



Synthesis and crystal structures of 4,4'-methylenebis(2,6-diethylaniline) and 4,4'-methylenebis(3-chloro-2,6-diethylaniline)

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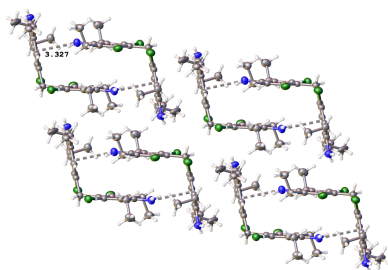
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Keywords: crystal structure; methylenedianiline; hydrogen bonding; synthesis; disorder.**CCDC references:** 2391630; 2391631**Supporting information:** this article has supporting information at journals.iucr.org/e

The title compounds 4,4'-methylenebis(2,6-diethylaniline) ($C_{21}H_{30}N_2$, **1**) and 4,4'-methylenebis(3-chloro-2,6-diethylaniline) ($C_{21}H_{28}Cl_2N_2$, **2**) are of significant interest as curing agents for a wide range of resins and as building blocks for sterically demanding compounds in the synthesis of ligands for catalysis. This paper describes their synthesis and the preparation of single crystals, with their structures determined through single-crystal X-ray analysis. The presence of the chlorine substituent slightly affects the twist angle between the two aromatic components. The molecules of compound **1** form a network structure through intermolecular $N-H \cdots N$ bonds and $C-H \cdots \pi$ interactions, while in the crystal structure of compound **2**, the molecules are assembled solely through $N-H \cdots \pi$ interactions. Consequently, despite their chemical similarity, it is the precise structural data that enables us to explain their differing reactivity and opens up the possibility of evaluating steric properties for the development of new materials and ligands.

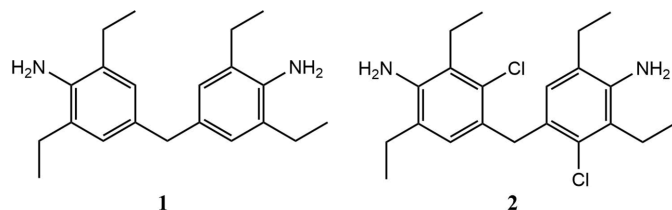
1. Chemical context

Aromatic diamines are widely utilized as hardeners for polyurethanes (Ueda *et al.*, 2017), epoxy resins (Yu *et al.*, 2020; Costa *et al.*, 2005), cyanate ester (Bauer & Bauer, 2001) and phthalonitriles (Bulgakov *et al.*, 2021). The reactivity of amines is primarily determined by their nucleophilicity, which depends on the electronic structure and geometry of the molecules. In the production of castings and composite materials, the gelation time of the binder is a critical factor: the longer the gelation time at low temperatures, the more time is available for the impregnation of the reinforcing filler or the moulding of products. Hardeners with low activity at room temperature also facilitate the manufacture of prepregs with a long shelf life. To decrease the reaction rate, the nucleophilicity of the diamine can be diminished by introducing electron-withdrawing substituents. An example of such a modification is 4,4'-diaminodiphenylsulfone (DDS), which exhibits significantly lower activity than 4,4'-methylenedianiline (MDA) (Kong & Park, 2003). Another strategy for reducing the activity of amines involves the introduction of sterically bulky groups in the *ortho* position, as seen in 4,4'-methylenebis(2,6-diethylaniline) (MDEA). An even more pronounced effect can be achieved by additionally incorporating chlorine into the 3-position of the aromatic ring, specifically in 4,4'-methylenebis(3-chloro-2,6-diethylaniline) (MCDEA). In terms of reactivity with epoxy monomers, these compounds follow the order: MDEA > DDS > MCDEA (Lahlali *et al.*, 2005, 2006). Similarly, in reactions with isocyanates, the order of reactivity



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is MDA > MDEA > MCDEA (Voelker *et al.*, 1988). Notably, a molecule carrying a methyl group in the *meta* position instead of chlorine, exhibits approximately the same activity as MDEA. Consequently, this raises the question of which factor – steric or electronic – primarily influences the activity of hardeners. Therefore, studying the molecular structure is particularly intriguing for comparing the steric hindrance of the amine in MDEA and MCDEA.



Another potential application of *ortho*-substituted anilines is the synthesis of N-heterocyclic carbenes, including polymeric structures (Peng *et al.*, 2018). The properties of carbenes as ligands are typically assessed based on the electronic and steric characteristics of the substituents on the nitrogen atoms (Cavallo *et al.*, 2005). The steric hindrance is quantitatively described by the parameter $V_{bur}\%$, which represents the volume of a sphere occupied by the ligand (Gomez-Suarez *et al.*, 2017). Molecular structure data obtained from X-ray crystallography can be utilized to calculate $V_{bur}\%$ and more accurately predict the catalytic properties of molecules through machine learning (Escayola *et al.*, 2024). Consequently, the parameters outlined in this paper will be valuable for various scientists seeking to understand the reactivity of curing agents and predict the properties of structures based on the compounds discussed.

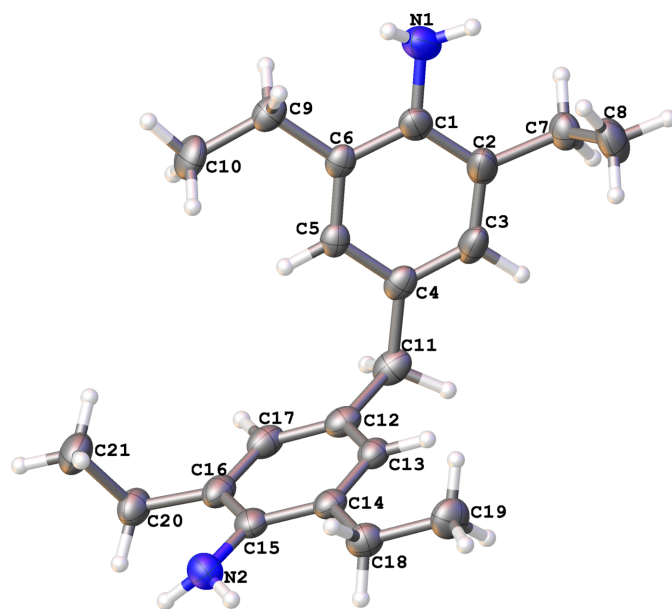


Figure 1
The molecular structure of **1**, with displacement ellipsoids drawn at the 50% probability level. The minor occupancy components are omitted for clarity.

2. Structural commentary

The molecular structures of the title compounds **1** and **2**, which differ in the presence of an additional chlorine substituent at *meta* position to amine, are illustrated in Figs. 1 and 2, respectively. In both cases, the organic molecules occupy general positions. Depending on the presence of chlorine substituents, the ring systems are twisted to a greater or lesser extent. In molecule **1**, the dihedral angle between the two aromatic parts is as large as $64.13(6)^\circ$, while the corresponding angle in molecule **2** is $39.59(8)^\circ$. The dihedral angle between those parts approximately equal for both structures [$80.24(4)$ and $77.72(6)^\circ$ for the **1** and **2**, respectively]. In molecule **2**, the chlorine atom in the C12–C17 ring is disordered with an occupancy ratio of 0.920(2):0.08(2).

In the unsubstituted molecule **1** the ethyl groups (C9–C10 and C18–C19) are nearly co-planar with their phenyl rings and the ethyl group connected with the C2 atom is disordered. The major components (73.6%) for C7–C8 and ethyl group C20–C21 are directed almost orthogonal [$88.6(4)^\circ$ and $80.9(2)^\circ$, respectively] to the plane of the phenyl rings, while the minor components are slightly inclined with torsion angle $C1-C2-C7'-C8' = 153.5(12)^\circ$.

