Acta Crystallographica Section C

Crystal Structure Communications

ISSN 0108-2701

5-Amino-3-*tert*-butyl-1-(4-nitrophenyl)-1*H*-pyrazole forms hydrogenbonded sheets of alternating $R_2^2(20)$ and $R_6^6(32)$ rings

John N. Low, at Justo Cobo, Rodrigo Abonia, Jairo Quiroga and Christopher Glidewell at Christopher Glidewell

^aDepartment of Chemistry, University of Aberdeen, Meston Walk, Old Aberdeen AB24 3UE, Scotland, ^bDepartamento de Química Inorgánica y Orgánica, Universidad de Jaén, 23071 Jaén, Spain, ^cGrupo de Investigación de Compuestos Heterocíclicos, Departamento de Química, Universidad de Valle, AA 25360, Cali, Colombia, and ^dSchool of Chemistry, University of St Andrews, Fife KY16 9ST, Scotland

Correspondence e-mail: cg@st-andrews.ac.uk

Received 7 January 2004 Accepted 9 January 2004 Online 10 February 2004

Molecules of the title compound, $C_{13}H_{16}N_4O_2$, are linked by one N-H···O hydrogen bond [H···O = 2.47 Å, N···O = 3.326 (2) Å and N-H···O = 166°] and one N-H···N hydrogen bond [H···N = 2.19 Å, N···N = 3.063 (2) Å and N-H···N = 173°] into sheets containing alternating R_2^2 (20) and R_6^6 (32) rings, both types of which are centrosymmetric.

Comment

The title compound, (I), is a frequently used intermediate for the synthesis of interesting fused pyrazolo[3,4-b]pyridine or similar derivatives. In the molecules of the title compound (Fig. 1), the dihedral angle between the aryl and heterocyclic rings is 33.7 (2)°, possibly as a result, in part, of the tendency of atom H12 and the amine group bonded to atom C5 to avoid one another. On the other hand, the nitro group is almost coplanar with the adjacent aryl ring, with a dihedral angle between the aryl plane and the $C-NO_2$ plane of only 2.2 (2)°. As well as the dihedral angle between the rings, the bond lengths provide no evidence for any delocalization of the lone pair at atom N1 into the nitro group to form the polarized structure (Ia). In particular, the C-N distances exocyclic to the aryl ring (Table 1) are typical of their type (Allen et al., 1987), while the C-C bond distances in the aryl ring span the small range 1.378 (3)-1.396 (3) Å, with no sign of quinonoidtype bond fixation. However, there is marked bond fixation within the heterocyclic ring (Table 1), where the C3-C4 and C4-C5 distances differ significantly, as do the N2-C3 and N1-C5 distances. Consistent with the pyramidal nature of atom N5, the C5—N5 bond is long for its type. The dihedral angle between the two rings and the conformation adopted by the *tert*-butyl group render the molecules chiral; however, the space group accommodates equal numbers of the two enantiomeric forms.

$$Me_3C$$
 N
 NO_2
 NH_2
 NH_2
 NH_2
 NH_2
 NH_2
 NH_2
 NH_2

The molecules of (I) are linked into sheets by one $N-H\cdots O$ hydrogen bond and one $N-H\cdots N$ hydrogen bond (Table 2), and the sheet formation is readily analysed by considering the action of each of these hydrogen bonds in turn.

