

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

(11-Methylpyrido[2,3-*b*][1,4]benzodiazepin-6-yl)(phenyl)methanone

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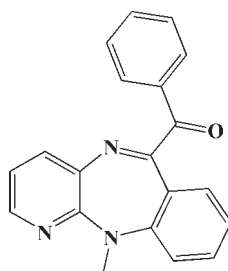
Received 21 July 2010; accepted 30 July 2010

Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.048; wR factor = 0.131; data-to-parameter ratio = 16.5.

In the title compound, $\text{C}_{20}\text{H}_{15}\text{N}_3\text{O}$, the diazepine ring adopts a boat conformation. The dihedral angle between pyridine and benzene rings is $55.2(1)^\circ$. The benzoyl phenyl ring forms dihedral angles of $49.4(1)$ and $75.9(1)^\circ$, respectively, with the pyridine and benzene rings. In the crystal, molecules are linked into centrosymmetric dimers by pairs of $\text{C}-\text{H}\cdots\text{N}$ hydrogen bonds.

Related literature

For general background to pyridobenzodiazepine derivatives, see: Eberlein *et al.* (1987); Horton *et al.* (2003); Shi *et al.* (2008, 2010); Tahtaoui *et al.* (2004). For a related structure, see: Spirlet *et al.* (2003).



Experimental

Crystal data

$\text{C}_{20}\text{H}_{15}\text{N}_3\text{O}$
 $M_r = 313.35$
Monoclinic, $P2_1/c$
 $a = 8.4442(17)$ Å
 $b = 16.503(3)$ Å
 $c = 11.682(2)$ Å
 $\beta = 98.14(3)^\circ$

$V = 1611.6(6)$ Å³
 $Z = 4$
Mo $K\alpha$ radiation
 $\mu = 0.08$ mm⁻¹
 $T = 293$ K
 $0.37 \times 0.30 \times 0.19$ mm

Data collection

Rigaku R-Axis RAPID
diffractometer
Absorption correction: multi-scan
(*ABSCOR*; Higashi, 1995)
 $T_{\min} = 0.389$, $T_{\max} = 0.431$

14767 measured reflections
3587 independent reflections
2492 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.048$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.048$
 $wR(F^2) = 0.131$
 $S = 1.05$
3587 reflections

217 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.22$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.16$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C15}-\text{H15}\cdots\text{N1}^i$	0.93	2.56	3.463 (2)	163

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

Data collection: *PROCESS-AUTO* (Rigaku, 1998); cell refinement: *PROCESS-AUTO*; data reduction: *CrystalStructure* (Rigaku/MS, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *DIAMOND* (Brandenburg, 2000); software used to prepare material for publication: *SHELXL97*.

This project was sponsored by the Scientific Research Foundation for Returned Overseas Chinese Scholars, State Education Ministry (grant No. 20071108) and the Scientific Research Foundation for the Returned Overseas Team, Chinese Education Ministry.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CI5138).

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supporting information

Acta Cryst. (2010). E66, o2292 [https://doi.org/10.1107/S1600536810030564]

(11-Methylpyrido[2,3-*b*][1,4]benzodiazepin-6-yl)(phenyl)methanone**Fuqiang Shi, Long Zhang, Jin-Feng Wang and Ya-Feng Li****S1. Comment**

Pyridobenzodiazepine derivatives possess biological and pharmacological activities (Horton *et al.*, 2003). In most of the reported pyridobenzodiazepines amino or aryl or alkyl group is attached at the C6-position of the heterocyclic nucleus (Eberlein *et al.*, 1987; Tahtaoui *et al.*, 2004; Shi *et al.*, 2008, 2010) while the attachment of a ketone group has not been reported. We report here the crystal structure of the title compound which contains a benzoyl group at the C6-position.

Bond lengths and angles in the title molecule (Fig.1) are comparable with those observed in a related structure (Spirlet *et al.*, 2003). The diazepine ring displays a boat conformation. The dihedral angle between pyridine and C15-C20 benzene rings is 55.2 (1)°. The benzoyl phenyl ring forms dihedral angles of 49.4 (1)° and 75.9 (1)°, respectively, with the pyridine and benzene ring of the benzodiazepine ring system.

