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Assignment of the NMR Signals for *cis,cis,cis*-RuCl₂(Me₂SO)₂(Me₃Bzm)₂.

The B12H₃ I signal at 3.95 ppm (assigned to ligand "b") has intense NOE cross-peaks to two signals, "b" B2H I and "b" B7H I. "b" B7H I was assigned from its second NOE to a methyl signal at 2.16 ppm, "b" B11H₃ I. "b" B11H₃ I has an intense NOE to a methyl signal at 1.70 ppm assigned to "b" B10H₃ I. The unassigned signal at 6.07 ppm is "b" B4H I, which possesses an expected NOE to "b" B10H₃ I. This same pattern was seen for the second set of I signals. "a" B12H₃ I at 3.90 ppm has an NOE to two signals, "a" B2H I and "a" B7H I. The methyl signal with an NOE to "a" B7H I was assigned to "a" B11H₃ I at 2.11 ppm. This signal in turn had an NOE to another methyl signal (1.71 ppm) assigned to "a" B10H₃ I. "a" B10H₃ I had an NOE to the final unassigned I signal (6.73 ppm), assigning it to "a" B4H I. B12H₃ II resonating at 3.92 ppm (assigned to ligand "b") has NOE's to signals at 8.90 ppm and 7.17 ppm, "b" B2H II and "b" B7H II, respectively. The more upfield signal has an additional NOE to a methyl signal at 2.31 ppm distinguishing it as "b" B7H II. The 2.31 ppm signal is "b" B11H₃ II. "b" B11H₃ II has an NOE to "b" B10H₃ II (1.85 ppm). This signal in turn has an NOE to the 6.11 ppm signal, "b" B4H II. The final B12H signal ("a" II) at 3.36 ppm has NOE's to "a" B2H II and "a" B7H II. As per the above arguments, "a" B7H II was assigned to the more upfield signal at 7.06 ppm and "a" B2H II to the more downfield signal at 7.43 ppm. The more upfield signal has an NOE to a methyl signal at 2.41 ppm, assigning this signal to "a" B11H₃ II. The final unassigned methyl signal at 2.48 ppm is "a" B10H₃ II. The expected NOE between "a" B10H₃ II and "a" B11H₃ II cannot be seen due to the close proximity of the NOE to the diagonal. As expected, "a" B10H₃ II has an NOE to a more downfield signal at 8.25 ppm, "a" B4H II. Of note, slightly less intense exchange-NOE's were also seen between rotamers. For example, an exchange-NOE is present between "a" B2H I and "a" B12H₃ II. These NOE's result from dipole-dipole interactions which remain after exchange between I and II. These exchange NOE's were distinguished by the lower intensity compared to direct NOE's.

