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[3aR-(3a α ,4 α ,7 α ,7a α)]-2-Acetyl-4,8,8-trimethylperhydro-4,7-methano-1,2-benzisoselenazol-3a-ol

THOMAS G. BACK,* BRIAN P. DYCK, SIQIAO NAN AND MASOOD PARVEZ*

Department of Chemistry, The University of Calgary, 2500 University Drive NW, Calgary, Alberta, Canada T2N 1N4. E-mail: parvez@acs.ucalgary.ca

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Abstract

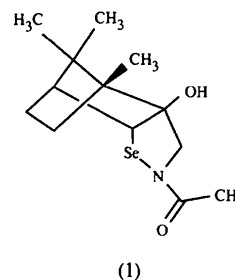
Crystals of the title compound, $C_{13}H_{21}NO_2Se$, are composed of independent molecules wherein the six-membered ring of the camphor-derived moiety is fused with a novel five-membered heterocyclic ring incorporating Se and N atoms. The bond lengths and angles are normal, with Se—N and Se—C distances of 1.865 (3) and 1.943 (3) Å, respectively, and an N—Se—C angle of 87.1 (2)°. The mean value of the C_{sp^3} — C_{sp^3} bond length in the camphor moiety is 1.542 (15) Å. The five-membered ring has an N-envelope conformation and the molecules are hydrogen bonded forming chains extended along the *b*-axis direction, with O...O and O—H...O separations of 2.769 (4) and 1.84 Å, respectively.

Comment

Glutathione peroxidase (GSH-Px) is a selenium-containing enzyme that catalytically destroys hydroperoxides formed as normal by-products of oxidative metabolism (Hoekstra, 1974; Shamberger, 1983; Burk, 1994; Tappel, 1984; Flohé, 1985). This protects living organisms from damage that ensues from free radicals that would otherwise be generated from such hydroperoxides. It has been established that the redox properties of selenocysteine residues in GSH-Px are responsible for its catalytic activity (Ganther, 1975; Ganther & Kraus, 1984). A variety of simpler synthetic selenium and tellurium compounds have been investigated for similar abilities to destroy peroxide species (Reich & Jasperse, 1987; Wilson *et al.*, 1989; Iwaoka & Tomoda, 1994; Engman *et al.*, 1992; Engman, Stern *et al.*, 1994; Vessman *et al.*, 1995; Engman, Andersson *et al.*, 1994; Detty *et al.*, 1994; Fong & Schiesser, 1995; Jacquemin *et al.*, 1992; Galet *et al.*, 1994). One such compound, Ebselen, has been studied for its therapeutic potential as an anti-inflammatory agent, and in connection with a variety of other disease states related to oxidative stress and ensuing free-radical formation (Fong & Schiesser, 1995; Jacquemin *et al.*, 1992; Galet *et al.*, 1994). Ebselen and related compounds, however, func-

tion by mechanisms that are significantly different from that of GSH-Px itself (Engman *et al.*, 1992; Fischer & Dereu, 1987; Haenen *et al.*, 1990).

We recently reported that the novel (1*R*)-(+)-camphor-derived cyclic selenenamide (1) catalytically destroys hydroperoxides *via* an identical mechanism to that used by GSH-Px (Back & Dyck, 1997). Since it is of interest to design and test structural variations of GSH-Px mimetic (1) in attempts to improve its catalytic properties, it is important to know the precise structure of this lead compound. In connection with our efforts in this area, we now report the structure and absolute configuration of (1) determined by X-ray crystallography.



The title compound (Fig. 1) contains a novel five-membered heterocyclic ring containing Se and N atoms, which is being reported for the first time in a crystal structure, fused with the six-membered ring of a camphor-derived moiety. A search of the Cambridge Structural Database (Allen & Kennard, 1993) did not reveal any structure containing the heterocyclic ring present in (1). The bond lengths and angles are normal, with Se—N and Se—C distances of 1.865 (3)

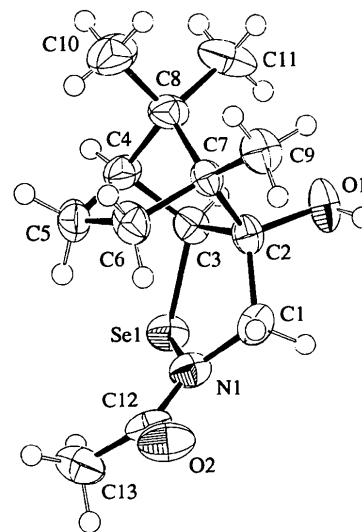


Fig. 1. ORTEP (Johnson, 1976) drawing of the title compound. Displacement ellipsoids are plotted at the 50% probability level.

