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Crystal structure, Hirshfeld surface analysis and DFT studies of 4-methyl-2-({[4-(trifluoromethyl)phenyl]imino}methyl)phenol

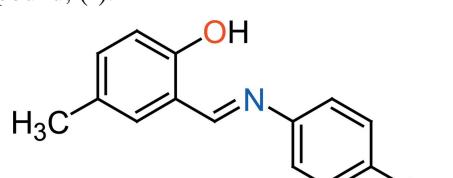
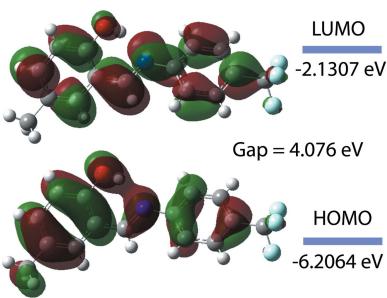
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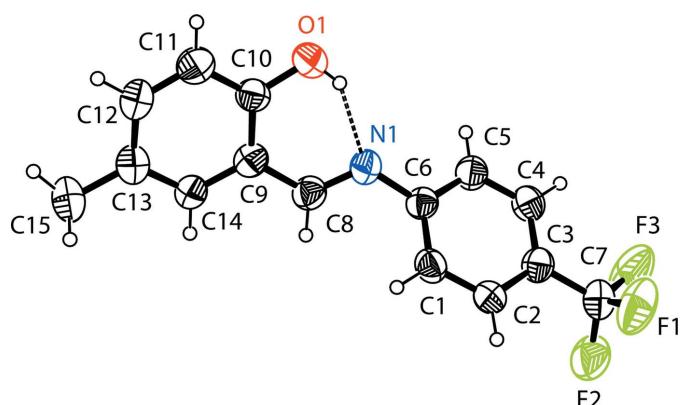
The title compound, $C_{15}H_{12}F_3NO$, crystallizes with one molecule in the asymmetric unit. The configuration of the $C\equiv N$ bond is *E* and there is an intramolecular O—H···N hydrogen bond present, forming an *S*(6) ring motif. The dihedral angle between the mean planes of the phenol and the 4-trifluoromethylphenyl rings is $44.77(3)^\circ$. In the crystal, molecules are linked by C—H···O interactions, forming polymeric chains extending along the *a*-axis direction. The Hirshfeld surface analysis indicates that the most important contributions to the crystal packing are from C···H/H···C (29.2%), H···H (28.6%), F···H/H···F (25.6%), O···H/H···O (5.7%) and F···F (4.6%) interactions. The density functional theory (DFT) optimized structure at the B3LYP/6-311 G(d,p) level is compared with the experimentally determined molecular structure in the solid state. The HOMO–LUMO behaviour was elucidated to determine the energy gap. The crystal studied was refined as an inversion twin.

1. Chemical context

Over the past 25 years, there has been extensive research on the synthesis and use of Schiff base compounds in organic and inorganic chemistry as they have important medicinal and pharmaceutical applications. These compounds show biological activities including antibacterial, antifungal, anti-cancer and herbicidal activities (Desai *et al.*, 2001; Singh & Dash, 1988; Karia & Parsania, 1999). Schiff bases are also becoming increasingly important in the dye and plastics industries, as well as in liquid-crystal technology and for the mechanistic investigation of drugs used in pharmacology, biochemistry and physiology (Sheikhshoae & Sharif, 2006). The present work is a part of an ongoing structural study of Schiff bases and their use in the synthesis of new organic, excited-state proton-transfer compounds and fluorescent chemosensors (Faizi *et al.*, 2016, 2018; Kumar *et al.*, 2018; Mukherjee *et al.*, 2018). We report here on the synthesis and crystal structure as well as the Hirshfeld surface analysis of the new compound, (I).



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**Figure 1**

The molecular structure of (I) with the atom-numbering scheme. Displacement ellipsoids are drawn at the 40% probability level. The intramolecular $\text{O}-\text{H}\cdots\text{N}$ hydrogen bond (Table 1) is shown as a dashed line.

The results of calculations by density functional theory (DFT) carried out at the B3LYP/6-311 G(d,p) level are compared with the experimentally determined molecular structure of (I) in the solid state.

