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5,5′-Dimethyl-2,2′-bipyridine

Zeinab Khoshtarkib,[a] Amin Ebadi,[b] Roya Ahmadi[a] and Robabeh Alizadeh[c]*

[a]Islamic Azad University, Shahr-e-Rey Branch, Tehran, Iran, [b]Department of Chemistry, Islamic Azad University, Kazerun Branch, Kazerun, Fars, Iran, and [c]Damghan University of Basic Sciences, School of Chemistry, Damghan, Iran
Correspondence e-mail: alizadeh@dubs.ac.ir

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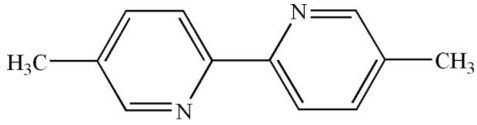
Key indicators: single-crystal X-ray study; T = 298 K; mean σ(C–C) = 0.003 Å; R factor = 0.066; wR factor = 0.205; data-to-parameter ratio = 21.2.

The asymmetric unit of the title compound, $C_{12}H_{12}N_2$, contains two half-molecules related by an inversion center, the planes of their pyridine rings being oriented at a dihedral angle of 69.62 (4)°. In the crystal structure, a π–π contact between the pyridine rings [centroid–centroid distance = 3.895 (3) Å] may stabilize the structure. A weak C—H$\cdots\pi$ interaction is also found.

Related literature

For related structures, see: Ahmadi *et al.* (2008); Albada *et al.* (2004); Amani *et al.* (2007); Kalateh *et al.* (2008); Khalighi *et al.* (2008); Maheshwari *et al.* (2007); Tadayon Pour *et al.* (2008). For bond-length data, see: Allen *et al.* (1987).



Experimental

Crystal data

$C_{12}H_{12}N_2$
M_r = 184.24
Triclinic, $P\overline{1}$
a = 6.409 (4) Å
b = 7.312 (5) Å
c = 11.533 (8) Å
α = 96.04 (5)°
β = 91.16 (4)°
γ = 105.03 (5)°
V = 518.4 (6) Å^3
Z = 2
Mo $K\alpha$ radiation
μ = 0.07 mm^{-1}
T = 298 K
0.50 × 0.41 × 0.29 mm

Data collection

Bruker SMART CCD area-detector diffractometer
Absorption correction: none
6191 measured reflections
2739 independent reflections
2067 reflections with $I > 2\sigma(I)$
R_{int} = 0.082

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.066
$wR(F^2)$ = 0.205
S = 1.08
2739 reflections
129 parameters
H-atom parameters constrained
$\Delta\rho_{max}$ = 0.27 e Å^{-3}
$\Delta\rho_{min}$ = −0.24 e Å^{-3}

Table 1
Hydrogen-bond geometry (Å, °).

D—H$\cdots A$	D—H	H$\cdots A$	$D\cdots A$	D—H$\cdots A$
C9—H9$\cdots Cg1^{i}$	0.93	2.78	3.669 (3)	160

Symmetry code: (i) $-x+1, -y+2, -z+2$. $Cg1$ is the centroid of the N1,C1–C4,C6 ring.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HK2693).

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supporting information

Acta Cryst. (2009). E**65**, o1586 [doi:10.1107/S1600536809019151]

5,5′-Dimethyl-2,2′-bipyridine

Zeinab Khoshtarkib, Amin Ebadi, Roya Ahmadi and Robabeh Alizadeh

S1. Comment

5,5′-Dimethyl-2,2′-bipyridine, (5,5′-dmbipy), is a good bidentate ligand, and numerous complexes with 5,5′-dmbipy have been prepared, such as that of zinc (Khalighi *et al.*, 2008), mercury (Tadayon Pour *et al.*, 2008), iron (Amani *et al.*, 2007), indium (Kalateh *et al.*, 2008), cadmium (Ahmadi *et al.*, 2008), copper (Albada *et al.*, 2004) and platin (Maheshwari *et al.*, 2007). We report herein the crystal structure of the title compound.

