

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

1,1'-(Butane-1,4-diyl)bis[2-(pyridin-2-yl)-1H-benzimidazole]

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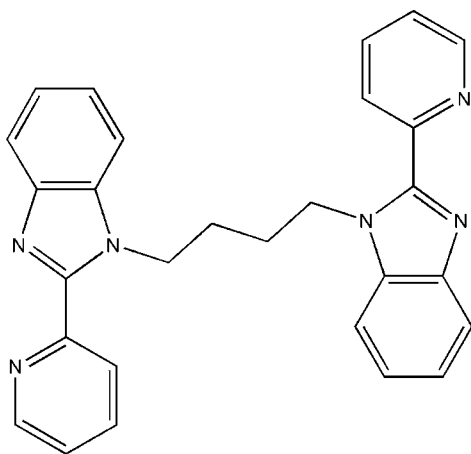
Received 9 April 2012; accepted 12 April 2012

Key indicators: single-crystal X-ray study; $T = 298$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.064; wR factor = 0.146; data-to-parameter ratio = 16.6.

The complete molecule of the title compound, $\text{C}_{28}\text{H}_{24}\text{N}_6$, is generated by inversion symmetry with the inversion centre located at the mid-point of the central C–C bond of the butanediyli unit. The benzimidazole and pyridine rings are almost coplanar, the dihedral angle between their mean planes being 6.86 (11)°.

Related literature

For the synthesis, see: Liu *et al.* (2010). For background to this study, see: Barnett & Champness (2003); Tong *et al.* (2009).



Experimental

Crystal data

$\text{C}_{28}\text{H}_{24}\text{N}_6$
 $M_r = 444.53$
 Monoclinic, $P2_1/n$
 $a = 6.5617$ (7) Å
 $b = 13.9716$ (13) Å
 $c = 12.3351$ (8) Å
 $\beta = 96.466$ (7)°

$V = 1123.66$ (17) Å³
 $Z = 2$
 Mo $K\alpha$ radiation
 $\mu = 0.08$ mm⁻¹
 $T = 298$ K
 $0.42 \times 0.18 \times 0.15$ mm

Data collection

Rigaku R-Axis RAPID
 diffractometer
 Absorption correction: multi-scan
 (ABSCOR; Higashi, 1995)
 $T_{\min} = 0.695$, $T_{\max} = 0.856$

6268 measured reflections
 2684 independent reflections
 1314 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.055$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.064$
 $wR(F^2) = 0.146$
 $S = 1.02$
 2684 reflections
 162 parameters

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\max} = 0.34$ e Å⁻³
 $\Delta\rho_{\min} = -0.20$ e Å⁻³

Data collection: *RAPID-AUTO* (Rigaku, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku/MSK, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

The paper was sponsored by the K. C. Wong Magna Fund in Ningbo University, the Project of Nonprofit Technology & Research in Zhejiang (2011 C37010), the Ningbo Municipal Natural Science Foundation (grant No. 2010 A610160), the Subject Object (No. xk1070) and the Foundation (No. XYL11009) of Ningbo University.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: FF2063).

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supplementary materials

Acta Cryst. (2012). E68, o1434 [doi:10.1107/S1600536812015899]

1,1'-(Butane-1,4-diyl)bis[2-(pyridin-2-yl)-1H-benzimidazole]**Shao-Chuan Zhou and Hong-Zhen Xie****Comment**

The long spacer ligands, particularly the flexible N-bridging donors have been investigated as auxiliary ligands for the construction of novel MOFs [Liu *et al.*, 2010; Barnett *et al.*, 2003; Tong *et al.*, 2009]. As part of our ongoing studies, the title compound was synthesized and characterized by X-ray diffraction.

The complete molecule of the title compound, C₂₈H₂₄N₆, is generated by crystallographic inversion symmetry and the central C—C bond of the butanediy unit is bisected by the inversion symmetry. The dihedral angle between the benzimidazole ring system and the pyridine ring is 6.86 (11)°, which indicates that they are almost coplanar.

