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Glycine zinc sulfate pentahydrate: redetermination at 10 K from time-of-flight neutron Laue diffraction

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Single crystals of glycine zinc sulfate pentahydrate [systematic name: hexaaquazinc tetraaquadiglycinezinc bis(sulfate)], [Zn(H₂O)₆][Zn(C₂H₅NO₂)₂- $(H_2O)_4](SO_4)_2$, have been grown by isothermal evaporation from aqueous solution at room temperature and characterized by single-crystal neutron diffraction. The unit cell contains two unique ZnO₆ octahedra on sites of symmetry $\overline{1}$ and two SO₄ tetrahedra with site symmetry 1; the octahedra comprise one [tetraaqua-diglycine zinc]²⁺ ion (centred on one Zn atom) and one $[hexaaquazinc]^{2+}$ ion (centred on the other Zn atom); the glycine zwitterion, $NH_3^+CH_2COO^-$, adopts a monodentate coordination to the first Zn atom. All other atoms sit on general positions of site symmetry 1. Glycine forms centrosymmetric closed cyclic dimers due to N-H. O hydrogen bonds between the amine and carboxylate groups of adjacent zwitterions and exhibits torsion angles varying from ideal planarity by no more than 1.2°, the smallest values for any known glycine zwitterion not otherwise constrained by a mirror plane. This work confirms the H-atom locations estimated in three earlier singlecrystal X-ray diffraction studies with the addition of independently refined fractional coordinates and U_{ii} parameters, which provide accurate internuclear X-H (X = N, O) bond lengths and consequently a more accurate and precise depiction of the hydrogen-bond framework.

1. Chemical context

Numerous coordination compounds of glycine (Glyc) with divalent metal sulfates are known. For the case of zinc, there is an anhydrous species, 2Glyc-ZnSO₄ (Moldobaev & Nogoev, 1970) and two hydrates, Glyc-ZnSO₄·3H₂O and Glyc--ZnSO₄·5H₂O. The trihydrate is dimorphic, occurring either as an orthorhombic crystal (space group $Pca2_1$) or as a monoclinic crystal $(P2_1/n)$ depending on the synthesis route (Fleck & Bohatý, 2004). The monoclinic form is isotypic with compounds of general formula $Glyc \cdot M(II)SO_4 \cdot 3H_2O$ where M(II) = Mg, Co or Fe (Oguey *et al.*, 2013*a*,*b*, 2014). Compounds with the general formula $Glyc \cdot M(II)SO_4 \cdot 5H_2O$ are known only as isotypic triclinic crystals (P1) for M(II) =Mg, Mn, Co, Fe and Zn (Lindqvist & Rosenstein, 1960; Elayaraja et al., 2007; Fleck & Bohatý, 2006; Tepavitcharova et al., 2012). Solubility data have been published for a purported Glyc·NiSO₄·5H₂O (Moldobaev et al., 1970; Alymkulova & Salyeva, 1987). We have collected as-yet unpublished X-ray powder-diffraction data from this species, showing that it is isotypic with the other known members of the series. The existence of Glyc·CuSO₄·5H₂O has been reported by Thilagavathi et al. (2012) but their work is in error, and quite describes unambiguously the well-known material CuSO₄·5H₂O.



We recently carried out the first neutron diffraction study of Glyc·MgSO₄·3H₂O and Glyc·MgSO₄·5H₂O using perdeuterated powder specimens (Howard *et al.*, 2016). Glyc-MgSO₄·5H₂O tends to form masses of crystals that are both of poor quality and are too small for single-crystal neutron diffraction study; however, this is not the case for Glyc-MnSO₄·5H₂O and Glyc·ZnSO₄·5H₂O, where fine tabular to blocky single crystals with volumes substantially in excess of 10 mm³ are formed with ease (Fig. 1). The objective of this work was to carry out the first single-crystal *neutron* diffraction study of any Glyc· $M(II)SO_4$ ·5H₂O compound, specifically using a specimen with M(II) = Zn.

Optical and mechanical properties of the title compound were reported by Balakrishnan & Ramamurthy (2007),



2. Structural commentary

Although the stoichiometry of the material is accurately reflected in its common name, glycine zinc sulfate pentahydrate, the presence of two symmetry-inequivalent Zn sites means that the crystallographically proper structural composition is the 'double' formula $[Gly \cdot ZnSO_4 \cdot 5H_2O]_2$, or more precisely $[Zn(H_2O)_6][Zn(H_2O)_4(C_2H_5NO_2)_2](SO_4)_2$; the unit cell contains *one* of these units.

