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Magnetic non-equivalence in phosphate esters. Solvent effects on the ¹H NMR spectrum of dibenzyl 2-pyridylphosphate

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Abstract

Methylene protons of benzyl groups in dibenzyl 2-pyridylphosphate show strong magnetic non-equivalence in $CDCl_3$, while the effect is absent in benzene- d_6 solutions. The non-equivalence is explained by the intramolecular $N \rightarrow P$ interactions leading to the "quasi trigonal bipyramidal" geometry of the substrate, observed in solid state, as determined by X-ray diffraction.

Keywords: NMR spectroscopy, Solvent effect, Phosphate ester

1. Introduction

The magnetic non-equivalence of the methylene protons (H_A, H_B) located in the vicinity of a chiral or prochiral centre is a well-recognized phenomenon [1] and includes some examples of the system 1:



In organophosphorus chemistry (1, Z = P), observation on "doubling the resonance signals" of the β - and γ -methyl groups in the ester functions of organic phosphates was reported over thirty years ago [2]. Williamson and Griffin [3] observed the magnetic non-equivalence of the O-methylene protons in the ¹H NMR spectra of three out of nine compounds of the type $(EtO)_2P(X)Y$ and explained the different chemical shifts of those protons in terms of the ethoxyl groups being attached to the P atom bearing three different substituents. For eight phosphonium ions of the structure $(PhCH_2)_2PXY^+$ the benzyl methylene hydrogens of five substrates showed non-equivalence in the ¹H NMR spectra recorded in CDCl₃ [4]. For one substrate the effect disappeared in DMSO- d_6 solution, while the temperature variation showed a negligible effect on the spectra. The most comprehensive

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study on the subject was carried out by Hall et al. [5], who examined ¹H NMR spectra of 22 organophosphorus substrates in various solvents and at a variable temperature. Only three non-chiral compounds of the type (EtO)₂P(X)Y showed the magnetic non-equivalence of the ethoxyl methylene groups; for MeP(S)(OEt)₂ the chemical shift difference varied in a series of solvents from $\Delta\delta_{AB} = 10.0$ Hz (in C₆D₆) to $\Delta\delta_{AB} = 6.3$ Hz (in DMSO-d₆). The results were interpreted in terms of "intrinsic" non-equivalence, combined with the conformational distributions.

We have recently found that the ¹H NMR spectra of some dibenzyl arylphosphates differ dramatically when recorded in two different solvents. In this paper we report the magnetic non-equivalence of the benzyl methylene groups in two closely related esters, dibenzyl phenyl (2a) and dibenzyl 2-pyridyl (2b) phosphate.



2. Experimental

2.1. Preparation of 2a

Triethylamine (3.49 ml, 25 mmol) was added dropwise, with stirring and cooling to 5°C, to a solution of phenol (2.35 g, 25 mmol) and dibenzyl phosphite (6.56 g, 25 mmol) in dry tetrachloromethane (16 ml). The reaction mixture was stirred overnight at room temperature, filtered, and the solvent was removed from the filtrate under reduced pressure. The product was purified by column chromatography (silica gel, CHCl₃/EtOH, 9:1) yielding pure **2a**, a colourless oil (54%). ¹H NMR (CDCl₃) δ 5.10 (4H, d, $J_{HP} = 8.3$ Hz), 7.11–7.36 (15H, m); ³¹P NMR δ – 5.58. Analysis. Calculated for $C_{20}H_{19}O_4P$: C, 67.8; H, 5.4. Found: C, 66.9; H, 5.5

