

Received 28 October 2017
Accepted 2 November 2017

Edited by H. Stoeckli-Evans, University of
Neuchâtel, Switzerland

Keywords: crystal structure; molecular conformation; hydrogen bonding; polymorphism.

CCDC references: 1583686; 1583685

Supporting information: this article has supporting information at journals.iucr.org/e

Crystal structures of (*E*)-1-{3-[(5-fluoro-2-hydroxybenzylidene)amino]phenyl}ethanone and of a fourth polymorph of (*E*)-1-{3-[(2-hydroxy-3-methoxybenzylidene)amino]phenyl}ethanone

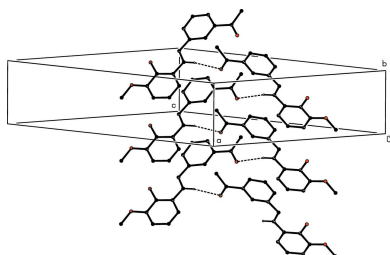
Marisiddaiah Girisha,^a Hemmige S. Yathirajan,^{a*} Ravindranath S. Rathore^b and Christopher Glidewell^{c*}

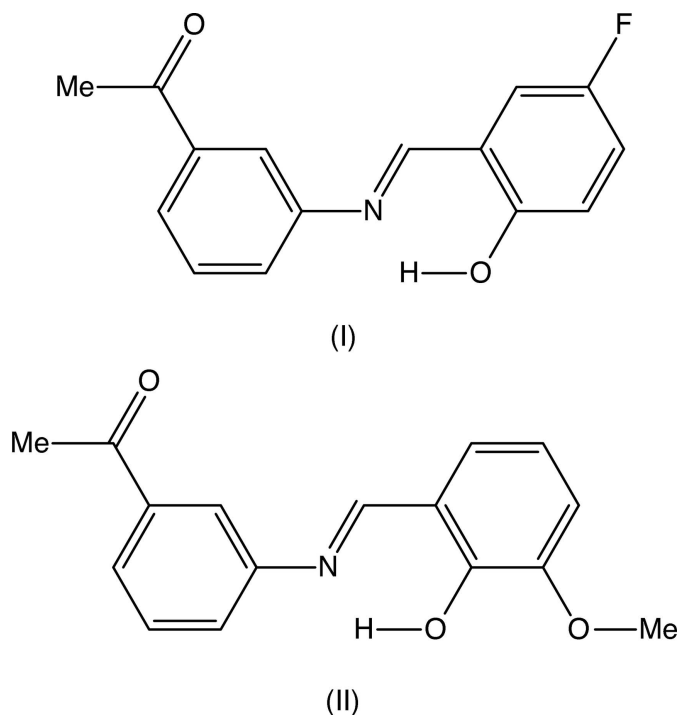
^aDepartment of Studies in Chemistry, University of Mysore, Manasagangotri, Mysuru 570 006, India, ^bCentre for Biological Sciences (Bioinformatics), School of Earth, Biological and Environmental Sciences, Central University of South Bihar, Patna 800 014, India, and ^cSchool of Chemistry, University of St Andrews, St Andrews, Fife KY16 9ST, UK.
*Correspondence e-mail: yathirajan@hotmail.com, cg@st-andrews.ac.uk

In the molecules of both (*E*)-1-{3-[(5-fluoro-2-hydroxybenzylidene)amino]phenyl}ethanone, C₁₅H₁₂FNO₂, (I), and (*E*)-1-{3-[(2-hydroxy-3-methoxybenzylidene)amino]phenyl}ethanone, C₁₆H₁₅NO₃, (II), which crystallizes with $Z' = 2$ in space group *Pca*2₁, there are intramolecular O—H···N hydrogen bonds, and the non-H atoms in each molecule are essentially coplanar. In the crystal of (I), molecules are linked by a single C—H···O hydrogen bond to form a *C*(8) chain, whereas in the crystal of (II), molecules are linked by three C—H···O hydrogen bonds to form sheets within which orthogonal *C*₂²(16) and *C*₂²(17) chains can be identified. Comparisons are made with some related structures.

1. Chemical context

Schiff bases of general type *RR'C=NR''* can exhibit very wide structural diversity and have found a wide range of applications (Jia & Li, 2015), ranging from anti-bacterial, anti-fungal and anti-tumour activity (Rani *et al.*, 2015), *via* catalysis (Kumar *et al.*, 2009), to use as organic photovoltaic materials (Jeevadason *et al.*, 2014). The extensive patent literature on their medicinal applications has recently been reviewed (Hameed *et al.*, 2017). With this great diversity of use in mind, we report herein on the molecular and supramolecular structures of two closely related Schiff bases, (*E*)-1-{3-[(5-fluoro-2-hydroxybenzylidene)amino]phenyl}ethanone (I) and (*E*)-1-{3-[(2-hydroxy-3-methoxybenzylidene)amino]phenyl}ethanone (II). Compounds (I) and (II) were prepared by straightforward condensation reactions between 3-acetylaniline (3-aminoacetophenone) and the appropriately substituted salicylaldehydes. Their molecular constitutions differ only in the identity and location of a single substituent, 5-fluoro in (I) *versus* 3-methoxy in (II), but their crystallization behaviour is different. Compound (I) crystallizes in the monoclinic space group *P*₂₁/*n* with $Z' = 1$ (Fig. 1), while compound (II) crystallizes in the orthorhombic space group *Pca*2₁ with $Z' = 2$ (Figs. 2 and 3), and it will be convenient to refer to the molecules of (II) which contain the atoms N11 and N21 as molecules of types 1 and 2, respectively. Compound (II), in fact, represents the fourth polymorphic form of this compound to be identified. Three other forms, one in *Pna*2₁ with $Z' = 2$, and two others in *P*₂₁2₁2₁, each with $Z' = 1$, have recently been reported (Zbačnik *et al.*, 2015).





2. Structural commentary

In each of compounds (I) (Fig. 1) and (II) (Figs. 2 and 3), the non-H atoms are almost coplanar. Thus in (I), the r.m.s. deviation of the non-H atoms from their mean plane is only 0.085 Å, with a maximum individual deviation from the plane of 0.196 (2) Å for the acetyl atom C18. Similarly, in compound (II), the r.m.s. deviations of the non-H atoms from the mean planes of the two molecules are 0.086 and 0.071 Å for molecules 1 and 2, respectively, with corresponding maximum deviations of 0.225 (5) and 0.211 (5) Å for atoms C118 and C218, respectively. In all of the molecules there is an intramolecular O—H...N hydrogen bond (Tables 1 and 2); although this probably influences the orientation of the hydroxylated ring relative to the central spacer unit, it will not have any influence on the orientation of the acetylphenyl ring

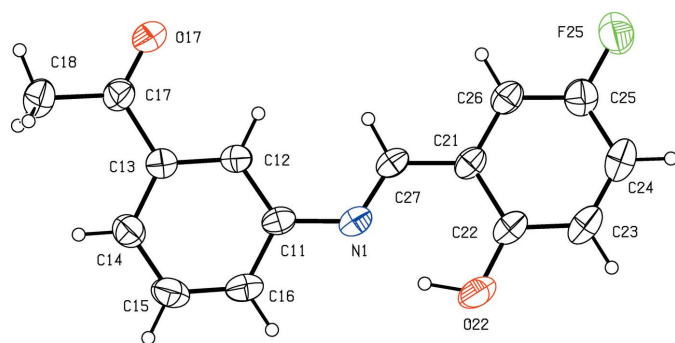


Figure 1
The molecular structure of compound (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level.

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

| <i>D</i> —H... <i>A</i> | <i>D</i> —H | H... <i>A</i> | <i>D</i> ... <i>A</i> | <i>D</i> —H... <i>A</i> |
|----------------------------|-------------|---------------|-----------------------|-------------------------|
| O22—H22...N1 | 0.98 (3) | 1.72 (3) | 2.607 (2) | 148 (3) |
| C27—H27...O17 ⁱ | 0.93 | 2.58 | 3.475 (3) | 163 |

Symmetry code: (i) $-x + \frac{1}{2}, y - \frac{1}{2}, -z + \frac{3}{2}$.