and 1.943 (3) Å, respectively, and an N—Se—C angle of 87.1 (2)° (Allen *et al.*, 1992). The mean value of the C_{sp³}—C_{sp³} bond length in the camphor moiety is 1.542 (15) Å, while the C_{sp³}—C_{sp²}, N—C_{sp³}, N—C_{sp²}, O—C_{sp³} and O=C bond lengths are 1.496 (7), 1.463 (5), 1.348 (5), 1.425 (4) and 1.243 (4) Å, respectively. The five-membered ring has an N1-envelope conformation, with N1 0.619(6) Å out of the plane of the remaining atoms of the ring. The selenenamido moiety comprised of the Se1, N1, O2, C1, C12, C13 atoms is almost planar, with a maximum deviation of 0.050 (5) Å. The molecules are linked in the solid by hydrogen bonds extending along the *b*-axis direction (details are in Table 2).

Experimental

The title compound was prepared according to a procedure described previously (Back & Dyck, 1997). Suitable crystals were grown by slow evaporation from an ether–hexane solution at room temperature.

Crystal data

C ₁₃ H ₂₁ NO ₂ Se	Cu Kα radiation
<i>M_r</i> = 302.27	λ = 1.54178 Å
Orthorhombic	Cell parameters from 25 reflections
<i>P</i> 2 ₁ 2 ₁ 2 ₁	θ = 20–30°
<i>a</i> = 6.8684 (10) Å	μ = 3.684 mm ⁻¹
<i>b</i> = 11.537 (2) Å	<i>T</i> = 293 (1) K
<i>c</i> = 17.079 (3) Å	Needle
<i>V</i> = 1353.4 (4) Å ³	0.30 × 0.12 × 0.10 mm
<i>Z</i> = 4	Colorless
<i>D_x</i> = 1.483 Mg m ⁻³	
<i>D_m</i> not measured	

Data collection

Enraf–Nonius CAD-4 diffractometer	2303 reflections with <i>I</i> > 2σ(<i>I</i>)
ω/2θ scans	<i>R</i> _{int} = 0.049
Absorption correction: empirical via ψ scan of three reflections (North <i>et al.</i> , 1968)	θ _{max} = 68.0°
<i>T</i> _{min} = 0.605, <i>T</i> _{max} = 0.692	<i>h</i> = 0 → 8
2802 measured reflections	<i>k</i> = 0 → 13
2466 independent reflections	<i>l</i> = -20 → 20
	3 standard reflections every 200 reflections
	intensity decay: 4.62%

Refinement

Refinement on <i>F</i> ²	(Δ/σ) _{max} = 0.003
<i>R</i> (<i>F</i>) = 0.033	Δρ _{max} = 0.295 e Å ⁻³
<i>wR</i> (<i>F</i> ²) = 0.085	Δρ _{min} = -0.529 e Å ⁻³
<i>S</i> = 1.107	Extinction correction: none
2466 reflections	Scattering factors from <i>International Tables for Crystallography</i> (Vol. C)
156 parameters	Absolute structure: Flack (1983)
H atoms riding	Flack parameter = 0.03 (3)
<i>w</i> = 1/[σ ² (<i>F</i> _o ²) + (0.05 <i>P</i>) ² + 0.5 <i>P</i>]	
where <i>P</i> = (<i>F</i> _o ² + 2 <i>F</i> _c ²)/3	

Table 1. Selected geometric parameters (Å, °)

Se1—N1	1.865 (3)	O2—C12	1.243 (5)
Se1—C3	1.943 (3)	N1—C12	1.348 (5)
O1—C2	1.425 (4)	N1—C1	1.463 (5)
N1—Se1—C3	87.1 (2)	C12—N1—Se1	126.4 (3)
C12—N1—C1	123.8 (3)	C1—N1—Se1	109.5 (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>
O1—H1...O2'	0.95	1.84	2.769 (4)	166

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

Data collection: *CAD-4 Software* (Schagen *et al.*, 1989). Cell refinement: *CAD-4 Software*. Data reduction: *TEXSAN* (Molecular Structure Corporation, 1994). Program(s) used to solve structure: *SAPI91* (Fan, 1991). Program(s) used to refine structure: *SHELXL93* (Sheldrick, 1993). Molecular graphics: *TEXSAN*. Software used to prepare material for publication: *SHELXL93*.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: FG1405). Services for accessing these data are described at the back of the journal.

