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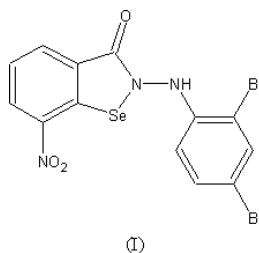
Key indicators

Single-crystal X-ray study
 $T = 293\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.005\text{ \AA}$
Disorder in main residue
 R factor = 0.033
 wR factor = 0.085
Data-to-parameter ratio = 14.4For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.2-(2,4-Dibromophenylamino)-7-nitro-
benzisoselenazol-3-one

The crystal structure determination of the title compound, $\text{C}_{13}\text{H}_7\text{Br}_2\text{N}_3\text{O}_3\text{Se}$, was undertaken in the course of a study on 1,2-benzoselenazin-4-one heterocycle homologues of ebselen, a well known anti-oxidizing agent. It allowed unambiguous identification of the molecular formula of the compound. The Br atom in the 4 position is split into two sites with occupation factors of 55 and 45%. The dihedral angle between the phenyl and benzisoselenazole mean planes is $85.0(5)^\circ$. Analysis of the crystal packing shows infinite chains of molecules along the b axis, linked together by an $\text{N}-\text{H}\cdots\text{O}$ intermolecular hydrogen bond.

Comment

Although it has attractive anti-inflammatory activity and low toxicity, ebselen (Natterman/RP, 1981; Dupont *et al.*, 1990) exhibits some physicochemical failings. Its low solubility in organic solvents and its high melting point (453–455 K) induce a feeble metabolism. Moreover, a photochemical reaction is observed with the time and enjoins particular form of conditioning; the drug capsules must have an opaque envelope to protect the drug from light. In the course of synthesizing new molecules of benzo[*e*]-1,2-selenazin-4-one family which could be potential anti-oxidizing agents, the title compound, (I), was obtained and the unexpected structure determined by X-ray diffraction.



In the final cycles of refinement, the position of atom Br2 was split into two sites, Br2A and Br2B, with occupation factors of 0.45 and 0.55, respectively. The introduction of this disordered distribution greatly improved the residual densities in the final difference map and also the conventional R and $wR2$ values. The distances and angles are similar to those found in 7-nitro-2-phenyl-1,2-benzisoselenazol-3(2H)-one, (II) (Dupont *et al.*, 1988). In (II) the dihedral angle between the phenyl and heterocycle mean planes is $13.2(1)^\circ$, whereas in (I) the corresponding planes are nearly orthogonal [$85.0(5)^\circ$]. The cohesion of the crystal is the result of an $\text{N}-\text{H}\cdots\text{O}$ hydrogen bond, which links molecules indefinitely along the b axis (Fig. 2), and of van der Waals interactions. There is also a short intramolecular $\text{N}3\cdots\text{Br}1$ contact

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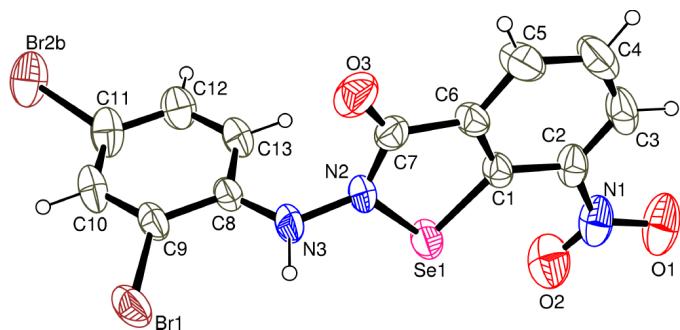


Figure 1
The molecular structure of (I), with the atom-labelling scheme. Displacement ellipsoids are shown at 50% probability levels. Atom Br2A (occupation factor = 45%) has been omitted.

[3.074 (3) Å], but, as the N—H···Br angle is rather narrow [103 (3)°], it could hardly be considered as a hydrogen bond. The packing shows five other short intermolecular contacts: Br1···Br2Aⁱ 3.629 (6), Br1···Br2Bⁱ 3.524 (5), Br1···O2ⁱⁱ 3.376 (4), Br2A···O1ⁱⁱⁱ 3.261 (7) and Br2B···O1ⁱⁱⁱ 3.368 (6) Å [symmetry codes: (i) 1−*x*, −1/2+*y*, 1/2−*z*; (ii) 1/2−*x*, 1/2+*y*, *z*; (iii) −*x*, 1/2+*y*, 3/2−*z*].

Experimental

The title compound was prepared by treatment of 2-methylseleno-3-nitro-3'-*N*-phenylbenzoylhydrazide with 3 equivalents of bromine in a solution of dichloroethane (Messali, 2001). An orange single crystal was obtained by slow evaporation of a toluene solution.

