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5,5',6,6'-Tetramethyl-3,3'-bi-1,2,4-triazine

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Key indicators: single-crystal X-ray study; T = 293 K; mean σ (C–C) = 0.003 Å; R factor = 0.064; wR factor = 0.272; data-to-parameter ratio = 13.1.

In the title compound, $C_{10}H_{12}N_6$, the two 5,6-dimethyl-1,2,4-triazine halves of the molecule are related by a centre of symmetry. The two triazine rings are coplanar to within a maximum deviation of 0.013 (2) Å from the mean plane of the ring atoms. In the crystal, molecules form layers parallel to the (100) crystallographic plane. Adjacent layers are held together via a $C-H\cdots\pi$ interaction involving molecules related by an a-glide plane.

Related literature

For background information, see: Branowska & Rykowski (2002); Branowska (2003); Boger & Weinrab (1987); Pabst *et al.* (1998). For the synthesis, see: Dedichen (1936, 1937). For a related structure, see: Breu & Range (1993).

Experimental

Crystal data

 $C_{10}H_{12}N_6$ V = 1100.59 (18) Å³ $M_r = 216.26$ Z = 4 Orthorhombic, Pbca Cu Kα radiation a = 8.1167 (7) Å $\mu = 0.71 \text{ mm}^{-1}$ b = 10.6662 (12) Å T = 293 K c = 12.7127 (11) Å $0.20 \times 0.20 \times 0.10 \text{ mm}$

Data collection

Kuma KM4 four-circle diffractometer Absorption correction: ψ scan (North *et al.*, 1968) $T_{\rm min} = 0.830, T_{\rm max} = 0.929$ 1637 measured reflections

1205 independent reflections 910 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.100$ 2 standard reflections every 100 reflections intensity decay: 1.3%

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.064$ $wR(F^2) = 0.272$ S = 1.161205 reflections

92 parameters All H-atom parameters refined $\Delta \rho_{\rm max} = 0.29$ e Å $^{-3}$ $\Delta \rho_{\rm min} = -0.24$ e Å $^{-3}$

Table 1Hydrogen-bond geometry (Å, °).

CgA is the centroid of the triazine ring.

$D-H\cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D-\mathrm{H}\cdots A$
$C51-H511\cdots CgA^{i}$	1.09 (5)	2.96 (4)	3.616 (3)	119 (3)
Symmetry code: (i) $x -$	$\frac{1}{2}$, y , $-z + \frac{3}{2}$.			_

Data collection: KM4B8 (Gałdecki et al., 1996); cell refinement: KM4B8; data reduction: DATAPROC (Gałdecki et al., 1995); program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: SHELXL97 and WinGX (Farrugia, 1999).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: FY2014).

References

Boger, D. L. & Weinrab, S. N. (1987). Hetero Diels-Alder Methodology in Organic Synthesis. Organic Chemistry: A Series of Monographs, Vol. 47, pp. 323–335. San Diego: Academic Press.

Branowska, D. (2003). Synthesis, pp. 2096–2100.

Branowska, D. & Rykowski, A. (2002). Synlett, pp. 1892–1894.

Breu, J. & Range, K.-J. (1993). Acta Cryst. C49, 1541-1543.

Dedichen, G. (1936). Avh. Nor. Vidensk. Akad. Oslo Mat. Naturvidensk. Kl. 5,

Dedichen, G. (1937). Chem. Abstr 31, 4985-4988.

Farrugia, L. J. (1997). J. Appl. Cryst. 30, 565.

Farrugia, L. J. (1999). J. Appl. Cryst. 32, 837-838.

Gałdecki, Z., Kowalski, A., Kucharczyk, D. & Uszyński, L. (1996). KM4B8. Kuma Diffraction, Wrocław, Poland.

Gałdecki, Z., Kowalski, A. & Uszyński, L. (1995). DATAPROC. Kuma Diffraction, Wrocław, Poland.

