

C(18)—C(11)—C(12)—C(13)	—61.0 (6)
C(11)—C(12)—C(13)=C(14)	—21.8 (8)
C(12)—C(13)=C(14)—C(15)	12.7 (7)
C(13)=C(14)—C(15)—C(16)	69.5 (6)
C(14)—C(15)—C(16)—C(17)	—60.7 (6)
C(15)—C(16)—C(17)=C(18)	—22.1 (8)
C(16)—C(17)=C(18)—C(11)	13.0 (7)
C(17)=C(18)—C(11)—C(12)	69.9 (6)

Table 3. Geometrical characteristics of hydrogen bonds (Å, °)

	N—H...S	H...S	N—H...S
(Thiourea) ₃ —1,5-cyclooctadiene* (173 K)	3.366(3)—3.457(3)	2.44(4)—2.66(3)	151(3)—172(3)
Pure thiourea† (110 K)	3.348(4)—3.417(6)	2.354(8)—2.415(10)	169.5(5)—170.5(7)
Pure thiourea† (298 K)	3.394(4)—3.526(8)	2.397(6)—3.055(12)	133.7(7)—168.9(5)

* This work.

† Elcombe & Taylor (1968).

The structure was solved by direct methods and difference Fourier synthesis using *SHELXS86* (Sheldrick, 1985). The parameters were refined with *SHELX76* (Sheldrick, 1976).

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Lists of structure factors, anisotropic displacement parameters, H-atom coordinates, complete geometry and least-squares-planes data have been deposited with the IUCr (Reference: BK1026). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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1-Cyclohexyl-2-cyclohexylamino-5(4*H*)-imidazolone

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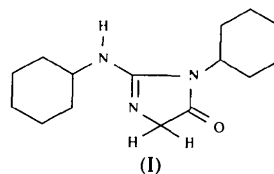
(Received 22 February 1994; accepted 6 June 1994)

Abstract

The structure of the title compound, C₁₅H₂₅N₃O, a product of the reaction of glycine ethyl ester with *N,N'*-dicyclohexylcarbodiimide, was elucidated by X-ray diffraction. The imidazolone ring is planar and the two cyclohexyl rings adopt chair conformations.

Comment

It has been reported (Muramatsu, 1961) that glycine ethyl ester reacts with *N,N'*-dicyclohexylcarbodiimide to form 1-cyclohexyl-2-cyclohexylamino-5(4*H*)-imidazolone, (I). The product was studied by IR spectroscopy and the structure proposed on the basis of analogy with guanidine formation observed for related reactions (DeTar, Silverstein & Rogers, 1966).



The title compound was synthesized by the method of Merrifield, Gisin & Bach (1977) and identified by comparison of its melting point with that given by Muramatsu (1961). A drawing of the molecule showing displacement ellipsoids is presented in Fig. 1.

Atom C4 is involved in three C—N bonds [C4—N1 1.405 (3), C4—N2 1.288 (2) and C4—N3 1.348 (6) Å] which are different from the bond lengths observed for fully delocalized π bonds ($d_{\text{opt}} = 1.334$ Å; Krygowski, 1993). The distances are similar to the values expected for localized bonds (Allen *et al.*, 1987). This suggests low (if any) delocalization of π electrons in this region.

The title structure has an intramolecular hydrogen bond: N3...N2ⁱ 3.058 (8), H3N...N2ⁱ 2.23 (5) Å, N3—H3N...N2ⁱ 159 (4)° [symmetry code: (i) $y - \frac{1}{2}$, $x - \frac{1}{2}$, $z - \frac{1}{2}$]. Its energy is estimated as 2.8 kJ using the empirical formula of Krygowski *et al.* (1991).

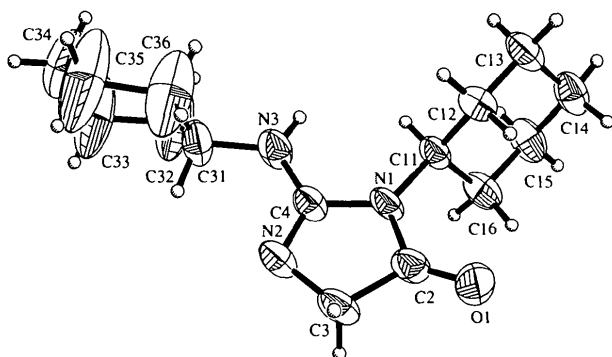


Fig. 1. View of the title molecule showing the atomic numbering system. Displacement ellipsoids are plotted at the 50% probability level.