The steric hindrance exerted by the chlorine atom in molecule **2** causes the ethyl groups to rotate out of the plane of the phenyl ring. The ethyl groups connected with the C12–C17

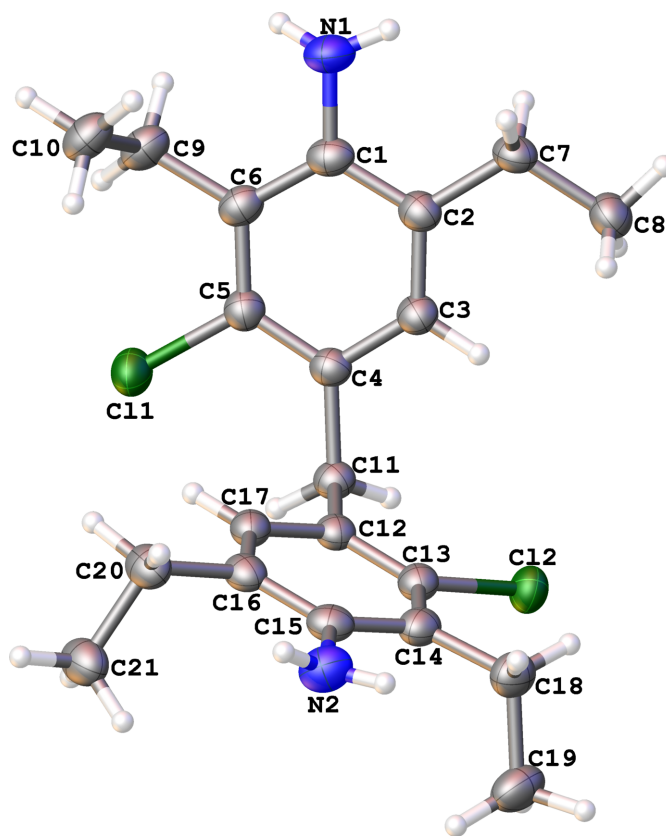


Figure 2
The molecular structure of **2**, with displacement ellipsoids drawn at the 50% probability level. The minor occupancy components are omitted for clarity.

Table 1

Hydrogen-bond geometry (Å, °) for **1**.

Cg_1 is the centroid of the C1–C6 ring.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N2-H2A\cdots N1^i$	0.88	2.62	3.472 (2)	165
$C17-H17\cdots Cg1^{ii}$	0.95	2.92	3.8568 (17)	171

Symmetry codes: (i) $x, y + 1, z$; (ii) $-x + 2, -y + 1, -z + 1$.

phenyl ring are parallel to each other with the torsion angle C19–C18–C20–C21 being $0.60 (17)^\circ$ while the torsion angle C8–C7–C9–C10 between other ethyl fragments is $116.7 (3)^\circ$.

3. Supramolecular features

In the crystal structure of **1**, the molecules form infinite chains extending along [010] via $N2-H2A\cdots N1$ hydrogen bonds [$N2\cdots N1 = 3.472 (2) \text{ \AA}$; Fig. 3, Table 1]. Additionally, the molecules are grafted together in a herringbone-like manner by $C-H\cdots\pi$ interactions [$3.8568 (17) \text{ \AA}$] involving the phenyl H17 atom and the centroids of the C1–C6 phenyl rings of adjacent molecules.

In contrast, these types of interaction are absent in the crystal structure of **2**. The molecules form centrosymmetric dimers by $N-H\cdots\pi$ interactions [$3.327 (2) \text{ \AA}$] between $N2-H2A$ and the C1–C6 ring centroid (Table 2, Fig. 4). Cohesion of the packing is provided by $C7-H7B\cdots\pi$ [$3.579 (2) \text{ \AA}$] and $C20-H20B\cdots\pi$ [$3.434 (2) \text{ \AA}$] interactions in the [100] direction and weak van der Waals interactions between the dimers.

4. Database survey

A search of the Cambridge Structural Database (CSD, version 5.45 updated to November 2023; Groom *et al.*, 2016) for **1** and **2** revealed that these structures had not been published previously. However, a similar uncharged structure without substituents in the ring, 4,4'-methylenebis(aniline), had been described in two independent experiments [CSD refcodes CEHCOH (Bel'skii *et al.*, 1983) and CEHCOH01 (Gibson *et al.*, 2010)]. Despite the similarity in molecular structures and

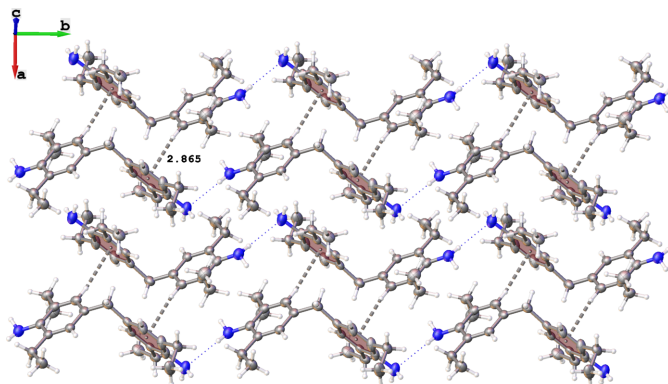


Figure 3
Fragment of the crystal packing of **1**.

Table 2

Hydrogen-bond geometry (Å, °) for **2**.

Cg_1 and Cg_2 are the centroids of the C1–C6 and C12–C17 rings, respectively.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N2-H2A\cdots Cg1^i$	0.88	2.63	3.327 (2)	137
$C7-H7B\cdots Cg2^{ii}$	0.99	2.83	3.579 (2)	133
$C20-H20B\cdots Cg2^i$	0.99	2.81	3.434 (2)	132

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

number of analogues interactions between these and the current study, such a herringbone-like packing motif does not occur in the previously published structures that can be explained by the rotational degree of freedom.

For the closest analogue with methyl substituents in the phenyl ring, 4,4'-methylenebis(2,6-dimethylaniline), a search resulted in 14 hits with CSD refcodes AWAYAZ–AWAYAZ13 (Bhattacharya & Saha, 2011). According to X-ray investigations, this structure was discovered in two polymorphic modifications in different space groups, $C2/c$ and $P\bar{1}$; additionally, it was established in the original work that both polymorphs crystallize simultaneously from the solution in the presence of additional reagents. In general, the weakness of intermolecular interactions is proved by the discovery of weak $N-H\cdots N$ interactions in one polymorphic form and the absence of such interactions in other structures.

The comparison of structures **1** and **2** and previously published analogues revealed that the twist angle between the two aromatic parts for the structure **1** [$115.87 (6)^\circ$] lies within the characteristic range [$44.21 (6)$ – $134.35 (5)^\circ$], while the corresponding angle for the structure of **2** is smaller at $39.59 (8)^\circ$.

