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2,6-Dibromo-4,4,8,8-tetraethylpyrazabole

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Abstract. $(BrC_3N_2H_2)_2B_2(C_2H_5)_4$, monoclinic, $P2_1/a$, $a = 15.623 (14), b = 12.420 (10), c = 9.927 (6) \text{ Å}, \beta =$ 92.48 (3)°, Z = 4, $D_c = 1.476$, $D_m = 1.49$ g cm⁻³ (flotation), μ (Mo K α) = 45.1 cm⁻¹. Each B atom is coordinated to two ethyl groups and a N of each pyrazolyl group. The two pyrazolyl rings are coplanar within experimental limits.

Introduction. Crystals of 2,6-dibromo-4,4,8,8-tetraethylpyrazabole were received from S. Trofimenko in a form suitable for crystal-structure analysis. Preliminary Weissenberg and precession photographs established the space group $P2_1/a$ (h0l, h = 2n; 0k0, k = 2n). A cube-like crystal of maximum dimension 0.4 mm was mounted on a Picker automated diffractometer for data collection. Cell dimensions were refined during the alignment procedure. Data were collected with a θ -2 θ

Table 1. Positional parameters ($\times 10^3$, for Br $\times 10^4$) for 2,6-dibromo-4,4,8,8-tetraethylpyrazabole

The estimated sendered deviations are since in momentum in

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Table 2. Bond distances (Å) and angles (°)

units of the last significant figure of the coordinate.				Br(1) - C(2)	1.92 (6)	C(1)-C(2)-C(3)	113 (4)
				Br(2) - C(5)	1.97 (5)	C(2)-C(3)-N(2)	106 (4)
			_	C(5) - C(6)	1.31 (6)	C(3)-N(2)-N(1)	111 (3)
	х	y	Z	C(5) - C(4)	1.45 (6)	N(2) - N(1) - C(1)	109 (3)
Br(1)	4347 (4)	1825 (5)	-6467 (5)	C(2) - C(1)	1.47 (7)	N(1)-C(1)-C(2)	100 (4)
Br(2)	928 (4)	638 (5)	2252 (5)	C(2) - C(3)	1.33 (7)	C(4) - C(5) - C(6)	122 (4)
N(1)	342 (2)	136 (3)	-278 (3)	C(4) - N(3)	1.48 (6)	C(5)-C(6)-N(4)	101 (4)
N(2)	266 (2)	158 (3)	-350 (3)	C(6) - N(4)	1.42 (5)	C(6) - N(4) - N(3)	110 (3)
N(3)	188 (2)	125 (3)	-140 (3)	C(3) - N(2)	1.35 (5)	N(4) - N(3) - C(4)	113 (3)
N(4)	260 (2)	103 (3)	-68 (3)	C(1) - N(1)	1.42 (6)	N(3)-C(4)-C(5)	94 (4)
C(1)	411 (3)	142 (4)	-364 (4)	N(1) - N(2)	1.38 (5)	N(2)-B(1)-N(3)	97 (3)
C(2)	365 (3)	165 (4)	-493 (5)	N(3) - N(4)	1.34 (5)	N(2)-B(1)-C(7)	103 (4)
C(3)	281 (3)	178 (4)	-481 (4)	N(1) - B(2)	1.72 (6)	N(2)-B(1)-C(9)	109 (4)
C(4)	111 (3)	118 (4)	-59 (5)	N(2) - B(1)	1.67 (7)	N(3)-B(1)-C(7)	101 (3)
C(5)	159 (3)	89 (4)	64 (4)	N(4) - B(2)	1.69 (7)	N(3)-B(1)-C(9)	114 (4)
C(6)	242 (3)	74 (4)	66 (4)	N(3) - B(1)	1.67 (6)	C(7)-B(1)-C(9)	128 (5)
C(7)	166 (3)	316 (4)	-256 (4)	B(1) - C(7)	1.76 (9)	N(1)-B(2)-N(4)	95 (3)
C(8)	161 (4)	385 (4)	-379 (5)	B(1) - C(9)	1.47 (8)	N(1)-B(2)-C(11)	114 (4)
C(9)	106 (3)	144 (5)	-378 (4)	B(2) - C(11)	1.50 (8)	N(1)-B(2)-C(13)	102 (3)
C(10)	112 (4)	-6 (4)	-390 (4)	B(2) - C(13)	1.74 (9)	N(4)-B(2)-C(11)	114 (4)
C(11)	421 (3)	177 (5)	-36 (4)	C(7)–C(8)	1.49 (7)	N(4) - B(2) - C(13)	104 (4)
C(12)	384 (4)	291 (5)	-30 (5)	C(9) - C(10)	1.50 (8)	C(11)-B(2)-C(13)	123 (5)
C(13)	386 (3)	-37 (4)	-126 (4)	C(11)–C(12)	1.52 (9)	N(1) - N(2) - B(1)	129 (3)
C(14)	329 (4)	-115 (4)	-207 (5)	C(13)–C(14)	1.53 (7)	N(4)-N(3)-B(1)	133 (4)
B(1)	168 (4)	177 (5)	-294 (5)			N(2)-N(1)-B(2)	133 (3)
B(2)	364 (4)	100 (5)	-112 (6)			N(3)-N(4)-B(2)	132 (3)



