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#### Key indicators

Single-crystal X-ray study T = 123 KMean  $\sigma$ (C–C) = 0.004 Å R factor = 0.059 wR factor = 0.123 Data-to-parameter ratio = 18.1

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

# 3-Nitro-1-(triisopropylsilyl)-1H-pyrrole

The nitration of 1-(triisopropylsilyl)-1*H*-pyrrole leads to a mixture of products following partial acid cleavage of the triisopropylsilyl protecting group. Structural determination showed the isolated products to be the title compound,  $C_{13}H_{24}N_2O_2Si$ , and 2,4-dinitropyrrole. In the solid state, the title compound exists as discrete molecules with only weak  $C-H\cdots$ nitro hydrogen bonds between them.

#### Comment

The quest for N-alkyl-substituted pyrroles bearing a nitro group at position three has led us to the use of the cleavable and bulky triisoproplysilyl (TIPS) group. This work is in connection with our research into minor-groove binding compounds (Khalaf et al., 2004). These compounds are analogues of naturally occurring distamycin and netropsin, which bind primarily to the adenine-thymine-rich minor groove of DNA. 3-Nitropyrrole (Bray et al., 1990) was required as a precursor; however, the straightforward nitration of pyrrole favours position two. To prevent the nitration from occurring at position two of the pyrrole, a TIPS protecting group was first attached to the nitrogen of the pyrrole ring. This led to the nitro group being directed to position three (as anticipated), giving rise to the desired product (I) (Fig. 1) in 32% yield. However, during the course of the reaction, and due to the presence of acetic acid, the TIPS group cleaved. The removal of the TIPS group allowed the nitration reaction to occur once again. However, this time it occurred at position two, leading to the formation of the undesired product, 2,4-dinitropyrrole (II), in 19% yield.



Despite the large number of substituted pyrroles reported in the Cambridge Structural Database (Version 5.27 with updates to May 2006; Allen, 2002), a search found only nine relevant 3-nitro derivatives. All of the ring bond lengths and angles (Table 1) of (I) fall within the ranges found for these nine structures, with the exception that in (I) the C2-N1-C5

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#### Figure 1

The molecular structure of (I), with displacement ellipsoids drawn at the 50% probability level.

angle is slightly below the literature range  $[106.8 (2)^{\circ} cf. 107.7-110.2^{\circ}]$ . This is presumably due to the bulk and inductive effects of the attached TIPS group as no such relevant *N*-silyl substituent was found in the database. All of the literature nitro groups are approximately coplanar with their pyrrole rings; it can be seen from Table 1 that this is also the case for (I).

In the absence of obvious hydrogen-bonding groups, nitro aromatics often display nitro-to-nitro interactions of the type described by Wozniak *et al.* (1994); however, none is found in (I). The only intermolecular interactions of any note are weak  $C-H\cdots O$  contacts utilizing both  $sp^2$  and  $sp^3$  CH groups (see Table 2). This can be rationalized as a consequence of the bulky triisopropylsilyl group, which ensures that the ring systems are widely separated from each other.

## **Experimental**

A solution of cupric nitrate trihydrate (2.70 g, 11.2 mmol) in acetic anhydride (20 ml) was cooled to 273 K and 1-(triisopropylsilyl)-1*H*pyrrole (2.50 g, 11.2 mmol) was added dropwise with stirring. The ice bath was removed and stirring was continued for 1 h at room temperature. The reaction mixture was poured slowly over a saturated sodium hydrogen carbonate solution with stirring. After extraction with diethyl ether the organic layer was collected, dried (MgSO<sub>4</sub>) and the solvent was removed under reduced pressure. The residue was subjected to flash chromatography on silica gel, using ethyl acetate/hexane (1:12) to elute (I). The solvents were removed under reduced pressure and a slow stream of air was passed over the Crystal data

 $C_{13}H_{24}N_2O_2Si$  Z = 4 

  $M_r = 268.43$   $D_x = 1.144 \text{ Mg m}^{-3}$  

 Monoclinic,  $P2_1/n$  Mo K $\alpha$  radiation

 a = 9.6924 (5) Å
  $\mu = 0.15 \text{ mm}^{-1}$  

 b = 15.9437 (10) Å
 T = 123 (2) K

 c = 10.1267 (6) Å
 Cut fragment, colourless

  $\beta = 95.089$  (4)°
  $0.30 \times 0.12 \times 0.08 \text{ mm}$  

 V = 1558.74 (16) Å<sup>3</sup>
  $\Lambda^3$ 

Data collection

Nonius KappaCCD diffractometer  $\omega$  and  $\varphi$  scans Absorption correction: none 13088 measured reflections

Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.059$   $wR(F^2) = 0.123$  S = 1.01 3060 reflections 169 parameters H-atom parameters constrained

#### 3060 independent reflections 1685 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.108$ $\theta_{\text{max}} = 26.0^{\circ}$

$$\begin{split} w &= 1/[\sigma^2(F_o^2) + (0.0471P)^2 \\ &+ 0.0035P] \\ \text{where } P &= (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{\text{max}} &= 0.001 \\ \Delta\rho_{\text{max}} &= 0.34 \text{ e } \text{ Å}^{-3} \\ \Delta\rho_{\text{min}} &= -0.36 \text{ e } \text{ Å}^{-3} \end{split}$$

Table 1

Selected geometric parameters (Å, °).

Si1-N1	1.805 (2)	C2-C3	1.365 (3)
N1-C2	1.365 (3)	C3-C4	1.410 (4)
N1-C5	1.395 (3)	C4-C5	1.358 (4)
C2-N1-C5	106.8 (2)	C5-C4-C3	105.3 (2)
C3-C2-N1	108.4 (2)	C4-C5-N1	110.3 (2)
C2-C3-C4	109.1 (2)		
O1-N2-C3-C2	-0.5 (4)	O2-N2-C3-C2	178.9 (2)

Table 2

H	yd	lrogen-	bond	geometry	(A,	°)	).	
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$D - H \cdots A$	<i>D</i> -H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$C5-H5\cdots O2^{i}$ $C10-H10\cdots O1^{ii}$	0.95 1.00	2.47 2.55	3.403 (4) 3.245 (4)	166 126
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Symmetry codes: (i)  $x + \frac{1}{2}, -y + \frac{1}{2}, z + \frac{1}{2}$ ; (ii) -x + 1, -y, -z.

All H atoms were positioned geometrically,  $C-H = 0.95 (Csp^2)$ , 0.98 (CH<sub>3</sub>) or 1.00 Å (Csp<sup>3</sup>), and refined using a riding model  $[U_{iso}(H) = 1.5U_{eq}(C)$  for CH<sub>3</sub> and  $1.2U_{eq}(C)$  for all others].

Data collection: *COLLECT* (Hooft, 1988) and *DENZO* (Otwinowski & Minor, 1997); cell refinement: *DENZO* and *COLLECT*; data reduction: *DENZO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

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