5. Synthesis and crystallization

The title compounds were prepared as follows:

4,4'-Methylenebis(2,6-diethylaniline) (**1**)

A mixture of 2,6-diethylaniline (14.92 g, 0.1 mol), paraformaldehyde (0.15 g, 0.05 mol) and 36% hydrochloric acid (8.6 mL, 0.1 mol) in water (100 mL) in a round-bottom flask was heated to 353 K for 3 h in an oil bath under argon. The

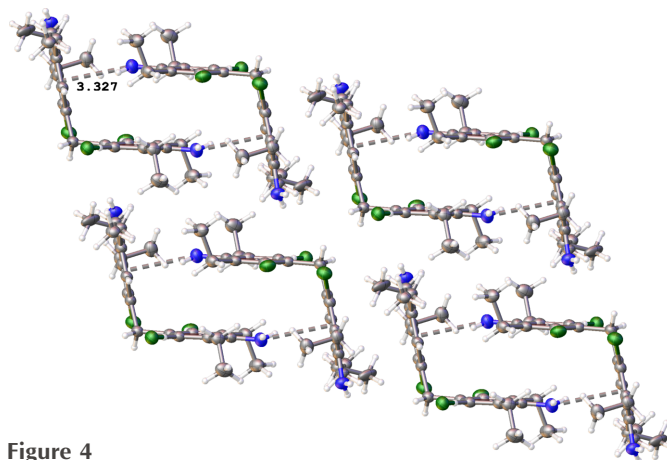


Figure 4
Fragment of the crystal packing of **2**.

Table 3
Experimental details.

	1	2
Crystal data		
Chemical formula	C ₂₁ H ₃₀ N ₂	C ₂₁ H ₂₈ Cl ₂ N ₂
<i>M_r</i>	310.47	379.35
Crystal system, space group	Monoclinic, <i>P</i> 2 ₁ / <i>n</i>	Triclinic, <i>P</i> $\bar{1}$
Temperature (K)	150	100
<i>a</i> , <i>b</i> , <i>c</i> (Å)	8.9895 (3), 11.8589 (3), 17.6765 (5)	8.993 (3), 9.6540 (13), 12.216 (4)
α , β , γ (°)	90, 102.4188 (11), 90	69.352 (11), 77.257 (12), 87.670 (8)
<i>V</i> (Å ³)	1840.32 (9)	967.2 (5)
<i>Z</i>	4	2
Radiation type	Cu <i>K</i> α	Synchrotron, $\lambda = 0.75268$ Å
μ (mm ⁻¹)	0.49	0.40
Crystal size (mm)	0.32 × 0.21 × 0.13	0.19 × 0.12 × 0.05
Data collection		
Diffractometer	Bruker D8 Venture	Rayonix SX165 CCD
Absorption correction	Multi-scan (<i>SADABS</i> ; Krause <i>et al.</i> , 2015)	Empirical (using intensity measurements) [<i>XDS</i> (Kabsch, 2010)]
<i>T</i> _{min} , <i>T</i> _{max}	0.639, 0.753	0.001, 1.000
No. of measured, independent and observed [<i>I</i> > 2 σ (<i>I</i>)] reflections	22931, 3455, 3212	17607, 4844, 3879
<i>R</i> _{int}	0.035	0.053
(<i>sin</i> θ / λ) _{max} (Å ⁻¹)	0.608	0.675
Refinement		
<i>R</i> [<i>F</i> ² > 2 σ (<i>F</i> ²)], <i>wR</i> (<i>F</i> ²), <i>S</i>	0.050, 0.135, 1.03	0.051, 0.147, 1.05
No. of reflections	3455	4844
No. of parameters	236	243
No. of restraints	1	0
H-atom treatment	H-atom parameters constrained	H-atom parameters constrained
$\Delta\rho_{\text{max}}$, $\Delta\rho_{\text{min}}$ (e Å ⁻³)	0.51, -0.33	0.64, -0.56

Computer programs: *APEX3* v2017.3.0 (Bruker, 2016), *Marccd* (Doyle, 2011), *XDS* (Kabsch, 2010), *SAINT* (Bruker, 2016), *SHELXT2014/5* (Sheldrick, 2015a), *SHELXL2019/3* (Sheldrick, 2015b), *OLEX2* 1.3 (Dolomanov *et al.*, 2009).

reaction mixture was cooled to room temperature and sodium hydroxide (4.40 g, 0.11 mol) was added. The precipitate was filtered and dried at 343 K in an oven in air for 12 h. Yield 14.52 g (94%). Single crystals suitable for X-ray analysis were grown by slow cooling (363–303 K, 5 K h⁻¹) of a solution of the substance in a DMSO/water (80:20, *v:v*) mixture.

¹H NMR (600 MHz, DMSO-*d*₆) δ 6.64 (*s*, 4H), 4.29 (*s*, 4H), 3.56 (*s*, 2H), 2.42 (*q*, *J* = 7.5 Hz, 8H), 1.09 (*t*, *J* = 7.5 Hz, 12H).

4,4'-Methylenebis(3-chloro-2,6-diethylaniline) (2)

A mixture of 3-chloro-2,6-diethylaniline (18.38 g, 0.1 mol), paraformaldehyde (0.15 g, 0.05 mol) and 36% hydrochloric acid (8.6 mL, 0.1 mol) in water (130 mL) in round-bottom flask was heated to 353 K for 3 h in an oil bath under argon. The reaction mixture was cooled to room temperature and sodium hydroxide (4.40 g, 0.11 mol) was added. The precipitate was filtered and dried at 343 K in an oven in air for 12 h. Yield 17.55 g (93%). Single crystals suitable for X-ray analysis were grown by slow evaporation of the solvent from a solution of the substance in toluene.

¹H NMR (600 MHz, DMSO-*d*₆) δ 6.51 (*s*, 2H), 4.70 (*s*, 4H), 3.84 (*s*, 2H), 2.72 (*q*, *J* = 7.4 Hz, 4H), 2.37 (*q*, *J* = 7.5 Hz, 4H), 1.08–0.99 (*m*, 12H).

6. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 3. X-ray diffraction studies for (2)

were carried out on the Belok'beamline (Svetogorov *et al.*, 2020) of the National Research Center "Kurchatov Institute" (Moscow, Russian Federation) using a Rayonix SX165 CCD detector. All hydrogen atoms in the structures of **1** and **2** were placed in calculated positions and refined using a riding model [*U*_{iso}(H) = 1.2–1.5*U*_{eq}(parent atom)]. In the structure of **2**, one chlorine atom was found to be disordered over two positions with a refined occupancy ratio of 0.920 (2):0.080 (2). The ethyl group connected to the C2 atom was found to be disordered with occupancy ratios of 0.736 (11)/0.264 (11) in the structure of **1**. A SADI instruction was used to restrain the C7–C2 and C7'–C2 bonds in **1**.

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Synthesis and crystal structures of 4,4'-methylenebis(2,6-diethylaniline) and 4,4'-methylenebis(3-chloro-2,6-diethylaniline)

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Computing details

4,4'-(Methanediyl)bis(2,6-diethylaniline) (1)

Crystal data

$C_{21}H_{30}N_2$

$M_r = 310.47$

Monoclinic, $P2_1/n$

$a = 8.9895$ (3) Å

$b = 11.8589$ (3) Å

$c = 17.6765$ (5) Å

$\beta = 102.4188$ (11)°

$V = 1840.32$ (9) Å³

$Z = 4$

$F(000) = 680$

$D_x = 1.121$ Mg m⁻³

Cu $K\alpha$ radiation, $\lambda = 1.54178$ Å

Cell parameters from 9874 reflections

$\theta = 4.5$ – 69.6 °

$\mu = 0.49$ mm⁻¹

$T = 150$ K

Prism, colourless

$0.32 \times 0.21 \times 0.13$ mm

Data collection

Bruker D8 Venture
diffractometer

Radiation source: microfocus sealed X-ray tube

φ and ω scans

Absorption correction: multi-scan
(SADABS; Krause *et al.*, 2015)

