

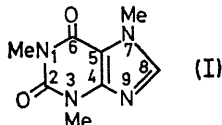
## Self-association of Caffeine in Aqueous Solution: $^1\text{H}$ Nuclear Magnetic Resonance Study

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**Summary** A possible mode of self-association of caffeine in aqueous solution is suggested on the basis of the concentration dependence of chemical shifts.

ALTHOUGH conclusive evidence has been presented for the self-association of caffeine<sup>1-3</sup> and of other xanthine derivatives<sup>1</sup> in aqueous solutions, the mechanism and mode of



association of these medically important molecules have not been elucidated. We report a study of the self-

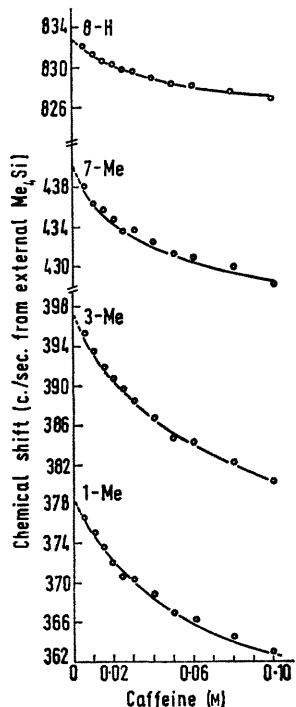


FIGURE 1. Concentration dependence of caffeine proton chemical shifts.

association of caffeine (I) by  $^1\text{H}$  n.m.r. spectroscopy, a tool which has been widely used for the study of the structurally analogous purine bases and nucleosides.<sup>4,5</sup>

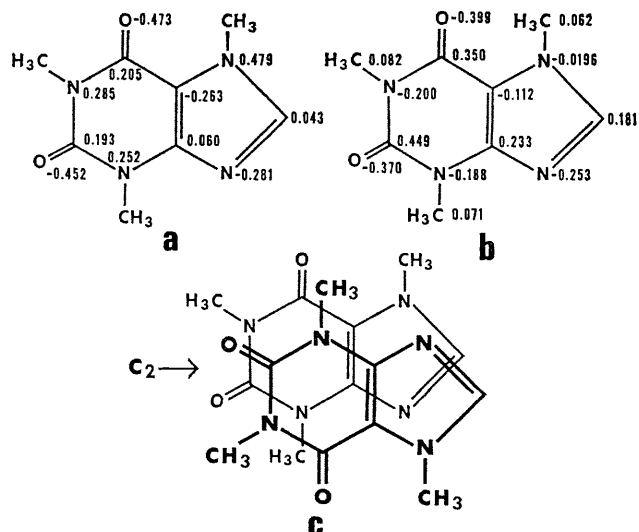


FIGURE 2. (a)  $\pi$ -Electron charge distribution and (b) total atom charge distribution calculated by the CNDO/2 method. (c) A possible mode of association for caffeine.

The high resolution 100 MHz  $^1\text{H}$  n.m.r. spectrum of a 0.01 M- $\text{D}_2\text{O}$  solution of caffeine has four signals of relative intensities 1:3:3:3 at *ca.* 831, 436, 393, and 375 c./sec. (from external  $\text{Me}_4\text{Si}$ ). These signals have been assigned<sup>6</sup> to the proton at C-8 and to the 7-, 3-, and 1-methyl protons, respectively. We have confirmed the assignment of the 7-methyl signal from observation of the nuclear Overhauser effect<sup>7</sup> for the 8-vinyl proton upon irradiation of the methyl signal at 436 c./sec.

The  $^1\text{H}$  n.m.r. spectrum of caffeine was examined over the concentration range of 0.005 to 0.1 M. Figure 1 shows the concentration dependence of chemical shifts of the various protons. The 1- and 3-methyl signals experience upfield shifts of nearly equal magnitude; signals of the 7-methyl and the C-8 protons also undergo significant upfield shifts. The chemical-shift data were treated by the method of Bangerter and Chan.<sup>8</sup> Based on the assumption that association is limited to dimerization, the method

requires visual extrapolation of chemical shift data to zero concentration. A digital computer then performs a least-squares search for best values of the dimer shift,  $\delta_{\text{dimer}} - \delta_{\text{monomer}}$ , and the dimerization constant,  $K$  (see Table). The close agreement of the dimer shifts of the 1- and 3-

*Dimer shifts and dimerization constant for caffeine in D<sub>2</sub>O at 30°*

Protons	$\delta_{\text{dimer}} - \delta_{\text{monomer}}$ (Hz)	$K$ (l. mole <sup>-1</sup> )	
1-Me	-37.7	5.3	Average $K$ 8.6
3-Me	-36.9	7.4	
7-Me	-21.2	12.5	
8-H	-11.3	9.3	

methyl protons indicates that these protons move into similar magnetic environments, different from those of the 7-methyl and 8-H protons.

We have calculated the  $\pi$ -electron charge distribution

(Figure 2a) and also the total atom charge distribution (Figure 2b) for caffeine by the CNDO/2 method. Assuming a parallel stacking interaction, these charge distributions suggest that the molecules are stacked directly over one another and oriented so that the complex has a  $C_2$  symmetry axis, as shown in Figure 2c. Such an arrangement would place the 1- and 3-methyl groups under the shielding region associated with the carbonyl groups of the adjacent molecule with the result that the methyl protons on the pyrimidine ring will experience a larger shielding than those on the five-membered ring. The observed chemical shift behaviour is consistent with this.

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