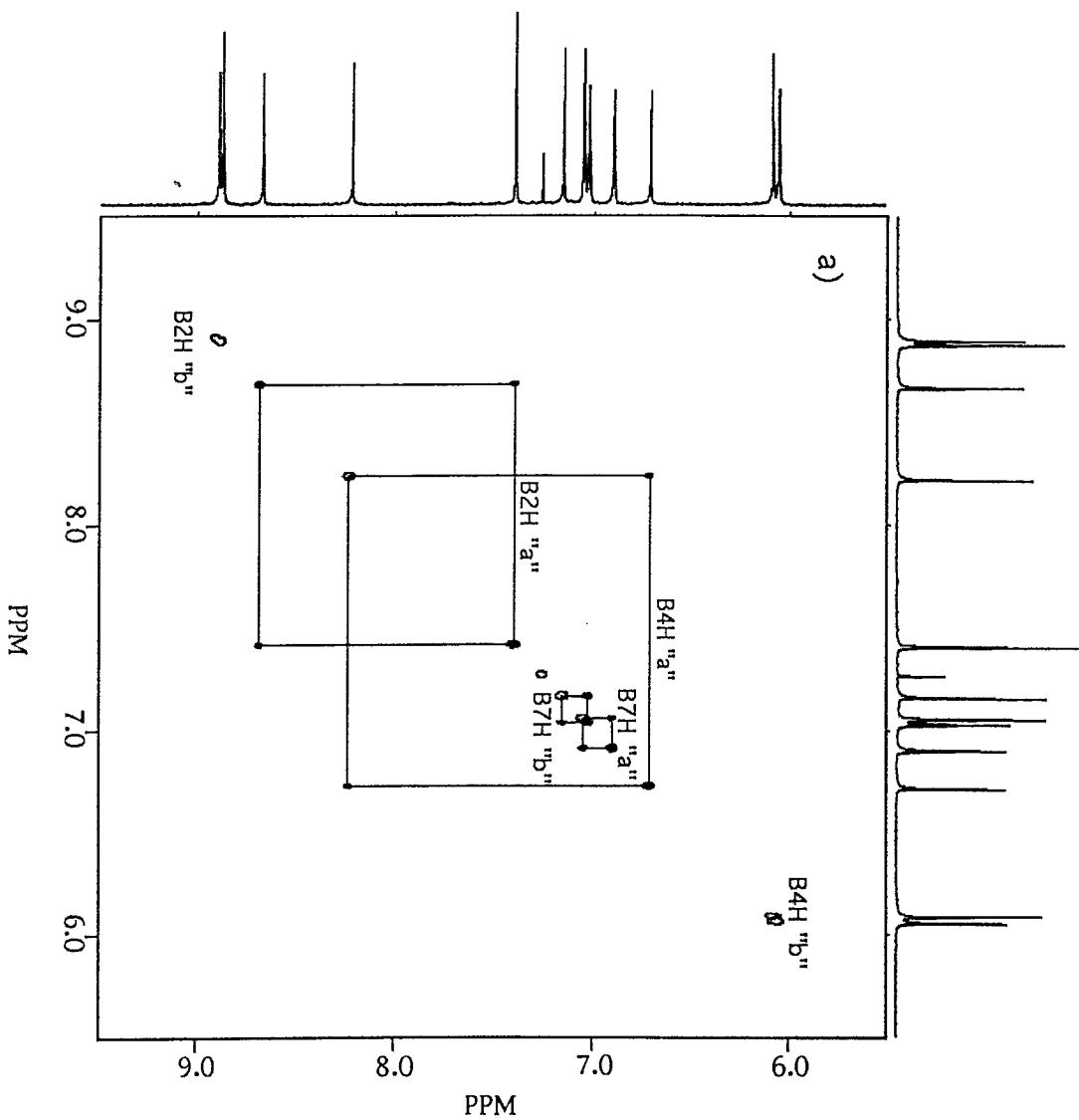
P-2545-m2

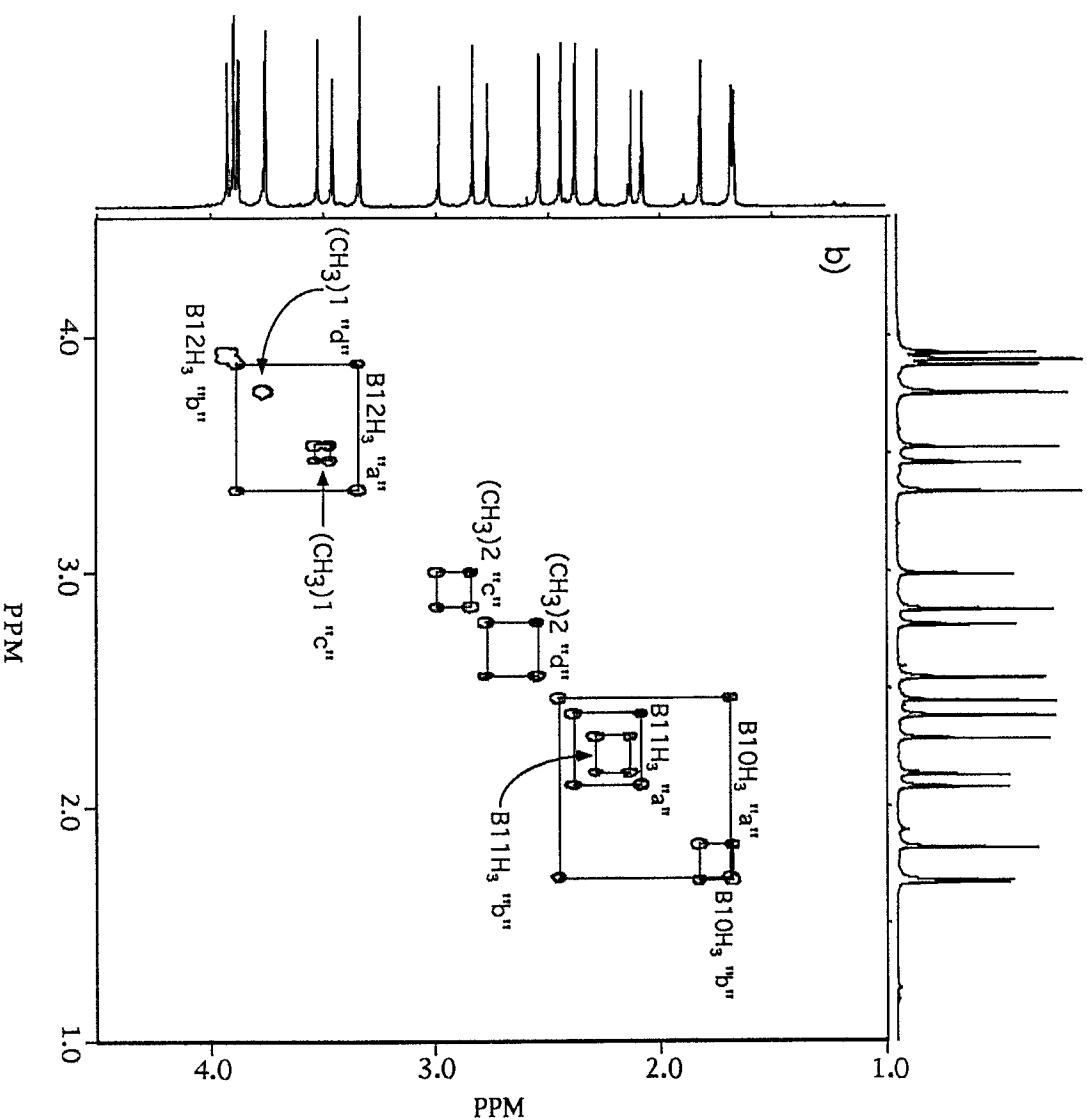
38

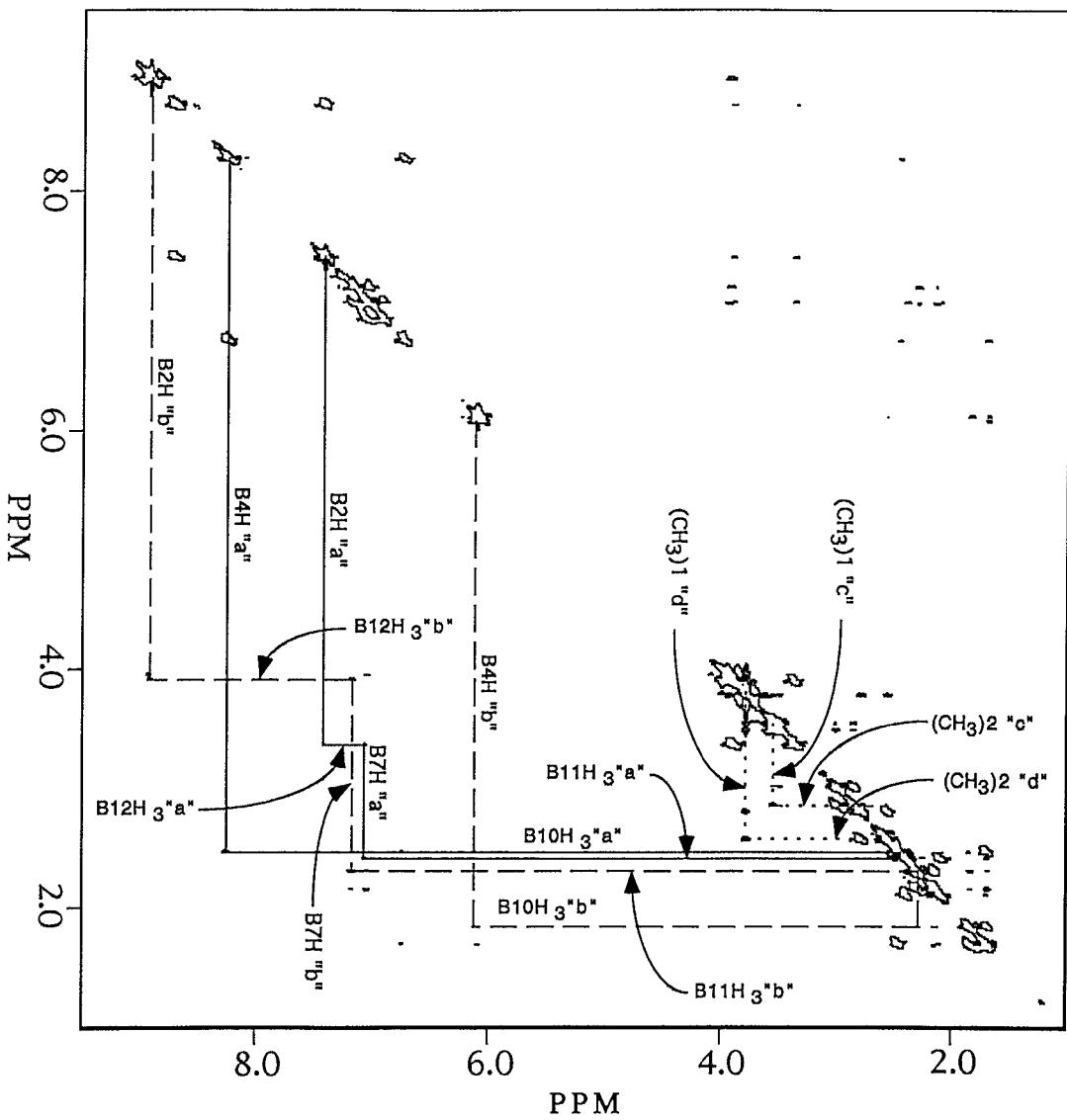
Supplementary Figure Captions

Figure S1. 2D ^1H Homonuclear EXSY spectrum of *cis,cis,cis*-RuCl₂(DMSO)₂(Me₃Bzm)₂ (**2**): a) downfield correlations, and b) upfield correlations. See Schemes 1 and 2 for Me₃Bzm notation and ligand positions, respectively.

