Table 2. Lattice parameters and thermal-expansion data of the compound Ni<sub>11</sub>Zr<sub>9</sub> at different temperatures

Heat treatment and water quenching		a (Å)		c (Å)			
Temperature (K)	Time (min)	Observed	Calculated from linear equation	Observed	Calculated from linear equation	$\begin{array}{c} \alpha_a \times 10^6 \\ (K^{-1}) \end{array} \qquad \alpha_a \end{array}$	$\begin{array}{c} \alpha_c  imes 10^6 \ (K^{-1}) \end{array}$
298	_	9-9646 (58)	9.9661	6.6940 (42)	6.6944	28.05 (1)	29.71 (4)
(As cast)							
1003	70	10-1626 (58)	10.1631	6.8357 (42)	6.8346	27.50(1)	29.10 (4)
1093	60	10.1892 (58)	10.1883	6.8538 (42)	6.8525	27.43 (1)	29.02 (4)
1223	50	10.2250 (58)	10.2246	6.8758 (42)	6.8783	27.33(1)	28.93 (4)
1293	40	10.2558 (58)	10.2442	6.8950 (42)	6.8923	27.25(1)	28.85 (4)
1423	60	10.2703 (58)	10.2805	6.9125 (42)	6.9181	27.21(1)	28.77 (4)
1493	30	10-2996 (58)	10.3001	6.9359 (42)	6.9320	27·14 (1)	28·68 (4)

Kirkpatrick & Larsen (1961) have shown the existence of this phase with a b.c.t. structure but the reported parameters appear to be tentative. The correctness of the crystal structure of this phase has to be decided by comparison with the phase  $Pt_{11}Zr_9$  (which has 40 atoms per unit cell and is b.c.t. with a = 10.257 and c = 6.888 Å) reported by Panda & Bhan (1974). Comparison shows that  $Ni_{11}Zr_9$  is isomorphous with the  $Pt_{11}Zr_9$  phase. This is expected as Ni and Pt are both members of the  $T^{10}$  group with ground-state electron configurations of their free atoms of  $3d^8 3s^2$  and  $5d^8 6s^2$  respectively and both have similar alloying behavior (Panda & Bhan, 1973).

## Thermal-expansion coefficient

The X-ray photographs of the alloy  $Ni_{55}Zr_{45}$  at each temperature were indexed and cell dimensions were determined using a least-squares treatment (Cohen, 1935, 1936). The standard errors in the lattice parameters so obtained were calculated by the method suggested by Jette & Foote (1935).

The lattice parameters vary linearly with temperature and the dependence may be expressed analytically as  $a_T =$ 9.8828 (14) + 2.795 (13) × 10<sup>-4</sup> T; and  $c_T =$  6.6351 (34) + 1.989 (31) × 10<sup>-4</sup> T; where  $a_T$  and  $c_T$  are the lattice parameters at TK.

The linear coefficients of thermal expansion for both parameters are calculated by  $\alpha_a = (1/a)(da_T/dT)$  and  $a_c = (1/c)(dc_T/dT)$  and are listed in Table 2.

The American Institute of Physics Handbook (1972) only gives the values of  $\alpha$  for pure Ni and Zr at 293 K, but the thermal-expansion coefficient of an alloy cannot be predicted from those of its constituents (Lonsdale, 1968).

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On the 'riding-model' correction for bond lengths. By R. SRINIVASAN and N. R. JAGANNATHAN, Department of Crystallography and Biophysics,\* University of Madras, Guindy Campus, Madras-600025, India

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#### Abstract

It is usual in structural reports based on X-ray or neutron diffraction methods to correct bond lengths for thermal-

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motion effects. Two common procedures adopted are the riding model [Busing & Levy (1964). Acta Cryst. 17, 142–146], and the rigid-body model [Schomaker & Trueblood (1968). Acta Cryst. B24, 63–76], the specific choice being dependent on the circumstances. It would appear that no critical assessment is available in the literature as to how

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sound and reliable these corrections are. This paper attempts to provide such an assessment, specifically for the ridingmodel correction, purely from an empirical point of view, by comparing neutron diffraction data with wide-line proton magnetic resonance (PMR) and microwave data.