The Zn1 coordination octahedron consists of tetraaquadiglycine zinc(II) with the glycine zwitterion (NH₃⁺CH₂COO⁻) coordinating to Zn by one of the carboxylate oxygen atoms (Fig. 2); the inversion centre results in an all-trans configuration for these units. The Zn2 octahedron has the form hexaaquazinc(II); the sulfate tetrahedra are isolated, accepting hydrogen bonds primarily (but not exclusively) from Zn-coordinating water molecules (Fig. 3). The S-O bond lengths (Table 1) reflect the number of hydrogen bonds accepted by each apical oxygen atom with a statistical significance which was not apparent from the powder refinement of Howard et al. (2016) but which are in excellent agreement with the single-crystal X-ray study of Tepavitcharova et al. (2012).

Bond lengths and angles of the glycine zwitterion agree very



Figure 1

Microphotograph of a representative Glyc-ZnSO₄·5H₂O single crystal viewed along the *a* axis; insets show details of less well-developed facets (*e.g.*, $\overline{111}$ and 102). Drawings with each face labelled by the Miller index are shown on the right and a quantitative representation of the model is included in the CIF data. Figure produced and CIF code exported using *WinXMorph* (Kaminsky, 2005, 2007).

well with other determinations of related compounds made by X-ray single-crystal diffraction at higher temperatures and extremely well with the determinations in α glycine at room temperature by neutron single-crystal diffraction (Jönsson & Kvick, 1972; Power et al., 1976), particularly in respect of their mean N-H bond lengths (1.039 Å) and mean C-H bond lengths (1.090 Å). The glycine zwitterion is remarkably planar, with torsion angles O7-C1-C2-N1 = $-1.18(3)^{\circ}$ and O8-C1- $C2-N1 = 179.23 (2)^{\circ}$, even by comparison with, for example, glycine nickel bromide tetrahydrate (Fleck & Bohatý, 2005), glycine lithium chromate monohydrate and glycine lithium molybdate (Fleck et al., 2006), where torsion angles are in the range 169-176°. Only in glycine magnesium chloride tetrahydrate,

Table 1

Comparison of bond lengths (Å), polyhedral volumes (Å³) and various distortion metrics (*cf.*, Robinson *et al.*, 1971) in Glyc-ZnSO₄·5H₂O from this work and the three preceding single-crystal X-ray diffraction studies.

The distortion index and quadratic elongation are dimensionless, whereas the bond-angle variance is in units of degrees squared.

	This work	Balamurugan et al. (2011)	Tepavitcharova et al. (2012)	Oguey et al. (2013c)
	Single-crystal neutron	Single-crystal X-ray	Single-crystal X-ray	Single-crystal X-ray
	T = 10 K	T = 293 K	T = 150 K	T = 153 K
S-O1*	1.474 (5)	1.472 (2)	1.472 (1)	1.473 (2)
S-O2	1.484 (4)	1.478 (2)	1.482 (1)	1.485 (2)
S-O3*	1.473 (4)	1.472 (2)	1.477 (1)	1.481 (2)
S-O4	1.480 (5)	1.481 (2)	1.484 (1)	1.479 (2)
Mean S–O	1.478	1.476	1.479	1.479
SO ₄ volume	1.656	1.649	1.659	1.661
Distortion index	0.0028	0.0025	0.0027	0.0022
Quadratic elongation	1.000	1.000	1.000	1.000
Bond-angle variance	0.410	0.268	0.320	0.420
Zn1-O5	2.039 (2)	2.024 (3)	2.032 (1)	2.035 (2)
Zn1-O6	2.093 (2)	2.101 (3)	2.098 (1)	2.098 (2)
$Zn1-O7^{\dagger}$	2.173 (2)	2.181 (3)	2.177 (1)	2.176 (2)
Mean Zn1–O	2.102	2.102	2.102	2.103
ZnO ₆ volume	12.338	12.339	12.339	12.336
Distortion index	0.0227	0.0251	0.0238	0.0232
Quadratic elongation	1.003	1.003	1.003	1.003
Bond-angle variance	6.308	4.815	5.975	6.292
Zn2-O9	2.129 (3)	2.141 (3)	2.133 (1)	2.135 (2)
Zn2-O10	2.067 (3)	2.071 (3)	2.070 (1)	2.072 (2)
Zn2-O11	2.075 (2)	2.063 (3)	2.065 (2)	2.065 (2)
Mean Zn2–O	2.090	2.092	2.089	2.091
ZnO_6 volume	12.127	12.176	12.123	12.145
Distortion index	0.0124	0.0156	0.0139	0.0142
Quadratic elongation	1.003	1.002	1.002	1.002
Bond-angle variance	7.982	5.942	6.617	6.541
C1-07	1.272 (4)	1.272 (5)	1.274 (2)	1.278 (3)
C1-O8	1.240 (3)	1.228 (5)	1.236 (2)	1.234 (3)
C1-C2	1.523 (4)	1.516 (5)	1.525 (3)	1.522 (3)
C2-N1	1.481 (2)	1.478 (5)	1.480 (2)	1.480 (3)

