

The proton-transfer compounds of strychnine with achiral salicylic acids: strychninium 3,5-dinitrosalicylate and the strychninium 5-nitrosalicylate bis(5-nitrosalicylic acid) adduct

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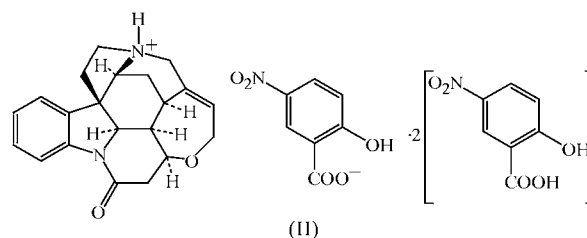
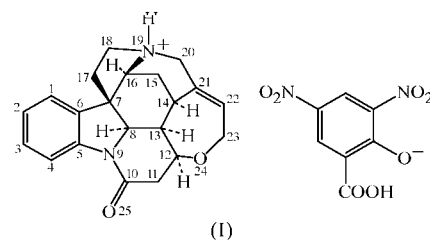
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In the crystal structures of the proton-transfer compounds of strychnine with 3,5-dinitrosalicylic acid, namely strychninium 3,5-dinitrosalicylate, $C_{21}H_{23}N_2O_2^+ \cdot C_7H_3N_2O_7^-$, (I), and 5-nitrosalicylic acid, namely strychninium–5-nitrosalicylate–5-nitrosalicylic acid (1/1/2), $C_{21}H_{23}N_2O_2^+ \cdot C_7H_4NO_5^- \cdot 2C_7H_5NO_5$, (II), protonation of one of the N atoms of the strychnine molecule occurs and this group is subsequently involved in intermolecular hydrogen-bonding interactions. In (I), this is four-centred, the primary being with an adjacent strychninium carbonyl O-atom acceptor in a side-to-side interaction giving linear chains. Other interactions are with the phenolate and nitro O-atom acceptors of the anionic species, resulting in a one-dimensional polymer structure. In (II), the $N^+ - H$ interaction is three-centred, the hydrogen bonding involving carboxyl O-atom acceptors of the anion and both acid adduct species, giving unique discrete hetero-tetramer units. The structure of (II) also features π -bonding interactions between the two acid adduct molecules.

Comment

Strychnine and brucine have been variously employed on a hit-or-miss basis as resolving agents for a range of chiral organic compounds, and the crystal structures of a large number of complexes with strychnine and brucine, together with their absolute configurations, have been determined. The complexes include those with acidic species, in which atom N19 of the strychnine or brucine molecule ($pK_{a2} = 11.7$) is protonated, e.g. the *N*-benzoyl-, *N*-phthaloyl- and *N*-acetyl-protected amino acids (Gould & Walkinshaw, 1984; Gould, Taylor & Walkinshaw, 1984; Gould *et al.*, 1985; Białońska & Ciunik, 2004a; Quinkert *et al.*, 1986; Kuwata *et al.*, 1993), and other chiral acid types (Gould *et al.*, 1987, 2002; Boiadjev *et al.*, 1992; Wright *et al.*, 1994; Bao *et al.*, 1996; Costente *et al.*, 1996; Dijkstra, Gould, Parsons, Taylor & Walkinshaw, 1998; Andersson *et al.*, 1999; Allenmark & Skogsberg, 2000; Białońska *et al.*, 2005). Other structures with neutral chiral guest species are known, e.g. with alcohols, lactones, cyanohydrins and ketones (Toda *et al.*, 1981, 1985; Tanaka *et al.*, 2001; Chandramohan & Ravikumar, 1999; Pinkerton *et al.*, 1993; Yamagishi *et al.*, 1992).



Although strychnine and brucine are both physicochemically and structurally similar and configurationally identical, brucine has proved to be the better of the two for optical resolution. This appears to be because of the methoxy groups in the 2- and 3-positions of the aromatic ring influencing the solid-state packing of the brucine molecules, which commonly form undulating parallel chain structures (Gould & Walkinshaw, 1984; Dijkstra *et al.*, 1998; Białońska *et al.*, 2005). These recognize compatible molecular guest species which occupy the interstitial cavities between the chains and associate with the host through hydrogen bonding. Water or other molecules of solvation may also act, if needed, in a space-filling and/or in a proton-donor or -acceptor capacity. This is apparent in the isomorphous crystals of brucine–ethanol–water (1/1/2) (Glover *et al.*, 1985) and brucine–propan-2-ol–water (1/1/2) (Białońska & Ciunik, 2004b), and in brucine–acetone (1/1) structures (Białońska & Ciunik, 2004b). Strychnine is less regular as a host structure for organic molecule recognition, often giving isolated molecular complexes or forming double-layer polymeric structures (Gould & Walkinshaw, 1984; Dijkstra, Gould, Parsons, Taylor & Walkinshaw, 1998).

