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# Crystal structure of borated *N*,*N*,*N'*,*N'*-tetramethyldiaminomethane

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In the title compound, {[(dimethylamino)methyl]dimethylamine}trihydridoboron,  $C_5H_{17}BN_2$ , the tetrahedral geometry of the N-C-N unit is slightly disorted. As a result of the bulky amine substituents, a wider N-C-N angle of 113.6 (1)° is observed. The bond lengths between the N atom and methyl groups are slighly elongated to 1.481 (2) and 1.482 (2) Å at the borated N atom, whereas the distances between the other N atom and its methyl groups are only 1.461 (2) and 1.462 (2) Å. The studied crystal was twinned. The twin data refinement was subsequently carried out with a scale factor of 0.263 (1). The two lattices of the twin domains were rotated by 179.84°.

Keywords: crystal structure; borane; amine; twin.

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#### 1. Related literature

For background to boranes, see: Falbe & Regitz (1999). Burg & Schlesinger (1937) reported the first borane amine complex. A feature of boranes is their metal character and pronounced Lewis acidity (Huheey *et al.*, 1995). This Lewis acidity is used to enable the  $\alpha$ -deprotonation of tertiary amines (Kessar *et al.*, 1991; Ebden *et al.*, 1995). Our group frequently uses methods to deprotonate compounds in  $\alpha$ -position (Strohmann & Gessner, 2007; Gessner & Strohmann, 2012). For crystal structures containing the borated *N*,*N*,*N'*,*N'*-tetramethyldiaminomethane motif, see: Fang *et al.* (1994); Hanic & Šubrtová (1969); Flores-Parra *et al.* (1999); Rojas-Lima *et al.* (2000). For comparison with other structures with dimethylaminoborane moiety, see: Gollas *et al.* (2013); Bera *et al.* (2011); Ramachandran *et al.* (2004); Netz *et al.* (2005). For diborated tetramethylethylenediamine, see: Chitsaz *et al.* (2001).

#### 2. Experimental

2.1. Crystal data

 $C_{5}H_{17}BN_{2}$   $M_{r} = 116.01$ Triclinic,  $P\overline{1}$  a = 6.0464 (8) Å b = 7.6987 (9) Å c = 9.5896 (11) Å  $\alpha = 69.602 (10)^{\circ}$   $\beta = 76.519 (11)^{\circ}$ 

2.2. Data collection

AgilentXcalibur, Sapphire3 diffractometer Absorption correction: multi-scan (*CrysAlis PRO*; Agilent, 2014) *T*<sub>min</sub> = 0.983, *T*<sub>max</sub> = 1.000

**2.3. Refinement**  $R[F^2 > 2\sigma(F^2)] = 0.041$  $wR(F^2) = 0.099$ S = 0.873232 reflections 90 parameters  $\gamma = 74.912 (10)^{\circ}$   $V = 398.95 (9) \text{ Å}^3$  Z = 2Mo K $\alpha$  radiation  $\mu = 0.06 \text{ mm}^{-1}$  T = 173 K $0.2 \times 0.15 \times 0.15 \text{ mm}$ 

3232 measured reflections 3232 independent reflections 1828 reflections with  $I > 2\sigma(I)$ 

H atoms treated by a mixture of independent and constrained refinement 
$$\begin{split} &\Delta\rho_{max}=0.18~e~{\rm \AA}^{-3}\\ &\Delta\rho_{min}=-0.22~e~{\rm \AA}^{-3} \end{split}$$

Data collection: *CrysAlis PRO* (Agilent, 2014); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SHELXS96* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL96* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *OLEX2* (Dolomanov *et al.*, 2009).

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Supporting information for this paper is available from the IUCr electronic archives (Reference: ZQ2234).

