs between ${}_{2}S_{2}$ and ${}_{2}N_{2}$, plus its attached hydrogen atom, when $\chi_1 = 240^\circ$ and $\psi = 126^\circ$. The S-H distance is only 2.23 Å. The energy of this interaction is about 2 kcal/mol, and the statistical weight is s_2 . Weak three-bond interactions, represented by s_3 , occur between ${}_3S_2$ and both ${}_1N_{3n_1-1}$ and ${}_2C'_1$ when χ_1 = 240° . The energy is 0.2 kcal/mol. The resulting statistical weight matrix is shown in eq 7. The dimensions of this matrix

are 4 × 12, rather than 4 × 3, because we wish to be able to recognize the dependence of rotation about the ${}_{3}C^{\beta}{}_{1-3}S_{2}$ bond on the state adopted by φ, ψ . The rectangular matrix denoted



Figure 6. Atoms considered in the formulation of ${}_{3}\mathbf{U}_{2}$.



Figure 7. Conformational energy as a function of χ_2 for the three values of χ_1 and four values of φ, ψ . The φ, ψ are given by the same type as lines used in Figure 5.

 ${}_{2}\mathbf{U}_{1} \ominus {}_{3}\mathbf{U}_{1}$, and defined in ref 23, is identical with ${}_{3}\mathbf{U}_{1}$.

Specification of the dihedral angle, χ_2 , about the ${}_3C^{\beta}{}_{1-3}S_2$ bond determines the position of ${}_3S_3$. Interactions contributing to the statistical weights embodied in ${}_3U_2$ are the torsional potential about the ${}_3C^{\beta}{}_{1-3}S_2$ bond and the nonbonded interaction between ${}_3S_3$ and all other atoms in Figure 6 except ${}_3C^{\beta}{}_1$ and ${}_3S_2$. The resulting energies are shown in Figure 7 for the four states assigned to φ , ψ and the three states for χ_1 . The energy in the vicinity of $\chi_2 = 0^\circ$ is seen to be essentially independent of χ_1 and the state assigned by φ , ψ .

The three-bond interaction between ${}_{3}S_{3}$ and ${}_{1}C^{\alpha}{}_{3n_{1}}$ is slightly attractive (0 to -0.1 kcal/mol) when χ_{2} is in either of the gauche states. This interaction is so small that it will be ignored. Repulsive interactions involving ${}_{3}S_{3}$ and atoms in the amide units may occur when $\chi_{2} = \pm 120^{\circ}$. These repulsions

cause the minima for the gauche states to frequently be displaced from $\pm 120^{\circ}$.

Interaction occurs between ${}_{3}S_{3}$ and ${}_{2}C'_{1}$ (separation 3.04 Å) when χ_{1} , $\chi_{2} = 0^{\circ}$, 110°. An additional interaction occurs between ${}_{3}S_{3}$ and the oxygen atom attached to ${}_{2}C'_{1}$ (separation 2.64 Å) when $\psi = 275^{\circ}$. The ${}_{3}S_{3}$ and ${}_{2}N_{2}$ atoms are separated by only 2.75 Å when $\psi = 328^{\circ}$. These latter two interactions can be relieved somewhat by an appropriate adjustment of ψ . The relevant statistical weights are denoted by ${}_{s45}$, ${}_{s4}$, ${}_{s4s6}$, and ${}_{s4}$ for the cases where φ , ψ are in the β_{1} , β_{2} , β_{3} , and α_{1} states, respectively. The energies corresponding to ${}_{s4-s6}$ are estimated at 0.2, 0.7, and 1.0 kcal/mol, respectively.

No repulsive interactions occur when $\chi_1, \chi_2 = 0^\circ, 250^\circ$. The energy in this conformation may actually be negative by as much as 0.2 kcal/mol due to favorable nonbonded interactions. The statistical weight is denoted by s_7 . No significant interactions occur when $\chi_1, \chi_2 = 120^\circ$, 110° unless φ, ψ are in the β_2 state. A slightly higher energy (up to 0.2 kcal/mol) may be appropriate for $\beta_2 g^+ g^+$ due to the interaction between $_3S_3$ and the oxygen atom attached to $_1C'_{3n_1-2}$. The statistical weight is s_8 .