The asymmetric unit of the title compound (Fig.1) contains two halves molecules, in which the bond lengths (Allen *et al.*, 1987) and angles are within normal ranges. Rings A (N1/C1-C4/C6) and B (N2/C7-C10/C12) are, of course, planar and they are oriented at a dihedral angle of A/B = 69.62 (4)°.

In the crystal structure (Fig. 2), the π–π contact between the pyridine rings, Cg2—Cg2^{i} [symmetry code: (i) 1 - x, 2 - y, 2 - z, where Cg2 is centroid of the ring B (N2/C7-C10/C12)] may stabilize the structure, with a centroid-centroid distance of 3.895 (3) Å. There also exists a weak C—H$\cdots\pi$ interaction (Table 1).

S2. Experimental

For the preparation of the title compound, a solution of 5,5′-dimethyl-2,2′ -bipyridine (0.15 g, 0.80 mmol) in methanol (15 ml) was added to a solution of $BaCl_2.2H_2O$, (0.10 g, 0.40 mmol) in water (5 ml) and the resulting colorless solution was stirred for 10 min at 313 K. Then, it was left to evaporate slowly at room temperature. After one week, colorless prismatic crystals of the title compound were isolated.

S3. Refinement

H atoms were positioned geometrically, with C-H = 0.93 and 0.96 Å for aromatic and methyl H, respectively, and constrained to ride on their parent atoms, with $U_{iso}(H) = 1.2U_{eq}(C)$.

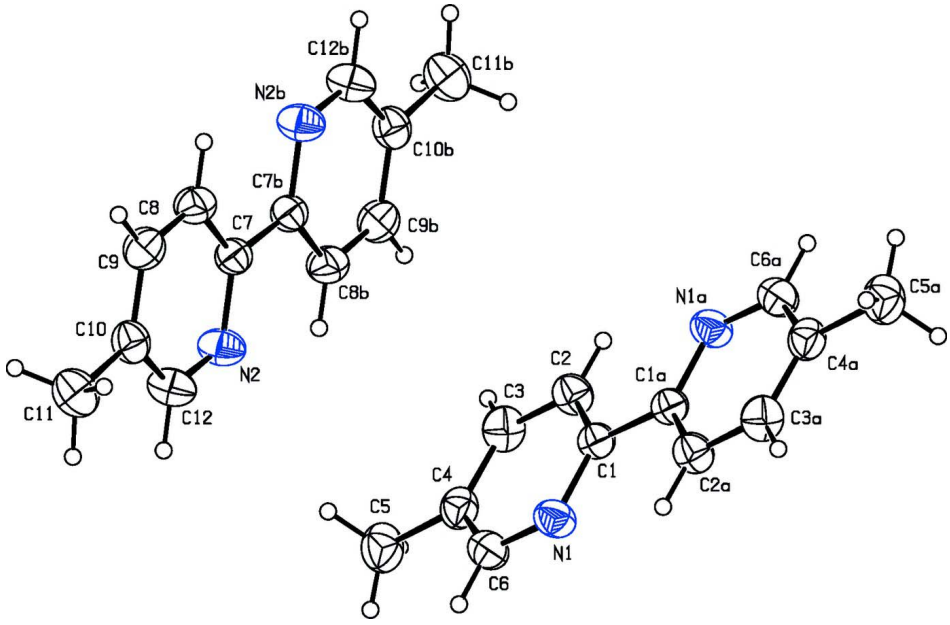

Figure 1

The molecular structure of the title molecule, with the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level [symmetry codes: (a) 1 - x, 2 - y, 1 - z, (b) -x, 2 - y, 2 - z].

