

Tris(4,4',5,5'-tetramethyl-2,2'-biimidazole)-zinc(II) dinitrate

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Key indicators

Single-crystal X-ray study

T = 293 K

Mean $\sigma(\text{C}-\text{C}) = 0.006 \text{ \AA}$

R factor = 0.043

wR factor = 0.111

Data-to-parameter ratio = 16.1

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

A new six-coordinate monomeric Zn^{II} complex, $[\text{Zn}(\text{L})_3](\text{NO}_3)_2$, where L is 4,4',5,5'-tetramethyl-2,2'-biimidazole ($\text{C}_{10}\text{H}_{14}\text{N}_4$), has been synthesized and characterized by X-ray diffraction techniques. The Zn^{II} atom, on a twofold axis, has a ZnN_6 distorted octahedral coordination environment. 4,4',5,5'-Tetramethyl-2,2'-biimidazole is bonded as a chelating ligand through its unprotonated N atoms. These discrete monomeric units possess excellent hydrogen-bonding sites which further self-assemble into a three-dimensional supramolecular architecture through intermolecular $\text{N}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds.

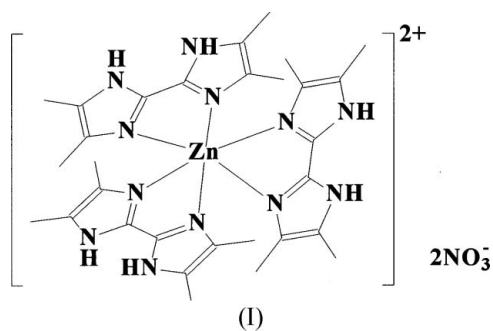
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Comment

The unique strength, directionality and complementarity of non-covalent interactions, such as hydrogen bonding and coordination bonding, play an important role in the creation of a variety of molecular architectures (Tadokoro *et al.*, 1999; Lehn, 1995). The building blocks possessing such non-covalent interaction sites can produce one-, two- and three-dimensional molecular arrangements with long-range order (Subramanian & Zaworotko, 1994; Ermer, 1988). We present here a new three-dimensional hydrogen-bonding network with discrete monomeric Zn^{II} building blocks, $[\text{Zn}(\text{L})_3](\text{NO}_3)_2$, (I), where L is 4,4',5,5'-tetramethyl-2,2'-biimidazole.



X-ray structure analysis reveals that (I) is a monomeric Zn^{II} complex (Fig. 1). The coordination geometry around the Zn^{II} atom, which lies on a twofold rotation axis, is octahedral, with $\text{Zn}-\text{N}$ distances in the range 2.149 (3)–2.208 (2) Å. The 4,4',5,5'-tetramethyl-2,2'-biimidazole ligand is bonded as a chelating ligand through its unprotonated N atoms. A striking structural feature of (I) is the formation of a three-dimensional supramolecular architecture through intermolecular $\text{N}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{O}$ hydrogen-bonding interactions (Table 1).

Experimental

Complex (I) was prepared by reacting an acetonitrile solution (15 ml) of zinc(II) nitrate with 4,4',5,5'-tetramethyl-2,2'-biimidazole in a 1:3 ratio. Red block-shaped crystals were recovered by filtration and dried in air. The yield was ca 78% (based on Zn). IR (KBr, cm⁻¹): 3232 (*m*), 2980 (*m*), 1696 (*s*), 1517 (*s*), 1456 (*m*), 1200 (*m*), 841 (*m*), 751 (*w*), 624 (*w*) cm⁻¹.

Crystal data

[Zn(C₁₀H₁₄N₄)₃](NO₃)₂
M_r = 760.15
 Hexagonal, *P*6₄
a = 13.900 (2) Å
c = 16.809 (3) Å
V = 2812.7 (8) Å³
Z = 3
D_x = 1.346 Mg m⁻³

Mo *K*α radiation
 Cell parameters from 6512 reflections
 θ = 2.9–24.1°
 μ = 0.72 mm⁻¹
T = 293 (2) K
 Block, red
 0.20 × 0.20 × 0.15 mm

Data collection

Bruker SMART CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (*SADABS*; Sheldrick, 1998)
T_{min} = 0.868, *T_{max}* = 0.898
 17035 measured reflections

4093 independent reflections
 3093 reflections with *I* > 2σ(*I*)
R_{int} = 0.116
 θ_{max} = 28.3°
h = -12 → 18
k = -18 → 15
l = -17 → 21

Refinement

Refinement on *F*²
R[*F*² > 2σ(*F*²)] = 0.043
wR(*F*²) = 0.111
S = 0.96
 4093 reflections
 254 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0565P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{max} < 0.001$
 $\Delta\rho_{max} = 0.68 \text{ e \AA}^{-3}$
 $\Delta\rho_{min} = -0.31 \text{ e \AA}^{-3}$
 Absolute structure: Flack (1983),
 1683 Friedel pairs
 Flack parameter = -0.027 (13)

Table 1

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>
N3—H3A...O1 ⁱ	0.86	2.01	2.852 (4)	167
N4—H4A...O3 ^l	0.86	2.01	2.818 (5)	156
N6—H6A...O2 ⁱⁱ	0.86	2.03	2.849 (5)	157
N6—H6A...O3 ⁱⁱ	0.86	2.60	3.276 (5)	136
C9—H9A...O3 ⁱⁱⁱ	0.96	2.47	3.403 (7)	163
C9—H9B...O1 ^{iv}	0.96	2.51	3.431 (7)	162
C10—H10C...O2 ^v	0.96	2.46	3.235 (6)	138

Symmetry codes: (i) *x* - *y*, *x*, *z* - $\frac{1}{3}$; (ii) *y*, 1 - *x* + *y*, $\frac{1}{3}$ + *z*; (iii) 1 - *x*, 1 - *y*, *z*; (iv) 1 - *x* + *y*, 2 - *x*, *z* - $\frac{1}{3}$; (v) *x* - 1, *y*, *z*.

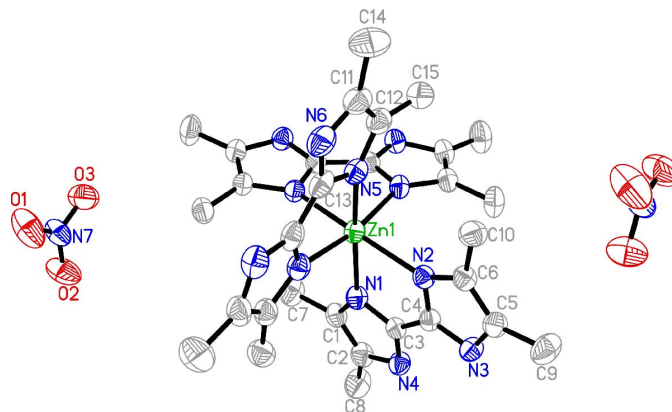


Figure 1
 A view of the title compound, with displacement ellipsoids drawn at the 30% probability level. H atoms have been omitted. Unlabeled atoms are related to labeled atoms by -*x*, -*y*, *z*.

All H atoms were placed in calculated positions and allowed to ride on their parent atoms [*C*—*H* = 0.93 Å and *U_{iso}*(*H*) = 1.2*U_{eq}*(*C*)].

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINTE* (Bruker, 1998); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997*a*); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997*a*); molecular graphics: *SHELXTL* (Sheldrick, 1997*b*); software used to prepare material for publication: *SHELXTL*.

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