North, A. C. T., Phillips, D. C. & Mathews, F. S. (1968). *Acta Cryst.* A24, 351–350

Pabst, R. G., Schmid, K. & Sauer, J. (1998). *Tetrahedron Lett.* **39**, 6691–6695. Sheldrick, G. M. (2008). *Acta Cryst.* A**64**, 112–122.

supporting information

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5,5',6,6'-Tetramethyl-3,3'-bi-1,2,4-triazine

Ewa Wolińska, Zbigniew Karczmarzyk, Andrzej Rykowski and Waldemar Wysocki

S1. Comment

In the course of our research we widely explored the synthesis of cycloalkeno[c]fused 2,2'-bipyridines using Diels—Alder reactions of 5,5'-bi-1,2,4-triazine derivatives as dienes (Branowska & Rykowski, 2002; Branowska, 2003). However, when we turned our attention to the synthesis of 5,5',6,6'-tetrasubstituted-2,2'-bipyridines, these dienes did not appear useful. Considering the mechanism of the Diels—Alder reaction of 5,5'-bi-1,2,4-triazine, it is clear that to obtain 5,5',6,6'-tetrasubstituted-2,2'-bipyridines, the substituents in positions 5 and 5' of the product have to originate from an unsymmetrical dienofile. Unfortunately, application of such a dienofile can lead to a mixture of 5,5',6,6'- and 3,3',6,6'-tetrasubstituted-2,2'-bipyridines (Boger & Weinrab, 1987). To solve the problem with selectivity, we envisaged that 3,3'-bi-1,2,4-triazines with substituents in 5 and 5' positions can be structurally ideal diene partners in the Diels—Alder synthesis of 5,5',6,6'-tetrasubstituted-2,2'-bipyridines (Pabst *et al.*, 1998). The title compound 5,5',6,6'-tetramethyl-3,3'-bi-1,2,4-triazine was synthesized and its X-ray structure was determined as a part of this research.

The two 5,6-dimethyl-1,2,4-triazine parts of the molecule (I) are related by a crystallographic center of symmetry and possess the *trans* conformation, with the triazine rings being coplanar to within a 0.013 (2) Å maximum deviation from the mean plane. The geometry and conformation of (I) are very similar to those observed in the related structure of 5,5',6,6'-tetraphenyl-3,3'-bi-1,2,4-triazine (Breu & Range, 1993).

In the crystal structure, the molecules of (I) form molecular layers parallel to the (100) crystallographic plane (Fig. 2), with the molecular mean planes being inclined to this plane at an angle of 34.8 (5)°. The layers are held together via C—H··· π interaction involving the C51—H151 atoms of the methyl group and the triazine ring from the molecule related by an a-glide plane.

S2. Experimental

The title compound, (I), was prepared by the condensation of oxalhydrazidine with 2,3-butanedione according to the procedure of Dedichen (1936, 1937). Crystals suitable for X-ray diffraction analysis were grown by slow evaporation of a benzene solution.

S3. Refinement

All H atoms were located in a difference Fourier map and their coordinates were refined freely with isotropic displacement parameters $U_{iso}(H) = 1.5 U_{eq}(C)$. Refined C—H distances were in the range 0.96 (5)–1.09 (5) Å.

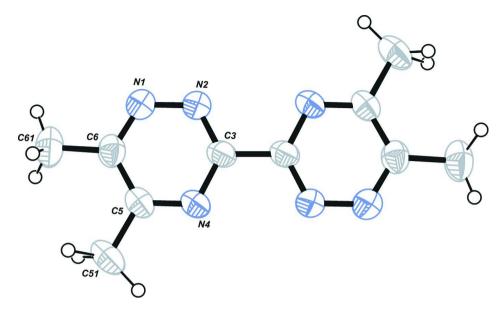


Figure 1The molecular structure of (I), with atom labels and 50% probability displacement ellipsoids for non-H atoms.

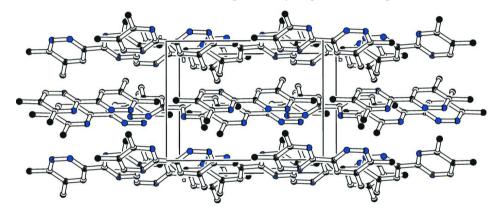


Figure 2

A view of the molecular packing in (I). H atoms are omitted for clarity.

