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(E)-4-Chloro-N-(2,4,6-trimethylbenzylidene)aniline

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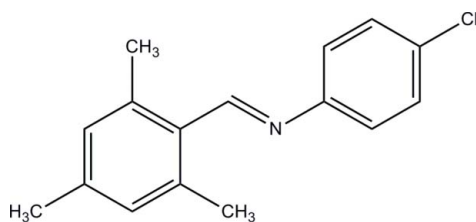
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 Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.046; wR factor = 0.136; data-to-parameter ratio = 15.3.

In the title compound, $\text{C}_{16}\text{H}_{16}\text{ClN}$, the dihedral angle between the benzene rings is $24.61(13)^\circ$. In the crystal, only van der Waals interactions occur between neighbouring molecules.

Related literature

For related structures, see: Nie (2008); Cui *et al.* (2009); Sun *et al.* (2011a,b).



Experimental

Crystal data

$\text{C}_{16}\text{H}_{16}\text{ClN}$
 $M_r = 257.75$
 Monoclinic, $P2_1/c$
 $a = 7.198(2)$ Å

$b = 12.398(4)$ Å
 $c = 15.865(5)$ Å
 $\beta = 102.296(4)^\circ$
 $V = 1383.3(7)$ Å³

$Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.26$ mm⁻¹

$T = 293$ K
 $0.31 \times 0.30 \times 0.29$ mm

Data collection

Bruker APEXII CCD diffractometer
 Absorption correction: multi-scan (SADABS; Bruker, 2004)
 $T_{\min} = 0.924$, $T_{\max} = 0.929$

7071 measured reflections
 2553 independent reflections
 1593 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.042$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.046$
 $wR(F^2) = 0.136$
 $S = 1.04$
 2553 reflections

167 parameters
 H-atom parameters constrained
 $\Delta\rho_{\max} = 0.19$ e Å⁻³
 $\Delta\rho_{\min} = -0.16$ e Å⁻³

Data collection: APEX2 (Bruker, 2004); cell refinement: SAINT (Bruker, 2004); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXTL.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB5944).

References

- Bruker (2004). APEX2, SAINT and SADABS. Bruker AXS Inc., Madison, Wisconsin, USA.
 Cui, C., Meng, Q. & Wang, Y. (2009). *Acta Cryst.* **E65**, o2472.
 Nie, Y. (2008). *Acta Cryst.* **E64**, o471.
 Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
 Sun, L.-X., Yu, Y.-D. & Wei, G.-Y. (2011a). *Acta Cryst.* **E67**, o1564.
 Sun, L.-X., Yu, Y.-D. & Wei, G.-Y. (2011b). *Acta Cryst.* **E67**, o1578.

supporting information

Acta Cryst. (2011). E67, o1999 [doi:10.1107/S1600536811027024]

(*E*)-4-Chloro-*N*-(2,4,6-trimethylbenzylidene)aniline

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S1. Comment

We report here the crystal structure of the title new Schiff base compound, (I). In (I) (Fig. 1), the geometric parameters of the title compound agree well with reported similar structures (Nie, 2008; Cui *et al.*, 2009; Sun *et al.*, 2011a,b). The dihedral angle between the two aromatic rings in the Schiff base molecule is 24.61 (13)°.

S2. Experimental

A mixture of 2,4,6-trimethylbenzaldehyde (5 mmol), 4-chloroaniline (5 mmol) and methanol (40 ml) was refluxed for 2 h. It was then allowed to cool and filtered. Recrystallization of the crude product from methanol yielded colorless blocks of (I).

S3. Refinement

H atoms were positioned geometrically and refined using the riding-model approximation, with C—H = 0.93–0.96 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ or $1.5U_{\text{eq}}(\text{methyl C})$.

