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4,4′-Bis(trimethylsilyl)-2,2′-bipyridine

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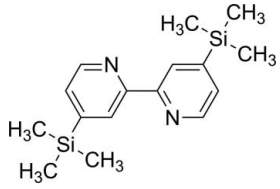
Key indicators: single-crystal X-ray study; T = 223 K; mean σ(C–C) = 0.003 Å; R factor = 0.046; wR factor = 0.125; data-to-parameter ratio = 17.4.

In the molecule of title compound, $C_{16}H_{24}N_2Si_2$, the pyridine rings are nearly planar (r.m.s. deviation = 0.002 Å).

Related literature

For the structure of 5,5′-bis(trimethylsilyl)-2,2′-bipyridines, see: Stange *et al.* (2000). For the structure of 4-trimethylsilylpyridine, see: Postigo & Rossi (2001). For synthetic procedure to obtain 4,4′-bis(methoxyl)-2,2′-bipyridine, see: Wenkert & Woodward (1983).



Experimental

Crystal data

$C_{16}H_{24}N_2Si_2$ M_r = 300.55
Monoclinic, $P2_1/c$
a = 13.154 (4) Å
b = 6.4599 (16) Å
c = 11.280 (3) Å
β = 111.222 (6)°
V = 893.5 (4) Å^3
Z = 2
Mo $K\alpha$ radiation
μ = 0.19 mm^{-1}
T = 223 K
0.50 × 0.30 × 0.20 mm

Data collection

Rigaku Saturn CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
T_{min} = 0.869, T_{max} = 0.963
4311 measured reflections
1649 independent reflections
1364 reflections with $I > 2\sigma(I)$
R_{int} = 0.027

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.046
$wR(F^2)$ = 0.125
S = 1.08
1649 reflections
95 parameters
2 restraints
H-atom parameters constrained
$\Delta\rho_{max}$ = 0.25 e Å^{-3}
$\Delta\rho_{min}$ = −0.26 e Å^{-3}

Data collection: *CrystalClear* (Rigaku, 2005); 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: RK2303).

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supporting information

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4,4′-Bis(trimethylsilyl)-2,2′-bipyridine

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S1. Comment

Derivatives of 2,2′-bipyridine have received much attention due to their potential to form polypyridyl metal complexes, particularly of ruthenium and rhenium which have diverse applications. The photochemical and redox properties of these complexes can be varied through appropriate substitution on the pyridine rings. The derivatization of a 2,2′-bipyridine ligand with electron donating/withdrawing groups in the 4,4′-positions has been a popular means of controlling the redox potential of transition metal bipyridine complexes. The 4,4′-disubstitution can also offers no steric complications on complexation. The research about synthesis and properties of pyridine rings interconnected with strong electron-donating groups, as well as trimethysilyl group, was recently reported (Stange *et al.*, 2000; Postigo & Rossi, 2001). However, there are no reports on 4,4′-bis(trimethylsilyl)-2,2′-bipyridine. Herein, we report crystal structure of the title compound.

The molecule is placed in centre of symmetry and nealy flat (Fig. 1) as the C—Si is co-planar with the aromatic rings. The torsion angle for N1—C4—C4^{i}—C5^{i} = 0°. In crystal, molecules are connected by weak non-classical intermolecular C3–H3···N1^{ii} hydrogen bonds with parameters C3···N1^{ii} = 3.626 (2) Å, H3···N1^{ii} = 2.714 Å and angle C3–H3···N1^{ii} = 164.3°. Symmetry codes: (i) -*x* + 1, -*y* + 1, -*z*; (ii) -*x* + 1, *y* - 1/2, -*z* + 1/2.

S2. Experimental

All the reagents and solvents empolyed were commercially available. The title compound was synthesized by using 2,2′-bipyridine as the starting material with successive polystepreactions (Wenkert & Woodward, 1983). The final product was dissolved in the solution of methanol and methylene chloride, which diffused slowly. After seven days, colourless block-shaped crystals were obtained which were suitable for X-ray analysis.

S3. Refinement

All H atoms were placed in geometrically idealized positions. H atoms of bipyridine constrained to ride on their parent atoms with C—H = 0.94 Å and refined with U_{iso}(H) = 1.2U_{iso}(C). The H atoms of methyl groups constrained to ride on their parent atoms with C—H = 0.97 Å and refined with U_{iso}(H) = 1.5U_{iso}(C).