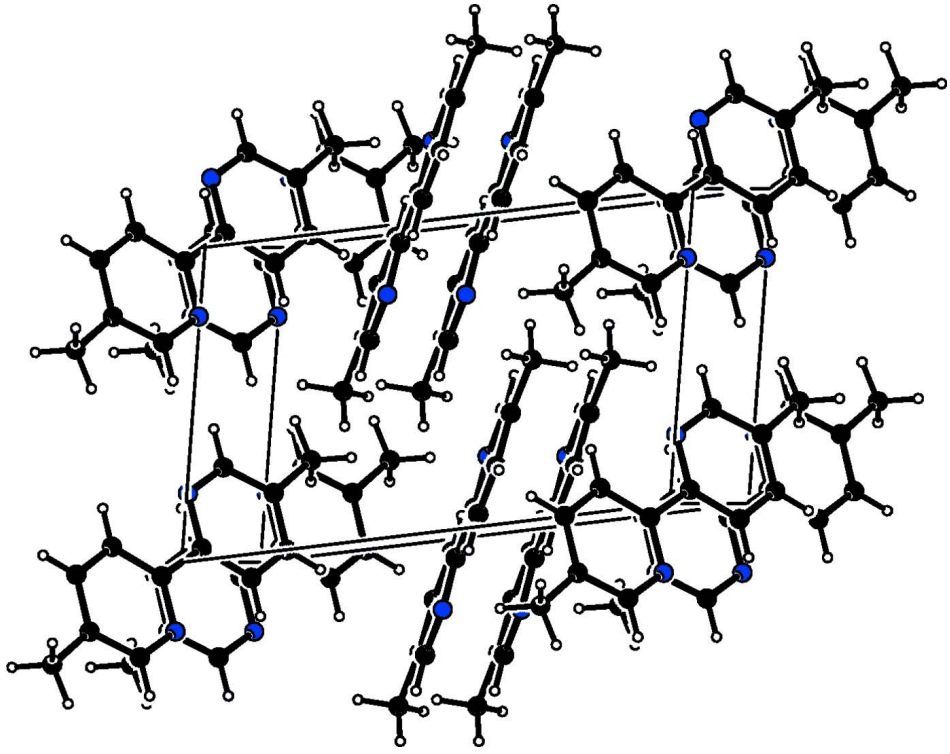

Figure 2

A partial packing diagram of the title compound.

5,5′-Dimethyl-2,2′-bipyridine

Crystal data

$C_{12}H_{12}N_2$
$M_r = 184.24$
Triclinic, $P\overline{1}$
Hall symbol: -P 1
$a = 6.409$ (4) Å
$b = 7.312$ (5) Å
$c = 11.533$ (8) Å
$\alpha = 96.04$ (5)°
$\beta = 91.16$ (4)°
$\gamma = 105.03$ (5)°
$V = 518.4$ (6) Å^3
$Z = 2$
$F(000) = 196$
$D_x = 1.180$ Mg m^{-3}
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 1133 reflections
$\theta = 1.8$–29.3°
$\mu = 0.07$ mm^{-1}
$T = 298$ K
Prism, colorless
0.50 × 0.41 × 0.29 mm

Data collection

Bruker SMART CCD area-detector diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
φ and ω scans
6191 measured reflections
2739 independent reflections
2067 reflections with $I > 2\sigma(I)$
$R_{int} = 0.082$
$\theta_{max} = 29.3°$, $\theta_{min} = 1.8°$
$h = -8 \rightarrow 8$
$k = -10 \rightarrow 10$
$l = -15 \rightarrow 15$

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)] = 0.066$
$wR(F^2) = 0.205$
$S = 1.08$
2739 reflections
129 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
$w = 1/[\sigma^2(F_o^2) + (0.1057P)^2 + 0.0566P]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max} = 0.002$
$\Delta\rho_{max} = 0.27$ e Å^{-3}
$\Delta\rho_{min} = -0.24$ e Å^{-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 (Å^2)

	x	y	z	U_{iso}*/U_{eq}
N1	0.4778 (2)	0.76393 (19)	0.53822 (13)	0.0604 (4)
N2	0.0741 (2)	0.77876 (19)	0.99037 (14)	0.0656 (4)
C1	0.4223 (2)	0.92544 (19)	0.52837 (12)	0.0486 (3)
C2	0.2280 (2)	0.9529 (2)	0.56860 (14)	0.0579 (4)

