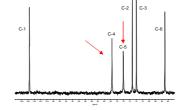
α-CYCLOALTRIN: MOLECULAR SHAPE AND HYDRATION PROPERTIES

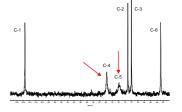
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 α -Cycloaltrin (1)^[1] adopts in the solid state^[2] a unique C_3 symmetrical conformation with nearly perfect 4C_1 and 1C_4 chairs in an alternating sequence (1a). 1H and 1S NMR data in D_2O at 30°C and 4°C (cf. Fig. 1) indicate a dynamic conformational equilibrium of at least two different altrose geometries with almost equal energies in the macrocycle.

Fig. 1: 13 C NMR (75 MHz) spectra of α -CA (1) in D₂O at 30°C (left) and 4°C (right) indicate a pronounced low-temperature broadening of the C-4 and C-5 signals, and hence, a dynamic equilibrium between at least two different altrose conformations.





Unlike the conformation 1a (Fig. 2) which features side-on indentations only, the intermediate all- $^{O}S_{2}$ structure 1b in the $^{4}C_{1} = ^{O}S_{2} = ^{1}C_{4}$ equilibrium features a "through-going" central cavity (Fig. 2). We have investigated the dynamic processes in 1 by means of molecular dynamics (MD) simulations in aqueous solution, focusing in particular on the hydration properties and water exchange phenomena of 1a and 1b, defining the first hydration shells of both geometries (Fig. 2).

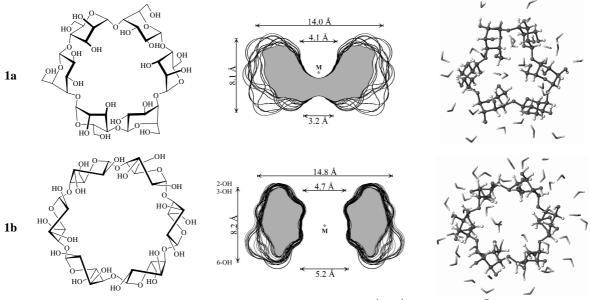


Fig. 2: Superimposed surface cross section cuts (center) of the alternating ${}^4C_1/{}^1C_4$ (1a) and all- 0S_2 conformation (1b) of α -cycloaltrin with approximate molecular dimensions in Å. On the right, typical MD derived snap-shot geometries of the first hydration shell of both geometries with 31 and 41 water molecules hydrogen bonded to 1a and 1b are displayed.

^[1] Y. Nogami, K. Fujita, K. Ohta, K. Nasu, H. Shimada, C. Shinohara, and T. Koga, J. Incl. Phenom. Mol. Recogn. Chem. 1996, 25, 57-60.

^[2] Y. Nogami, K. Nasu, T. Koga, K. Ohta, K. Fujita, S. Immel, H. J. Lindner, G. E. Schmitt, and F. W. Lichtenthaler, *Angew. Chem.* **1997**, *109*, 1987-1991; *Angew. Chem., Int. Ed. Engl.* **1997**, *35*, 1899-1902.