2.2. Preparation of 2b

Prepared as **2a** from 2-hydroxypyridine (2.38 g, 25 mmol) and dibenzyl phosphite (6.56 g, 25 mmol) and purified by column chromatography (silica gel, CHCl₃/EtOH, 4:1) and recrystallized from ether; white solid (83%); m.p. $61.5-62.5^{\circ}$ C. ¹H NMR (C₆D₆) δ 5.24 (4H, d, $J_{HP} = 8.2$ Hz), 6.42 (1H, dd, $J_{HH} = 7.2$, 5.1 Hz), 6.74 (1H, d, $J_{HH} = 8.2$ Hz), 6.92 (1H, dt, $J_{HH} = 8.2$, 7.2, Hz), 7.05–7.12 (6H, m), 7.25–7.28 (4H, m), 7.99 (1H, dd, $J_{HH} = 5.1$, 2.0 Hz); ³¹P NMR δ – 5.50; ¹³C NMR (CDCl₃) δ 70.1 (d, $J_{CP} = 5.4$ Hz), 113.2 (d, $J_{CP} = 7.7$ Hz), 120.8 (s), 128.0 (s), 128.5 (s), 135.7 (d, $J_{CP} = 7.7$ Hz), 140.0 (s), 147.9 (s), 157. 6(s).

NMR spectra (structural assignments and nonequivalency) were recorded on a Bruker AC300 spectrometer at a probe temperature of 30° C. Chemical shifts are given relative to SiMe₄ (¹H, ¹³C) as an internal standard and 85% H₃PO₄ (³¹P) as an external standard. Variabletemperature measurements were carried out using a Varian Unity 400 spectrometer incorporating a Varian VT control unit. A temperature array was set between 25 and 60°C, using 5°C increments with a 300 s delay for thermal equilibration. NMR simulations were carried out using a Varian Unity 400 spectrometer with an IPX SPARC station data system; the LAOCOON program incorporated in the Varian VNMR software package was used.

2.3. Crystal structure determination

Accurate unit cell parameters were obtained by the least-squares method from the position of 25 selected centred reflections on an Enraf Nonius CAD4 diffractometer with Mo $K\alpha$ radiation using a graphite monochromator. Corrections for the observed crystal decay were applied, and intensities were also corrected for Lorentz and polarization effects. No absorption correction was applied. The structure was solved by standard Patterson and Fourier methods [6]. Refinement was done by a full-matrix least-squares method [7]. Atomic scattering factors were taken from the literature [8].

Data/parameter	Value used
Empirical formula	C ₁₉ H ₁₈ NO ₄ P
Molecular weight (g mol ⁻¹)	355.33
Crystal dimension (mm)	$0.21 \times 0.23 \times 0.23$
Space group	PT
Cell dimensions: a (Å)	5.822 (1)
b (Å)	9.188 (1)
c (Å)	17.389 (2)
α (deg)	76.82 (1)
β (deg)	85.07 (1)
γ (deg)	79.51 (1)
Z	2
Volume (Å ³)	889.6
D (calc.) (g cm ⁻³)	1.308
μ (cm ⁻¹)	1.35
Radiation (λ, \dot{A})	0.7107
T (°C)	22
F (000)	372.00
Scan type $(\omega/2\theta)$	1:0
Scan range (deg)	$3 \leq \theta \leq 30$
Zone collected: h	-8, 8
k	0, 12
1	-24, 24
Maximum scan speed (deg min ⁻¹)	5.49
Maximum scan time (s)	60
Scan angle $(\omega + 0.34 \tan \theta)$ (deg)	0.70
Aperture size (mm)	1.4×4.0
Reflections collected	5376
Decay (%)	11.8 (corrected)
Unique reflections used (> $3\sigma(I)$)	2872
R _{int}	0.017
Parameters refined	227
Maximum positional shift/esd	0.002
Residual electron density ($e Å^3$)	
Maximum	0.304
Minimum	-0.418
$U_{\rm iso}({\rm H})$ (Å ²)	0.134 (5)
R	0.072
R_w	0.056

Table 1 Crystal data and processing parameters for ester 2b

The non-hydrogen atoms were refined anisotropically and the hydrogen atoms were placed in idealized positions with a common isotropic thermal parameter that was also included in the refinement. Convergence was reached using a $\sigma^{-2}(F_0)$ weighting scheme. Crystal data and processing parameters are given in Table 1. The tables showing the structure factors, coordinates, and anisotropic thermal parameters have been deposited with the B.L.L.D. as Supplementary Publication No. SUP 26562 (16 pages).