Amine atom N5 in the molecule at (x, y, z) acts as a hydrogen-bond donor, via atom H5B, to nitro atom O142 in the molecule at (1 - x, 1 - y, 1 - z), thereby generating a centrosymmetric $R_2^2(20)$ (Bernstein et al., 1995) dimer centred at $(\frac{1}{2}, \frac{1}{2}, \frac{1}{2})$. The action of the N-H···N hydrogen bond is to link each of these dimers to four adjacent dimers, thereby forming a continuous sheet. Amine atoms N5 at (x, y, z) and (1-x, 1-y, 1-z) both lie in the dimer centred at $(\frac{1}{2}, \frac{1}{2}, \frac{1}{2})$, and these atoms act as hydrogen-bond donors, this time via atom H5A, to atoms N52 in the molecules at $(x, \frac{1}{2} - y, -\frac{1}{2} + z)$ and $(1-x, \frac{1}{2}+y, \frac{3}{2}-z)$, respectively, which themselves lie in the dimers centred at $(\frac{1}{2}, 0, 0)$ and $(\frac{1}{2}, 1, 1)$, respectively. Likewise, atoms N52 at (x, y, z) and (1 - x, 1 - y, 1 - z)accept hydrogen bonds from atoms N5 in the molecules at (x, y) $\frac{1}{2} - y$, $\frac{1}{2} + z$) and (1 - x, $\frac{1}{2} + y$, $\frac{1}{2} - z$), which themselves lie in the dimers centred at $(\frac{1}{2}, 0, 1)$ and $(\frac{1}{2}, 1, 0)$. Propagation by the space group of these two hydrogen bonds then generates a (100) sheet built from alternating $R_2^2(20)$ and $R_6^6(32)$ rings, where both ring types are centrosymmetric (Fig. 2). If the $R_2^2(20)$ dimers are regarded as the nodes of the resulting net then this is of (4,4)-type (Batten & Robson, 1998).

There are no direction-specific interactions between adjacent sheets; in particular, there are neither $C-H\cdots\pi_{arene}$ hydrogen bonds nor aromatic $\pi-\pi$ stacking interactions

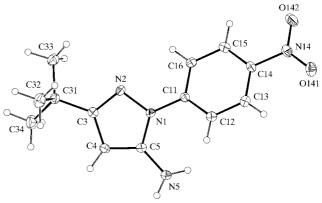


Figure 1The molecule of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level.

[‡] Postal address: Department of Electrical Engineering and Physics, University of Dundee, Dundee DD1 4HN, Scotland.

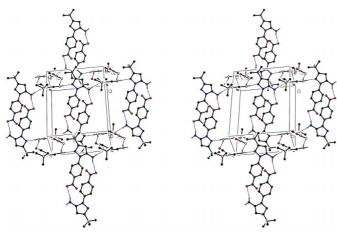


Figure 2 A stereoview of part of the crystal structure of (I), showing the formation of a (100) sheet built from alternating $R_2^2(20)$ and $R_6^6(32)$ rings.

present in the structure. The $R_6^6(32)$ rings are largely occupied by a pair of tert-butyl groups, one pendent from each of the adjacent (100) sheets, and there are no solvent-accessible voids in the structure.

Experimental

A suspension of equimolar amounts (16 mmol) of 4-nitrophenylhydrazine and 4,4-dimethyl-3-oxopentanenitrile in concentrated HCl (10 ml) and water (20 ml) was stirred at room temperature for 10 min. The volume was then reduced to half the original, concentrated HCl (20 ml) was added, and the resulting mixture was heated with stirring for 15 min. The mixture was then cooled and neutralized with aqueous ammonia. The solid product, (I), was collected and recrystallized from water (yield 90%, m.p. 445 K).

Crystal data

$C_{13}H_{16}N_4O_2$	$D_x = 1.315 \text{ Mg m}^{-3}$
$M_r = 260.30$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 3001
a = 5.9581 (2) Å	reflections
b = 18.3064 (11) Å	$\theta = 3.4-27.5^{\circ}$
c = 12.0776 (7) Å	$\mu = 0.09 \text{ mm}^{-1}$
$\beta = 93.4150 (12)^{\circ}$	T = 120 (2) K
$V = 1314.98 (12) \text{ Å}^3$	Lath, yellow
Z = 4	$0.20 \times 0.10 \times 0.04 \text{ mm}$

Data collection	
Nonius KappaCCD diffractometer φ scans, and ω scans with κ offsets Absorption correction: multi-scan (SORTAV; Blessing, 1995, 1997) $T_{\min} = 0.977, T_{\max} = 0.996$ 14 938 measured reflections 3001 independent reflections	1700 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.071$ $\theta_{\text{max}} = 27.5^{\circ}$ $h = -7 \rightarrow 7$ $k = -23 \rightarrow 21$ $l = -15 \rightarrow 15$
Refinement	

