

Intermolecular C—H···O and C—H···X interactions in substituted spiroacenaphthylene structures

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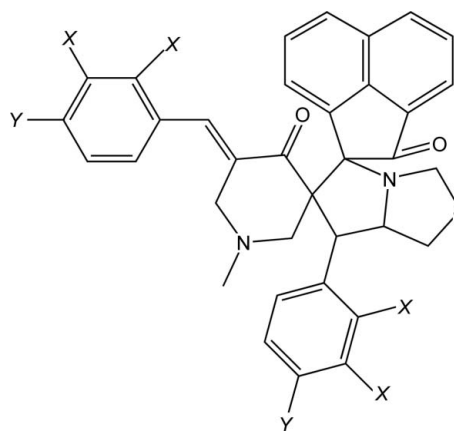
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In the three spiroacenaphthylene structures 5''-[(*E*)-2,3-dichlorobenzylidene]-7'-(2,3-dichlorophenyl)-1''-methyl-dispiro[acenaphthylene-1,5'-pyrrolo[1,2-*c*][1,3]thiazole-6',3''-piperidine]-2,4''-dione, C₃₅H₂₆Cl₄N₂O₂S, (I), 5''-[(*E*)-4-fluorobenzylidene]-7'-(4-fluorophenyl)-1''-methyl-dispiro[acenaphthylene-1,5'-pyrrolo[1,2-*c*][1,3]thiazole-6',3''-piperidine]-2,4''-dione, C₃₅H₂₈F₂N₂O₂S, (II), and 5''-[(*E*)-4-bromobenzylidene]-7'-(4-bromophenyl)-1''-methyl-dispiro[acenaphthylene-1,5'-pyrrolo[1,2-*c*][1,3]thiazole-6',3''-piperidine]-2,4''-dione, C₃₅H₂₈Br₂N₂O₂S, (III), the substituted aryl groups are 2,3-dichloro-, 4-fluoro- and 4-bromophenyl, respectively. The six-membered piperidine ring in all three structures adopts a half-chair conformation, the thiazolidine ring adopts a slightly twisted envelope and the pyrrolidine ring an envelope conformation; in each case, the C atom linking the rings is the flap atom. In all three structures, weak intramolecular C—H···O interactions are present. The crystal packing is stabilized through a number of intermolecular C—H···O and C—H···X interactions, where X = Cl in (I) and F or S in (II), and C—H···O interactions are observed predominantly in (III). In all three structures, molecules are linked through centrosymmetric ring motifs, further tailored through a relay of C—H···X [Cl in (I), Br in (II) and O in (III)] interactions.

Comment

The piperidine ring is a distinct structural feature in a variety of alkaloid natural products and drug candidates. During the past decade, thousands of piperidine compounds have been reported in clinical and preclinical studies (Watson *et al.*, 2000). Piperidinones, though less prominent, are also precursors to a host of biologically active compounds and natural alkaloids, prior to their conversion to piperidines. Spiro systems are also of interest as they exhibit a wide range of biological activities (Kobayashi *et al.*, 1991; James *et al.*, 1991).

Spiropyrrolidines have attracted much attention as potential antileukaemic and anticonvulsant agents (Abou Gharbia *et al.*, 1979), with antiviral (Lundahl *et al.*, 1972) and local anaesthetic (Kornett & Thio, 1976) activities. Our interest in preparing pharmacologically active piperidones led us to the title 5''-arylmethyl-7'-aryl-1''-methyl-dispiro[acenaphthylene-1,5'-pyrrolo[1,2-*c*][1,3]thiazole-6',3''-piperidine]-2,4''-dione compounds, where aryl is 2,3-dichlorophenyl in (I), 4-fluorophenyl in (II) and 4-bromophenyl in (III), and especially the study of their conformational features. There are few structures in the Cambridge Structural Database (CSD, Version 5.26; Allen, 2002) containing pyrrolothiazine ring systems and this is the first report with the attached five-membered ring. Hence, these structures are presumed to be interesting and contain rarely studied moieties.



