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4-[2-[Methyl(2-pyridyl)amino]ethoxy]-benzaldehyde

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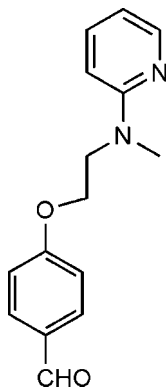
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Key indicators: single-crystal X-ray study; $T = 294$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.035; wR factor = 0.101; data-to-parameter ratio = 8.5.

In the title compound, $\text{C}_{15}\text{H}_{16}\text{N}_2\text{O}_2$, the dihedral angle between the aromatic ring planes is $81.37(13)^\circ$. In the crystal structure, $\pi-\pi$ stacking [centroid-centroid separation = $3.890(2)$ Å] helps to establish the packing.

Related literature

For the synthesis, see: Cantello *et al.* (1994).

Experimental

Crystal data

$\text{C}_{15}\text{H}_{16}\text{N}_2\text{O}_2$
 $M_r = 256.30$
 Monoclinic, $P2_1$
 $a = 6.1995(19)$ Å
 $b = 9.416(3)$ Å
 $c = 11.714(3)$ Å
 $\beta = 98.328(5)^\circ$

$V = 676.6(3)$ Å³
 $Z = 2$
 Mo $K\alpha$ radiation
 $\mu = 0.09$ mm⁻¹
 $T = 294(2)$ K
 $0.24 \times 0.22 \times 0.10$ mm

Data collection

Bruker SMART 1000 CCD diffractometer
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.980$, $T_{\max} = 0.992$

3897 measured reflections
 1477 independent reflections
 980 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.036$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.035$
 $wR(F^2) = 0.101$
 $S = 0.98$
 1477 reflections
 174 parameters

1 restraint
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.11$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.11$ e Å⁻³

Data collection: SMART (Bruker, 1997); cell refinement: SAINT (Bruker, 1997); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 1997); software used to prepare material for publication: SHELXTL.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB2594).

References

- Bruker (1997). SMART (Version 5.611), SAINT (Version 5.01) and SHELXTL (Version 6.10). Bruker AXS Inc., Madison, Wisconsin, USA.
 Cantello, B. C. C., Cawthorne, M. A., Cottam, G. P., Duff, P. T., Haigh, D., Hindley, R. M., Lister, C. A., Smith, S. A. & Thurlby, P. L. (1994). *J. Med. Chem.* **37**, 3977–3985.
 Sheldrick, G. M. (1996). SADABS. University of Göttingen, Germany.
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supplementary materials

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4-{2-[Methyl(2-pyridyl)amino]ethoxy}benzaldehyde

J. Hou, C. Zheng, H. Li, X. Zhang and C. Li

Comment

The title compound, (I), (Fig. 1) is an intermediate in the synthesis of the anti-diabetic drug rosiglitazone.

The dihedral angle between the N1/C1—C5 pyridine ring and C9—C14 phenyl ring is 81.37 (13)°. In the crystal (Fig. 2), weak aromatic π - π stacking helps to establish the packing with a centroid-centroid separation of 3.890 (2) Å for the N1/C1—C5 and C9ⁱ—C14ⁱ ($i = 1 - x, 1/2 + y, 1 - z$) rings.

Experimental

The title compound was prepared according to the method of Cantello *et al.* (1994). Colourless blocks of (I) were obtained by recrystallization from dichromethane.

Refinement

Anomalous dispersion was negligible and Friedel pairs were merged before refinement.

All the H atoms were positioned geometrically (C—H = 0.93–0.97 Å) and refined as riding with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ or $1.5U_{\text{eq}}(\text{methyl C})$.

Figures

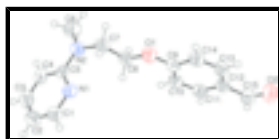


Fig. 1. The molecular structure of (I). Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii.

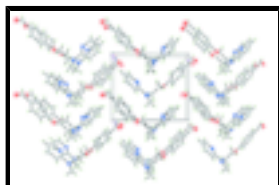


Fig. 2. The crystal packing for (I).