2. Structural commentary

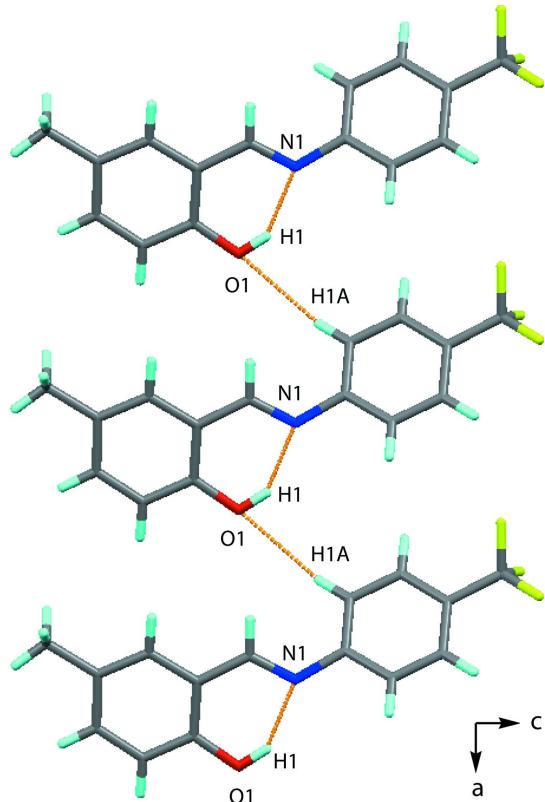
The molecular structure of the title compound, (I), is illustrated in Fig. 1. There is an intramolecular $\text{O}-\text{H}\cdots\text{N}$

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{O1}-\text{H1}\cdots\text{N1}$	0.82	1.90	2.620 (7)	146
$\text{C1}-\text{H1A}\cdots\text{O1}^i$	0.93	2.60	3.463 (7)	154

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

hydrogen bond present (Table 1 and Fig. 1), forming an $S(6)$ ring motif; this is a common feature in related imine–phenol compounds. The imine group displays a $\text{C9}-\text{C8}-\text{N1}-\text{C6}$ torsion angle of $170.1(4)^\circ$ while the mean plane of the phenol ring ($\text{C9}-\text{C14}$) is inclined to that of the 4-trifluoromethylphenyl group ($\text{C1}-\text{C6}$) by $44.77(3)^\circ$. The configuration of the $\text{C8}=\text{N1}$ bond is *E*. The $\text{C10}-\text{O1}$ bond length [1.357 (8) \AA (experimental) and 1.342 \AA (calculated)] indicates single-bond character (Ozeryanskii *et al.*, 2006), while the imine $\text{C8}=\text{N1}$ bond length [1.283 (8) \AA (experimental) and 1.290 \AA (calculated)] indicates double-bond character. All these data support the existence of the phenol–imine tautomer for (I) in its crystalline state. These features are similar to those observed in related 4-dimethylamino-*N*-salicylideneanilines (Pizzala *et al.*, 2000).

**Figure 2**

A view along the b axis of the polymeric chain formed via $\text{C}-\text{H}\cdots\text{O}$ interactions (see Table 1 for details).

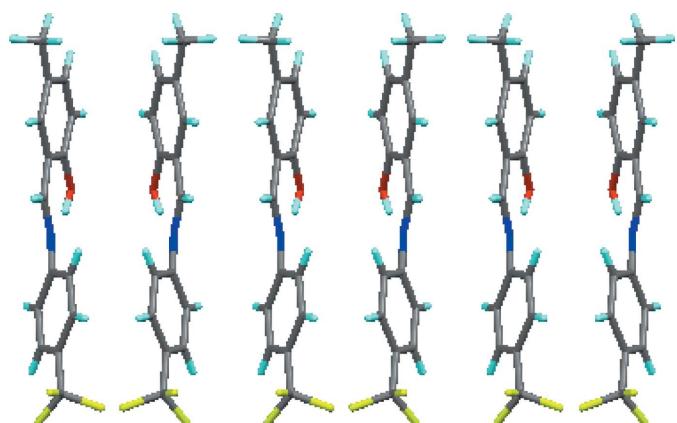


Figure 3
A view of the crystal packing along the a axis.

3. Supramolecular features

In the crystal of (I), molecules are linked by intermolecular C—H···O interactions, forming chains extending along the *a*-axis direction (Fig. 2 and Table 1). The crystal packing along the *a*-axis direction is shown in Fig. 3.

4. Hirshfeld surface analysis and two-dimensional fingerprint plots

In order to visualize the role of weak intermolecular interactions in the crystal, a Hirshfeld surface (HS) analysis (Spackman & Jayatilaka, 2009) was carried out along with the associated two-dimensional fingerprint plots (McKinnon *et al.*, 2007) generated using *CrystalExplorer17.5* (Turner *et al.*, 2017). The three-dimensional d_{norm} (Fig. 4*a*) and shape-index

(Fig. 4*c*) surfaces of (I) are shown with a standard surface resolution and a fixed colour scale of −0.1805 to 1.0413 a.u. The darkest red spots on the Hirshfeld surface indicate contact points with atoms participating in intramolecular C—H···O (Fig. 4*b*) interactions that involve C1—H1A and the oxygen atom O1 of the phenol group (Table 1). As illustrated in Fig. 5*a*, the corresponding fingerprint plots for (I) have characteristic pseudo-symmetrical wings along the d_e and d_i diagonal axes. The presence of C—H···O interactions in the crystal is indicated by the pair of characteristic wings in the fingerprint plot delineated into C···H/H···C (Fig. 5*b*) contacts (29.2% contributions to the Hirshfeld surface). In Fig. 5*c*, the widely scattered points in the fingerprint plot are related to H···H contacts, which make a contribution of 28.6% to the Hirshfeld surface. There are also F···H/H···F (25.6%; Fig. 5*d*), O···H/H···O (5.7%; Fig. 5*e*) and F···F

