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## Jing-Min Shi,* Zhe Liu, Jian-Jun Lu and Lian-Dong Liu

Department of Chemistry, Shandong Normal University, Jinan 250014, People's Republic of China

Correspondence e-mail:
shijingmin@beelink.com

## Key indicators

Single-crystal X-ray study
$T=293 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.008 \AA$
$R$ factor $=0.030$
$w R$ factor $=0.075$
Data-to-parameter ratio $=18.1$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## Diiodidobis(4-methylpyridine $\boldsymbol{N}$-oxide- $\boldsymbol{\kappa} \mathrm{O}$ )zinc(II)

The title mononuclear complex, $\left[\mathrm{Zn}(\mathrm{I})_{2}\left(\mathrm{C}_{6} \mathrm{H}_{7} \mathrm{NO}\right)_{2}\right]$, lies on a special position of site symmetry 2 . The Zn atom is coordinated by two I and two O atoms in a tetrahedral geometry. There is a $\pi-\pi$ stacking interaction of the 4-methylpyridine $N$-oxide units.

## Comment

In the title molecular complex, (I), (Fig. 1), the Zn atom is coordinated by two I atoms and the O atoms from two 4-methylpyridine $N$-oxide ligands. Atom Zn 1 lies on a twofold axis in a distorted tetrahedral environment. The 4-methylpyridine $N$-oxide units are stacked over each other at a distance of about $3.66 \AA$. Such $\pi-\pi$ stacking causes the molecules to pack as columns along the $b$ axis.


## Experimental

4-Methylpyridin $N$-dioxide ( $0.0625 \mathrm{~g}, 0.573 \mathrm{mmol}$ ) was added to an aqueous solution ( 10 ml ) containing $\mathrm{Zn}\left(\mathrm{ClO}_{4}\right)_{2} \cdot 6 \mathrm{H}_{2} \mathrm{O}(0.1035 \mathrm{~g}$, $0.278 \mathrm{mmol})$ and $\mathrm{NaI}(0.0870 \mathrm{~g}, 0.580 \mathrm{mmol})$. Colourless crystals of (I) were obtained after the solution was allowed to stand at room temperature for three weeks.

## Crystal data

| $\left[\mathrm{Zn}(\mathrm{I})_{2}\left(\mathrm{C}_{6} \mathrm{H}_{7} \mathrm{NO}\right)_{2}\right]$ | $D_{x}=2.052 \mathrm{Mg} \mathrm{m}^{-3}$ |
| :---: | :---: |
| $M_{r}=537.42$ | Mo $K \alpha$ radiation |
| Monoclinic, C2/c | Cell parameters from 1828 |
| $a=19.385$ (5) A | reflections |
| $b=7.5134$ (19) $\AA$ | $\theta=2.6-26.0^{\circ}$ |
| $c=14.859$ (4) $\AA$ | $\mu=4.96 \mathrm{~mm}^{-1}$ |
| $\beta=126.487$ (3) ${ }^{\circ}$ | $T=293$ (2) K |
| $V=1739.9$ (8) $\AA^{3}$ | Prism, colourless |
| $Z=4$ | $0.21 \times 0.12 \times 0.09 \mathrm{~mm}$ |
| Data collection |  |
| Bruker SMART area-detector diffractometer | 1597 independent reflections 1371 reflections with $I>2 \sigma(I)$ |
| $\varphi$ and $\omega$ scans | $R_{\text {int }}=0.029$ |
| Absorption correction: multi-scan | $\theta_{\text {max }}=25.5^{\circ}$ |
| (SADABS; Sheldrick, 1996) | $h=-23 \rightarrow 20$ |
| $T_{\text {min }}=0.406, T_{\text {max }}=0.637$ | $k=-9 \rightarrow 8$ |
| 4279 measured reflections | $l=-17 \rightarrow 17$ |

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## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.030$
$w R\left(F^{2}\right)=0.075$
$S=1.04$
1597 reflections
88 parameters

H -atom parameters constrained
$w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.037 P)^{2}\right]$
where $P=\left(F_{\mathrm{o}}{ }^{2}+2 F_{\mathrm{c}}{ }^{2}\right) / 3$
$(\Delta / \sigma)_{\max }=0.001$
$\Delta \rho_{\text {max }}=0.59 \mathrm{e}^{-3}$
$\Delta \rho_{\text {min }}=-0.49 \mathrm{e}^{-3}$

Table 1
Selected geometric parameters $\left(\AA^{\circ},{ }^{\circ}\right)$.

| $\mathrm{Zn} 1-\mathrm{O} 1$ | $1.992(3)$ | $\mathrm{Zn} 1-\mathrm{I} 1$ | $2.5399(6)$ |
| :--- | :--- | :--- | :--- |
|  |  |  |  |
| $\mathrm{O} 1-\mathrm{Zn} 1-\mathrm{O} 1^{\mathrm{i}}$ | $102.7(2)$ | $\mathrm{O} 1-\mathrm{Zn} 1-\mathrm{I} 1^{\mathrm{i}}$ | $108.00(9)$ |
| $\mathrm{O} 1-\mathrm{Zn} 1-\mathrm{I} 1$ | $108.34(9)$ | $\mathrm{I} 1-\mathrm{Zn} 1-1^{\mathrm{i}}$ | $120.11(3)$ |

Symmetry code: (i) $-x+1, y,-z+\frac{3}{2}$.

All H atoms were placed in calculated positions and included in the final cycles of refinement using a riding model $(\mathrm{C}-\mathrm{H}=0.93 \AA$ for aromatic H atoms and $0.96 \AA$ for methyl H atoms); $U_{\text {iso }}(\mathrm{H})=$ $1.2 U_{\mathrm{eq}}(\mathrm{C})$ for aromatic H atoms and $1.5 U_{\mathrm{eq}}(\mathrm{C})$ for methyl H atoms.

Data collection: SMART (Bruker, 1997); cell refinement: SAINT (Bruker, 1997); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Sheldrick, 2001); software used to prepare material for publication: SHELXTL.

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## Figure 1

A view of complex (I), with the atom-numbering scheme, showing $30 \%$ probability displacement ellipsoids [symmetry code: (i) $1-x, y, \frac{3}{2}-z$ ]. H atoms are shown as small spheres of arbitary radii.

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