*Denotes sulfate O atoms accepting two hydrogen bonds instead of three. [†]Denotes carboxylate oxygen ligand instead of water oxygen.



Figure 2

Local coordination environment of the Zn1 atom. Displacement ellipsoids are drawn at the 50% probability level for H and 90% for all other atoms. Dashed lines indicate N– $H \cdots O$ hydrogen bonds. [Symmetry code: (i) 1 - x, -y, 1 - z.]

Local coordination environment of the Zn2 atom. Displacement ellipsoids are drawn at the 50% probability level for H and 90% for all other atoms. Dashed lines indicate O– H···O hydrogen bonds. [Symmetry code: (i) 1 - x, -y, 1 - z.]

Table 2 Comparison of X-H(D) bond lengths (Å) from earlier work (*a*-*e*) with our own (*f*).

Element symbols indicate the cation in each compound. 'X-ray' denotes single-crystal X-ray diffraction; 'NPD' denotes a neutron powder diffraction experiment on a deuterated analogue carried out at 10 K; 'neutron' indicates single-crystal neutron diffraction on a protonated analogue carried out at 10 K. Note that the atom symbols employed in our work are the same as those used by Elayaraja *et al.* (2007) and by Howard *et al.* (2016). Although other authors have used different atom labels – and indeed use them inconsistently in their own reports – we list equivalent contacts in this table.

	Mg, X-ray ^(a)	Mg, NPD ^(b)	Co, X-ray ^(c)	Mg, X-ray ^(c)	Zn, X-ray ^(c)	Zn, X-ray ^(d)	Zn, X-ray ^(e)	Zn, neutron ^(f)
N-H1N	0.87 (4)	1.008 (4)	0.847 (1)	0.849(1)	0.881 (2)	0.85 (2)	0.910 (2)	1.033 (7)
N-H2N	0.87 (4)	0.982 (4)	0.907(1)	0.898 (1)	0.904 (1)	0.83 (3)	0.911(2)	1.028 (8)
N-H3N	0.87 (5)	0.991 (5)	0.904 (1)	0.902(1)	0.946 (1)	absent	0.910(2)	1.022 (6)
Average N-H	0.87	0.995	0.877	0.874	0.892	0.84	0.911	1.030
C-H2A	0.970 (4)	1.077 (4)	0.961 (1)	0.960(1)	0.967 (2)	0.970 (3)	0.990 (2)	1.085 (6)
C-H2B	0.970 (3)	1.083 (4)	0.901 (1)	1.014 (1)	1.050 (2)	0.970 (3)	0.990 (2)	1.091 (7)
Average C-H	0.970	1.080	0.931	0.987	1.009	0.970	0.990	1.088
O5-H5A	0.84 (3)	0.975 (5)	0.880(1)	0.789(1)	0.879 (2)	0.85 (2)	0.83 (3)	0.973 (7)
O5-H5B	0.85 (3)	0.946 (5)	0.914 (1)	0.930(1)	0.838(1)	0.85 (3)	0.85 (3)	0.997 (7)
O6-H6A	0.84 (2)	0.987 (5)	0.964 (1)	0.875 (1)	0.864 (1)	0.83 (3)	0.86 (3)	0.981 (6)
O6-H6B	0.83 (3)	0.988 (5)	0.906 (1)	0.897 (1)	0.886(1)	0.84 (3)	0.85 (2)	0.985 (6)
O9-H9A	0.83 (2)	0.977 (5)	0.864 (1)	0.871 (1)	0.881 (2)	0.87 (3)	0.86 (2)	0.979 (5)
O9-H9B	0.84 (2)	0.984 (4)	0.884 (1)	0.901 (1)	0.964 (1)	0.87 (2)	0.87 (3)	0.966 (6)
O10-H10A	0.84 (4)	0.954 (5)	0.972 (1)	0.911 (1)	0.887(1)	0.82 (2)	0.87 (2)	0.977 (8)
O10-H10B	0.84 (3)	0.972 (5)	0.855 (1)	0.821 (1)	0.913 (1)	0.84 (2)	0.85 (2)	0.978 (6)
O11-H11A	0.84 (3)	1.002 (5)	0.822(1)	0.884 (1)	0.808(1)	0.83 (3)	0.86 (2)	0.966 (6)
O11-H11B	0.83 (3)	0.965 (5)	0.906 (1)	0.859 (1)	0.900 (1)	0.84 (2)	0.84 (2)	0.966 (6)
Average O-H	0.84	0.975	0.897	0.874	0.882	0.85	0.85	0.977