3-(5,6-dimethyl-1,2,4-triazin-3-yl)-5,6-dimethyl-1,2,4-triazine

Crystal data

 $C_{10}H_{12}N_6$ $M_r = 216.26$ Orthorhombic, PbcaHall symbol: -P 2ac 2ab a = 8.1167 (7) Å b = 10.6662 (12) Å c = 12.7127 (11) Å V = 1100.59 (18) Å³ Z = 4F(000) = 456 $D_{\rm x}=1.305~{
m Mg~m^{-3}}$ Melting point = 441–442 K Cu $K\alpha$ radiation, $\lambda=1.54178~{
m \AA}$ Cell parameters from 25 reflections $\theta=11.5$ –22.4° $\mu=0.71~{
m mm^{-1}}$ $T=293~{
m K}$ Prism, yellow $0.20\times0.20\times0.10~{
m mm}$

Data collection

Kuma KM4 four-circle diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

 $\omega/2\theta$ scans

Absorption correction: ψ scan (North *et al.*, 1968)

 $T_{\min} = 0.830, T_{\max} = 0.929$

1637 measured reflections

Refinement

Refinement on F^2

Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.064$

 $wR(F^2) = 0.272$

S = 1.16

1205 reflections 92 parameters

0 restraints

Primary atom site location: structure-invariant

direct methods

1205 independent reflections 910 reflections with $I > 2\sigma(I)$

 $R_{\rm int} = 0.100$

 $\theta_{\text{max}} = 80.2^{\circ}, \, \theta_{\text{min}} = 7.0^{\circ}$

 $h = -1 \rightarrow 10$

 $k = -1 \rightarrow 13$

 $l = -1 \rightarrow 16$

2 standard reflections every 100 reflections

intensity decay: 1.3%

Secondary atom site location: difference Fourier

map

Hydrogen site location: difference Fourier map

All H-atom parameters refined

 $w = 1/[\sigma^2(F_0^2) + (0.2P)^2]$

where $P = (F_0^2 + 2F_c^2)/3$

 $(\Delta/\sigma)_{\text{max}} \leq 0.001$

 $\Delta \rho_{\rm max} = 0.29 {\rm e \ \AA^{-3}}$

 $\Delta \rho_{\min} = -0.24 \text{ e Å}^{-3}$

Extinction correction: *SHELXL97* (Sheldrick, 2008), Fc*=kFc[1+0.001xFc $^2\lambda^3$ /sin(2 θ)]^{-1/4}

Extinction coefficient: 0.032 (7)

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R-factor wR and goodness of fit S are based on F^2 , conventional R-factors R are based on F, with F set to zero for negative F^2 . The threshold expression of $F^2 > 2 \operatorname{sigma}(F^2)$ is used only for calculating R-factors(gt) etc. and is not relevant to the choice of reflections for refinement. R-factors based on F^2 are statistically about twice as large as those based on F, and R- factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\hat{A}^2)

	x	у	Z	$U_{ m iso}$ */ $U_{ m eq}$
N1	0.4275 (4)	0.2701 (2)	0.50737 (19)	0.0750 (8)
N2	0.4248 (3)	0.1532(2)	0.4693 (2)	0.0733 (8)
N4	0.58882 (19)	0.08158 (18)	0.61103 (13)	0.0491 (6)
C3	0.5036(2)	0.0648 (2)	0.52195 (15)	0.0488 (7)
C5	0.5959(2)	0.1973 (2)	0.64669 (16)	0.0501 (7)
C6	0.5112 (3)	0.2946 (2)	0.59397 (18)	0.0541 (7)
C51	0.6953 (4)	0.2213 (3)	0.7429 (2)	0.0750 (9)
H511	0.771 (6)	0.142 (5)	0.767(3)	0.113*
H512	0.762 (7)	0.297 (4)	0.732(3)	0.113*
H513	0.621 (6)	0.241 (6)	0.803(3)	0.113*
C61	0.5097 (4)	0.4263 (3)	0.6326 (3)	0.0733 (9)
H611	0.482 (6)	0.431 (4)	0.709 (4)	0.110*
H612	0.427 (5)	0.476 (6)	0.599 (4)	0.110*
H613	0.617 (5)	0.462 (5)	0.618(3)	0.110*