$T_{\min} = 0.639$, $T_{\max} = 0.753$

22931 measured reflections

3455 independent reflections

3212 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.035$

$\theta_{\max} = 69.6$ °, $\theta_{\min} = 4.5$ °

$h = -10 \rightarrow 10$

$k = -14 \rightarrow 14$

$l = -21 \rightarrow 21$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.050$

$wR(F^2) = 0.135$

$S = 1.03$

3455 reflections

236 parameters

1 restraint

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0668P)^2 + 0.8039P]$

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

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

$\Delta\rho_{\max} = 0.51$ e Å⁻³

$\Delta\rho_{\min} = -0.33$ e Å⁻³

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.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
N1	0.50801 (18)	0.22887 (12)	0.44320 (9)	0.0495 (4)	
H1A	0.444518	0.203843	0.401577	0.073 (7)*	
H1B	0.454778	0.243338	0.478419	0.042 (5)*	
N2	0.76330 (16)	1.02220 (12)	0.52830 (8)	0.0434 (3)	
H2A	0.702084	1.069094	0.497865	0.052*	
H2B	0.841893	1.056893	0.557216	0.052*	
C1	0.59366 (18)	0.32315 (12)	0.42982 (9)	0.0371 (3)	
C2	0.6281 (2)	0.34245 (13)	0.35674 (9)	0.0450 (4)	
C3	0.7176 (2)	0.43384 (13)	0.34743 (8)	0.0422 (4)	
H3	0.738899	0.447362	0.297852	0.051*	
C4	0.77715 (16)	0.50617 (12)	0.40722 (8)	0.0322 (3)	
C5	0.74348 (15)	0.48530 (12)	0.47942 (8)	0.0288 (3)	
H5	0.782842	0.534865	0.521151	0.035*	
C6	0.65472 (15)	0.39491 (11)	0.49237 (8)	0.0293 (3)	
C7	0.5780 (5)	0.2673 (2)	0.28671 (14)	0.0371 (8)	0.736 (11)
H7A	0.572874	0.188130	0.303718	0.045*	0.736 (11)
H7B	0.653349	0.271536	0.253388	0.045*	0.736 (11)
C7'	0.493 (2)	0.2769 (8)	0.2921 (5)	0.056 (4)	0.264 (11)
H7'A	0.522766	0.196888	0.288739	0.068*	0.264 (11)
H7'B	0.396857	0.278633	0.310534	0.068*	0.264 (11)
C8	0.4216 (5)	0.3041 (3)	0.2405 (3)	0.0527 (11)	0.736 (11)
H8A	0.391557	0.255842	0.194700	0.079*	0.736 (11)
H8B	0.426673	0.382675	0.224074	0.079*	0.736 (11)
H8C	0.346445	0.297264	0.272966	0.079*	0.736 (11)
C8'	0.4671 (12)	0.3274 (9)	0.2140 (6)	0.047 (2)	0.264 (11)
H8'A	0.403094	0.276881	0.176766	0.070*	0.264 (11)
H8'B	0.564933	0.338586	0.199170	0.070*	0.264 (11)
H8'C	0.415683	0.400219	0.214345	0.070*	0.264 (11)
C9	0.62289 (17)	0.36943 (13)	0.57138 (8)	0.0355 (3)	
H9A	0.659524	0.292198	0.586402	0.043*	
H9B	0.511194	0.369814	0.566853	0.043*	
C10	0.6940 (2)	0.44930 (16)	0.63601 (9)	0.0455 (4)	
H10A	0.659453	0.526391	0.622011	0.068*	
H10B	0.805186	0.445821	0.643944	0.068*	
H10C	0.663280	0.427331	0.683880	0.068*	
C11	0.88289 (18)	0.60062 (13)	0.39550 (9)	0.0391 (4)	
H11A	0.988886	0.577452	0.418229	0.047*	
H11B	0.874584	0.611130	0.339183	0.047*	
C12	0.85318 (16)	0.71251 (12)	0.43033 (8)	0.0339 (3)	

C13	0.73112 (15)	0.78002 (12)	0.39480 (8)	0.0314 (3)
H13	0.666330	0.754365	0.348255	0.038*
C14	0.70101 (16)	0.88340 (12)	0.42509 (8)	0.0317 (3)
C15	0.79580 (17)	0.92047 (13)	0.49506 (8)	0.0342 (3)
C16	0.91879 (17)	0.85307 (14)	0.53197 (8)	0.0373 (3)
C17	0.94554 (16)	0.75095 (14)	0.49856 (9)	0.0374 (3)
H17	1.029684	0.706187	0.523247	0.045*
C18	0.57200 (17)	0.95838 (13)	0.38505 (9)	0.0385 (3)
H18A	0.615497	1.031988	0.374583	0.046*
H18B	0.504104	0.972245	0.421362	0.046*
C19	0.47564 (19)	0.91440 (15)	0.30954 (10)	0.0464 (4)
H19A	0.539667	0.904872	0.271582	0.070*
H19B	0.394034	0.968336	0.289597	0.070*
H19C	0.431041	0.841598	0.318726	0.070*
C20	1.0202 (2)	0.88664 (17)	0.60861 (9)	0.0477 (4)
H20A	1.119249	0.847402	0.614784	0.057*
H20B	1.039622	0.968796	0.608464	0.057*
C21	0.9492 (3)	0.8578 (2)	0.67641 (11)	0.0614 (5)
H21A	0.852117	0.897807	0.671087	0.092*
H21B	1.018192	0.880692	0.724765	0.092*
H21C	0.931458	0.776368	0.677302	0.092*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N1	0.0589 (9)	0.0356 (7)	0.0489 (8)	−0.0072 (6)	0.0002 (7)	0.0029 (6)
N2	0.0480 (8)	0.0408 (7)	0.0411 (7)	−0.0051 (6)	0.0087 (6)	−0.0037 (6)
C1	0.0444 (8)	0.0276 (7)	0.0353 (8)	0.0053 (6)	0.0000 (6)	0.0041 (6)
C2	0.0686 (11)	0.0323 (8)	0.0296 (8)	0.0050 (7)	0.0005 (7)	−0.0005 (6)
C3	0.0659 (10)	0.0359 (8)	0.0257 (7)	0.0120 (7)	0.0121 (7)	0.0035 (6)
C4	0.0348 (7)	0.0317 (7)	0.0309 (7)	0.0117 (6)	0.0088 (6)	0.0056 (5)
C5	0.0284 (6)	0.0307 (7)	0.0264 (6)	0.0086 (5)	0.0037 (5)	0.0006 (5)
C6	0.0293 (7)	0.0300 (7)	0.0277 (7)	0.0101 (5)	0.0044 (5)	0.0036 (5)
C7	0.0406 (18)	0.0373 (12)	0.0329 (12)	0.0057 (11)	0.0069 (10)	−0.0055 (8)
C7'	0.089 (10)	0.039 (4)	0.037 (4)	−0.018 (5)	0.003 (5)	−0.003 (3)
C8	0.056 (2)	0.0475 (17)	0.045 (2)	0.0010 (14)	−0.0112 (16)	−0.0127 (15)
C8'	0.050 (5)	0.049 (4)	0.041 (5)	−0.003 (4)	0.007 (3)	−0.007 (4)
C9	0.0344 (7)	0.0403 (8)	0.0328 (7)	0.0075 (6)	0.0099 (6)	0.0074 (6)
C10	0.0533 (9)	0.0575 (10)	0.0291 (7)	0.0086 (8)	0.0163 (7)	−0.0005 (7)
C11	0.0380 (8)	0.0419 (8)	0.0403 (8)	0.0090 (6)	0.0151 (6)	0.0086 (6)
C12	0.0308 (7)	0.0361 (8)	0.0367 (7)	0.0000 (6)	0.0114 (6)	0.0091 (6)
C13	0.0295 (7)	0.0355 (7)	0.0292 (7)	−0.0026 (5)	0.0065 (5)	0.0048 (5)
C14	0.0302 (7)	0.0339 (7)	0.0316 (7)	−0.0024 (6)	0.0081 (5)	0.0067 (6)
C15	0.0368 (7)	0.0351 (7)	0.0325 (7)	−0.0064 (6)	0.0112 (6)	0.0036 (6)
C16	0.0334 (7)	0.0455 (9)	0.0322 (7)	−0.0071 (6)	0.0053 (6)	0.0088 (6)
C17	0.0304 (7)	0.0434 (8)	0.0373 (8)	0.0004 (6)	0.0050 (6)	0.0128 (6)
C18	0.0373 (8)	0.0365 (8)	0.0411 (8)	0.0042 (6)	0.0073 (6)	0.0052 (6)
C19	0.0415 (8)	0.0498 (9)	0.0434 (9)	0.0082 (7)	−0.0010 (7)	0.0077 (7)

