

S = 0.78

6093 reflections

484 parameters

H atoms treated by a
mixture of independent
and constrained refinement

Extinction correction: none

Scattering factors from
*International Tables for
Crystallography* (Vol. C)

Absolute structure:

Flack (1983)

Flack parameter = -0.06 (6)

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Table 1. Selected geometric parameters (Å, °)

C1A—F1A	1.365 (5)	C16A—C17A	1.368 (6)
C1B—F1B	1.366 (5)	C16A—O1A	1.384 (5)
C4A—C7A	1.518 (6)	C16B—C17B	1.383 (6)
C4B—C7B	1.518 (5)	C16B—O1B	1.392 (5)
C7A—C11A	1.537 (5)	C17A—O2A	1.375 (5)
C7A—C8A	1.541 (5)	C17B—O2B	1.370 (4)
C7B—C8B	1.538 (5)	C19A—O1A	1.417 (6)
C7B—C11B	1.539 (5)	C19A—O2A	1.445 (5)
C8A—C9A	1.517 (6)	C19B—O2B	1.418 (5)
C8B—C9B	1.516 (5)	C19B—O1B	1.452 (5)
C9A—N1A	1.496 (5)	C11···H1O4 ⁱ	2.35 (5)
C9B—N1B	1.490 (5)	C12···H2N2	1.94 (5)
C10A—N1A	1.475 (5)	C12···H1N1	2.18 (4)
C10A—C11A	1.523 (6)	N1A—H1N1	0.89 (4)
C10B—N1B	1.483 (5)	N1A—H2N1	1.12 (5)
C10B—C11B	1.542 (5)	N1B—H1N2	0.84 (4)
C11A—C12A	1.524 (5)	N1B—H2N2	1.22 (5)
C11B—C12B	1.501 (5)	O4—H1O4	0.79 (5)
C12A—O3A	1.428 (4)	O4—H2O4	0.99 (7)
C12B—O3B	1.428 (4)	O4···H2N1	1.69 (5)
C13A—O3A	1.381 (4)	O4···N1A	2.772 (6)
C13B—O3B	1.371 (4)		
C4A—C7A—C11A	114.5 (4)	C16A—C17A—O2A	110.0 (4)
C4A—C7A—C8A	111.3 (3)	C18A—C17A—O2A	127.1 (4)
C4B—C7B—C8B	112.5 (3)	O2B—C17B—C18B	128.3 (4)
C4B—C7B—C11B	109.7 (3)	O2B—C17B—C16B	109.9 (3)
O3A—C12A—C11A	105.8 (3)	O1A—C19A—O2A	107.2 (4)
O3B—C12B—C11B	108.7 (3)	O2B—C19B—O1B	107.0 (3)
O3A—C13A—C14A	115.7 (4)	H1N1—N1A—H2N1	112 (4)
O3A—C13A—C18A	122.8 (4)	H1N2—N1B—H2N2	114 (4)
O3B—C13B—C14B	123.3 (4)	H1O4—O4—H2O4	94 (5)
O3B—C13B—C18B	115.2 (4)	N1A—H1N1···C12	174 (4)
C15A—C16A—O1A	128.4 (4)	N1A—H2N1···O4	159 (4)
C17A—C16A—O1A	109.7 (4)	N1B—H2N2···C12	152 (3)
C15B—C16B—O1B	128.7 (4)	O4—H1O4···C11 ⁱⁱ	164 (4)
C17B—C16B—O1B	109.2 (4)	O4—H2O4···C11	143 (6)

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

The positional and isotropic displacement parameters of the H atoms on the piperidinium N atoms and on the water molecule were refined. Other H atoms were placed at calculated positions and were refined with a riding model. All non-H atoms were refined anisotropically.

Data collection: *SMART* (Bruker, 1997). Cell refinement: *SMART*. Data reduction: *SAINT-Plus* (Bruker, 1997). Program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990). Program(s) used to refine structure: *SHELXTL/PC* (Sheldrick, 1997). Molecular graphics: *SHELXTL/PC*. Software used to prepare material for publication: *SHELXTL/PC*.

I thank Schein Pharmaceutical, Danbury, Connecticut, USA, for the sample of the title compound and Christopher Pernin and Paul Jene for assistance.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: FR1162). Services for accessing these data are described at the back of the journal.

Acta Cryst. (1999). **C55**, 434–436

Fromm's methylphenyldithiodimethylketuret

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(Received 9 July 1998; accepted 15 October 1998)

Abstract

The 'keturet' obtained by reacting 1-methyl-1-phenyldithiobiuret with acetone is shown to be 6,6-dimethyl-2-methylanilino-5,6-dihydro-4H-1,3,5-thiadiazine-4-thione, C₁₂H₁₅N₃S₂. The molecule contains a six-membered thiadiazine ring with a distorted half-chair conformation, rather than a previously proposed four-membered ring structure. Molecular dimensions are normal, and the molecules (two per asymmetric unit) are linked by two N—H···S hydrogen bonds, one fairly strong [H···S 2.39 (3) Å] and one weaker [H···S 2.65 (4) Å].

