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***N'*-[*E*]-(*1*-Methyl-*1H*-pyrrol-*2*-yl)methyl- idene]pyridine-*4*-carbohydrazide. Corrigendum**

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The name of one of the authors in the paper by Hussain *et al.* [*Acta Cryst.* (2010), **E66**, o1881] is corrected.

In the paper by Hussain *et al.* (2010), the last author is incorrectly given as 'Muhammad Mazhar'. The correct name of the last author should be 'Mazhar Hussain' as above.

References

Hussain, A., Tahir, M. N., Shafiq, Z., Yaqub, M. & Mazhar, M. (2010). *Acta Cryst.* **E66**, o1881.

N'-[*E*-(1-Methyl-1*H*-pyrrol-2-yl)methylidene]pyridine-4-carbohydrazide

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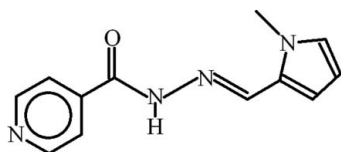
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Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.045; wR factor = 0.127; data-to-parameter ratio = 18.1.

In the title compound, $\text{C}_{12}\text{H}_{12}\text{N}_4\text{O}$, the pyridine and pyrrole rings are inclined at an angle of 29.22 (8) $^\circ$ and an intramolecular $\text{C}-\text{H}\cdots\text{N}$ interaction generates a $S(6)$ ring. In the crystal, molecules are linked by $\text{N}-\text{H}\cdots\text{N}$ hydrogen bonds, forming (010) $C(7)$ chains. The chains are cross-linked by weak $\text{C}-\text{H}\cdots\text{O}$ interactions, which generate $R_2^2(18)$ ring motifs within an infinite sheet. Finally, two $\text{C}-\text{H}\cdots\pi$ interactions are present, where the $\text{C}-\text{H}$ groups are from the pyridine ring and π is the pyrrole ring.

Related literature

For background information on Schiff bases containing heterocyclic rings and for related structures, see: Shafiq *et al.*, (2009*a,b*); Hussain *et al.* (2010) For graph-set notation, see: Bernstein *et al.* (1995).



Experimental

Crystal data

$\text{C}_{12}\text{H}_{12}\text{N}_4\text{O}$

$M_r = 228.26$

Monoclinic, $P2_1/n$

$a = 8.2134$ (3) Å

$b = 10.6740$ (4) Å

$c = 13.1332$ (4) Å

$\beta = 96.938$ (2) $^\circ$

$V = 1142.95$ (7) Å³

$Z = 4$

Mo $K\alpha$ radiation

$\mu = 0.09$ mm⁻¹

$T = 296$ K

$0.24 \times 0.18 \times 0.15$ mm

Data collection

Bruker Kappa APEXII CCD

diffractometer

Absorption correction: multi-scan

(*SADABS*; Bruker, 2005)

$T_{\min} = 0.980$, $T_{\max} = 0.985$

12030 measured reflections

2803 independent reflections

2023 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.029$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.045$

$wR(F^2) = 0.127$

$S = 1.04$

2803 reflections

155 parameters

H-atom parameters constrained

$\Delta\rho_{\text{max}} = 0.28$ e Å⁻³

$\Delta\rho_{\text{min}} = -0.21$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, $^\circ$).

Cg1 is the centroid of the C8–C11/N4 ring.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N2}-\text{H2}\cdots\text{N1}^{\text{i}}$	0.86	2.19	3.0205 (18)	163
$\text{C4}-\text{H4}\cdots\text{O1}^{\text{ii}}$	0.93	2.54	3.3821 (19)	150
$\text{C12}-\text{H12B}\cdots\text{O1}^{\text{iii}}$	0.96	2.55	3.450 (2)	156
$\text{C12}-\text{H12C}\cdots\text{N3}$	0.96	2.36	3.025 (2)	126
$\text{C2}-\text{H2A}\cdots\text{Cg1}^{\text{iv}}$	0.93	2.83	3.3258 (16)	114
$\text{C5}-\text{H5}\cdots\text{Cg1}^{\text{v}}$	0.93	2.71	3.4669 (17)	139

Symmetry codes: (i) $-x + \frac{1}{2}, y + \frac{1}{2}, -z + \frac{1}{2}$; (ii) $x - \frac{1}{2}, -y + \frac{1}{2}, z - \frac{1}{2}$; (iii) $-x + 2, -y + 1, -z + 1$; (iv) $-x + 1, -y + 1, -z + 1$; (v) $-x + \frac{3}{2}, y - \frac{1}{2}, -z + \frac{1}{2}$.

Data collection: *APEX2* (Bruker, 2009); cell refinement: *S SAINT* (Bruker, 2009); data reduction: *S SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997) and *PLATON* (Spek, 2009); software used to prepare material for publication: *WinGX* (Farrugia, 1999) and *PLATON*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB5530).

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