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Key indicators

Single-crystal X-ray study
 $T = 292$ K
Mean $\sigma(C-C) = 0.011$ Å
 R factor = 0.071
 wR factor = 0.235
Data-to-parameter ratio = 14.9For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>. μ -Benzene-1,4-dicarboxylato-bis[aquachloro-
(2-phenyl-1*H*-1,3,7,8-tetraazacyclopenta[*l*]-
phenanthrene)zinc(II)]

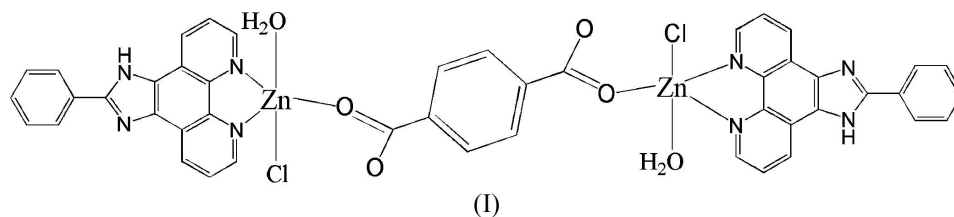
In the title compound, $[Zn_2(C_8H_4O_4)Cl_2(C_{19}H_{12}N_4)_2(H_2O)_2]$ or $[Zn_2(BDC)Cl_2(L)_2(H_2O)_2]$ (where BDC is benzene-1,4-dicarboxylate and *L* is 2-phenyl-1*H*-1,3,7,8-tetraaza-cyclopenta[*l*]phenanthrene), the two Zn^{II} atoms are bridged by the BDC ligand to form a dinuclear compound in which each Zn^{II} atom is five-coordinated by two N atoms from one *L* ligand, one chloride ion, and two O atoms, one from the BDC and one from the water molecule. The molecule is centrosymmetric. Neighbouring dinuclear complexes interact through π - π contacts to form stacks, leading to a two-dimensional supramolecular structure.

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Comment

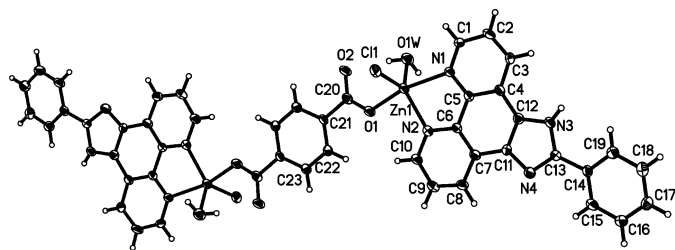
The synthesis and design of supramolecular complexes exhibiting novel structures and properties has provided exciting new prospects for chemists (Eddaoudi *et al.*, 2001). A common and successful approach to the preparation of supramolecular frameworks is *via* direct chemical combination of functional inorganic and organic components. 1,10-Phenanthroline (phen) and its derivatives are one type of important organic ligand widely used in the construction of metal-organic complexes. 2-Phenyl-1*H*-1,3,7,8-tetraazacyclopenta[*l*]phenanthrene (*L*) is one such phen derivative and, to date, the chemistry of supramolecular architectures based on *L* has received little attention (Collins *et al.*, 1998). We present here a new compound $[Zn_2(BDC)Cl_2(L)_2(H_2O)_2]$, (I), incorporating the *L* ligand (BDC is benzene-1,4-dicarboxylate).



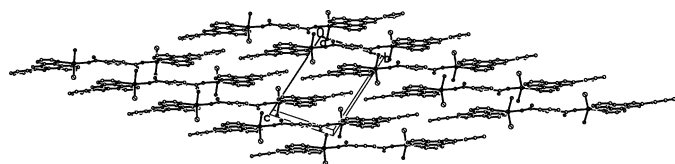
Selected bond lengths and angles for (I) are given in Table 1. There is a centre of symmetry at the centre of the BDC ligand. In compound (I), the Zn^{II} atom is five-coordinated by two N atoms from one *L* ligand, one chloride ion, and two O atoms from one BDC and one water molecule (Fig. 1). The two Zn^{II} atoms are bridged by the BDC ligand to form a dinuclear compound.

The water molecule, NH group, chloride ion and carboxylate O atoms form hydrogen bonds (Table 2).

Neighbouring dinuclear complexes interact through π - π contacts, leading to a two-dimensional supramolecular structure (Fig. 2). There are two types of π - π interactions: one


Figure 1

The molecular structure of (I), showing the atomic numbering scheme and displacement ellipsoids drawn at the 50% probability level. Unlabelled atoms are related to labelled atoms by $(-x, 1 - y, 1 - z)$.