(I) X = Cl, Y = H
(II) X = H, Y = F
(III) X = H, Y = Br

The six-membered piperidine rings in all three compounds (Figs. 1–3) adopt the half-chair conformation, with deviations of atoms N1 and C5 from the least-squares plane defined by atoms C2/C3/C4/C6 of 0.613 (3) and 0.509 (3) Å in (I),

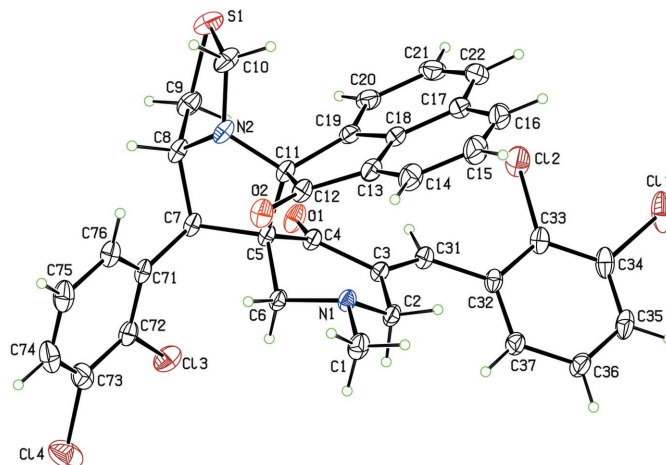


Figure 1
The molecular structure of (I), showing 30% probability displacement ellipsoids and the atom-numbering scheme.

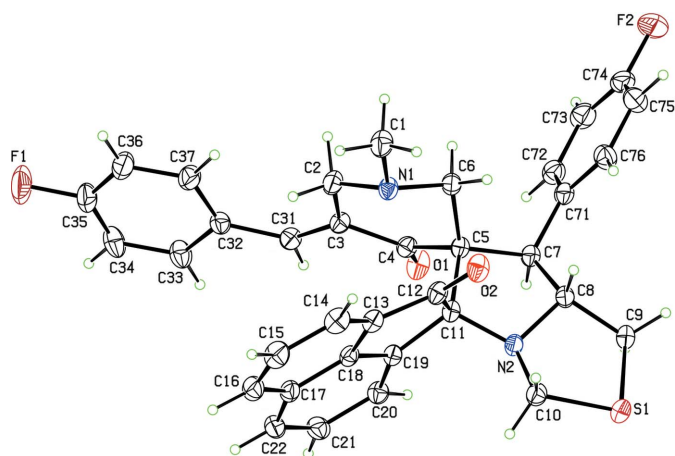


Figure 2
The molecular structure of (II), showing 30% probability displacement ellipsoids and the atom-numbering scheme.

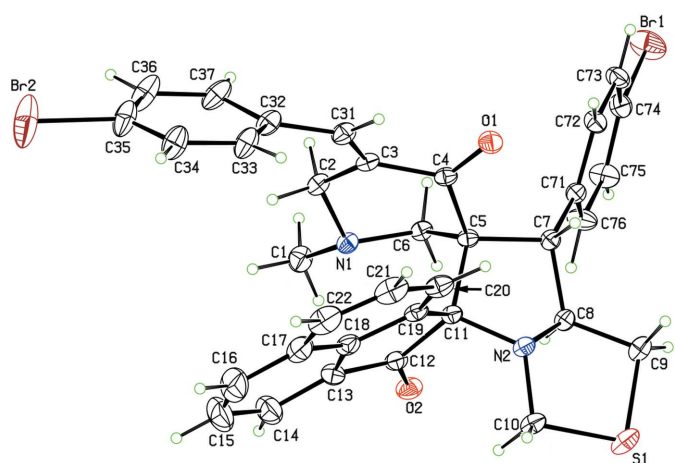


Figure 3
The molecular structure of (III), showing 30% probability displacement ellipsoids and the atom-numbering scheme.