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

| <i>D</i> —H... <i>A</i> | <i>D</i> —H | H... <i>A</i> | <i>D</i> ... <i>A</i> | <i>D</i> —H... <i>A</i> |
|--------------------------------|-------------|---------------|-----------------------|-------------------------|
| O122—H122...N11 | 1.06 (6) | 1.68 (6) | 2.604 (4) | 142 (5) |
| O222—H222...N21 | 0.92 (6) | 1.79 (6) | 2.603 (5) | 147 (5) |
| C116—H116...O223 ⁱ | 0.93 | 2.50 | 3.347 (6) | 152 |
| C127—H127...O217 | 0.93 | 2.59 | 3.496 (5) | 164 |
| C227—H227...O117 ⁱⁱ | 0.93 | 2.58 | 3.487 (5) | 164 |

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

relative to the rest of the molecule. In the two molecules of (II), the deviation of the methoxy C atoms C128 and C228 from the planes of their adjacent aryl rings are 0.107 (9) and 0.049 (11) Å, respectively. Consistent with this, the pair of exocyclic C—C—O angles at each of the atoms C123 and C223 differ by *ca* 10°, as is generally observed in planar alkoxyarene derivatives (Seip & Seip, 1973; Ferguson *et al.*, 1996). The dihedral angle between the mean planes of the two molecules in (II) is 80.74 (3)°.

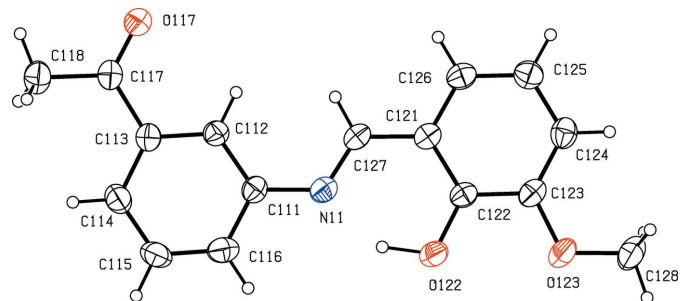


Figure 2
The structure of molecule 1 in compound (II), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level.

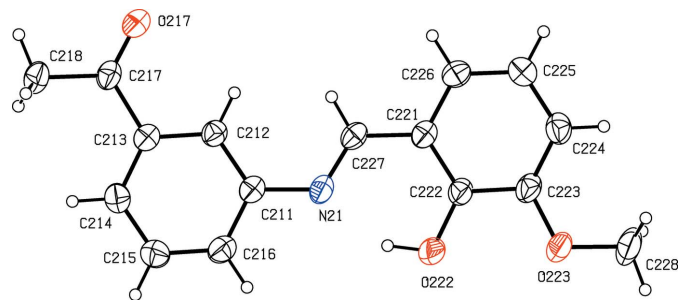


Figure 3
The structure of molecule 2 in compound (II), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level.

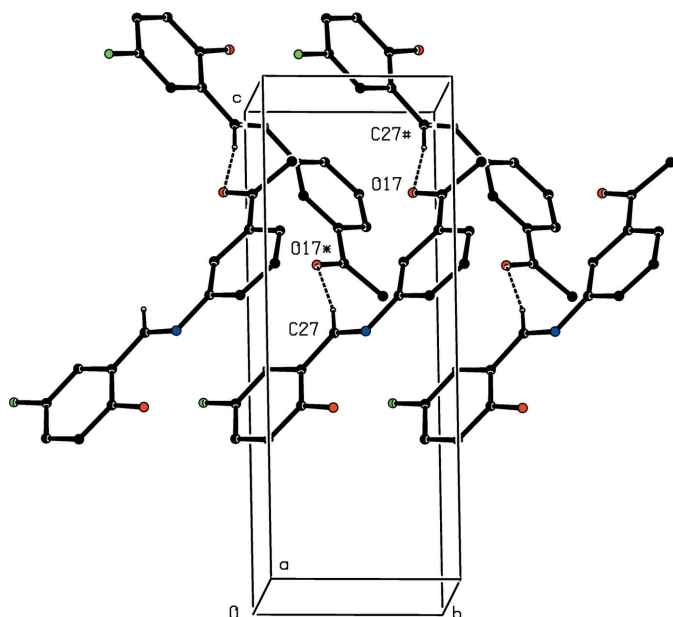


Figure 4
Part of the crystal structure of compound (I), showing the formation of a hydrogen-bonded $C(8)$ chain running parallel to the $[010]$ direction. For the sake of clarity, the H atoms not involved in the motif shown have been omitted. Hydrogen bonds are drawn as dashed lines and the atoms marked with an asterisk (*) or a hash (#) are at the symmetry positions $(\frac{1}{2} - x, -\frac{1}{2} + y, \frac{3}{2} - z)$ and $(\frac{1}{2} - x, \frac{1}{2} + y, \frac{3}{2} - z)$, respectively.

3. Supramolecular features

The supramolecular assembly in compound (I) is very simple, as shown in Fig. 4. In addition to the intramolecular hydrogen bond noted above, there is a single $C-H \cdots O$ hydrogen bond (Table 1), which links molecules related by a 2_1 screw axis into $C(8)$ chains running parallel to the $[010]$ direction. Two chains of this type, related to one another by inversion, pass through each unit cell, but there are no direction-specific interactions between adjacent chains.

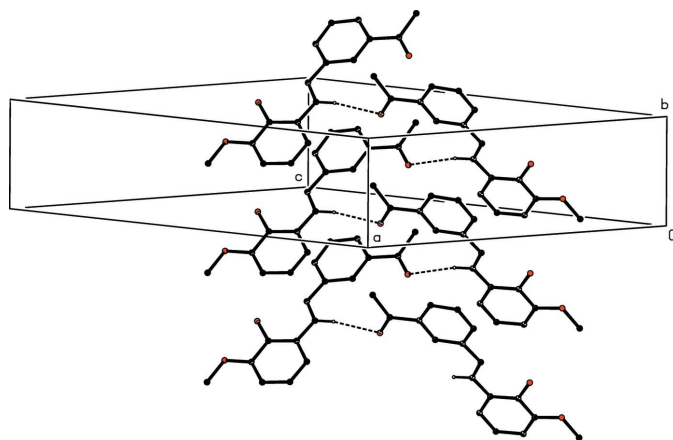


Figure 5
Part of the crystal structure of compound (II), showing the formation of a hydrogen-bonded $C_2(16)$ chain running parallel to the $[010]$ direction. For the sake of clarity, the H atoms not involved in the motif shown have been omitted, and the hydrogen bonds are drawn as dashed lines.

There are three $C-H \cdots O$ hydrogen bonds in the structure of compound (II) (Table 2): one of these links the two molecules within the selected asymmetric unit and the two others link these bimolecular aggregates into complex sheets, whose formation is readily analysed in terms of two one-dimensional sub-structures (Ferguson *et al.*, 1998*a,b*; Gregson *et al.*, 2000). The hydrogen bond having atom C227 as the donor links bimolecular aggregates related by translation to form a $C_2(16)$ chain running parallel to the $[010]$ direction (Fig. 5), and that having atom C116 as the donor links aggregates related by a 2_1 screw axis into $C_2(17)$ chains running parallel to the $[001]$ direction (Fig. 6). The combination of the orthogonal chains along $[010]$ and $[001]$ generates a sheet lying parallel to (100) . Two sheets of this type, related to one another by the glide planes, pass through each unit cell but there are no direction-specific interactions between adjacent sheets.

4. Database survey

The structures of Schiff bases derived from hydroxyaryl aldehydes have recently been the subject of a general survey, in which a number of structural errors, often involving misplaced H atoms, were pointed out (Blagus *et al.*, 2010).

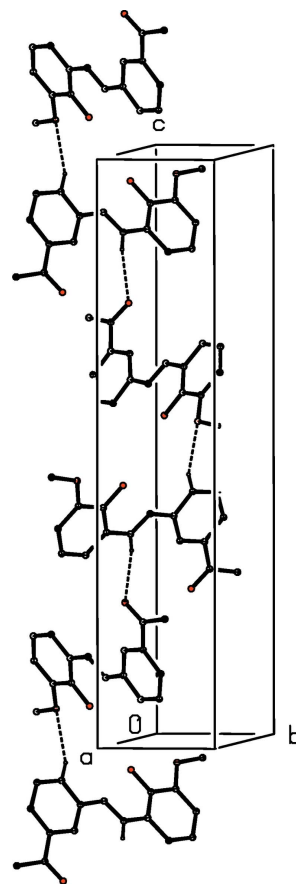


Figure 6
Part of the crystal structure of compound (II), showing the formation of a hydrogen-bonded $C_2(17)$ chain running parallel to the $[001]$ direction. For the sake of clarity, the H atoms not involved in the motif shown have been omitted, and the hydrogen bonds are drawn as dashed lines.