Fig. 1. A projection of 2,6-dibromo-4,4,8,8-tetraethylpyrazabole onto the xz plane.

and polarization factors. Absorption corrections were not made.

The structure was solved with conventional heavyatom techniques. Full-matrix refinement of the positional parameters and isotropic temperature factors for all 22 non-hydrogen atoms gave R = 17.2%. A difference Fourier map revealed anisotropic motion about the Br atoms and ethyl groups. The temperature factors for these ten atoms were converted to their anisotropic equivalents, and six cycles of refinement of all positional and thermal parameters gave a final R of 6.2%. On the final cycle the maximum ratio of a parameter to its corresponding standard deviation was 0.05. A final difference map showed no peak higher than 0.36 e Å⁻³. Unit weights were used throughout.

Atomic coordinates are given in Table 1, bond lengths and angles in Table 2.* Fig. 1 shows a projection of the structure onto the xz plane of the cell.

Discussion. Di- and tripyrazolyl borates have been studied crystallographically in various coordinations to heavy atoms (Trofimenko, 1971, 1972). In the tripyrazolyl borate structures the pyrazolyl rings have been shown to display various modes of skew about the boron-heavy-metal axis (Avitabile, Ganis & Nemiroff, 1971; Holt, Holt & Watson, 1973; Holt, Holt, Cavalito & Watson, 1976; Churchill, Gold & Maw, 1970; Cotton, Frenz & Shaver, 1973; Davies & Payne, 1974; Holt & Holt, 1973). Dipyrazolyl borate has been studied in a situation where the structure was constrained as a result of hydrogen bonding (Kosky, Ganis

Table 3. Equations of the least-squares planes, and
deviations (Å) of atoms from the planes

Plane 1: Br(1), N(1), N(2), C(1), C(2), C(3)

 $\begin{array}{c} 0.169x + 0.121y + 0.193z = 0.169 \\ |Br(1) 0.004, N(1) - 0.004, N(2) - 0.006, C(1) 0.017, \\ C(2) - 0.031, C(3) 0.021 | \end{array}$

Plane 2: Br(2), N(3), N(4), C(4), C(5), C(6)

 $\begin{array}{c} 0.158x + 0.120y + 0.244z = 0.145\\ [Br(2)\ 0.008,\ N(3)\ -0.002,\ N(4)\ 0.021,\ C(4)\ -0.010,\\ C(5)\ 0.007,\ C(6)\ -0.024]\end{array}$

& Avitabile, 1971). The reported structure was undertaken to provide an observation of a pyrazabole in normal unstrained circumstances.

Coordination about each B atom is distorted tetrahedral [97 (3)–128 (5)°] with each B atom being bonded to a N of each of two pyrazolyl groups and also to two ethyl groups. The B–N distances are longer than those found in other pyrazolyl borate structures [1.67 (6)– 1.72 (6) Å] as opposed to the reported range of 1.50– 1.60 Å. Each B has one significantly shorter bond (1.47–1.50 Å) and one longer bond (1.74–1.76 Å) to C. The bromopyrazolyl rings are planar (maximum deviation from plane: 0.03 Å) (Table 3). The angle between the planes of the two rings is 177°. Calculations show that B(1) and B(2) lie 0.15, 0.20 Å and -0.08, 0.04 Å respectively from the planes of the rings.