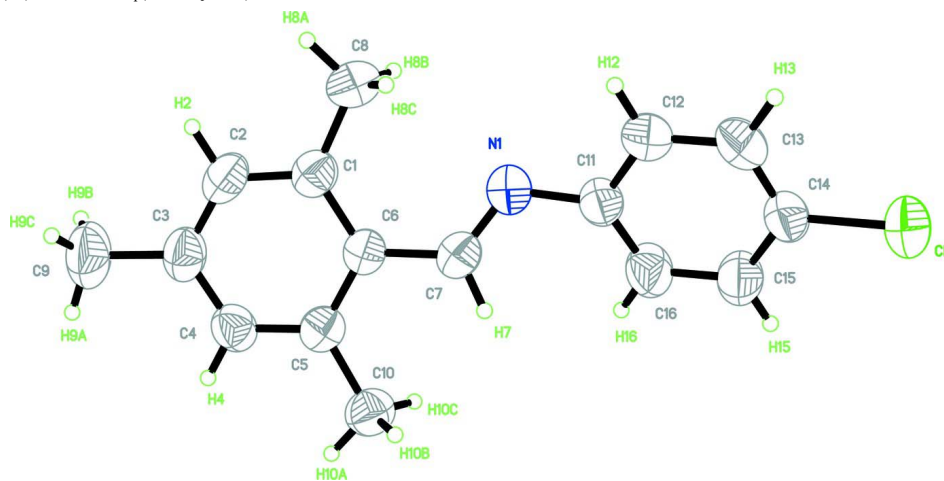


Figure 1

The molecular structure of the title compound with 50% probability displacement ellipsoids for non-hydrogen atoms.

(*E*)-4-Chloro-*N*-(2,4,6-trimethylbenzylidene)aniline

Crystal data

$\text{C}_{16}\text{H}_{16}\text{ClN}$

$M_r = 257.75$

Monoclinic, $P2_1/c$

Hall symbol: $-P\ 2ybc$

$a = 7.198\ (2)\ \text{\AA}$

$b = 12.398\ (4)\ \text{\AA}$

$c = 15.865\ (5)\ \text{\AA}$

$\beta = 102.296\ (4)^\circ$

$V = 1383.3 (7) \text{ \AA}^3$
 $Z = 4$
 $F(000) = 544$
 $D_x = 1.238 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
 Cell parameters from 1950 reflections

$\theta = 2.6\text{--}24.8^\circ$
 $\mu = 0.26 \text{ mm}^{-1}$
 $T = 293 \text{ K}$
 Block, colorless
 $0.31 \times 0.30 \times 0.29 \text{ mm}$

Data collection

Bruker APEXII CCD
 diffractometer
 Radiation source: fine-focus sealed tube
 Graphite monochromator
 φ and ω scans
 Absorption correction: multi-scan
 (SADABS; Bruker, 2004)
 $T_{\min} = 0.924$, $T_{\max} = 0.929$

7071 measured reflections
 2553 independent reflections
 1593 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.042$
 $\theta_{\max} = 25.5^\circ$, $\theta_{\min} = 2.6^\circ$
 $h = -8 \rightarrow 8$
 $k = -15 \rightarrow 12$
 $l = -19 \rightarrow 19$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.046$
 $wR(F^2) = 0.136$
 $S = 1.04$
 2553 reflections
 167 parameters
 0 restraints
 Primary atom site location: structure-invariant
 direct methods
 Secondary atom site location: difference Fourier
 map

Hydrogen site location: inferred from
 neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0476P)^2 + 0.5504P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.19 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.16 \text{ e \AA}^{-3}$
 Extinction correction: SHELXL97 (Sheldrick,
 2008), $F_c^* = kF_c [1 + 0.001x F_c^2 \lambda^3 / \sin(2\theta)]^{-1/4}$
 Extinction coefficient: 0.024 (3)

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s 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 $F^2 > \sigma(F^2)$ 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 (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.2061 (3)	0.3177 (2)	-0.07427 (15)	0.0528 (6)
C2	0.1376 (4)	0.3161 (2)	-0.16278 (16)	0.0607 (7)
H2	0.1204	0.2497	-0.1906	0.073*
C3	0.0937 (4)	0.4085 (2)	-0.21154 (16)	0.0608 (7)
C4	0.1175 (3)	0.5063 (2)	-0.16925 (16)	0.0568 (7)
H4	0.0864	0.5692	-0.2010	0.068*
C5	0.1867 (3)	0.51398 (19)	-0.08065 (15)	0.0489 (6)
C6	0.2323 (3)	0.41907 (19)	-0.03188 (14)	0.0451 (6)
C7	0.3158 (3)	0.4292 (2)	0.06073 (15)	0.0504 (6)

