Free-energy Barrier of Conformational Inversion in Bilirubin

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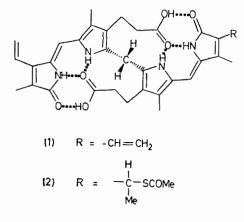
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Summary Bilirubin exists in chloroform solution as two enantiomeric conformations (1 and its mirror image) separated by a free-energy barrier of 17.9 ± 0.5 kcal mol⁻¹.

It has long been recognised that bilirubin is stabilised by intramolecular hydrogen bonds.¹ To account for certain spectral data and chemical anomalies Knell *et al.*² have recently proposed the dissymmetric conformation (1) as a secondary structure of the pigment in non-polar solvents and in the solid state. The n.m.r. study reported here strongly supports such an assumption, showing that two interconvertible enantiomeric forms of bilirubin (probably 1 and its mirror image) must occur in chloroform solution; it also allows the free-energy barrier of conformational inversion to be calculated.

The ¹H n.m.r. spectrum (100 MHz; hexamethyldisiloxane reference) of (2)³ in anhydrous CDCl₃ at room temperature showed the methyl signals of the Me-COS- and Me-CH-(Ar)-S- groups as a doublet (centred at δ 2·18, $\Delta \nu$ 3·1 Hz)

and a doublet of doublets (centred at $\delta 1.50$, $\Delta v 3.4$ Hz, J 7.0 Hz), respectively. As the temperature was raised to 53 ± 3 °C, the doublet at 2.18 merged to a broad singlet and



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the system of peaks at δ 1.50 became a doublet (still with J 7.0 Hz, † thus indicating the occurrence of two exchanging populations of conformers; further, the broadened lineshapes observed over this temperature range showed an approximately equal distribution of the molecules between the two populations.

These data are best explained by assuming a restricted rotation of the dipyrromethene units of (2) with respect to each other; such a restriction causes four stereoisomeric forms of (2) (two pairs of enantiomers) to exist as a consequence of the presence of two chiral elements in the molecule: *i.e.* the 'helical' conformation $(R,S)^4$ of the tetrapyrrolic skeleton and the asymmetric carbon atom (R', S') in the side chain. The rate constant at the colescence temperature for the interconversion of two diastereomeric conformers of (2) was then calculated⁵ from the above spectral data [equations (1) and (2)].

$$(RR')[\text{or } (SS')] \xrightarrow[k_{-1}]{k_{-1}} (SR')[\text{or } (RS')]$$
(1)

$$k_1 = ca. k_{-1} = k = 7.2 \pm 0.4 \text{ s}^{-1} (T_o = 53 \pm 3 \text{ °C}) (2)$$

The free-energy of activation obtained from k using the Evring equation ($\Delta G^{\ddagger} = 17.9 \pm 0.5$ kcal mol⁻¹, the transmission coefficient being taken as unity)⁵ represents the barrier of conformational inversion in (2) and, by implication, in bilirubin itself (1). In fact, bilirubin in chloroform shows the same u.v. spectrum and i.r. absorption bands [in the range 3600-2900 (NH, OH) and 1700-1600 cm⁻¹ (amide and carboxy CO groups)] as compound (2).³

It must be pointed out that the above value of the freeenergy of activation is consistent with the cleavage of four strong and two weaker hydrogen bonds per molecule.^{2,6}

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† There should be two different coalescence temperatures, one for each methyl group; in practice they were not distinguishable.

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⁶ G. C. Pimentel and A. L. McClellan, 'The Hydrogen Bond,' Freeman, San Francisco, 1960, pp. 224, 348.