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### supporting information

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### Crystal structure of borated N,N,N',N'-tetramethyldiaminomethane

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#### S1. Comment

Boranes are a useful substance in today's chemistry. In 1979, Herbert C. Brown received the Nobel Prize in chemistry for his studies on these interesting compounds (Falbe & Regitz, 1999). Special about boranes is their metal character and pronounced Lewis acidity (Huheey *et al.*, 1995). This Lewis acidity is used to enable the  $\alpha$ -deprotonation of tertiary amines (Kessar *et al.*, 1991; Ebden *et al.*, 1995). Our group frequently uses methods to deprotonate compounds in  $\alpha$ -position (Gessner & Strohmann, 2007; Gessner & Strohmann, 2012). Here, BH<sub>3</sub> was added to *N*,*N*,*N'*,*N'*-tetramethyldiaminomethane (TMMDA) in order to deplete the nitrogen's +M-effect, smoothing the way for  $\alpha$ -lithiation. We isolated and structurally characterized the borated TMMDA for the first time. Lithiation of the product however was not successful.

The title compound crystallizes in the triclinic crystal system with space group P-1. The N–C–N-bonds are not equidistant. Longer N–C-bonds are observed for the borated nitrogen [N1–C1 1.4806 (17) Å, N1–C2 1.4818 (17) Å, N1–C3 1.5039 (16) Å] than for the other [N2–C3 1.4393 (17) Å, N2–C4 1.4612 (18) Å, N2–C5 1.4622 (18) Å]. Furthermore, bond angles at the nitrogen atoms differ. Due to steric hindrance of the methyl groups, angles on N2 are found to be broader [C3–N2–C4 112.82 (11)°, C3–N2–C5 112.85 (12)°, C4–N2–C5 110.31 (11)°] than the ideal sp<sup>3</sup>-angle of 109.5°. A torsion angle of 179.82 (13)° can be observed for B1–N1–C3–N2, placing B1 as far away from C3 and its hydrogen atoms as possible.

#### S1.1. Experimental

BH<sub>3</sub>-solution (11.7 mL, 1 M in thf, 11.7 mmol) was added to N,N,N',N'-tetramethyldiaminomethane (9.79 mmol, 1.00 g) at 0 °C. The reaction mixture was stirred for 3 h at room temperature after which saturated K<sub>2</sub>CO<sub>3</sub>-solution (7 mL) was added. The reaction was allowed to continue for 72 h at room temperature. After extraction with Et<sub>2</sub>O (3x10 mL), the combined extracts were dried (Na<sub>2</sub>SO<sub>4</sub>). After evaporation under reduced pressure, colorless crystals precipitated (694 mg, 5.98 mmol, 61% yield).

#### S1.2. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 1. Hydrogen atoms linked to carbon were placed and refined by using the riding model (C–H = 0.95-0.99 Å,  $U_{iso}(H) = 1.2 U_{eq}(C)$  and  $U_{iso}(H) = 1.5 U_{eq}(C)$  for terminal groups). Hydrogen atoms linked to boron were taken from difference Fourier maps.

Twin domains were found in the crystal and refined to a ratio of 0.26/0.74. The two lattices were rotated by 179.84°. HKLF5 reflection file was used for refinement.





Molecular structure of the title compound with displacement ellipsoids drawn at 50% probability level.



#### Figure 2

Molecular packing viewed along the *a* axis.

#### {[(Dimethylamino)methyl]dimethylamine}trihydridoboron

Crystal data C<sub>5</sub>H<sub>17</sub>BN<sub>2</sub>  $M_r = 116.01$ Triclinic, *P*1 a = 6.0464 (8) Å b = 7.6987 (9) Å c = 9.5896 (11) Å a = 69.602 (10)°  $\beta = 76.519$  (11)°  $\gamma = 74.912$  (10)° V = 398.95 (9) Å<sup>3</sup>

#### Data collection

AgilentXcalibur, Sapphire3 diffractometer Radiation source: Enhance (Mo) X-ray Source Graphite monochromator Z = 2 F(000) = 132  $D_x = 0.966 \text{ Mg m}^{-3}$ Mo K\alpha radiation, \lambda = 0.71073 Å Cell parameters from 1858 reflections  $\theta = 3.0-28.3^{\circ}$   $\mu = 0.06 \text{ mm}^{-1}$  T = 173 KBlock, clear light colourless  $0.2 \times 0.15 \times 0.15 \text{ mm}$ 

