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# **Crystal Structure Communications**

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5-[N-(1H-Benzotriazol-1-ylmethyl)-amino]-3-tert-butyl-1-phenylpyrazole: sheets built from N— $H \cdots N$ , C— $H \cdots N$  and C— $H \cdots \pi$ (pyrazole) interactions

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In the title compound,  $C_{20}H_{22}N_6$ , the molecules are linked into a chain of rings by N-H···N [H···N 2.16 Å, N···N 2.950 (3) Å and N-H···N 149°] and C-H···N [H···N 2.55 Å, C···N 3.481 (3) Å and C-H···N 165°] hydrogen bonds, and these chains are linked into sheets by means of C-H··· $\pi$ (pyrazole) interactions.

### Comment

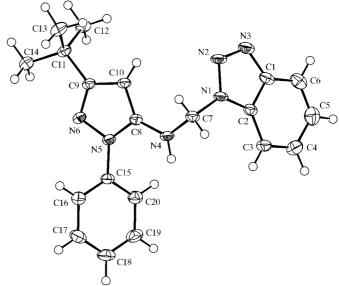
The title compound, (I) (Fig. 1), was prepared as a precursor for the synthesis of new hydrogenated pyrazolo-pyridines using the benzotriazole methodology (Katritzky *et al.*, 1995, 1998; Abonia *et al.*, 2001).

The C-C bond distances (Table 1) in the fused arene ring show some evidence for bond fixation, with bonds C3-C4 and C5-C6 being shorter than the neighbouring bonds; likewise, the N2-N3 bond is markedly shorter than both N1-N2 and N5-N6. The pattern of C-N distances is of interest; the C7-N1 and C7-N4 distances, which involve planar three-connected N and four-connected C atoms, are significantly different; likewise, the C1-N3 and C9-N6 distances, which involve two-connected N and planar three-connected C atoms, differ significantly, consistent with the occurrence of single and double bonds, respectively. On the other hand, the three distances C2-N1, C8-N4 and C8-N5, which involve two planar three-connected atoms, span only a small range.

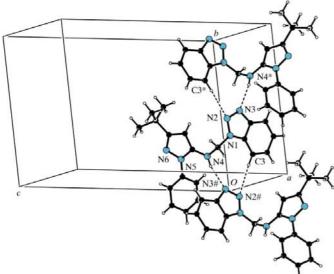
The supramolecular structure of (I) is determined by a combination of  $N-H\cdots N$  and  $C-H\cdots N$  hydrogen bonds and  $C-H\cdots \pi(pyrazole)$  interactions (Table 2). The amino

atom N4 at (x, y, z) acts as hydrogen-bond donor to triazole N3 at  $(2 - x, -\frac{1}{2} + y, \frac{1}{2} - z)$ , while N4 at  $(2 - x, -\frac{1}{2} + y, \frac{1}{2} - z)$ , in turn, acts as donor to N3 at (x, -1 + y, z), so producing a C(6) chain generated by the  $2_1$  screw axis along  $(1, y, \frac{1}{4})$ . This is

augmented by a fairly weak  $C-H\cdots N$  hydrogen bond; C3 at (x, y, z) acts as donor to triazole N2 at  $(2-x, -\frac{1}{2}+y, \frac{1}{2}-z)$ , producing a C(5) chain generated by the same screw axis as before, so that the resulting chain of rings running parallel to the [010] direction (Fig. 2) has the descriptor  $C(5)C(6)[R_2^2(9)]$  (Bernstein *et al.*, 1995). Two chains of this type run through each unit cell, in the domains 0.06 < z < 0.44 and 0.56 < z < 0.94, and within each domain, adjacent chains are linked into a (001) sheet by  $C-H\cdots\pi$  (pyrazole) interactions. Atom C19 in the molecule at (x, y, z), which lies in the hydrogen-bonded chain along  $(1, y, \frac{1}{4})$ , forms a short  $C-H\cdots\pi$  contact (Table 2) with the centroid, Cg2, of the pyrazole ring of the molecule at  $(1-x, -\frac{1}{2}+y, \frac{1}{2}-z)$ , which itself lies in the hydrogen-bonded



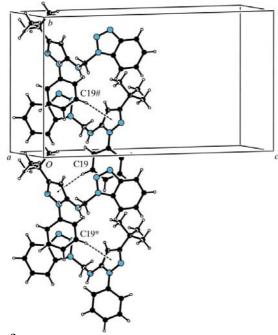
**Figure 1**The molecule of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level.



**Figure 2** Part of the crystal structure of (I), showing the formation of a chain of rings along [010]. Atoms marked with an asterisk (\*) or hash (#) are at the symmetry positions  $(2-x, -\frac{1}{2}+y, \frac{1}{2}-z)$  and  $(2-x, \frac{1}{2}+y, \frac{1}{2}-z)$ , respectively.

chain along  $(\frac{1}{2}, y, \frac{1}{4})$  (Fig. 3). Propagation of this interaction thus links all of the [010] chains in a given domain of z into a single sheet. Two sheets run through each unit cell, but there are no direction-specific interactions between adjacent sheets.

