

N,N'*-Bis[4-(trifluoromethyl)benzylidene]-butane-1,4-diamine*Mohammad Hossein Habibi,^{a*}
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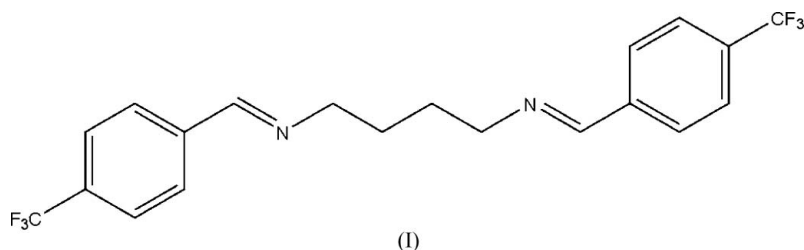
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Key indicatorsSingle-crystal X-ray study
T = 150 K
Mean σ (C–C) = 0.002 Å
R factor = 0.048
wR factor = 0.130
Data-to-parameter ratio = 17.4For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The title compound, C₂₀H₁₈F₆N₂, a Schiff base, was synthesized by the reaction of 1,4-diaminobutane and 4-(trifluoromethyl)benzaldehyde in chloroform solution and was crystallized from a mixture of ethanol and dichloromethane (1:1). It lies across a crystallographic inversion centre. The crystal packing is stabilized by C–H...F hydrogen bonds and π – π interactions.

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Schiff bases have been used widely as ligands in the formation of transition metal complexes. Many such complexes have been structurally characterized, but only a relatively small number of free Schiff bases have been characterized (Calligaris & Randaccio, 1987). Schiff base compounds can be classified by their photochromic and thermochromic characteristics (Cohen *et al.*, 1964). Most Schiff bases possess antibacterial, anticancer and antitoxic activities (Williams, 1972). As an extension of our work (Habibi *et al.*, 2006) on the structural characterization of Schiff base compounds, the title compound, (I), is reported here.



The asymmetric unit is composed of one half-molecule of (I). There is a centre of inversion at the mid-point of the central C–C bond. The C=N unit adopts a *trans* configuration (Fig. 1). The bond lengths and angles in (I) are within expected ranges (You *et al.*, 2004, Munro & Camp, 2003). The benzene rings and the C=N imine bonds are coplanar, as indicated by the C2–N1–C3–C4 torsion angle [178.71 (13)°]. The crystals are held together by hydrogen bonds between secondary CH groups and F atoms in neighboring molecules (Fig. 2 and Table 1). The benzene rings are stacked with centroid–centroid separations of 3.820 (12) and 3.872 (11) Å, indicating π – π interactions.

Experimental

The title compound was prepared by the reaction of 1,4-diaminobutane (0.5 mmol, 44 mg) and 4-(trifluoromethyl)benzaldehyde (1 mmol, 174 mg) dissolved in chloroform (10 ml). The mixture was

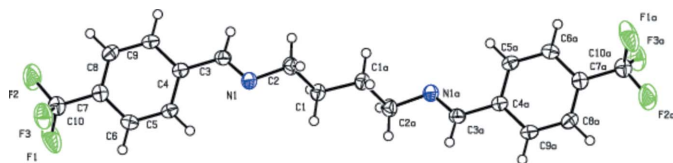


Figure 1

The molecular structure of (I), shown with 50% probability displacement ellipsoids for non-H atoms. [Symmetry code: (a) $-x, 1 - y, -z$.]

stirred at room temperature for 4 h. Single crystals suitable for X-ray structure analysis were obtained from an ethanol–dichloromethane (1:1) solution.

Crystal data

$C_{20}H_{18}F_6N_2$ $V = 930.5 (2) \text{ \AA}^3$
 $M_r = 400.36$ $Z = 2$
 Monoclinic, $P2_1/c$ $Mo K\alpha$ radiation
 $a = 11.0107 (16) \text{ \AA}$ $\mu = 0.13 \text{ mm}^{-1}$
 $b = 9.8541 (14) \text{ \AA}$ $T = 150 (2) \text{ K}$
 $c = 8.6457 (13) \text{ \AA}$ $0.40 \times 0.40 \times 0.20 \text{ mm}$
 $\beta = 97.266 (2)^\circ$

Data collection

Bruker SMART 1K CCD area-detector diffractometer 7930 measured reflections
 Absorption correction: multi-scan (SADABS; Sheldrick, 2004) 2233 independent reflections
 $T_{min} = 0.951, T_{max} = 0.975$ 1930 reflections with $I > 2\sigma(I)$
 $R_{int} = 0.021$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.048$ $\Delta\rho_{max} = 0.62 \text{ e \AA}^{-3}$
 $wR(F^2) = 0.130$ $\Delta\rho_{min} = -0.47 \text{ e \AA}^{-3}$
 $S = 1.06$ Absolute structure: direct methods
 2233 reflections Flack parameter: ?
 128 parameters Rogers parameter: ?
 H-atom parameters constrained

Table 1

Hydrogen-bond geometry ($\text{\AA}, ^\circ$).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$C5-H5\cdots N1$	0.95	2.61	2.875 (2)	97
$C1-H1B\cdots F2^i$	0.99	2.65	3.582 (2)	158

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

All H atoms were placed in idealized positions and constrained to ride on their parent atoms, with $C-H = 0.95$ or 0.99 \AA and $U_{iso}(H) = 1.2U_{eq}(C)$.

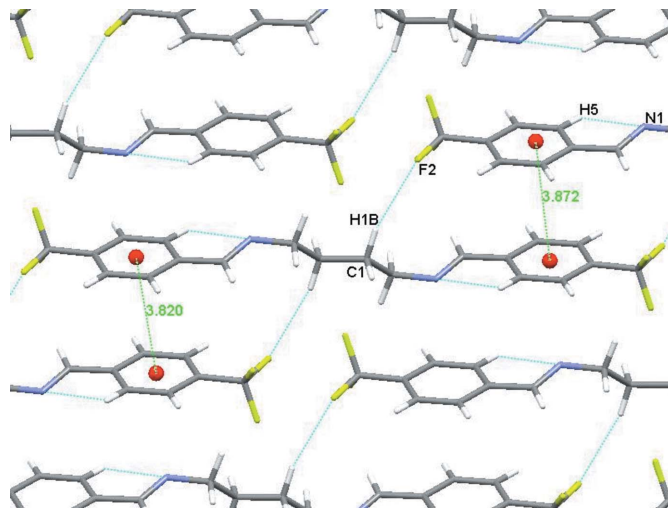


Figure 2

The packing of (I), showing one layer of molecules connected by $C-H\cdots F$ hydrogen bond (turquoise dotted lines) and $\pi-\pi$ interactions (green dotted lines).

Data collection: SMART (Bruker, 2001); cell refinement: SAINT (Bruker, 2001); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Sheldrick, 2001); software used to prepare material for publication: SHELXTL and local programs.

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