A weak repulsive interaction occurs between ${}_{3}S_{3}$ and the hydrogen atom attached to ${}_{1}N_{3n_{1}-1}$ when $\chi_{1} = 120^{\circ}$, $\chi_{2} = 250^{\circ}$, and $\varphi = 90 \pm 4^{\circ}$. This interaction is alleviated by a slight decrease in φ . A stronger interaction occurs between ${}_{3}S_{3}$ and ${}_{1}C'_{3n_{1}-2}$ (separation 2.70 Å) when $\chi_{1} = 120^{\circ}$, $\chi_{2} = 250^{\circ}$, and $\varphi = 26^{\circ}$. The statistical weights are denoted by s_{9} and s_{10} and the corresponding energies are estimated to be 0.2 and 1 kcal/mol, respectively.

Severe interaction occurs between ${}_{3}S_{3}$ and the hydrogen atom attached to ${}_{1}N_{3n_{1}-1}$ (separation 2.13-2.14 Å) when $\varphi =$ 90 ± 4°, $\chi_{1} = 240^{\circ}$, and $\chi_{2} = 110^{\circ}$. This interaction is reflected in s_{11} , with an energy of 2-3 kcal/mol.

Repulsive interactions exist whenever χ_1 , χ_2 are in the g⁻g⁻ state. Interaction occurs between ${}_3S_3$ and the oxygen atom attached to ${}_2C'_1$ when $\psi = 300 \pm 28^\circ$, with the interaction becoming of increasing severity as ψ decreases through this range. The statistical weights are denoted by s_{12} - s_{14} with estimated energies of 0.5, 1-2, and >5 kcal/mol. The interactions are between ${}_3S_3$ and both ${}_2N_2$ and its hydrogen atom when ψ is 126°. The estimated energy is >3 kcal/mol. The statistical weight matrix ${}_3U_2$ is shown in eq 8.



Specification of the dihedral angle, χ_3 , about the ${}_3S_{2-3}S_3$ bond determines the position of ${}_3C^{\beta}_4$. Interactions contributing to the statistical weights embodied in ${}_3U_3$ are the torsional potential about the ${}_3S_{2-3}S_3$ bond and the nonbonded interactions between ${}_3C^{\beta}_4$ and ${}_3C^{\beta}_1$ and between ${}_3C^{\beta}_4$ and ${}_1C^{\alpha}_{3n_1}$. The resulting energies are shown in Figure 8 for the three states of



Figure 8. Conformational energies as a function of χ_3 for the indicated values of χ_2 .

Table III. Energies Associated with the s_i

Statistical wt	Estd energy. kcal/mol	Statistical wt	Estd energy, kcal/mol
S I	0.6	59	0.2
\$2	2	\$10	1
s 3	0.2	511	2
54	0.2	\$12	0.5
\$5	0.7	S13	1.5
56	1	514	5
\$7	-0.2	\$15	3
58	0.2	s ₁₆	-0.1

the ${}_{3}C^{\beta}{}_{1-3}S_{2}$ bond. Essentially the same result is found when the gauche states for χ_{2} are located at ±110° or ±120°. Minima occur near ±100° for χ_{3} in Figure 8. The lowest energies are obtained when χ_{2} and χ_{3} adopt gauche placements of opposite signs, due to favorable nonbonded interactions generated between the ${}_{3}C^{\beta}_{4}$ and ${}_{1}C^{\alpha}{}_{3n_{1}}$ atoms in this conformation. The energy associated with s_{16} in ${}_{3}U_{3}$ is about -0.1 kcal/mol. The energy in the $g^{+}g^{+}$ and $g^{-}g^{-}$ states is calculated to be only about 0.02 kcal/mol lower than that in the tg^{\pm} states, and consequently it will be assigned a statistical weight of unity.

$${}_{3}U_{3} = {\begin{array}{c} t \\ g^{+} \\ g^{-} \\ g^{-} \\ s_{16} \\ s_{16}$$

The interactions embodied in ${}_{3}U_{4}$ and ${}_{3}U_{5}$ have already been discussed in connection with preceding statistical weight matrices. The resulting matrices are shown in eq 10 and 11.

$${}_{3}\mathbf{U}_{4} = \frac{g^{+}}{g^{-}} \begin{bmatrix} t & g^{+} & g^{-} \\ 1 & 1 & s_{16} \\ 1 & s_{16} & 1 \end{bmatrix}$$
(10)



Mattice / Cystine Residue

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Figure 9. A priori probabilities for the bonds in branch 3 when RT = 0.6 kcal/mol.