In the crystal structure, the molecules are linked into dimers by C15—H15⋯N1 hydrogen bonds (Table 1).

S2. Experimental

Polyphosphoric acid (254 mg, 0.75 mmol), N-2-methyl-N-2-phenylpyridine-2,3-diamine (100 mg, 0.5 mmol) and 2-phenylacetic acid (102 mg, 0.75 mmol) were dissolved in POCl₃ (5 ml). The solution was heated at 368 K in an oil bath for 7 h and the solution was poured into ice-water (20 ml), treated with 5 N NaOH to pH 9-10, and then extracted with EtOAc (3 × 20 ml). The combined organic phase was washed with saturated NaHCO₃ and brine, dried with anhydrous Na₂SO₄, concentrated *in vacuo*, and purified by flash chromatography with petroleum ether/EtOAc (10:1, v/v) as eluent to afford a mixture of 6-benzyl-11-methylpyrido[2,3-*b*][1,4]benzodiazepine and 6-benzoyl-11-methylpyrido[2,3-*b*][1,4]benzodiazepine. The mixture was dissolved in dichloromethane (5 ml), stirred for 24 h under oxygen at room temperature and 6-benzyl-11-methylpyrido[2,3-*b*][1,4]benzodiazepine disappeared. The reagent was concentrated *in vacuo*, purified by flash chromatography (yield: 140 mg, 95%) and then crystallized from dichloromethane to obtain colourless crystals of the title compound suitable for X-ray analysis.

S3. Refinement

H atoms were positioned geometrically [C—H = 0.93–0.96 Å] and treated as riding with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C}_{\text{methyl}})$ and $1.2U_{\text{eq}}(\text{C})$.

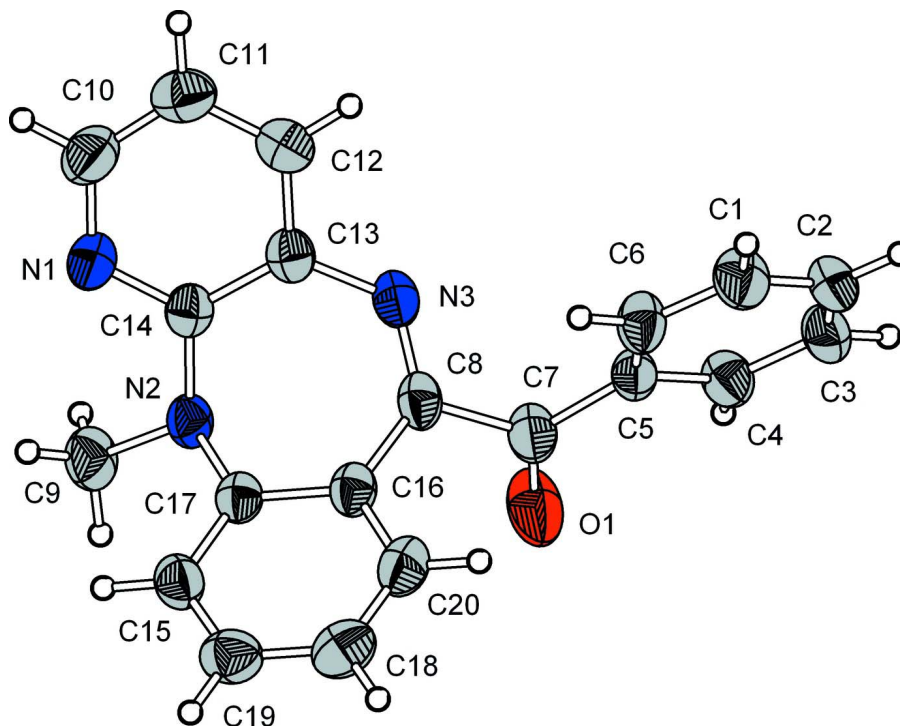


Figure 1

The molecular structure of the title compound, with the atom-labelling scheme. Displacement ellipsoids are shown at the 50% probability level.