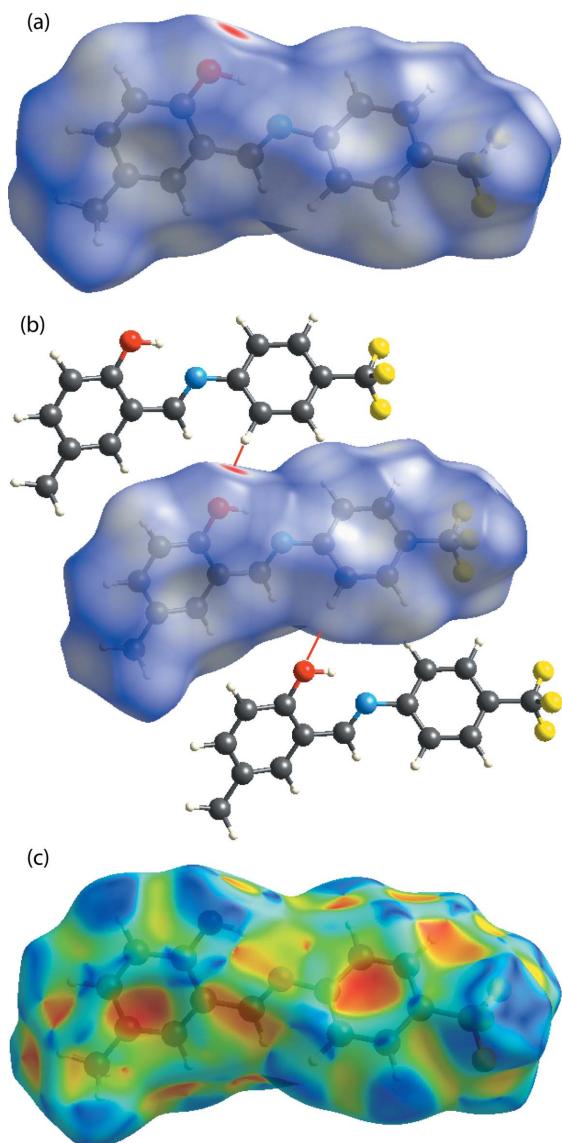


Figure 4

A view of the Hirshfeld surface of (I) mapped over (a) d_{norm} , (b) intermolecular C—H···O interactions and (c) shape-index.

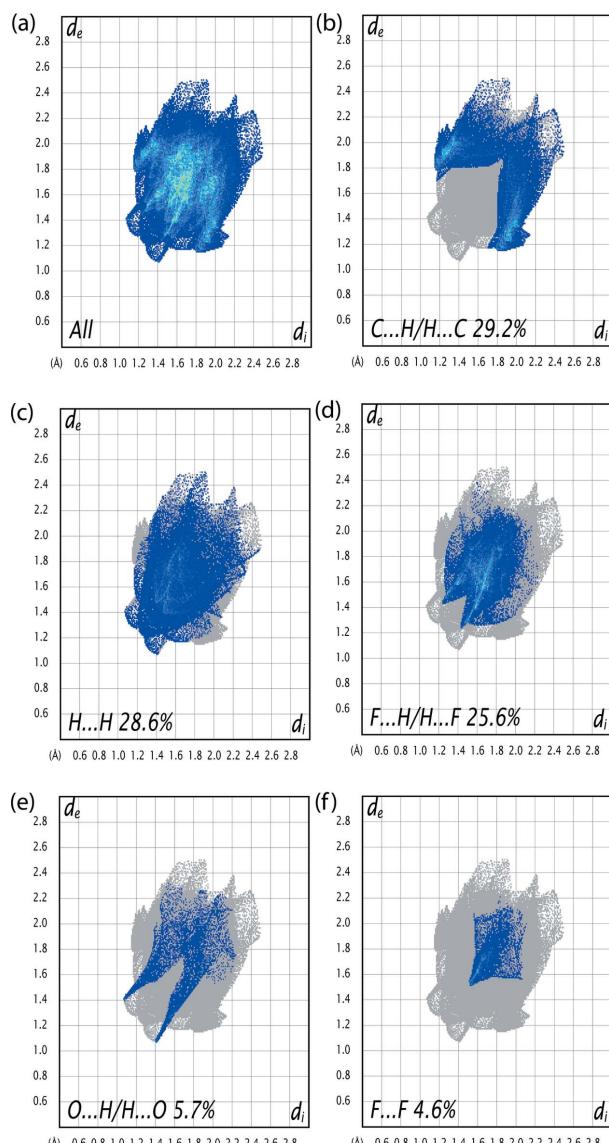


Figure 5

(a) The overall two-dimensional finger print plot for (I) and those delineated into: (b) C···H/H···C (29.2%), (c) H···H (28.6%), (d) F···H/H···F (25.6%), (e) O···H/H···O (5.7%) and (f) F···F (4.6%) contacts.