C20	0.0445 (9)	0.0587 (10)	0.0363 (8)	-0.0091 (8)	0.0009 (7)	0.0062 (7)
C21	0.0733 (13)	0.0726 (13)	0.0371 (9)	-0.0096 (11)	0.0091 (9)	0.0061 (9)

Geometric parameters (Å, °)

N1—C1	1.406 (2)	C9—H9A	0.9900
N1—H1A	0.8801	C9—H9B	0.9900
N1—H1B	0.8799	C10—H10A	0.9800
N2—C15	1.400 (2)	C10—H10B	0.9800
N2—H2A	0.8799	C10—H10C	0.9800
N2—H2B	0.8799	C11—C12	1.511 (2)
C1—C6	1.410 (2)	C11—H11A	0.9900
C1—C2	1.410 (2)	C11—H11B	0.9900
C2—C3	1.380 (2)	C12—C17	1.386 (2)
C2—C7	1.513 (3)	C12—C13	1.394 (2)
C2—C7'	1.668 (9)	C13—C14	1.388 (2)
C3—C4	1.377 (2)	C13—H13	0.9500
C3—H3	0.9500	C14—C15	1.413 (2)
C4—C5	1.3957 (19)	C14—C18	1.511 (2)
C4—C11	1.512 (2)	C15—C16	1.406 (2)
C5—C6	1.384 (2)	C16—C17	1.391 (2)
C5—H5	0.9500	C16—C20	1.514 (2)
C6—C9	1.5151 (19)	C17—H17	0.9500
C7—C8	1.530 (5)	C18—C19	1.519 (2)
C7—H7A	0.9900	C18—H18A	0.9900
C7—H7B	0.9900	C18—H18B	0.9900
C7'—C8'	1.476 (15)	C19—H19A	0.9800
C7'—H7'A	0.9900	C19—H19B	0.9800
C7'—H7'B	0.9900	C19—H19C	0.9800
C8—H8A	0.9800	C20—C21	1.513 (2)
C8—H8B	0.9800	C20—H20A	0.9900
C8—H8C	0.9800	C20—H20B	0.9900
C8'—H8'A	0.9800	C21—H21A	0.9800
C8'—H8'B	0.9800	C21—H21B	0.9800
C8'—H8'C	0.9800	C21—H21C	0.9800
C9—C10	1.516 (2)		
C1—N1—H1A	114.1	H9A—C9—H9B	107.4
C1—N1—H1B	111.4	C9—C10—H10A	109.5
H1A—N1—H1B	107.7	C9—C10—H10B	109.5
C15—N2—H2A	116.6	H10A—C10—H10B	109.5
C15—N2—H2B	115.6	C9—C10—H10C	109.5
H2A—N2—H2B	112.2	H10A—C10—H10C	109.5
N1—C1—C6	118.89 (14)	H10B—C10—H10C	109.5
N1—C1—C2	121.28 (14)	C12—C11—C4	115.05 (12)
C6—C1—C2	119.68 (14)	C12—C11—H11A	108.5
C3—C2—C1	119.02 (14)	C4—C11—H11A	108.5
C3—C2—C7	116.33 (19)	C12—C11—H11B	108.5

C1—C2—C7	124.6 (2)	C4—C11—H11B	108.5
C3—C2—C7'	129.8 (3)	H11A—C11—H11B	107.5
C1—C2—C7'	106.3 (5)	C17—C12—C13	117.99 (14)
C4—C3—C2	122.54 (14)	C17—C12—C11	121.21 (13)
C4—C3—H3	118.7	C13—C12—C11	120.79 (13)
C2—C3—H3	118.7	C14—C13—C12	122.25 (13)
C3—C4—C5	117.81 (14)	C14—C13—H13	118.9
C3—C4—C11	120.56 (13)	C12—C13—H13	118.9
C5—C4—C11	121.54 (13)	C13—C14—C15	118.80 (13)
C6—C5—C4	122.31 (13)	C13—C14—C18	122.21 (13)
C6—C5—H5	118.8	C15—C14—C18	118.99 (13)
C4—C5—H5	118.8	N2—C15—C16	120.62 (14)
C5—C6—C1	118.61 (13)	N2—C15—C14	119.65 (14)
C5—C6—C9	122.46 (13)	C16—C15—C14	119.70 (14)
C1—C6—C9	118.93 (13)	C17—C16—C15	119.23 (14)
C2—C7—C8	110.3 (3)	C17—C16—C20	118.89 (15)
C2—C7—H7A	109.6	C15—C16—C20	121.84 (15)
C8—C7—H7A	109.6	C12—C17—C16	122.01 (14)
C2—C7—H7B	109.6	C12—C17—H17	119.0
C8—C7—H7B	109.6	C16—C17—H17	119.0
H7A—C7—H7B	108.1	C14—C18—C19	116.41 (13)
C8'—C7'—C2	113.4 (8)	C14—C18—H18A	108.2
C8'—C7'—H7'A	108.9	C19—C18—H18A	108.2
C2—C7'—H7'A	108.9	C14—C18—H18B	108.2
C8'—C7'—H7'B	108.9	C19—C18—H18B	108.2
C2—C7'—H7'B	108.9	H18A—C18—H18B	107.3
H7'A—C7'—H7'B	107.7	C18—C19—H19A	109.5
C7—C8—H8A	109.5	C18—C19—H19B	109.5
C7—C8—H8B	109.5	H19A—C19—H19B	109.5
H8A—C8—H8B	109.5	C18—C19—H19C	109.5
C7—C8—H8C	109.5	H19A—C19—H19C	109.5
H8A—C8—H8C	109.5	H19B—C19—H19C	109.5
H8B—C8—H8C	109.5	C21—C20—C16	111.99 (15)
C7'—C8'—H8'A	109.5	C21—C20—H20A	109.2
C7'—C8'—H8'B	109.5	C16—C20—H20A	109.2
H8'A—C8'—H8'B	109.5	C21—C20—H20B	109.2
C7'—C8'—H8'C	109.5	C16—C20—H20B	109.2
H8'A—C8'—H8'C	109.5	H20A—C20—H20B	107.9
H8'B—C8'—H8'C	109.5	C20—C21—H21A	109.5
C6—C9—C10	115.95 (13)	C20—C21—H21B	109.5
C6—C9—H9A	108.3	H21A—C21—H21B	109.5
C10—C9—H9A	108.3	C20—C21—H21C	109.5
C6—C9—H9B	108.3	H21A—C21—H21C	109.5
C10—C9—H9B	108.3	H21B—C21—H21C	109.5
N1—C1—C2—C3	-177.50 (15)	C3—C4—C11—C12	137.06 (15)
C6—C1—C2—C3	-2.0 (2)	C5—C4—C11—C12	-46.59 (19)
N1—C1—C2—C7	0.4 (3)	C4—C11—C12—C17	103.56 (16)