Figure 2

View of the two-dimensional supramolecular structure of (I), generated by intermolecular π - π interactions.

exists between two *L* ligands (*ca* 3.442 Å), while the other exists between the *L* and BDC ligands (*ca* 3.709 Å). It is clear that aromatic-aromatic interactions help to stabilize this and related architectures (Noveron *et al.*, 2002).

Experimental

The *L* ligand was synthesized according to the reported method (Collins *et al.*, 1998). Complex (I) was self-assembled from *L*, $\text{ZnCl}_2 \cdot \text{H}_2\text{O}$ and H_2BDC . A methanol solution (10 ml) of *L* (0.5 mmol) was added slowly to an aqueous solution (10 ml) of $\text{ZnCl}_2 \cdot \text{H}_2\text{O}$ (0.5 mmol) and H_2BDC (1 mmol) with stirring. The resulting solution was filtered and the filtrate allowed to stand in air at room temperature for several days, yielding pale-yellow crystals of (I) (68% yield based on Zn).

Crystal data

$[\text{Zn}_2(\text{C}_8\text{H}_4\text{O}_4)\text{Cl}_2(\text{C}_{19}\text{H}_{12}\text{N}_4)_2 \cdot (\text{H}_2\text{O})_2]$	$\gamma = 100.49 (3)^\circ$
$M_r = 994.44$	$V = 988.0 (4) \text{ \AA}^3$
Triclinic, $P\bar{1}$	$Z = 1$
$a = 9.0702 (18) \text{ \AA}$	$D_x = 1.671 \text{ Mg m}^{-3}$
$b = 9.1815 (18) \text{ \AA}$	Mo $K\alpha$ radiation
$c = 12.642 (3) \text{ \AA}$	$\mu = 1.42 \text{ mm}^{-1}$
$\alpha = 103.54 (3)^\circ$	$T = 292 (2) \text{ K}$
$\beta = 98.01 (3)^\circ$	Block, pale yellow
	$0.34 \times 0.30 \times 0.21 \text{ mm}$

Data collection

Rigaku R-AXIS RAPID diffractometer	9673 measured reflections
ω scan	4441 independent reflections
Absorption correction: multi-scan (ABSCOR; Higashi, 1995)	2309 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.651, T_{\max} = 0.744$	$R_{\text{int}} = 0.093$
	$\theta_{\max} = 27.5^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.071$
 $wR(F^2) = 0.235$
 $S = 0.98$
 4441 reflections
 298 parameters

H atoms treated by a mixture of independent and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.1295P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 1.08 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.85 \text{ e \AA}^{-3}$

Table 1

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

N1–Zn1	2.157 (5)	O1W–Zn1	2.084 (6)
N2–Zn1	2.088 (6)	Zn1–Cl1	2.275 (2)
O1–Zn1	2.051 (5)		
O1–Zn1–O1W	87.8 (2)	N2–Zn1–N1	78.2 (2)
O1–Zn1–N2	90.7 (2)	O1–Zn1–Cl1	99.75 (17)
O1W–Zn1–N2	127.5 (2)	O1W–Zn1–Cl1	117.30 (18)
O1–Zn1–N1	163.8 (2)	N2–Zn1–Cl1	114.64 (18)
O1W–Zn1–N1	89.7 (2)	N1–Zn1–Cl1	95.60 (17)

Table 2

Hydrogen-bond geometry ($\text{\AA}, ^\circ$).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
O1W–HW11 \cdots O2	0.84 (6)	1.96 (4)	2.769 (7)	160 (9)
N3–H3B \cdots Cl1 ⁱ	0.81 (11)	2.57 (11)	3.264 (6)	144 (9)

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

All H atoms on C atoms were positioned geometrically and refined as riding, with $C-H = 0.93 \text{ \AA}$ and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. Aqua H atoms were located in difference Fourier maps and refined with $O-H = 0.85 \text{ \AA}$ and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{O})$. The N-bound H atom was located in a difference Fourier map and refined freely, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N})$. The highest residual density peak was located 0.87 \AA from atom H2.

Data collection: *PROCESS-AUTO* (Rigaku, 1998); cell refinement: *PROCESS-AUTO*; data reduction: *PROCESS-AUTO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL-Plus* (Siemens, 1990); software used to prepare material for publication: *SHELXL97*.

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