Figure S2. 2D ^1H Homonuclear ROESY spectrum of *cis,cis,cis*-RuCl₂(Me₂SO)₂(Me₃Bzm)₂ (**2**). Assignment paths for II ligands "a" (—), "b" (- - -), "c" (.....), and "d" (.....) are depicted. See Scheme 1 for Me₃Bzm notation and NOE connectivity path. Scheme 2 shows ligand positions.







P-2545-m6

Table of Bond Distances of 1 in Angstroms

Atom 1	Atom 2	Distance	Atom 1	Atom 2	Distance
=====	=====	=====	=====	=====	=====
Ru	C11	2.4148(7)	N3	C10	1.343(3)
Ru	C12	2.4233(6)	N3	C11	1.454(3)
Ru	S1	2.2576(7)	N3	C12	1.380(3)
Ru	S2	2.2541(6)	C5	C6	1.381(4)
Ru	N1	2.110(2)	C6	C7	1.368(4)
Ru	N2	2.124(2)	C7	C8	1.361(4)
S1	O1	1.479(2)	C8	C9	1.378(4)
S1	C1	1.781(3)	C12	C13	1.397(3)
S1	C2	1.783(3)	C12	C19	1.390(3)
S2	O2	1.478(2)	C13	C14	1.379(3)
S2	C3	1.782(3)	C14	C15	1.507(3)
S2	C4	1.781(4)	C14	C17	1.416(3)
N1	C5	1.349(3)	C16	C17	1.506(3)
N1	C9	1.344(3)	C17	C18	1.384(3)
N2	C10	1.325(3)	C18	C19	1.398(3)
N2	C19	1.399(3)			

Numbers in parentheses are estimated standard deviations.

