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Key indicators

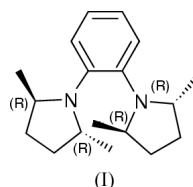
Single-crystal X-ray study
 $T = 123$ K
Mean $\sigma(\text{C}-\text{C}) = 0.002$ Å
 R factor = 0.033
 wR factor = 0.084
Data-to-parameter ratio = 12.6For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.1,2-Bis[(2*R*,5*R*)-2,5-dimethylpyrrolidin-1-yl]-
benzeneThe title compound, $\text{C}_{18}\text{H}_{28}\text{N}_2$, with a crystallographic twofold
rotation axis, can function as a C_2 -symmetric diamine ligand.

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Comment

 C_2 -Symmetric diphosphine ligands, such as Me-DuPHOS [1,2-
bis(2,5-dimethylphospholano)benzene; Burk, 1991], have
been used extensively in enantioselective hydrogenation
reactions (Burk *et al.*, 1993). Chiral diamines have been shown
to act as effective chiral modifiers of the osmium tetroxide
dihydroxylation of alkenes (Johnson & Sharpless, 2000). The
nitrogen analogue, (I), of Me-DuPHOS has been prepared
and evaluated as a ligand for use in the osmium tetroxide
dihydroxylation of stilbene. An X-ray structure determination
of (I) was undertaken to confirm the stereochemistry (Fig. 1).
The molecule has crystallographic twofold rotation symmetry.

Experimental

1-Amino-2-[(2*R*,5*R*)-2,5-dimethylpyrrolidin-1-yl]benzene (400 mg,
2.11 mmol) was added to a solution of (2*S*,5*S*)-2,5-hexanediol cyclic
sulfate (Burk *et al.*, 1993) (379 mg, 2.11 mmol) in tetrahydrofuran
(THF, 100 ml). The resulting orange solution was refluxed for 2 d
after the addition of 80% sodium hydride (633 mg, 21.10 mmol). As
thin-layer chromatography indicated the presence of unreacted
aminophenyl pyrrolidine, more cyclic sulfate (379 mg, 2.11 mmol)
dissolved in THF (30 ml) was added. The mixture was stirred at reflux
for a further 2 d and then quenched with 10% NH_4Cl (30 ml). The
THF was removed under reduced pressure and the aqueous mixture
extracted with dichloromethane (3×100 ml). The combined organic
extract was washed with water (100 ml) and brine (100 ml), dried
(MgSO_4), filtered and evaporated to give a brown semi-solid. Puri-
fication using column chromatography (10% ethyl acetate/hexane)
gave the diamine (I) as a colourless solid (408 mg, 71%); m.p. 381–
385 K. Diethyl ether (10 ml) was added to the diamine and single
crystals suitable for X-ray structure determination were obtained by
slow evaporation of the solvent.

Crystal data

 $\text{C}_{18}\text{H}_{28}\text{N}_2$
 $M_r = 272.42$
Orthorhombic, $P2_21$
 $a = 8.5423$ (2) Å
 $b = 8.8892$ (2) Å
 $c = 10.6486$ (2) Å
 $V = 808.59$ (3) Å³
 $Z = 2$
 $D_x = 1.119$ Mg m⁻³Mo $K\alpha$ radiation
Cell parameters from 11 522
reflections
 $\theta = 3.8$ – 28.3°
 $\mu = 0.07$ mm⁻¹
 $T = 123$ (2) K
Prism, colourless
 $0.25 \times 0.20 \times 0.18$ mm

Data collection

Nonius KappaCCD diffractometer
 Thick-slice φ and ω scans
 11 552 measured reflections
 1174 independent reflections
 1094 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.043$
 $\theta_{\text{max}} = 28.3^\circ$
 $h = -11 \rightarrow 11$
 $k = -11 \rightarrow 11$
 $l = -14 \rightarrow 14$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.033$
 $wR(F^2) = 0.084$
 $S = 1.07$
 1174 reflections
 93 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0398P)^2 + 0.1166P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.17 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.16 \text{ e } \text{\AA}^{-3}$

H atoms were included in the riding-model approximation [methine C–H = 1.00 Å, methylene C–H = 0.99 Å, methyl C–H = 0.98 Å and aromatic C–H = 0.95 Å; $U_{\text{iso}}(\text{H})$ is $1.2U_{\text{eq}}(\text{C})$]. The absolute configuration cannot be determined reliably from the diffraction data and was assumed from the synthesis; Friedel opposites were merged in the final cycles of refinement.

Data collection: *COLLECT* (Nonius, 1997–2002); cell refinement: *HKL SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *HKL DENZO* (Otwinowski & Minor, 1997) and *SCALEPACK*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* publication routines (Farrugia, 1999).

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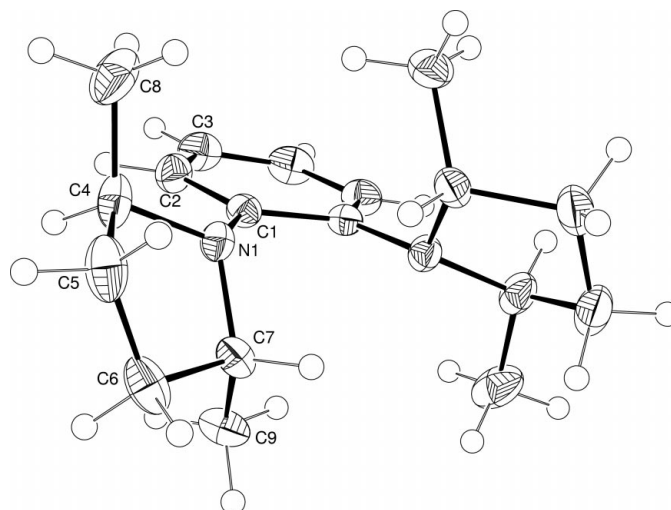


Figure 1
View of (I) (50% probability displacement ellipsoids)

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