Fig. 1. ¹H NMR spectrum of the methylene protons of dibenzyl phenylphosphate (**2a**) in: (a) chloroform-d; (b) benzene- d_6 .

3. Results and discussion

Sections of the ¹H NMR spectra (300 MHz) of **2a** and **2b** in CDCl₃ and in C_6D_6 are shown in Figs. 1 and 2, respectively. For each substrate in one solvent a "deceptively simple" signal of the methylene protons was observed: for 2a in CDCl₃ (d, $J_{HP} = 8.3$ Hz), and for **2b** in C₆D₆ (d, $J_{\rm HP} = 8.2$ Hz). On the other hand, **2a** in C₆D₆ showed the pattern expected for two ABX systems (two dd, $J_{HP} = 8.6$, 8.8 Hz; $J_{gem} = 11.7$ Hz), and for **2b** the ABX pattern (two dd, $J_{HP} = 7.8, 8.2$ Hz, $J_{\text{gem}} = 11.7$ Hz) was observed in CDCl₃. The reverse solvent effect indicated essentially different structural effects responsible for the observed behaviour. Ester 2a belongs to a group of phosphate derivatives where no specific, intramolecular interactions are possible, and numerous compounds of that type have been studied before. Magnetic non-equivalence, as demonstrated in the ¹H NMR spectra, seems to be a very subtle function of the substituents at phosphorus, as well as of solvation effects. For example, no non-equivalence of



Fig. 2. ¹H NMR spectrum of the methylene protons of dibenzyl 2-pyridylphosphate (2b) in: (a) chloroform-d; (b) benzene- d_6 .

methylene protons was observed for diethyl phenylphosphate [2], but it was observed for diethyl phosphorochloridate [5]. It was concluded that for a substantial difference in the chemical shifts $(\Delta \delta_{AB})$ to exist, "... the potentially anisotropic sensor protons must be influenced by a highly unsymmetrically substituted phosphorus atom." [5]. We believe that 2a belongs to that "general" type of phosphate, where the non-equivalence results from the fact that each OCH₂ group is located in the vicinity of phosphorus bearing three different substituents (phosphoryl oxygen, phenoxy group, and the second benzyloxy substituent). In a lowpolarity solvent (C_6D_6) the environment of the individual methylene protons is sufficiently different, so the expected signal, consisting of two doublets of doublets, is observed. In the more polar chloroform, in agreement with previous observations of the decrease of $\Delta \delta_{AB}$ with the solvent's polarity [5], the difference is too small to give rise to an observable ABX pattern, and the "deceptively simple' signal is obtained.

We propose, however, that the pyridyl ester **2b** belongs to a different group of substrates, and that

the observed non-equivalence is a consequence of specific, highly solvent-dependent, intramolecular interactions. Since the non-equivalence is observed in a more polar but non-aromatic solvent ($CDCl_3$), and not in a less polar but aromatic solvent (C_6D_6), we conclude that the specific solvation effects are capable of breaking those intramolecular interactions that are responsible for the non-equivalence. Because of such a dramatic difference in the ¹H NMR spectra of 2b in CDCl₃ and in C₆D₆, we were interested in the magnitude of the solvent effects on the chemical shift difference ($\Delta \delta_{AB}$) in the CH_AH_B system of 2b. In order to obtain the $\Delta \delta_{AB}$ value, simple simulation experiments were carried out for the methylene protons. For CDCl₃ as a medium, the methylene protons' signal can be described as an A_2B_2X system. Simulation of that system requires estimated values of both $J_{H(A)P}$ and $J_{H(B)P}$ coupling constants. Those values should, in principle, be obtainable from the H-coupled ³¹P NMR spectrum of 2b in CDCl₃; a signal consisting of two triplets is expected as a first-order approximation. The experimental spectrum revealed, however, partial overlap of signals yielding a broad quintet, from which the $J_{\rm HP}$ values could not be accurately determined. The ¹H NMR spectrum (Fig. 2), on the other hand, allowed us to estimate both coupling constants as $J_{H(A)P} = 8.12$ Hz, and $J_{H(B)P} = 7.76$ Hz. Using those two J values, simulation of the A₂B₂X system resulted in a spectrum which corresponded with the experimental spectrum to a reasonable degree (Fig. 3). From the simulation experiment a chemical shift difference of $\Delta \delta_{AB} = 23$ Hz (in CDCl₃) was obtained indicating a significant difference in the molecular environment of the two methylene protons. In addition, the geminal coupling constant, $J_{AB} = -10.5$ Hz was obtained, which is in reasonable agreement with the value estimated from the experimental spectrum, and with typical J_{gem} values reported in the literature [9]. The temperature effect on the non-equivalence in 2b was studied in CDCl₃ up to 60°C and was found to be negligible, in accordance with earlier reports for other systems [4,5,10].