(a) Elayaraja et al. (2007); (b) Howard et al. (2016); (c) Tepavitcharova et al. (2012); (d) Balamurugan et al. (2011); (e) Oguey et al. (2013c); (f) this work.

where the glycine zwitterion lies on a mirror plane, are values of 180° realized (Fleck & Bohatý, 2005). In α -glycine, the equivalent torsion angles are -19.60 (3) and 161.28 (2)°.

In respect of the heavy atoms, agreement in the interatomic distances and angles between the single-crystal X-ray and single-crystal neutron refinements (Table 1) is excellent, with some differences emerging in respect of the room-temperature refinement by Balamurugan *et al.* (2011). However, the neutron data provide a substantial improvement in accuracy with respect to the X-ray data in the hydrogen atom's fractional coordinates and U_{ij} parameters. This is not surprising since neutrons are sensitive to the nuclear positions and X-rays to the electron density; in the covalent X-H bond the centroid of the H-atom's electron-density distribution is displaced towards the heavy atom by 0.1 Å, yielding X-H distances from 10–15% shorter than the true internuclear separation (Coppens, 1997). Table 2 compares X-H bond lengths from a range of Gly·M(II)SO₄·5H₂O crystals obtained by single-crystal X-ray diffraction and by neutron powder diffraction. In the work of Elayaraja *et al.* (2007), methyl



Figure 4

Packing of polyhedra in the structure of Glyc-ZnSO₄·5H₂O viewed along a (left) and along c (right). ZnO₆ octahedra are green, SO₄ tetrahedra are yellow.

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Table 3	
Hydrogen-bond	geometry (Å, °).

$D - H \cdot \cdot \cdot A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$O5-H5A\cdots O4^{i}$	0.973 (7)	1.793 (7)	2.755 (4)	169.0 (7)
$O5-H5B\cdots O8^{ii}$	0.997 (7)	1.656 (8)	2.642 (4)	168.9 (6)
$O6-H6A\cdots O3^{iii}$	0.981 (6)	1.722 (6)	2.696 (3)	170.8 (5)
$O6-H6B\cdots O4^{iv}$	0.985 (6)	1.751 (5)	2.729 (3)	171.8 (7)
$O9-H9A\cdotsO1^{iv}$	0.979 (5)	1.732 (5)	2.707 (3)	173.8 (6)
O9−H9B···O2	0.966 (6)	1.895 (6)	2.811 (3)	157.2 (6)
$O10-H10A\cdots O3^{iii}$	0.977 (8)	1.740 (8)	2.713 (4)	173.0 (7)
$O10-H10B\cdots O2^{v}$	0.979 (6)	1.811 (7)	2.745 (4)	158.5 (7)
$O11 - H11A \cdots O2^{vi}$	0.966 (6)	1.772 (6)	2.726 (3)	168.6 (6)
$O11 - H11B \cdots O1$	0.966 (6)	1.824 (6)	2.750 (3)	159.5 (7)
$C2-H2A\cdots O2^{vi}$	1.085 (6)	2.682 (9)	3.351 (4)	119.4 (6)
$C2-H2A\cdots O5^{vii}$	1.085 (6)	2.716 (8)	3.489 (3)	127.9 (6)
$C2-H2B\cdots O10$	1.091 (7)	2.579 (8)	3.649 (4)	166.7 (7)
$N1 - H1N \cdot \cdot \cdot O7^{vii}$	1.033 (7)	1.853 (7)	2.848 (3)	160.8 (7)
$N1 - H2N \cdots O4$	1.027 (8)	1.961 (7)	2.877 (3)	147.0 (7)
$N1 - H3N \cdots O6$	1.022 (6)	2.216 (7)	3.066 (3)	139.5 (5)