4-{2-[Methyl(2-pyridyl)amino]ethoxy}benzaldehyde

Crystal data

C₁₅H₁₆N₂O₂

$M_r = 256.30$

$F_{000} = 272$

$D_x = 1.258 \text{ Mg m}^{-3}$

supplementary materials

Monoclinic, $P2_1$

Hall symbol: P 2yb

$a = 6.1995 (19) \text{ \AA}$

$b = 9.416 (3) \text{ \AA}$

$c = 11.714 (3) \text{ \AA}$

$\beta = 98.328 (5)^\circ$

$V = 676.6 (3) \text{ \AA}^3$

$Z = 2$

Melting point: 338-339 K

Mo $K\alpha$ radiation

$\lambda = 0.71073 \text{ \AA}$

Cell parameters from 1123 reflections

$\theta = 2.8\text{--}23.7^\circ$

$\mu = 0.09 \text{ mm}^{-1}$

$T = 294 (2) \text{ K}$

Block, colorless

$0.24 \times 0.22 \times 0.10 \text{ mm}$

Data collection

Bruker SMART 1000 CCD
diffractometer

Radiation source: fine-focus sealed tube

Monochromator: graphite

$T = 294(2) \text{ K}$

φ and ω scans

Absorption correction: multi-scan
(SADABS; Sheldrick, 1996)

$T_{\min} = 0.980$, $T_{\max} = 0.992$

3897 measured reflections

1477 independent reflections

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

$R_{\text{int}} = 0.036$

$\theta_{\max} = 26.5^\circ$

$\theta_{\min} = 1.8^\circ$

$h = -7 \rightarrow 7$

$k = -6 \rightarrow 11$

$l = -11 \rightarrow 14$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.035$

$wR(F^2) = 0.101$

$S = 0.98$

1477 reflections

174 parameters

1 restraint

Primary atom site location: structure-invariant direct
methods

Secondary atom site location: difference Fourier map

Hydrogen site location: inferred from neighbouring
sites

H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.058P)^2]$$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} < 0.001$

$\Delta\rho_{\max} = 0.11 \text{ e \AA}^{-3}$

$\Delta\rho_{\min} = -0.11 \text{ e \AA}^{-3}$

Extinction correction: SHELXL97 (Sheldrick, 1997),

$$F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$$

Extinction coefficient: 0.144 (13)

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > 2\sigma(F^2)$ is used only for calculat-

ing R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.4167 (3)	0.5026 (2)	0.66454 (15)	0.0590 (5)
O2	0.1555 (4)	0.1192 (3)	1.0759 (2)	0.0905 (8)
N1	0.6212 (3)	0.4415 (2)	0.33536 (18)	0.0537 (6)
N2	0.7653 (3)	0.6252 (2)	0.45508 (18)	0.0554 (6)
C1	0.6382 (5)	0.3521 (3)	0.2477 (2)	0.0672 (8)
H1	0.5234	0.2897	0.2258	0.081*
C2	0.8123 (6)	0.3465 (4)	0.1885 (3)	0.0751 (10)
H2	0.8161	0.2825	0.1284	0.090*
C3	0.9817 (5)	0.4388 (4)	0.2208 (3)	0.0731 (9)
H3	1.1034	0.4373	0.1829	0.088*
C4	0.9714 (4)	0.5337 (3)	0.3093 (2)	0.0571 (8)
H4	1.0853	0.5968	0.3312	0.068*
C5	0.7866 (4)	0.5338 (3)	0.36600 (19)	0.0437 (6)
C6	0.9392 (5)	0.7225 (4)	0.4982 (3)	0.0798 (10)
H6A	1.0639	0.6696	0.5332	0.120*
H6B	0.8910	0.7843	0.5546	0.120*
H6C	0.9780	0.7780	0.4355	0.120*
C7	0.5769 (4)	0.6150 (3)	0.5166 (2)	0.0552 (7)
H7A	0.4488	0.5907	0.4624	0.066*
H7B	0.5512	0.7067	0.5499	0.066*
C8	0.6100 (4)	0.5051 (3)	0.6109 (2)	0.0530 (7)
H8A	0.6349	0.4125	0.5788	0.064*
H8B	0.7354	0.5293	0.6670	0.064*
C9	0.4029 (4)	0.4081 (3)	0.7518 (2)	0.0467 (6)
C10	0.5581 (4)	0.3040 (3)	0.7876 (2)	0.0524 (7)
H10	0.6831	0.2961	0.7527	0.063*
C11	0.5236 (4)	0.2129 (3)	0.8756 (2)	0.0533 (7)
H11	0.6271	0.1434	0.8992	0.064*
C12	0.3387 (4)	0.2219 (3)	0.9301 (2)	0.0492 (6)
C13	0.1869 (4)	0.3291 (3)	0.8948 (2)	0.0554 (7)
H13	0.0641	0.3386	0.9312	0.066*
C14	0.2173 (4)	0.4204 (3)	0.8069 (2)	0.0527 (7)
H14	0.1146	0.4906	0.7839	0.063*
C15	0.3057 (5)	0.1201 (4)	1.0211 (2)	0.0661 (8)
H15	0.4108	0.0497	1.0380	0.079*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0572 (11)	0.0594 (12)	0.0629 (11)	0.0153 (9)	0.0170 (8)	0.0084 (11)
O2	0.0982 (16)	0.0866 (17)	0.0965 (16)	0.0028 (16)	0.0467 (14)	0.0149 (16)
N1	0.0592 (14)	0.0494 (13)	0.0526 (13)	-0.0098 (12)	0.0081 (10)	-0.0044 (12)