More recently, the structures of a number of neutral and proton-transfer compounds of strychnine and brucine with achiral organic molecules have been determined, e.g. with 4-nitrophenol (Guo *et al.* 2001), fumaric and maleic acids (Dijkstra, Gould, Parsons & Walkinshaw, 1998), 4-hydroxybenzoic acid (Sada *et al.*, 1998), 3-nitrobenzoic acid (Oshikawa *et al.*, 2002) and 8-aminonaphthalene-2-sulfonic acid (Smith, Wermuth, Healy & Young, 2005). Because it was observed by Oshikawa *et al.* (2002) that brucine has a recognitive affinity for *meta*-substituted benzoic acids, we therefore considered that the analogous acids 3,5-dinitrosalicylic acid (DNSA),

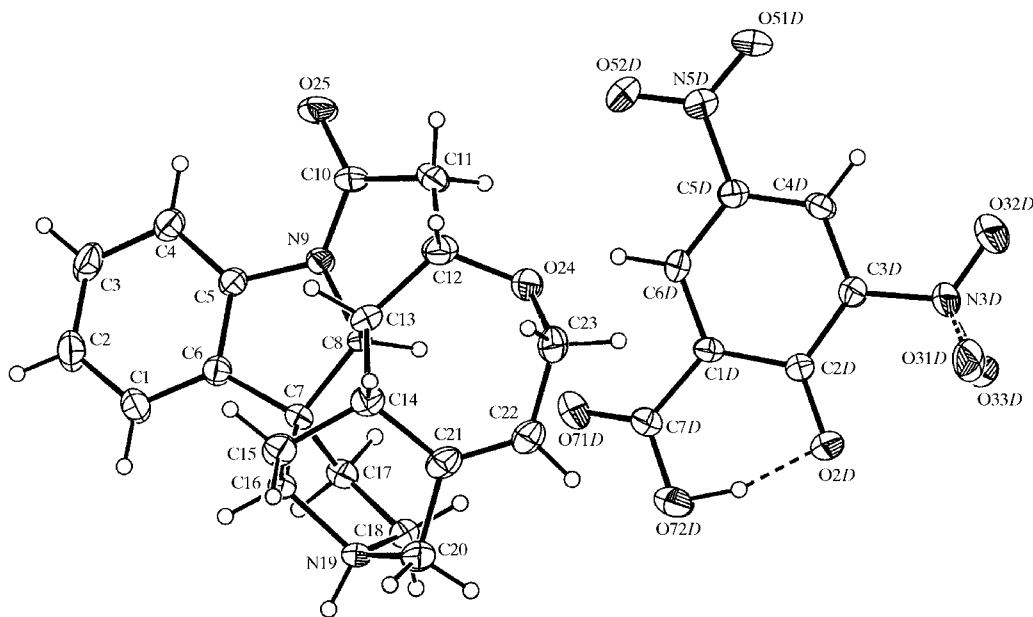


Figure 1

The molecular configuration and atom-numbering scheme for the strychninium cation and the DNSA anion species in compound (I). Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radii. Atoms O31D and O33D are disordered sites (see *Comment*).

5-nitrosalicylic acid (5-NSA), 5-sulfosalicylic acid (5-SSA) and 3-nitrophthalic acid (NPA) were likely candidates for similar recognition by brucine and possibly strychnine. This has proved to be the case with brucine, where good crystalline products were obtained within one week for DNSA, two weeks for 5-NSA and 5-SSA, and several weeks with NPA. The crystal structures of all four compounds have now been determined (Smith, Wermuth & Healy, 2005; Smith, Wermuth, Young & Healy, 2005). However, with strychnine, no complex was obtained with 5-SSA or NPA, although good crystals of the compounds with DNSA and 5-NSA were formed, albeit more slowly than with brucine. The crystal structures of these two compounds, strychninium 3,5-dinitrosalicylate, (I), and the adduct strychninium–5-nitrosalicylate–5-nitrosalicylic acid (1/1/2), (II), respectively, are reported here.