Table 2

Comparison of selected observed (X-ray data) and calculated (DFT) geometric parameters (\AA , $^\circ$) for (I).

Parameter	X-ray	B3LYP/6-311G(d,p)
O1—C10	1.357 (8)	1.342
N1—C8	1.283 (8)	1.290
C3—C7	1.497 (9)	1.502
C6—N1	1.410 (7)	1.404
C8—C9	1.430 (9)	1.446
N1—C8—C9	123.9 (6)	122.6
C8—N1—C6	122.2 (5)	121.0
O1—C10—C9	122.1 (5)	122.3

(4.6%; Fig. 5f) contacts, with smaller contributions from N···H/H···N (2.4%), O···C/C···O (2.2%), F···C/C···F (0.8%) and O···N/N···O (0.2%) contacts.

5. DFT calculations

The optimized structure of (I) in the gas phase was generated theoretically *via* density functional theory (DFT) using the standard B3LYP functional and the 6-311G(d,p) basis-set calculations (Becke, 1993) as implemented in *GAUSSIAN 09* (Frisch *et al.*, 2009). The theoretical and experimental results are in good agreement (Table 2). The C8=N1 bond length is 1.283 (8) \AA (experimental) and 1.290 \AA (calculated) and the C10—O1 bond length is 1.357 (8) \AA (experimental) and 1.342 \AA (calculated).

The highest-occupied molecular orbital (HOMO) and the lowest-unoccupied molecular orbital (LUMO) are very important aspects as many electronic, optical and chemical reactivity properties of compounds can be predicted from these frontier molecular orbitals (Tanak, 2019). A molecule with a small HOMO–LUMO bandgap is more polarizable than one with a large gap and is considered a ‘soft’ molecule because of its high polarizability while molecules with a large bandgap are considered to be ‘hard’ molecules. To better understand the nature of (I), the electron affinity (A =

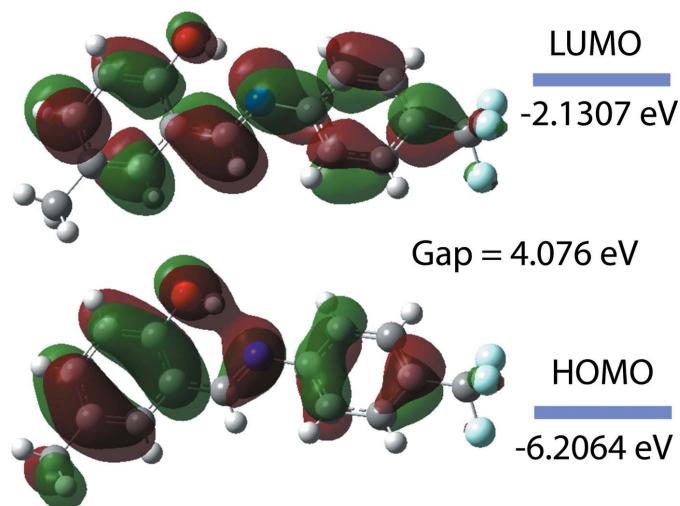


Figure 6
The energy band gap of (I).

Table 3
Interaction energies for (I).

Molecular Energy (a.u.) (eV)	Compound (I)
Total Energy TE (eV)	-27438.7489
E_{HOMO} (eV)	-6.2064
E_{LUMO} (eV)	-2.1307
Gap, ΔE (eV)	4.076
Dipole moment, μ (Debye)	4.466
Ionization potential, I (eV)	6.2064
Electron affinity, A	2.1307
Electronegativity, χ	4.1685
Hardness, η	2.038
Electrophilicity index, ω	4.2631
Softness, σ	0.245
Fraction of electrons transferred, ΔN	0.695

$-E_{\text{HOMO}}$), the ionization potential ($I = -E_{\text{LUMO}}$), the HOMO–LUMO energy gap (ΔE), the chemical hardness (η) and softness (S) (based on the E_{HOMO} and E_{LUMO} energies; Tanak, 2019) were calculated (Table 3). Based on the relatively large ΔE and η values, the title compound can be classified as a hard molecule.

The electron distribution of the HOMO and LUMO energy levels is shown in Fig. 6. The DFT study shows that the HOMO and LUMO are localized in a plane extending over the whole 4-methyl-2-[(4-trifluoromethylphenylimino)methyl]phenol unit. From the frontier orbital analysis, it can be concluded that a HOMO-to-LUMO excitation of (I) would be a $\pi-\pi^*$ transition that would weaken the imine bond and drive the production of an excited-state keto–amine tautomer from the enol–imine ground state observed in the solid state. The calculated band gap of (I) is 4.076 eV, which is similar to that reported for other Schiff base materials, such as for example (*E*)-2-[(3-chlorophenyl)imino]methyl]-6-methylphenol (energy gap = 4.069 eV; Faizi *et al.*, 2019) and (*E*)-2-[(2-hydroxy-5-methoxybenzylidene)amino]benzonitrile (energy gap = 3.520 eV; Saracoğlu *et al.*, 2020).