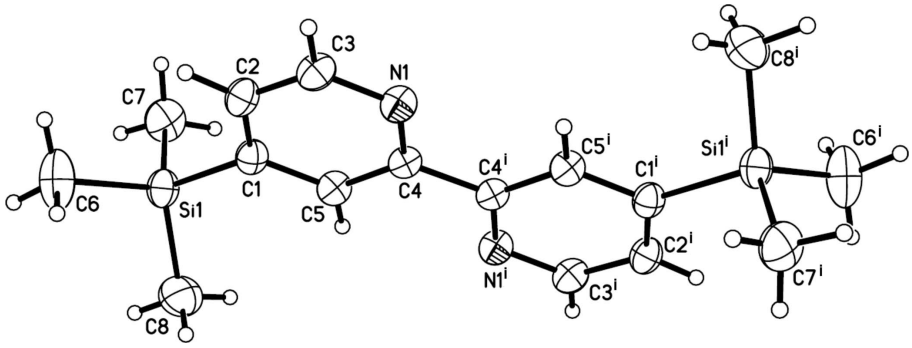

Figure 1

Molecular structure of title compound with the atom numbering scheme. Displacement ellipsoids are drawn at 40% probability level. H atoms are presented as a small spheres of arbitrary radius. Symmetry code: (i) -*x* + 1, -*y* + 1, -*z*.

4,4′-Bis(trimethylsilyl)-2,2′-bipyridine

Crystal data

$C_{16}H_{24}N_2Si_2$
$M_r = 300.55$
Monoclinic, $P2_1/c$
Hall symbol: -P 2ybc
$a = 13.154$ (4) Å
$b = 6.4599$ (16) Å
$c = 11.280$ (3) Å
$\beta = 111.222$ (6)°
$V = 893.5$ (4) Å^3
$Z = 2$
$F(000) = 324$
$D_x = 1.117$ Mg m^{-3}
Mo $K\alpha$ radiation, $\lambda = 0.71075$ Å
Cell parameters from 3568 reflections
$\theta = 3.2$–27.5°
$\mu = 0.19$ mm^{-1}
$T = 223$ K
Block, colourless
0.50 × 0.30 × 0.20 mm

Data collection

Rigaku Saturn CCD diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
Detector resolution: 14.63 pixels mm^{-1}
ω scans
Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
$T_{min} = 0.869$, $T_{max} = 0.963$
4311 measured reflections
1649 independent reflections
1364 reflections with $I > 2\sigma(I)$
$R_{int} = 0.027$
$\theta_{max} = 25.5°$
$h = -13 \rightarrow 15$
$k = -7 \rightarrow 7$
$l = -13 \rightarrow 12$

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)] = 0.046$
$wR(F^2) = 0.125$
$S = 1.08$
1649 reflections
95 parameters
2 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
$w = 1/[\sigma^2(F_o^2) + (0.0734P)^2 + 0.0832P]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max} < 0.001$
$\Delta\rho_{max} = 0.25$ e Å^{-3}
$\Delta\rho_{min} = -0.26$ e Å^{-3}

Special details

Geometry. All s.u.'s (except the s.u. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell s.u.'s are taken into account individually in the estimation of s.u.'s in distances, angles and torsion angles; correlations between s.u.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell s.u.'s is used for estimating s.u.'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 (Å²)

	x	*y*	*z*	U_{iso}*/U_{eq}
Si1	0.80366 (4)	0.08423 (9)	0.02023 (5)	0.0405 (2)
N1	0.49321 (13)	0.2932 (2)	0.10461 (14)	0.0366 (4)
C1	0.68120 (15)	0.1675 (3)	0.05735 (17)	0.0356 (5)
C2	0.64022 (16)	0.0538 (3)	0.13516 (18)	0.0383 (5)
H2	0.6754	−0.0686	0.1740	0.046*
C3	0.54791 (17)	0.1206 (3)	0.15535 (18)	0.0391 (5)
H3	0.5221	0.0400	0.2079	0.047*
C4	0.53132 (15)	0.4054 (3)	0.02861 (17)	0.0311 (4)
C5	0.62411 (16)	0.3477 (3)	0.00480 (17)	0.0353 (5)
H5	0.6487	0.4314	−0.0475	0.042*
C6	0.8938 (2)	−0.0751 (5)	0.1542 (3)	0.0724 (9)
H6A	0.8573	−0.2038	0.1584	0.109*
H6B	0.9101	0.0003	0.2333	0.109*
H6C	0.9611	−0.1049	0.1408	0.109*
C7	0.7569 (2)	−0.0682 (4)	−0.1293 (2)	0.0576 (6)
H7A	0.8197	−0.1205	−0.1457	0.086*
H7B	0.7140	0.0195	−0.1993	0.086*
H7C	0.7127	−0.1834	−0.1210	0.086*
C8	0.8778 (2)	0.3196 (4)	0.0009 (3)	0.0687 (7)
H8A	0.8948	0.4060	0.0760	0.103*
H8B	0.8323	0.3964	−0.0732	0.103*
H8C	0.9448	0.2790	−0.0101	0.103*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Si1	0.0418 (4)	0.0433 (4)	0.0397 (4)	0.0085 (2)	0.0189 (3)	0.0001 (2)
N1	0.0437 (9)	0.0376 (9)	0.0332 (8)	0.0016 (7)	0.0195 (7)	0.0033 (7)
C1	0.0387 (10)	0.0352 (10)	0.0323 (9)	0.0035 (8)	0.0120 (8)	−0.0037 (8)
C2	0.0451 (11)	0.0361 (10)	0.0339 (10)	0.0059 (9)	0.0144 (9)	0.0028 (8)
C3	0.0496 (12)	0.0373 (11)	0.0347 (10)	−0.0001 (9)	0.0202 (9)	0.0051 (8)
C4	0.0368 (10)	0.0307 (9)	0.0277 (9)	−0.0011 (8)	0.0139 (8)	−0.0013 (7)
C5	0.0410 (10)	0.0377 (10)	0.0317 (9)	0.0005 (8)	0.0185 (8)	−0.0008 (8)
C6	0.0688 (17)	0.093 (2)	0.0590 (15)	0.0392 (15)	0.0270 (14)	0.0160 (14)
C7	0.0642 (15)	0.0625 (15)	0.0532 (13)	0.0052 (12)	0.0297 (12)	−0.0119 (11)
C8	0.0544 (14)	0.0656 (16)	0.097 (2)	−0.0045 (13)	0.0409 (14)	−0.0089 (15)