(11-Methylpyrido[2,3-*b*][1,4]benzodiazepin-6-yl)(phenyl)methanone

Crystal data

$C_{20}H_{15}N_3O$

$M_r = 313.35$

Monoclinic, $P2_1/c$

Hall symbol: $-P\ 2_1/c$

$a = 8.4442(17)\ \text{\AA}$

$b = 16.503(3)\ \text{\AA}$

$c = 11.682(2)\ \text{\AA}$

$\beta = 98.14(3)^\circ$

$V = 1611.6(6)\ \text{\AA}^3$

$Z = 4$

$F(000) = 656$

$D_x = 1.291\ \text{Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073\ \text{\AA}$

Cell parameters from 2000 reflections

$\theta = 3.0\text{--}27.5^\circ$

$\mu = 0.08\ \text{mm}^{-1}$

$T = 293\ \text{K}$

Block, yellow

$0.37 \times 0.30 \times 0.19\ \text{mm}$

Data collection

Rigaku R-AXIS RAPID

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution: $10.00\ \text{pixels mm}^{-1}$

ω scans

Absorption correction: multi-scan

(*ABSCOR*; Higashi, 1995)

$T_{\min} = 0.389$, $T_{\max} = 0.431$

14767 measured reflections

3587 independent reflections

2492 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.048$

$\theta_{\max} = 27.5^\circ$, $\theta_{\min} = 3.0^\circ$

$h = -10 \rightarrow 9$

$k = -21 \rightarrow 21$

$l = -14 \rightarrow 15$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.048$

$wR(F^2) = 0.131$

$S = 1.05$

3587 reflections

217 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

$w = 1/[\sigma^2(F_o^2) + (0.0594P)^2 + 0.2193P]$

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

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

$\Delta\rho_{\max} = 0.22 \text{ e } \text{\AA}^{-3}$

$\Delta\rho_{\min} = -0.16 \text{ e } \text{\AA}^{-3}$

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 $F^2 > \sigma(F^2)$ 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	0.84497 (15)	0.36871 (8)	-0.05524 (17)	0.0774 (5)
N2	0.66756 (14)	0.13174 (8)	0.03808 (12)	0.0394 (3)
N3	0.99137 (15)	0.19569 (8)	0.04732 (13)	0.0432 (3)
C1	1.3716 (2)	0.31549 (11)	-0.12859 (17)	0.0529 (5)
H1	1.4411	0.2799	-0.1580	0.063*
C2	1.4173 (2)	0.39435 (11)	-0.10549 (16)	0.0497 (4)
H2	1.5176	0.4119	-0.1192	0.060*
C3	1.3153 (2)	0.44719 (11)	-0.06219 (16)	0.0500 (5)
H3	1.3468	0.5004	-0.0457	0.060*
C4	1.16610 (19)	0.42134 (10)	-0.04322 (16)	0.0464 (4)
H4	1.0970	0.4575	-0.0146	0.056*
C5	1.11770 (18)	0.34200 (9)	-0.06626 (14)	0.0387 (4)
C6	1.2226 (2)	0.28871 (10)	-0.10835 (16)	0.0472 (4)
H6	1.1930	0.2350	-0.1230	0.057*
C7	0.9509 (2)	0.31859 (10)	-0.05454 (17)	0.0468 (4)
C8	0.90799 (18)	0.23004 (9)	-0.03891 (15)	0.0402 (4)
C9	0.5167 (2)	0.11410 (12)	0.08085 (17)	0.0526 (5)
H9A	0.4339	0.1483	0.0421	0.079*
H9B	0.5284	0.1241	0.1626	0.079*
H9C	0.4887	0.0583	0.0660	0.079*
C10	0.9020 (2)	-0.02679 (11)	0.18587 (17)	0.0571 (5)
H10	0.8833	-0.0755	0.2217	0.069*
C11	1.0573 (2)	-0.00296 (11)	0.18578 (16)	0.0536 (5)
H11	1.1418	-0.0357	0.2177	0.064*
C12	1.0848 (2)	0.07088 (11)	0.13706 (15)	0.0478 (4)