supporting information

Atomic displacement parameters (\mathring{A}^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N1	0.1042 (18)	0.0492 (13)	0.0717 (14)	0.0018 (11)	-0.0267 (12)	0.0010 (10)
N2	0.1037 (18)	0.0482 (12)	0.0680 (13)	0.0020 (10)	-0.0359(11)	-0.0007(9)
N4	0.0483 (9)	0.0560 (12)	0.0431 (9)	-0.0048(6)	-0.0053(6)	-0.0018(6)
C3	0.0512 (10)	0.0529 (13)	0.0424 (10)	-0.0044(8)	-0.0061(7)	0.0006 (8)
C5	0.0484 (10)	0.0574 (13)	0.0445 (10)	-0.0097(7)	0.0017 (7)	-0.0064(8)
C6	0.0610 (12)	0.0471 (12)	0.0543 (11)	-0.0093 (8)	0.0071 (8)	-0.0029(8)
C51	0.0804 (16)	0.0834 (19)	0.0613 (14)	-0.0116 (15)	-0.0183 (12)	-0.0177 (13)
C61	0.092(2)	0.0515 (15)	0.0760 (18)	-0.0118 (12)	0.0119 (14)	-0.0092 (12)

Geometric parameters (Å, °)

N1—C6	1.320 (3)	C6—C61	1.488 (3)
N1—N2	1.337 (3)	C51—H511	1.09 (5)
N2—C3	1.322 (3)	C51—H512	0.98 (5)
N4—C5	1.316 (3)	C51—H513	1.00 (5)
N4—C3	1.339 (2)	C61—H611	1.00 (5)
C3—C3 ⁱ	1.492 (4)	C61—H612	0.96 (5)
C5—C6	1.414 (4)	C61—H613	0.97 (5)
C5—C51	1.488 (3)		
C6—N1—N2	119.7 (2)	C5—C51—H511	114 (2)
C3—N2—N1	118.3 (2)	C5—C51—H512	109 (3)
C5—N4—C3	116.05 (19)	H511—C51—H512	112 (4)
N2—C3—N4	125.6 (2)	C5—C51—H513	110 (3)
N2—C3—C3 ⁱ	116.9 (2)	H511—C51—H513	107 (4)
N4—C3—C3 ⁱ	117.4 (2)	H512—C51—H513	106 (4)
N4—C5—C6	120.27 (19)	C6—C61—H611	112 (3)
N4—C5—C51	118.0 (2)	C6—C61—H612	113 (3)
C6—C5—C51	121.8 (2)	H611—C61—H612	105 (4)
N1—C6—C5	120.0(2)	C6—C61—H613	108 (3)
N1—C6—C61	117.3 (2)	H611—C61—H613	111 (4)
C5—C6—C61	122.7 (2)	H612—C61—H613	109 (4)
C6—N1—N2—C3	1.8 (4)	N2—N1—C6—C5	-0.8 (4)
N1—N2—C3—N4	-0.7(4)	N2—N1—C6—C61	180.0(2)
N1—N2—C3—C3 ⁱ	179.2 (2)	N4—C5—C6—N1	-1.4(3)
C5—N4—C3—N2	-1.4(3)	C51—C5—C6—N1	178.4 (3)
C5—N4—C3—C3 ⁱ	178.6 (2)	N4—C5—C6—C61	177.8 (2)
C3—N4—C5—C6	2.4(3)	C51—C5—C6—C61	-2.4(3)
C3—N4—C5—C51	-177.4(2)		

Symmetry code: (i) -x+1, -y, -z+1.

supporting information

Hydrogen-bond geometry (Å, °)

CgA is the centroid of the triazine ring.

D— H ··· A	D—H	$H\cdots A$	D··· A	<i>D</i> —H··· <i>A</i>
C51—H511···CgA ⁱⁱ	1.09 (5)	2.96 (4)	3.616 (3)	119 (3)

Symmetry code: (ii) x-1/2, y, -z+3/2.