Acta Crystallographica Section C Crystal Structure Communications

ISSN 0108-2701

# Twinned dichlorobis(2,6-dichloropyrazine-N<sup>4</sup>)zinc(II)

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Received 20 June 2001 Accepted 17 July 2001

For crystals of the title compound,  $[ZnCl_2(C_4H_2Cl_2N_2)_2]$ , metric considerations suggest a monoclinic unit cell. However, the symmetry of the diffraction pattern shows the crystals to be triclinic, with very similar lattice constants *a* and *b*, and *a* and *β*. All crystals examined were twinned. The discrepancy between the reliability indices for merging in the monoclinic lattice metric symmetry and the triclinic crystal symmetry constitutes an indicator for the volume ratio between the components. The asymmetric unit contains three independent molecules. At the molecular level, the compound shows a distorted tetrahedral coordination around one Zn atom.

### Comment

Our interest in intermolecular interactions recently prompted us to synthesize and study the first one-dimensional chain

polymers of the type  $[Zn(\mu Cl_{2}py_{2}]_{\infty}$ , (II) (py is 3,5-dichloropyridine or 3,5-dibromopyridine; Hu & Englert, 2001). 2,6-Dichloropyrazine, the ligand involved in the present study, was expected to have similar steric requirements to the 3,5-dihalopyridines. Furthermore, coordination of a metal should only occur in the 4- and not in the 1position. Despite these similarities between the dihalopyridine and dihalopyrazine ligands, reaction of zinc dichloride with the latter does not result in the formation of а chain polymer, but gives the title tetrahedral mononuclear



#### Figure 1

The molecular view of (I) showing the three independent molecules. Displacement ellipsoids are at the 50% probability level and H atoms are shown as small spheres of arbitrary radii.

complex, (I), underlining the rare nature of the abovementioned type of coordination polymers.



We studied five crystals of (I) by X-ray diffraction and obtained four data sets of comparable accuracy; the intensity data from the fifth sample proved to be less satisfactory in terms of internal consistency and will not be reported. All crystals were twinned. A metrically possible transformation to a monoclinic centred cell is incompatible with the symmetry of the diffraction pattern. The twinning can be classified as metric merohedry (Nespolo & Ferraris, 2000) with monoclinic lattice metric symmetry (LMS). Metric merohedry means that all reflections of the twin components will overlap. In other words, the reciprocal lattice of the twin mimics that of a single crystal and, in the limiting case of a very small twin component, twinning might be completely overlooked. In the present case, the twin law interchanges a and b according to the matrix (010, 100, 001). Internal agreement factors  $(R_{int})$  for merging the intensity data under the LMS differ strongly between independent data sets and we wish to point out that this variation represents a warning sign for merohedral twinning. A synopsis of  $R_{int}$ , obtained under both the triclinic crystal symmetry and the monoclinic LMS, and the volume ratio between the twin components, is provided in Table 1. The discrepancy between  $R_{int}$  for merging in the LMS and the crystal symmetry may serve as an indicator for the volume ratio between the twin individuals. In the following discussion, in Table 2 and in the supplementary material), numerical values and their s.u.'s refer to the data set with the smallest contribution of a second individual.

The asymmetric unit of (I) contains three independent molecules (Fig. 1). They are similar not only in terms of bond lengths and angles, but also with respect to the dihedral angles subtended by the planes through the pyrazine ligands



#### Figure 2

A packing diagram for (I), viewed along the c axis.

 $[87.40 (18), 87.39 (18) \text{ and } 89.64 (18)^{\circ}]$ . A packing diagram is shown in Fig. 2. No particularly short intermolecular interactions occur, the shortest Cl···Cl distance being *ca* 3.39 Å.

## **Experimental**

Colourless crystals of (I) were grown by slow evaporation from a solution of ZnCl<sub>2</sub> and 2,6-Cl<sub>2</sub>-pyrazine (1:2) in ethanol under a flow of nitrogen.