0.553 (3) and 0.562 (3) Å in (II), and 0.545 (3) and 0.565 (3) Å in (III), respectively. The differences in the deviations are due to steric hindrance of the different substituents at the C3 and C5 positions of the piperidine ring. The olefinic double bond in all three structures has an *E* conformation and the pyrrolidine ring is in an envelope conformation. The thiazolidine ring is in a twisted envelope conformation with atom C8 at the flap, with a pseudo-twofold axis passing through atom S1 and the C8–N2 bond. The puckering parameters (Cremer & Pople, 1975) and the smallest displacement asymmetry parameters (Nardelli, 1983) are $q_2 = 0.356$ (2) Å, $\varphi = -78.0$ (4)° and $\Delta s(\text{C8}) = 0.035$ (1) Å for the thiazolidine ring, and $q_2 = 0.263$ (2) Å, $\varphi = -178.2$ (4)° and $\Delta s(\text{C8}) = 0.010$ (1) Å for the pyrrolidine ring of (I); $q_2 = 0.419$ (2) Å, $\varphi = -93.3$ (3)° and $\Delta s(\text{S1}) = 0.016$ (1) Å for the thiazolidine ring, and $q_2 = 0.456$ (2) Å, $\varphi = -0.4$ (3)° and $\Delta s(\text{C8}) = 0.001$ (1) Å for the pyrrolidine ring of (II); and $q_2 = 0.440$ (3) Å, $\varphi = -75.0$ (4)° and $\Delta s(\text{C8}) = 0.029$ (1) Å for the thiazolidine ring, and $q_2 = 0.460$ (3) Å, $\varphi = 144.6$ (4)° and $\Delta s(\text{C8}) = 0.011$ (1) Å for the pyrrolidine ring of (III).

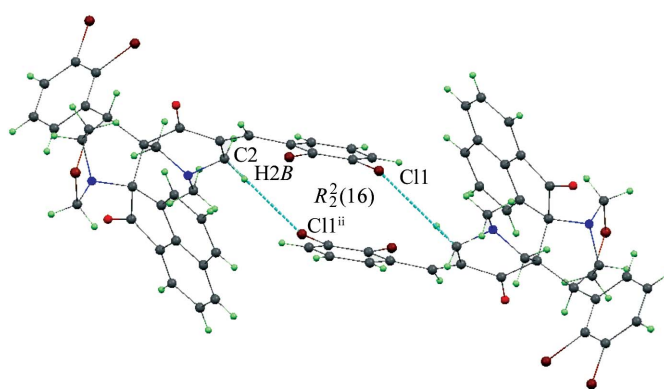


Figure 4
A partial packing view for (I), showing the $R_2^2(16)$ ring. Dashed lines indicate hydrogen bonds. [Symmetry code: (ii) $-x + 1, -y + 1, -z$]

The aryl rings in all three structures are not coplanar with the mean plane of the piperidone ring; the torsion angle C3–C31–C32–C37 is 32.4 (3)° in (I), 9.6 (4)° in (II) and -44.0 (5)° in (III). This lack of coplanarity is caused by nonbonded interactions between one of the *ortho* H atoms in the aryl ring and the equatorial H atoms at the 2-position of the piperidone ring (H37/H2A or H2B). Steric repulsions are reduced by the expansion of the C3–C31–C32 angle [129.57 (19)° in (I), 132.7 (2)° in (II) and 127.1 (3)° in (III)]. The dihedral angle between the dichlorophenyl rings is 58.4 (1)° in (I), that between the fluorophenyl rings is 87.2 (1)° in (II) and that between the bromophenyl rings is 86.1 (1)° in (III), and these rings make angles of 56.5 (1) and 65.2 (1)° in (I), 45.4 (1) and 43.1 (1)° in (II), and 48.7 (1) and 48.4 (1)° in (III) with their respective acenaphthylene group.

The C–C bond lengths and C–C–C angles in the acenaphthylene groups of (I)–(III) compare with those of related structures (Hazell, 1976; Hazell & Hazell, 1977; Hazell & Weigelt, 1976; Jones *et al.*, 1992; Sundar *et al.*, 2002). The C8–N2 bond length is 1.451 (3) Å in (I), 1.448 (3) Å in (II) and 1.455 (3) Å in (III), and these values are comparable with the $\text{Csp}^2\text{–Nsp}^2$ distances found in similar structures (Sussman & Wodak, 1973; Wodak, 1975). All three title structures feature a weak intramolecular C–H...O interaction linking pyrrolidine and piperidine rings.