We recently examined the neutron diffraction data for a number of small organic molecules with a view to arriving at the most appropriate geometry of groups such as NH<sup>+</sup><sub>1</sub>, CH<sub>1</sub> and CH<sub>2</sub>. This was undertaken in connection with the theoretical calculation of second moments for the interpretation of wide-line PMR absorption spectra. The internuclear distances enter the expression for the second moment as the inverse sixth power (Van Vleck, 1948) and hence precise coordinates of the protons (in the case of PMR) are important in these calculations. One of the assumptions in the riding-model correction for a bond length is that one of the atoms is light compared to the other to justify the riding concept. H attached to C, N or O in organic molecules would seem to qualify well in this respect. H-atom location from X-ray data tends to be less precise and there have been several instances where unrealistically large values of the second moment for organic compounds were estimated by use of X-ray coordinates (Watton, Reynhardt, Sandhu & Petch, 1977; Jagannathan & Srinivasan, 1982a). This obviously results from the fact that X-ray data normally yield lengths which are too short for bonds such as C-H and N-H, especially in groups such as CH<sub>3</sub> and NH<sub>4</sub><sup>+</sup> (Tomiie, 1958; McDonald, 1960; Banyard & March, 1961; Nahringbauer, 1967; Khan, Boux & James, 1972). Our analysis thus concentrates on neutron diffraction data, since these yield more reliable coordinates. Also we shall confine ourselves to riding-model corrections, since the data available on rigid-body corrections are rather meagre for a critical analysis.

Initial statistical analysis (Jagannathan & Srinivasan, 1982b) carried out on about 80 organic compounds containing these groups gave bond-length averages for C-H of about 1.090 Å, if only uncorrected values are used. In many cases, bond lengths corrected by the riding model were also available and the average corrected C-H length was about 1.126 Å. The standard deviations in these averages were, however, rather large ( $\sim > 0.015$  Å). This was obviously because no special effort was made to screen the structures in relation to accuracy of analysis and applicability of the riding-model correction. The latter aspect has been discussed by Brown (1968).

Thus, to make the analysis more effective the current investigation is restricted to a select group of 16 carbohydrates which have been studied by neutron diffraction. These are reasonably accurate structures as judged by the final R factors and the e.s.d.'s in bond lengths and other parameters (Table 1). Also, the analysis is restricted to the three groups CH<sub>3</sub>, CH<sub>2</sub> and CH. An additional factor governing the choice of these compounds is that wide-line PMR studies are available for three sugars, and these studies show that the structures are essentially 'rigid' even at room temperature. This reduces the possible inaccuracy in the second moment as would be the case if group rotations exist. Thus the experimental second moment is interpreted more reliably in terms of the model geometry from the statistical analysis.

# Table 1. Compounds included in the statistical analysis

Serial		R	
number	Compound	factor	Reference
(1)	α-L-Xylopyranose	0.036	( <i>a</i> )
(2)	$\beta$ -D-Fructopyranose	0.053	(b)
(3)	$\beta$ -L-Lyxopyranose	0.048	(c)
(4)	3-Amino-1,6-anhydro-3- deoxy-β-D-glucopyranose	0.056	( <i>d</i> )
(5)	Methyl a-D-mannopyranoside	0.030	(e)
(6)	Methyl a-D-glucopyranoside	0.046	(e)
(7)	$\beta$ -D,L-Arabinopyranose	0.058	(í)
(8)	Methyl α-D-galactopyranoside monohydrate	0.055	(g)
(9)	Methyl $\beta$ -D-galactopyranoside	0.058	(g)
(10)	a-L-Sorbopyranose	0.044	$(\tilde{h})$
(11)	a-d-Glucose	0.048	(i)
(12)	α-L-Rhamnose monohydrate	0.042	Ű
(13)	$\beta$ -L-Arabinose	0.035	(k)
(14)	Methyl $\beta$ -D-xylopyranoside	0.045	(k)
(15)	Methyl $\alpha$ -D-altropyranoside	0.089	$\tilde{U}$
(16)	Sucrose	0.033	(m)

References: (a) Takagi & Jeffrey (1979c); (b) Takagi & Jeffrey (1977b); (c) Nordenson, Takagi & Jeffrey (1978); (d) Noordik & Jeffrey (1977); (e) Jeffrey, McMullan & Takagi (1977); (f) Takagi & Jeffrey (1979b); (g) Takagi & Jeffrey (1979a); (h) Nordenson, Takagi & Jeffrey (1979); (j) Takagi & Jeffrey (1979); (j) Takagi & Jeffrey (1978); (k) Takagi & Jeffrey (1977a); (l) Poppleton, Jeffrey & Williams (1975); (m) Brown & Levy (1973).

