

Through-space Interaction and Chromophore Shape: the Chiral 13-*cis*-Retinal Schiff's Base Dimer

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The hydrogen-bridged dimer of a chiral 13-*cis*-retinal Schiff's base has been prepared and characterised by UV and circular dichroism (CD) spectroscopy; the CD spectrum is almost enantiomeric to that of the corresponding all-*trans*-retinal Schiff's base dimer pointing to the importance of chromophore shape to through-space interaction.

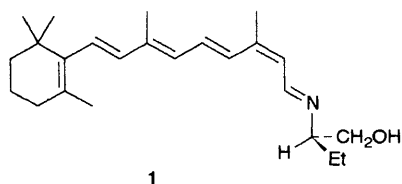
Of the different retinals known the all-*trans*-, 11-*cis*- and 13-*cis*-isomers merit, and have received, special attention because of their role as specialized molecular tools in basic biochemical processes: 11-*cis*- and all-*trans*-retinal as the photoreceptor and primary photoproduct, respectively, of the visual process in the vertebrate visual pigment rhodopsin; 13-*cis*- and all-*trans*-retinal both as components of the dark-adapted form of the proton-pumping pigment bacteriorhodopsin in the purple membrane of *Halobacterium halobium*. Molecular shapes and their changes during and after electronic excitation, probably play a most important role in bringing these processes about.¹

Information regarding the conformational flexibility, or rigidity, of these molecules comes from many different sources. The picture that emerges from these studies is generally consistent with the result of force-field or quantum-mechanical calculations, *viz.* that of a polyene chain preferring a planar arrangement of the conjugated double bonds,

significant deviations from this geometry being caused only by severe steric interactions. Thus, all retinals and retinal Schiff's bases are twisted about the C(6)–C(7) bond, and only the 11-*cis*-isomer has another highly twisted bond, C(12)–C(13), resulting in the existence of two conformational isomers, *viz.* the 12-*s-cis* and the 12-*s-trans* form.

We have studied the chiral dimers formed from chiral all-*trans*-² and 11-*cis*-retinal Schiff's bases³ using UV and CD spectroscopy and found the results consistent with a structure in which the two retinal moieties are projecting out from a rather rigid ten-membered ring system held together by two hydrogen bonds. We now report on the aggregation behaviour of the chiral 13-*cis*-retinal Schiff's base **1**.

We also present the results of a theoretical analysis which confirms the experimental finding that the chromophores in the dimer formed from **1**, due to the *cis*-configured double bond, have a close to mirror-image like relationship to the one formed from the homochiral all-*trans* retinal Schiff's base.



tion⁶ were joined *via* two hydrogen bonds to form a ten-membered ring in a boat-like arrangement found earlier⁷ to represent the minimum energy conformation. The dimer was then allowed to relax in the MMP2 force field. The geometry obtained is shown in Fig. 2, together with the structures of the all-*trans*- and 11-*cis*-retinal Schiff's base dimers calculated in a completely analogous manner. The only symmetry element of these aggregates is a C_2 -axis making them dissymmetric and non-superimposable on their mirror-images. The asymmetric centre which is the carbon next to the azomethine nitrogen is identical in all three aggregates; still the over-all shapes differ widely due to the different shapes of the retinylidene chromophores. Indeed, the two dimers derived from the all-*trans*- and the 13-*cis*-retinal isomers have an almost mirror image-like relationship with respect to the chromophores (see Fig. 2). The inverse relationship of the CD spectra of these two dimers, with rotational strengths of opposite sign for the two exciton states should therefore really not come as a surprise. For additional support of the proposed structures we have calculated the spectroscopic data to be expected for the dimers by a perturbational treatment involving the energies of the seven lowest CI states obtained from CNDO/S and the Coulombic interaction between atomic charges obtained as a multipole expansion of the corresponding CI transition densities. Energies and oscillator and rotatory strengths were then obtained from the eigenstates of the diagonalized interaction matrix. The resulting CD spectra are compared in Fig. 3 with the experimental spectra. There is

very good agreement for the all-*trans*- and 13-*cis*-retinal Schiff's base dimers, except for a 10 to 15 nm blue shift, of the long wavelength couplets; for 11-*cis*-retinal, however, the results completely disagree, the most probable reason being the almost parallel orientation of the transition moments which leads to sign inversion upon minute geometry changes in this dimer[†] and the inability of the geometry optimization procedure to account for differences as subtle as these.

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References

- 1 See, e.g. in L. Stryer, *Biochemistry*, Freeman, New York, 1988.
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- 3 V. Haas, U. Wingen and V. Buss, *J. Chem. Soc., Chem. Commun.*, 1989, 1053.
- 4 V. Buss, V. Haas and U. Wingen, *Z. Naturforsch., B*, 1989, **44**, 333.
- 5 C. J. Simmons, R. H. Liu, M. Denny and K. Seff, *Acta Crystallogr., Sect. B*, 1981, **37**, 2197.
- 6 A *cis*- or *Z*-configured azomethine bond was postulated for 13-*cis*-retinal Schiff's base in bacteriorhodopsin on the basis of ¹³C NMR spectra (G. S. Harbison, S. O. Smith, J. A. Pardo, C. Winkel, J. Lugtenburg, J. Herzfeld, R. Mathies and R. Griffin, *Proc. Natl. Acad. Sci., USA*, 1984, **81**, 1706). This is, however, a steric effect allowing the chromophore to fit into the same protein pocket as all-*trans*-retinal.
- 7 M. Klein, U. Wingen and V. Buss, *J. Am. Chem. Soc.*, 1987, **109**, 6486.

† The geometry adopted for the 11-*cis*-retinal chromophore has both the C(6)–C(7) and C(12)–C(13) bonds twisted in the same direction (+57 and +158°, respectively) from the planar *s-cis* conformation. There is another feasible geometry, with the C(6)–C(7) bond twisted in the opposite direction, which, however, leads to a dimer energy which is 2.2 kcal mol⁻¹ higher than the one described. The CD spectrum of this second dimer is essentially unchanged with respect to the long wavelength couplet.