Refinement on F^2	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.052$	$w = 1/[\sigma^2(F_o^2) + (0.0613P)^2]$
$wR(F^2) = 0.124$	where $P = (F_o^2 + 2F_c^2)/3$
S = 0.96	$(\Delta/\sigma)_{\rm max} < 0.001$
3001 reflections	$\Delta \rho_{\text{max}} = 0.22 \text{ e Å}^{-3}$
175 parameters	$\Delta \rho_{\min} = -0.28 \text{ e Å}^{-3}$

Space group $P2_1/c$ was uniquely assigned from the systematic absences. All H atoms were located from difference maps, which showed clearly the pyramidal nature of the amine group based on

Table 1 Selected geometric parameters (Å, °).

N1-N2	1.393 (2)	N1-C11	1.415 (2)
N2-C3	1.327 (2)	C5-N5	1.366(2)
C3-C4	1.411 (3)	C14-N14	1.466 (2)
C4-C5	1.373 (3)	N14-O141	1.226(2)
C5-N1	1.370(2)	N14-O142	1.229 (2)
N2-N1-C11-C12	147.35 (17)	C13-C14-N14-O141	2.8 (3)
C5-N1-C11-C12	-36.0(3)	C13-C14-N14-O142	-178.3(2)
	* *		

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

D $ H$ $\cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - H \cdot \cdot \cdot A$
N5−H5 <i>A</i> ···N2 ⁱ	0.88	2.19	3.063 (2)	173
N5−H5 <i>B</i> ···O142 ⁱⁱ	0.88	2.47	3.326 (2)	166

Symmetry codes: (i) $x, \frac{1}{2} - y, z - \frac{1}{2}$; (ii) 1 - x, 1 - y, 1 - z.

atom N5, and all H atoms were then treated as riding atoms, with C-H distances of 0.95 (aromatic) or 0.98 Å (methyl), and N-H distances of 0.88 Å.

Data collection: KappaCCD Server Software (Nonius, 1997); cell refinement and data reduction: DENZO-SMN (Otwinowski & Minor, 1997); program(s) used to solve structure: OSCAIL (McArdle, 2003) and SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: PLATON (Spek, 2003); software used to prepare material for publication: SHELXL97 and PRPKAPPA (Ferguson, 1999).

X-ray data were collected at the EPSRC X-ray Crystallographic Service, University of Southampton, England; the authors thank the staff for all their help and advice. JNL thanks NCR Self-Service, Dundee, for grants that have provided computing facilities for this work. JC thanks the Ministerio de Educación, Cultura y Deportes (Programa de Cooperación con Iberoamérica, AECI) of Spain. RA and JQ thank COLCIENCIAS and the Universidad de Valle for financial support.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: SK1695). Services for accessing these data are described at the back of the journal.

References

Allen, F. H., Kennard, O., Watson, D. G., Brammer, L., Orpen, A. G. & Taylor, R. (1987). J. Chem. Soc. Perkin Trans. 2, pp. S1-19.

Batten, S. R. & Robson, R. (1998). Angew. Chem. Int. Ed. 37, 1460-1494. Bernstein, J., Davis, R. E., Shimoni, L. & Chang, N.-L. (1995). Angew. Chem. Int. Ed. Engl. 34, 1555-1573.

Blessing, R. H. (1995). Acta Cryst. A51, 33-37.

Blessing, R. H. (1997). J. Appl. Cryst. 30, 421-426.

Ferguson, G. (1999). PRPKAPPA. University of Guelph, Canada.

McArdle, P. (2003). OSCAIL for Windows. Version 10. Crystallography Centre, Chemistry Department, NUI Galway, Ireland.

Nonius (1997). KappaCCD Server Software. Windows 3.11 Version. Nonius BV. Delft. The Netherlands.

Otwinowski, Z. & Minor, W. (1997). Methods in Enzymology, Vol. 276, Macromolecular Crystallography, Part A, edited by C. W. Carter Jr & R. M. Sweet, pp. 307-326. New York: Academic Press.

Sheldrick, G. M. (1997). SHELXS97 and SHELXL97. University of Göttingen, Germany.

Spek, A. L. (2003). J. Appl. Cryst. 36, 7-13.