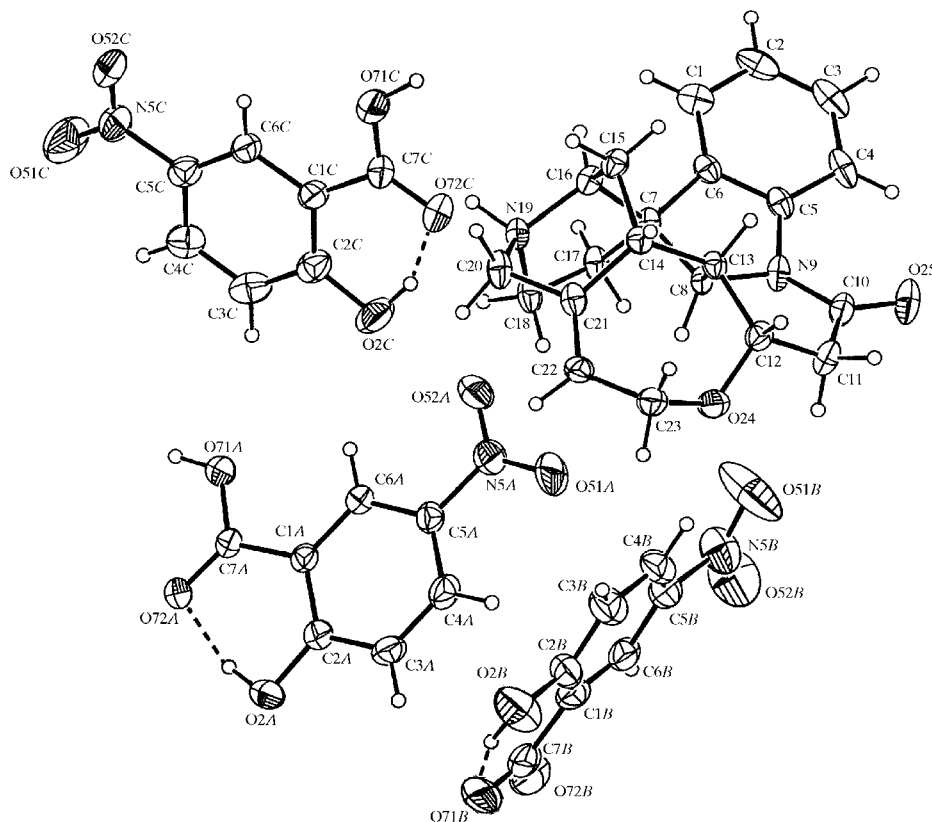
Compounds (I) and (II) are both anhydrous, which is consistent with the structures of the proton-transfer compounds of both DNSA and 5-NSA with Lewis bases, where water or other molecules of solvation are seldom incorporated (Smith *et al.*, 2002, 2003; Smith, Hartono *et al.*, 2005), although the brucine complex with DNSA is a monohydrate (Smith, Wermuth & Healy, 2005). The major difference between (I) and (II) is the presence in (II) of two additional adduct molecules of 5-NSA acid, adduct formation being unusual among 5-NSA compounds, as well as among examples of brucine or strychnine complexes.

In the structures of both (I) and (II) (Figs. 1 and 2), the expected proton transfer to N19 of the strychnine molecule occurs and this group is subsequently involved in intermolecular $N^+ - H \cdots O$ hydrogen-bonding interactions with two O-atom acceptors of the anion species, and, in the case of (I), a strychnine carbonyl O atom (Tables 1 and 2).

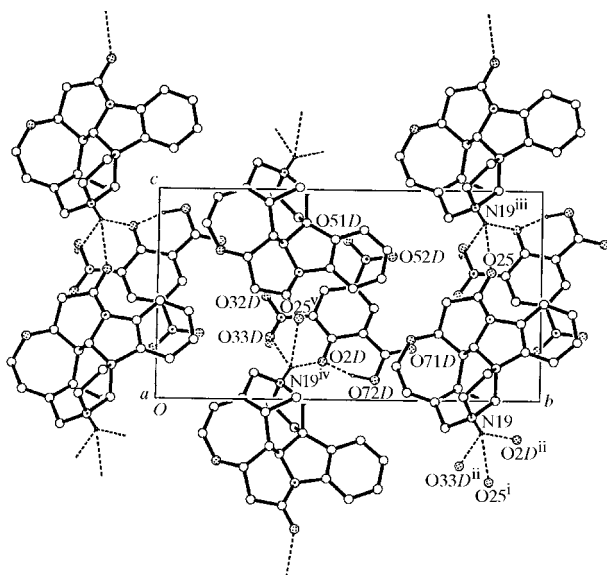
In each structure, the absolute configuration of the parent strychnine molecule (Peerdeman, 1956) is invoked. This includes the ‘apparent’ change in configuration at C7 (*R* to *S*), which is a consequence of the change in the hierarchy of the protonated N19 group in the Cahn–Ingold–Prelog system and the introduction of a new chiral centre at N19 (*S*) (Smith, Wermuth, Healy & Young, 2005). This gives the overall absolute stereochemistry for the strychninium cations in (I) and (II) (as with all proton-transfer compounds of both brucine and strychnine) as C7(*S*), C8(*S*), C12(*S*), C13(*R*), C14(*R*), C16(*S*) and N19(*S*).