C6—C1—C2—C7	175.9 (2)	C4—C11—C12—C13	-76.11 (17)
N1—C1—C2—C7'	24.7 (6)	C17—C12—C13—C14	0.6 (2)
C6—C1—C2—C7'	-159.8 (6)	C11—C12—C13—C14	-179.74 (12)
C1—C2—C3—C4	1.2 (3)	C12—C13—C14—C15	-1.2 (2)
C7—C2—C3—C4	-176.90 (19)	C12—C13—C14—C18	177.80 (13)
C7'—C2—C3—C4	153.0 (9)	C13—C14—C15—N2	-177.22 (12)
C2—C3—C4—C5	-0.5 (2)	C18—C14—C15—N2	3.8 (2)
C2—C3—C4—C11	175.98 (15)	C13—C14—C15—C16	0.7 (2)
C3—C4—C5—C6	0.6 (2)	C18—C14—C15—C16	-178.32 (13)
C11—C4—C5—C6	-175.81 (12)	N2—C15—C16—C17	178.22 (13)
C4—C5—C6—C1	-1.5 (2)	C14—C15—C16—C17	0.4 (2)
C4—C5—C6—C9	177.61 (12)	N2—C15—C16—C20	0.3 (2)
N1—C1—C6—C5	177.72 (13)	C14—C15—C16—C20	-177.59 (13)
C2—C1—C6—C5	2.1 (2)	C13—C12—C17—C16	0.5 (2)
N1—C1—C6—C9	-1.4 (2)	C11—C12—C17—C16	-179.18 (13)
C2—C1—C6—C9	-176.97 (13)	C15—C16—C17—C12	-1.0 (2)
C3—C2—C7—C8	-93.4 (5)	C20—C16—C17—C12	177.05 (14)
C1—C2—C7—C8	88.6 (4)	C13—C14—C18—C19	0.3 (2)
C3—C2—C7'—C8'	-1.0 (18)	C15—C14—C18—C19	179.22 (13)
C1—C2—C7'—C8'	153.5 (12)	C17—C16—C20—C21	-97.07 (19)
C5—C6—C9—C10	0.1 (2)	C15—C16—C20—C21	80.9 (2)
C1—C6—C9—C10	179.21 (13)		

Hydrogen-bond geometry (\AA , $^\circ$)

Cg1 is the centroid of the C1–C6 ring.

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
N2—H2 <i>A</i> \cdots N1 ⁱ	0.88	2.62	3.472 (2)	165
C17—H17 \cdots Cg1 ⁱⁱ	0.95	2.92	3.8568 (17)	171

Symmetry codes: (i) $x, y+1, z$; (ii) $-x+2, -y+1, -z+1$.**4,4'-(MethAnediyl)bis(3-chloro-2,6-diethylaniline) (2)***Crystal data* $\text{C}_{21}\text{H}_{28}\text{Cl}_2\text{N}_2$ $M_r = 379.35$ Triclinic, $P\bar{1}$ $a = 8.993$ (3) \AA $b = 9.6540$ (13) \AA $c = 12.216$ (4) \AA $\alpha = 69.352$ (11) $^\circ$ $\beta = 77.257$ (12) $^\circ$ $\gamma = 87.670$ (8) $^\circ$ $V = 967.2$ (5) \AA^3 $Z = 2$ $F(000) = 404$ $D_x = 1.303$ Mg m^{-3} Synchrotron radiation, $\lambda = 0.75268$ \AA

Cell parameters from 5979 reflections

 $\theta = 1.7$ – 31.0° $\mu = 0.40$ mm^{-1} $T = 100$ K

Plate, colourless

 $0.19 \times 0.12 \times 0.05$ mm*Data collection*Rayonix SX165 CCD
diffractometer φ scansAbsorption correction: empirical (using
intensity measurements)[*XDS* (Kabsch, 2010)] $T_{\min} = 0.001$, $T_{\max} = 1.000$

17607 measured reflections
 4844 independent reflections
 3879 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.053$

$\theta_{\text{max}} = 30.5^\circ$, $\theta_{\text{min}} = 2.4^\circ$
 $h = -12 \rightarrow 12$
 $k = -13 \rightarrow 13$
 $l = -16 \rightarrow 16$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.051$
 $wR(F^2) = 0.147$
 $S = 1.05$
 4844 reflections
 243 parameters
 0 restraints
 Primary atom site location: structure-invariant
 direct methods

Secondary atom site location: difference Fourier
 map
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0847P)^2 + 0.2296P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.64 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.56 \text{ e } \text{\AA}^{-3}$
 Extinction correction: SHELXL-2019/2
 (Sheldrick 2015b),
 $F_c^* = kF_c[1 + 0.001x F_c^2 \lambda^3 / \sin(2\theta)]^{-1/4}$
 Extinction coefficient: 0.090 (12)

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.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
C11	0.60707 (6)	1.01312 (5)	0.11839 (5)	0.04260 (17)	
C12	0.57096 (6)	0.36463 (6)	0.14187 (5)	0.03812 (19)	0.920 (2)
C12'	0.4357 (7)	0.8207 (7)	0.3210 (6)	0.0426 (19)	0.080 (2)
N1	1.1478 (2)	0.8869 (2)	0.18791 (17)	0.0425 (4)	
H1A	1.162855	0.983508	0.163730	0.051*	
H1B	1.220754	0.824611	0.182099	0.051*	
N2	0.2511 (2)	0.3094 (2)	0.56093 (16)	0.0444 (4)	
H2A	0.232317	0.332820	0.625941	0.053*	
H2B	0.268619	0.216227	0.569476	0.053*	
C1	1.0157 (2)	0.8455 (2)	0.16291 (16)	0.0348 (4)	
C2	1.0027 (2)	0.7044 (2)	0.15586 (16)	0.0335 (4)	
C3	0.8665 (2)	0.6623 (2)	0.13837 (16)	0.0319 (4)	
H3	0.857145	0.566189	0.135205	0.038*	
C4	0.7421 (2)	0.7542 (2)	0.12515 (15)	0.0307 (3)	
C5	0.7613 (2)	0.8934 (2)	0.13066 (16)	0.0326 (4)	
C6	0.8947 (2)	0.9433 (2)	0.14874 (16)	0.0340 (4)	
C7	1.1390 (2)	0.6070 (2)	0.16344 (18)	0.0382 (4)	
H7A	1.229620	0.666353	0.106943	0.046*	
H7B	1.157666	0.578413	0.245303	0.046*	
C8	1.1257 (3)	0.4681 (3)	0.1365 (2)	0.0443 (5)	
H8A	1.105026	0.494073	0.056275	0.066*	
H8B	1.041972	0.403709	0.196063	0.066*	
H8C	1.221399	0.416063	0.139488	0.066*	

