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#### Key indicators

Single-crystal X-ray study T = 173 K Mean  $\sigma$ (C–C) = 0.003 Å R factor = 0.042 wR factor = 0.112 Data-to-parameter ratio = 16.3

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

# (1,2-Bis{[3-(2-pyridyl)pyrazol-1-yl]methyl}benzene)perchloratosodium

In the title compound,  $[Na(ClO_4)(C_{24}H_{20}N_6)]$ , the sodium cation is in a square-pyramidal geometry, which is distorted from the ideal ( $\tau = 0.19$ ), with the four N-donor atoms of the bis-bidentate chelating units from the ligand arranged equatorially, and with the perchlorate anion at the apical site.

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## Comment

The coordination chemistry of the ligand L1 has been well documented for the first-row transition metals; it forms tetrahedral cages and open-chain dinuclear structures (Fleming *et al.*, 1998; Paul *et al.*, 2002). Several complexes of alkali metals with N-donor ligands are observed in the literature, although they usually employ highly preorganized ligands, where the cavity size is comparable to that of the alkali metal ion and the coordination geometry is particularly unsuited to a transition metal. Examples include ligands such as torands (Bell *et al.*, 1992, 1993), macrocycles (Constable *et al.*, 1988) and cryptands (Lehn & DeVains, 1992). However, a helical structure containing K<sup>+</sup> ions has been reported, based on a hexadentate N-donor ligand which consists of two terdentate binding domains (Psillakis *et al.*, 1997).





The molecular structure of the title compound, (I), is shown in Fig. 1. It is a relatively unusual example of an alkali metal complex with an open-chain N-donor chelating ligand. The sodium cation is surrounded by four nitrogen donors from L1and a single perchlorate anion, thus yielding a neutral complex. The average Na-N bond length is 2.487 Å, which is comparable to the average Na-N bond length found in six-

 $\bigcirc$  2003 International Union of Crystallography Printed in Great Britain – all rights reserved coordinate chelate complexes with the bidentate phen unit (Qian et al., 1994). Although there are many sodium complexes reported with cryptands (Echegoyen et al., 1991; McKee et al., 1992), the comparison of bond lengths can be complex, since it is thought that they are, in part, dependent on the rigidity of the ligand and the coordination number of the Na<sup>+</sup> ion (Caron et al., 1985). The N atoms of the ligand unit are not bound to the sodium ion in a plane. One of the pyrazole-pyridine units is approximately planar, but the other has an angle of  $17.1 (2)^{\circ}$  between the rings. The approximate angle between the bidentate arms is 131.9  $(14)^{\circ}$ , thus enabling the perchlorate anion to coordinate comfortably to the Na<sup>+</sup> ion in the apical position. The Na-O bond length is 2.3359 (18) Å.

For a five-coordinate geometry, there is a structural continuum between the ideal square-pyramidal (SP) and trigonal-bipyramidal (TBP) extremes. The structural index parameter  $\tau$  (degree of trigonality) can be defined as  $\tau =$  $(\beta - \alpha)/60$ , where  $\beta$  and  $\alpha$  are the largest and smallest angles, respectively, at the central atom.  $\tau$  is equal to zero for a perfect SP geometry, and equal to unity for a perfect TBP geometry (Addison *et al.*, 1984). For the title compound,  $\tau = 0.19$ , indicating that the coordination geometry of sodium is closer to, but distorted from, ideal square pyramidal.

# **Experimental**

Crystals of the title compound were obtained via a serendipitous route upon reaction of (CH<sub>3</sub>CO<sub>2</sub>)<sub>3</sub>Mn·2H<sub>2</sub>O and L1 in a 2:3 ratio in methanol. An aqueous solution of NaClO4 was added, to afford a white precipitate, which was collected by filtration. The title compound was isolated upon recrystallization of the precipitate from MeCN/Et<sub>2</sub>O.

#### Crystal data

$[Na(ClO_4)(C_{24}H_{20}N_6)]$	Z = 2
$M_r = 514.90$	$D_x = 1.471 \text{ Mg m}^{-3}$
Triclinic, P1	Mo $K\alpha$ radiation
a = 9.353 (4)  Å	Cell parameters from 54
b = 10.558 (3) Å	reflections
c = 12.794 (4) Å	$\theta = 0.8  1.0^{\circ}$
$\alpha = 100.26 \ (3)^{\circ}$	$\mu = 0.23 \text{ mm}^{-1}$
$\beta = 107.64 \ (3)^{\circ}$	T = 173 (2)  K
$\gamma = 97.33 \ (3)^{\circ}$	Block, colourless
$V = 1162.3 (8) \text{ Å}^3$	$0.20\times0.15\times0.12$ mm
Data collection	

