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Key indicators

Single-crystal X-ray study T = 133 KMean $\sigma(C-C) = 0.002 \text{ Å}$ R factor = 0.040 wR factor = 0.117 Data-to-parameter ratio = 20.3

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. In the title compound, $C_{18}H_{30}N_6$, the central triazine ring is planar, with C–N bond lengths 1.3415 (14)–1.3504 (13) Å, N–C–N angles 125.58 (10)–125.97 (9)° and C–N–C angles 113.82 (9)–114.33 (9)°.

2,4,6-Tripiperidino-1,3,5-triazine

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Comment

Cyclotrimerization of organic nitriles is a well known method for the industrial synthesis of 1,3,5-triazines (Martin et al., 1978). These triazines have wide applications in pharmaceutical and other industries and they are also used as monomers for the synthesis of heat-resistant and thermostable polymers. Generally this cyclotrimerization is catalysed by hydrochloric acid (Grundmann et al., 1952) but various other catalysts (Michurin et al., 1969; Zilberman et al., 1969, 1971) can also be used. The formation of 2,4,6-tripiperidino-1,3,5-triazine, (I), has been reported (Shishulina et al., 1999) together with the crystal structure of its 2:2:1 solvated adduct with 2.6-bis(trichloromethyl)-1,4,3,5-oxathiadiazine 4,4-dioxide and benzene. The formation of (I) in this molecular adduct resulted from the reaction of piperidinonitrile with 6-piperidino-2-trichloroacetylimino-4-trichloromethyl-1,2,3,5-oxathiadiazine 2oxide in benzene. However, (I) could not be isolated as a pure compound from the adduct. We report here the synthesis of (I) by metal-carbonyl-catalysed cyclotrimerization of piperidinonitrile, together with its crystal structure.



The crystal structure of (I) (Fig. 1) can be compared with that of the triazine in the 2:2:1 adduct solvate (Shishulina *et al.*, 1999) and also with the simplest 1,3,5-triazine, namely (HCN)₃ (Smith & Rae, 1978).

The six-membered triazine ring in (I) is essentially planar (0.008 Å mean deviation from the least-squares plane). The piperidine rings are rotated out of this plane, as defined by torsion angles of 14.03 (13) (N1-C2-N6-C18), -5.34 (15) (N3-C1-N4-C8) and -4.87 (15)° (N2-C3-N5-C9). The CN bond distances (see Table 1) are similar to those reported for the same triazine in the solvate complex. The triazine (HCN)₃ shows one independent CN bond distance in the same range [1.338 (4) Å; Smith & Rae, 1978]. The exocyclic CN bond distances for the triazine (I) [1.3535 (15)-1.3713 (13) Å] are significantly larger than those in the solvate

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Figure 1

The molecular structure of the title compound. Displacement ellipsoids are drawn at the 30% probability level. H atoms have been omitted for clarity.

complex (1.325–1.333 Å). The narrow CNC angles (ca 114°) and the wide NCN angles ($ca \ 126^{\circ}$) are similar to the CNC $[114.8 (10)^{\circ}]$ and the NCN $[125.2 (10)^{\circ}]$ angles of (HCN)₃.

