

MEASUREMENT OF LONG-RANGE HETERONUCLEAR COUPLINGS : APPLICATION TO OLIGOSACCHARIDE CONFORMATION

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ABSTRACT : $^3J_{(C,H)}$ values have been measured using 2D-J heteronuclear-resolved n.m.r. spectroscopy to estimate interglycosidic conformations in various oligosaccharides with a degree of polymerization ranging from 6 to 30.

One of the most important n.m.r. parameters in conformational studies of all classes of molecules is the vicinal or three-bond coupling constant (3J) between nuclei like ^{13}C and 1H in pathways which could involve heteroatoms (oxygen, nitrogen, ...). These couplings show a clear correlation with the dihedral angle θ between the two planes defined by three atoms, as was first noticed by Lemieux *et al.*¹ and Schwarcz *et al.*² for carbohydrates. This relation is often represented by the general formula :

$$^3J = A \cos^2 \theta - B \cos \theta + C$$

Coefficients A, B and C depend mainly on the nature of the four atoms in the pathway. Such a relation is derived from the first Karplus relation established for H-C-C-H fragments.³ Its applicability has been proved both theoretically and by measurements on conformationally rigid molecules for a large number of molecular fragments.⁴

In the carbohydrate field, the measurement of such couplings has been obtained through studies of ^{13}C -enriched and selectively deuterated compounds.⁵⁻¹⁰ The selective 2D-J heteronuclear experiment proposed by Bax and Freeman¹¹ and variously used by different authors ¹²⁻¹⁷ has also been performed on mono-, di- and some oligo-saccharides in natural abundance compounds.

In order to expand the data base (with the ultimate purpose of establishing accurate Karplus-type relationships between $^3J_{(C,H)}$ values and the corresponding dihedral angles) and to obtain experimental data which can be safely used in conformational analysis, we now report our results on a series of oligosaccharides with a degree of polymerization ranging from 6 to 30, using a modified version of Bax and Freeman sequence¹⁵.

The relative orientation of a pair of contiguous sugar residues is described in the

usual way by the torsion angles ϕ and ψ around the glycosidic bonds C1'--O and O--Cn respectively in the fragment H1'--C1'--O--Cn--Hn (primed numbers refer to the glycone and unprimed to the aglycone).

The $^3J(C,H)$ values measured are listed in Table 1. An example of resolution obtained by this technique is shown on Figure 1 with the **1a-1c** compounds. The solvent was CDCl₃. The spectra were obtained on Bruker AM 300 or AM 400.

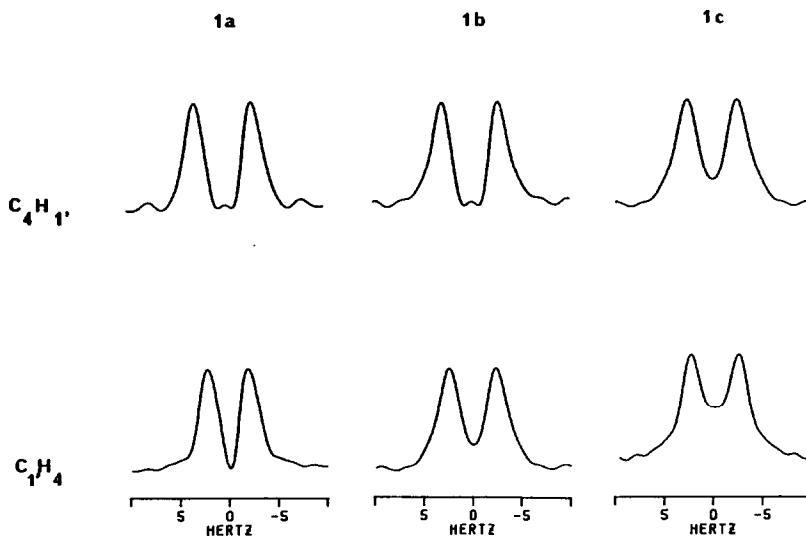
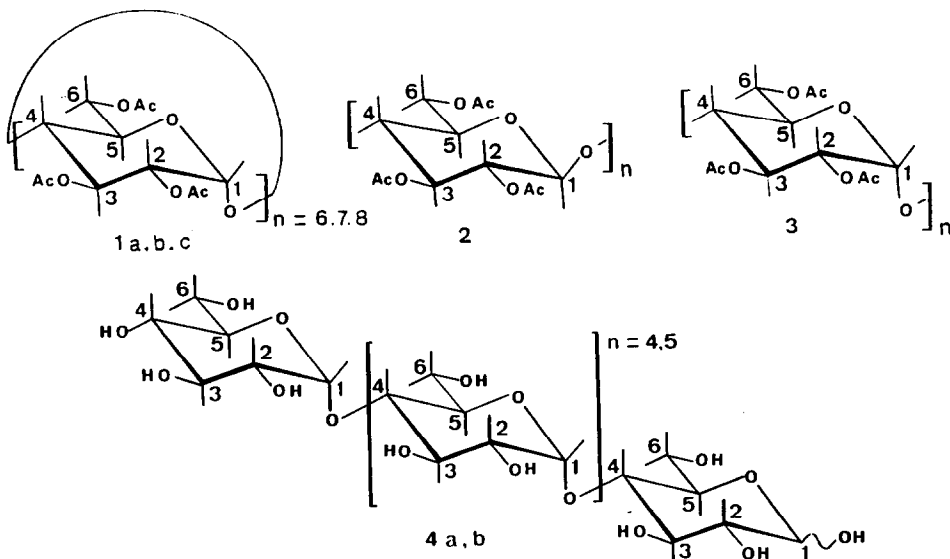