Experimental

Crystal data

C₁₅H₂₅N₃O

M_r = 263.38

Trigonal

R3c

a = 17.158 (2) Å

α = 116.66 (2)°

V = 2344.1 (5) Å³

Z = 6

D_x = 1.119 Mg m⁻³

Cu Kα radiation

λ = 1.5418 Å

Cell parameters from 25

reflections

θ = 10–15°

μ = 0.560 mm⁻¹

T = 293 (2) K

Prism

0.33 × 0.30 × 0.28 mm

Colourless

Data collection

Kuma KM-4 diffractometer

ω/2θ scans

Absorption correction:

none

4024 measured reflections

2705 independent reflections

1688 observed reflections

[*I* > 2σ(*I*)]

R_{int} = 0.0415

θ_{max} = 70.07°

h = -20 → 0

k = -8 → 20

l = -9 → 15

2 standard reflections

monitored every 100

reflections

intensity decay: <5%

Refinement

Refinement on *F*²

R(*F*) = 0.0493

wR(*F*²) = 0.1482

S = 1.080

2705 reflections

191 parameters

w = 1/[σ²(*F_o*²) + (0.0878*P*)² + 0.0449*P*]

where *P* = (*F_o*² + 2*F_c*²)/3

(Δ/σ)_{max} = -0.031

Δρ_{max} = 0.198 e Å⁻³

Δρ_{min} = -0.281 e Å⁻³

Atomic scattering factors

from *International Tables*

for *Crystallography* (1992,

Vol. C, Tables 4.2.6.8 and

6.1.1.4)

Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters (Å²)

$$U_{eq} = (1/3)\sum_i\sum_j U_{ij}a_i^*a_j^*a_i \cdot a_j$$

	<i>x</i>	<i>y</i>	<i>z</i>	<i>U_{eq}</i>
N1	0.1626 (3)	0.2728 (2)	0.7415 (2)	0.0417 (6)
C2	0.2453 (3)	0.4137 (3)	0.8226 (3)	0.0472 (7)
O1	0.2191 (3)	0.4428 (3)	0.7720 (3)	0.0668 (7)
C3	0.3715 (2)	0.5180 (2)	0.98431 (15)	0.0519 (8)

N2	0.3478 (2)	0.4253 (2)	0.98449 (15)	0.0476 (6)
C4	0.2280 (2)	0.2896 (2)	0.84543 (15)	0.0411 (6)
C11	0.0300 (2)	0.1344 (2)	0.57589 (15)	0.0410 (7)
C12	-0.1028 (3)	0.0877 (4)	0.5060 (3)	0.0483 (7)
C13	-0.2391 (4)	-0.0569 (4)	0.3354 (4)	0.0578 (9)
C14	-0.2028 (4)	-0.0405 (4)	0.2732 (4)	0.0586 (9)
C15	-0.0709 (4)	0.0062 (4)	0.3433 (4)	0.0613 (9)
C16	0.0667 (4)	0.1493 (4)	0.5142 (3)	0.0535 (8)
N3	0.1634 (3)	0.1677 (3)	0.7964 (3)	0.0542 (7)
C31	0.2190 (4)	0.1799 (4)	0.9013 (4)	0.0605 (9)
C32	0.2150 (10)	0.0805 (10)	0.8543 (9)	0.124 (2)
C33	0.2781 (13)	0.0963 (13)	0.9675 (12)	0.149 (3)
C34	0.1876 (11)	0.0535 (10)	0.9730 (11)	0.134 (2)
C35	0.1876 (18)	0.1513 (19)	1.0189 (17)	0.195 (4)
C36	0.1400 (13)	0.1527 (14)	0.9189 (13)	0.148 (3)

Table 2. Selected geometric parameters (Å, °)

N1—C2	1.388 (4)	C13—C14	1.522 (12)
N1—C4	1.405 (3)	C14—C15	1.509 (10)
N1—C11	1.478 (2)	C15—C16	1.525 (6)
C2—O1	1.203 (9)	N3—C31	1.461 (10)
C2—C3	1.505 (5)	C31—C36	1.44 (3)
C3—N2	1.456 (5)	C31—C32	1.48 (2)
N2—C4	1.288 (2)	C32—C33	1.55 (3)
C4—N3	1.348 (6)	C33—C34	1.46 (3)
C11—C16	1.509 (8)	C34—C35	1.50 (4)
C11—C12	1.523 (7)	C35—C36	1.55 (4)
C12—C13	1.525 (6)		
C2—N1—C4	107.3 (3)	C11—C12—C13	110.5 (3)
C2—N1—C11	124.1 (3)	C14—C13—C12	111.1 (3)
C4—N1—C11	128.6 (2)	C15—C14—C13	111.0 (6)
O1—C2—N1	126.6 (3)	C14—C15—C16	112.0 (3)
O1—C2—C3	128.8 (3)	C11—C16—C15	111.2 (3)
N1—C2—C3	104.6 (3)	C4—N3—C31	121.0 (3)
N2—C3—C2	106.8 (2)	C36—C31—N3	112.2 (8)
C4—N2—C3	105.7 (3)	C36—C31—C32	112.9 (13)
N2—C4—N3	124.7 (2)	N3—C31—C32	109.4 (7)
N2—C4—N1	115.5 (1)	C31—C32—C33	109.7 (14)
N3—C4—N1	119.8 (2)	C34—C33—C32	110.3 (15)
N1—C11—C16	111.6 (2)	C33—C34—C35	110.0 (2)
N1—C11—C12	111.3 (2)	C34—C35—C36	112.0 (2)
C16—C11—C12	112.1 (2)	C31—C36—C35	113.0 (18)