Calculations were performed on an XDS Sigma 7 with the XRAY system (1976).

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^{*} Lists of structure factors and thermal parameters have been deposited with the British Library Lending Division as Supplementary Publication No. SUP 32491 (4 pp.). Copies may be obtained through The Executive Secretary, International Union of Crystallography, 13 White Friars, Chester CH1 INZ, England.

2,6-DIBROMO-4,4,8,8-TETRAETHYLPYRAZABOLE

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$18(17 \rightarrow 16)$ abeo-Yohimbine Acetate Methanol Solvate

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N(1)

C(2)

C(3) N(4)

C(5) C(6)

C(7)

C(8)

C(9) C(10)

C(11)

C(12)

C(13)

C(14)

C(15) C(16)

C(17)

C(18)

C(19)

C(20) C(21)

O(22)

C(23)

O(24)

C(25)

C(26) O(27) O(28)

C(29)

O(30)

C(31)

Abstract. $C_{23}H_{28}N_2O_4$. CH₄O, monoclinic, $P2_1$, a = 6.734 (5), b = 14.465 (9), c = 12.114 (10) Å, $\beta = 105.2$ (2)°, Z = 2, final R value 0.07. The C/D ring junction is *trans*-quinolizidine; the D/E ring junction is *trans*. The six-membered rings C and D have half-chair and chair conformations, the five-membered ring E has a half-chair conformation.

Introduction. In a study of ring closure by mercury oxidation, demethylcorynantheine led mostly to a new derivative of the $18(17 \rightarrow 16)abeo$ -yohimbine type (Djakouré, Jarreau & Goutarel, 1975). Recently, the reaction mechanism has been established (Boivin, Païs & Goutarel, 1976). The present X-ray structure determination was undertaken to ascertain the stereo-chemistry at C(20).

Data were collected from a crystal $0.3 \times 0.3 \times 0.6$ mm on a Philips PW 1100 diffractometer with graphitemonochromated Cu Karadiation ($\lambda = 1.5418$ Å), for θ between 3 and 68°. 2266 independent reflexions were measured, of which 1972 were considered observed with $I > 3\sigma(I)$, $\sigma(I)$ being derived from counting statistics.

The structure was solved by straightforward application of MULTAN (Germain, Main & Woolfson,

Table 1. Final coordinates $(\times 10^4)$ and their e.s.d.'s

x	у	z
7051 (5)	506 (0)	2066 (3)
5242 (6)	952 (3)	1495 (4)
5071 (6)	1965 (3)	1319 (4)
2823 (5)	2205 (3)	1022 (3)
1661 (8)	1610(4)	97 (4)
1599 (7)	610 (4)	479 (4)
3709 (7)	326 (3)	1162 (4)
4553 (7)	-565 (3)	1532 (4)
3710 (9)	-1458 (4)	1485 (5)
4995 (11)	-2189(4)	1964 (5)
7056 (11)	-2034 (4)	2477 (5)
7923 (8)	-1154 (4)	2546 (4)
6652 (7)	-430(3)	2086 (4)
6248 (6)	2540(3)	2335 (4)
5969 (6)	3544 (3)	2032 (4)
6896 (6)	4314 (3)	2901 (4)
9196 (7)	4416 (4)	2987 (4)
5636 (7)	5178 (3)	2372 (4)
3802 (7)	4842 (3)	1401 (4)
3675 (6)	3805 (3)	1636 (3)
2539 (7)	3189 (3)	681 (4)
10114 (5)	5098 (3)	3844 (3)
10842 (7)	5876 (4)	3504 (5)
10665 (6)	6071 (3)	2531 (4)
11889 (11)	6476 (5)	4479 (7)
6696 (8)	4035 (3)	4068 (4)
7942 (6)	3618(3)	4765 (3)
4874 (5)	4303 (3)	4243 (3)
4452 (12)	3957 (6)	5301 (5)
11023 (5)	1604 (3)	2838 (4)
12360 (11)	12.56 (9)	3789 (7)

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