The structure of the isomeric compound (II) (see *Scheme* above) has recently been reported in space group  $P2_1/c$  (Low *et al.*, 2002). The supramolecular structure was described in terms of a single hydrogen bond, N51—H51···N43<sup>i</sup> [symmetry



**Figure 3** Part of the crystal structure of (I), showing the linking of the [010] chains by  $C-H\cdots\pi$ (pyrazole) interactions. Atoms marked with an asterisk (\*) or hash (#) are at the symmetry positions  $(1-x, -\frac{1}{2}+y, \frac{1}{2}-z)$  and  $(1-x, \frac{1}{2}+y, \frac{1}{2}-z)$ , respectively.

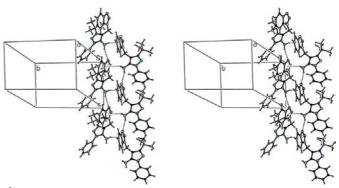


Figure 4 Stereoview of part of the crystal structure of (II), showing the formation of a molecular ladder built from  $N-H\cdots N$  and  $N-H\cdots \pi$  (triazole) interactions.

code: (i) x, y, 1+z], forming C(8) chains by translation. However, re-examination of the structure of (II) in the light of the above analysis for (I), shows that there is also a short N—  $H \cdots \pi$ (triazole) interaction, involving the N51—H52 bond, which does not participate in conventional hydrogen bonding, and the centroid (Cg2) of the triazole ring;  $H52 \cdots Cg2^{ii}$  2.72 Å, N51··· $Cg2^{ii}$  3.477 (2) Å and N51—H52··· $Cg2^{ii}$  146° [symmetry code: (ii) x,  $\frac{1}{2} - y$ ,  $\frac{1}{2} + z$ ]. This interaction produces a zigzag chain parallel to [001], generated by the c-glide plane at  $y = -\frac{1}{4}$ . The combination of N—H···N and N—H··· $\pi$  interactions thus generates a double chain, or molecular ladder (Fig. 4).

Since  $C-H\cdots\pi$  interactions most frequently involve benzenoid rings as acceptors (Desiraju & Steiner, 1999; Suezawa *et al.*, 2001), with rather few examples involving

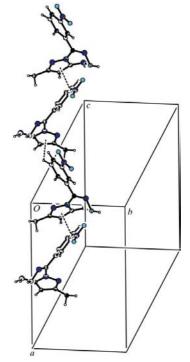


Figure 5 Part of the crystal structure of (IV), showing the formation of a  $C-H\cdots\pi$ (pyrazole) chain consisting of type 1 molecules only.

## organic compounds

heterocycles, we have briefly investigated the occurrence of  $C-H\cdots\pi$ (pyrazole) interactions by examining the structures of pyrazoles in the Cambridge Structural Database (CSD; Allen & Kennard, 1993) containing the fragment (III), namely a pyrazole ring carrying the same substituent atoms as found in (I). Nine examples were retrieved; the structures of three of these contained only  $C-H\cdots\pi$  (arene) interactions, those of five others contained no  $C-H \cdot \cdot \cdot \pi$  interactions at all, but the structure of the ninth [CSD refcode TIXPEV, (IV); Ramm et al., 1996], which crystallizes in space group  $P2_1/a$  with Z'=2, displays quite short  $C-H\cdots\pi(pyrazole)$  interactions, such that the two independent molecules each form simple chains. For the type 1 molecules, the chain is generated by the a-glide plane at  $y = -\frac{1}{4} [H \cdot \cdot \cdot Cg1^{iii} 2.84 \text{ Å}, C \cdot \cdot \cdot Cg1^{iii} 3.645 (2) \text{ Å} and$  $C-H \cdot \cdot \cdot Cg1^{iii} 127^{\circ}$ ; symmetry code: (iii)  $-\frac{1}{2} + x$ ,  $-\frac{1}{2} - y$ , z] (Fig. 5), while for the type 2 molecules, the chain is generated by the a-glide plane at  $y = \frac{1}{4} [H \cdot \cdot \cdot Cg2^{iv} \ 2.69 \text{ Å}, \ C \cdot \cdot \cdot \cdot Cg2^{iv}]$ 3.537(2) Å and C-H···Cg2<sup>iv</sup> 145°; symmetry code: (iv)  $\frac{1}{2} + x$ ,

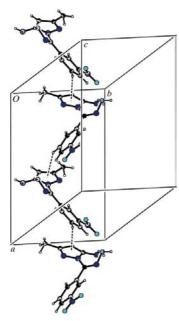


Figure 6 Part of the crystal structure of (IV), showing the formation of a C- $H \cdot \cdot \cdot \pi$ (pyrazole) chain consisting of type 2 molecules only.

