## organic papers

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#### Key indicators

Single-crystal X-ray study T = 150 KMean  $\sigma$ (C–C) = 0.002 Å Disorder in main residue R factor = 0.059 wR factor = 0.192 Data-to-parameter ratio = 21.9

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

# [3,5-Bis(trifluoromethyl)phenyl]{(*E*)-2-[(*E*)-3,5-bis(trifluoromethyl)phenylimino]-1-methylpropylidene}amine

At 150 K, the two imine groups of the title compound,  $C_{20}H_{12}F_{12}N_2$ , are mutually *trans*. The molecule has crystal-lographic inversion symmetry.

#### Comment

The title compound, (I), is of interest as a bidentate ligand for catalytically active platinum(II) compounds (Johansson *et al.*, 1999), in which the imine groups are mutually *cis*.



The molecule of (I) is centrosymmetric in the crystal structure. Thus the central diimine group is *trans* and planar. The dihedral angle between the least-squares planes of this group and the benzene ring is 77.25 (12)°. Atom C3 lies 0.0291 (16) Å out of the least-squares plane of the diimine group and N1 lies 0.1344 (13) Å out of the plane of the benzene ring. Thus, there is a slight pyramidalization of *ipso*-C atom C3, which lies 0.0471 (14) Å above the plane of its bonded neighbouring atoms (N1, C4 and C8).

#### **Experimental**

The title compound was prepared by the method of Johansson *et al.* (1999). Suitable crystals were grown from chloroform by slow evaporation.



#### Figure 1

View of the molecule of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms are represented by circles of arbitrary size. Symmetry code (i) is as in Table 1. Both disorder components are shown.

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## Crystal data

 $C_{20}H_{12}F_{12}N_2$  $M_r = 508.32$ Triclinic,  $P\overline{1}$ a = 8.1406 (12) Åb = 8.2225 (12) Åc = 8.7087 (13) Å $\alpha = 82.044 \ (12)^{\circ}$  $\beta = 71.590 (11)^{\circ}$  $\gamma = 68.264 \ (11)^{\circ}$  $V = 513.62 (13) \text{ Å}^3$ Data collection

Stoe IPDS-II diffractometer  $\omega$  scans Absorption correction: none 10945 measured reflections 3638 independent reflections 2477 reflections with  $I > 2\sigma(I)$ 

### Refinement

| Refinement on $F^2$             | $w = 1/[\sigma^2(F_o^2) + (0.1038P)^2]$                   |
|---------------------------------|-----------------------------------------------------------|
| $R[F^2 > 2\sigma(F^2)] = 0.059$ | + 0.0549P]                                                |
| $wR(F^2) = 0.192$               | where $P = (F_o^2 + 2F_c^2)/3$                            |
| S = 1.15                        | $(\Delta/\sigma)_{\rm max} < 0.001$                       |
| 3638 reflections                | $\Delta \rho_{\rm max} = 0.62 \ {\rm e} \ {\rm \AA}^{-3}$ |
| 166 parameters                  | $\Delta \rho_{\rm min} = -0.45 \text{ e} \text{ Å}^{-3}$  |
| H-atom parameters constrained   | Extinction correction: SHELXL97                           |
|                                 | Extinction coefficient: 0.065 (17)                        |

Z = 1

 $D_{\rm r} = 1.643 {\rm Mg} {\rm m}^{-3}$ 

Cell parameters from 8717

Mo  $K\alpha$  radiation

reflections

 $\mu = 0.17 \text{ mm}^{-1}$ 

T = 150 (2) K

 $R_{\rm int} = 0.036$ 

 $\theta_{\rm max} = 32.3^{\circ}$ 

 $h = -12 \rightarrow 12$  $k = -12 \rightarrow 12$ 

 $l = -13 \rightarrow 13$ 

Lath, colourless

 $0.60 \times 0.30 \times 0.10 \text{ mm}$ 

 $\theta = 2.5 - 32.3^{\circ}$ 

## Table 1

Selected geometric parameters (Å, °).

| N1-C1                              | 1.2731 (19)              | C1-C2                | 1.497 (2)    |
|------------------------------------|--------------------------|----------------------|--------------|
| N1-C3                              | 1.4079 (18)              | C1-C1 <sup>i</sup>   | 1.505 (3)    |
| C1 N1 C3                           | 121 02 (13)              | $C^2$ $C^1$ $C^{1i}$ | 117 77 (15)  |
| N1 C1 C2                           | 121.92(13)<br>126.47(13) | $C_2 - C_1 - C_1$    | 110 56 (13)  |
| N1 - C1 - C2<br>$N1 - C1 - C1^{i}$ | 120.47(15)<br>115.74(16) | C4 - C3 - N1         | 120.81 (13)  |
| M-ei-ei                            | 115.74 (10)              | 04–05–101            | 120.01 (13)  |
| C3-N1-C1-C2                        | 2.5 (2)                  | C1-N1-C3-C8          | -107.89 (17) |
| C3-N1-C1-C1 <sup>i</sup>           | -179.06 (14)             | C1-N1-C3-C4          | 78.82 (19)   |
| Summatry and a (i) y               | 2 1                      |                      |              |

Symmetry code: (i) -x, 2 - y, 1 - z.

All H atoms were initially located in a difference Fourier map. The methyl H atoms were constrained to an ideal geometry, with C-H distances of 0.98 Å, but each group was allowed to rotate freely about its X-C bond. All other H atoms were placed in geometrically idealized positions, with C-H distances of 0.95 Å.  $U_{iso}(H)$  values were set at  $1.2U_{eq}(C)$  for all H atoms. The C10 CF<sub>3</sub> group was disordered over two sites with relative occupancies of 0.887 (2):0.113 (2).





Data collection: X-AREA (Stoe & Cie, 2001); cell refinement: X-AREA; data reduction: X-RED (Stoe & Cie, 2001); program(s) used to solve structure: X-STEP32 (Stoe & Cie, 2001) and SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: WinGX (Farrugia, 1999) and SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: WinGX.

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