H2	0.1916	1.0661	0.5601	0.069*
C3	0.0896 (3)	0.8105 (3)	0.62124 (16)	0.0655 (4)
H3	−0.0407	0.8276	0.6480	0.079*
C4	0.1445 (3)	0.6443 (2)	0.63387 (13)	0.0594 (4)
C5	0.0011 (4)	0.4859 (3)	0.69263 (18)	0.0797 (6)
H5C	0.0475	0.3716	0.6761	0.096*
H5B	−0.1459	0.4638	0.6637	0.096*
H5A	0.0105	0.5218	0.7755	0.096*
C6	0.3403 (3)	0.6293 (2)	0.58998 (16)	0.0641 (4)
H6	0.3793	0.5168	0.5971	0.077*
C7	0.0828 (2)	0.96057 (19)	1.02576 (12)	0.0513 (4)
C8	0.2406 (3)	1.0695 (2)	1.10747 (14)	0.0632 (4)
H8	0.2442	1.1958	1.1315	0.076*
C9	0.3905 (3)	0.9892 (3)	1.15227 (15)	0.0679 (5)
H9	0.4973	1.0614	1.2066	0.082*
C10	0.3838 (3)	0.8018 (2)	1.11714 (14)	0.0600 (4)
C11	0.5426 (3)	0.7065 (3)	1.1636 (2)	0.0793 (5)
H11C	0.5070	0.5755	1.1305	0.095*
H11B	0.5374	0.7130	1.2470	0.095*
H11A	0.6856	0.7699	1.1431	0.095*
C12	0.2214 (3)	0.7049 (2)	1.03580 (18)	0.0686 (5)
H12	0.2146	0.5782	1.0109	0.082*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N1	0.0614 (8)	0.0537 (7)	0.0719 (8)	0.0235 (6)	0.0113 (6)	0.0111 (6)
N2	0.0615 (8)	0.0483 (7)	0.0819 (9)	0.0103 (6)	−0.0066 (7)	−0.0034 (6)
C1	0.0470 (7)	0.0513 (7)	0.0491 (7)	0.0170 (6)	−0.0020 (5)	0.0031 (5)
C2	0.0500 (8)	0.0626 (9)	0.0669 (9)	0.0231 (6)	0.0037 (6)	0.0129 (7)
C3	0.0508 (8)	0.0790 (11)	0.0696 (9)	0.0200 (8)	0.0091 (7)	0.0131 (8)
C4	0.0607 (9)	0.0615 (9)	0.0513 (7)	0.0076 (7)	0.0001 (6)	0.0067 (6)
C5	0.0809 (13)	0.0784 (12)	0.0723 (11)	0.0034 (10)	0.0095 (9)	0.0176 (9)
C6	0.0734 (10)	0.0539 (8)	0.0690 (9)	0.0212 (7)	0.0104 (8)	0.0125 (7)
C7	0.0527 (7)	0.0454 (7)	0.0523 (7)	0.0062 (6)	0.0075 (6)	0.0052 (5)
C8	0.0722 (10)	0.0521 (8)	0.0608 (8)	0.0129 (7)	−0.0068 (7)	−0.0031 (6)
C9	0.0727 (10)	0.0664 (10)	0.0595 (9)	0.0131 (8)	−0.0108 (7)	0.0003 (7)
C10	0.0567 (8)	0.0625 (9)	0.0620 (8)	0.0137 (7)	0.0086 (7)	0.0158 (7)
C11	0.0705 (11)	0.0809 (13)	0.0911 (13)	0.0247 (10)	−0.0004 (10)	0.0194 (10)
C12	0.0640 (10)	0.0500 (8)	0.0896 (12)	0.0137 (7)	−0.0015 (8)	0.0017 (8)

Geometric parameters (Å, º)