6. Database survey

A search of the Cambridge Structural Database (CSD, version 5.40, update of November 2018; Groom *et al.*, 2016) for the (*Z*)-1-phenyl-*N*-[3-(trifluoromethylphenyl)methanimine skeleton yielded seven matches. Metal complexes with ligands analogous to (I) are the ruthenium complex chloro-(1-methyl-4-(propan-2-yl)benzene)-(2-((4-(trifluoromethyl)phenyl)imino)methyl)phenolato)ruthenium(II) (BIHCED; Cassells *et al.*, 2018), the rhodium complex (η^5 -pentamethylcyclopentadienyl)chlorido[2-((4-(trifluoromethyl)phenyl)imino)methyl]phenolato]rhodium(III) (BICIH; Cassells *et al.*, 2018) and the iridium complex (η^5 -pentamethylcyclopentadienyl)chlorido[2-((4-(trifluoromethyl)phenyl)imino)methyl]phenolato]iridium(III) (BIHCON; Cassells *et al.*, 2018). Other similar ligands are incorporated into the titanium complex dichloridobis(3,5-di-*tert*-butyl-*N*-(4-trifluoromethylphenyl)salicylaldiminato)titanium(IV) toluene solvate (INOTUA; Mason *et al.*, 2002) and the copper complex bis[4-trifluoromethylphenyl][(2-oxo-3*H*-naphth-3-ylidene)methyl]aminato}-

copper(II) (POPFEF; Fernández *et al.*, 1994). Two vanadium complexes with ligands similar to that in (I) are dichlorido-{2-[*N*-(4-trifluoromethylphenyl)iminomethyl]phenolato}bis(tetrahydrofuran)vanadium(III) (YOGSUJ; Wu *et al.*, 2008) and chloridobis{2-[*N*-(4-trifluoromethylphenyl)iminomethyl]phenolato}(tetrahydrofuran)vanadium(III) (YOGTOE; Wu *et al.*, 2008). Similar uncomplexed Schiff base molecules are *N*-[3,5-bis(trifluoromethyl)phenyl]-3-methoxysalicylaldimine (Karadayı *et al.*, 2015), 2-[3,5-bis(trifluoromethyl)phenyl]-carboimidoylphenol (Yıldız *et al.*, 2015), 2-[3,5-bis(trifluoromethyl)phenyl]carboimidoylphenol (Ünver *et al.*, 2016), (*E*)-3-[3-(trifluoromethyl)phenylimino]methylbenzene-1,2-diol (Koşar *et al.*, 2010), 2-fluoro-*N*-(3-nitrobenzylidene)-5-(trifluoromethyl)aniline (Yang *et al.*, 2007), (*E*)-2-methyl-6-[3-(trifluoromethyl)phenyliminomethyl]phenol (Akkaya *et al.*, 2007), (*E*)-2-[(4-chlorophenyl)iminomethyl]-4-(trifluoromethoxy)phenol (Tüfekçi *et al.*, 2009) and (*E*)-4-methyl-2-[3-(trifluoromethyl)phenyliminomethyl]phenol (Gül *et al.*, 2007). The C≡N bond lengths in these structures vary from 1.270 (3) to 1.295 (5) Å and the C—O bond lengths from 1.336 (5) to 1.366 (2) Å. The molecular configurations of these structures are also not planar, with dihedral angles between the phenyl rings varying between 5.00 (5) and 47.62 (9)°. It is likely that the intramolecular O—H···N hydrogen bond, where the imine N atom acts as an hydrogen-bond acceptor, is an important prerequisite for the tautomeric shift toward the phenol-imine form. In fact, in all eight structures of the phenol-imine tautomers, hydrogen bonds of this type are observed.

7. Synthesis and crystallization

The title compound was prepared by combining solutions of 2-hydroxy-5-methylbenzaldehyde (38.0 mg, 0.28 mmol) in ethanol (15 mL) and 4-trifluoromethylphenylamine (42.0 mg, 0.28 mmol) in ethanol (15 mL) and stirring the mixture for 8 h under reflux. Single crystals suitable for X-ray analysis were obtained by slow evaporation of an ethanol solution (yield 65%, m.p. 425–427 K).

8. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 4. All C-bound H atoms were positioned geometrically and refined using a riding model with C—H = 0.93–0.97 Å and with $U_{\text{iso}}(\text{H}) = 1.2\text{--}1.5U_{\text{eq}}(\text{C})$. The hydrogen atom of the phenol group was located in a difference map and also included as a riding contributor with O—H = 0.82 Å and $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$. During refinement, the twin transformation matrix (−1.0 0.0 0.0, 0.0 −1.0 0.0, 0.0 0.0 −1.0), was used.

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Table 4
Experimental details.

