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Key indicators

Single-crystal X-ray study

T = 160 K

Mean $\sigma(\text{C}-\text{C}) = 0.003 \text{ \AA}$

R factor = 0.026

wR factor = 0.068

Data-to-parameter ratio = 12.8

For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.Diaquabis(*N,N*-diethylnicotinamide)bis(4-nitrobenzoato)zinc(II)

The title compound, $[\text{Zn}(\text{C}_7\text{H}_4\text{NO}_4)_2(\text{C}_{10}\text{H}_{14}\text{N}_2\text{O})_2(\text{H}_2\text{O})_2]$, crystallizes as centrosymmetric mononuclear molecules with octahedrally coordinated zinc, all pairs of equivalent ligands being mutually *trans*. Intramolecular hydrogen bonding links each aqua ligand with an uncoordinated carboxylate O atom, and intermolecular hydrogen bonding between aqua ligands and the carbonyl O atoms of *N,N*-diethylnicotinamide ligands link the molecules together into chains.

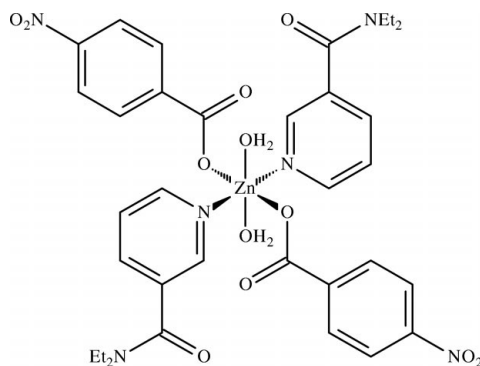
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Comment

The title compound, (I), was prepared as part of a series of studies of the coordination chemistry of the ligand *N,N*-diethylnicotinamide (DENA). It has a molecular structure (Fig. 1) very similar to that of the corresponding complex in which 2-hydroxybenzoate (salicylate) replaces 4-nitrobenzoate, reported in a parallel publication (Necefoglu *et al.*, 2001). Coordination of zinc is close to regular octahedral, with all pairs of equivalent ligands mutually *trans* across a crystallographic inversion centre.



(I)

Each aqua ligand is involved as donor in two hydrogen bonds. One of these is intramolecular, to an uncoordinated carboxylate O atom, lengthening the $\text{C}=\text{O}$ nominal double bond to almost the same as $\text{C}-\text{O}$ for the coordinated O atom. The other is intermolecular, linking the molecules together in chains along the *a* axis.

Experimental

The title compound was prepared by the reaction of 4.34 g (0.01 mol) of $[\text{Zn}(\text{OH}_2)_2(4\text{-O}_2\text{NC}_6\text{H}_4\text{COO})_2]$ and 2.56 g (0.02 mol) of diethylnicotinamide in 100 ml water. The solution was filtered and set aside for crystallization at ambient temperature for a few days. Suitable colourless crystals were obtained.

Crystal data

[Zn(C₇H₄NO₄)₂(C₁₀H₁₄N₂O)₂·(H₂O)₂]
M_r = 790.09
 Triclinic, P1̄
a = 7.211 (3) Å
b = 8.588 (4) Å
c = 16.515 (7) Å
 α = 96.00 (2)°
 β = 100.79 (2)°
 γ = 112.09 (3)°
V = 913.6 (7) Å³

Z = 1
D_x = 1.436 Mg m⁻³
 Mo Kα radiation
 Cell parameters from 36 reflections
 θ = 10.7–12.5°
 μ = 0.74 mm⁻¹
T = 160 (2) K
 Block, colourless
 0.42 × 0.27 × 0.19 mm

Data collection

Stoe–Siemens four-circle diffractometer
 ω/θ scans with on-line profile fitting (Clegg, 1981)
 Absorption correction: ψ scan (XPREP; Sheldrick, 1997)
T_{min} = 0.821, *T_{max}* = 0.959
 6298 measured reflections
 3214 independent reflections

2889 reflections with *I* > 2σ(*I*)
R_{int} = 0.020
 θ_{max} = 25.0°
h = -8 → 8
k = -10 → 10
l = -19 → 19
 5 standard reflections
 frequency: 60 min
 intensity decay: 1%

Refinement

Refinement on *F*²
R [*F*² > 2σ(*F*²)] = 0.026
wR (*F*²) = 0.068
S = 1.10
 3214 reflections
 251 parameters
 H atoms treated by a mixture of independent and constrained refinement

w = 1/[σ²(*F_o*²) + (0.0275*P*)² + 0.4095*P*]
 where *P* = (*F_o*² + 2*F_c*²)/3
 (Δ/σ)_{max} < 0.001
 Δρ_{max} = 0.33 e Å⁻³
 Δρ_{min} = -0.26 e Å⁻³

Table 1 Selected geometric parameters (Å, °).

