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## An-Yu Zhou, Zuo-Xiang Wang,\* Yan Lan, Lu-Tong Yuan and Chun-Yi Liu

Department of Chemistry and Chemical Engineering, Southeast University, Nanjing 210096, People's Republic of China

Correspondence e-mail: wangzx0908@yahoo.com.cn

#### **Key indicators**

Single-crystal X-ray study T = 293 K Mean  $\sigma$ (C–C) = 0.005 Å R factor = 0.058 wR factor = 0.129 Data-to-parameter ratio = 14.3

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

# Bis[4-(4-methoxyphenyl)-3-methyl-5-(2-pyridyl)-4*H*-1,2,4-triazole- $\kappa N^1$ ]bis(salicylato- $\kappa O$ )copper(II) methanol disolvate

In the title complex,  $[Cu(C_7H_5O_3)_2(C_{15}H_{14}N_4O)_2]\cdot 2CH_3OH$ , the  $Cu^{II}$  cation is located on a centre of inversion and exists in a distorted octahedral geometry defined by a  $CuN_2N_2'O_2$  donor set.

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## Comment

The coordination chemistry of substituted 1,2,4-triazoles has received considerable attention in recent years (Bencini *et al.*, 1987; Koningsbruggen *et al.*, 1997; Moliner *et al.*, 1998, 2001; Klingele & Brooker, 2003). This arises, in part, because the ligand donor ability is optimal to give  $Fe^{II}$  spin-crossover complexes, which could be used as molecular-based memory devices, displays and optical switches (Garcia *et al.*, 1997; Kahn & Martinez, 1998). Recently, we have prepared some new substituted 1,2,4-triazole derivatives (Wang *et al.*, 2005; Liu *et al.*, 2005) and their transition metal complexes. Here, we report the crystal structure analysis of the title complex, (I).



The crystal structure of (I) (Fig. 1 and Table 1) contains centrosymmetric  $Cu^{II}$  complexes in which individual Cu centres exist in a  $CuN_2N'_2O_2$  donor set that defines a distorted octahedral geometry. From symmetry, the four N atoms coordinate in a plane around the Cu atom. The O atoms derived from the two carboxylate groups interact more weakly in the axial positions; the Cu1-O1 distance is 2.442 (2) Å.

The N1 atom of the 1,2,4-triazole ring and the carboxylate atom O1 each form a hydrogen bond with the OH atom of the solvent methanol molecule, indicating that this H atom is bifurcated; details are presented in Table 2.

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## Experimental

The title complex was synthesized by the reaction of 3-methyl-4-(p-methoxyphenyl)-5-(2-pyridyl)-1,2,4-triazole (1.20 g, 3.0 mmol) with copper(II) disalicylate (0.51 g, 1.5 mmol) dissolved in methanol (20 ml) at room temperature. Single crystals of (I) suitable for X-ray diffraction were obtained by evaporation of the methanol solution at room temperature.

#### Crystal data

| $[Cu(C_7H_5O_3)_2(C_{15}H_{14}N_4O)_2]$ | $D_x = 1.409 \text{ Mg m}^{-3}$           |
|---|---|
| 2CH <sub>4</sub> O                      | Mo $K\alpha$ radiation                    |
| $M_r = 934.45$                          | Cell parameters from 895                  |
| Monoclinic, $P2_1/n$                    | reflections                               |
| a = 8.643 (3)  Å                        | $\theta = 2.7-25.6^{\circ}$               |
| b = 14.638 (2) Å                        | $\mu = 0.56 \text{ mm}^{-1}$              |
| c = 17.715 (3) Å                        | T = 293 (2) K                             |
| $\beta = 100.579 \ (5)^{\circ}$         | Block, blue                               |
| $V = 2203.3 (9) \text{ Å}^3$            | $0.33 \times 0.24 \times 0.22 \text{ mm}$ |
| Z = 2                                   |   |

#### Data collection

| Bruker SMART APEX CCD area-          | 4265 independent reflections           |
|--------------------------------------|--|
| detector diffractometer              | 2913 reflections with $I > 2\sigma(I)$ |
| $\varphi$ and $\omega$ scans         | $R_{\rm int} = 0.055$                  |
| Absorption correction: multi-scan    | $\theta_{\rm max} = 26.0^{\circ}$      |
| (SADABS; Bruker, 2000)               | $h = -10 \rightarrow 10$               |
| $T_{\min} = 0.849, T_{\max} = 0.881$ | $k = -18 \rightarrow 15$               |
| 11519 measured reflections           | $l = -18 \rightarrow 21$               |

#### Refinement

| Refinement on $F^2$             | H-atom parameters constrained                              |
|---------------------------------|--|
| $R[F^2 > 2\sigma(F^2)] = 0.058$ | $w = 1/[\sigma^2(F_o^2) + (0.0614P)^2]$                    |
| $wR(F^2) = 0.129$               | where $P = (F_0^2 + 2F_c^2)/3$                             |
| S = 1.03                        | $(\Delta/\sigma)_{\rm max} < 0.001$                        |
| 4265 reflections                | $\Delta \rho_{\rm max} = 0.42 \ {\rm e} \ {\rm \AA}^{-3}$  |
| 298 parameters                  | $\Delta \rho_{\rm min} = -0.32 \text{ e } \text{\AA}^{-3}$ |

#### Table 1

#### Selected geometric parameters (Å, °).

| 1.249 (4) |
|-----------|
|           |
| 1.339 (4) |
| 1.339 (4) |
| 1.249 (4) |
|           |
| 123.6 (3) |
| 118.3 (3) |
| 118.2 (3) |
|           |

### Table 2

Hydrogen-bond geometry (Å, °).

| $\overline{D - \mathbf{H} \cdots A}$   | D-H  | $H \cdots A$ | $D \cdots A$ | $D - \mathbf{H} \cdots A$ |
|--|------|--------------|--------------|---------------------------|
| $\begin{array}{c} O5 - H5D \cdots O1^{i} \\ O5 - H5D \cdots N1^{ii} \end{array}$ | 0.89 | 2.29         | 2.792 (3)    | 116                       |
|  | 0.89 | 2.47         | 3.302 (4)    | 156                       |

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



#### Figure 1

The molecular structure of the cation in (I), showing the atomic labelling. Displacement ellipsoids are shown at the 50% probability level. H atoms have been omitted for clarity. Unlabelled atoms and those with symmetry code (i) are at (1 - x, 1 - y, 2 - z).

All H atoms were allowed to ride on their parent atoms at distances of 0.93 (aromatic H), 0.96 (methyl H), 0.89 (O–H methanol) and 0.96 Å (O–H phenol), and with  $U_{\rm iso}$ (H) values of  $1.2U_{\rm eq}$ (parent atom) for aromatic H and methanol O–H, and  $1.5U_{\rm eq}$ (parent atom) for methyl H and phenol O–H.

Data collection: *SMART* (Bruker, 2000); cell refinement: *SMART*; data reduction: *SAINT* (Bruker, 2000); program(s) used to solve structure: *SHELXTL* (Bruker, 2000); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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