Crystal data	
Chemical formula	C ₁₅ H ₁₂ F ₃ NO
M _r	279.26
Crystal system, space group	Orthorhombic, Pca2 ₁
Temperature (K)	296
<i>a</i> , <i>b</i> , <i>c</i> (Å)	6.2592 (5), 7.2229 (6), 28.551 (3)
<i>V</i> (Å ³)	1290.77 (19)
<i>Z</i>	4
Radiation type	Mo $\text{K}\alpha$
μ (mm ^{−1})	0.12
Crystal size (mm)	0.72 × 0.55 × 0.22
Data collection	
Diffractometer	Stoe IPDS 2
Absorption correction	Integration (X-RED32; Stoe & Cie, 2002)
<i>T</i> _{min} , <i>T</i> _{max}	0.936, 0.982
No. of measured, independent and observed [<i>I</i> > 2σ(<i>I</i>)] reflections	6209, 2135, 1517
<i>R</i> _{int}	0.111
(sin θ/λ) _{max} (Å ^{−1})	0.622
Refinement	
<i>R</i> [F ² > 2σ(F ²)], <i>wR</i> (F ²), <i>S</i>	0.073, 0.213, 0.99
No. of reflections	2135
No. of parameters	182
No. of restraints	1
H-atom treatment	H-atom parameters constrained
Δρ _{max} , Δρ _{min} (e Å ^{−3})	0.24, −0.22
Absolute structure	Refined as an inversion twin

Computer programs: *X-AREA* and *X-RED32* (Stoe & Cie, 2002), *SHELXT2018/3* (Sheldrick, 2015a), *SHELXL2018/3* (Sheldrick, 2015b), *OLEX2* (Dolomanov *et al.*, 2009), *Mercury* (Macrae *et al.*, 2020), *WinGX* (Farrugia, 2012), *PLATON* (Spek, 2020) and *pubCIF* (Westrip, 2010).

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Crystal structure, Hirshfeld surface analysis and DFT studies of 4-methyl-2-({[4-(trifluoromethyl)phenyl]imino}methyl)phenol

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Computing details

Data collection: *X-AREA* (Stoe & Cie, 2002); cell refinement: *X-AREA*; data reduction: *X-RED32* (Stoe & Cie, 2002); program(s) used to solve structure: *SHELXT2018/3* (Sheldrick, 2015a); program(s) used to refine structure: *SHELXL2018/3* (Sheldrick, 2015b); molecular graphics: *OLEX2* (Dolomanov *et al.*, 2009) and *Mercury* (Macrae *et al.*, 2020); software used to prepare material for publication: *WinGX* (Farrugia, 2012), *PLATON* (Spek, 2020), *SHELXL2018* (Sheldrick, 2015b) and *publCIF* (Westrip, 2010).

4-Methyl-2-({[4-(trifluoromethyl)phenyl]imino}methyl)phenol

Crystal data

$C_{15}H_{12}F_3NO$
 $M_r = 279.26$
Orthorhombic, $Pca2_1$
 $a = 6.2592 (5)$ Å
 $b = 7.2229 (6)$ Å
 $c = 28.551 (3)$ Å
 $V = 1290.77 (19)$ Å³
 $Z = 4$
 $F(000) = 576$

$D_x = 1.437$ Mg m⁻³
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 7251 reflections
 $\theta = 2.8\text{--}26.8^\circ$
 $\mu = 0.12$ mm⁻¹
 $T = 296$ K
Prism, yellow
0.72 × 0.55 × 0.22 mm

Data collection

Stoe IPDS 2
diffractometer
Radiation source: sealed X-ray tube, 12 x 0.4
mm long-fine focus
Plane graphite monochromator
Detector resolution: 6.67 pixels mm⁻¹
rotation method scans
Absorption correction: integration
(X-RED32; Stoe & Cie, 2002)

$T_{\min} = 0.936$, $T_{\max} = 0.982$
6209 measured reflections
2135 independent reflections
1517 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.111$
 $\theta_{\max} = 26.2^\circ$, $\theta_{\min} = 2.8^\circ$
 $h = -7 \rightarrow 6$
 $k = -8 \rightarrow 8$
 $l = -28 \rightarrow 34$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.073$
 $wR(F^2) = 0.213$
 $S = 0.99$
2135 reflections

182 parameters
1 restraint
Primary atom site location: dual
Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from neighbouring sites

H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.1465P)^2]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

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

$$\Delta\rho_{\max} = 0.24 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\min} = -0.22 \text{ e \AA}^{-3}$$

Absolute structure: Refined as an inversion twin

Absolute structure parameter: 0 (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.

