organic papers

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Key indicators

Single-crystal X-ray study T = 298 KMean $\sigma(C-C) = 0.003 \text{ Å}$ R factor = 0.054 wR factor = 0.142 Data-to-parameter ratio = 15.1

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

N,N,N',N'-Tetrakis(2-pyridylmethyl)-1,2diaminoethane: a multidentate ligand

The reaction of 1,2-diaminoethane and 2-pyridinecarboxyaldehyde in the presence of a reducing agent, sodium triacetoxyborohydride, leads to the title compound, $C_{26}H_{28}N_6$. The molecule has a centre of symmetry. Received 14 January 2004 Accepted 16 January 2004 Online 23 January 2004

Comment

The title compound, (I), reveals a wide variety of coordination modes in binding to metal ions, having six N donor atoms, and there are many studies of metal complexes with (I). This compound acts not only as a hexadentate ligand, but also as a pentadentate or a bridging ligand, forming a dinuclear complex in some cases. One of the previous synthetic methods for producing (I) was based on the reaction of 2-picolyl chloride hydrochloride and ethylenediamine, although the yield was not very high. Another synthetic route was an alkylation of ethylenediamine with 2-(chloromethyl)pyridinium chloride in the presence of hexadecyltrimethylammonium chloride as a phase-transfer catalyst, and this gave the corresponding tetrakis(2-pyridylmethyl)alkanediamines in good yields (Sato et al., 1992; Tamura et al., 2000). This report describes a new synthetic method via a reductive amination reaction, and the crystal structure of the product, viz. the title compound, (I), is presented.



As depicted in the Scheme, compound (I) can be obtained from the reductive amination of 2-pyridinecarboxyaldehyde, in 1,2-dichloroethane as a solvent, with sodium triacetoxyborohydride, which is used as a general reducing agent for the reductive amination of aldehydes and ketones (Abdel-Magid *et al.*, 1996). The product was easily purified by chromatography on a short silica-gel column and subsequent recrystallization from *n*-hexane–ethyl acetate. The yield of (I) by the present method (46%) is comparable with that of the reported method using a phase-transfer catalyst (68%). The present method would be convenient for preparing N,N,N',N'tetrakis(2-pyridylmethyl)- α,ω -alkanediamine.

The structure of (I), with the atom-numbering scheme, is shown in Fig. 1. This is the first X-ray crystallographic study of the unprotonated form of (I). The ring angles at N in the

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pyridine rings are significantly smaller [117.41 (15) and 117.54 (16) at N2 and N3, respectively] than those of the protonated form (ca 123°; Gunatilleke & Norman, 2003). Other structural parameters of (I) are comparable with the values reported for related compounds (Mandel & Douglas, 1989).

Experimental

To 1,2-dichloroethane (15 ml) in a flask were added 1,2-diaminoethane (0.3 g, 4.0 mmol) and 2-pyridinecarboxyaldehyde (2.1 g, 19 mmol), and the mixture was stirred for 30 min at room temperature (293 K) under an argon atmosphere. Small amounts of sodium triacetoxyborohydride (3.5 g, 17 mmol) and CH₂ClCH₂Cl (10 ml) were added and the solution was stirred for 24 h at room temperature under an argon atmosphere, yielding a yellow solution. After neutralization with saturated aqueous NaHCO₃ (25 ml) to give a weakly alkaline solution, the product was extracted with CHCl₃ (20 ml), which had been dried by adding anhydrous Na₂SO₄ (6 g). The crude product was purified by silica-gel column chromatography with toluene–ethanol mixed eluent (20:1 v/v). A pale-vellow oily residue containing toluene was obtained by evaporation of the solvent under reduced pressure, and this residue was subjected to successive drying in vacuo. The resulting crystalline solid was recrystallized from *n*-hexane–ethyl acetate and dried *in vacuo* in a desiccator. Colourless crystals of (I) were obtained in 46% yield [m.p. 383.65-384.15 K; literature value (Anderegg & Wenk, 1967): m.p. 383.15–384.65 K]. Spectroscopic analysis: ¹H NMR (CDCl₃, TMS_{int}, δ, p.p.m.): 8.48 (4*H*, *d*, ${}^{1}J$ = 5.0 Hz, ${}^{2}J$ = 1.8 Hz), 7.57 (4*H*, *t*, ${}^{1}J$ = 7.7 Hz, ${}^{2}J = 1.8$ Hz), 7.45 (4*H*, *d*, *J* = 8.1 Hz), 7.11 (4*H*, *t*, ${}^{1}J = 7.4$ Hz, ${}^{2}J =$ 1.3 Hz), 3.78 (8*H*, *s*), 2.76 (4*H*, *s*); FAB-MS (NBA): m/z = 425 $([M + H]^+, 100\%).$