Detector resolution: 16.0560 pixels mm<sup>-1</sup>  $\omega$  scans Absorption correction: multi-scan (*CrysAlis PRO*; Agilent, 2014)

$T_{\min} = 0.983, \ T_{\max} = 1.000$	$\theta_{\text{max}} = 27.0^{\circ}, \ \theta_{\text{min}} = 3.1^{\circ}$
3232 measured reflections	$h = -7 \rightarrow 7$
3232 independent reflections	$k = -9 \rightarrow 9$
1828 reflections with $I > 2\sigma(I)$	$l = -12 \rightarrow 12$
Refinement	

Refinement on $F^2$	Primary atom site location: structure-invariant
Least-squares matrix: full	direct methods
$R[F^2 > 2\sigma(F^2)] = 0.041$	Hydrogen site location: mixed
$wR(F^2) = 0.099$	H atoms treated by a mixture of independent
S = 0.87	and constrained refinement
3232 reflections	$w = 1/[\sigma^2(F_o^2) + (0.0498P)^2]$
90 parameters	where $P = (F_o^2 + 2F_c^2)/3$
0 restraints	$(\Delta/\sigma)_{\rm max} = 0.001$
	$\Delta \rho_{\rm max} = 0.18 \text{ e} \text{ Å}^{-3}$
	$\Delta \rho_{\min} = -0.22 \text{ e} \text{ Å}^{-3}$

#### Special details

**Geometry**. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes. **Refinement**. Refined as a 2-component twin. 1. Twinned data refinement Scales: 0.7367 (11) 0.2633 (11) 2. Fixed  $U_{iso}$  At 1.2 times of: All C(H,H) groups At 1.5 times of: All C(H,H,H) groups 3.a Secondary CH2 refined with riding coordinates: C3(H3A,H3B) 3.b Idealized Me refined as rotating group: C2(H2A,H2B,H2C), C1(H1A,H1B,H1C), C5(H5A,H5B,H5C), C4(H4A,H4B,H4C)

	x	У	Ζ	$U_{ m iso}*/U_{ m eq}$
N1	0.17275 (18)	0.17659 (15)	0.69988 (12)	0.0182 (3)
C3	0.2336 (2)	-0.00477 (18)	0.82279 (15)	0.0214 (3)
H3A	0.2756	0.0246	0.9048	0.026*
H3B	0.0951	-0.0634	0.8648	0.026*
N2	0.42225 (19)	-0.13895 (16)	0.77307 (13)	0.0226 (3)
C2	0.1062 (3)	0.1356 (2)	0.57701 (16)	0.0274 (4)
H2A	0.2389	0.0568	0.5326	0.041*
H2B	0.0591	0.2545	0.4995	0.041*
H2C	-0.0233	0.0685	0.6177	0.041*
C1	0.3751 (2)	0.2702 (2)	0.63738 (18)	0.0293 (4)
H1A	0.4230	0.2959	0.7179	0.044*
H1B	0.3328	0.3894	0.5589	0.044*
H1C	0.5036	0.1873	0.5939	0.044*
C5	0.3510 (3)	-0.3102 (2)	0.78120 (19)	0.0340 (4)
H5A	0.2924	-0.3745	0.8861	0.051*
H5B	0.4844	-0.3946	0.7432	0.051*
H5C	0.2284	-0.2762	0.7197	0.051*
B1	-0.0405 (3)	0.3123 (3)	0.7713 (2)	0.0289 (4)
C4	0.6200 (3)	-0.1878 (2)	0.85213 (18)	0.0333 (4)
H4A	0.6693	-0.0721	0.8429	0.050*
H4B	0.7483	-0.2698	0.8077	0.050*

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters  $(Å^2)$ 

## supporting information

H4C	0.5749	-0.2542	0.9588	0.050*	
H1D	-0.083 (2)	0.4427 (19)	0.6758 (15)	0.032 (4)*	
H1E	-0.189 (2)	0.2332 (19)	0.8155 (16)	0.034 (4)*	
H1F	0.023 (2)	0.3383 (19)	0.8634 (17)	0.036 (4)*	