Zn—O1	2.1373 (15)	C11—O3	1.264 (2)
Zn—O3	2.1149 (14)	C11—O4	1.255 (2)
Zn—N1	2.1527 (16)		
O3—Zn—O1	87.54 (6)	O1—Zn—N1	86.71 (6)
O3—Zn—N1	88.54 (6)	C11—O3—Zn	125.41 (11)

Table 2 Hydrogen-bonding geometry (Å, °).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
O1—H1A...O4 ⁱ	0.86 (3)	1.82 (3)	2.659 (2)	164 (3)
O1—H1B...O2 ⁱⁱ	0.81 (3)	1.95 (3)	2.765 (2)	173 (2)

Symmetry codes: (i) 1 - *x*, 1 - *y*, 1 - *z*; (ii) 2 - *x*, 1 - *y*, 1 - *z*.

H atoms were placed geometrically and refined with a riding model (including free rotation about C—C bonds), and with *U*_{iso} constrained to be 1.2 (1.5 for methyl groups) times *U*_{eq} of the carrier atom, except for aqua ligand H atoms, which were refined freely with individual *U*_{iso}.

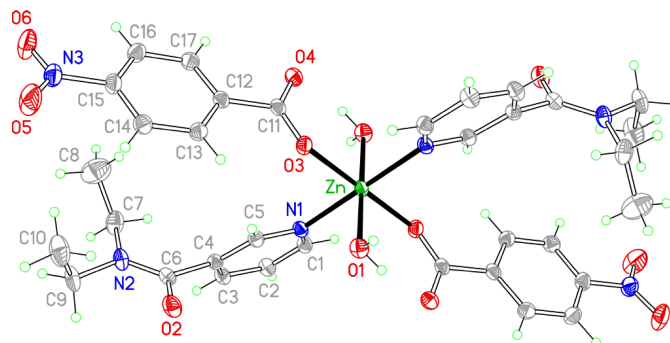


Figure 1 The molecular structure of (I) with atom labels and 50% probability ellipsoids for non-H atoms.

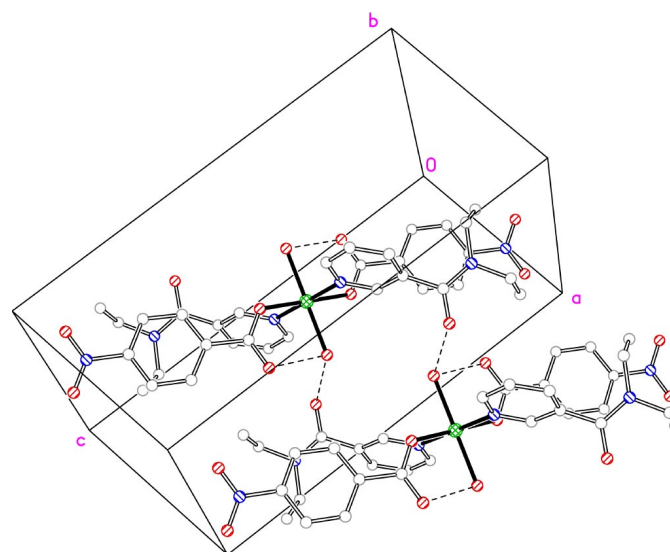


Figure 2 View of two molecules and the unit-cell outline, showing intramolecular and intermolecular hydrogen bonding as dashed lines.

Data collection: *DIF4* (Stoe & Cie, 1988); cell refinement: *DIF4*; data reduction: local programs; program(s) used to solve structure: *SHELXTL* (Sheldrick, 1997); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL* and local programs.

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References

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