Crystal data

| C ₂₆ H ₂₈ N ₆ | Z = 1 | |
|--|---|--|
| $M_r = 424.54$ | $D_x = 1.179 \text{ Mg m}^{-3}$ | |
| Triclinic, P1 | Mo $K\alpha$ radiation | |
| a = 6.228(1) Å | Cell parameters from 954 | |
| b = 9.482(2) Å | reflections | |
| c = 10.760 (2) Å | $\theta = 2.7 - 22.9^{\circ}$ | |
| $\alpha = 77.66 \ (3)^{\circ}$ | $\mu = 0.07 \text{ mm}^{-1}$ | |
| $\beta = 87.74 \ (3)^{\circ}$ | T = 298 (2) K | |
| $\gamma = 74.47 \ (3)^{\circ}$ | Prism, colourless | |
| $V = 598.0 (2) \text{ Å}^3$ | $0.48 \times 0.45 \times 0.35 \text{ mm}$ | |

Data collection

Bruker SMART APEX CCD areadetector diffractometer $R_{\rm int}=0.018$ ω and ω scans Absorption correction: multi-scan $\theta_{\rm max} = 27.9^{\circ}$ (SADABS; Sheldrick, 1996) $h = -8 \rightarrow 8$ $k = -12 \rightarrow 10$ $T_{\rm min}=0.966,\;T_{\rm max}=0.977$ $l=-14\rightarrow13$ 4392 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.054$ $wR(F^2) = 0.142$ S=1.022824 reflections 187 parameters Only coordinates of H atoms refined

2824 independent reflections 1800 reflections with $I > 2\sigma(I)$

 $w = 1/[\sigma^2(F_o^2) + (0.0689P)^2]$ + 0.0224P] where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.12 \text{ e } \text{\AA}^{-3}$ $\Delta \rho_{\rm min} = -0.12 \text{ e} \text{ Å}^{-3}$



Figure 1

The molecular structure of (I), with 50% probability displacement ellipsoids and the atom-numbering scheme. Atoms labelled with a prime are at symmetry position (-x, 1 - y, 1 - z).

Table 1

Selected geometric parameters (Å, °).

| Commentation and as (i) | 1 1 | | |
|-------------------------|-------------|--------------------|-------------|
| N3-C9-C8-N1 | -130.8 (1) | N2-C3-C2-N1 | -131.5 (2) |
| C3-N2-C7 | 117.41 (15) | C9-N3-C13 | 117.54 (16) |
| C3-C2 | 1.504 (2) | C11-C12 | 1.360 (3) |
| C3-C4 | 1.379 (2) | C5-C6 | 1.361 (3) |
| N3-C13 | 1.337 (2) | C10-C11 | 1.368 (3) |
| N3-C9 | 1.331 (2) | C13-C12 | 1.361 (3) |
| N2-C7 | 1.337 (2) | C7-C6 | 1.366 (3) |
| N2-C3 | 1.330 (2) | C4-C5 | 1.384 (3) |
| N1-C2 | 1.467 (2) | C1-C1 ⁱ | 1.507 (3) |
| N1-C1 | 1.464 (2) | C9-C8 | 1.499 (2) |
| N1-C8 | 1.459 (2) | C9-C10 | 1.376 (2) |

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

All H atoms were located in difference Fourier maps and refined with $U_{iso}(H) = 1.2U_{eq}$ (parent atom). The C-H bond lengths are in the range 0.92 (2)–1.03 (2) Å.

Data collection: SMART-W2K/NT (Bruker, 2003); cell refinement: SMART-W2K/NT; data reduction: SAINT-W2K/NT (Bruker, 2003); program(s) used to solve structure: SHELXTL-NT (Bruker, 2003); program(s) used to refine structure: SHELXTL-NT; molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: SHELXTL-NT.

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