In (I), the linear strychninium framework is formed through side-to-side hydrogen-bonding interactions involving the protonated N19 group and an adjacent strychnine carbonyl O atom [$N19-H19 \cdots O25^i = 3.148(4) \text{ \AA}$; see Table 1 for symmetry code], extending along the *c*-axis direction (Fig. 3). Atom N19 is then involved in a proximal association with both a phenolic-O and a disordered nitro-O acceptor of a glide-related DNSA anion [$N19 \cdots O2D^{ii} = 2.857(5)$ and $N19 \cdots O33D^{ii} = 3.00(3) \text{ \AA}$; see Table 1 for symmetry code]. This generates a linear polymer structure in which, surprisingly, there are no intermolecular associations involving the O-atom acceptors of the DNSA carboxyl group.

In (II), an unusual discrete hetero-tetramer is formed, comprising the strychninium cation, the 5-NSA anion (molecule *B*) and the two 5-NSA acid adduct molecules (*A* and *C*) (Fig. 4). The three-centre association with $N19^+ - H$ involves carboxyl-O acceptors of the anion [$N19 \cdots O72B^i = 3.223(8) \text{ \AA}$; see Table 2 for symmetry code] and an adduct molecule *A* [$N19 \cdots O72A^{ii} = 2.958(5) \text{ \AA}$. Completing a cyclic $R_2^2(6)$ association is the carboxyl H atom of the adduct molecule *A* [$O71A \cdots O72B^{ii} = 2.553(6) \text{ \AA}$]. The second adduct

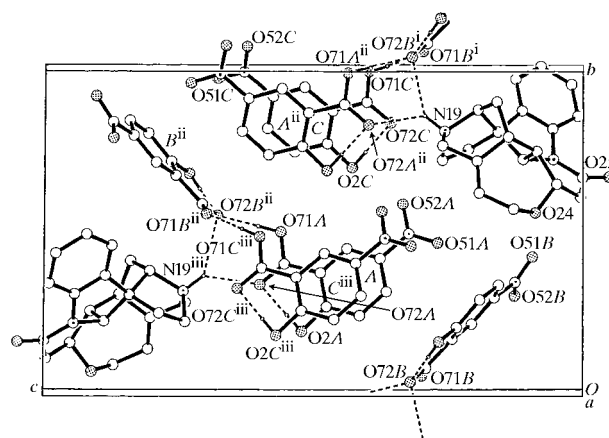

Figure 2

The molecular configuration and atom-numbering scheme for the strychninium cation, the 5-NSA anion and the two 5-NSA acid adduct species in compound (II). Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radii.


Figure 3

The packing of (I) in the unit cell, viewed down the *a* axis. Non-interacting H atoms have been omitted. Hydrogen-bonding associations are shown as broken lines. [Symmetry codes: (iii) $x, y, 1 + z$; (iv) $-x, y - \frac{1}{2}, -z$; (v) $-x, y - \frac{1}{2}, 1 - z$; for others, see Table 1.]

molecule is linearly hydrogen bonded to the anion molecule through the carboxyl groups [2.709 (8) Å], such that both adduct molecules form π -associated stacks [ring centroid


Figure 4

The packing of (II) in the unit cell, viewed down the *c* axis. The discrete hydrogen-bonded tetrameric units are shown (dashed lines). Also illustrated are the π -bonding associations between the two 5-NSA adduct molecules (*A* and *C*). [Symmetry codes: (iii) $1 - x, y - \frac{1}{2}, 1 - z$; for others, see Table 2.]

separations: $CgA \cdots CgC = 3.72 (1)$ (intra) and $4.09 (1)$ Å (inter); CgA and CgC denote the centroids of the rings in molecules *A* and *C*, respectively]. Both adduct formation and π -stacking effects are previously unknown among structures of 5-NSA compounds (Smith, Hartono *et al.*, 2005). The tetramer units are unassociated, except for an unusual side-on

interaction between the intramolecularly hydrogen-bonded phenol group of an adduct molecule *C* and a nitro O atom of an adjacent adduct molecule *A* [$O2C \cdots O52A = 2.854(6) \text{ \AA}$] (Figs. 2 and 4).

