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Key indicators

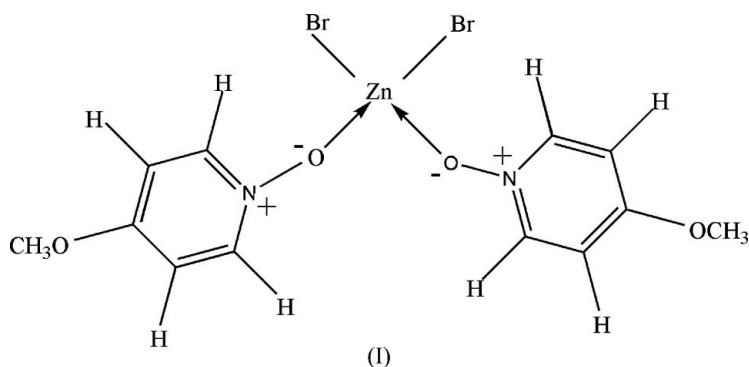
Single-crystal X-ray study
 $T = 298\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.007\text{ \AA}$
 R factor = 0.041
 wR factor = 0.102
Data-to-parameter ratio = 14.8For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Dibromobis(4-methoxypyridine *N*-oxide- κO)zinc(II)

In the title mononuclear complex, $[\text{ZnBr}_2(\text{C}_6\text{H}_7\text{NO}_2)_2]$, the Zn^{II} atom is coordinated by two Br and two O atoms in a distorted tetrahedral geometry. There are π - π stacking interactions between the 4-methoxypyridine *N*-oxide units.

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Comment

Pyridine *N*-oxide and its derivatives usually act as bridging and terminal ligands, and a large number of complexes have been prepared with those ligands (Watson, 1969; Shi, 2005). We report here the synthesis and the structure of the title Zn^{II} complex, (I) (Fig. 1).



The Zn^{II} atom assumes a distorted ZnBr_2O_2 tetrahedral coordination geometry (Table 1), where the dihedral angle between the $\text{Zn}/\text{Br}/\text{Br}$ and $\text{Zn}/\text{O}/\text{O}$ planes is $88.18(10)^\circ$. There are significant π - π stacking interactions between neighbouring pyridine rings; the relevant distances are $\text{Cg}1 \cdots \text{Cg}1^i = 4.335(3)\text{ \AA}$ and $\text{Cg}1 \cdots 1^i_{\text{perp}} = 3.563\text{ \AA}$, and $\text{Cg}2 \cdots \text{Cg}2^{\text{ii}} =$

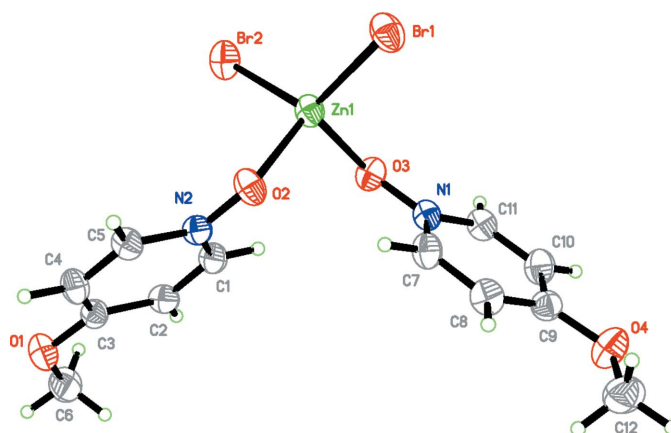


Figure 1
View of complex (I), showing the atom-numbering scheme and 30% probability displacement ellipsoids.

3.590 (3) Å and $Cg2 \cdot \cdot 2_{\text{perp}}^{\text{ii}} = 3.559$ Å [symmetry codes: (i) $1 - x, -y, 1 - z$; (ii) $2 - x, 1 - y, -z$; $Cg1$ and $Cg2$ are the centroids of the N1/C7–C11 and N2/C1–C5 rings, respectively; $CgI \cdot \cdot J_{\text{perp}}$ is the perpendicular distance from CgI to ring J].

Experimental

4-Methoxypyridine *N*-oxide (0.1025 g, 0.819 mmol, in 15 ml H₂O) was added to an aqueous solution (10 ml) containing Zn(ClO₄)₂·6H₂O (0.1573 g, 0.422 mmol) and NaBr (0.0873 g, 0.848 mmol). The resulting solution was stirred for a few minutes. Colourless single crystals were obtained after the solution had been allowed to stand at room temperature for three weeks.

Crystal data

| | |
|--|---------------------------------------|
| [ZnBr ₂ (C ₆ H ₇ NO ₂) ₂] | $Z = 2$ |
| $M_r = 475.44$ | $D_x = 1.933$ Mg m ⁻³ |
| Triclinic, $P\bar{1}$ | Mo $K\alpha$ radiation |
| $a = 7.1783$ (18) Å | Cell parameters from 1347 reflections |
| $b = 8.501$ (2) Å | $\theta = 2.5$ – 23.8° |
| $c = 14.108$ (3) Å | $\mu = 6.41$ mm ⁻¹ |
| $\alpha = 78.294$ (3)° | $T = 298$ (2) K |
| $\beta = 76.293$ (3)° | Prism, colourless |
| $\gamma = 83.317$ (4)° | $0.23 \times 0.15 \times 0.10$ mm |
| $V = 816.9$ (3) Å ³ | |

Data collection

| | |
|---|--|
| Bruker SMART CCD area-detector diffractometer | 2846 independent reflections |
| φ and ω scans | 2115 reflections with $I > 2\sigma(I)$ |
| Absorption correction: multi-scan (SADABS; Sheldrick, 1996) | $R_{\text{int}} = 0.021$ |
| $T_{\text{min}} = 0.298$, $T_{\text{max}} = 0.527$ | $\theta_{\text{max}} = 25.1^\circ$ |
| 4287 measured reflections | $h = -8 \rightarrow 8$ |
| | $k = -10 \rightarrow 6$ |
| | $l = -16 \rightarrow 16$ |

Refinement

| | |
|---------------------------------|---|
| Refinement on F^2 | H-atom parameters constrained |
| $R[F^2 > 2\sigma(F^2)] = 0.041$ | $w = 1/[\sigma^2(F_o^2) + (0.0532P)^2]$ |
| $wR(F^2) = 0.102$ | where $P = (F_o^2 + 2F_c^2)/3$ |
| $S = 0.99$ | $(\Delta/\sigma)_{\text{max}} < 0.001$ |
| 2846 reflections | $\Delta\rho_{\text{max}} = 0.62$ e Å ⁻³ |
| 192 parameters | $\Delta\rho_{\text{min}} = -0.41$ e Å ⁻³ |

Table 1

Selected geometric parameters (Å, °).

| | | | |
|------------|-------------|-------------|-------------|
| Zn1–O3 | 1.980 (3) | Zn1–Br1 | 2.3267 (9) |
| Zn1–O2 | 2.022 (3) | Zn1–Br2 | 2.3565 (10) |
| O3–Zn1–O2 | 104.29 (14) | O3–Zn1–Br2 | 104.88 (10) |
| O3–Zn1–Br1 | 110.95 (10) | O2–Zn1–Br2 | 106.35 (10) |
| O2–Zn1–Br1 | 106.89 (10) | Br1–Zn1–Br2 | 122.08 (3) |

All H atoms were placed in calculated positions and refined using a riding model, with C–H = 0.93 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ for pyridine H atoms, and C–H = 0.96 Å and $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$ 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 (Bruker, 2000); software used to prepare material for publication: SHELXTL.

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