H7	0.3552	0.4976	0.0811	0.061*
C8	0.2484 (5)	0.2115 (2)	-0.02813 (18)	0.0752 (8)
H8A	0.2283	0.1538	-0.0695	0.113*
H8B	0.1655	0.2022	0.0114	0.113*
H8C	0.3782	0.2107	0.0031	0.113*
C9	0.0207 (5)	0.4020 (3)	-0.30862 (17)	0.0902 (10)
H9A	0.0000	0.4735	-0.3319	0.135*
H9B	-0.0967	0.3626	-0.3210	0.135*
H9C	0.1129	0.3657	-0.3342	0.135*
C10	0.2072 (4)	0.6250 (2)	-0.04014 (17)	0.0625 (7)
H10A	0.1588	0.6781	-0.0833	0.094*
H10B	0.3390	0.6392	-0.0163	0.094*
H10C	0.1367	0.6281	0.0048	0.094*
C11	0.4309 (4)	0.3741 (2)	0.20165 (15)	0.0544 (6)
C12	0.5466 (4)	0.2942 (2)	0.24460 (18)	0.0742 (9)
H12	0.5584	0.2297	0.2162	0.089*
C13	0.6456 (4)	0.3073 (2)	0.32874 (18)	0.0729 (8)
H13	0.7257	0.2532	0.3561	0.087*
C14	0.6243 (4)	0.4007 (2)	0.37115 (16)	0.0589 (7)
C15	0.5060 (4)	0.4801 (2)	0.33155 (17)	0.0704 (8)
H15	0.4903	0.5429	0.3613	0.085*
C16	0.4093 (4)	0.4668 (2)	0.24666 (17)	0.0679 (8)
H16	0.3289	0.5211	0.2198	0.081*
Cl1	0.74755 (12)	0.41813 (6)	0.47717 (5)	0.0868 (3)
N1	0.3387 (3)	0.35391 (18)	0.11479 (13)	0.0642 (6)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0509 (15)	0.0545 (15)	0.0543 (15)	-0.0032 (12)	0.0145 (12)	-0.0052 (12)
C2	0.0647 (18)	0.0620 (17)	0.0567 (16)	-0.0099 (13)	0.0161 (13)	-0.0195 (13)
C3	0.0560 (16)	0.0773 (19)	0.0477 (14)	-0.0106 (14)	0.0076 (12)	-0.0068 (14)
C4	0.0553 (16)	0.0606 (16)	0.0514 (15)	-0.0046 (13)	0.0042 (12)	0.0065 (12)
C5	0.0395 (13)	0.0540 (15)	0.0532 (14)	-0.0032 (11)	0.0097 (11)	-0.0038 (11)
C6	0.0386 (13)	0.0526 (14)	0.0451 (13)	-0.0037 (11)	0.0113 (10)	-0.0036 (11)
C7	0.0478 (14)	0.0525 (15)	0.0507 (14)	-0.0031 (12)	0.0097 (11)	-0.0070 (12)
C8	0.102 (2)	0.0502 (16)	0.0739 (19)	0.0005 (15)	0.0191 (17)	-0.0075 (13)
C9	0.098 (2)	0.118 (3)	0.0494 (16)	-0.013 (2)	0.0029 (16)	-0.0095 (17)
C10	0.0644 (18)	0.0555 (16)	0.0653 (17)	0.0012 (13)	0.0087 (13)	-0.0035 (13)
C11	0.0577 (16)	0.0568 (15)	0.0468 (14)	-0.0031 (13)	0.0067 (12)	0.0032 (12)
C12	0.109 (2)	0.0492 (15)	0.0595 (17)	0.0074 (16)	0.0064 (16)	0.0013 (13)
C13	0.091 (2)	0.0588 (17)	0.0618 (17)	0.0151 (16)	-0.0005 (16)	0.0133 (14)
C14	0.0611 (17)	0.0606 (17)	0.0495 (15)	-0.0013 (13)	-0.0004 (12)	0.0066 (12)
C15	0.087 (2)	0.0643 (18)	0.0535 (16)	0.0166 (16)	0.0004 (15)	-0.0049 (13)
C16	0.075 (2)	0.0669 (18)	0.0560 (16)	0.0206 (15)	0.0018 (14)	0.0017 (14)
Cl1	0.1009 (7)	0.0845 (6)	0.0590 (5)	-0.0013 (5)	-0.0189 (4)	0.0027 (4)
N1	0.0795 (16)	0.0600 (14)	0.0491 (12)	-0.0032 (12)	0.0049 (11)	0.0004 (11)

Geometric parameters (Å, °)