Comment

Condensation of 1-aryldithiobiurets with ketones, under the influence of dry hydrogen chloride, gives compounds called 'keturets' (Fromm, 1893), some of which can be formulated as hexahydrotriazine derivatives (Fairfull & Peak, 1955). Reaction of 1-methyl-1-phenyldithiobiuret, (I), with acetone gives a keturet which cannot be a hexahydrotriazine (since the disubstitution on N1

Experimental

The 'keturet' was prepared from 1-methyl-1-phenyldithio-biuret and acetone, and crystallized from ethanol, using the procedure previously described by Fromm & Junius (1895) and Fairfull & Peak (1955) (yield 69%, m.p. 424–425 K).

Crystal data

C₁₂H₁₅N₃S₂
M_r = 265.39
 Triclinic
P $\bar{1}$
a = 9.871 (2) Å
b = 11.044 (2) Å
c = 13.117 (3) Å
 α = 87.403 (8)°
 β = 89.625 (4)°
 γ = 67.3872 (11)°
V = 1318.6 (4) Å³
Z = 4
D_r = 1.337 Mg m⁻³
D_m not measured

Mo *K*α radiation
 λ = 0.71069 Å
 Cell parameters from 5212 reflections
 θ = 2–30°
 μ = 0.385 mm⁻¹
T = 180 K
 Plate
 0.35 × 0.20 × 0.05 mm
 Colorless

Data collection

Rigaku/ADSC CCD diffractometer
 CCD scans
 Absorption correction: multi-scan (*d*TREK*; Molecular Structure Corporation, 1997*a*)
T_{min} = 0.85, *T_{max}* = 0.98
 12 370 measured reflections

5927 independent reflections
 2257 reflections with *I* > 3σ(*I*)
R_{int} = 0.050
 θ_{\max} = 30.07°
h = -11 → 13
k = -11 → 15
l = -15 → 18

Refinement

Refinement on *F*²
R(*F*) = 0.040
wR(*F*²) = 0.077
S = 1.020
 5927 reflections
 315 parameters
 H atoms: see below
w = 1/[σ²(*F_o*²)]
 (Δ/σ)_{max} = 0.005

Δρ_{max} = 1.00 e Å⁻³ (0.8 Å from H10, bonded to C6)
 Δρ_{min} = -0.90 e Å⁻³ (0.4 Å from H8, bonded to C6)
 Extinction correction: none
 Scattering factors from *International Tables for Crystallography* (Vol. C)

Table 1. Selected geometric parameters (Å, °)

S1—C2	1.774 (3)	S3—C14	1.775 (3)
S1—C3	1.822 (4)	S3—C15	1.842 (3)
S2—C1	1.711 (3)	S4—C13	1.689 (3)
N1—C1	1.324 (4)	N4—C13	1.350 (4)
N2—C2	1.307 (4)	N5—C14	1.307 (4)
C2—S1—C3	96.7 (2)	C14—S3—C15	95.7 (2)

Table 2. Hydrogen-bonding geometry (Å, °)

D—H...A	D—H	H...A	D...A	D—H...A
N1—H1...S4 ⁱ	1.02 (3)	2.39 (3)	3.353 (3)	157 (2)
N4—H16...S2 ⁱⁱ	0.96 (3)	2.65 (4)	3.522 (3)	152 (3)

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

The *U_{ij}* values are quite small (low-temperature data), but physically very reasonable, with atoms near the centers of the molecules having the lowest displacement parameters and those on the periphery the highest. The highest correlation

coefficient in the refinement is 0.51, for *U₂₂* and *U₁₂* of atom C12 (see *Comment* for a discussion of the pseudosymmetry). The higher than usual value of *R_{int}* resulted from lower than usual crystal quality. All parameters were refined for the two H atoms bonded to N; the other H atoms were placed on calculated sites, with C—H = 0.98 Å and *U*(H) equal to 1.2 times *U* of the C atom to which they were bonded. Methyl group H atoms were placed from a difference synthesis.

Data collection: *d*TREK* (Molecular Structure Corporation, 1997*a*). Cell refinement: *d*TREK*. Data reduction: *d*TREK*. Program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1993). Program(s) used to refine structure: *TEXSAN* (Molecular Structure Corporation, 1997*b*). Software used to prepare material for publication: *TEXSAN*.

We thank the Natural Sciences and Engineering Research Council of Canada for financial support.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: FR1159). Services for accessing these data are described at the back of the journal.

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Acta Cryst. (1999). **C55**, 436–439

Two dioxane derivatives of anthraquinone

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(Received 17 September 1998; accepted 6 November 1998)

Abstract

The structures of 1-methoxy-4-(2-methylprop-2-enyl-oxy)-2-[(2*R*,6*R*)-4,4,6-trimethyl-1,3-dioxan-2-yl]anthraquinone, C₂₆H₂₈O₆, and 4-hydroxy-3-(2-methylprop-2-