H12	1.1890	0.0897	0.1390	0.057*
C13	0.95740 (19)	0.11717 (9)	0.08517 (14)	0.0391 (4)
C14	0.80223 (18)	0.08588 (9)	0.08663 (13)	0.0376 (4)
N1	0.77531 (18)	0.01626 (8)	0.13702 (13)	0.0494 (4)
C15	0.53710 (19)	0.11522 (10)	-0.16328 (16)	0.0447 (4)
H15	0.4594	0.0820	-0.1390	0.054*
C16	0.77505 (18)	0.19642 (9)	-0.12099 (14)	0.0383 (4)
C17	0.65890 (17)	0.14649 (9)	-0.08261 (14)	0.0358 (3)
C18	0.6450 (2)	0.18248 (12)	-0.31775 (16)	0.0563 (5)
H18	0.6406	0.1941	-0.3960	0.068*
C19	0.5310 (2)	0.13338 (11)	-0.27957 (16)	0.0530 (5)
H19	0.4489	0.1122	-0.3326	0.064*
C20	0.7655 (2)	0.21400 (11)	-0.23854 (16)	0.0497 (4)
H20	0.8419	0.2476	-0.2639	0.060*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0463 (7)	0.0440 (8)	0.1469 (16)	0.0030 (6)	0.0313 (8)	0.0024 (8)
N2	0.0350 (7)	0.0437 (8)	0.0418 (7)	-0.0032 (5)	0.0136 (6)	0.0033 (6)
N3	0.0388 (7)	0.0397 (8)	0.0519 (9)	-0.0071 (6)	0.0095 (6)	-0.0011 (6)
C1	0.0422 (9)	0.0514 (11)	0.0686 (12)	0.0050 (8)	0.0196 (9)	0.0014 (9)
C2	0.0360 (9)	0.0570 (11)	0.0557 (11)	-0.0089 (7)	0.0053 (8)	0.0049 (8)
C3	0.0440 (9)	0.0414 (9)	0.0628 (12)	-0.0123 (7)	0.0009 (8)	-0.0029 (8)
C4	0.0409 (9)	0.0368 (9)	0.0616 (11)	0.0001 (7)	0.0076 (8)	-0.0049 (8)
C5	0.0356 (8)	0.0331 (8)	0.0482 (9)	-0.0025 (6)	0.0086 (7)	0.0025 (7)
C6	0.0478 (9)	0.0337 (9)	0.0626 (11)	-0.0027 (7)	0.0171 (8)	-0.0018 (8)
C7	0.0408 (9)	0.0385 (9)	0.0631 (11)	-0.0020 (7)	0.0143 (8)	0.0009 (8)
C8	0.0360 (8)	0.0366 (8)	0.0513 (10)	-0.0042 (6)	0.0175 (7)	-0.0004 (7)
C9	0.0435 (9)	0.0603 (12)	0.0586 (11)	-0.0028 (8)	0.0231 (9)	0.0047 (9)
C10	0.0689 (13)	0.0437 (10)	0.0566 (11)	-0.0028 (9)	0.0016 (10)	0.0092 (8)
C11	0.0584 (11)	0.0482 (10)	0.0516 (11)	0.0088 (8)	-0.0018 (9)	-0.0011 (8)
C12	0.0415 (9)	0.0511 (10)	0.0498 (10)	0.0009 (7)	0.0027 (8)	-0.0041 (8)
C13	0.0405 (8)	0.0392 (9)	0.0384 (8)	-0.0045 (6)	0.0088 (7)	-0.0039 (7)
C14	0.0424 (9)	0.0367 (8)	0.0346 (8)	-0.0035 (6)	0.0088 (7)	-0.0019 (6)
N1	0.0536 (9)	0.0425 (8)	0.0520 (9)	-0.0076 (6)	0.0064 (7)	0.0090 (7)
C15	0.0399 (9)	0.0382 (9)	0.0556 (11)	-0.0039 (7)	0.0056 (8)	-0.0008 (7)
C16	0.0383 (8)	0.0357 (8)	0.0432 (9)	-0.0006 (6)	0.0132 (7)	0.0016 (7)
C17	0.0361 (8)	0.0312 (8)	0.0414 (9)	0.0015 (6)	0.0099 (7)	-0.0003 (6)
C18	0.0690 (12)	0.0573 (12)	0.0423 (10)	0.0052 (9)	0.0075 (9)	0.0054 (8)
C19	0.0570 (11)	0.0507 (11)	0.0477 (10)	0.0012 (8)	-0.0047 (9)	-0.0041 (8)
C20	0.0555 (10)	0.0461 (10)	0.0498 (11)	-0.0017 (8)	0.0156 (9)	0.0083 (8)

