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4-(2-Oxa-6-azaspiro[3.3]hept-6-yl)-benzonitrile

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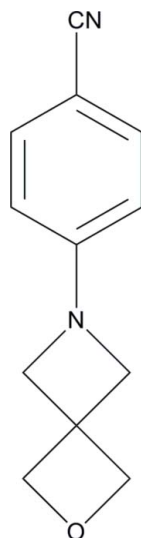
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 Key indicators: single-crystal X-ray study; $T = 113$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.041; wR factor = 0.099; data-to-parameter ratio = 16.4.

In the title compound, $\text{C}_{12}\text{H}_{12}\text{N}_2\text{O}$, the azetidine ring (r.m.s. deviation = 0.021 Å) and the oxetane ring (r.m.s. deviation = 0.014 Å) are nearly perpendicular to each other [dihedral angle = 89.7 (1)°]. The azetidine ring is twisted out of the plane of the benzene ring by 18.3 (1)°. In the crystal structure, molecules are linked to form chains along the c axis by $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds.

Related literature

The title compound is a key intermediate to synthesize (pyrrolo[3,4- c]pyrazol-3-yl)benzamide derivatives. For the anti-tumor effect of these derivatives, see: Fancelli *et al.* (2005).



Experimental

Crystal data

$\text{C}_{12}\text{H}_{12}\text{N}_2\text{O}$	$V = 1002.2$ (7) Å ³
$M_r = 200.24$	$Z = 4$
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation
$a = 9.484$ (4) Å	$\mu = 0.09$ mm ⁻¹
$b = 11.033$ (4) Å	$T = 113$ K
$c = 10.419$ (4) Å	$0.60 \times 0.60 \times 0.27$ mm
$\beta = 113.186$ (5)°	

Data collection

Rigaku AFC10/Saturn 724-Plus diffractometer	2234 independent reflections
7662 measured reflections	1872 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.029$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.041$	136 parameters
$wR(F^2) = 0.099$	H-atom parameters constrained
$S = 1.00$	$\Delta\rho_{\text{max}} = 0.30$ e Å ⁻³
2234 reflections	$\Delta\rho_{\text{min}} = -0.16$ e Å ⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{C4}-\text{H4}\cdots\text{O1}^i$	0.95	2.47	3.371 (2)	159

 Symmetry code: (i) $x, y, z + 1$.

Data collection: *CrystalClear* (Rigaku, 2008); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *PLATON* (Spek, 2009); software used to prepare material for publication: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CI5062).

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supporting information

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4-(2-Oxa-6-azaspiro[3.3]hept-6-yl)benzotrile

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S1. Experimental

A DMSO solution of 2-oxa-6-azaspiro[3,3]heptane (0.99 g, 0.01 mol) with 4-fluorobenzotrile (1.21 g, 0.01 mol) was heated to reflux for 5 h, then water (100 ml) was added into the solution. The mixture was extracted with CH₂Cl₂. Then the solvent was removed to give a red powder. Single crystals were obtained from a CH₂Cl₂ solution after 3 days.

S2. Refinement

H atoms were placed in the calculated positions (C—H = 0.95–0.99 Å) and refined using a riding model with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

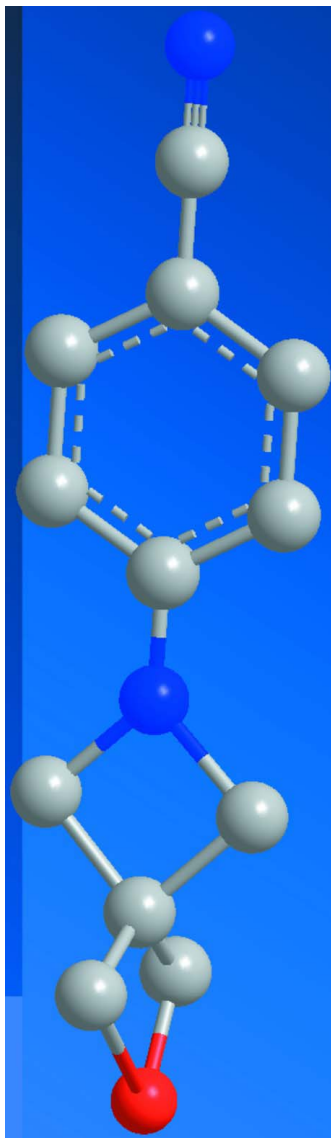


Figure 1

?