References

- Allen, F. H. & Kennard, O. (1993). *Chem. Des. Autom. News*, **8**, 131–137.
- Allen, F. H., Kennard, O., Watson, D. G., Brammer, L., Orpen, A. G. & Taylor, R. (1992). *International Tables for Crystallography*, Vol. C, edited by A. J. C. Wilson, pp. 691–706. Dordrecht/Boston/London: Kluwer Academic Publishers.
- Back, T. G. & Dyck, B. P. (1997). *J. Am. Chem. Soc.* **119**, 2079–2083.
- Burk, R. F. (1994). In *Selenium in Biology and Human Health*. New York: Springer-Verlag.
- Detty, M. R., Friedman, A. E. & Oseroff, A. R. (1994). *J. Org. Chem.* **59**, 8245–8250.
- Engman, L., Andersson, C., Morgenstern, R., Cotgreave, I. A., Andersson, C. M. & Hallberg, A. (1994). *Tetrahedron*, **50**, 2929–2938.
- Engman, L., Stern, D., Cotgreave, I. A. & Andersson, C. M. (1992). *J. Am. Chem. Soc.* **114**, 9737–9743.
- Engman, L., Stern, D., Pelcman, M. & Andersson, C. M. (1994). *J. Org. Chem.* **59**, 1973–1979.
- Fan, H.-F. (1991). *SAPI91. Structure Analysis Program with Intelligent Control*. Rigaku Corporation, Tokyo, Japan.
- Fischer, H. & Dereu, N. (1987). *Bull. Soc. Chim. Belg.* **96**, 757–768.
- Flack, H. D. (1983). *Acta Cryst.* **A39**, 876–881.
- Flohé, L. (1985). *Curr. Top. Cell Regul.* **27**, 473–478.
- Fong, M. C. & Schiesser, C. H. (1995). *Tetrahedron Lett.* **36**, 7329–7332.
- Galet, V., Bernier, J.-L., Hénichart, J.-P., Lesieur, D., Abadie, C., Rochette, L., Lindenbaum, A., Chalas, J., Renaud de la Faverie, J.-F., Pfeiffer, B. & Renard, P. (1994). *J. Med. Chem.* **37**, 2903–2911.
- Ganther, H. E. (1975). *Chem. Scr.* **8a**, 79–84.
- Ganther, H. E. & Kraus, R. J. (1984). *Methods in Enzymology*, edited by S. P. Colowick & N. O. Kaplan, pp. 593–602. New York: Academic Press.
- Haenen, G. R. M. M., De Rooij, B. M., Vermeulen, N. P. E. & Bast, A. (1990). *Mol. Pharmacol.* **37**, 412–422.

- Hoekstra, W. G. (1974). *Trace Element Metabolism in Animals 2*, edited by W. G. Hoekstra, J. W. Suttie, H. E. Ganther & W. Mertz. Baltimore: University Park Press.
- Iwaoka, M. & Tomoda, S. (1994). *J. Am. Chem. Soc.* **116**, 2557–2561.
- Jacquemin, P. V., Christiaens, L. E., Renson, M. J., Evers, M. J. & Dereu, N. (1992). *Tetrahedron Lett.* **33**, 3863–3866.
- Johnson, C. K. (1976). *ORTEP*. Report ORNL-5138. Oak Ridge National Laboratory, Tennessee, USA.
- Molecular Structure Corporation (1994). *TEXSAN. Single Crystal Structure Analysis Software*. MSC, 3200 Research Forest Drive, The Woodlands, TX 77381, USA.
- North, A. C. T., Phillips, D. C. & Mathews, F. S. (1968). *Acta Cryst.* **A24**, 351–359.
- Reich, H. J. & Jasperse, C. P. (1987). *J. Am. Chem. Soc.* **109**, 5549–5551.
- Schagen, J. D., Straver, L., van Meurs, F. & Williams, G. (1989). *CAD-4 Software*. Delft Instruments X-ray Diffraction, PO Box 811, 2600 AV Delft, The Netherlands.
- Shamberger, R. J. (1983). In *Biochemistry of Selenium*. New York: Plenum Press.
- Sheldrick, G. M. (1993). *SHELXL93. Program for the Refinement of Crystal Structures*. University of Göttingen, Germany.
- Tappel, A. L. (1984). *Curr. Top. Cell Regul.* **24**, 87–97.
- Vessman, K., Ekström, M., Berglund, M., Andersson, C. M. & Engman, L. (1995). *J. Org. Chem.* **60**, 4461–4467.
- Wilson, S. R., Zucker, P. A., Huang, R.-R. C. & Spector, A. (1989). *J. Am. Chem. Soc.* **111**, 5936–5939.