Refinement. Refined as a two-component inversion twin

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
N1	-0.2713 (8)	-0.7682 (6)	-0.49830 (18)	0.0645 (11)
O1	0.0688 (7)	-0.6785 (7)	-0.54674 (17)	0.0809 (12)
H1	0.000741	-0.705484	-0.523169	0.121*
F2	-0.8837 (9)	-0.7711 (8)	-0.3228 (2)	0.1216 (18)
C10	-0.0518 (9)	-0.7125 (6)	-0.5853 (2)	0.0630 (13)
C6	-0.3839 (8)	-0.7631 (6)	-0.4556 (2)	0.0570 (12)
F1	-0.6915 (9)	-0.5479 (6)	-0.30622 (18)	0.1272 (19)
C5	-0.2780 (10)	-0.8240 (7)	-0.4157 (2)	0.0655 (14)
H5	-0.141666	-0.874119	-0.418207	0.079*
C3	-0.5758 (10)	-0.7369 (7)	-0.3684 (2)	0.0626 (13)
C4	-0.3737 (10)	-0.8105 (7)	-0.3725 (2)	0.0657 (14)
H4	-0.301758	-0.851192	-0.345952	0.079*
C14	-0.3760 (10)	-0.8129 (7)	-0.6233 (2)	0.0640 (13)
H14	-0.513415	-0.861146	-0.621131	0.077*
C7	-0.6847 (12)	-0.7213 (8)	-0.3219 (2)	0.0744 (16)
C9	-0.2636 (9)	-0.7789 (6)	-0.5821 (2)	0.0628 (13)
C8	-0.3645 (10)	-0.8001 (7)	-0.5375 (2)	0.0657 (14)
H8	-0.505930	-0.839357	-0.536897	0.079*
F3	-0.5971 (13)	-0.8224 (11)	-0.29008 (19)	0.160 (3)
C12	-0.0825 (12)	-0.7083 (8)	-0.6683 (2)	0.0719 (16)
H12	-0.020903	-0.683218	-0.697211	0.086*
C1	-0.5870 (9)	-0.6869 (7)	-0.4518 (2)	0.0638 (13)
H1A	-0.658405	-0.646274	-0.478455	0.077*
C11	0.0350 (11)	-0.6758 (7)	-0.6286 (3)	0.0745 (15)
H11	0.173005	-0.628998	-0.630903	0.089*
C2	-0.6836 (10)	-0.6716 (7)	-0.40787 (19)	0.0627 (13)
H2	-0.818272	-0.618523	-0.404942	0.075*
C13	-0.2928 (12)	-0.7782 (7)	-0.6671 (2)	0.0706 (15)
C15	-0.4146 (15)	-0.8082 (10)	-0.7111 (3)	0.089 (2)
H15A	-0.326864	-0.775423	-0.737439	0.134*
H15B	-0.454828	-0.936105	-0.713450	0.134*
H15C	-0.540560	-0.732414	-0.710949	0.134*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N1	0.069 (3)	0.072 (2)	0.053 (3)	-0.0016 (18)	0.003 (2)	0.002 (2)
O1	0.067 (3)	0.109 (3)	0.067 (3)	-0.016 (2)	-0.006 (2)	-0.002 (2)
F2	0.110 (3)	0.159 (4)	0.096 (3)	-0.050 (3)	0.031 (3)	-0.024 (3)
C10	0.058 (3)	0.065 (3)	0.066 (3)	0.001 (2)	-0.002 (3)	-0.001 (2)
C6	0.058 (3)	0.056 (2)	0.057 (3)	-0.0059 (19)	0.004 (2)	0.003 (2)
F1	0.176 (5)	0.098 (3)	0.108 (3)	-0.023 (3)	0.049 (4)	-0.037 (2)
C5	0.068 (3)	0.068 (3)	0.060 (3)	0.006 (2)	-0.001 (3)	-0.002 (2)
C3	0.070 (4)	0.062 (3)	0.056 (3)	-0.002 (2)	0.003 (3)	0.006 (2)
C4	0.074 (4)	0.065 (3)	0.059 (3)	0.001 (2)	-0.006 (3)	0.009 (2)
C14	0.063 (3)	0.071 (2)	0.059 (3)	0.000 (2)	0.004 (3)	-0.005 (2)
C7	0.092 (4)	0.073 (3)	0.058 (3)	-0.003 (3)	0.000 (3)	-0.003 (3)
C9	0.065 (3)	0.059 (2)	0.064 (3)	0.002 (2)	0.005 (3)	0.005 (2)
C8	0.065 (4)	0.067 (3)	0.064 (4)	-0.005 (2)	0.005 (3)	0.002 (2)
F3	0.200 (7)	0.207 (6)	0.072 (3)	0.102 (5)	0.042 (3)	0.052 (3)
C12	0.089 (5)	0.069 (3)	0.058 (3)	0.003 (3)	0.007 (3)	0.000 (2)
C1	0.066 (3)	0.072 (3)	0.054 (3)	-0.001 (2)	-0.008 (2)	0.005 (2)
C11	0.074 (4)	0.072 (3)	0.078 (4)	-0.001 (2)	0.006 (3)	0.002 (3)
C2	0.060 (3)	0.069 (3)	0.060 (3)	0.003 (2)	-0.006 (3)	0.001 (2)
C13	0.085 (4)	0.067 (3)	0.060 (3)	0.001 (3)	-0.002 (3)	0.000 (2)
C15	0.113 (6)	0.095 (4)	0.060 (4)	-0.007 (4)	-0.011 (3)	-0.002 (3)

Geometric parameters (\AA , $^\circ$)