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N2	0.0621 (13)	0.0538 (14)	0.0506 (12)	-0.0115 (13)	0.0093 (10)	-0.0076 (13)
C1	0.089 (2)	0.0523 (18)	0.0591 (18)	-0.0121 (17)	0.0078 (16)	-0.0050 (16)
C2	0.104 (3)	0.064 (2)	0.0608 (19)	0.013 (2)	0.0237 (18)	-0.0032 (17)
C3	0.075 (2)	0.090 (2)	0.0601 (18)	0.022 (2)	0.0290 (16)	0.018 (2)
C4	0.0484 (14)	0.0660 (19)	0.0570 (17)	-0.0023 (14)	0.0086 (12)	0.0161 (16)
C5	0.0480 (13)	0.0414 (14)	0.0402 (13)	-0.0013 (12)	0.0018 (10)	0.0092 (13)
C6	0.088 (2)	0.074 (2)	0.0747 (18)	-0.027 (2)	0.0024 (17)	-0.0139 (19)
C7	0.0631 (16)	0.0519 (16)	0.0516 (14)	0.0118 (15)	0.0113 (12)	0.0012 (14)
C8	0.0527 (14)	0.0530 (16)	0.0548 (15)	0.0069 (13)	0.0128 (11)	-0.0026 (15)
C9	0.0500 (14)	0.0436 (15)	0.0460 (14)	0.0049 (12)	0.0053 (11)	-0.0079 (13)
C10	0.0495 (14)	0.0536 (16)	0.0566 (16)	0.0088 (13)	0.0161 (12)	-0.0047 (14)
C11	0.0547 (15)	0.0506 (16)	0.0534 (15)	0.0110 (14)	0.0032 (12)	-0.0025 (15)
C12	0.0504 (14)	0.0507 (15)	0.0461 (14)	-0.0032 (14)	0.0053 (11)	-0.0066 (14)
C13	0.0473 (15)	0.0605 (18)	0.0593 (17)	-0.0024 (14)	0.0110 (12)	-0.0060 (15)
C14	0.0425 (14)	0.0538 (18)	0.0614 (17)	0.0072 (13)	0.0065 (13)	-0.0025 (15)
C15	0.0736 (19)	0.0639 (19)	0.0610 (17)	-0.0011 (17)	0.0107 (15)	0.0025 (18)

Geometric parameters (Å, °)

O1—C9	1.367 (3)	C6—H6C	0.9600
O1—C8	1.431 (3)	C7—C8	1.506 (4)
O2—C15	1.205 (3)	C7—H7A	0.9700
N1—C1	1.343 (3)	C7—H7B	0.9700
N1—C5	1.353 (3)	C8—H8A	0.9700
N2—C5	1.373 (3)	C8—H8B	0.9700
N2—C6	1.449 (4)	C9—C10	1.394 (3)
N2—C7	1.462 (3)	C9—C14	1.403 (4)
C1—C2	1.366 (4)	C10—C11	1.382 (4)
C1—H1	0.9300	C10—H10	0.9300
C2—C3	1.373 (5)	C11—C12	1.393 (3)
C2—H2	0.9300	C11—H11	0.9300
C3—C4	1.376 (4)	C12—C13	1.401 (4)
C3—H3	0.9300	C12—C15	1.470 (4)
C4—C5	1.405 (3)	C13—C14	1.375 (4)
C4—H4	0.9300	C13—H13	0.9300
C6—H6A	0.9600	C14—H14	0.9300
C6—H6B	0.9600	C15—H15	0.9300
C9—O1—C8	119.0 (2)	C8—C7—H7B	109.2
C1—N1—C5	117.7 (2)	H7A—C7—H7B	107.9
C5—N2—C6	120.8 (2)	O1—C8—C7	107.4 (2)
C5—N2—C7	120.3 (2)	O1—C8—H8A	110.2
C6—N2—C7	118.5 (2)	C7—C8—H8A	110.2
N1—C1—C2	124.6 (3)	O1—C8—H8B	110.2
N1—C1—H1	117.7	C7—C8—H8B	110.2
C2—C1—H1	117.7	H8A—C8—H8B	108.5
C1—C2—C3	117.7 (3)	O1—C9—C10	124.5 (2)
C1—C2—H2	121.2	O1—C9—C14	115.7 (2)
C3—C2—H2	121.2	C10—C9—C14	119.7 (2)
C2—C3—C4	120.1 (3)	C11—C10—C9	119.1 (2)