Table 3
Experimental details.

| | (I) | (II) |
|---|--|---|
| Crystal data | | |
| Chemical formula | C ₁₅ H ₁₂ FNO ₂ | C ₁₆ H ₁₅ NO ₃ |
| <i>M_r</i> | 257.26 | 269.29 |
| Crystal system, space group | Monoclinic, <i>P</i> 2 ₁ / <i>n</i> | Orthorhombic, <i>Pca</i> 2 ₁ |
| Temperature (K) | 294 | 294 |
| <i>a</i> , <i>b</i> , <i>c</i> (Å) | 14.9527 (5), 5.5152 (2), 16.6918 (5) | 19.1904 (4), 5.33856 (12), 26.5678 (6) |
| α , β , γ (°) | 90, 114.739 (2), 90 | 90, 90, 90 |
| <i>V</i> (Å ³) | 1250.19 (7) | 2721.85 (10) |
| <i>Z</i> | 4 | 8 |
| Radiation type | Cu <i>K</i> α | Cu <i>K</i> α |
| μ (mm ⁻¹) | 0.84 | 0.75 |
| Crystal size (mm) | 0.15 × 0.15 × 0.10 | 0.10 × 0.10 × 0.05 |
| Data collection | | |
| Diffractometer | Bruker APEX3 | Bruker APEX3 |
| Absorption correction | Multi-scan (<i>SADABS</i> ; Bruker, 2016) | Multi-scan (<i>SADABS</i> ; Bruker, 2016) |
| <i>T</i> _{min} , <i>T</i> _{max} | 0.848, 0.919 | 0.907, 0.963 |
| No. of measured, independent and observed [<i>I</i> > 2 σ (<i>I</i>)] reflections | 15703, 2452, 1764 | 51776, 5393, 3796 |
| <i>R</i> _{int} | 0.041 | 0.117 |
| (<i>sin</i> θ / λ) _{max} (Å ⁻¹) | 0.619 | 0.619 |
| Refinement | | |
| <i>R</i> [<i>F</i> ² > 2 σ (<i>F</i> ²)], <i>wR</i> (<i>F</i> ²), <i>S</i> | 0.046, 0.122, 1.04 | 0.048, 0.113, 1.02 |
| No. of reflections | 2452 | 5393 |
| No. of parameters | 176 | 371 |
| No. of restraints | 0 | 1 |
| H-atom treatment | H atoms treated by a mixture of independent and constrained refinement | H atoms treated by a mixture of independent and constrained refinement |
| $\Delta\rho_{\max}$, $\Delta\rho_{\min}$ (e Å ⁻³) | 0.15, -0.12 | 0.11, -0.13 |
| Absolute structure | – | Flack <i>x</i> determined using 1493 quotients [(<i>I</i> ⁺) – (<i>I</i> [–])] / [(<i>I</i> ⁺) + (<i>I</i> [–])] (Parsons <i>et al.</i> , 2013) |
| Absolute structure parameter | – | –0.04 (16) |

Computer programs: *APEX3* and *SAINT* (Bruker, 2016), *SHELXT2014* (Sheldrick, 2015a), *SHELXL2014* (Sheldrick, 2015b) and *PLATON* (Spek, 2009).

Closely related to the present structures are those of (*E*)-1-[3-[(2-hydroxy-3-methoxybenzylidene)amino]phenyl]ethanone (III) (De *et al.*, 2009), and of the previously recorded polymorphs of (II) (Zbačnik *et al.*, 2015).

Compound (III) is isomorphous with compound (I): as in (I), the structure of (III) contains an intramolecular O–H···N hydrogen bond and the non-H atoms are effectively coplanar. The structure of (III) also contains an intermolecular C–H···O hydrogen bond, although this is nowhere mentioned in the original report (De *et al.*, 2009); this interaction forms *C*(8) chains along [010], exactly the same as those in the structure of (I), so that (I) and (III) are, in fact, isostructural despite their different patterns of substitution.

Three other polymorphic forms of compound (II) have recently been reported and are described as forms I, II and III, respectively (Zbačnik *et al.*, 2015). Form I is orthorhombic in space group *Pna*2₁ with *Z*' = 2, and forms II and III both crystallize in space group *P*2₁2₁2₁ with *Z*' = 1, so that the *Pca*2₁ form reported here can be regarded as form IV. All three forms, I–III, can be crystallized from ethanol solutions under different conditions and a crucial factor in determining which polymorph is obtained appears to be the filtration process used prior to crystallization. By contrast, the form described here was crystallized from a solution in dichloromethane. In all of the molecules in forms I–III, there is an intramolecular

O–H···N hydrogen bond and, in every case, the non-H atoms are effectively co-planar as found here for (I) and (II). The supramolecular assembly differs in all three polymorphs I–III: form II contains no intermolecular hydrogen bonds; in form III two C–H···O hydrogen bonds generate a *C*(8)*C*(10)[*R*₂¹(6)] chain of rings; and in form I, three C–H···O hydrogen bonds generate sheets in which the component sub-structures both involve molecules related by an *n*-glide plane, in contrast to the sheets found for form IV reported here.

5. Synthesis and crystallization

For the synthesis of compounds (I) and (II), 3-acetyl aniline (0.740 mmol) and a catalytic quantity of acetic acid were added to solution of the appropriate aldehyde, 5-fluoro-salicylaldehyde for (I) or 3-methoxysalicylaldehyde for (II) (0.740 mmol) in ethanol (20 cm³), and these mixtures were then heated under reflux for 5 h. The mixtures were then cooled to ambient temperature and the solvent was removed under reduced pressure. The solid residues were then washed with cold ethanol and dried under reduced pressure. Crystals suitable for single crystal X-ray diffraction were grown by slow evaporation, at ambient temperature and in the presence of air, of solutions in dimethylsulfoxide for (I) and in dichloro-

methane for (II): m.p. for (I) 362–364 K and m.p. for (II) 352–354 K.

6. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 3. For compound (II), one bad outlier reflection (8,1,3) was omitted from the data set before the final refinements. All H atoms were located in difference-Fourier maps. The C-bound H atoms were subsequently treated as riding atoms in geometrically idealized positions: C–H 0.93–0.96 Å with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C-methyl})$ and $1.2U_{\text{eq}}(\text{C})$ for other C-bound H atoms. The methyl groups were permitted to rotate but not to tilt. For the H atoms bonded to O atoms, the atomic coordinates were refined with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$, giving the O–H distances shown in Tables 1 and 2. The correct orientation of the structure of (II) relative to the polar axis direction was established by means of the Flack x parameter (Flack, 1983), $x = -0.04$ (16) calculated (Parsons *et al.*, 2013) using 1493 quotients of the type $[(I^+) - (I^-)] / [(I^+) + (I^-)]$, and by means of the Hooft y parameter (Hooft *et al.*, 2010), $y = -0.03$ (16). In the final analysis of variance for (I) there was a large value, 1.859, of $K = [\text{mean}(F_o^2) / \text{mean}(F_c^2)]$ for the group of 4258 very weak reflections having $F_o/F_c(\text{max})$ in the range $0.000 < F_o/F_c(\text{max}) < 0.008$; the corresponding value for (II) was 2.1539 for 565 reflections having $F_o/F_c(\text{max})$ in the range $0.000 < F_o/F_c(\text{max}) < 0.009$.

Acknowledgements

MG thanks the UGC (India) for the award of a Rajeev Gandhi fellowship and HSY thanks the University of Mysore for research facilities.

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supporting information

Acta Cryst. (2017). E73, 1835-1839 [https://doi.org/10.1107/S2056989017015985]

Crystal structures of (*E*)-1-{3-[(5-fluoro-2-hydroxybenzylidene)amino]phenyl}-ethanone and of a fourth polymorph of (*E*)-1-{3-[(2-hydroxy-3-methoxybenzylidene)amino]phenyl}ethanone

Marisiddaiah Girisha, Hemmige S. Yathirajan, Ravindranath S. Rathore and Christopher Glidewell

Computing details

For both structures, data collection: *APEX3* (Bruker, 2016); cell refinement: *SAINTE* (Bruker, 2016); data reduction: *SAINTE* (Bruker, 2016); program(s) used to solve structure: *SHELXT2014* (Sheldrick, 2015a); program(s) used to refine structure: *SHELXL2014* (Sheldrick, 2015b); molecular graphics: *PLATON* (Spek, 2009); software used to prepare material for publication: *SHELXL2014* (Sheldrick, 2015b) and *PLATON* (Spek, 2009).