Experimental

Metal carbonyls $M(CO)_6$ (M = Cr, Mo, W) effectively catalyse the cyclotrimerization of N-cyanopiperidine to (I), in good to excellent yields, in toluene under heating conditions. The best results are obtained in the reaction catalysed by W(CO)₆. Metal carbonyls are recovered in almost quantitative yield in all reactions, and the triazine is obtained in pure form and in good to excellent yields. A solution of piperidinonitrile (1.10 g, 10 mmol) and $W(CO)_6$ (5% mol) in toluene (5 ml) was heated for 3-5 d. All reactions were monitored by thinlayer chromatography. The resulting pale-brown solution was concentrated and the product was separated by low-temperature column chromatography (SiO₂, 271 K, 10% diethyl ether in petroleum ether). Evaporation of the solvents of the second fraction and recrystallization from dichloromethane at 253 K yielded the title compound as colourless crystals. Analysis calculated for C₁₈H₃₀N₆: C 65.42, H 9.15, N 25.43%; found: C 65.40, H 9.20, N 25.39%. M.p. 492 K, cf. 423 K for the solvate complex. ¹H NMR (CDCl₃): δ 1.43– 1.49 (m, 12H, piperidine ring), 1.52–1.57 (m, 12H, piperidine ring), 3.62–3.64 (m, 6H, piperidine ring); ¹³C NMR: δ 25.1 (s, piperidine ring), 25.8 (s, piperidine ring), 44.1 (s, piperidine ring), 165.4 [s, triazine ring; cf. Tornieporth-Oetting et al. (1992)]; MS (70 eV, EI) m/ z 330 (molecular ion peak; 100%) (M^+); IR (KBr, cm⁻¹): ν (C=N) 1533 (cf. 1725 and 1625 cm⁻¹ for the solvate complex; Shishulina et al., 1999).

Crystal data

$C_{18}H_{30}N_6$	Z = 2
$M_r = 330.48$	$D_x = 1.228 \text{ Mg m}^{-3}$
Triclinic, $P\overline{1}$	Mo $K\alpha$ radiation
a = 8.3527 (8) Å	Cell parameters from 5270
b = 9.8192 (10) Å	reflections
c = 12.0219 (12) Å	$\theta = 2 - 30^{\circ}$
$\alpha = 98.377 \ (3)^{\circ}$	$\mu = 0.08 \text{ mm}^{-1}$
$\beta = 101.455 \ (3)^{\circ}$	T = 133 (2) K
$\gamma = 108.361 \ (3)^{\circ}$	Cut tablet, colourless
$V = 893.88 (15) \text{ Å}^3$	$0.38 \times 0.22 \times 0.13 \text{ mm}$

Data collection

Bruker SMART 1000 CCD	$R_{\rm int} = 0.031$
diffractometer	$\theta_{\rm max} = 28.3^{\circ}$
ω and φ scans	$h = -11 \rightarrow 11$
12 279 measured reflections	$k = -13 \rightarrow 13$
4407 independent reflections	$l = -16 \rightarrow 16$
3407 reflections with $I > 2\sigma(I)$	
Refinement	

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.065P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.040$	+ 0.0581P]
$wR(F^2) = 0.117$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.05	$(\Delta/\sigma)_{\rm max} < 0.001$
4407 reflections	$\Delta \rho_{\rm max} = 0.27 \ {\rm e} \ {\rm \AA}^{-3}$
217 parameters	$\Delta \rho_{\rm min} = -0.17 {\rm e} {\rm \AA}^{-3}$
H-atom parameters constrained	

Table 1

Selected geometric parameters (Å, °).

C1-N3	1.3422 (14)	C2-N6	1.3713 (13)
C1-N1	1.3504 (13)	C3-N3	1.3465 (13)
C1-N4	1.3614 (13)	C3-N2	1.3482 (13)
C2-N2	1.3415 (14)	C3-N5	1.3535 (15)
C2-N1	1.3480 (13)		
N3-C1-N1	125.97 (9)	C2-N1-C1	113.82 (9)
N2-C2-N1	125.97 (9)	C2-N2-C3	114.33 (9)
N3-C3-N2	125.58 (10)	C1-N3-C3	114.27 (9)
N3-C1-N4-C4	-178.34 (9)	N1-C2-N6-C18	14.03 (13)
N2-C3-N5-C9	176.35 (9)		

H atoms were positioned at calculated positions with C-H = 0.99 Å and $U_{iso}(H) = 1.2U_{eq}(C)$, and refined using a riding model.

Data collection: SMART (Bruker, 1998); cell refinement: SAINT (Bruker, 1998); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1990); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: XP (Siemens, 1994); software used to prepare material for publication: SHELXL97.

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