P-2545-m7

Table of Bond Angles of 1 in Degrees

Atom 1	Atom 2	Atom 3	Angle	Atom 1	Atom 2	Atom 3	Angle
=====	=====	=====	=====	=====	=====	=====	=====
C11	Ru	C12	88.89(2)	Ru	N1	C9	122.6(2)
C11	Ru	S1	93.27(2)	C5	N1	C9	116.7(2)
C11	Ru	S2	87.70(2)	Ru	N2	C10	123.2(1)
C11	Ru	N1	176.20(6)	Ru	N2	C19	131.4(1)
C11	Ru	N2	87.29(5)	C10	N2	C19	105.0(2)
C12	Ru	S1	89.95(2)	C10	N3	C11	126.4(2)
C12	Ru	S2	174.96(3)	C10	N3	C12	107.0(2)
C12	Ru	N1	88.07(5)	C11	N3	C12	126.6(2)
C12	Ru	N2	86.32(5)	N1	C5	C6	122.4(2)
S1	Ru	S2	93.95(2)	C5	C6	C7	119.7(3)
S1	Ru	N1	89.01(5)	C6	C7	C8	118.5(3)
S1	Ru	N2	176.21(5)	C7	C8	C9	119.6(3)
S2	Ru	N1	95.18(5)	N1	C9	C8	123.0(2)
S2	Ru	N2	89.81(5)	N2	C10	N3	113.0(2)
N1	Ru	N2	90.23(7)	N3	C12	C13	131.9(2)
Ru	S1	O1	117.28(9)	N3	C12	C19	106.3(2)
Ru	S1	C1	113.8(1)	C13	C12	C19	121.8(2)
Ru	S1	C2	113.7(1)	C12	C13	C14	118.2(2)
O1	S1	C1	105.3(1)	C13	C14	C15	119.6(2)
O1	S1	C2	106.4(1)	C13	C14	C17	120.4(2)
C1	S1	C2	98.4(2)	C15	C14	C17	120.0(2)
Ru	S2	O2	120.15(8)	C14	C17	C16	120.2(2)
Ru	S2	C3	113.7(1)	C14	C17	C18	120.8(2)
Ru	S2	C4	111.1(1)	C16	C17	C18	118.9(2)
O2	S2	C3	105.3(1)	C17	C18	C19	118.7(2)
O2	S2	C4	105.3(1)	N2	C19	C12	108.7(2)
C3	S2	C4	98.9(2)	N2	C19	C18	131.3(2)
Ru	N1	C5	120.6(2)	C12	C19	C18	120.0(2)

P-2545-m8

TABLE : General displacement parameter expressions, B(Å).

	B(1,1)	B(2,2)	B(3,3)	B(1,2)	B(1,3)	B(2,3)	B _{eq}
Ru	2.009(5)	3.322(6)	2.217(5)	-0.244(6)	-0.108(4)	0.288(6)	2.537(3)
C11	2.42(2)	6.13(3)	5.24(3)	0.71(2)	-0.31(2)	1.42(3)	4.64(1)
C12	5.25(3)	4.96(3)	2.47(2)	-1.35(2)	-0.32(2)	-0.01(2)	4.27(1)
S1	2.70(2)	4.42(3)	2.84(2)	0.24(2)	0.12(2)	-0.42(2)	3.33(1)
S2	3.16(2)	4.46(3)	2.83(2)	-1.11(2)	0.08(2)	0.54(2)	3.50(1)
O1	3.68(7)	6.0(1)	2.97(6)	0.05(7)	-0.75(6)	-0.60(7)	4.27(4)
O2	5.85(9)	7.4(1)	3.30(7)	-2.64(8)	-0.17(7)	1.76(7)	5.55(5)
N1	2.70(6)	3.03(7)	2.99(7)	-0.08(6)	0.08(6)	0.12(6)	2.92(3)
N2	2.35(6)	3.13(7)	3.06(7)	-0.28(6)	-0.13(6)	0.25(6)	2.87(3)
N3	3.09(7)	2.86(7)	3.19(7)	0.01(6)	0.06(6)	0.39(6)	3.06(4)
C1	7.4(2)	4.8(1)	5.5(1)	1.2(1)	0.9(1)	-1.2(1)	5.88(8)
C2	3.39(9)	9.8(2)	4.1(1)	0.5(1)	1.38(8)	-1.0(1)	5.71(8)
C3	3.00(9)	6.1(1)	5.5(1)	-1.5(1)	-0.05(9)	-0.7(1)	4.87(6)
C4	5.9(1)	3.9(1)	6.0(1)	-0.8(1)	1.1(1)	0.6(1)	5.21(7)
C5	3.41(9)	3.18(9)	3.61(9)	-0.08(8)	0.80(7)	-0.01(8)	3.37(5)
C6	3.7(1)	4.0(1)	5.6(1)	0.50(9)	1.49(9)	-0.2(1)	4.36(6)
C7	3.6(1)	4.6(1)	6.6(2)	1.3(1)	0.3(1)	0.5(1)	4.93(6)
C8	4.1(1)	4.8(1)	4.9(1)	1.2(1)	-0.1(1)	1.2(1)	4.64(6)
C9	3.19(9)	4.5(1)	3.58(9)	0.37(9)	0.28(8)	0.89(9)	3.76(5)
C10	2.72(8)	3.30(9)	3.12(8)	-0.01(7)	-0.19(7)	0.19(8)	3.08(4)
C11	4.4(1)	3.3(1)	3.64(9)	0.06(9)	0.20(8)	0.65(9)	3.79(5)
C12	2.81(8)	2.83(8)	2.94(8)	-0.04(7)	0.23(7)	-0.17(7)	2.86(4)
C13	3.27(8)	2.91(9)	3.84(9)	-0.53(8)	0.53(7)	-0.16(8)	3.33(4)
C14	2.80(8)	3.20(9)	3.82(9)	-0.37(7)	0.32(7)	-0.81(8)	3.27(4)