C9	0.9082 (3)	1.0940 (2)	0.15711 (18)	0.0405 (4)	
H9A	0.842955	1.162596	0.108485	0.049*	
H9B	1.015074	1.132801	0.123406	0.049*	
C10	0.8605 (3)	1.0892 (2)	0.28658 (19)	0.0459 (5)	
H10A	0.870197	1.189198	0.288216	0.069*	
H10B	0.926583	1.023404	0.334555	0.069*	
H10C	0.754312	1.052072	0.319926	0.069*	
C11	0.5968 (2)	0.6974 (2)	0.10932 (16)	0.0318 (4)	
H11A	0.622748	0.646743	0.050261	0.038*	
H11B	0.533729	0.782290	0.077199	0.038*	
C12	0.5054 (2)	0.5907 (2)	0.22658 (16)	0.0301 (3)	
C13	0.4873 (2)	0.4395 (2)	0.25223 (16)	0.0312 (4)	
H13	0.531639	0.400676	0.191570	0.037*	0.080 (2)
C14	0.4072 (2)	0.3412 (2)	0.36264 (17)	0.0343 (4)	
C15	0.3368 (2)	0.4002 (2)	0.45127 (16)	0.0345 (4)	
C16	0.3530 (2)	0.5527 (2)	0.42903 (17)	0.0338 (4)	
C17	0.4371 (2)	0.6427 (2)	0.31877 (16)	0.0316 (4)	
H17	0.449079	0.745242	0.304869	0.038*	0.920 (2)
C18	0.3987 (3)	0.1759 (2)	0.3901 (2)	0.0413 (4)	
H18A	0.403956	0.124330	0.474962	0.050*	
H18B	0.488723	0.148971	0.339552	0.050*	
C19	0.2542 (3)	0.1209 (3)	0.3690 (2)	0.0524 (6)	
H19A	0.250468	0.167582	0.284280	0.079*	
H19B	0.164323	0.146679	0.418874	0.079*	
H19C	0.255270	0.013057	0.390275	0.079*	
C20	0.2797 (2)	0.6213 (3)	0.52112 (19)	0.0410 (4)	
H20A	0.334078	0.716805	0.502372	0.049*	
H20B	0.291900	0.555093	0.601287	0.049*	
C21	0.1098 (3)	0.6484 (3)	0.5257 (2)	0.0481 (5)	
H21A	0.096024	0.709184	0.445546	0.072*	
H21B	0.071381	0.700143	0.581735	0.072*	
H21C	0.053358	0.553298	0.552671	0.072*	

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C11	0.0462 (3)	0.0385 (3)	0.0499 (3)	0.0102 (2)	-0.0180 (2)	-0.0203 (2)
C12	0.0439 (3)	0.0365 (3)	0.0389 (3)	0.0056 (2)	-0.0091 (2)	-0.0197 (2)
C12'	0.035 (3)	0.045 (3)	0.058 (4)	0.005 (2)	-0.017 (3)	-0.027 (3)
N1	0.0361 (9)	0.0487 (10)	0.0464 (9)	-0.0057 (7)	-0.0128 (7)	-0.0182 (8)
N2	0.0486 (10)	0.0485 (10)	0.0345 (8)	-0.0073 (8)	-0.0085 (8)	-0.0122 (7)
C1	0.0338 (9)	0.0434 (10)	0.0289 (8)	-0.0044 (7)	-0.0075 (7)	-0.0137 (7)
C2	0.0320 (9)	0.0408 (9)	0.0305 (8)	0.0035 (7)	-0.0093 (7)	-0.0149 (7)
C3	0.0337 (9)	0.0362 (8)	0.0300 (8)	0.0027 (7)	-0.0103 (7)	-0.0148 (7)
C4	0.0298 (8)	0.0356 (8)	0.0274 (8)	0.0002 (7)	-0.0070 (7)	-0.0116 (7)
C5	0.0348 (9)	0.0340 (8)	0.0304 (8)	0.0029 (7)	-0.0092 (7)	-0.0120 (7)
C6	0.0381 (9)	0.0350 (9)	0.0293 (8)	-0.0039 (7)	-0.0073 (7)	-0.0114 (7)
C7	0.0323 (9)	0.0482 (10)	0.0388 (10)	0.0058 (8)	-0.0127 (8)	-0.0187 (8)

C8	0.0408 (10)	0.0486 (11)	0.0487 (11)	0.0104 (9)	-0.0149 (9)	-0.0216 (9)
C9	0.0483 (11)	0.0367 (9)	0.0371 (10)	-0.0071 (8)	-0.0092 (9)	-0.0132 (8)
C10	0.0581 (13)	0.0417 (10)	0.0409 (10)	-0.0098 (9)	-0.0078 (10)	-0.0187 (9)
C11	0.0316 (8)	0.0352 (8)	0.0312 (8)	0.0010 (7)	-0.0100 (7)	-0.0130 (7)
C12	0.0274 (8)	0.0348 (8)	0.0323 (8)	0.0031 (6)	-0.0115 (7)	-0.0142 (7)
C13	0.0305 (8)	0.0357 (8)	0.0324 (8)	0.0031 (7)	-0.0118 (7)	-0.0152 (7)
C14	0.0333 (9)	0.0349 (9)	0.0385 (9)	0.0022 (7)	-0.0145 (8)	-0.0137 (7)
C15	0.0304 (8)	0.0433 (10)	0.0310 (8)	-0.0009 (7)	-0.0117 (7)	-0.0115 (7)
C16	0.0297 (8)	0.0444 (10)	0.0340 (9)	0.0043 (7)	-0.0126 (7)	-0.0188 (8)
C17	0.0300 (8)	0.0353 (8)	0.0353 (9)	0.0037 (7)	-0.0130 (7)	-0.0161 (7)
C18	0.0446 (11)	0.0336 (9)	0.0457 (11)	0.0016 (8)	-0.0141 (9)	-0.0116 (8)
C19	0.0611 (14)	0.0389 (10)	0.0621 (14)	-0.0052 (10)	-0.0253 (12)	-0.0160 (10)
C20	0.0382 (10)	0.0556 (12)	0.0385 (10)	0.0054 (9)	-0.0122 (8)	-0.0258 (9)
C21	0.0411 (11)	0.0651 (14)	0.0501 (12)	0.0083 (10)	-0.0120 (9)	-0.0344 (11)

Geometric parameters (Å, °)

C11—C5	1.7666 (19)	C10—H10A	0.9800
C12—C13	1.7614 (19)	C10—H10B	0.9800
C12'—C17	1.728 (6)	C10—H10C	0.9800
N1—C1	1.395 (2)	C11—C12	1.516 (3)
N1—H1A	0.8801	C11—H11A	0.9900
N1—H1B	0.8801	C11—H11B	0.9900
N2—C15	1.389 (3)	C12—C13	1.389 (2)
N2—H2A	0.8802	C12—C17	1.402 (2)
N2—H2B	0.8799	C13—C14	1.401 (3)
C1—C2	1.406 (3)	C13—H13	0.9500
C1—C6	1.410 (3)	C14—C15	1.415 (3)
C2—C3	1.389 (2)	C14—C18	1.511 (3)
C2—C7	1.513 (3)	C15—C16	1.407 (3)
C3—C4	1.401 (2)	C16—C17	1.386 (3)
C3—H3	0.9500	C16—C20	1.517 (3)
C4—C5	1.388 (2)	C17—H17	0.9500
C4—C11	1.513 (2)	C18—C19	1.529 (3)
C5—C6	1.400 (3)	C18—H18A	0.9900
C6—C9	1.506 (3)	C18—H18B	0.9900
C7—C8	1.504 (3)	C19—H19A	0.9800
C7—H7A	0.9900	C19—H19B	0.9800
C7—H7B	0.9900	C19—H19C	0.9800
C8—H8A	0.9800	C20—C21	1.531 (3)
C8—H8B	0.9800	C20—H20A	0.9900
C8—H8C	0.9800	C20—H20B	0.9900
C9—C10	1.529 (3)	C21—H21A	0.9800
C9—H9A	0.9900	C21—H21B	0.9800
C9—H9B	0.9900	C21—H21C	0.9800
C1—N1—H1A	113.1	C12—C11—H11A	109.3
C1—N1—H1B	111.5	C4—C11—H11B	109.3