4-(2-Oxa-6-azaspiro[3.3]hept-6-yl)benzotrile*Crystal data* $C_{12}H_{12}N_2O$ $M_r = 200.24$ Monoclinic, $P2_1/n$ $a = 9.484 (4) \text{ \AA}$ $b = 11.033 (4) \text{ \AA}$ $c = 10.419 (4) \text{ \AA}$ $\beta = 113.186 (5)^\circ$ $V = 1002.2 (7) \text{ \AA}^3$ $Z = 4$ $F(000) = 424$ $D_x = 1.327 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 2956 reflections

 $\theta = 3.1\text{--}27.5^\circ$ $\mu = 0.09 \text{ mm}^{-1}$ $T = 113 \text{ K}$

Prism, colourless

 $0.60 \times 0.60 \times 0.27 \text{ mm}$

Data collection

Rigaku AFC10/Saturn 724-Plus
diffractometer
Radiation source: rotating anode
Graphite monochromator
Detector resolution: 28.5714 pixels mm⁻¹
 φ and ω scans
7662 measured reflections

2234 independent reflections
1872 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.029$
 $\theta_{\text{max}} = 27.5^\circ$, $\theta_{\text{min}} = 3.1^\circ$
 $h = -12 \rightarrow 12$
 $k = -14 \rightarrow 11$
 $l = -12 \rightarrow 12$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.041$
 $wR(F^2) = 0.099$
 $S = 1.00$
2234 reflections
136 parameters
0 restraints
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
 $w = 1/[\sigma^2(F_o^2) + (0.0456P)^2 + 0.36P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.30 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.16 \text{ e } \text{\AA}^{-3}$

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds 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 $F^2 > \sigma(F^2)$ is used only for calculating 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)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.53098 (12)	0.21116 (9)	-0.00129 (10)	0.0272 (3)
N1	0.81838 (14)	0.46062 (12)	1.07793 (13)	0.0275 (3)
N2	0.58259 (13)	0.37004 (10)	0.38573 (11)	0.0186 (3)
C1	0.74599 (15)	0.47583 (12)	0.59326 (14)	0.0178 (3)
H1	0.7933	0.5192	0.5421	0.021*
C2	0.79050 (15)	0.49646 (12)	0.73352 (14)	0.0186 (3)
H2	0.8679	0.5547	0.7788	0.022*
C3	0.72270 (15)	0.43219 (12)	0.81086 (14)	0.0177 (3)
C4	0.60810 (15)	0.34638 (12)	0.74350 (14)	0.0183 (3)
H4	0.5624	0.3022	0.7954	0.022*
C5	0.56183 (15)	0.32601 (12)	0.60278 (14)	0.0178 (3)
H5	0.4835	0.2683	0.5577	0.021*
C6	0.62984 (14)	0.39026 (12)	0.52463 (13)	0.0165 (3)
C7	0.49702 (15)	0.26581 (12)	0.30534 (14)	0.0178 (3)
H7A	0.3842	0.2751	0.2702	0.021*
H7B	0.5303	0.1871	0.3534	0.021*
C8	0.56407 (14)	0.29329 (12)	0.19522 (14)	0.0167 (3)

C9	0.65782 (15)	0.39702 (12)	0.28976 (13)	0.0177 (3)
H9A	0.7699	0.3827	0.3302	0.021*
H9B	0.6331	0.4782	0.2461	0.021*
C10	0.45979 (16)	0.31151 (12)	0.04116 (14)	0.0204 (3)
H10A	0.3499	0.2979	0.0208	0.024*
H10B	0.4749	0.3907	0.0037	0.024*
C11	0.63751 (16)	0.19237 (13)	0.14221 (14)	0.0226 (3)
H11A	0.7455	0.2092	0.1575	0.027*
H11B	0.6281	0.1114	0.1790	0.027*
C12	0.77569 (15)	0.44870 (12)	0.95891 (15)	0.0202 (3)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0353 (6)	0.0266 (6)	0.0202 (5)	0.0027 (5)	0.0113 (5)	-0.0041 (4)
N1	0.0257 (7)	0.0323 (7)	0.0231 (7)	-0.0015 (5)	0.0082 (5)	-0.0038 (6)
N2	0.0203 (6)	0.0187 (6)	0.0180 (6)	-0.0073 (4)	0.0086 (5)	-0.0027 (5)
C1	0.0179 (6)	0.0147 (6)	0.0217 (7)	-0.0007 (5)	0.0086 (5)	0.0028 (5)
C2	0.0156 (6)	0.0151 (6)	0.0223 (7)	0.0007 (5)	0.0043 (5)	-0.0017 (5)
C3	0.0165 (6)	0.0176 (6)	0.0177 (7)	0.0041 (5)	0.0054 (5)	-0.0004 (5)
C4	0.0177 (6)	0.0181 (6)	0.0212 (7)	0.0025 (5)	0.0098 (5)	0.0022 (5)
C5	0.0157 (6)	0.0163 (6)	0.0212 (7)	-0.0017 (5)	0.0069 (5)	-0.0012 (5)
C6	0.0151 (6)	0.0147 (6)	0.0188 (7)	0.0023 (5)	0.0058 (5)	0.0008 (5)
C7	0.0184 (6)	0.0157 (6)	0.0195 (7)	-0.0029 (5)	0.0076 (5)	-0.0014 (5)
C8	0.0161 (6)	0.0149 (6)	0.0193 (7)	0.0004 (5)	0.0071 (5)	0.0002 (5)
C9	0.0171 (6)	0.0186 (6)	0.0172 (6)	-0.0020 (5)	0.0065 (5)	0.0004 (5)
C10	0.0210 (7)	0.0184 (6)	0.0195 (7)	-0.0010 (5)	0.0057 (5)	-0.0002 (6)
C11	0.0257 (7)	0.0215 (7)	0.0227 (7)	0.0046 (6)	0.0117 (6)	0.0016 (6)
C12	0.0175 (6)	0.0195 (6)	0.0235 (8)	0.0024 (5)	0.0079 (5)	-0.0010 (6)