Geometric parameters (Å, °)

O1—C7	1.218 (2)	C9—H9B	0.96
N2—C14	1.416 (2)	C9—H9C	0.96
N2—C17	1.422 (2)	C10—N1	1.342 (2)

N2—C9	1.462 (2)	C10—C11	1.369 (3)
N3—C8	1.277 (2)	C10—H10	0.93
N3—C13	1.412 (2)	C11—C12	1.379 (3)
C1—C2	1.373 (2)	C11—H11	0.93
C1—C6	1.385 (2)	C12—C13	1.387 (2)
C1—H1	0.93	C12—H12	0.93
C2—C3	1.372 (3)	C13—C14	1.411 (2)
C2—H2	0.93	C14—N1	1.325 (2)
C3—C4	1.377 (2)	C15—C19	1.385 (3)
C3—H3	0.93	C15—C17	1.392 (2)
C4—C5	1.387 (2)	C15—H15	0.93
C4—H4	0.93	C16—C20	1.395 (2)
C5—C6	1.387 (2)	C16—C17	1.402 (2)
C5—C7	1.485 (2)	C18—C20	1.377 (3)
C6—H6	0.93	C18—C19	1.380 (3)
C7—C8	1.523 (2)	C18—H18	0.93
C8—C16	1.477 (2)	C19—H19	0.93
C9—H9A	0.96	C20—H20	0.93
C14—N2—C17	114.46 (13)	N1—C10—C11	123.60 (17)
C14—N2—C9	116.49 (13)	N1—C10—H10	118.2
C17—N2—C9	116.60 (13)	C11—C10—H10	118.2
C8—N3—C13	122.74 (13)	C10—C11—C12	118.14 (16)
C2—C1—C6	120.38 (17)	C10—C11—H11	120.9
C2—C1—H1	119.8	C12—C11—H11	120.9
C6—C1—H1	119.8	C11—C12—C13	120.11 (16)
C3—C2—C1	120.07 (16)	C11—C12—H12	119.9
C3—C2—H2	120.0	C13—C12—H12	119.9
C1—C2—H2	120.0	C12—C13—C14	117.22 (15)
C2—C3—C4	119.92 (16)	C12—C13—N3	117.58 (14)
C2—C3—H3	120.0	C14—C13—N3	124.75 (14)
C4—C3—H3	120.0	N1—C14—C13	122.73 (15)
C3—C4—C5	120.82 (16)	N1—C14—N2	117.57 (14)
C3—C4—H4	119.6	C13—C14—N2	119.61 (14)
C5—C4—H4	119.6	C14—N1—C10	118.11 (15)
C4—C5—C6	118.82 (15)	C19—C15—C17	120.35 (16)
C4—C5—C7	119.02 (15)	C19—C15—H15	119.8
C6—C5—C7	121.96 (15)	C17—C15—H15	119.8
C1—C6—C5	119.96 (16)	C20—C16—C17	119.45 (15)
C1—C6—H6	120.0	C20—C16—C8	119.57 (15)
C5—C6—H6	120.0	C17—C16—C8	120.98 (14)
O1—C7—C5	121.87 (15)	C15—C17—C16	118.99 (15)
O1—C7—C8	117.75 (15)	C15—C17—N2	122.48 (14)
C5—C7—C8	120.38 (14)	C16—C17—N2	118.51 (14)
N3—C8—C16	128.95 (14)	C20—C18—C19	119.19 (17)
N3—C8—C7	113.99 (14)	C20—C18—H18	120.4
C16—C8—C7	117.03 (14)	C19—C18—H18	120.4
N2—C9—H9A	109.5	C18—C19—C15	120.88 (17)

N2—C9—H9B	109.5	C18—C19—H19	119.6
H9A—C9—H9B	109.5	C15—C19—H19	119.6
N2—C9—H9C	109.5	C18—C20—C16	121.15 (17)
H9A—C9—H9C	109.5	C18—C20—H20	119.4
H9B—C9—H9C	109.5	C16—C20—H20	119.4

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
C15—H15 \cdots N1 ⁱ	0.93	2.56	3.463 (2)	163

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