H1A—N1—H1B	124.1	C12—C11—H11B	109.3
C15—N2—H2A	123.3	H11A—C11—H11B	107.9
C15—N2—H2B	109.2	C13—C12—C17	116.00 (17)
H2A—N2—H2B	117.7	C13—C12—C11	124.17 (16)
N1—C1—C2	119.48 (18)	C17—C12—C11	119.80 (16)
N1—C1—C6	119.95 (18)	C12—C13—C14	123.61 (17)
C2—C1—C6	120.54 (17)	C12—C13—C12	118.97 (14)
C3—C2—C1	118.41 (17)	C14—C13—C12	117.41 (14)
C3—C2—C7	122.75 (17)	C12—C13—H13	118.2
C1—C2—C7	118.81 (17)	C14—C13—H13	118.2
C2—C3—C4	123.40 (17)	C13—C14—C15	118.00 (17)
C2—C3—H3	118.3	C13—C14—C18	122.08 (18)
C4—C3—H3	118.3	C15—C14—C18	119.89 (18)
C5—C4—C3	116.10 (16)	N2—C15—C16	119.08 (18)
C5—C4—C11	124.22 (16)	N2—C15—C14	120.80 (18)
C3—C4—C11	119.66 (16)	C16—C15—C14	120.12 (18)
C4—C5—C6	123.76 (17)	C17—C16—C15	118.63 (17)
C4—C5—C11	118.40 (14)	C17—C16—C20	119.26 (18)
C6—C5—C11	117.81 (14)	C15—C16—C20	122.10 (18)
C5—C6—C1	117.77 (17)	C16—C17—C12	123.59 (17)
C5—C6—C9	121.62 (18)	C16—C17—C12'	107.0 (3)
C1—C6—C9	120.57 (17)	C12—C17—C12'	129.4 (3)
C8—C7—C2	115.95 (16)	C16—C17—H17	118.2
C8—C7—H7A	108.3	C12—C17—H17	118.2
C2—C7—H7A	108.3	C14—C18—C19	113.90 (17)
C8—C7—H7B	108.3	C14—C18—H18A	108.8
C2—C7—H7B	108.3	C19—C18—H18A	108.8
H7A—C7—H7B	107.4	C14—C18—H18B	108.8
C7—C8—H8A	109.5	C19—C18—H18B	108.8
C7—C8—H8B	109.5	H18A—C18—H18B	107.7
H8A—C8—H8B	109.5	C18—C19—H19A	109.5
C7—C8—H8C	109.5	C18—C19—H19B	109.5
H8A—C8—H8C	109.5	H19A—C19—H19B	109.5
H8B—C8—H8C	109.5	C18—C19—H19C	109.5
C6—C9—C10	111.93 (17)	H19A—C19—H19C	109.5
C6—C9—H9A	109.2	H19B—C19—H19C	109.5
C10—C9—H9A	109.2	C16—C20—C21	113.45 (16)
C6—C9—H9B	109.2	C16—C20—H20A	108.9
C10—C9—H9B	109.2	C21—C20—H20A	108.9
H9A—C9—H9B	107.9	C16—C20—H20B	108.9
C9—C10—H10A	109.5	C21—C20—H20B	108.9
C9—C10—H10B	109.5	H20A—C20—H20B	107.7
H10A—C10—H10B	109.5	C20—C21—H21A	109.5
C9—C10—H10C	109.5	C20—C21—H21B	109.5
H10A—C10—H10C	109.5	H21A—C21—H21B	109.5
H10B—C10—H10C	109.5	C20—C21—H21C	109.5
C4—C11—C12	111.77 (14)	H21A—C21—H21C	109.5
C4—C11—H11A	109.3	H21B—C21—H21C	109.5

N1—C1—C2—C3	176.01 (17)	C17—C12—C13—C14	-0.3 (2)
C6—C1—C2—C3	-1.9 (3)	C11—C12—C13—C14	177.54 (16)
N1—C1—C2—C7	-6.0 (3)	C17—C12—C13—C12	-179.42 (12)
C6—C1—C2—C7	176.03 (17)	C11—C12—C13—C12	-1.5 (2)
C1—C2—C3—C4	1.2 (3)	C12—C13—C14—C15	2.1 (3)
C7—C2—C3—C4	-176.68 (17)	C12—C13—C14—C15	-178.78 (13)
C2—C3—C4—C5	-0.1 (3)	C12—C13—C14—C18	-175.87 (16)
C2—C3—C4—C11	-178.58 (17)	C12—C13—C14—C18	3.2 (2)
C3—C4—C5—C6	-0.3 (3)	C13—C14—C15—N2	177.64 (16)
C11—C4—C5—C6	178.12 (17)	C18—C14—C15—N2	-4.3 (3)
C3—C4—C5—C11	-178.45 (13)	C13—C14—C15—C16	-2.3 (3)
C11—C4—C5—C11	-0.1 (2)	C18—C14—C15—C16	175.74 (16)
C4—C5—C6—C1	-0.5 (3)	N2—C15—C16—C17	-179.18 (16)
C11—C5—C6—C1	177.74 (14)	C14—C15—C16—C17	0.8 (3)
C4—C5—C6—C9	-178.43 (18)	N2—C15—C16—C20	0.0 (3)
C11—C5—C6—C9	-0.2 (2)	C14—C15—C16—C20	179.95 (16)
N1—C1—C6—C5	-176.37 (17)	C15—C16—C17—C12	1.1 (3)
C2—C1—C6—C5	1.6 (3)	C20—C16—C17—C12	-178.07 (15)
N1—C1—C6—C9	1.6 (3)	C15—C16—C17—C12'	-179.6 (2)
C2—C1—C6—C9	179.58 (17)	C20—C16—C17—C12'	1.2 (3)
C3—C2—C7—C8	6.9 (3)	C13—C12—C17—C16	-1.4 (2)
C1—C2—C7—C8	-170.95 (19)	C11—C12—C17—C16	-179.33 (16)
C5—C6—C9—C10	91.6 (2)	C13—C12—C17—C12'	179.5 (3)
C1—C6—C9—C10	-86.4 (2)	C11—C12—C17—C12'	1.6 (3)
C5—C4—C11—C12	-102.4 (2)	C13—C14—C18—C19	-96.4 (2)
C3—C4—C11—C12	76.0 (2)	C15—C14—C18—C19	85.6 (2)
C4—C11—C12—C13	-109.95 (18)	C17—C16—C20—C21	99.5 (2)
C4—C11—C12—C17	67.9 (2)	C15—C16—C20—C21	-79.7 (2)

Hydrogen-bond geometry (\AA , $^\circ$)

*Cg*1 and *Cg*2 are the centroids of the C1–C6 and C12–C17 rings, respectively.

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
N2—H2 <i>A</i> \cdots <i>Cg</i> 1 ⁱ	0.88	2.63	3.327 (2)	137
C7—H7 <i>B</i> \cdots <i>Cg</i> 2 ⁱⁱ	0.99	2.83	3.579 (2)	133
C20—H20 <i>B</i> \cdots <i>Cg</i> 2 ⁱ	0.99	2.81	3.434 (2)	132

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