In order to account for the observed magnetic non-equivalence of the methylene protons in **2b**, the following hypotheses were considered.



Fig. 3. ¹H NMR spectrum of the methylene protons of dibenzyl 2-pyridylphosphate (**2b**) in chloroform-d: (a) experimental spectrum; (b) simulate spectrum.

(i) Intermolecular interactions

Hudson et al. [11] found that dialky α hydroxyalkylphosphonates exist in the solid state as intermolecularly hydrogen-bonded dimers, and suggested that the bonding may persist in solvents of low polarity. Although the same authors showed that the observed chemical shift non-equivalence results not from H-bonding but from the chirality of the α -C atom, they pointed out that the restricted rotation due to the intermolecular association might also be responsible for the effect. Following the same argument, the non-equivalence in 2b could, in principle, be the result of a dimeric association via two $N \rightarrow P$ intermolecular interactions operating in the CDCl₃ solution. The ¹H NMR spectrum of 2b in CDCl₃ does not, however, show any changes with concentration, so the "dimerization" hypothesis was therefore disregarded.

(ii) Intramolecular ring stacking

Self-association of nucleotides via ring stacking of nucleobases is an important phenomenon [12], and since the molecule of **2b** contains three aromatic groups in a flexible framework, such interactions are conceivable for 2b. A Dreiding model of 2b revealed a possibility of such ring stacking with the pyridyl ring located between two phenyl groups. Such a structure would impose non-equivalence of the molecular environment of the methylene hydrogens, with one hydrogen of each group located "inside", and the other "outside", the ring-stacked system. This explanation was also discarded because of the observed solvent effect on the chemical shift of the pyridyl protons in **2b.** When pyridine itself is transferred from $CDCl_3$ to C_6D_6 , the signals of β -H and γ -H undergo a much stronger upfield shift than that of α -H; the effect was explained [13] by a differential shielding of the α, β, γ -hydrogens by solvating benzene molecules (the effect being absent in CDCl₃). When moving 2b from $CDCl_3$ to C_6D_6 , we observed almost the same change of the $\delta_{\rm H}$ values for the pyridyl hydrogens as for pyridine, so the explanation postulated for 2b in CDCl₃, a structure involving the pyridyl group "solvated" intramolecularly by two phenyl rings (a situation similar to that existing in benzene solution), had to be discarded.

(iii) Intramolecular $N \rightarrow P$ donor-acceptor interaction.

We have previously demonstrated [14] the existence of intramolecular interactions between nitrogen and phosphorus atoms in the 8-quinolylphosphate system leading to the distortion of the geometry at phosphorus due to a cyclic arrangement of the N-C-C-O-P linkage. The possibility of a similar (in this case, four-membered) $N \rightarrow P$ interaction in 2b was investigated by determining the structure of 2b by single crystal X-ray diffraction. Fig. 4 depicts a perspective view of the molecule with atomic nomenclature, while Table 2 lists selected bond lengths, bond angles, and dihedral angles.

