

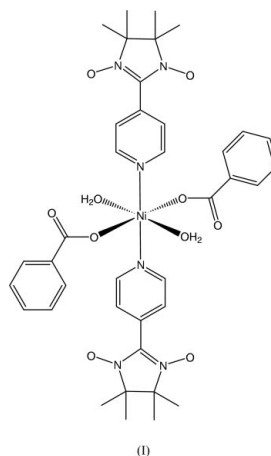
Diaquadibenzoatobis[2-(4-pyridyl- $\kappa$ N)-  
4,4,5,5-tetramethylimidazoline-1-oxyl  
3-oxide]nickel(II)Qihua Zhao,<sup>a</sup> Xiaofeng Wang,<sup>a</sup>  
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## Key indicators

Single-crystal X-ray study  
 $T = 223$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å  
 $R$  factor = 0.049  
 $wR$  factor = 0.123  
Data-to-parameter ratio = 21.4For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.The Ni atom in  $[\text{Ni}(\text{O}_2\text{CCH}_2)_2(\text{PPN})_2(\text{H}_2\text{O})_2]$ , where PPN is 2-(*p*-pyridyl)-4,4,5,5-tetramethylimidazoline-1-oxyl 3-oxide ( $\text{C}_{12}\text{H}_{16}\text{N}_3\text{O}_2$ ), is located on a centre of inversion and exists in a distorted octahedral geometry defined by an  $\text{N}_2\text{O}_4$  donor set. Complex molecules aggregate into chains mediated by hydrogen-bonding interactions.

## Comment

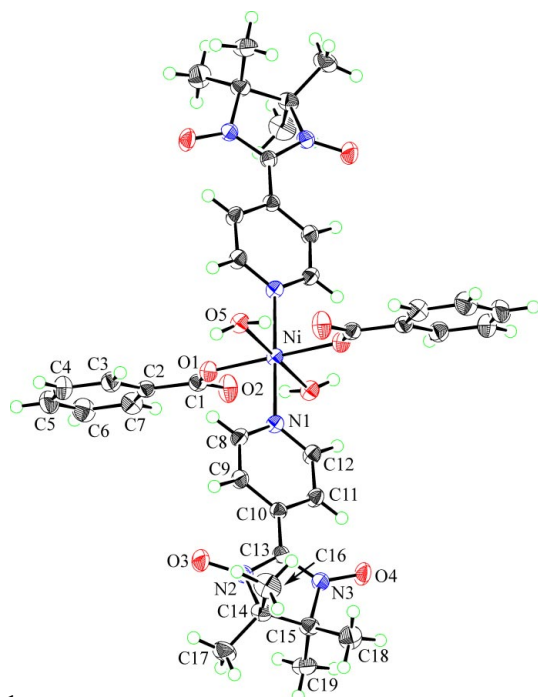
The title compound, (I), is isomorphous with the  $\text{Mn}^{\text{II}}$  species that was investigated for novel magnetic characteristics owing to the presence of the radical ligand, PPN, *i.e.* 2-(*p*-pyridyl)-4,4,5,5-tetramethylimidazoline-1-oxyl 3-oxide (Fettouhi *et al.*, 1999). The Ni atom is situated on a centre of inversion and exists in a distorted octahedral geometry defined by an all-*trans*- $\text{N}_2\text{O}_4$  donor set (Fig. 1 and Table 1).

The benzoate ligand is effectively monodentate as the  $\text{Ni} \cdots \text{O}2$  distance is 3.3110 (14) Å. There are both intra- and intermolecular hydrogen-bonding interactions involving the coordinated water molecules and the O2 atoms. The parameters associated with the intramolecular interaction are  $\text{H} \cdots \text{O}2 = 1.78$  Å,  $\text{O}2 \cdots \text{O}5 = 2.620$  (2) Å and angle at H = 171°. The intermolecular interactions occur between translationally related molecules, along the **a** direction, and are characterized by  $\text{H} \cdots \text{O}2^i = 2.07$  Å,  $\text{O}5 \cdots \text{O}2^i = 2.8468$  (19) Å and the angle at H = 154 [symmetry code: (i) 1 + x, y, z]. These interactions lead to the formation of 12-membered  $[-\text{Ni}-\text{O}-\text{H} \cdots \text{O}-\text{C}-\text{O}-]_2$  rings and the chains that are thereby formed are separated by regions occupied by the PPN ligands (Fig. 2). Links between the chains are facilitated by  $\text{C}11-\text{H} \cdots \text{O}4^{\text{ii}}$  contacts, so that  $\text{H} \cdots \text{O}4^{\text{ii}} = 2.47$  Å,  $\text{C}11 \cdots \text{O}4^{\text{ii}} = 3.260$  (2) Å and the angle subtended at H = 142° [symmetry

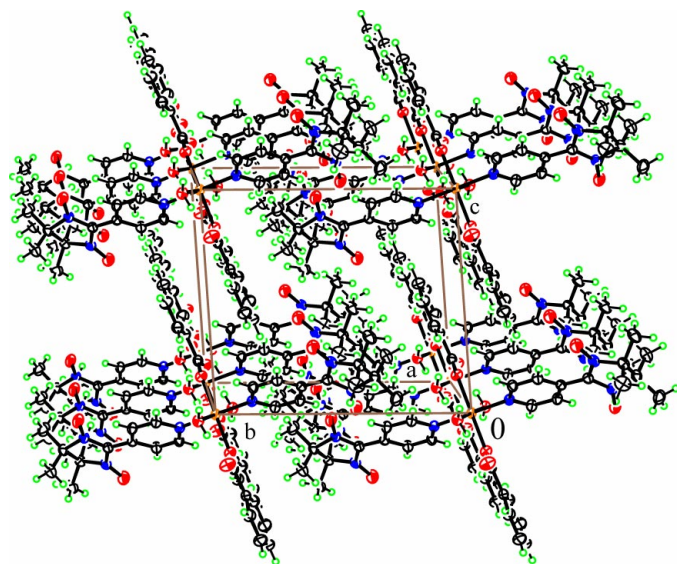
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**Figure 1**  
The molecular structure and crystallographic numbering scheme for (I). The unlabelled half of the molecule is generated by the symmetry operation  $-x, -y, -z$ . Displacement ellipsoids are drawn at the 50% probability level.