#### Crystal data

$[ZnCl_2(C_4H_2Cl_2N_2)_2]$	Z = 6
$M_r = 434.22$	$D_x = 1.903 \text{ Mg m}^{-3}$
Triclinic, P1	Mo $K\alpha$ radiation
a = 12.4478 (7) Å	Cell parameters from 31 443
b = 12.4880 (8) Å	reflections
c = 15.6692 (9) Å	$\theta = 1.4-28.3^{\circ}$
$\alpha = 75.954 (1)^{\circ}$	$\mu = 2.67 \text{ mm}^{-1}$
$\beta = 76.134 (1)^{\circ}$	T = 293 (2)  K
$\gamma = 79.438 (1)^{\circ}$	Block, colourless
V = 2273.7 (2) Å <sup>3</sup>	$0.36 \times 0.30 \times 0.20 \text{ mm}$
Data collection	
Bruker SMART APEX CCD	11 257 independent reflections
area-detector diffractometer	6480 reflections with $I > 2\sigma(I)$
$\omega$ scans	$R_{\rm int} = 0.046$
Absorption correction: empirical	$\theta_{\rm max} = 28.3^{\circ}$
(SADABS; Sheldrick, 1996)	$h = -16 \rightarrow 16$
$T_{\rm min} = 0.398, T_{\rm max} = 0.587$	$k = -16 \rightarrow 16$

 $l = -20 \rightarrow 20$ 

31 443 measured reflections

Refinement

Refinement on $F^2$ $R[F^2 > 2\sigma(F^2)] = 0.038$ $wR(F^2) = 0.077$	H-atom parameters constrained $w = 1/[\sigma^2(F_o^2) + (0.0309P)^2]$ where $P = (F^2 + 2F^2)/3$
S = 0.81 11 257 reflections	$(\Delta/\sigma)_{\text{max}} = 0.001$ $\Delta\rho_{\text{max}} = 0.50 \text{ e} \text{ Å}^{-3}$
515 parameters	$\Delta \rho_{\rm min} = -0.37 \ {\rm e} \ {\rm A}^{-3}$

#### Table 1

Internal agreement factors  $(R_{int})$  – a hint for twinning.

Data set	$R_{\rm int}(\rm LMS)$	$R_{\rm int}({ m crystal})$	Volume fraction of 2nd twin individual
1	0.343	0.046	0.1016 (4)
2	0.154	0.061	0.3673 (8)
3	0.075	0.047	0.4765 (8)
4	0.067	0.048	0.4974 (9)

Table 2				
Selected	geometric parameters	(Å.	°).	

Zn1-N1	2.084 (3)	Zn1-Cl1	2.1833 (11)
Zn1-N3	2.084 (3)	Zn1-Cl2	2.1905 (11)
Zn2-N5	2.094 (3)	Zn2-Cl7	2.1757 (10)
Zn2-N7	2.086 (3)	Zn2-Cl8	2.1894 (10)
Zn3-N9	2.085 (3)	Zn3-Cl13	2.1924 (10)
Zn3-N11	2.095 (3)	Zn3-Cl14	2.1764 (11)
N1-Zn1-N3	101.30 (11)	Cl1-Zn1-Cl2	127.86 (5)
N5-Zn2-N7	101.26 (11)	Cl7-Zn2-Cl8	127.53 (4)
N9-Zn3-N11	101.53 (11)	Cl13-Zn3-Cl14	127.47 (4)

All H atoms were introduced in idealized positions and included as riding (C-H = 0.98 Å), with  $U_{iso}(H) = 1.3U_{eq}(non-H)$ .

Data collection: SMART (Bruker, 2001); cell refinement: SMART; data reduction: SAINT-Plus (Bruker, 1999); program(s) used to solve structure: SHELXTL (Sheldrick, 1997); program(s) used to refine structure: SHELXTL; molecular graphics: PLATON (Spek, 1990); software used to prepare material for publication: SHELXTL.

Support from the DFG and the Fonds der Chemischen Industrie is gratefully acknowledged.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: GD1164). Services for accessing these data are described at the back of the journal.

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