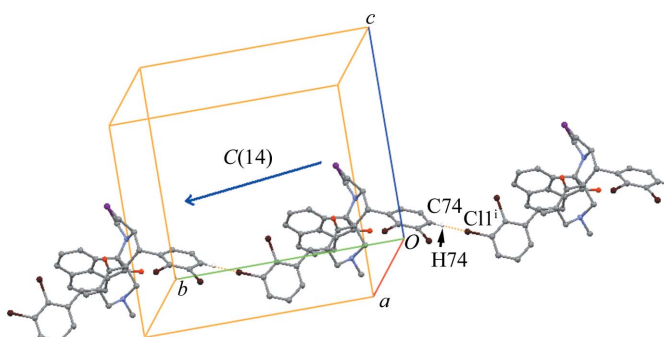
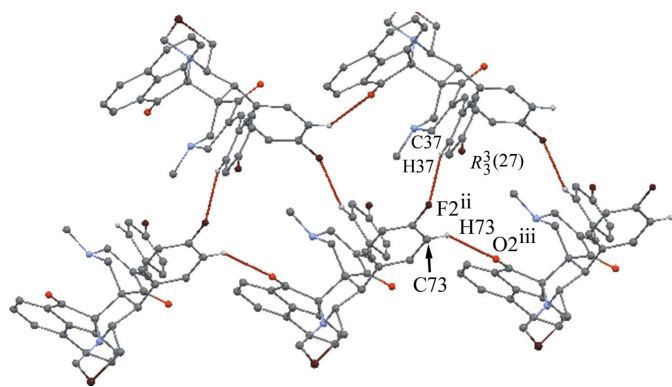


Figure 5
A partial packing view for (I), showing the C(14) chain extending along the *b* axis of the unit cell. Dashed lines indicate hydrogen bonds. Some H atoms have been omitted for clarity. [Symmetry code: (i) $x, y - 1, z$]

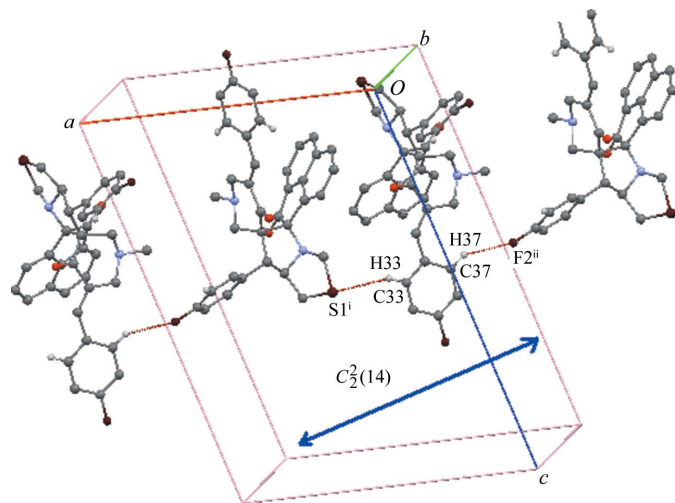
**Figure 6**

A partial packing view for (II), showing the unusual $R_3^2(27)$ ring. Dashed lines indicate hydrogen bonds. Some H atoms have been omitted for clarity. [Symmetry codes: (ii) $-x - \frac{1}{2}, y + \frac{1}{2}, -z + \frac{1}{2}$; (iii) $x, y - 1, z$.]

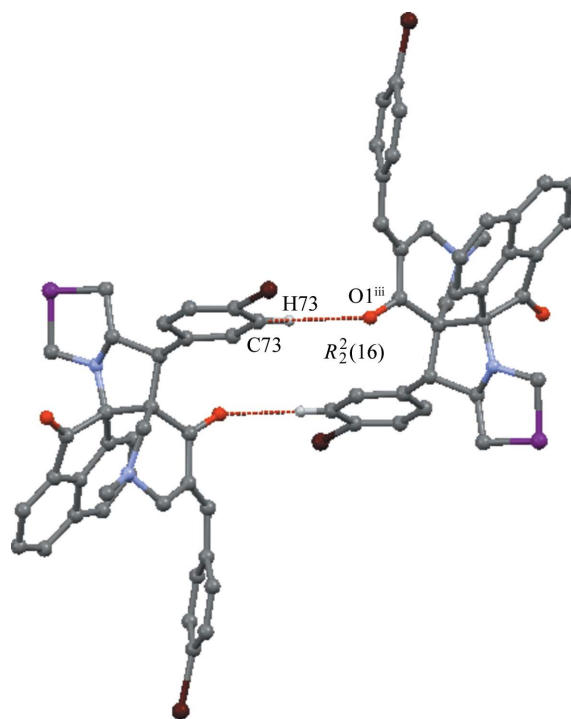
In all three structures, due to the lack of suitable donor and acceptor atoms for classical hydrogen bonding, the crystal structures are stabilized through intermolecular C—H \cdots X interactions [$X = \text{Cl}$ in (I), F in (II) and O in (III)]. These C—H \cdots X interactions lead to primary ring motifs around inversion centres, which are further connected through chain motifs (Bernstein *et al.*, 1995).