Experimental

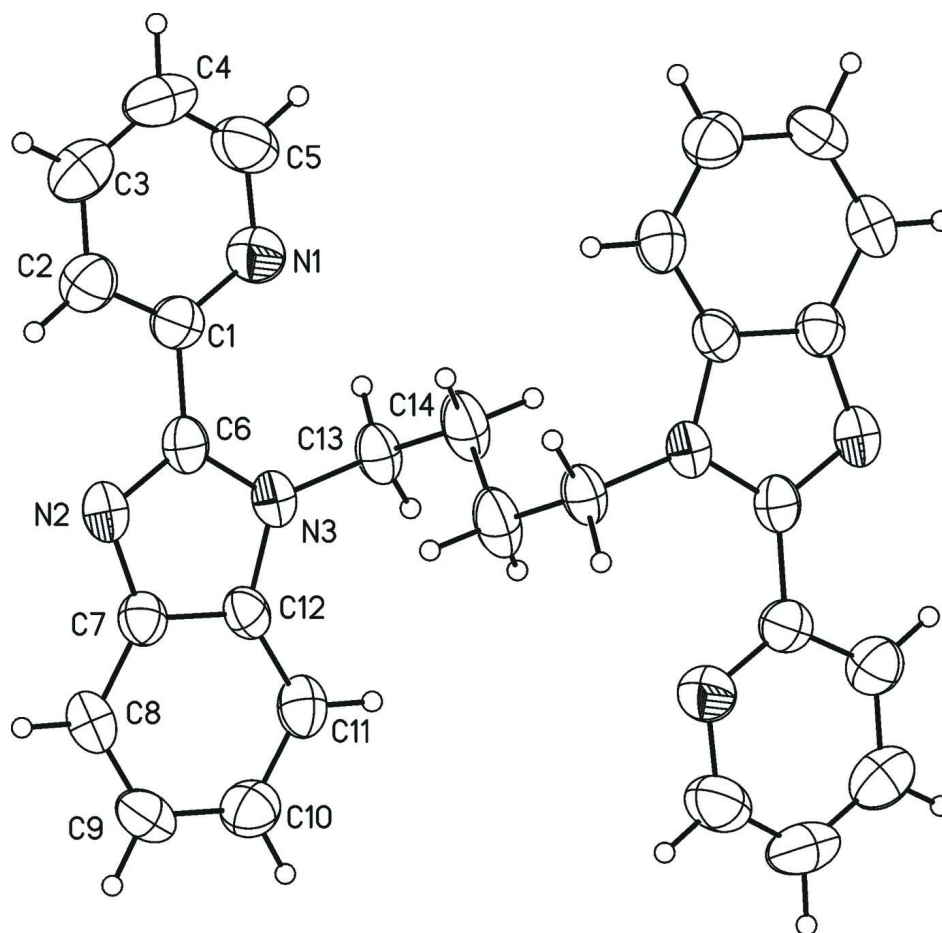
According to the literature [Liu *et al.*, 2010], 2-(2-pyridyl)benzimidazole (7.80 g) and NaOH (1.68 g) in DMSO (20 ml) were stirred at 60°C for 0.5 h, and then 1,4-dibromobutane (4.32 g) was added. The mixture was stirred at 60°C for 12 h, and then poured into 400 ml of ice water after being cooled to room temperature. The yellow solid was obtained and isolated by filtration after drying in air. The above products were recrystallized in methanol and yellow crystals of the title compounds were obtained.

Refinement

The H atoms bonded to C except for C14 were placed at calculated positions and refined in riding mode with $U_{\text{iso}}(\text{H})=1.2U_{\text{eq}}(\text{C})$. The H atoms of C14 were located at difference Fourier maps and refined freely.

Computing details

Data collection: *RAPID-AUTO* (Rigaku, 1998); cell refinement: *RAPID-AUTO* (Rigaku, 1998); data reduction: *CrystalStructure* (Rigaku/MSK, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97* (Sheldrick, 2008).

**Figure 1**

ORTEP view of complex molecule of (I). Displacement ellipsoids are drawn at the 45% probability level. H atoms were omitted for clarity.

1,1'-(butane-1,4-diyl)bis[2-(pyridin-2-yl)-1H-benzimidazole]

Crystal data

$C_{28}H_{24}N_6$

$M_r = 444.53$

Monoclinic, $P2_1/n$

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

$a = 6.5617(7)\ \text{\AA}$

$b = 13.9716(13)\ \text{\AA}$

$c = 12.3351(8)\ \text{\AA}$

$\beta = 96.466(7)^\circ$

$V = 1123.66(17)\ \text{\AA}^3$

$Z = 2$

$F(000) = 468$

$D_x = 1.314\ \text{Mg m}^{-3}$

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

Cell parameters from 6230 reflections

$\theta = 3.1\text{--}29.7^\circ$

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

$T = 298\ \text{K}$

Block, yellow

$0.42 \times 0.18 \times 0.15\ \text{mm}$

Data collection

Rigaku R-AXIS RAPID
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω scans

Absorption correction: multi-scan

(*ABSCOR*; Higashi, 1995)