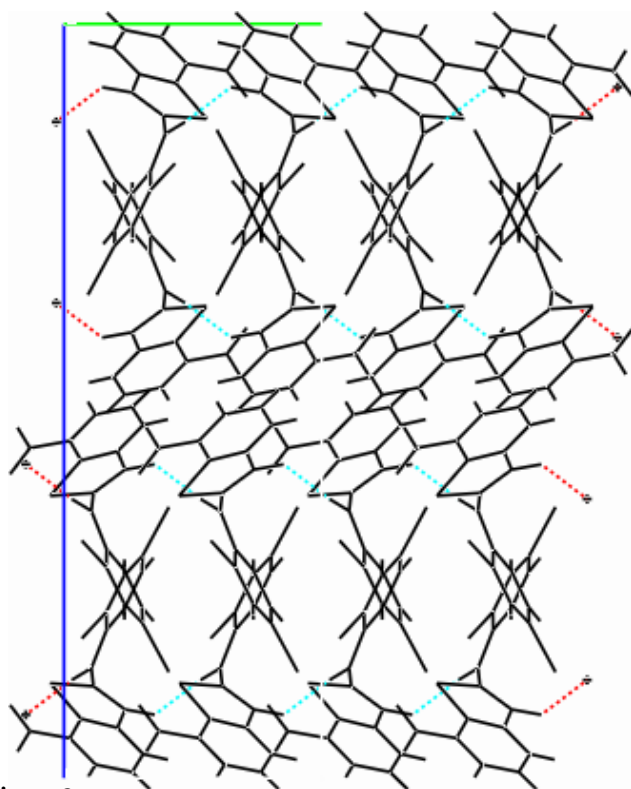


Figure 2
Projection of the structure along *a*, with the scheme of hydrogen bonds. The packing range is expanded horizontally along *b*, to show two unit cells.

Crystal data

C₁₃H₇Br₂N₃O₃Se
M_r = 492.00
 Orthorhombic, *Pbca*
a = 13.8685 (10) Å
b = 8.6075 (10) Å
c = 25.233 (2) Å
V = 3012.1 (5) Å³
Z = 8
D_x = 2.170 Mg m^{−3}

Cu *K*α radiation
 Cell parameters from 25 reflections
 θ = 30–42°
 μ = 9.75 mm^{−1}
T = 293 (2) K
 Prism, orange
 0.30 × 0.25 × 0.15 mm

Data collection

Enraf–Nonius CAD-4 diffractometer
 $\theta/2\theta$ scans
 Absorption correction: ψ scan (North *et al.*, 1968)
 T_{\min} = 0.12, T_{\max} = 0.39
 3587 measured reflections
 3075 independent reflections
 2634 reflections with $I > 2\sigma(I)$

R_{int} = 0.022
 θ_{max} = 74.2°
 h = −17 → 10
 k = 0 → 10
 l = 0 → 31
 3 standard reflections every 200 reflections
 intensity decay: 4%

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)]$ = 0.033
 $wR(F^2)$ = 0.085
 S = 1.10
 3075 reflections
 213 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0307P)^2 + 4.0625P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}}$ = 0.001
 $\Delta\rho_{\text{max}}$ = 0.44 e Å^{−3}
 $\Delta\rho_{\text{min}}$ = −0.48 e Å^{−3}

Table 1

Selected geometric parameters (Å, °).

Br1—C9	1.890 (4)	N1—C2	1.446 (5)
Se1—C1	1.855 (3)	N2—C7	1.355 (5)
Se1—N2	1.899 (3)	N2—N3	1.386 (4)
O1—N1	1.228 (4)	N3—C8	1.385 (5)
O2—N1	1.230 (5)	C11—Br2A	1.847 (6)
O3—C7	1.220 (4)	C11—Br2B	1.939 (6)
C1—Se1—N2	83.82 (14)	C7—N2—Se1	117.5 (2)
O1—N1—O2	123.9 (4)	N3—N2—Se1	121.6 (3)
C7—N2—N3	120.6 (3)	C8—N3—N2	119.3 (3)
C1—Se1—N2—C7	1.2 (3)	Se1—N2—N3—C8	−93.8 (4)
C1—Se1—N2—N3	−173.1 (3)		

Table 2

Hydrogen-bonding 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>
N3—H30···O3 ⁱ	0.93 (4)	1.98 (4)	2.826 (4)	151 (4)

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

Least-squares restraints were applied to refine the disordered positions of Br2A and Br2B. All H atoms, with the exception of the nitrogen-bound atom (H30), were included in the refinement in the riding-model approximation, with isotropic displacement parameters fixed at 1.2*U*_{eq} of the parent atom. H30 was refined isotropically, its displacement parameters being fixed at 1.2*U*_{eq} of the attached N atom.

Data collection: *CAD-4 EXPRESS* (Enraf–Nonius, 1992); cell refinement: *CAD-4 EXPRESS*; data reduction: *HELENA* (Spek, 1997); program(s) used to solve structure: *SIR92* (Altomare *et al.*,

1993); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPIII* (Burnett & Johnson, 1996) and *MERCURY* 1.1 (Bruno *et al.*, 2002); software used to prepare material for publication: *SHELXL97*.

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