P-2545-mq

45

TABLE : General displacement parameter expressions, $B(\text{\AA})$.

	$B(1,1)$	$B(2,2)$	$B(3,3)$	$B(1,2)$	$B(1,3)$	$B(2,3)$	B_{eq}
C15	3.04(9)	4.7(1)	5.6(1)	-0.83(9)	0.25(9)	-0.6(1)	4.46(6)
C16	2.95(9)	6.0(1)	4.1(1)	0.2(1)	-0.68(8)	-0.3(1)	4.41(6)
C17	2.70(8)	3.6(1)	3.05(8)	0.10(8)	-0.07(7)	-0.69(8)	3.14(4)
C18	2.75(8)	3.22(9)	2.87(8)	-0.15(7)	-0.02(7)	-0.02(7)	2.97(4)
C19	2.39(7)	2.83(8)	2.85(8)	-0.22(7)	0.13(6)	-0.16(7)	2.70(4)

The form of the anisotropic displacement parameter is:

$$\exp[-0.25\{h^2a^2B(1,1)+k^2b^2B(2,2)+l^2c^2B(3,3)+2hkabB(1,2)+2hlacB(1,3)+2klbcB(2,3)\}];$$

where a, b , and c are reciprocal lattice constants.

P-2545-m10

46

TABLE : Positional and thermal parameters of hydrogen atoms.

	X	Y	Z	B (Å ²)
	-	-	-	-----
H1	0.2689	0.5403	0.1760	7*
H2	0.1265	0.5440	0.1703	7*
H3	0.1992	0.5474	0.2287	7*
H4	0.0389	0.1866	0.2301	7*
H5	0.0615	0.3357	0.2610	7*
H6	-0.0111	0.3323	0.2027	7*
H7	-0.1100	0.1115	0.1307	6*
H8	-0.0950	0.0430	0.0730	6*
H9	-0.1190	-0.0583	0.1224	6*
H10	0.1941	-0.1700	0.0948	6*
H11	0.0623	-0.2261	0.1011	6*
H12	0.0863	-0.1247	0.0517	6*
H13	0.4166	0.1317	0.0527	4*
H14	0.6034	0.0122	0.0708	5*
H15	0.6529	-0.1019	0.1563	6*
H16	0.5099	-0.0979	0.2211	6*
H17	0.3254	0.0235	0.2008	4*
H18	0.1724	0.4787	0.0222	4*
H19	0.2399	0.6930	-0.0314	4*
H20	0.3304	0.7945	0.0043	4*
H21	0.3792	0.6871	-0.0383	4*
H22	0.5746	0.7046	0.0466	4*
H23	0.7754	0.7134	0.0885	5*
H24	0.7896	0.6782	0.1515	5*

P-2545-m11

47

TABLE : Positional and thermal parameters of hydrogen atoms.

(cont.)

	x	y	z	B (Å ²)
	-	-	-	-----
H25	0.8287	0.5623	0.1095	5*
H26	0.6770	0.3207	0.2094	5*
H27	0.7772	0.3577	0.1706	5*
H28	0.7414	0.4736	0.2133	5*
H29	0.4791	0.3183	0.1663	3*

For starred atoms $B=1.3B_{eq}$ of the bonded atom.