(*E*)-1-{3-[(5-Fluoro-2-hydroxybenzylidene)amino]phenyl}ethanone (I)

Crystal data

$C_{15}H_{12}FNO_2$

$M_r = 257.26$

Monoclinic, $P2_1/n$

$a = 14.9527$ (5) Å

$b = 5.5152$ (2) Å

$c = 16.6918$ (5) Å

$\beta = 114.739$ (2)°

$V = 1250.19$ (7) Å³

$Z = 4$

$F(000) = 536$

$D_x = 1.367$ Mg m⁻³

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

Cell parameters from 2452 reflections

$\theta = 3.3\text{--}72.5^\circ$

$\mu = 0.84$ mm⁻¹

$T = 294$ K

Block, yellow

$0.15 \times 0.15 \times 0.10$ mm

Data collection

Bruker APEX3
diffractometer

Radiation source: microfocus sealed tube

Multilayer mirror monochromator

φ and ω scans

Absorption correction: multi-scan
(SADABS; Bruker, 2016)

$T_{\min} = 0.848$, $T_{\max} = 0.919$

15703 measured reflections

2452 independent reflections

1764 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.041$

$\theta_{\max} = 72.5^\circ$, $\theta_{\min} = 3.3^\circ$

$h = -18 \rightarrow 18$

$k = -6 \rightarrow 6$

$l = -20 \rightarrow 20$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.046$

$wR(F^2) = 0.122$

$S = 1.03$

2452 reflections

176 parameters

0 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: mixed
H atoms treated by a mixture of independent
and constrained refinement

$$w = 1/[\sigma^2(F_o^2) + (0.0427P)^2 + 0.3677P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.15 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.12 \text{ e } \text{\AA}^{-3}$

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)

| | x | y | z | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|------|--------------|-------------|--------------|----------------------------------|
| N1 | 0.35562 (11) | 0.5843 (3) | 0.53906 (9) | 0.0566 (4) |
| C11 | 0.39602 (12) | 0.7592 (3) | 0.60710 (11) | 0.0529 (4) |
| C12 | 0.36799 (12) | 0.7897 (3) | 0.67598 (11) | 0.0539 (4) |
| H12 | 0.3219 | 0.6851 | 0.6812 | 0.065* |
| C13 | 0.40781 (12) | 0.9740 (3) | 0.73703 (11) | 0.0551 (4) |
| C14 | 0.47738 (14) | 1.1282 (4) | 0.72986 (13) | 0.0672 (5) |
| H14 | 0.5041 | 1.2531 | 0.7704 | 0.081* |
| C15 | 0.50686 (15) | 1.0962 (4) | 0.66268 (14) | 0.0750 (6) |
| H15 | 0.5540 | 1.1987 | 0.6583 | 0.090* |
| C16 | 0.46702 (14) | 0.9138 (4) | 0.60227 (12) | 0.0655 (5) |
| H16 | 0.4878 | 0.8932 | 0.5574 | 0.079* |
| C17 | 0.37489 (14) | 0.9972 (4) | 0.80988 (12) | 0.0635 (5) |
| O17 | 0.31869 (12) | 0.8495 (3) | 0.81717 (10) | 0.0859 (5) |
| C18 | 0.41085 (18) | 1.2043 (5) | 0.87234 (15) | 0.0900 (7) |
| H18A | 0.3840 | 1.1931 | 0.9153 | 0.135* |
| H18B | 0.3903 | 1.3539 | 0.8404 | 0.135* |
| H18C | 0.4814 | 1.1995 | 0.9016 | 0.135* |
| C27 | 0.29372 (13) | 0.4249 (3) | 0.53959 (10) | 0.0554 (4) |
| H27 | 0.2759 | 0.4216 | 0.5867 | 0.066* |
| C21 | 0.25071 (13) | 0.2505 (3) | 0.46968 (10) | 0.0541 (4) |
| C22 | 0.27440 (15) | 0.2459 (4) | 0.39638 (12) | 0.0632 (5) |
| O22 | 0.33757 (13) | 0.4080 (3) | 0.38812 (10) | 0.0858 (5) |
| H22 | 0.361 (2) | 0.512 (5) | 0.4408 (19) | 0.129* |
| C23 | 0.23177 (18) | 0.0728 (4) | 0.33148 (13) | 0.0774 (6) |
| H23 | 0.2473 | 0.0695 | 0.2830 | 0.093* |
| C24 | 0.16706 (17) | -0.0935 (4) | 0.33765 (13) | 0.0776 (6) |
| H24 | 0.1386 | -0.2094 | 0.2938 | 0.093* |
| C25 | 0.14467 (15) | -0.0872 (4) | 0.40930 (13) | 0.0701 (5) |
| F25 | 0.08044 (11) | -0.2522 (3) | 0.41531 (9) | 0.1050 (5) |
| C26 | 0.18467 (14) | 0.0810 (4) | 0.47482 (11) | 0.0631 (5) |
| H26 | 0.1678 | 0.0819 | 0.5225 | 0.076* |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|-------------|-------------|-------------|--------------|-------------|--------------|
| N1 | 0.0650 (9) | 0.0630 (9) | 0.0489 (8) | 0.0097 (7) | 0.0308 (7) | 0.0060 (7) |
| C11 | 0.0543 (9) | 0.0585 (10) | 0.0503 (9) | 0.0096 (8) | 0.0263 (7) | 0.0124 (8) |
| C12 | 0.0550 (9) | 0.0571 (10) | 0.0556 (9) | -0.0021 (8) | 0.0290 (8) | 0.0028 (8) |
| C13 | 0.0542 (9) | 0.0569 (10) | 0.0556 (10) | 0.0015 (8) | 0.0243 (8) | 0.0026 (8) |
| C14 | 0.0680 (12) | 0.0636 (11) | 0.0681 (11) | -0.0095 (9) | 0.0266 (9) | 0.0000 (9) |
| C15 | 0.0710 (13) | 0.0798 (14) | 0.0795 (13) | -0.0147 (11) | 0.0366 (11) | 0.0115 (12) |
| C16 | 0.0664 (11) | 0.0795 (13) | 0.0614 (11) | 0.0038 (10) | 0.0372 (9) | 0.0150 (10) |
| C17 | 0.0646 (11) | 0.0673 (12) | 0.0634 (11) | 0.0009 (10) | 0.0314 (9) | -0.0068 (9) |
| O17 | 0.1074 (11) | 0.0916 (11) | 0.0868 (10) | -0.0230 (9) | 0.0682 (9) | -0.0232 (8) |
| C18 | 0.0975 (16) | 0.0946 (17) | 0.0870 (15) | -0.0151 (14) | 0.0476 (13) | -0.0330 (13) |
| C27 | 0.0641 (10) | 0.0641 (11) | 0.0434 (8) | 0.0101 (9) | 0.0279 (8) | 0.0076 (8) |
| C21 | 0.0636 (10) | 0.0573 (10) | 0.0416 (8) | 0.0157 (8) | 0.0222 (7) | 0.0076 (7) |
| C22 | 0.0782 (12) | 0.0663 (11) | 0.0528 (10) | 0.0190 (10) | 0.0349 (9) | 0.0058 (9) |
| O22 | 0.1131 (12) | 0.0961 (11) | 0.0753 (9) | -0.0030 (9) | 0.0660 (9) | -0.0058 (8) |
| C23 | 0.1021 (16) | 0.0819 (14) | 0.0559 (11) | 0.0201 (13) | 0.0406 (11) | -0.0040 (11) |
| C24 | 0.0948 (15) | 0.0731 (13) | 0.0556 (11) | 0.0175 (12) | 0.0222 (10) | -0.0091 (10) |
| C25 | 0.0775 (13) | 0.0646 (12) | 0.0601 (11) | 0.0036 (10) | 0.0209 (10) | 0.0051 (10) |
| F25 | 0.1246 (11) | 0.1000 (10) | 0.0836 (9) | -0.0353 (9) | 0.0367 (8) | -0.0144 (8) |
| C26 | 0.0735 (12) | 0.0698 (12) | 0.0454 (9) | 0.0064 (10) | 0.0245 (8) | 0.0051 (9) |

