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60439 Frankfurt/Main, GermanyCorrespondence e-mail:
bolte@chemie.uni-frankfurt.de**Key indicators**

Single-crystal X-ray study

T = 173 K

Mean $\sigma(\text{C}-\text{C}) = 0.002 \text{ \AA}$

Disorder in main residue

R factor = 0.050

wR factor = 0.144

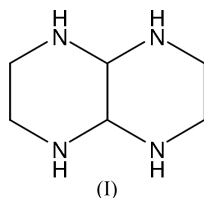
Data-to-parameter ratio = 10.0

For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.**Redetermination of *trans*-2,5,7,10-tetraaza-
bicyclo[4.4.0]decane**

The structure of the title compound, $\text{C}_6\text{H}_{14}\text{N}_4$, has previously been determined at room temperature by Bottcher, Buchkremer-Hermanns, Honle & von Schnering [*Z. Kristallogr.* (1987), **181**, 223–226], who found only one position for the amine H atom. We report here the low-temperature structure of the title compound, showing that the amine H atom is disordered over two positions. This H atom must be located either in the equatorial or in the axial position so that $\text{N}-\text{H}\cdots\text{N}$ hydrogen bonds can be formed. A mirror plane contains the bond that the two rings have in common and a twofold rotation axis runs through the mid-points of all three C–C bonds.

Comment

A perspective view of the title compound, (I), is shown in Fig. 1. Bond lengths and angles can be regarded as normal (Cambridge Structural Database, *CONQUEST* Version 1.6 plus three updates; *MOGUL* Version 1.0; Allen, 2002). The molecule possesses crystallographic C_{2h} symmetry. A mirror plane contains the bond that the two rings have in common and a twofold rotation axis runs through the mid-points of all three C–C bonds. As a result, the asymmetric unit comprises a quarter of the molecule.



The structure of (I) has already been determined at room temperature by Bottcher *et al.* (1987). These authors have

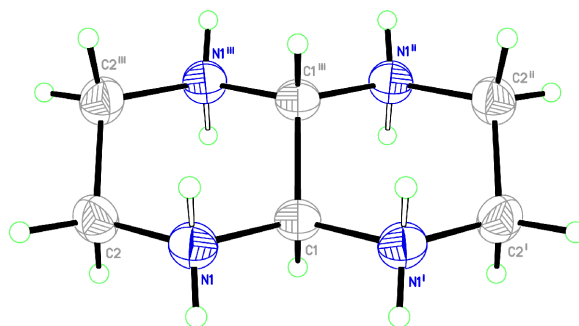


Figure 1
Perspective view of the title compound with the atom numbering; displacement ellipsoids are at the 50% probability level. Both positions are shown for each of the disordered H atoms. [Symmetry codes: (i) $x, 1 - y, z$; (ii) $-x, 1 - y, 1 - z$; (iii) $-x, y, 1 - z$.]

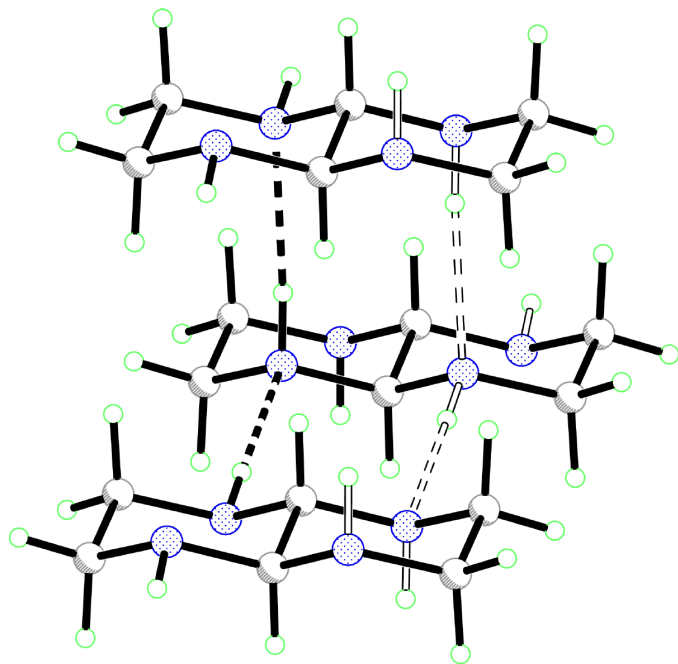


Figure 2
Hydrogen-bond (dashed lines) pattern of the title compound.

located the positions of the H atoms in a ΔF synthesis and refined the H atoms isotropically. They located the amine H atom in an equatorial position. However, they have not reported existence and nature of hydrogen bonds.