$T_{\min} = 0.695$, $T_{\max} = 0.856$

6268 measured reflections

2684 independent reflections

1314 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.055$
 $\theta_{\text{max}} = 29.3^\circ$, $\theta_{\text{min}} = 2.9^\circ$

$h = -7 \rightarrow 8$
 $k = -14 \rightarrow 18$
 $l = -16 \rightarrow 15$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.064$
 $wR(F^2) = 0.146$
 $S = 1.02$
 2684 reflections
 162 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 atoms treated by a mixture of independent
 and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0451P)^2 + 0.0549P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.34 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.20 \text{ e } \text{\AA}^{-3}$

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>	$U_{\text{iso}}^*/U_{\text{eq}}$
N1	-0.2449 (3)	0.71889 (16)	0.85672 (16)	0.0593 (6)
N2	-0.0037 (3)	0.59178 (15)	0.65051 (13)	0.0501 (6)
N3	0.1079 (3)	0.59573 (14)	0.82913 (13)	0.0436 (5)
C1	-0.2217 (4)	0.68146 (17)	0.75907 (18)	0.0465 (6)
C2	-0.3657 (4)	0.6937 (2)	0.6693 (2)	0.0572 (7)
H2	-0.3442	0.6675	0.6022	0.069*
C3	-0.5399 (4)	0.7445 (2)	0.6803 (2)	0.0663 (8)
H3	-0.6393	0.7525	0.6210	0.080*
C4	-0.5664 (5)	0.7834 (2)	0.7785 (3)	0.0693 (8)
H4	-0.6835	0.8186	0.7877	0.083*
C5	-0.4172 (5)	0.7695 (2)	0.8636 (2)	0.0702 (9)
H5	-0.4360	0.7968	0.9305	0.084*
C6	-0.0367 (4)	0.62376 (17)	0.74647 (17)	0.0442 (6)
C7	0.1763 (4)	0.53894 (18)	0.66954 (17)	0.0437 (6)
C8	0.2838 (4)	0.48915 (19)	0.59693 (18)	0.0534 (7)
H8	0.2371	0.4871	0.5229	0.064*
C9	0.4599 (4)	0.4432 (2)	0.6369 (2)	0.0567 (7)
H9	0.5342	0.4099	0.5893	0.068*
C10	0.5305 (4)	0.4451 (2)	0.7475 (2)	0.0580 (7)
H10	0.6509	0.4131	0.7725	0.070*
C11	0.4249 (4)	0.49381 (19)	0.82047 (18)	0.0525 (7)
H11	0.4713	0.4953	0.8945	0.063*

C12	0.2482 (4)	0.54014 (17)	0.77956 (16)	0.0415 (6)
C13	0.1227 (4)	0.61541 (18)	0.94648 (15)	0.0469 (7)
H13B	0.0880	0.6820	0.9573	0.056*
H13A	0.2633	0.6057	0.9782	0.056*
C14	-0.0179 (6)	0.5522 (2)	1.0057 (2)	0.0620 (9)
H14B	0.005 (3)	0.5697 (16)	1.0838 (18)	0.054 (7)*
H14A	-0.173 (5)	0.544 (2)	0.978 (2)	0.099 (11)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N1	0.0668 (17)	0.0549 (15)	0.0575 (12)	0.0066 (13)	0.0128 (11)	-0.0001 (11)
N2	0.0581 (14)	0.0578 (14)	0.0344 (10)	-0.0035 (12)	0.0047 (9)	0.0026 (9)
N3	0.0506 (13)	0.0476 (12)	0.0328 (10)	-0.0050 (11)	0.0055 (9)	0.0015 (9)
C1	0.0516 (17)	0.0354 (14)	0.0543 (14)	-0.0035 (13)	0.0138 (12)	0.0039 (12)
C2	0.0607 (19)	0.0516 (17)	0.0591 (15)	0.0017 (15)	0.0058 (13)	0.0040 (14)
C3	0.061 (2)	0.0537 (18)	0.0818 (19)	0.0009 (16)	-0.0008 (16)	0.0112 (16)
C4	0.0522 (19)	0.0499 (18)	0.108 (2)	0.0101 (15)	0.0175 (17)	0.0112 (18)
C5	0.085 (2)	0.056 (2)	0.0748 (19)	0.0079 (18)	0.0293 (18)	-0.0059 (15)
C6	0.0505 (16)	0.0444 (15)	0.0381 (12)	-0.0069 (13)	0.0066 (11)	0.0047 (11)
C7	0.0423 (15)	0.0469 (15)	0.0423 (13)	-0.0043 (12)	0.0069 (11)	0.0043 (11)
C8	0.0600 (18)	0.0629 (19)	0.0387 (12)	-0.0069 (16)	0.0115 (12)	-0.0005 (12)
C9	0.0580 (19)	0.0595 (18)	0.0566 (15)	0.0009 (15)	0.0234 (13)	-0.0076 (14)
C10	0.0474 (17)	0.0620 (18)	0.0649 (16)	0.0016 (14)	0.0077 (13)	0.0059 (14)
C11	0.0534 (17)	0.0628 (18)	0.0399 (12)	-0.0063 (15)	-0.0005 (11)	0.0032 (13)
C12	0.0472 (15)	0.0411 (14)	0.0381 (12)	-0.0051 (13)	0.0129 (11)	0.0016 (11)
C13	0.0600 (16)	0.0503 (16)	0.0303 (11)	-0.0095 (13)	0.0046 (10)	-0.0044 (11)
C14	0.095 (3)	0.0579 (17)	0.0345 (13)	-0.0119 (19)	0.0139 (14)	-0.0042 (14)