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

| | | | |
|-------------|-------------|---------------|-------------|
| N1—C27 | 1.279 (2) | C18—H18B | 0.9600 |
| N1—C11 | 1.418 (2) | C18—H18C | 0.9600 |
| C11—C12 | 1.389 (2) | C27—C21 | 1.440 (2) |
| C11—C16 | 1.390 (2) | C27—H27 | 0.9300 |
| C12—C13 | 1.385 (2) | C21—C26 | 1.388 (3) |
| C12—H12 | 0.9300 | C21—C22 | 1.409 (2) |
| C13—C14 | 1.387 (2) | C22—O22 | 1.349 (2) |
| C13—C17 | 1.496 (2) | C22—C23 | 1.383 (3) |
| C14—C15 | 1.377 (3) | O22—H22 | 0.98 (3) |
| C14—H14 | 0.9300 | C23—C24 | 1.368 (3) |
| C15—C16 | 1.371 (3) | C23—H23 | 0.9300 |
| C15—H15 | 0.9300 | C24—C25 | 1.371 (3) |
| C16—H16 | 0.9300 | C24—H24 | 0.9300 |
| C17—O17 | 1.213 (2) | C25—F25 | 1.357 (2) |
| C17—C18 | 1.487 (3) | C25—C26 | 1.366 (3) |
| C18—H18A | 0.9600 | C26—H26 | 0.9300 |
| C27—N1—C11 | 121.93 (14) | C17—C18—H18C | 109.5 |
| C12—C11—C16 | 118.43 (17) | H18A—C18—H18C | 109.5 |
| C12—C11—N1 | 124.77 (15) | H18B—C18—H18C | 109.5 |
| C16—C11—N1 | 116.78 (15) | N1—C27—C21 | 122.19 (15) |
| C13—C12—C11 | 120.82 (16) | N1—C27—H27 | 118.9 |
| C13—C12—H12 | 119.6 | C21—C27—H27 | 118.9 |

| | | | |
|-----------------|--------------|-----------------|--------------|
| C11—C12—H12 | 119.6 | C26—C21—C22 | 119.03 (17) |
| C12—C13—C14 | 119.56 (16) | C26—C21—C27 | 119.20 (15) |
| C12—C13—C17 | 118.34 (16) | C22—C21—C27 | 121.76 (17) |
| C14—C13—C17 | 122.09 (17) | O22—C22—C23 | 119.35 (17) |
| C15—C14—C13 | 119.92 (19) | O22—C22—C21 | 121.21 (17) |
| C15—C14—H14 | 120.0 | C23—C22—C21 | 119.4 (2) |
| C13—C14—H14 | 120.0 | C22—O22—H22 | 107.5 (17) |
| C16—C15—C14 | 120.28 (18) | C24—C23—C22 | 120.82 (18) |
| C16—C15—H15 | 119.9 | C24—C23—H23 | 119.6 |
| C14—C15—H15 | 119.9 | C22—C23—H23 | 119.6 |
| C15—C16—C11 | 120.97 (17) | C23—C24—C25 | 119.2 (2) |
| C15—C16—H16 | 119.5 | C23—C24—H24 | 120.4 |
| C11—C16—H16 | 119.5 | C25—C24—H24 | 120.4 |
| O17—C17—C18 | 120.53 (17) | F25—C25—C26 | 118.85 (18) |
| O17—C17—C13 | 120.10 (17) | F25—C25—C24 | 119.1 (2) |
| C18—C17—C13 | 119.37 (18) | C26—C25—C24 | 122.1 (2) |
| C17—C18—H18A | 109.5 | C25—C26—C21 | 119.48 (17) |
| C17—C18—H18B | 109.5 | C25—C26—H26 | 120.3 |
| H18A—C18—H18B | 109.5 | C21—C26—H26 | 120.3 |
| | | | |
| C27—N1—C11—C12 | -5.5 (3) | C11—N1—C27—C21 | 178.53 (15) |
| C27—N1—C11—C16 | 176.20 (16) | N1—C27—C21—C26 | 179.41 (16) |
| C16—C11—C12—C13 | 1.8 (2) | N1—C27—C21—C22 | -0.1 (3) |
| N1—C11—C12—C13 | -176.54 (16) | C26—C21—C22—O22 | 179.49 (17) |
| C11—C12—C13—C14 | -0.7 (3) | C27—C21—C22—O22 | -1.0 (3) |
| C11—C12—C13—C17 | -179.86 (16) | C26—C21—C22—C23 | -0.2 (3) |
| C12—C13—C14—C15 | -0.5 (3) | C27—C21—C22—C23 | 179.31 (17) |
| C17—C13—C14—C15 | 178.60 (18) | O22—C22—C23—C24 | -179.71 (18) |
| C13—C14—C15—C16 | 0.6 (3) | C21—C22—C23—C24 | 0.0 (3) |
| C14—C15—C16—C11 | 0.4 (3) | C22—C23—C24—C25 | 0.0 (3) |
| C12—C11—C16—C15 | -1.6 (3) | C23—C24—C25—F25 | 179.98 (19) |
| N1—C11—C16—C15 | 176.83 (17) | C23—C24—C25—C26 | 0.3 (3) |
| C12—C13—C17—O17 | 3.8 (3) | F25—C25—C26—C21 | 179.80 (16) |
| C14—C13—C17—O17 | -175.26 (19) | C24—C25—C26—C21 | -0.5 (3) |
| C12—C13—C17—C18 | -175.50 (18) | C22—C21—C26—C25 | 0.5 (3) |
| C14—C13—C17—C18 | 5.4 (3) | C27—C21—C26—C25 | -179.09 (16) |

Hydrogen-bond geometry (Å, °)

| <i>D</i> —H... <i>A</i> | <i>D</i> —H | H... <i>A</i> | <i>D</i> ... <i>A</i> | <i>D</i> —H... <i>A</i> |
|----------------------------|-------------|---------------|-----------------------|-------------------------|
| O22—H22...N1 | 0.98 (3) | 1.72 (3) | 2.607 (2) | 148 (3) |
| C27—H27...O17 ⁱ | 0.93 | 2.58 | 3.475 (3) | 163 |

Symmetry code: (i) $-x+1/2, y-1/2, -z+3/2$.

(E)-1-[3-[(2-Hydroxy-3-methoxybenzylidene)amino]phenyl]ethanone (II)*Crystal data*

| | |
|----------------------------------|---|
| $C_{16}H_{15}NO_3$ | $D_x = 1.314 \text{ Mg m}^{-3}$ |
| $M_r = 269.29$ | Cu $K\alpha$ radiation, $\lambda = 1.54184 \text{ \AA}$ |
| Orthorhombic, $Pca2_1$ | Cell parameters from 5394 reflections |
| $a = 19.1904 (4) \text{ \AA}$ | $\theta = 3.3\text{--}72.6^\circ$ |
| $b = 5.33856 (12) \text{ \AA}$ | $\mu = 0.75 \text{ mm}^{-1}$ |
| $c = 26.5678 (6) \text{ \AA}$ | $T = 294 \text{ K}$ |
| $V = 2721.85 (10) \text{ \AA}^3$ | Block, yellow |
| $Z = 8$ | $0.10 \times 0.10 \times 0.05 \text{ mm}$ |
| $F(000) = 1136$ | |

Data collection

| | |
|---|--|
| Bruker APEX3 | 51776 measured reflections |
| diffractometer | 5393 independent reflections |
| Radiation source: microfocus sealed tube | 3796 reflections with $I > 2\sigma(I)$ |
| Multilayer mirror monochromator | $R_{\text{int}} = 0.117$ |
| φ and ω scans | $\theta_{\text{max}} = 72.6^\circ$, $\theta_{\text{min}} = 3.3^\circ$ |
| Absorption correction: multi-scan | $h = -23 \rightarrow 23$ |
| (SADABS; Bruker, 2016) | $k = -6 \rightarrow 6$ |
| $T_{\text{min}} = 0.907$, $T_{\text{max}} = 0.963$ | $l = -32 \rightarrow 32$ |

Refinement

| | |
|--|---|
| Refinement on F^2 | Hydrogen site location: mixed |
| Least-squares matrix: full | H atoms treated by a mixture of independent |
| $R[F^2 > 2\sigma(F^2)] = 0.048$ | and constrained refinement |
| $wR(F^2) = 0.113$ | $w = 1/[\sigma^2(F_o^2) + (0.0463P)^2 + 0.4368P]$ |
| $S = 1.02$ | where $P = (F_o^2 + 2F_c^2)/3$ |
| 5393 reflections | $(\Delta/\sigma)_{\text{max}} < 0.001$ |
| 371 parameters | $\Delta\rho_{\text{max}} = 0.11 \text{ e \AA}^{-3}$ |
| 1 restraint | $\Delta\rho_{\text{min}} = -0.13 \text{ e \AA}^{-3}$ |
| Primary atom site location: structure-invariant | Absolute structure: Flack x determined using |
| direct methods | 1493 quotients $[(I^+)-(I^-)]/[(I^+)+(I^-)]$ (Parsons <i>et al.</i> , 2013) |
| Secondary atom site location: difference Fourier | Absolute structure parameter: $-0.04 (16)$ |
| map | |

