Preliminary communication

An n.m.r. study of the effects of fluorine substituents on the association between lysozyme and derivatives of 2-amino-2-deoxy-D-glucose

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Previous studies from this laboratory¹ have shown that fluorine nuclear magnetic resonance (¹⁹F n.m r.) parameters are generally more sensitive to changes in chemical environment than are proton (¹H) n m.r parameters. In order to evaluate the potential of ¹⁹F n m r. in studies of biochemically relevant systems, we have investigated the molecular association between lysozyme (salt-free, from hen egg-white, purchased from Worthington Biochemical Corporation) and some specifically fluorinated derivatives of 2-amino-2-deoxy-D-glucopyranose (D-glucosamine), this enzyme was chosen as a model system because of the detailed X-ray² and ¹H n.m r data³ that were already available in the literature; since this work was completed, two other laboratories have also made ¹⁹F n m.r. studies^{4,5}.

Four, N-trifluoroacetyl derivatives (1-4) of D-glucosamine were prepared by standard methods⁶. The ¹⁹F chemical shifts of various concentrations (0 10-0 02 M) of these substances were measured in the presence of lysozyme (3 0 mM), 2-hydroxy-1, 1,1-trifluoroethane was used as an internal reference.

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Analysis⁷ of the chemical-shift changes observed for 1 during such ¹⁹F n m.r experiments gave values (Table I) for the bound chemical-shift, Δ_B , and the binding constant, K_B . However, the lack of detectable changes in the shift of the ¹⁹F resonances of 2, 3, or 4 effectively precluded any calculations of the binding for these substances. Nevertheless, separate series of competition experiments clearly indicated that both 3 and 4 compete directly, although less strongly, for the same binding site as 7, a known inhibitor. Competition experiments between 1 and 2 indicated that, if 2 binds, it does so less strongly than 1

TABLE I $\begin{tabular}{ll} VALUES OF BINDING CONSTANTS (K_B) AND BOUND CHEMICAL-SHIFTS (Δ_B) FOR DERIVATIVES OF D-GLUCOSAMINES <math display="block"> \end{tabular}$

Compound	K _B (mole ⁻¹)	$\Delta_{\mathrm{B}}(p p m.)$
1	97	0 95
2	a	a
3	a	a
4	a	а
2 3 4 5 6	62 ^b 30 ^b	0 68 b
6	30 b	0 51 ^b
7	22.7	0 73
	19.2 ^c	
8	30 ^c	0 55 ^C 0.54 ^C

^a Within experimental error, there was no appreciable broadening or chemical shift of the ¹⁹F resonance of these compounds 23 a function of concentration. ^b Data taken from Ref. 3a. ^c Data taken from Ref. 7

In addition to the binding constant for 1, we have also determined the rates of association (k_1) and dissociation (k_{-1}) of its complex with lysozyme, the experiments were performed with a highly selective, pulse n m r spectrometer⁸, and the data were analysed by the method of Sykes^{3C} These rate constants, together with the literature values for 5, are listed in Table II In spite of the substantial experimental errors that are associated with such measurements, it seems that k_1 is higher for 1 than for 5

TABLE II ${\tt RATES\ OF\ ASSOCIATION\ (k_1)\ FOR\ THE\ INTERACTION\ OF\ D\text{-}GLUCOSAMINE\ DERIVATIVES\ WITH\ LYSOZYME }$

Compound	k ₁	k_1
1	1.1 × 10 ⁶	1.1 × 10 ⁴
5 a	3 5 × 10 ⁵	8 5 × 10 ³

a Data taken from Ref. 3d.

In addition to the above numerical data, several important details of perspective clearly emerge from this study. First, although, for this system, ¹⁹F chemical shifts are more sensitive to *inter*molecular changes in environment than are ¹H chemical shifts, the enhancement is not as marked as it is for *intra*molecular changes. Second, although a change in the chemical shift of a probe site of a small, reversible-inhibitor substance can be readily interpreted in terms of a molecular association, absence of any observed shift does not necessarily indicate an absence of molecular association. Finally, the expectation that the introduction of a fluorine substituent can cause changes in the nature of a molecular association, clearly demonstrated here, implies that specifically fluorinated substrates cannot *necessarily* be used as direct models for normal (non-fluorinated) substrates.

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