We have redetermined the structure at low temperature and have found that the amine H atom is disordered over two positions, either equatorial or axial. If this H atom were located in all molecules at the same position, a short H...H contact to a neighbouring N—H group would arise and no hydrogen bond would be possible. As a result, this H atom must occupy different positions in hydrogen-bonded molecules (Fig. 2).

Experimental

In an attempt to crystallize t -Bu₃SiBr (Wiberg *et al.*, 1997) from CH₃CN, single crystals of the title compound suitable for X-ray diffraction were obtained.

Crystal data

C ₆ H ₁₄ N ₄	$D_x = 1.282 \text{ Mg m}^{-3}$
$M_r = 142.21$	Mo $K\alpha$ radiation
Monoclinic, $C2/m$	Cell parameters from 2510 reflections
$a = 5.283 \text{ (2) \AA}$	$\theta = 4.2\text{--}27.7^\circ$
$b = 15.937 \text{ (4) \AA}$	$\mu = 0.09 \text{ mm}^{-1}$
$c = 4.6205 \text{ (18) \AA}$	$T = 173 \text{ (2) K}$
$\beta = 108.78 \text{ (3)^\circ}$	Rod, colourless
$V = 368.3 \text{ (2) \AA}^3$	$0.32 \times 0.14 \times 0.13 \text{ mm}$
$Z = 2$	

Data collection

Stoe IPDS-II two-circle diffractometer	362 reflections with $I > 2\sigma(I)$
ω scans	$R_{\text{int}} = 0.069$
Absorption correction: none	$\theta_{\text{max}} = 27.8^\circ$
2406 measured reflections	$h = -6 \rightarrow 6$
449 independent reflections	$k = -18 \rightarrow 20$
	$l = -6 \rightarrow 5$

Refinement

Refinement on F^2	All H-atom parameters refined
$R[F^2 > 2\sigma(F^2)] = 0.050$	$w = 1/[\sigma^2(F_o^2) + (0.0894P)^2]$
$wR(F^2) = 0.144$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 0.99$	$(\Delta/\sigma)_{\text{max}} < 0.001$
449 reflections	$\Delta\rho_{\text{max}} = 0.26 \text{ e \AA}^{-3}$
45 parameters	$\Delta\rho_{\text{min}} = -0.27 \text{ e \AA}^{-3}$

Table 1

Selected geometric parameters (\AA , $^\circ$).

N1—C1	1.4591 (13)	N1—C2	1.4669 (14)
C1—N1—C2	109.91 (10)	N1—C1—C1 ⁱⁱ	110.25 (10)
N1 ⁱ —C1—N1	108.77 (13)	N1—C2—C2 ⁱⁱⁱ	111.18 (9)

Symmetry codes: (i) $x, 1 - y, z$; (ii) $-x, 1 - y, 1 - z$; (iii) $-x, y, 1 - z$.

Table 2

Hydrogen-bonding geometry (\AA , $^\circ$).

$D\text{—}H\cdots A$	$D\text{—}H$	$H\cdots A$	$D\cdots A$	$D\text{—}H\cdots A$
N1—H1 ⁱ ...N1 ^{iv}	0.80 (4)	2.36 (4)	3.146 (3)	167 (3)
N1—H1 ⁱⁱ ...N1 ^v	0.93 (3)	2.27 (3)	3.182 (3)	168 (2)

Symmetry codes: (iv) $1 - x, y, 2 - z$; (v) $-x, y, 2 - z$.

All H atoms were located in a difference map and refined isotropically. The two sites for the disordered H atoms bonded to N1 refined to relative occupancies of 0.54 (4) and 0.46 (4) for the equatorial and axial H atoms, respectively.

Data collection: *X-AREA* (Stoe & Cie, 2001); cell refinement: *X-AREA*; data reduction: *X-AREA*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* in *SHELXTL-Plus* (Sheldrick, 1991); software used to prepare material for publication: *SHELXL97* and *PLATON* (Spek, 2003).

References

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