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)

| | x | y | z | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|------|--------------|------------|--------------|----------------------------------|
| N11 | 0.73276 (17) | 0.3158 (7) | 0.39187 (12) | 0.0502 (8) |
| C111 | 0.6779 (2) | 0.4896 (8) | 0.38707 (15) | 0.0466 (10) |
| C112 | 0.6379 (2) | 0.5213 (8) | 0.34388 (15) | 0.0475 (10) |
| H112 | 0.6453 | 0.4169 | 0.3164 | 0.057* |
| C113 | 0.5874 (2) | 0.7053 (8) | 0.34117 (15) | 0.0485 (10) |
| C114 | 0.5762 (3) | 0.8623 (9) | 0.38243 (18) | 0.0610 (11) |
| H114 | 0.5431 | 0.9893 | 0.3807 | 0.073* |

| | | | | |
|------|--------------|--------------|--------------|-------------|
| C115 | 0.6145 (3) | 0.8276 (10) | 0.42579 (18) | 0.0673 (13) |
| H115 | 0.6066 | 0.9297 | 0.4536 | 0.081* |
| C116 | 0.6641 (2) | 0.6437 (9) | 0.42814 (17) | 0.0613 (12) |
| H116 | 0.6891 | 0.6212 | 0.4578 | 0.074* |
| C117 | 0.5463 (2) | 0.7282 (9) | 0.29372 (17) | 0.0535 (11) |
| O117 | 0.55324 (18) | 0.5744 (7) | 0.26015 (13) | 0.0739 (10) |
| C118 | 0.4965 (3) | 0.9408 (11) | 0.28736 (19) | 0.0750 (15) |
| H11A | 0.4741 | 0.9280 | 0.2552 | 0.112* |
| H11B | 0.5214 | 1.0965 | 0.2893 | 0.112* |
| H11C | 0.4620 | 0.9346 | 0.3135 | 0.112* |
| C127 | 0.7464 (2) | 0.1542 (8) | 0.35777 (15) | 0.0495 (10) |
| H127 | 0.7185 | 0.1499 | 0.3292 | 0.059* |
| C121 | 0.8030 (2) | -0.0210 (8) | 0.36174 (14) | 0.0453 (10) |
| C122 | 0.8458 (2) | -0.0276 (8) | 0.40460 (14) | 0.0459 (9) |
| C123 | 0.8976 (2) | -0.2119 (9) | 0.40885 (15) | 0.0520 (11) |
| C124 | 0.9068 (2) | -0.3823 (9) | 0.37035 (16) | 0.0577 (11) |
| H124 | 0.9407 | -0.5059 | 0.3733 | 0.069* |
| C125 | 0.8660 (2) | -0.3715 (9) | 0.32720 (17) | 0.0584 (11) |
| H125 | 0.8737 | -0.4843 | 0.3011 | 0.070* |
| C126 | 0.8145 (2) | -0.1951 (8) | 0.32308 (16) | 0.0555 (11) |
| H126 | 0.7870 | -0.1906 | 0.2943 | 0.067* |
| O122 | 0.83809 (16) | 0.1383 (6) | 0.44236 (10) | 0.0596 (8) |
| H122 | 0.798 (3) | 0.267 (10) | 0.433 (2) | 0.089* |
| O123 | 0.93514 (16) | -0.2049 (7) | 0.45239 (12) | 0.0740 (10) |
| C128 | 0.9848 (3) | -0.3972 (10) | 0.4596 (2) | 0.0769 (15) |
| H12A | 0.9620 | -0.5571 | 0.4583 | 0.115* |
| H12B | 1.0068 | -0.3765 | 0.4917 | 0.115* |
| H12C | 1.0193 | -0.3886 | 0.4335 | 0.115* |
| N21 | 0.49089 (17) | -0.1872 (7) | 0.10711 (13) | 0.0520 (9) |
| C211 | 0.5456 (2) | -0.0084 (9) | 0.11255 (15) | 0.0477 (10) |
| C212 | 0.58545 (19) | 0.0190 (9) | 0.15570 (15) | 0.0486 (10) |
| H212 | 0.5781 | -0.0874 | 0.1829 | 0.058* |
| C213 | 0.6363 (2) | 0.2045 (8) | 0.15866 (15) | 0.0482 (10) |
| C214 | 0.6471 (2) | 0.3607 (9) | 0.11801 (17) | 0.0579 (11) |
| H214 | 0.6803 | 0.4875 | 0.1199 | 0.069* |
| C215 | 0.6087 (2) | 0.3286 (9) | 0.07452 (17) | 0.0634 (12) |
| H215 | 0.6170 | 0.4309 | 0.0468 | 0.076* |
| C216 | 0.5582 (2) | 0.1458 (9) | 0.07194 (17) | 0.0589 (12) |
| H216 | 0.5324 | 0.1262 | 0.0426 | 0.071* |
| C217 | 0.6772 (2) | 0.2248 (9) | 0.20644 (17) | 0.0540 (11) |
| O217 | 0.67061 (18) | 0.0695 (7) | 0.23932 (13) | 0.0769 (10) |
| C218 | 0.7275 (3) | 0.4370 (10) | 0.21282 (19) | 0.0759 (15) |
| H21A | 0.7023 | 0.5924 | 0.2138 | 0.114* |
| H21B | 0.7595 | 0.4390 | 0.1851 | 0.114* |
| H21C | 0.7528 | 0.4158 | 0.2437 | 0.114* |
| C227 | 0.4769 (2) | -0.3452 (9) | 0.14214 (16) | 0.0494 (11) |
| H227 | 0.5037 | -0.3431 | 0.1713 | 0.059* |
| C221 | 0.4212 (2) | -0.5259 (9) | 0.13805 (15) | 0.0483 (10) |

| | | | | |
|------|--------------|--------------|--------------|-------------|
| C222 | 0.3792 (2) | -0.5367 (9) | 0.09519 (15) | 0.0544 (11) |
| C223 | 0.3269 (2) | -0.7193 (10) | 0.09223 (16) | 0.0629 (13) |
| C224 | 0.3180 (3) | -0.8857 (9) | 0.13109 (18) | 0.0626 (13) |
| H224 | 0.2842 | -1.0098 | 0.1286 | 0.075* |
| C225 | 0.3588 (2) | -0.8708 (9) | 0.17392 (18) | 0.0600 (12) |
| H225 | 0.3517 | -0.9833 | 0.2002 | 0.072* |
| C226 | 0.4092 (2) | -0.6926 (8) | 0.17789 (17) | 0.0553 (11) |
| H226 | 0.4358 | -0.6813 | 0.2071 | 0.066* |
| O222 | 0.38787 (19) | -0.3768 (8) | 0.05600 (12) | 0.0797 (12) |
| H222 | 0.425 (3) | -0.274 (12) | 0.063 (2) | 0.119* |
| O223 | 0.28893 (19) | -0.7128 (8) | 0.04865 (12) | 0.0957 (14) |
| C228 | 0.2377 (3) | -0.9052 (14) | 0.0424 (2) | 0.109 (2) |
| H22A | 0.2603 | -1.0655 | 0.0410 | 0.163* |
| H22B | 0.2125 | -0.8774 | 0.0117 | 0.163* |
| H22C | 0.2059 | -0.9018 | 0.0703 | 0.163* |