C1—N1	1.334 (2)	C7—N2	1.335 (2)
C1—C2	1.393 (2)	C7—C8	1.389 (2)
C1—C1^{i}	1.491 (3)	C7—C7^{ii}	1.475 (3)
C2—C3	1.384 (3)	C8—C9	1.368 (3)
C2—H2	0.9300	C8—H8	0.9300

C3—C4	1.371 (3)	C9—C10	1.377 (3)
C3—H3	0.9300	C9—H9	0.9300
C4—C6	1.389 (3)	C10—C12	1.380 (3)
C4—C5	1.511 (3)	C10—C11	1.494 (3)
C5—H5C	0.9600	C11—H11C	0.9600
C5—H5B	0.9600	C11—H11B	0.9600
C5—H5A	0.9600	C11—H11A	0.9600
C6—N1	1.340 (2)	C12—N2	1.327 (2)
C6—H6	0.9300	C12—H12	0.9300

N1—C1—C2	121.57 (15)	N2—C7—C7^{ii}	116.86 (16)
N1—C1—C1^{i}	116.75 (16)	C8—C7—C7^{ii}	121.84 (17)
C2—C1—C1^{i}	121.68 (16)	C9—C8—C7	119.40 (16)
C3—C2—C1	119.51 (15)	C9—C8—H8	120.3
C3—C2—H2	120.2	C7—C8—H8	120.3
C1—C2—H2	120.2	C8—C9—C10	120.16 (16)
C4—C3—C2	119.99 (16)	C8—C9—H9	119.9
C4—C3—H3	120.0	C10—C9—H9	119.9
C2—C3—H3	120.0	C9—C10—C12	116.32 (17)
C3—C4—C6	116.29 (16)	C9—C10—C11	122.45 (18)
C3—C4—C5	122.09 (17)	C12—C10—C11	121.22 (17)
C6—C4—C5	121.62 (17)	C10—C11—H11C	109.5
C4—C5—H5C	109.5	C10—C11—H11B	109.5
C4—C5—H5B	109.5	H11C—C11—H11B	109.5
H5C—C5—H5B	109.5	C10—C11—H11A	109.5
C4—C5—H5A	109.5	H11C—C11—H11A	109.5
H5C—C5—H5A	109.5	H11B—C11—H11A	109.5
H5B—C5—H5A	109.5	N2—C12—C10	124.98 (16)
N1—C6—C4	125.26 (16)	N2—C12—H12	117.5
N1—C6—H6	117.4	C10—C12—H12	117.5
C4—C6—H6	117.4	C1—N1—C6	117.36 (15)
N2—C7—C8	121.30 (15)	C12—N2—C7	117.84 (15)

N1—C1—C2—C3	0.7 (2)	C8—C9—C10—C12	0.4 (3)
C1^{i}—C1—C2—C3	−179.66 (16)	C8—C9—C10—C11	−179.67 (17)
C1—C2—C3—C4	0.3 (3)	C9—C10—C12—N2	−0.2 (3)
C2—C3—C4—C6	−0.9 (3)	C11—C10—C12—N2	179.92 (17)
C2—C3—C4—C5	178.87 (16)	C2—C1—N1—C6	−0.9 (2)
C3—C4—C6—N1	0.7 (3)	C1^{i}—C1—N1—C6	179.44 (15)
C5—C4—C6—N1	−179.06 (16)	C4—C6—N1—C1	0.2 (3)
N2—C7—C8—C9	0.2 (3)	C10—C12—N2—C7	−0.1 (3)
C7^{ii}—C7—C8—C9	−179.88 (17)	C8—C7—N2—C12	0.0 (3)
C7—C8—C9—C10	−0.5 (3)	C7^{ii}—C7—N2—C12	−179.87 (17)

Symmetry codes: (i) −*x*+1, −*y*+2, −*z*+1; (ii) −*x*, −*y*+2, −*z*+2.

Hydrogen-bond geometry (Å, º)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
C9—H9···*Cg*1^{iii}	0.93	2.78	3.669 (3)	160

Symmetry code: (iii) −*x*+1, −*y*+2, −*z*+2.