In (I), the molecules are connected through a C2—H2B \cdots Cl1ⁱⁱ interaction, leading to an $R_2^2(16)$ ring around the inversion centres of the unit cell (Fig. 4), and this ring, translated by a unit cell along the *b* axis connected by an intermolecular C74—H74 \cdots Cl1ⁱ interaction (see Table 1 for symmetry codes), leads to a $C(14)$ chain extending along the *b* axis (Fig. 5). These chains or rings do not have linking C—H \cdots O or C—H \cdots π interactions.

In (II), one C73—H73 \cdots O2ⁱⁱⁱ and two C37—H37 \cdots F2ⁱⁱ interactions lead to an unusual $R_3^2(27)$ ring (Fig. 6), which is connected through a secondary $C_2^2(14)$ chain extending along

**Figure 7**

A partial packing view for (II), showing the $C_2^2(14)$ chain extending along the *ab* diagonal of the unit cell. Dashed lines indicate hydrogen bonds. Some H atoms have been omitted for clarity. [Symmetry codes: (i) $-x + \frac{1}{2}, y - \frac{1}{2}, -z + \frac{1}{2}$; (ii) $-x - \frac{1}{2}, y + \frac{1}{2}, -z + \frac{1}{2}$.]

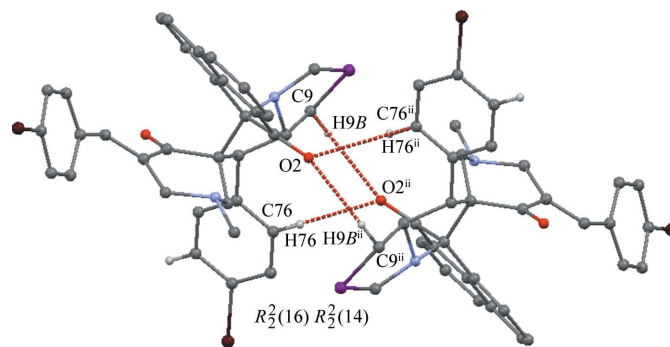
**Figure 8**

A partial packing view for (III), showing the $R_2^2(16)$ ring. Dashed lines indicate hydrogen bonds. Some H atoms have been omitted for clarity. [Symmetry code: (iii) $-x + 1, -y + 1, -z + 1$.]

the *ab* diagonal of the unit cell (Fig. 7), formed through C37—H37 \cdots F2ⁱⁱ and C33—H33 \cdots S1ⁱ interactions (see Table 2 for symmetry codes).

In (III), all the intermolecular interactions are of the C—H \cdots O type (Table 3). The crystal packing features two dimeric $R_2^2(16)$ and $R_2^2(14)R_2^2(16)$ ring motifs formed through C73—H73 \cdots O1ⁱⁱⁱ and C76—H76 \cdots O2ⁱⁱ/C9—H9B \cdots O2ⁱⁱ hydrogen bonds (Figs. 8 and 9). These C—H \cdots O hydrogen-bonded dimers are arranged in tandem and form a zigzag $C(8)$ chain extending along the *b* axis through a C34—H34 \cdots O1ⁱ interaction (Fig. 10) (see Table 3 for symmetry codes).

Thus, all three title spiroacenaphthylene crystal structures are stabilized through nonclassical C—H \cdots X interactions with primary ring and secondary chain motifs.

**Figure 9**

A partial packing view for (III), showing the $R_2^2(14)R_2^2(16)$ ring. Dashed lines indicate hydrogen bonds. Some H atoms have been omitted for clarity. [Symmetry code: (ii) $-x, -y + 1, -z + 1$.]