#### Data collection

Bruker SMART CCD area-detector	5304 independent reflect
diffractometer	3677 reflections with $I >$
Narrow-frame $\omega$ scans	$R_{\rm int} = 0.036$
Absorption correction: multi-scan	$\theta_{\rm max} = 27.5^{\circ}$
(SADABS; Sheldrick, 1996)	$h = -12 \rightarrow 12$
$T_{\min} = 0.824, T_{\max} = 0.970$	$k = -13 \rightarrow 13$
12 525 measured reflections	$l = -16 \rightarrow 16$

## Refinement

Refinement on  $F^2$  $R[F^2 > 2\sigma(F^2)] = 0.042$  $wR(F^2) = 0.112$ S = 1.015304 reflections 325 parameters

ctions  $> 2\sigma(I)$ 

H-atom parameters constrained  $w = 1/[\sigma^2(F_o^2) + (0.0581P)^2]$ where  $P = (F_o^2 + 2F_c^2)/3$  $(\Delta/\sigma)_{\rm max} = 0.001$  $\Delta \rho_{\rm max} = 0.30 \ {\rm e} \ {\rm \AA}^{-3}$  $\Delta \rho_{\rm min} = -0.35 \text{ e} \text{ Å}^{-3}$ 



#### Figure 1

The molecular structure of (I), showing 50% probability displacement ellipsoids. H atoms have been omitted for clarity.

# Table 1

Selected geometric parameters (Å, °).

Na1-O11	2.3359 (18)	Na1-N41	2.499 (2)
Na1-N11	2.462 (2)	Na1-N51	2.5046 (19)
Na1-N21	2.4805 (18)		
O11-Na1-N11	90.60 (7)	N21-Na1-N41	111.12 (7)
O11-Na1-N21	92.44 (6)	O11-Na1-N51	103.64 (6)
N11-Na1-N21	68.75 (6)	N11-Na1-N51	103.28 (7)
O11-Na1-N41	117.95 (6)	N21-Na1-N51	162.31 (6)
N11-Na1-N41	151.16 (6)	N41-Na1-N51	67.84 (6)

All H atoms were constrained to ideal geometries, with C-H distances of 0.93 and 0.97 Å, and assigned isotropic displacement parameters equal to  $1.2U_{eq}$  of their parent C atoms.

Data collection: SMART (Bruker, 1998); cell refinement: SAINT (Bruker, 1998); data reduction: SAINT and SHELXTL (Bruker, 1998); program(s) used to solve structure: SHELXTL; program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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# References

Addison, A. W., Rao, T. N., Reedijk, J., Van Rijn, J. & Verschoor, G. C. (1984). J. Chem. Soc. Dalton Trans. pp. 1349-1356.

Bell, T. W., Cragg, P. J., Drew, M. G. B., Firestone, A. & Kwok, D.-I. A. (1992). Angew. Chem. Int. Ed. Engl. 31, 345-347.

Bell, T. W., Cragg, P. J., Drew, M. G. B., Firestone, A., Kwok, D.-I. A., Liu, J., Ludwig, R. T. & Papoulis, A. T. (1993). Pure Appl. Chem. 65, 361-366.

Bruker (1998). SMART, SAINT and SHELXTL. Bruker AXS Inc., Madison, Wisconsin, USA.

- Caron, A., Guilhelm, J., Riche, C., Pascard, C., Alpha, B., Lehn, J.-M. & Rodriguez-Ubis, J.-C. (1985). Helv. Chim. Acta, 68, 1577-1582.
- Constable, E., Doyle, M. J., Healy, J. & Raithby, P. R. (1988). J. Chem. Soc. Chem. Commun. pp. 1262-1264.

Echegoyen, L., DeCian, A., Fischer, J. & Lehn, J.-M. (1991). Angew. Chem. Int. Ed. Engl. 30, 838-840.

Fleming, J. S., Mann, K. L. V., Carraz, C.-A., Psillakis, E., Jeffery, J. C., McCleverty, J. A. & Ward, M. D. (1998). Angew. Chem. Int. Ed. Engl. 37, 1279-1281.

- Lehn, J.-M. & DeVains, J. B. R. (1992). *Helv. Chim. Acta*, **75**, 1221–1236. McKee, V., Dorrity, M. R. J., Malone, J. F., Marrs, D. & Nelson, J. (1992). *J. Chem. Soc. Chem. Commun.* pp. 383–386. Paul, R. L., Bell, Z. R., Jeffery, J. C., McCleverty, J. A. & Ward, M. D. (2002).
- Proc. Natl Acad. Sci. USA, 99, 4883-4888.
- Psillakis, E., Jeffery, J. C., McCleverty, J. A. & Ward, M. D. (1997). Chem. Commun. pp. 479-478.
- Qian, C., Wang, B., Xin, Y. & Lin, Y. (1994). J. Chem. Soc. Dalton Trans. pp. 2109–2112.
- Sheldrick, G. M. (1996). SADABS. University of Göttingen, Germany.