Figure 1 : 2D F1-dimension slices of C-1 and C-4 corresponding to H-4 (bottom) and H-1' (top) selective inversion on compounds **1a**, **1b** and **1c**. Coupling constant values have been measured without any correction.



These values were used to estimate the torsion angle about the glycosidic bond via an equation with the following parameters $A = 5.7$, $B = 0.6$, $C = 0.5$. These were established for a series of conformationally rigid carbohydrate derivatives¹⁸ and they lead to a Karplus-type curve which fits fairly well our values obtained previously¹⁵. They are also very close to the parameters used by B. Mulloy *et al.*¹⁷.

The torsion angles determined for the acetylated cyclomalto-dextrines are all in the region 0-35° as shown below.

Interglycosidic carbon-proton coupling constants for oligosaccharides.

Compound	Linkage	$^3J_{C,H}$	Value(Hz) ^a	Estimated Torsion angle ^b
Peracetate cyclomaltohexaose (1a)	$\alpha(1,4)$	C-4,H-1'	5.6	0
		C-1',H-4	4.0	33
Peracetate cyclomaltoheptaose (1b)	$\alpha(1,4)$	C-4,H-1'	5.5	0
		C-1',H-4	4.6	25
Peracetate cyclomaltooctaose (1c)	$\alpha(1,4)$	C-4,H-1'	4.8	22
		C-1',H-4	4.5	26
Cellulose triacetate (DP 30) (2)	$\beta(1,4)$	C-4,H-1'	5.6	0
		C-1',H-4	5.7	0
Amylose triacetate (DP 30) (3)	$\alpha(1,4)$	C-4,H-1'	2.9	45
		C-1',H-4	4.6	25
Maltose (4a)	$\alpha(1,4)$	C-4,H-1' ^c	4.3	29
		C-1',H-4 ^c	5.0	19
Maltose (4b)	$\alpha(1,4)$	C-4,H-1' ^c	4.4	28
		C-1',H-4 ^c	5.1	18

^a Coupling constants are given at ± 0.5 Hz. This lead to estimate the torsion angle to $\pm 10^\circ$.

^b As several solutions exist for a given coupling constant only the absolute value in the zero region is presented.

^c Coupling constants corresponding to central residues

They show a slight variation of the torsion angles in parallel to the increase of the annular aperture in going from the cyclic hexasaccharide (1a) to the octasaccharide (1c). This is characterized by a great similarity between the cyclic hexa- and heptasaccharides, as already mentioned by Lindner *et al.*¹⁹ for the OH-form of the cyclomalto-dextrines, and also by an adjustment of the ϕ torsion angle for the octasaccharide in order to assume a larger radius of the macrocycle.

The conformation of cellulose triacetate (2) in $CDCl_3$ can be described by torsion angles close to zero. This result is consistent with the torsion angles found in the solid state ($\phi = 15^\circ$, $\psi = -22^\circ$) from X-ray and electron diffraction analyses.²⁰ These lead to the presence in solution of chains with a two-fold axis.

The results for amylose triacetate (3) could indicate that in solution the molecule has a relatively high degree of freedom as the dihedral angles estimated in our study differ from

those published for the crystalline structure²¹ (-39° and -50° for the ϕ and ψ angles respectively).

At last within the accuracy of our data, the coupling constants corresponding to the central residues for maltohexaose (4a) and maltoheptaose (4b) are similar indicating a very close conformation for these two oligosaccharides of the amylose family.

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