C1—C2	1.386 (3)	C9—H9B	0.9600
C1—C6	1.419 (3)	C9—H9C	0.9600
C1—C8	1.506 (3)	C10—H10A	0.9600
C2—C3	1.380 (4)	C10—H10B	0.9600
C2—H2	0.9300	C10—H10C	0.9600
C3—C4	1.379 (3)	C11—C12	1.378 (4)
C3—C9	1.520 (3)	C11—C16	1.379 (3)
C4—C5	1.391 (3)	C11—N1	1.419 (3)
C4—H4	0.9300	C12—C13	1.382 (4)
C5—C6	1.408 (3)	C12—H12	0.9300
C5—C10	1.513 (3)	C13—C14	1.364 (4)
C6—C7	1.469 (3)	C13—H13	0.9300
C7—N1	1.254 (3)	C14—C15	1.365 (4)
C7—H7	0.9300	C14—C11	1.740 (3)
C8—H8A	0.9600	C15—C16	1.388 (4)
C8—H8B	0.9600	C15—H15	0.9300
C8—H8C	0.9600	C16—H16	0.9300
C9—H9A	0.9600		
C2—C1—C6	118.4 (2)	H9A—C9—H9B	109.5
C2—C1—C8	118.1 (2)	C3—C9—H9C	109.5
C6—C1—C8	123.5 (2)	H9A—C9—H9C	109.5
C3—C2—C1	123.1 (2)	H9B—C9—H9C	109.5
C3—C2—H2	118.5	C5—C10—H10A	109.5
C1—C2—H2	118.5	C5—C10—H10B	109.5
C4—C3—C2	117.8 (2)	H10A—C10—H10B	109.5
C4—C3—C9	121.3 (3)	C5—C10—H10C	109.5
C2—C3—C9	120.8 (2)	H10A—C10—H10C	109.5
C3—C4—C5	122.2 (2)	H10B—C10—H10C	109.5
C3—C4—H4	118.9	C12—C11—C16	117.8 (2)
C5—C4—H4	118.9	C12—C11—N1	117.5 (2)
C4—C5—C6	119.3 (2)	C16—C11—N1	124.7 (2)
C4—C5—C10	118.3 (2)	C11—C12—C13	121.7 (3)
C6—C5—C10	122.4 (2)	C11—C12—H12	119.1
C5—C6—C1	119.2 (2)	C13—C12—H12	119.1
C5—C6—C7	118.4 (2)	C14—C13—C12	119.2 (2)
C1—C6—C7	122.3 (2)	C14—C13—H13	120.4
N1—C7—C6	125.9 (2)	C12—C13—H13	120.4
N1—C7—H7	117.0	C13—C14—C15	120.7 (2)
C6—C7—H7	117.0	C13—C14—C11	119.7 (2)
C1—C8—H8A	109.5	C15—C14—C11	119.7 (2)
C1—C8—H8B	109.5	C14—C15—C16	119.7 (3)
H8A—C8—H8B	109.5	C14—C15—H15	120.1
C1—C8—H8C	109.5	C16—C15—H15	120.1
H8A—C8—H8C	109.5	C11—C16—C15	120.9 (3)
H8B—C8—H8C	109.5	C11—C16—H16	119.6

C3—C9—H9A	109.5	C15—C16—H16	119.6
C3—C9—H9B	109.5	C7—N1—C11	119.9 (2)
C6—C1—C2—C3	0.0 (4)	C5—C6—C7—N1	169.1 (2)
C8—C1—C2—C3	179.8 (2)	C1—C6—C7—N1	-14.4 (4)
C1—C2—C3—C4	-0.8 (4)	C16—C11—C12—C13	-2.9 (4)
C1—C2—C3—C9	179.4 (2)	N1—C11—C12—C13	178.6 (3)
C2—C3—C4—C5	1.2 (4)	C11—C12—C13—C14	1.8 (5)
C9—C3—C4—C5	-179.0 (2)	C12—C13—C14—C15	0.4 (4)
C3—C4—C5—C6	-0.7 (4)	C12—C13—C14—C11	179.8 (2)
C3—C4—C5—C10	-179.8 (2)	C13—C14—C15—C16	-1.3 (4)
C4—C5—C6—C1	-0.2 (3)	C11—C14—C15—C16	179.3 (2)
C10—C5—C6—C1	178.9 (2)	C12—C11—C16—C15	2.0 (4)
C4—C5—C6—C7	176.4 (2)	N1—C11—C16—C15	-179.6 (3)
C10—C5—C6—C7	-4.5 (3)	C14—C15—C16—C11	0.1 (5)
C2—C1—C6—C5	0.5 (3)	C6—C7—N1—C11	176.6 (2)
C8—C1—C6—C5	-179.3 (2)	C12—C11—N1—C7	-143.8 (3)
C2—C1—C6—C7	-176.0 (2)	C16—C11—N1—C7	37.8 (4)
C8—C1—C6—C7	4.2 (4)		