C2—C3—H3	119.9	C11—C10—H10	120.5
C4—C3—H3	119.9	C9—C10—H10	120.5
C3—C4—C5	119.1 (3)	C10—C11—C12	122.1 (2)
C3—C4—H4	120.4	C10—C11—H11	119.0
C5—C4—H4	120.4	C12—C11—H11	119.0
N1—C5—N2	116.9 (2)	C11—C12—C13	118.1 (2)
N1—C5—C4	120.8 (2)	C11—C12—C15	120.2 (3)
N2—C5—C4	122.3 (2)	C13—C12—C15	121.8 (2)
N2—C6—H6A	109.5	C14—C13—C12	120.8 (2)
N2—C6—H6B	109.5	C14—C13—H13	119.6
H6A—C6—H6B	109.5	C12—C13—H13	119.6
N2—C6—H6C	109.5	C13—C14—C9	120.2 (2)
H6A—C6—H6C	109.5	C13—C14—H14	119.9
H6B—C6—H6C	109.5	C9—C14—H14	119.9
N2—C7—C8	111.9 (2)	O2—C15—C12	126.2 (3)
N2—C7—H7A	109.2	O2—C15—H15	116.9
C8—C7—H7A	109.2	C12—C15—H15	116.9
N2—C7—H7B	109.2		
C5—N1—C1—C2	1.1 (4)	N2—C7—C8—O1	-179.4 (2)
N1—C1—C2—C3	-0.1 (5)	C8—O1—C9—C10	-5.3 (4)
C1—C2—C3—C4	-0.7 (5)	C8—O1—C9—C14	174.6 (2)
C2—C3—C4—C5	0.4 (4)	O1—C9—C10—C11	-178.9 (2)
C1—N1—C5—N2	179.4 (2)	C14—C9—C10—C11	1.3 (4)
C1—N1—C5—C4	-1.3 (3)	C9—C10—C11—C12	-0.2 (4)
C6—N2—C5—N1	176.5 (3)	C10—C11—C12—C13	-1.2 (4)
C7—N2—C5—N1	3.4 (3)	C10—C11—C12—C15	178.2 (3)
C6—N2—C5—C4	-2.7 (4)	C11—C12—C13—C14	1.6 (4)
C7—N2—C5—C4	-175.8 (2)	C15—C12—C13—C14	-177.8 (2)
C3—C4—C5—N1	0.7 (4)	C12—C13—C14—C9	-0.5 (4)
C3—C4—C5—N2	179.9 (2)	O1—C9—C14—C13	179.2 (2)
C5—N2—C7—C8	84.3 (3)	C10—C9—C14—C13	-0.9 (4)
C6—N2—C7—C8	-89.0 (3)	C11—C12—C15—O2	178.3 (3)
C9—O1—C8—C7	178.3 (2)	C13—C12—C15—O2	-2.3 (4)

Fig. 1

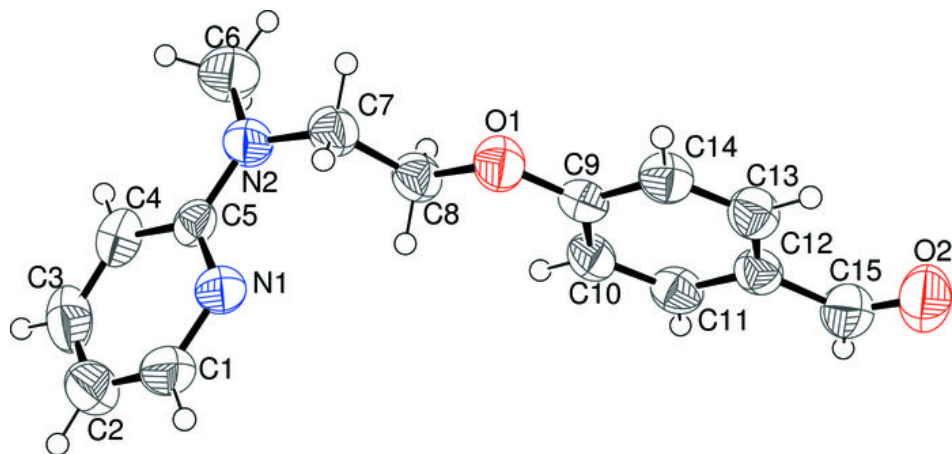


Fig. 2