Table 2. Average bond lengths (Å) and bond angles (°) for different groups (values in parentheses are e.s.d.'s)

Group	Туре	Uncorrected	Minimum riding	Riding
СН	(C-H)	1.100 (4)	1.104 (4)	1.123 (3)
CH <sub>2</sub> (ring)	(C-H)	1.093 (5)	1.097 (5)	$1 \cdot 118(5)$
	(H-C-H)	109.0 (5)	-	_
CH <sub>2</sub> (free)	(C-H)	1.090 (7)	1.097 (6)	1.127 (6)
	(H-C-H)	108.0 (7)		-
CH3	(C-H)	1.054 (16)	1.082 (9)	1.142 (17)
	(H-C-H)	108.6 (19)		- ` `
	(X-C-H)	110.3 (16)	-	_

Table 2 shows that the uncorrected C–H bond length in the CH<sub>3</sub> group has a mean value of 1.054 Å (with a standard error of  $\pm 0.016$  Å) as against means of 1.092 Å ( $\pm 0.004$  Å) and 1.100 Å ( $\pm 0.004$  Å) for CH<sub>2</sub> and CH groups. This would seem reasonable since the CH<sub>3</sub> group is a terminal group and is likely to be affected by rotational and other motional effects.

In the case of the CH<sub>2</sub> group a minor subdivision was felt necessary depending upon whether or not the group occurs in the sugar ring. The mean C-H bond lengths are not different, but there appears to be a significant difference in the angle H-C-H.

It is of interest to note that the mean corrected values are  $1 \cdot 142$  Å for the CH<sub>3</sub> group and  $1 \cdot 123$  Å for CH<sub>2</sub> and CH groups, although one would have expected that the corrected values would be the same. However, the indication is clear that the corrected value for the CH<sub>3</sub> group (which has larger motional effects) is higher than for the other groups. This

 Table 3. Observed second moments for some sugars compared with theoretical values

The units for second moment are  $G^2$  (1 G  $\doteq 10^{-4}$  T).

	Second moment using actual neutron diffraction data			
Compound	Average C-H in the structure (Å)	Theoretical second moment	Theoretical second moment with e C-H = $1 \cdot 127 \text{ Å}$	Observed experimental second moment
D-Glucose L-Xylose L-Sorbosc	1.097 1.099 1.095	8.9 14.1 14.5	7.6 10.6 13.6	8·5 (0·5)* 13·1 (0·4) 13·9 (0·7)

\* Values in the parentheses are e.s.d.'s.

 
 Table 4. C-H bond lengths for various groups from microwave data

Values in parentheses are e.s.d.'s.

Group	Bond length
CH <sub>3</sub>	1·095 (3) Å
CH,	1.090 (3)
CH	1.094 (4)
Overall	1.093 (3)

itself might be attributed to the fact that the correction is overdone. However, additional evidence arises as follows.

The wide-line PMR results for three of the sugars are given in Table 3. The theoretical values of the second moment listed correspond to two geometries for the  $CH_2$  and CHgroups, namely C-H lengths with and without correction. The uncorrected set corresponds to the actual neutron diffraction results for these compounds, while for the corrected set the C-H bond lengths were changed uniformly to 1.127 Å. The angles were kept as observed in the neutron diffraction analyses. This corresponds to using 1.097 and 1.127 Å respectively for the C-H bond lengths. It is clear from Table 3 that the agreement is poor for the corrected set and very much better for the uncorrected one.

Further independent evidence is obtained from microwave data. Table 4 lists the average C-H values in different groups derived from about 25 simple organic compounds. The geometry of these groups is more precisely determined from microwave data as is evident from the quoted e.s.d.'s. The overall mean of 1.093 (3) Å (Table 4) also indicates that the values in column 5 of Table 2 are overcorrected. Although we have not discussed angles, the microwave data broadly support the general results from diffraction data (for example, the H-C-H angle in the CH<sub>3</sub> group is ~108.7°, less than the tetrahedral value).

As regards column 4 of Table 2 where minimum corrected C-H lengths are listed, these are also rather unsatisfactory and inconsistent. For instance, the average for CH<sub>3</sub> is 1.082 (9) Å, distinctly less than 1.093 (3) Å (which may, for the present, be taken as standard), whereas the average taken over the CH, and CH groups is 1.099 (5) Å.

From these considerations we may conclude that bond lengths corrected by the riding model tend to yield overcorrected values. The uncorrected values, particularly for groups such as  $CH_2$  and CH which are relatively rigid, should be preferred and are probably close to the true values than values for groups such as  $CH_3$  where motional effects are large. The conclusion drawn above, although based on data for  $CH_3$ ,  $CH_2$  and CH groups, may be expected to hold good in general for other groups such as  $NH_4^+$ ,  $NH_3^+$ ,  $NH_2$ and OH.

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