Atomic displacement parameters (Å²)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|------|-------------|-----------|-------------|--------------|--------------|--------------|
| N11 | 0.0449 (19) | 0.061 (2) | 0.0451 (19) | -0.0023 (18) | -0.0030 (15) | 0.0026 (18) |
| C111 | 0.038 (2) | 0.052 (3) | 0.050 (2) | -0.003 (2) | 0.0001 (16) | 0.0036 (19) |
| C112 | 0.051 (3) | 0.048 (2) | 0.044 (2) | 0.001 (2) | -0.0012 (17) | 0.0011 (18) |
| C113 | 0.046 (2) | 0.046 (2) | 0.054 (2) | -0.0002 (19) | 0.0016 (18) | 0.004 (2) |
| C114 | 0.062 (3) | 0.054 (3) | 0.067 (3) | 0.008 (2) | 0.003 (2) | -0.008 (2) |
| C115 | 0.071 (3) | 0.072 (3) | 0.059 (3) | 0.002 (3) | 0.002 (2) | -0.024 (2) |
| C116 | 0.060 (3) | 0.075 (3) | 0.049 (2) | -0.007 (3) | -0.005 (2) | -0.007 (2) |
| C117 | 0.050 (2) | 0.051 (3) | 0.060 (3) | 0.006 (2) | 0.001 (2) | 0.005 (2) |
| O117 | 0.090 (2) | 0.069 (2) | 0.063 (2) | 0.0239 (19) | -0.0219 (17) | -0.0097 (19) |
| C118 | 0.077 (3) | 0.075 (4) | 0.074 (3) | 0.024 (3) | -0.004 (3) | 0.004 (3) |
| C127 | 0.045 (2) | 0.056 (3) | 0.047 (2) | -0.004 (2) | -0.0084 (18) | 0.005 (2) |
| C121 | 0.043 (2) | 0.048 (2) | 0.045 (2) | -0.008 (2) | -0.0013 (17) | 0.006 (2) |
| C122 | 0.045 (2) | 0.053 (2) | 0.040 (2) | -0.006 (2) | 0.0003 (16) | 0.0062 (18) |
| C123 | 0.043 (2) | 0.065 (3) | 0.047 (2) | -0.003 (2) | -0.0026 (18) | 0.011 (2) |
| C124 | 0.049 (2) | 0.062 (3) | 0.062 (3) | 0.004 (2) | 0.004 (2) | 0.010 (2) |
| C125 | 0.059 (3) | 0.057 (3) | 0.059 (3) | -0.004 (2) | 0.001 (2) | -0.002 (2) |
| C126 | 0.052 (2) | 0.062 (3) | 0.053 (3) | -0.008 (2) | -0.007 (2) | -0.001 (2) |
| O122 | 0.0591 (18) | 0.075 (2) | 0.0451 (16) | 0.0062 (16) | -0.0082 (14) | -0.0004 (15) |
| O123 | 0.069 (2) | 0.097 (3) | 0.0564 (19) | 0.024 (2) | -0.0184 (17) | 0.0036 (18) |
| C128 | 0.065 (3) | 0.089 (4) | 0.077 (4) | 0.016 (3) | -0.018 (3) | 0.014 (3) |
| N21 | 0.0439 (19) | 0.062 (2) | 0.050 (2) | -0.0086 (18) | -0.0011 (15) | -0.0127 (18) |
| C211 | 0.042 (2) | 0.053 (3) | 0.048 (2) | 0.003 (2) | -0.0002 (17) | -0.0074 (19) |
| C212 | 0.041 (2) | 0.058 (3) | 0.046 (2) | -0.001 (2) | 0.0011 (17) | -0.0033 (19) |
| C213 | 0.044 (2) | 0.053 (2) | 0.048 (2) | 0.003 (2) | -0.0008 (17) | -0.0063 (19) |
| C214 | 0.053 (3) | 0.055 (3) | 0.065 (3) | -0.006 (2) | 0.001 (2) | 0.001 (2) |
| C215 | 0.065 (3) | 0.063 (3) | 0.063 (3) | -0.002 (3) | -0.004 (2) | 0.010 (2) |
| C216 | 0.054 (3) | 0.069 (3) | 0.054 (3) | 0.000 (2) | -0.009 (2) | -0.002 (2) |
| C217 | 0.052 (2) | 0.055 (3) | 0.054 (3) | -0.007 (2) | -0.003 (2) | -0.012 (2) |
| O217 | 0.088 (3) | 0.078 (2) | 0.064 (2) | -0.023 (2) | -0.0262 (18) | 0.005 (2) |

| | | | | | | |
|------|-----------|-----------|-------------|------------|--------------|------------|
| C218 | 0.082 (4) | 0.078 (3) | 0.067 (3) | -0.031 (3) | -0.010 (3) | -0.011 (3) |
| C227 | 0.038 (2) | 0.059 (3) | 0.051 (3) | 0.006 (2) | -0.0052 (18) | -0.008 (2) |
| C221 | 0.039 (2) | 0.058 (3) | 0.047 (2) | 0.002 (2) | 0.0019 (17) | -0.010 (2) |
| C222 | 0.047 (2) | 0.073 (3) | 0.043 (2) | -0.015 (2) | 0.0048 (17) | -0.007 (2) |
| C223 | 0.055 (3) | 0.089 (4) | 0.044 (2) | -0.021 (3) | 0.005 (2) | -0.015 (2) |
| C224 | 0.060 (3) | 0.067 (3) | 0.061 (3) | -0.016 (2) | 0.012 (2) | -0.010 (2) |
| C225 | 0.061 (3) | 0.058 (3) | 0.061 (3) | -0.001 (2) | 0.005 (2) | 0.004 (2) |
| C226 | 0.052 (2) | 0.059 (3) | 0.054 (2) | 0.007 (2) | -0.007 (2) | -0.001 (2) |
| O222 | 0.079 (2) | 0.117 (3) | 0.0437 (17) | -0.045 (2) | -0.0040 (16) | 0.004 (2) |
| O223 | 0.093 (3) | 0.143 (4) | 0.0508 (18) | -0.067 (3) | -0.0126 (18) | 0.000 (2) |
| C228 | 0.097 (4) | 0.151 (6) | 0.078 (4) | -0.070 (4) | -0.010 (3) | -0.022 (4) |

Geometric parameters (Å, °)

| | | | |
|-----------|-----------|-----------|-----------|
| N11—C127 | 1.278 (5) | N21—C227 | 1.285 (5) |
| N11—C111 | 1.409 (5) | N21—C211 | 1.427 (5) |
| C111—C112 | 1.391 (5) | C211—C216 | 1.378 (6) |
| C111—C116 | 1.392 (6) | C211—C212 | 1.386 (5) |
| C112—C113 | 1.382 (6) | C212—C213 | 1.393 (6) |
| C112—H112 | 0.9300 | C212—H212 | 0.9300 |
| C113—C114 | 1.397 (6) | C213—C214 | 1.380 (6) |
| C113—C117 | 1.492 (6) | C213—C217 | 1.497 (6) |
| C114—C115 | 1.379 (7) | C214—C215 | 1.381 (6) |
| C114—H114 | 0.9300 | C214—H214 | 0.9300 |
| C115—C116 | 1.369 (6) | C215—C216 | 1.378 (6) |
| C115—H115 | 0.9300 | C215—H215 | 0.9300 |
| C116—H116 | 0.9300 | C216—H216 | 0.9300 |
| C117—O117 | 1.219 (5) | C217—O217 | 1.211 (5) |
| C117—C118 | 1.494 (7) | C217—C218 | 1.497 (6) |
| C118—H11A | 0.9600 | C218—H21A | 0.9600 |
| C118—H11B | 0.9600 | C218—H21B | 0.9600 |
| C118—H11C | 0.9600 | C218—H21C | 0.9600 |
| C127—C121 | 1.437 (5) | C227—C221 | 1.443 (6) |
| C127—H127 | 0.9300 | C227—H227 | 0.9300 |
| C121—C126 | 1.403 (6) | C221—C222 | 1.397 (6) |
| C121—C122 | 1.405 (5) | C221—C226 | 1.402 (6) |
| C122—O122 | 1.346 (5) | C222—O222 | 1.357 (6) |
| C122—C123 | 1.403 (6) | C222—C223 | 1.401 (6) |
| C123—O123 | 1.363 (5) | C223—O223 | 1.369 (5) |
| C123—C124 | 1.380 (6) | C223—C224 | 1.373 (7) |
| C124—C125 | 1.389 (6) | C224—C225 | 1.384 (7) |
| C124—H124 | 0.9300 | C224—H224 | 0.9300 |
| C125—C126 | 1.369 (6) | C225—C226 | 1.361 (6) |
| C125—H125 | 0.9300 | C225—H225 | 0.9300 |
| C126—H126 | 0.9300 | C226—H226 | 0.9300 |
| O122—H122 | 1.06 (5) | O222—H222 | 0.93 (6) |
| O123—C128 | 1.414 (5) | O223—C228 | 1.432 (6) |
| C128—H12A | 0.9600 | C228—H22A | 0.9600 |