Initially, we found the unit-cell parameters in the monoclinic system [*a* = 18.014 (5), *b* = 29.204 (8), *c* = 9.525 (4) Å, *β* = 110.66 (3)°], but could not solve the structure, so all the parameters were transformed to the trigonal cell. The structure was solved by direct methods and refined by full-matrix least-squares calculations. The H atoms were placed in calculated positions and refined with isotropic displacement parameters. The N3 H atom was obtained from a Δρ map and refined with an isotropic displacement parameter. Programs used were *SHELX76* (Sheldrick, 1976), *SHELXS86* (Sheldrick, 1990) and *SHELXL93* (Sheldrick, 1993).

We thank Professor T. M. Krygowski for helpful discussions.

Lists of structure factors, anisotropic displacement parameters, H-atom coordinates, complete geometry, including bond distances and angles involving H atoms, and torsion angles have been deposited with the IUCr (Reference: KA1083). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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1-[4-(Dimethylamino)phenyl]-3-(1-pyrenyl)hexamethyltrisilane

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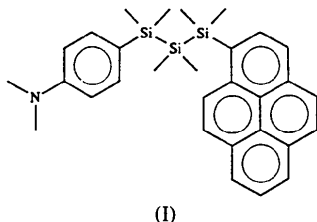
(Received 23 June 1994; accepted 12 September 1994)

Abstract

The trisilane chain C1—Si1—Si2—Si3—C23 of the title compound, 1-[4-(dimethylamino)phenyl]-1,1,2,2,3,3-hexamethyl-3-(1-pyrenyl)trisilane, C₃₀H₃₇NSi₃, adopts a +synclinal, –antiperiplanar conformation. The two aromatic systems make an angle of 56.2 (1)° with each other.

Comment

Donor–acceptor substituted oligosilanes can be used as model systems for the study of through-bond interactions. The structure determination of the title compound, (I), was undertaken in order to establish the mutual orientation of the two aromatic systems, pyrene and phenyl, and to determine the conformation of the trisilane chain.



The molecular structure with atomic labelling is depicted in Fig. 1. The trisilane chain adopts a +synclinal [C1—Si1—Si2—Si3 = 48.2 (2)°], –antiperiplanar

[Si1—Si2—Si3—C23 = –177.9 (1)°] conformation, with the two aromatic systems pointing away from each other.

The angle between the best planes through the pyrene and phenyl atoms is 56.2 (1)°. The maximum deviation from the best plane through the pyrene atoms is 0.030 (3) Å (for atom C1) and through the phenyl atoms the maximum deviation is 0.004 (3) Å. This structure is comparable to the recently reported structure of 1,3-bis(1-pyrenyl)hexamethyltrisilane (Van Meervelt, 1993), where the angle between the pyrene ring systems is 71° with a –antiperiplanar, +synclinal conformation for the trisilane chain.

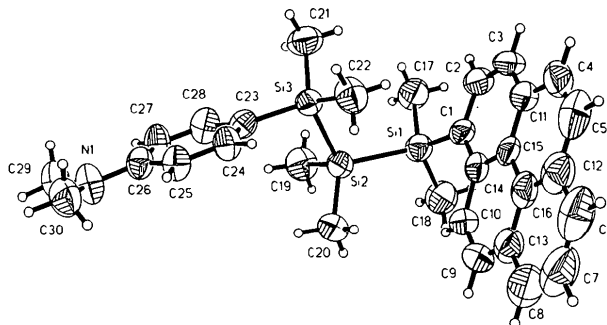


Fig. 1. View (SHELXTL/PC; Sheldrick 1990) of the title molecule with atomic labelling. Displacement ellipsoids are shown at 50% probability levels and H atoms are drawn as small circles of arbitrary size.

Experimental

Crystal data

C₃₀H₃₇NSi₃
M_r = 495.88
 Monoclinic
*P*2₁/*n*
a = 7.8980 (10) Å
b = 26.463 (4) Å
c = 14.099 (2) Å
 β = 99.69 (2)°
V = 2904.7 (7) Å³
Z = 4
D_x = 1.134 Mg m^{–3}

Cu *K*α radiation
 λ = 1.54178 Å
 Cell parameters from 25 reflections
 θ = 7–23°
 μ = 1.626 mm^{–1}
T = 289 (2) K
 Block
 0.20 × 0.20 × 0.10 mm
 Yellow
 Crystal source: grown from cyclohexane

Data collection

Siemens *P4/PC* diffractometer
 θ – 2θ scans
 Absorption correction: empirical
 T_{\min} = 0.479, T_{\max} = 0.991
 4081 measured reflections
 3030 independent reflections
 1827 observed reflections
 $[I > 2\sigma(I)]$

R_{int} = 0.0520
 θ_{max} = 50.41°
 h = –1 → 7
 k = –1 → 26
 l = –14 → 14
 3 standard reflections monitored every 100 reflections
 intensity decay: none