Geometric parameters (Å, °)

O1—C11	1.4525 (17)	C5—C6	1.4125 (19)
O1—C10	1.4534 (17)	C5—H5	0.95
N1—C12	1.1504 (18)	C7—C8	1.5451 (18)
N2—C6	1.3544 (17)	C7—H7A	0.99
N2—C7	1.4653 (17)	C7—H7B	0.99
N2—C9	1.4695 (17)	C8—C11	1.5274 (18)
C1—C2	1.3711 (19)	C8—C10	1.5314 (18)
C1—C6	1.4127 (18)	C8—C9	1.5431 (18)
C1—H1	0.95	C9—H9A	0.99
C2—C3	1.4049 (19)	C9—H9B	0.99
C2—H2	0.95	C10—H10A	0.99
C3—C4	1.4034 (19)	C10—H10B	0.99
C3—C12	1.4335 (19)	C11—H11A	0.99
C4—C5	1.3734 (19)	C11—H11B	0.99
C4—H4	0.95		

C11—O1—C10	90.79 (9)	H7A—C7—H7B	111.1
C6—N2—C7	128.02 (11)	C11—C8—C10	85.12 (10)
C6—N2—C9	130.44 (11)	C11—C8—C9	122.78 (11)
C7—N2—C9	94.47 (10)	C10—C8—C9	122.94 (11)
C2—C1—C6	120.13 (12)	C11—C8—C7	120.34 (11)
C2—C1—H1	119.9	C10—C8—C7	121.30 (11)
C6—C1—H1	119.9	C9—C8—C7	88.49 (10)
C1—C2—C3	120.61 (12)	N2—C9—C8	88.40 (10)
C1—C2—H2	119.7	N2—C9—H9A	113.9
C3—C2—H2	119.7	C8—C9—H9A	113.9
C4—C3—C2	119.53 (12)	N2—C9—H9B	113.9
C4—C3—C12	119.89 (12)	C8—C9—H9B	113.9
C2—C3—C12	120.49 (12)	H9A—C9—H9B	111.1
C5—C4—C3	120.22 (12)	O1—C10—C8	91.91 (10)
C5—C4—H4	119.9	O1—C10—H10A	113.3
C3—C4—H4	119.9	C8—C10—H10A	113.3
C4—C5—C6	120.45 (12)	O1—C10—H10B	113.3
C4—C5—H5	119.8	C8—C10—H10B	113.3
C6—C5—H5	119.8	H10A—C10—H10B	110.6
N2—C6—C5	119.95 (12)	O1—C11—C8	92.11 (10)
N2—C6—C1	121.00 (12)	O1—C11—H11A	113.3
C5—C6—C1	119.05 (12)	C8—C11—H11A	113.3
N2—C7—C8	88.47 (10)	O1—C11—H11B	113.3
N2—C7—H7A	113.9	C8—C11—H11B	113.3
C8—C7—H7A	113.9	H11A—C11—H11B	110.6
N2—C7—H7B	113.9	N1—C12—C3	179.26 (15)
C8—C7—H7B	113.9		
C6—C1—C2—C3	0.67 (19)	N2—C7—C8—C11	130.78 (12)
C1—C2—C3—C4	-0.17 (18)	N2—C7—C8—C10	-125.31 (12)
C1—C2—C3—C12	176.30 (12)	N2—C7—C8—C9	3.04 (9)
C2—C3—C4—C5	-0.45 (19)	C6—N2—C9—C8	154.66 (13)
C12—C3—C4—C5	-176.95 (12)	C7—N2—C9—C8	3.21 (10)
C3—C4—C5—C6	0.57 (19)	C11—C8—C9—N2	-128.76 (12)
C7—N2—C6—C5	-18.4 (2)	C10—C8—C9—N2	123.99 (12)
C9—N2—C6—C5	-161.23 (13)	C7—C8—C9—N2	-3.04 (9)
C7—N2—C6—C1	162.08 (12)	C11—O1—C10—C8	-2.22 (10)
C9—N2—C6—C1	19.3 (2)	C11—C8—C10—O1	2.12 (10)
C4—C5—C6—N2	-179.56 (12)	C9—C8—C10—O1	128.42 (12)
C4—C5—C6—C1	-0.07 (19)	C7—C8—C10—O1	-120.66 (12)
C2—C1—C6—N2	178.93 (12)	C10—O1—C11—C8	2.22 (10)
C2—C1—C6—C5	-0.55 (19)	C10—C8—C11—O1	-2.12 (10)
C6—N2—C7—C8	-155.71 (13)	C9—C8—C11—O1	-128.56 (12)
C9—N2—C7—C8	-3.21 (10)	C7—C8—C11—O1	121.54 (12)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H··· <i>A</i>	<i>D</i> —H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> —H··· <i>A</i>
C4—H4···O1 ⁱ	0.95	2.47	3.371 (2)	159

Symmetry code: (i) *x*, *y*, *z*+1.