Geometric parameters (Å, º)

Si1—C7	1.855 (2)	C4—C4^{i}	1.485 (3)
Si1—C6	1.858 (2)	C5—H5	0.9400
Si1—C8	1.861 (3)	C6—H6A	0.9700
Si1—C1	1.884 (2)	C6—H6B	0.9700
N1—C3	1.338 (2)	C6—H6C	0.9700
N1—C4	1.350 (2)	C7—H7A	0.9700
C1—C2	1.394 (3)	C7—H7B	0.9700
C1—C5	1.396 (3)	C7—H7C	0.9700
C2—C3	1.383 (3)	C8—H8A	0.9700
C2—H2	0.9400	C8—H8B	0.9700
C3—H3	0.9400	C8—H8C	0.9700
C4—C5	1.392 (3)		
C7—Si1—C6	110.44 (13)	C4—C5—H5	119.5
C7—Si1—C8	110.03 (13)	C1—C5—H5	119.5
C6—Si1—C8	109.91 (14)	Si1—C6—H6A	109.5
C7—Si1—C1	108.97 (10)	Si1—C6—H6B	109.5
C6—Si1—C1	108.84 (11)	H6A—C6—H6B	109.5
C8—Si1—C1	108.60 (10)	Si1—C6—H6C	109.5
C3—N1—C4	116.95 (17)	H6A—C6—H6C	109.5
C2—C1—C5	115.83 (18)	H6B—C6—H6C	109.5
C2—C1—Si1	123.10 (15)	Si1—C7—H7A	109.5
C5—C1—Si1	121.05 (15)	Si1—C7—H7B	109.5
C3—C2—C1	120.11 (18)	H7A—C7—H7B	109.5
C3—C2—H2	119.9	Si1—C7—H7C	109.5
C1—C2—H2	119.9	H7A—C7—H7C	109.5
N1—C3—C2	123.92 (18)	H7B—C7—H7C	109.5
N1—C3—H3	118.0	Si1—C8—H8A	109.5
C2—C3—H3	118.0	Si1—C8—H8B	109.5
N1—C4—C5	122.12 (17)	H8A—C8—H8B	109.5
N1—C4—C4^{i}	116.3 (2)	Si1—C8—H8C	109.5
C5—C4—C4^{i}	121.6 (2)	H8A—C8—H8C	109.5
C4—C5—C1	121.06 (18)	H8B—C8—H8C	109.5
C7—Si1—C1—C2	92.79 (18)	C4—N1—C3—C2	0.6 (3)
C6—Si1—C1—C2	−27.7 (2)	C1—C2—C3—N1	−0.4 (3)
C8—Si1—C1—C2	−147.35 (18)	C3—N1—C4—C5	−0.9 (3)
C7—Si1—C1—C5	−85.66 (18)	C3—N1—C4—C4^{i}	179.05 (18)
C6—Si1—C1—C5	153.85 (17)	N1—C4—C5—C1	1.1 (3)
C8—Si1—C1—C5	34.20 (19)	C4^{i}—C4—C5—C1	−178.86 (19)
C5—C1—C2—C3	0.5 (3)	C2—C1—C5—C4	−0.9 (3)
Si1—C1—C2—C3	−178.01 (14)	Si1—C1—C5—C4	177.70 (14)

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

Hydrogen-bond geometry (Å, º)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
C3—H3···N1^{ii}	0.94	2.71	3.626 (2)	164

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