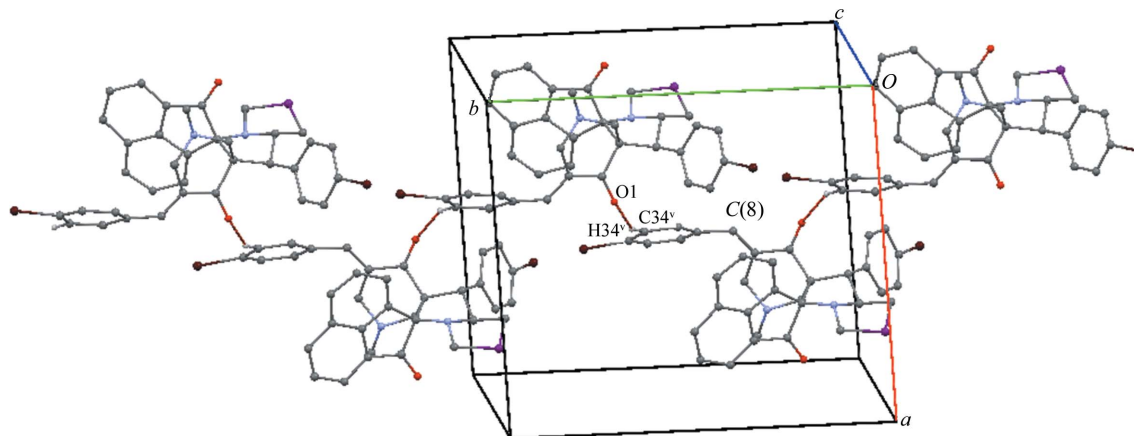


Figure 10
A partial packing view for (III), showing the C(8) chain extending along the *b* axis of the unit cell. Dashed lines indicate hydrogen bonds. Some H atoms have been omitted for clarity. [Symmetry code: (v) $-x + 1, y - \frac{1}{2}, -z + \frac{3}{2}$.]

Experimental

For the preparation of (I), a mixture of 3,5-bis[(*E*)-2,3-dichlorobenzylidene]-1-methyltetrahydropyridin-4(1*H*)-one (1 mmol), acenaphthenequinone (0.182 g, 1 mmol) and 1,3-thiazolane-4-carboxylic acid (0.133 g, 1 mmol) was dissolved in methanol (10 ml) and refluxed for 30 min. After completion of the reaction, as evident from thin-layer chromatography (TLC), the mixture was poured into water (50 ml) and the precipitated solid was filtered off and washed with water (100 ml) to obtain the pure product as a pale-yellow solid. The product was recrystallized from ethyl acetate to obtain suitable crystals of (I) for X-ray analysis (yield 92%, m.p. 477 K).

For the preparation of (II), a mixture of 3,5-bis[(*E*)-4-fluorobenzylidene]-1-methyltetrahydropyridin-4(1*H*)-one (1 mmol), acenaphthenequinone (0.182 g, 1 mmol) and 1,3-thiazolane-4-carboxylic acid (0.133 g, 1 mmol) was dissolved in methanol (10 ml) and refluxed for 30 min. After completion of the reaction, as evident from TLC, the mixture was poured into water (50 ml) and the precipitated solid was filtered off and washed with water (100 ml) to obtain the pure product as a pale-yellow solid. The product was recrystallized from ethyl acetate to obtain suitable crystals of (II) for X-ray analysis (yield 91%, m.p. 483 K).

For the preparation of (III), a mixture of 3,5-bis[(*E*)-4-bromobenzylidene]-1-methyltetrahydropyridin-4(1*H*)-one (1 mmol), acenaphthenequinone (0.182 g, 1 mmol) and 1,3-thiazolane-4-carboxylic acid (0.133 g, 1 mmol) was dissolved in methanol (10 ml) and refluxed for 30 min. After completion of the reaction, as evident from TLC, the mixture was poured into water (50 ml) and the precipitated solid was filtered off and washed with water (100 ml) to obtain the pure product as a pale-yellow solid. The product was recrystallized from ethyl acetate to obtain suitable crystals of (III) for X-ray analysis (yield 91%, m.p. 485 K).

Compound (I)

Crystal data

C₃₅H₂₆Cl₄N₂O₂S
M_r = 680.44
 Monoclinic, *P*2₁/*n*
a = 13.4722 (6) Å
b = 16.4269 (8) Å
c = 14.8673 (7) Å
 β = 108.243 (2)°
V = 3124.9 (3) Å³
Z = 4
 Mo Kα radiation
 μ = 0.48 mm⁻¹
T = 293 K
 0.19 × 0.16 × 0.11 mm

Data collection

Bruker Kappa APEXII area-detector diffractometer
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
T_{min} = 0.912, *T_{max}* = 0.948
 37631 measured reflections
 8258 independent reflections
 5375 reflections with *I* > 2σ(*I*)
R_{int} = 0.034