 $\frac{1}{2} - y$ , z] (Fig. 6), where Cg1 and Cg2 represent the ring centroids of the pyrazole rings in the molecules of types 1 and 2, respectively.

### **Experimental**

A solution of 5-amino-3-tert-butyl-1-phenylpyrazole (Grandberg et al., 1961; Abonia et al., 2002) (1.00 g, 4.65 mmol) and 1-hydroxymethylbenzotriazole (Burckhalter et al., 1952) (0.70 g, 4.69 mmol) in ethanol (5 ml) was heated under reflux for 3 min. After cooling, the precipitate which formed was filtered off and washed with ethanol, giving (I) as a white solid (84% yield; m.p. 453 K; analysis, found: C 69.3, H 6.4, N 24.2%;  $C_{20}H_{22}N_6$  requires: C 69.3, H 6.4, N 24.3%). Crystals suitable for single-crystal X-ray diffraction were grown from a solution in ethanol.

Crystal data

 $C_{20}H_{22}N_6$ Mo  $K\alpha$  radiation  $M_r = 346.44$ Cell parameters from 2418 Orthorhombic, P2<sub>1</sub>2<sub>1</sub>2<sub>1</sub> reflections a = 7.2380 (5) Å $\theta = 1.9 – 27.4^{\circ}$  $\mu = 0.08~\mathrm{mm}^{-1}$ b = 11.9743 (9) Åc = 21.2062 (16) ÅT = 120 (2) K $V = 1837.9 (2) \text{ Å}^3$ Plate, colourless  $0.30 \times 0.20 \times 0.07 \text{ mm}$ Z = 4 $D_x = 1.252 \text{ Mg m}^{-3}$ 

Data collection

Bruker SMART1000 CCD diffractometer  $R_{\rm int} = 0.030$   $\theta_{\rm max} = 27.4^{\circ}$  Absorption correction: multi-scan  $(SADABS; {\rm Bruker}, 1997)$   $k=-15 \rightarrow 15$   $T_{\rm min} = 0.979, T_{\rm max} = 0.995$   $l=-27 \rightarrow 27$  l=2704 measured reflections  $l=-27 \rightarrow 15$  Intensity decay: negligible 2418 independent reflections

Refinement

 $\begin{array}{lll} \mbox{Refinement on } F^2 & & w = 1/[\sigma^2(F_o^2) + (0.0692P)^2 \\ R[F^2 > 2\sigma(F^2)] = 0.044 & & + 0.2514P] \\ wR(F^2) = 0.108 & & where $P = (F_o^2 + 2F_c^2)/3$ \\ S = 1.07 & (\Delta/\sigma)_{\rm max} < 0.001 \\ 2418 \ {\rm reflections} & \Delta\rho_{\rm max} = 0.32 \ {\rm e \ \AA^{-3}} \\ 238 \ {\rm parameters} & \Delta\rho_{\rm min} = -0.25 \ {\rm e \ \AA^{-3}} \end{array}$ 

Table 1
Selected interatomic distances (Å).

C1-C2	1.397 (3)	N2-N3	1.305 (3)
C2-C3	1.408 (3)	N1-C7	1.468 (3)
C3-C4	1.371 (3)	C7-N4	1.422 (3)
C4-C5	1.415 (4)	N4-C8	1.372 (3)
C5-C6	1.373 (4)	C8-N5	1.359 (3)
C6-C1	1.397 (3)	N5-N6	1.383 (2)
C1-N3	1.376 (3)	N6-C9	1.327 (3)
C2-N1	1.361 (3)	C9-C10	1.412 (3)
N1-N2	1.361(2)	C10-C8	1.388 (3)

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

Cg2 is the centroid of the N5/N6/C8-C10 pyrazole ring.

$D$ $ H$ $\cdot \cdot \cdot A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D$ $ H$ $\cdot \cdot \cdot A$
$ N4-H4A\cdots N3^{i} $ $ C3-H3\cdots N2^{i} $ $ C19-H19\cdots Cg2^{ii} $	0.88	2.16	2.950 (3)	149
	0.95	2.55	3.481 (3)	165
	0.95	2.85	3.779 (3)	167

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

Compound (I) crystallized in the orthorhombic system; space group  $P2_12_12_1$  was assigned uniquely from the systematic absences. H atoms were treated as riding, with C-H distances in the range 0.95-0.99 Å and an N-H distance of 0.88 Å. The Friedel equivalent reflections were merged prior to the final refinements, as the absolute structure could not be assigned.

Data collection: *SMART* (Bruker, 1997); cell refinement: *SMART*; data reduction: *SHELXTL* (Bruker, 1997); program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics:

*PLATON* (Spek, 2002); software used to prepare material for publication: *SHELXL*97 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 which have provided computing facilities for this work.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: SK1550). Services for accessing these data are described at the back of the journal.

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