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1-Acetyl-2-thiohydantoin

JOSÉ S. CASAS,^a ALFONSO CASTIÑEIRAS,^a DELFINA COUCE,^b
NURIA PLAYÁ,^a JOSÉ SORDO^a AND JOSÉ M. VARELA^a

^aUniversidad de Santiago de Compostela, Departamento de Química Inorgánica, Facultad de Farmacia, Campus Universitario Sur, E-15706 Santiago de Compostela, Spain, and ^bUniversidad de Vigo, Departamento de Química Inorgánica, Lagoas – Marcosende, E-36200 Vigo, Spain.
E-mail: qiac01@usc.es

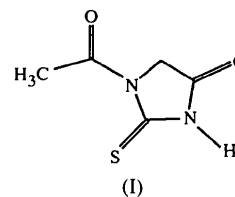
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Abstract

In the title compound (1-acetyl-4-oxoimidazolidine-2-thione, C₅H₆N₂O₂S), the plane of the acetyl group forms an angle of 6.7° with the essentially planar thiohydantoin ring. N—H···O hydrogen bonds create quasi-planar chains of molecules along the *y* axis.

Comment

The structure of the title compound, (I), has been established as part of a study of the synthesis and characterization of metal complexes of 2-thiohydantoin and its derivatives (Casas *et al.*, 1995).



The molecular structure of (I) is shown in Fig. 1. The N1—C1—N2—C3—C2 ring and the peripheral S, C4 and O3 atoms define a plane (r.m.s. deviation 0.013 Å), as does the *N*-acetyl N1—C4(=O3)—C5 fragment (r.m.s. deviation 0.006 Å). The angle between the two planes is 6.7°.

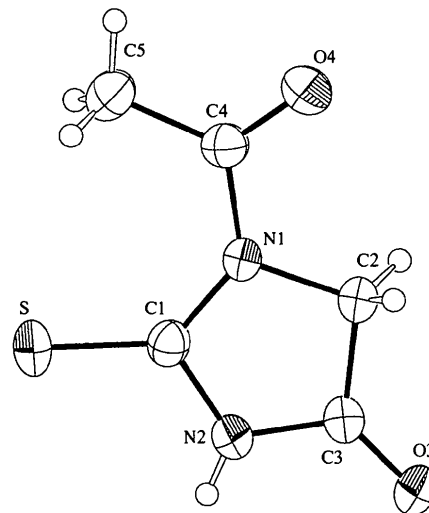


Fig. 1. The molecular structure of the title compound showing the atom-labelling scheme and 50% probability displacement ellipsoids.

The bond lengths and angles in the acetyl fragment are similar to those found in 1-acetyl-2-[1-(acetylthio)ethyl]thiohydantoin (MacKay *et al.*, 1992), although in this C2-substituted thiohydantoin, the angle between the thiohydantoin ring and the acetyl group is 12°. In (I), the thiohydantoin ring bond lengths differing most from those found in 2-thiohydantoin (Devillanova *et al.*, 1987; Walker *et al.*, 1969) are those of C1—N1 and N1—C2, which are longer in the acetyl derivative. The internal ring angle most affected by *N*-acetylation is C2—N1—C1, which widens slightly to approximately the same value as in 1-acetyl-2-[1-(acetylthio)ethyl]thiohydantoin (MacKay *et al.*, 1992). *N*-Acetylation also affects the external angles flanking the C=S group, with N1—C1—S becoming wider and N2—C1—S narrower.

The N2—H2 bond and the O3 atom are involved in a hydrogen bond [N2—H2 0.77 (3), H2···O3ⁱ 2.08 (4), N2···O3ⁱ 2.849 (3) Å and N2—H2···O3ⁱ 170 (3)°; sym-