Within the DNSA anion in (I), the structural features vary slightly from those of the majority of proton-transfer compounds (Smith *et al.*, 2003), particularly with regard to the conformation of the nitro-substituent groups. The proximal nitro group at C3 is more commonly involved in hydrogen bonding than the C5 nitro group and therefore usually shows a greater rotation out of the plane of the ring than the C5 group. In (I), this is also the case [$C2D-C3D-N3D-O33D = 167.4(5)^\circ$ and $C4D-C5D-N5D-O52D = 179.5(4)^\circ$]. However, the previously mentioned intermolecularly unassociated carboxyl group is non-coplanar [$C2D-C1D-C7D-O71D = -166.4(4)^\circ$], although it is involved in the intramolecular hydrogen bond. This hydrogen bond [$O2D \cdots O72D = 2.492(5) \text{ \AA}$] has the H atom located on the carboxyl O atom, rather than on the phenol group as is found in *ca* 70% of the proton-transfer compounds of DNSA (Smith *et al.*, 2002, 2003).

With the 5-NSA species in (II), despite the presence of adduct acid molecules and the associated π -stacking effects, structural features vary little from those previously reported (Smith, Hartono *et al.*, 2005), which includes near-coplanarity between the parent ring and both the carboxyl- and nitro-substituent groups, the invariable location of the intramolecular H atom on the phenol O atom, and a contraction of this intramolecular O \cdots O distance with deprotonation [$2.509(10) \text{ \AA}$ in molecule *B*, compared with $2.595(5) \text{ \AA}$ in molecule *A* and $2.605(6) \text{ \AA}$ in molecule *C*].

Experimental

The title compounds were synthesized by heating 1 mmol quantities of strychnine (strychnidin-10-one) and either 3,5-dinitrosalicylic acid (DNSA) or 5-nitrosalicylic acid (5-NSA) in 50% ethanol-water (50 ml) for 10 min under reflux. After concentration to *ca* 30 ml, partial room-temperature evaporation of the hot-filtered solutions gave yellow prisms of (I) (m.p. 468–470 K) and minor colourless prisms of (II) (m.p. 487–489 K).

Compound (I)

Crystal data

$C_{21}H_{23}N_2O_2^+ \cdot C_7H_3N_2O_7^-$
 $M_r = 562.53$
 Monoclinic, $P2_1$
 $a = 7.5036(15) \text{ \AA}$
 $b = 17.219(3) \text{ \AA}$
 $c = 9.4799(19) \text{ \AA}$
 $\beta = 96.905(3)^\circ$
 $V = 1216.0(4) \text{ \AA}^3$
 $Z = 2$
 $D_x = 1.536 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation
 Cell parameters from 2715 reflections
 $\theta = 2.4\text{--}27.3^\circ$
 $\mu = 0.12 \text{ mm}^{-1}$
 $T = 295(2) \text{ K}$
 Block, yellow
 $0.40 \times 0.30 \times 0.20 \text{ mm}$

Data collection

Bruker SMART CCD area-detector diffractometer
 φ and ω scans
 5865 measured reflections
 2222 independent reflections
 1909 reflections with $F^2 > 2\sigma(F^2)$
 $R_{\text{int}} = 0.071$
 $\theta_{\text{max}} = 25.0^\circ$
 $h = -8 \rightarrow 8$
 $k = -20 \rightarrow 18$
 $l = -10 \rightarrow 11$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.058$
 $wR(F^2) = 0.142$
 $S = 1.17$
 2222 reflections
 384 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0734P)^2 + 0.1261P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.021$
 $\Delta\rho_{\text{max}} = 0.22 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.25 \text{ e \AA}^{-3}$

Table 1

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

<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>
O72D—H72D \cdots O2D	1.02	1.50	2.492 (5)	165
N19—H19 \cdots O25 ⁱ	0.91 (5)	2.59 (5)	3.148 (4)	120 (4)
N19—H19 \cdots O2D ⁱⁱ	0.91 (5)	2.11 (5)	2.857 (5)	139 (4)
N19—H19 \cdots O33D ⁱⁱ	0.91 (5)	2.41 (6)	3.00 (3)	123 (4)