**Figure 2**  
The unit-cell contents for (I) (Farrugia, 1997).

code: (ii)  $-x, -y, 1 - z$ ]. Links between the PPN ligands also occur *via*  $C-H \cdots O$  interactions so that  $C17-H \cdots O3^{iii} = 2.55 \text{ \AA}$ ,  $C17 \cdots O3^{iii} = 3.496 (3) \text{ \AA}$  and the angle at H =  $165^\circ$ , and  $C16-H \cdots O4^{iv} = 2.54 \text{ \AA}$ ,  $C16 \cdots O4^{iv} = 3.474 (3) \text{ \AA}$  and the angle at H =  $162^\circ$  [symmetry codes: (iii)  $1 - x, -1 - y, 1 - z$ ; (iv)  $1 - x, -y, 1 - z$ ].

## Experimental

To a solution of  $Ni(O_2CC_5H_5)_2 \cdot 4H_2O$  (0.5 mmol) and benzoic acid (1 mmol) in ethanol (25 ml) was added a solid sample of PPN

(1 mmol). The solution was stirred for 30 min, after which a small quantity of green solid precipitated. After filtration, the clear solution, kept in the dark, was slowly evaporated at room temperature over a period of one month, yielding green crystals suitable for analysis.

## Crystal data

$[Ni(C_7H_5O_2)_2(C_{12}H_{16}N_3O_2)_2 \cdot (H_2O)_2]$   
 $M_r = 805.52$   
 Triclinic,  $P\bar{1}$   
 $a = 6.7001 (3) \text{ \AA}$   
 $b = 11.3354 (6) \text{ \AA}$   
 $c = 12.8740 (6) \text{ \AA}$   
 $\alpha = 86.319 (2)^\circ$   
 $\beta = 87.290 (2)^\circ$   
 $\gamma = 79.486 (2)^\circ$   
 $V = 958.73 (8) \text{ \AA}^3$

$Z = 1$   
 $D_x = 1.395 \text{ Mg m}^{-3}$   
 Mo  $K\alpha$  radiation  
 Cell parameters from 2446 reflections  
 $\theta = 2.4\text{--}28.9^\circ$   
 $\mu = 0.57 \text{ mm}^{-1}$   
 $T = 223 (2) \text{ K}$   
 Irregular block, green  
 $0.28 \times 0.14 \times 0.10 \text{ mm}$

## Data collection

Bruker AXS SMART CCD diffractometer  
 $\omega$  scans  
 Absorption correction: multi-scan (SADABS; Bruker, 2000; Blessing, 1995)  
 $T_{\min} = 0.725, T_{\max} = 0.944$   
 8055 measured reflections

5468 independent reflections  
 4462 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.022$   
 $\theta_{\max} = 30.0^\circ$   
 $h = -8 \rightarrow 9$   
 $k = -15 \rightarrow 14$   
 $l = -16 \rightarrow 18$

## Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.049$   
 $wR(F^2) = 0.123$   
 $S = 1.05$   
 5468 reflections  
 256 parameters

H atoms treated by a mixture of independent and constrained refinement  
 $w = 1/[\sigma^2(F_o^2) + (0.0644P)^2]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.61 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\min} = -0.25 \text{ e \AA}^{-3}$

**Table 1**

Selected geometric parameters ( $\text{\AA}, ^\circ$ ).

Ni—O1	2.0497 (13)	O2—C1	1.256 (2)
Ni—O5	2.1036 (13)	O3—N2	1.272 (2)
Ni—N1	2.1011 (15)	O4—N3	1.274 (2)
O1—C1	1.260 (2)		
O1—Ni—O5	88.52 (5)	O1—Ni—N1 <sup>i</sup>	90.01 (5)
O1—Ni—N1	89.99 (5)	O5—Ni—N1	93.10 (5)
O1—Ni—O5 <sup>i</sup>	91.48 (5)	O5—Ni—N1 <sup>i</sup>	86.90 (5)

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

The carbon-bound H atoms were included in the riding-model approximation, with C—H distances of  $0.94 \text{ \AA}$  ( $0.97 \text{ \AA}$  for methyl),  $U_{\text{iso}}(\text{phenyl-H}) = 1.2U_{\text{eq}}(\text{C})$  and  $U_{\text{iso}}(\text{methyl-H}) = 1.5U_{\text{eq}}(\text{C})$ . The water H atoms were located from a difference map and refined, subject to  $O-H$   $0.85 (1) \text{ \AA}$ ,  $H \cdots H$   $1.39 (1) \text{ \AA}$  and  $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$ .

Data collection: SMART (Bruker, 2000); cell refinement: SAINT (Bruker, 2000); data reduction: SAINT and SHELXTL (Bruker, 2000); program(s) used to solve structure: PATTY in DIRDIF92 (Beurskens *et al.*, 1992); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPII (Johnson, 1976); software used to prepare material for publication: SHELXTL.

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