A summary of the preliminary estimates of the energies associated with the various s_i is presented in Table III. The dimensions of ${}_{3}U_{5}$, ${}_{4}U_{2}$, and ${}_{5}U_{2}$ require that ${}_{4}U_{1}$ and ${}_{5}U_{1}$ each be a column consisting of 12 elements. Independence of events at adjacent α -carbon atoms⁵ requires that all of the elements be identical.

$${}_{4}\mathbf{U}_{1} = {}_{5}\mathbf{U}_{1} = {}_{4}\mathbf{U}_{1} \Theta {}_{5}\mathbf{U}_{1}$$

= col (1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1) (12)

Configuration Partition Function and average Bond Conformations. The configuration partition function, Z, for the molecule in Figure 1 is obtained from the statistical weight matrices via eq 13.

$$Z = {}_{1}U_{1}^{(n_{1})}({}_{2}U_{1} \ominus {}_{3}U_{1})[({}_{2}U_{2}^{(n_{2}-1)}) \\ \otimes [{}_{3}U_{2}^{(4)}({}_{4}U_{1} \ominus {}_{5}U_{1})]({}_{4}U_{2}^{(n_{4}-1)}) \otimes ({}_{5}U_{2}^{(n_{5}-1)})]]]$$
(13)

Symbolism of the type ${}_{1}U_{1}^{(n_{1})}$ denotes the product of n_{1} successive matrices, commencing with ${}_{1}U_{1}$, and \otimes denotes the direct product.⁴ The a priori probability that bond *i* in branch *j* is in state η , denoted ${}_{j}p_{\eta,i}$, and the a priori probability that this bond is in state η and the preceding bond is in state ξ , denoted ${}_{j}p_{\xi\eta,i}$, were calculated as Z'/Z.²³ The procedure for calculating Z' is identical with that used for Z except that appropriate elements of one of the statistical weight matrices are replaced by zeros.^{3,23} Table II shows the ${}_{1}p_{\eta,n_{1}}$ calculated using the energies in Table III, the w_{i} in Table II, and RT = 0.6 kcal/mol. They differ only slightly from the w_{i} . This result indicates that atoms beyond ${}_{3}C^{\beta}_{1}$ have only a minor effect upon the conformational preferences for φ , ψ , a result in harmony with the experimental unperturbed dimensions of several homopolypeptides containing $-CH_{2}R$ side chains.^{5-7,12}

A priori probabilities for the bonds in branch 3 are shown in Figure 9. Symmetry dictates that $_{3}p_{\eta;i} = _{3}p_{\eta;6-i}$. All bonds except the disulfide bond favor the trans conformation. The a priori probability that the disulfide bond will be in the g⁺ state is slightly greater than the a priori probability that it will be in the g⁻ state.

It is of interest to compare the results obtained from the present study with those predicted^{31,32} from theoretical studies of the electronic transitions in the disulfide group. The wavelength of the lowest energy electronic transition is predicted, and observed, to be extremely sensitive to the dihedral angle about the disulfide bond.³¹ If χ_3 is ±100°, as suggested by Figure 8, this band is predicted to be located at 260-265 nm.³¹



Figure 10. Conformational energy map for L-cysteine followed by L-proline. Contours are drawn at 1, 2, and 5 kcal/mol above the energy minimum, which occurs at $\varphi, \psi = 100^{\circ}, 320^{\circ}$. Minima are denoted by X's. The squares represent the rotational states used (see text).

Coleman and Blout³³ find that the lowest energy disulfide transition is at 262 ± 3 nm for N,N'-diacetyl-L-cystine bis(methylamide) in aqueous solution.