| | | | |
|----------------|-----------|----------------|-----------|
| C128—H12B | 0.9600 | C228—H22B | 0.9600 |
| C128—H12C | 0.9600 | C228—H22C | 0.9600 |
| C127—N11—C111 | 122.2 (4) | C227—N21—C211 | 121.3 (4) |
| C112—C111—C116 | 118.0 (4) | C216—C211—C212 | 119.2 (4) |
| C112—C111—N11 | 124.5 (4) | C216—C211—N21 | 116.7 (4) |
| C116—C111—N11 | 117.4 (4) | C212—C211—N21 | 124.1 (4) |
| C113—C112—C111 | 121.1 (4) | C211—C212—C213 | 120.5 (4) |
| C113—C112—H112 | 119.4 | C211—C212—H212 | 119.7 |
| C111—C112—H112 | 119.4 | C213—C212—H212 | 119.7 |
| C112—C113—C114 | 119.6 (4) | C214—C213—C212 | 119.4 (4) |
| C112—C113—C117 | 118.2 (4) | C214—C213—C217 | 122.7 (4) |
| C114—C113—C117 | 122.2 (4) | C212—C213—C217 | 117.9 (4) |
| C115—C114—C113 | 119.5 (4) | C213—C214—C215 | 120.0 (4) |
| C115—C114—H114 | 120.3 | C213—C214—H214 | 120.0 |
| C113—C114—H114 | 120.3 | C215—C214—H214 | 120.0 |
| C116—C115—C114 | 120.4 (4) | C216—C215—C214 | 120.3 (4) |
| C116—C115—H115 | 119.8 | C216—C215—H215 | 119.8 |
| C114—C115—H115 | 119.8 | C214—C215—H215 | 119.8 |
| C115—C116—C111 | 121.4 (4) | C215—C216—C211 | 120.5 (4) |
| C115—C116—H116 | 119.3 | C215—C216—H216 | 119.8 |
| C111—C116—H116 | 119.3 | C211—C216—H216 | 119.8 |
| O117—C117—C113 | 120.4 (4) | O217—C217—C213 | 120.5 (4) |
| O117—C117—C118 | 119.9 (4) | O217—C217—C218 | 120.3 (4) |
| C113—C117—C118 | 119.7 (4) | C213—C217—C218 | 119.3 (4) |
| C117—C118—H11A | 109.5 | C217—C218—H21A | 109.5 |
| C117—C118—H11B | 109.5 | C217—C218—H21B | 109.5 |
| H11A—C118—H11B | 109.5 | H21A—C218—H21B | 109.5 |
| C117—C118—H11C | 109.5 | C217—C218—H21C | 109.5 |
| H11A—C118—H11C | 109.5 | H21A—C218—H21C | 109.5 |
| H11B—C118—H11C | 109.5 | H21B—C218—H21C | 109.5 |
| N11—C127—C121 | 122.8 (4) | N21—C227—C221 | 122.6 (4) |
| N11—C127—H127 | 118.6 | N21—C227—H227 | 118.7 |
| C121—C127—H127 | 118.6 | C221—C227—H227 | 118.7 |
| C126—C121—C122 | 119.0 (4) | C222—C221—C226 | 119.7 (4) |
| C126—C121—C127 | 119.8 (4) | C222—C221—C227 | 121.1 (4) |
| C122—C121—C127 | 121.2 (4) | C226—C221—C227 | 119.3 (4) |
| O122—C122—C123 | 118.6 (3) | O222—C222—C221 | 122.0 (4) |
| O122—C122—C121 | 121.5 (4) | O222—C222—C223 | 118.8 (4) |
| C123—C122—C121 | 119.8 (4) | C221—C222—C223 | 119.2 (4) |
| O123—C123—C124 | 125.4 (4) | O223—C223—C224 | 125.9 (4) |
| O123—C123—C122 | 115.1 (4) | O223—C223—C222 | 114.3 (4) |
| C124—C123—C122 | 119.5 (4) | C224—C223—C222 | 119.8 (4) |
| C123—C124—C125 | 120.8 (4) | C223—C224—C225 | 120.7 (4) |
| C123—C124—H124 | 119.6 | C223—C224—H224 | 119.6 |
| C125—C124—H124 | 119.6 | C225—C224—H224 | 119.6 |
| C126—C125—C124 | 120.1 (4) | C226—C225—C224 | 120.4 (4) |
| C126—C125—H125 | 120.0 | C226—C225—H225 | 119.8 |

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|---------------------|------------|---------------------|------------|
| C124—C125—H125 | 120.0 | C224—C225—H225 | 119.8 |
| C125—C126—C121 | 120.8 (4) | C225—C226—C221 | 120.1 (4) |
| C125—C126—H126 | 119.6 | C225—C226—H226 | 119.9 |
| C121—C126—H126 | 119.6 | C221—C226—H226 | 119.9 |
| C122—O122—H122 | 109 (3) | C222—O222—H222 | 108 (4) |
| C123—O123—C128 | 116.8 (4) | C223—O223—C228 | 116.5 (4) |
| O123—C128—H12A | 109.5 | O223—C228—H22A | 109.5 |
| O123—C128—H12B | 109.5 | O223—C228—H22B | 109.5 |
| H12A—C128—H12B | 109.5 | H22A—C228—H22B | 109.5 |
| O123—C128—H12C | 109.5 | O223—C228—H22C | 109.5 |
| H12A—C128—H12C | 109.5 | H22A—C228—H22C | 109.5 |
| H12B—C128—H12C | 109.5 | H22B—C228—H22C | 109.5 |
| | | | |
| C127—N11—C111—C112 | -6.6 (6) | C227—N21—C211—C216 | 177.4 (4) |
| C127—N11—C111—C116 | 174.9 (4) | C227—N21—C211—C212 | -3.2 (6) |
| C116—C111—C112—C113 | 2.0 (6) | C216—C211—C212—C213 | 1.8 (6) |
| N11—C111—C112—C113 | -176.5 (4) | N21—C211—C212—C213 | -177.6 (4) |
| C111—C112—C113—C114 | -0.1 (6) | C211—C212—C213—C214 | -0.4 (6) |
| C111—C112—C113—C117 | -179.9 (4) | C211—C212—C213—C217 | 179.7 (4) |
| C112—C113—C114—C115 | -1.6 (7) | C212—C213—C214—C215 | -1.5 (7) |
| C117—C113—C114—C115 | 178.2 (4) | C217—C213—C214—C215 | 178.5 (4) |
| C113—C114—C115—C116 | 1.2 (8) | C213—C214—C215—C216 | 1.8 (7) |
| C114—C115—C116—C111 | 0.9 (7) | C214—C215—C216—C211 | -0.4 (7) |
| C112—C111—C116—C115 | -2.4 (6) | C212—C211—C216—C215 | -1.4 (7) |
| N11—C111—C116—C115 | 176.2 (4) | N21—C211—C216—C215 | 178.1 (4) |
| C112—C113—C117—O117 | 6.3 (6) | C214—C213—C217—O217 | -173.1 (4) |
| C114—C113—C117—O117 | -173.5 (4) | C212—C213—C217—O217 | 6.9 (6) |
| C112—C113—C117—C118 | -173.7 (4) | C214—C213—C217—C218 | 5.9 (7) |
| C114—C113—C117—C118 | 6.5 (7) | C212—C213—C217—C218 | -174.2 (4) |
| C111—N11—C127—C121 | 178.8 (4) | C211—N21—C227—C221 | 179.1 (4) |
| N11—C127—C121—C126 | 179.4 (4) | N21—C227—C221—C222 | -0.2 (6) |
| N11—C127—C121—C122 | 1.6 (6) | N21—C227—C221—C226 | 179.9 (4) |
| C126—C121—C122—O122 | 178.5 (4) | C226—C221—C222—O222 | 178.8 (4) |
| C127—C121—C122—O122 | -3.7 (6) | C227—C221—C222—O222 | -1.0 (7) |
| C126—C121—C122—C123 | -2.1 (6) | C226—C221—C222—C223 | -1.8 (6) |
| C127—C121—C122—C123 | 175.7 (4) | C227—C221—C222—C223 | 178.4 (4) |
| O122—C122—C123—O123 | 0.8 (6) | O222—C222—C223—O223 | -0.6 (7) |
| C121—C122—C123—O123 | -178.6 (4) | C221—C222—C223—O223 | 180.0 (4) |
| O122—C122—C123—C124 | -179.4 (4) | O222—C222—C223—C224 | 178.9 (5) |
| C121—C122—C123—C124 | 1.2 (6) | C221—C222—C223—C224 | -0.5 (7) |
| O123—C123—C124—C125 | -179.4 (4) | O223—C223—C224—C225 | -178.7 (5) |
| C122—C123—C124—C125 | 0.9 (7) | C222—C223—C224—C225 | 1.9 (7) |
| C123—C124—C125—C126 | -2.0 (7) | C223—C224—C225—C226 | -1.0 (7) |
| C124—C125—C126—C121 | 1.0 (7) | C224—C225—C226—C221 | -1.4 (7) |
| C122—C121—C126—C125 | 1.0 (6) | C222—C221—C226—C225 | 2.8 (7) |
| C127—C121—C126—C125 | -176.8 (4) | C227—C221—C226—C225 | -177.4 (4) |
| C124—C123—O123—C128 | -3.7 (7) | C224—C223—O223—C228 | -3.2 (8) |
| C122—C123—O123—C128 | 176.1 (4) | C222—C223—O223—C228 | 176.2 (5) |

Hydrogen-bond geometry (Å, °)

| <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> |
|---------------------------------------|-------------|---------------------|----------------------------|-------------------------------|
| O122—H122 \cdots N11 | 1.06 (6) | 1.68 (6) | 2.604 (4) | 142 (5) |
| O222—H222 \cdots N21 | 0.92 (6) | 1.79 (6) | 2.603 (5) | 147 (5) |
| C116—H116 \cdots O223 ⁱ | 0.93 | 2.50 | 3.347 (6) | 152 |
| C127—H127 \cdots O217 | 0.93 | 2.59 | 3.496 (5) | 164 |
| C227—H227 \cdots O117 ⁱⁱ | 0.93 | 2.58 | 3.487 (5) | 164 |

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