Geometric parameters (\AA , $^\circ$)

N1—C1	1.337 (3)	C7—C12	1.386 (3)
N1—C5	1.344 (3)	C7—C8	1.388 (3)
N2—C6	1.306 (3)	C8—C9	1.364 (3)
N2—C7	1.391 (3)	C8—H8	0.9300
N3—C6	1.370 (3)	C9—C10	1.390 (3)
N3—C12	1.397 (3)	C9—H9	0.9300
N3—C13	1.466 (2)	C10—C11	1.376 (3)
C1—C2	1.383 (3)	C10—H10	0.9300
C1—C6	1.480 (3)	C11—C12	1.373 (3)
C2—C3	1.365 (4)	C11—H11	0.9300
C2—H2	0.9300	C13—C14	1.523 (3)
C3—C4	1.357 (4)	C13—H13B	0.9700
C3—H3	0.9300	C13—H13A	0.9700
C4—C5	1.366 (4)	C14—C14 ⁱ	1.486 (6)
C4—H4	0.9300	C14—H14B	0.99 (2)
C5—H5	0.9300	C14—H14A	1.04 (3)
C1—N1—C5	116.4 (2)	C9—C8—H8	120.9
C6—N2—C7	104.66 (19)	C7—C8—H8	120.9

C6—N3—C12	105.58 (17)	C8—C9—C10	121.4 (2)
C6—N3—C13	130.1 (2)	C8—C9—H9	119.3
C12—N3—C13	124.27 (19)	C10—C9—H9	119.3
N1—C1—C2	122.4 (2)	C11—C10—C9	121.0 (3)
N1—C1—C6	119.0 (2)	C11—C10—H10	119.5
C2—C1—C6	118.5 (2)	C9—C10—H10	119.5
C3—C2—C1	119.3 (3)	C12—C11—C10	117.3 (2)
C3—C2—H2	120.4	C12—C11—H11	121.3
C1—C2—H2	120.4	C10—C11—H11	121.3
C4—C3—C2	119.3 (3)	C11—C12—C7	122.3 (2)
C4—C3—H3	120.3	C11—C12—N3	132.2 (2)
C2—C3—H3	120.3	C7—C12—N3	105.5 (2)
C3—C4—C5	118.4 (3)	N3—C13—C14	112.9 (2)
C3—C4—H4	120.8	N3—C13—H13B	109.0
C5—C4—H4	120.8	C14—C13—H13B	109.0
N1—C5—C4	124.2 (3)	N3—C13—H13A	109.0
N1—C5—H5	117.9	C14—C13—H13A	109.0
C4—C5—H5	117.9	H13B—C13—H13A	107.8
N2—C6—N3	113.9 (2)	C14 ⁱ —C14—C13	114.4 (3)
N2—C6—C1	120.2 (2)	C14 ⁱ —C14—H14B	109.1 (14)
N3—C6—C1	125.8 (2)	C13—C14—H14B	106.8 (13)
C12—C7—C8	119.8 (2)	C14 ⁱ —C14—H14A	90.9 (18)
C12—C7—N2	110.35 (19)	C13—C14—H14A	122.1 (16)
C8—C7—N2	129.9 (2)	H14B—C14—H14A	113 (2)
C9—C8—C7	118.3 (2)		

Symmetry code: (i) $-x, -y+1, -z+2$.