The lowest energy electronic transition is predicted^{31,32} to have negative rotatory strength when $\chi_3 = 100^\circ$ and positive rotatory strength when $\chi_3 = -100^\circ$. Experimentally, this band exhibits negative circular dichroism for solutions of L-cystine,³³ N.N'-diacetyl-L-cystine,³³ N.N'-diacetyl-L-cystine bis-(methylamide),³³ oxidized glutathione,^{33,34} S-ethylthio-Lcysteine,³⁴ S-ethylthio-L-cysteine,³⁴ S-propylthio-L-cysteine,³⁴ and S-methylthioglutathione.³⁴ These results, in conjunction with Figure 8, require that $_{3}p_{g^+;3} > _{3}p_{g^-;3}$. The a priori probabilities presented in Figure 9 are in agreement with this inequality.

The observed optical activity of the longest wavelength electronic transition in S-methylthio-L-cysteine, S-ethylthio-L-cysteine, and S-propylthio-L-cysteine decreases substantially with increasing temperature.³⁴ Circular dichroism spectra show that the maximum ellipticity at RT = 0.7 kcal/ mol is only 67-71% of the value at RT = 0.6 kcal/mol. The configuration partition function for N-acetyl-S-methylthio-L-cysteine methylamide can be written as

$$Z = {}_{1}U_{n_{1}}({}_{2}U_{1} \ominus {}_{3}U_{1})({}_{3}U_{2})({}_{3}U_{3})\begin{bmatrix}1\\1\end{bmatrix}$$
(14)

The $_{3}p_{g^{\pm};3}$ were calculated using the w_i and s_i for RT = 0.6 and 0.7 kcal/mol. This procedure indicates that $_{3}p_{g^{\pm};3} - _{3}p_{g^{-};3}$ at RT = 0.7 kcal/mol would be 68% of the value at RT = 0.6 kcal/mol, a result in excellent agreement with the observed temperature dependence of the circular dichroism.

An estimate of $_{3}p_{g^+;3}$ can be obtained from eq 15,

$$R_{\text{obsd}} = R[_{3}p_{g^{+};3} - (1 - _{3}p_{g^{+};3})]$$
(15)

where R_{obsd} is the observed rotatory strength and R is the rotatory strength predicted for $\chi_3 = 100^{\circ}$. Coleman and Blout³³ find that R_{obsd} is $-1.641 \pm 0.235 \times 10^{-40}$ erg cm³ for the 262.3-nm band of N.N'-diacetyl-L-cystine bis(methylamide). Webb et al.³² calculate that the net rotatory strength for the two closely situated, lowest energy electronic transitions should be -121×10^{-40} and -193×10^{-40} erg cm³ when χ_3 is 90 and 105°, respectively. Linear interpolation provides an estimate of -169×10^{-40} erg cm³ for R when χ_3 is 100°. Use of these values for R_{obsd} and R in eq 15 provides a result of 0.50485 ±

Designation	Energy, kcal/mol	arphi,deg	ψ, deg.	$\langle \varphi \rangle$. deg	$\langle \psi \rangle$, deg	Statistical wt
β_1	0.00	100	320	97	311	$0.49(w_1)$
β_2	0.14	20	310-320	30	310	$0.51 (w_2)$
	2.66	240	260	241	276	



Figure 11. Conformational energy as a function of χ_1 for L-cystine followed by L-proline.

0.0007 for $_{3}p_{g^{+};3}$. The value calculated from the rotational isomeric state treatment (0.5068) is in surprisingly good agreement with the estimate from eq 15.

The effect on $_{3}p_{g^{+};3}$ of doubling and halving the energies associated with the s_i was explored. A larger value of $_{3}p_{g^{+};3}$ is obtained with an increase in the energy associated with s_3 , s_4 . s_5 , s_6 , s_8 , or s_{11} or with a decrease in the energy associated with s_7 , s_9 , s_{10} , s_{12} , s_{13} , or s_{16} . Little effect is obtained upon variation of the energy associated with s_1 , s_2 , s_{14} , or s_{15} . A larger $_{3}p_{g^{+};3}$ is also obtained by increasing w_1 or w_3 or by decreasing w_2 . The result is insensitive to w_4 . These variations in the w_i and the energies associated with the s_i do not significantly improve the ability to reproduce the observed spectral characteristics of small molecules related to L-cystine.