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

Compound (II)

Crystal data

$C_{21}H_{23}N_2O_2^+ \cdot C_7H_4NO_5^-$
 $2C_7H_5NO_5$
 $M_r = 883.77$
 Monoclinic, $P2_1$
 $a = 7.5762(9) \text{ \AA}$
 $b = 12.3729(9) \text{ \AA}$
 $c = 20.891(4) \text{ \AA}$
 $\beta = 97.96(1)^\circ$
 $V = 1939.5(5) \text{ \AA}^3$
 $Z = 2$
 $D_x = 1.513 \text{ Mg m}^{-3}$

Cu $K\alpha$ radiation
 Cell parameters from 25 reflections
 $\theta = 20\text{--}30^\circ$
 $\mu = 1.01 \text{ mm}^{-1}$
 $T = 295(2) \text{ K}$
 Prismatic, colourless
 $0.30 \times 0.20 \times 0.20 \text{ mm}$

Data collection

Enraf-Nonius CAD-4F diffractometer
 $\omega/2\theta$ scans
 3960 measured reflections
 3856 independent reflections
 3427 reflections with $F^2 > 2\sigma(F^2)$
 $R_{\text{int}} = 0.045$

$\theta_{\text{max}} = 69.9^\circ$
 $h = -9 \rightarrow 9$
 $k = 0 \rightarrow 15$
 $l = 0 \rightarrow 25$
 3 standard reflections
 frequency: 160 min
 intensity decay: none

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.056$
 $wR(F^2) = 0.168$
 $S = 1.02$
 3856 reflections
 585 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.1204P)^2 + 0.567P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.002$
 $\Delta\rho_{\text{max}} = 0.57 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.35 \text{ e \AA}^{-3}$

Table 2

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

<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>
N19—H19 \cdots O72B ⁱ	0.99 (13)	2.52 (7)	3.233 (8)	128 (2)
N19—H19 \cdots O72A ⁱⁱ	0.99 (13)	2.19 (13)	2.958 (5)	133 (2)
O71A—H71A \cdots O72B ⁱⁱ	0.91	1.65	2.553 (6)	179
O71C—H71C \cdots O71B ⁱ	0.90	1.81	2.709 (8)	179
O2A—H2A \cdots O72A	0.91	1.69	2.595 (5)	180
O2B—H2B \cdots O72B	0.82	1.70	2.509 (10)	168
O2C—H2C \cdots O72C	0.90	1.71	2.605 (6)	179

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

H atoms potentially involved in hydrogen-bonding interactions were located by difference methods but, with the exception of the strychninium atom H19 (on N19), their positional and isotropic displacement parameters were not refined. Other H atoms were included in the refinement in calculated positions (aromatic C—H = 0.93 Å and aliphatic C—H = 0.97 Å) and treated using a riding model, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. The atom-numbering scheme for the strychninium species in (I) and (II) (Figs. 1 and 2, respectively) follows the original Robinson convention (Holmes, 1952). Friedel pairs were averaged and the absolute configuration determined for the parent strychnine (Peerdeman, 1956) was invoked, giving the overall Cahn–Ingold–Prelog absolute stereochemistry (Eliel, 1962) as C7(*S*), C8(*S*), C12(*S*), C13(*R*), C14(*R*), C16(*S*) and N19(*S*). One of the nitro O atoms in (I) was found to be disordered and was subsequently refined over two sites [O31*D*, with a site-occupancy factor of 0.44 (5), and O33*D*, with a site-occupancy factor of 0.56 (5)].

Data collection: *SMART* (Bruker, 2000) for (I); *CAD-4 Software* (Enraf–Nonius, 1989) for (II). Cell refinement: *SMART* for (I); *CAD-4 Software* for (II). Data reduction: *SAINT* (Bruker, 1999) for (I); *XCAD4* (Harms & Wocadlo, 1995) for (II). Structure solution: *SHELXTL* (Bruker, 1997) for (I); *SHELXS97* (Sheldrick, 1997) in *WinGX* (Farrugia, 1999) for (II). Structure refinement: *SHELXTL* for (I); *SHELXL97* (Sheldrick, 1997) in *WinGX* for (II). For both compounds, molecular graphics: *PLATON* (Spek, 2003); publication software: *PLATON*.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: TA1499). Services for accessing these data are described at the back of the journal.

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