Atomic displacement parameters  $(Å^2)$ 

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
N1	0.0196 (6)	0.0158 (6)	0.0184 (6)	-0.0027 (5)	-0.0032 (5)	-0.0047 (5)
C3	0.0245 (8)	0.0191 (7)	0.0183 (7)	-0.0020 (6)	-0.0041 (6)	-0.0041 (6)
N2	0.0222 (7)	0.0187 (6)	0.0260 (7)	-0.0015 (5)	-0.0044 (5)	-0.0072 (5)
C2	0.0311 (9)	0.0279 (9)	0.0229 (8)	0.0000 (7)	-0.0093 (7)	-0.0081 (7)
C1	0.0259 (9)	0.0240 (8)	0.0347 (9)	-0.0071 (7)	0.0007 (7)	-0.0066 (7)
C5	0.0367 (10)	0.0230 (8)	0.0421 (10)	-0.0019 (7)	-0.0055 (8)	-0.0131 (8)
B1	0.0284 (10)	0.0236 (9)	0.0273 (10)	0.0026 (8)	-0.0018 (8)	-0.0063 (8)
C4	0.0255 (9)	0.0317 (9)	0.0404 (10)	0.0024 (7)	-0.0102 (8)	-0.0107 (8)

Geometric parameters (Å, °)

N1—C3	1.5039 (16)	C1—H1A	0.9800
N1—C2	1.4818 (17)	C1—H1B	0.9800
N1—C1	1.4806 (17)	C1—H1C	0.9800
N1—B1	1.615 (2)	С5—Н5А	0.9800
С3—НЗА	0.9900	С5—Н5В	0.9800
С3—Н3В	0.9900	С5—Н5С	0.9800
C3—N2	1.4393 (17)	B1—H1D	1.117 (13)
N2—C5	1.4622 (18)	B1—H1E	1.131 (14)
N2—C4	1.4612 (18)	B1—H1F	1.137 (14)
C2—H2A	0.9800	C4—H4A	0.9800
C2—H2B	0.9800	C4—H4B	0.9800
C2—H2C	0.9800	C4—H4C	0.9800
C3—N1—B1	108.16 (10)	N1—C1—H1C	109.5
C2—N1—C3	109.62 (10)	H1A—C1—H1B	109.5
C2—N1—B1	110.43 (11)	H1A—C1—H1C	109.5
C1—N1—C3	110.11 (10)	H1B—C1—H1C	109.5
C1—N1—C2	108.59 (10)	N2—C5—H5A	109.5
C1—N1—B1	109.93 (12)	N2—C5—H5B	109.5
N1—C3—H3A	108.8	N2—C5—H5C	109.5
N1—C3—H3B	108.8	H5A—C5—H5B	109.5
НЗА—СЗ—НЗВ	107.7	H5A—C5—H5C	109.5
N2—C3—N1	113.61 (10)	H5B—C5—H5C	109.5
N2—C3—H3A	108.8	N1—B1—H1D	105.7 (7)
N2—C3—H3B	108.8	N1—B1—H1E	106.1 (7)
C3—N2—C5	112.85 (12)	N1—B1—H1F	106.4 (7)
C3—N2—C4	112.82 (11)	H1D—B1—H1E	111.1 (10)
C4—N2—C5	110.31 (11)	H1D—B1—H1F	113.7 (10)
N1—C2—H2A	109.5	H1E—B1—H1F	113.2 (10)

	100 5		100 5
NI-C2-H2B	109.5	N2—C4—H4A	109.5
N1—C2—H2C	109.5	N2—C4—H4B	109.5
H2A—C2—H2B	109.5	N2—C4—H4C	109.5
H2A—C2—H2C	109.5	H4A—C4—H4B	109.5
H2B—C2—H2C	109.5	H4A—C4—H4C	109.5
N1—C1—H1A	109.5	H4B—C4—H4C	109.5
N1—C1—H1B	109.5		
N1—C3—N2—C5	-114.53 (13)	C1—N1—C3—N2	-60.04 (14)
N1-C3-N2-C4	119.61 (12)	B1—N1—C3—N2	179.82 (13)
C2—N1—C3—N2	59.36 (14)		