Treatment when Proline Follows Cysteine. The conformational energy map for an L-alanyl residue is altered when it is followed by an L-prolyl residue.¹⁸ For this reason it can be anticipated that a different result will be obtained for the cystine configuration partition function when the amino acid sequence is -Cys-Pro-. The major cause of the change is the steric interactions generated by the δ (CH₂) of the prolyl residue. In order to estimate the changes required, the hydrogen atom bonded to $_2N_2$ was replaced by a methylene unit, using a C^{δ} -N bond length of 1.47 Å and a C'-N-C^{δ} angle of 126°.¹³

Reevaluation of the energies appropriate for consideration in ${}_{1}U_{n_{1}}$ yields Figure 10. This figure bears a close resemblance to the conformational energy map computed by Schimmel and Flory¹⁸ for an L-alanyl residue followed by an L-prolyl residue. The minimum energy in Figure 10 is at φ , $\psi = 100^{\circ}$, 320°, which is close to the location (100°, 300°) found for Ala-Pro.¹⁸ This minimum, and two local minima within 3 kcal/mol of the minimum, is denoted by \times in Figure 10 and summarized in Table IV. The statistical weights, $\langle \varphi \rangle$, and $\langle \psi \rangle$ were obtained by the same procedure applied for ${}_{1}U_{n_{1}}$ when the residue following cysteine is not L-prolyl. The β_{1} and β_{2} designations bear an obvious correspondence to two of the entries in Table II. No state is assigned to the region at $\varphi = 240^{\circ}$ because of its low



Figure 12. Conformational energy as a function of χ_2 for the three values of χ_1 and two values of φ , ψ used in Figure 11.



Figure 13. A priori probabilities for the bonds in branch 3 when L-proline follows L-cysteine in both polypeptide chains. RT = 0.6 kcal/mol.

statistical weight (0.002). The statistical weight matrix $_1U_{n_1}$ is therefore given by eq 5 with $w_1 = 0.49$, $w_2 = 0.51$, $w_3 = w_4 = 0$.

The energies appropriate for consideration in ${}_{3}U_{1}$ are shown

Table V. Changes Required for Cys(Pro)

$w_1 = 0.49$	
$w_2 = 0.51$	
$w_3 = w_4 = 0$	
$E_{s_5} = E_{s_7} = E_{s_8} = 0.0 \text{ kcal/mol}$	
$E_{s_{12}} = 1.5 \text{ kcal/mol}$	
$\langle \varphi \rangle, \langle \psi \rangle = 97^{\circ}, 311^{\circ}$ for β_1	
$\langle \varphi \rangle$, $\langle \psi \rangle = 30^{\circ}$, 310° for β_2	

in Figure 11. There is a close correspondence with the analogous two curves in Figure 5. Hence eq 7 can still be used for $3U_{1}$.

Figure 12 presents the energies which are reflected in ${}_{3}U_{2}$. Comparison with Figure 7 reveals a general similarity, but there are differences in detail. The interaction between ${}_{3}S_{3}$ and $_2N_2$ when $\chi_1, \chi_2 = 0^\circ$, 110° is relieved upon a reduction of ψ from 328 to 311°. For this reason the energy associated with s_5 is estimated to be 0.0 kcal/mol in Cys(Pro). The energies at $\chi_1, \chi_2 = 0^\circ$, 250° and for $\beta_2 g^+ g^+$ are sufficiently close to that obtained when $\chi_2 = 0^\circ$ to permit assignment of zero energy for the statistical weights denoted by s_7 and s_8 . Finally, the energy associated with the $\beta_1 g^- g^-$ state must be changed due to the increased interaction between ${}_{3}S_{3}$ and the oxygen atom bonded to ${}_{2}C'_{1}$ when ψ decreases from 328 to 311°. The energy associated with s_{12} is increased from 0.5 to 1.5 kcal/ mol. A summary of the changes required is presented in Table V. The energies associated with s_1 , s_2 . s_6 . s_{14} , and s_{15} are irrelevant since terms containing these statistical weights will be rejected. A priori probabilities for the bonds in branch 3 are shown in Figure 13. The value of $_{3}p_{g^+;3}$ is 0.4920. It can be made to exceed 0.5 by minor (0.1 kcal/mol) adjustment of the energies associated with the s_i . The adjustments are in the direction which serves to increase $_{3}p_{g+3}$ for the case where the amino acid residue following cysteine is not proline.

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