Refinement

R[*F*² > 2σ(*F*²)] = 0.047
wR(*F*²) = 0.134
S = 1.02
 8258 reflections
 397 parameters
 H-atom parameters constrained
 Δρ_{max} = 0.39 e Å⁻³
 Δρ_{min} = -0.51 e Å⁻³

Compound (II)

Crystal data

C₃₅H₂₈F₂N₂O₂S
M_r = 578.65
 Monoclinic, *P*2₁/*n*
a = 14.8802 (5) Å
b = 9.6789 (3) Å
c = 20.2209 (6) Å
 β = 104.155 (2)°
V = 2823.87 (15) Å³
Z = 4
 Mo Kα radiation
 μ = 0.16 mm⁻¹
T = 293 K
 0.21 × 0.14 × 0.11 mm

Data collection

Bruker Kappa APEXII area-detector diffractometer
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
T_{min} = 0.973, *T_{max}* = 0.982
 26085 measured reflections
 5494 independent reflections
 3862 reflections with *I* > 2σ(*I*)
R_{int} = 0.031

Refinement

R[*F*² > 2σ(*F*²)] = 0.043
wR(*F*²) = 0.140
S = 1.08
 5494 reflections
 381 parameters
 H-atom parameters constrained
 Δρ_{max} = 0.48 e Å⁻³
 Δρ_{min} = -0.22 e Å⁻³

Compound (III)

Crystal data

C₃₅H₂₈Br₂N₂O₂S
M_r = 700.47
 Monoclinic, *P*2₁/*c*
a = 14.2275 (7) Å
b = 15.6753 (8) Å
c = 15.3780 (7) Å
 β = 116.761 (2)°
V = 3062.3 (3) Å³
Z = 4
 Mo Kα radiation
 μ = 2.75 mm⁻¹
T = 293 K
 0.19 × 0.16 × 0.11 mm

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>
C74—H74...Cl1 ⁱ	0.93	2.82	3.497 (3)	130
C2—H2B...Cl1 ⁱⁱ	0.97	2.86	3.784 (2)	159
C9—H9B...O1	0.97	2.44	3.251 (3)	141

Symmetry codes: (i) $x, y - 1, z$; (ii) $-x + 1, -y + 1, -z$.**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>
C33—H33...S1 ⁱ	0.93	2.87	3.716 (3)	152
C37—H37...F2 ⁱⁱ	0.93	2.48	3.302 (3)	147
C73—H73...O2 ⁱⁱⁱ	0.93	2.68	3.454 (4)	141
C10—H10B...O2	0.97	2.59	3.202 (3)	122

Symmetry codes: (i) $-x + \frac{1}{2}, y - \frac{1}{2}, -z + \frac{1}{2}$; (ii) $-x - \frac{1}{2}, y + \frac{1}{2}, -z + \frac{1}{2}$; (iii) $x, y - 1, z$.**Table 3**
Hydrogen-bond geometry (Å, °) for (III).

<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>
C34—H34...O1 ⁱ	0.93	2.57	3.440 (5)	157
C76—H76...O2 ⁱⁱ	0.93	2.46	3.250 (5)	143
C73—H73...O1 ⁱⁱⁱ	0.93	2.45	3.361 (5)	166
C9—H9B...O2 ⁱⁱ	0.97	2.62	3.298 (3)	128
C10—H10A...O2	0.97	2.54	3.176 (4)	123

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

Data collection

Bruker Kappa APEXII area-detector diffractometer	30894 measured reflections
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	6169 independent reflections
$T_{\min} = 0.599, T_{\max} = 0.739$	3897 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.039$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.044$	H-atom parameters constrained
$wR(F^2) = 0.141$	$\Delta\rho_{\text{max}} = 0.78 \text{ e } \text{Å}^{-3}$
$S = 1.00$	$\Delta\rho_{\text{min}} = -0.98 \text{ e } \text{Å}^{-3}$
6169 reflections	
382 parameters	

H atoms were placed at calculated positions and allowed to ride on their carrier atoms, with C—H = 0.93 (aromatic), 0.96 (methyl), 0.97 (methylene) or 0.98 Å (methine), and with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$ for methyl H atoms or $1.2U_{\text{eq}}(\text{C})$ otherwise.

For all three compounds, 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: *PLATON* (Spek, 2009); software used to prepare material for publication: *SHELXL97*.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: GG3273). Services for accessing these data are described at the back of the journal.

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