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The authors thank the Instituto de Química de São Carlos for the data collection and CNPq, FINEP and FAPEMIG(Brazil) for financial support, and Dr Dalton L. F. Alves for helpful advice on the isolation of the title compound.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: FG1165). Services for accessing these data are described at the back of the journal.

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Acta Cryst. (1997). C53, 984-985

N-Cyano-*N*'-(4-methoxyphenyl)guanidine

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(Received 14 September 1995; accepted 21 February 1997)

Abstract

An X-ray structural analysis of the title compound, $C_9H_{10}N_4O$, shows a planar guanidine portion, the N-aryl group tilted $ca 57^{\circ}$ relative to the guanidine plane and E geometry about the formal 'imino' bond.

Comment

Monosubstituted guanidines such as nitroguanidine, (2), cyanoguanidine, (3), sulfaguanidine, (4), L-canavanine, (5), and N-methyl-N'-nitro-N-nitrosoguanidine, (6), which bear strongly electron-withdrawing groups, exist as neutral organic molecules preferring the 'imino' tautomeric form [e.g. $(NH_2)_2C=N-NO_2$] in solution and in the solid state (Patai & Rappoport, 1991). In a previous paper, we showed that disubstituted N-aryl-N'-cyanoguanidines in solution in Me₂SO also have the 'imino' structure seen for the title compound, (1) (Cunningham, Wan & Cox, 1994). While crystal structures have been reported for (3) (Hughes, 1940; Begley, Hubberstey & Moore, 1985; Hirshfield & Hope, 1980) and for the other guanidines listed above (Patai & Rappoport, 1991), none are known for N-cyano-N'-substituted guanidines. In this paper, we report the crystal structure of N-cyano-N'-(4-methoxyphenyl)guanidine, (1).



The X-ray crystal structure is shown in Fig. 1 (arbitrary numbering), where it can be seen that the imino tautomer is preferred in the solid state. The cyanoguanidine portion is essentially planar, as shown by the bond angles about C1, N1 and N2 (Table 1) and the C2-N3-C1-N1 and C2-N3-C1-N2 torsion angles. The N1-C1, N2-C1 and N3-C1 bonds which comprise the guanidine system are all of length 1.33(1) Å; these lengths are almost identical to those seen for unsubstituted cyanoguanidine (Begley, Hubberstey & Moore, 1985) and are between the values for C-N and C=N (Allen et al., 1987), indicating that the N-atom lone pairs and the C=N bond shown formally in (1) are completely delocalized over the guanidine system. The bond length (N4-C2) for the cyano group at 1.142 (3) Å is typical, suggesting that the cyano group is not involved in resonance with the guanidine system. Interestingly, the aryl group is twisted out of the guanidine plane by about 57° showing that resonance

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Fig. 1. The molecular structure of (1) with displacement ellipsoids drawn at the 50% probability level.

between the cyanoguanidine portion and the aromatic ring is minimal.

Fig. 1 also illustrates that the geometry about the formal C = N bond is E; this preference may be due to packing as much as to steric factors.

Experimental

The title compound was prepared by reaction of 4-methoxyaniline with sodium dicyanamide in aqueous hydrochloric acid and recrystallized from ethanol.

Crystal data

 $I > 3\sigma(I)$

| $C_9H_{10}N_4O$ | Mo $K\alpha$ radiation |
|--|--|
| $M_r = 190.21$ | $\lambda = 0.71073$ Å |
| Triclinic | Cell parameters from 25 |
| P_1^{-} | reflections |
| a = 6.904 (3) Å b = 8.431 (3) Å c = 9.555 (6) Å $\alpha = 111.21 (4)^{\circ}$ $\beta = 92.02 (4)^{\circ}$ $\gamma = 107.74 (4)^{\circ}$ $V = 487 (1) \text{ Å}^{3}$ | $\theta = 7-13^{\circ}$ $\mu = 0.085 \text{ mm}^{-1}$ T = 293 K Square prism $0.5 \times 0.3 \times 0.2 \text{ mm}$ Colourless |
| Z = 2 $D_x = 1.294 \text{ Mg m}^{-3}$ $D_m \text{ not measured}$ Data collection | |
| Enraf-Nonius CAD-4 | $\theta_{\text{max}} = 26.0^{\circ}$ |
| diffractometer | $h = 0 \rightarrow 8$ |
| $\omega/2\theta$ scans | $k = -10 \rightarrow 9$ |
| Absorption correction: none | $l = -11 \rightarrow 11$ |
| 2075 measured reflections | 1 standard reflection |
| 1909 independent reflections | frequency: 60 min |
| 1340 reflections with | intensity decay: 11.0% |

|--|

| Refinement on F | $(\Delta/\sigma)_{max} = 0.16$ |
|--------------------------------------|---|
| R = 0.058 | $\Delta\rho_{max} = 0.19 \text{ e } \text{\AA}^{-3}$ |
| wR = 0.076 | $\Delta\rho_{min} = -0.22 \text{ e } \text{\AA}^{-3}$ |
| S = 1.026 | Extinction correction: none |
| 1340 reflections | Scattering factors from <i>later</i> . |
| 3 = 1.020 1340 reflections | Scattering factors from Inter- |
| 139 parameters | national Tables for X-ray |
| $w = 1/[\sigma^2(F) + (0.5F)^2 + 5]$ | Crystallography (Vol. IV) |

Table 1. Selected geometric parameters (Å, °)

| N1—C1 N2—C1 N3—C1 | 1.328 (3) 1.317 (3) 1.342 (3) | N3—C2 N4—C2 | 1.314 (3) 1.142 (3) |
|--|--|--|--|
| C1N1C11 C1N1H1 C11N1H1 C1N2H2A C1N2H2B H2AN2H2B | 127.9 (2) 117 (2) 115 (2) 125 (2) 122 (2) 112 (3) | C1—N3—C2 N1—C1—N2 N1—C1—N3 N2—C1—N3 N3—C2—N4 | i 19.5 (2) 121.0 (2) 114.5 (2) 124.6 (2) 173.5 (3) |
| C1—N1—C11—C16 C2—N3—C1—N1 | 57 (1) 178 (1) | C2-N3-C1-N2 | 3(1) |

The structure was solved by non-default use of *MULTAN78* (Main *et al.*, 1978) when the correct solution yielded 12 of the atoms in the molecule. A structure factor/Fourier calculation revealed the missing atoms and with H atoms included (except those on N atoms), but not refined, isotropic refinement converged slowly to R = 0.20. Anisotropic refinement of the non-H atoms rapidly converged to the final R of 0.058. In the final refinement, the H atoms attached to N atoms were found from a difference Fourier and were refined isotropically.

Data collection: CAD-4 Software (Enraf-Nonius, 1989). Cell refinement: CAD-4 Software. Data reduction: SDP (Frenz, 1983). Program(s) used to solve structure: SDP and MUL-TAN78. Program(s) used to refine structure: SDP.

The authors are grateful to Zeneca FCMO for financial support (NCW).

Supplementary data for this paper are available from the IUCr electronic archives (Reference: BM1034). Services for accessing these data are described at the back of the journal.

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