N1—C8	1.283 (8)	C14—C9	1.391 (9)
N1—C6	1.410 (7)	C14—H14	0.9300
O1—C10	1.357 (8)	C7—F3	1.289 (8)
O1—H1	0.8200	C9—C8	1.430 (9)
F2—C7	1.297 (9)	C8—H8	0.9300
C10—C11	1.376 (10)	C12—C11	1.371 (10)
C10—C9	1.412 (8)	C12—C13	1.410 (10)
C6—C5	1.389 (8)	C12—H12	0.9300
C6—C1	1.389 (8)	C1—C2	1.397 (8)
F1—C7	1.331 (7)	C1—H1A	0.9300
C5—C4	1.374 (8)	C11—H11	0.9300
C5—H5	0.9300	C2—H2	0.9300
C3—C4	1.377 (9)	C13—C15	1.487 (10)
C3—C2	1.395 (8)	C15—H15A	0.9600
C3—C7	1.497 (9)	C15—H15B	0.9600
C4—H4	0.9300	C15—H15C	0.9600
C14—C13	1.377 (9)		
		C8—N1—C6	122.2 (5)
		C10—O1—H1	109.5
		O1—C10—C11	118.3 (5)
		O1—C10—C9	122.2 (6)
		C14—C9—C8	120.7 (5)
		C10—C9—C8	120.5 (6)
		N1—C8—C9	123.9 (6)
		N1—C8—H8	118.0

C11—C10—C9	119.5 (6)	C9—C8—H8	118.0
C5—C6—C1	119.9 (5)	C11—C12—C13	122.8 (6)
C5—C6—N1	117.6 (5)	C11—C12—H12	118.6
C1—C6—N1	122.3 (5)	C13—C12—H12	118.6
C4—C5—C6	120.3 (5)	C6—C1—C2	119.8 (5)
C4—C5—H5	119.8	C6—C1—H1A	120.1
C6—C5—H5	119.8	C2—C1—H1A	120.1
C4—C3—C2	120.4 (6)	C12—C11—C10	119.8 (6)
C4—C3—C7	121.5 (6)	C12—C11—H11	120.1
C2—C3—C7	118.1 (6)	C10—C11—H11	120.1
C5—C4—C3	120.3 (6)	C3—C2—C1	119.3 (5)
C5—C4—H4	119.9	C3—C2—H2	120.4
C3—C4—H4	119.9	C1—C2—H2	120.4
C13—C14—C9	122.9 (6)	C14—C13—C12	116.1 (6)
C13—C14—H14	118.6	C14—C13—C15	123.2 (7)
C9—C14—H14	118.6	C12—C13—C15	120.7 (6)
F3—C7—F2	105.4 (7)	C13—C15—H15A	109.5
F3—C7—F1	108.0 (7)	C13—C15—H15B	109.5
F2—C7—F1	103.7 (6)	H15A—C15—H15B	109.5
F3—C7—C3	112.9 (6)	C13—C15—H15C	109.5
F2—C7—C3	113.5 (5)	H15A—C15—H15C	109.5
F1—C7—C3	112.6 (5)	H15B—C15—H15C	109.5
C14—C9—C10	118.8 (5)		
C8—N1—C6—C5	147.0 (5)	O1—C10—C9—C8	4.5 (7)
C8—N1—C6—C1	−38.2 (7)	C11—C10—C9—C8	−173.6 (5)
C1—C6—C5—C4	0.5 (7)	C6—N1—C8—C9	170.1 (4)
N1—C6—C5—C4	175.5 (4)	C14—C9—C8—N1	−178.8 (5)
C6—C5—C4—C3	0.2 (8)	C10—C9—C8—N1	−2.6 (8)
C2—C3—C4—C5	−1.4 (8)	C5—C6—C1—C2	0.1 (7)
C7—C3—C4—C5	179.7 (5)	N1—C6—C1—C2	−174.7 (5)
C4—C3—C7—F3	−16.4 (9)	C13—C12—C11—C10	0.3 (8)
C2—C3—C7—F3	164.6 (7)	O1—C10—C11—C12	−179.9 (5)
C4—C3—C7—F2	−136.3 (6)	C9—C10—C11—C12	−1.7 (8)
C2—C3—C7—F2	44.7 (7)	C4—C3—C2—C1	2.0 (8)
C4—C3—C7—F1	106.3 (7)	C7—C3—C2—C1	−179.1 (5)
C2—C3—C7—F1	−72.7 (7)	C6—C1—C2—C3	−1.3 (8)
C13—C14—C9—C10	−2.5 (7)	C9—C14—C13—C12	1.1 (8)
C13—C14—C9—C8	173.8 (5)	C9—C14—C13—C15	−177.9 (5)
O1—C10—C9—C14	−179.1 (5)	C11—C12—C13—C14	0.0 (8)
C11—C10—C9—C14	2.7 (7)	C11—C12—C13—C15	179.0 (6)

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
O1—H1···N1	0.82	1.90	2.620 (7)	146

C1—H1A···O1 ⁱ	0.93	2.60	3.463 (7)	154
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Symmetry code: (i) $x-1, y, z$.