According to a general effect observed for phosphate triesters [14], the geometry at the phosphorus atom in **2b** deviates from a regular tetrahedron in a sense of smaller O-P-O angles involving ester P-O bonds (98.2-106.4°) and larger angles involving the phosphoryl group (114.6-118.3°). Superimposed on that effect, however, is a secondary effect involving the interaction between the



Fig. 4. Crystal structure of dibenzyl 2-pyridylphosphate (2b).

nucleophilic pyridyl nitrogen and the electrophilic phosphoryl centre. The $N \cdots P$ non-bonded distance in **2b** is 3.010 Å, while the corresponding sum of van der Waals radii is 3.4 Å [15]. This "short contact" clearly indicates $N \rightarrow P$ intramolecular interaction and is expected to change the geometry of the phosphorus centre towards

Table 2

Selected molecular parameters of dibenzyl 2-pyridylphosphate 2b

Bond distances (Å)		
P-O ₁	1.453(3)	
P-O ₂	1.595(3)	
P-O ₃	1.563(3)	
P-O ₄	1.542(3)	
$O_2 - C_1$	1.394(5)	
O ₃ -C ₆	1.462(4)	
$O_4 - C_{13}$	1.442(4)	
Bond angles (deg)		
$O_1 - P - O_2$	114.6(2)	
$O_1 - P - O_3$	118.3(2)	
O ₁ -P-O ₄	117.7(2)	
$O_2 - P - O_3$	98.7(2)	
O ₂ P-O ₄	106.4(2)	
$O_3 - P - O_4$	98.2(2)	
Torsional angles (deg)		
$P-O_2-C_1-N$	30.6(0.5)	
$O_3 - P - O_2 - C_1$	173.1(0.3)	
$O_4 - P - O_2 - C_1$	-85.6(0.4)	
$\frac{P-O_2-C_1-C_5}{2}$	-152.0(0.4)	

that of a trigonal bipyramidal. Analysis of the molecular parameters given in Table 1 confirm the postulated structural effect. The observed dihedral angles, $P-O_2-C_1-N$ (30.65°; ideal 0°) and $P-O_2-C_1-C_5$ (-152.04°; ideal-180°), are an indication of the tendency of nitrogen to locate itself in the vicinity of phosphorus in a quasi four-membered ring. The angles $O_3 - P - O_2 - C_1$ (173.04°; ideal 180°) and $O_4 - P - O_2 - C_1$ (-85.62°; ideal-90°) suggest the 'quasi apical' and 'quasi equatorial" positions of the two benzyloxy groups. These conclusions are corroborated by the slightly longer $P-O_3$ bond (1.563 Å) than the $P-O_4$ bond (1.542 Å), as well as by the bond angles $O_1 - P - O_2$ (114.6°), $O_1 - P - O_4$ (117.7°), and $O_2 - P - O_4$ (106.4°), tending towards the ideal value of 120°, and the angles $O_2 - P - O_3$ (98.7°) and $O_3 - P - O_4$ (98.2°) approaching the ideal value of 90°. As a result of these intramolecular effects, the geometry of 2b can be presented (in an exaggerated form) by Fig. 5.

In chloroform solution, 2b may retain its "quasi trigonal bypyramidal" geometry found in the solid state. That geometry necessarily results in a strong molecular non-equivalence of the methylene groups ("quasi apical" and "quasi equatorial"). Fast (on the NMR time-scale) pseudorotation should lead to an exchange of both benzyloxy substituents, but the individual H_A and H_B atoms of each methylene group will always exist in a different molecular environment, leading to the observed non-equivalence in the ¹H NMR spectrum. In benzene all aromatic groups in 2b (including the pyridyl ring) should be effectively solvated (intermolecular ring stacking), so the stabilization of the system via the donor-acceptor interactions leading to the geometry shown in Fig. 5 is no longer necessary. The chemical shift differences for the methylene hydrogens of two benzyloxy groups should



Fig. 5. "Quasi trigonal bypiramidal" geometry of 2b.

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