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

hydrogens were positioned geometrically and allowed to ride with $U_{iso}(H) = 1.2U_{eq}(C)$; water hydrogen atoms were refined with restraints; the N-H bond lengths were all restrained to be equal. Balamurugan *et al.* (2011) placed the majority of their hydrogen atoms geometrically, although failed to identify the third amine hydrogen atom; water and the two amine H atoms were refined isotropically whereas the two methyl hydrogen atoms were riding on the C atom. Tepavitcharova *et al.* (2012) placed the hydrogen atoms in Gly·ZnSO₄·5H₂O geometrically and treated all of them as riding on their associated heavy atom during refinement. Oguey *et al.* (2013*c*) allowed all water hydrogen atoms to refine isotropically but fixed the methyl and amine hydrogen atoms in this work were unrestrained and allowed to refine anisotropically.

Our values for the N-H and C-H bond lengths are in excellent agreement with other single-crystal neutron diffraction work, as noted in the preceding paragraph. Our values for the O-H bond lengths also agree well with those found in similar environments in hydrated M(II) coordination compounds, such as MgSO₄·11H₂O and MgCrO₄·11H₂O where the average O-H = 0.974 Å (Fortes *et al.*, 2013), MgSeO₄·9H₂O, O-H_{av} = 0.972 Å (Fortes *et al.*, 2015), and MgSeO₄·7H₂O, O-H_{av} = 0.974 Å (Fortes & Gutmann, 2014).

3. Supramolecular features

The overall three-dimensional framework is completed by a variety of hydrogen bonds with a range of strengths (Table 3). Fig. 4 shows the spatial relationship of the main structural elements. The majority of the hydrogen bonds are $O-H\cdots O$ contacts of medium strength ($1.66 < H \cdots O < 1.90$ Å) and high linearity ($\angle O-H \cdots O > 157^{\circ}$), characteristic of two-centred interactions. As expected, the $N-H \cdots O$ hydrogen bonds are weaker (*i.e.*, longer, $1.85 < H \cdots O < 2.22$ Å) and more strained ($\angle N-H \cdots O$ between approx. 140–160°). The methyl groups appear to participate in weak $C-H \cdots O$ hydrogen bonds (*cf.*,

Steiner & Desiraju, 1998). One C–H···O bond is evidently a two-centred interaction, being the shortest and most of linear contact of this kind, with H···O = 2.58 (1) Å and \angle C–H···O = 167 (1)°. The other, involving C–H2A, is clearly a two-centred interaction (*i.e.*, a bifurcated hydrogen bond) with 'arms' of roughly equal length, H···O $\simeq 2.7$ Å and C–H···O angles of 119 and 128° involving O2 and O5, respectively.

3.1. Glycine dimers

A hitherto unrecognized aspect of the supramolecular structure of Glyc·M(II)SO₄·5H₂O compounds is the presence of glycine dimers (Fig. 5). These occur as closed cyclic structures formed by N-H···O hydrogen bonds between the amine group of one glycine zwitterion and the Zn-coordinating carboxylate oxygen (O7) of another zwitterion, related to the first by an inversion centre. A similar cyclic dimer occurs in the structure of α -glycine. A direct comparison between the dimers in Glyc·ZnSO₄·5H₂O and in α -glycine is shown in Fig. 6; clearly, the main difference between these two dimers is the orientation of the carboxylate group, which is presumably due to the influence of a divalent metal being coordinated by



Figure 5

Connectivity between adjacent Zn1 octahedra is *via* a closed cyclic glycine dimer. As before, displacement ellipsoids are drawn at the 50% probability level for H and 90% for all other atoms. Dashed lines indicate N-H···O hydrogen bonds. [Symmetry codes: (i) 1 - x, 1 - y, 1 - z; (ii) 1 - x, -y, 1 - z; (iii) x, 1 + y, z.]



Figure 6

Comparison of the closed cyclic dimers involving zwitterionic glycine that occur in the crystal structures of α -glycine (top) and in Glyc-ZnSO₄·5H₂O (bottom).

the O7 carboxylate oxygen. Experimental studies of aqueous solutions indicate that only glycine monomers exist in the liquid phase (Huang *et al.*, 2008). However, there has been widespread disagreement on this matter from computational studies, which indicate either that there are no dimers (Hamad & Catlow, 2011), substantial quantities of closed zwitterionic dimers (Friant-Michel & Ruiz-López, 2010), or a small fraction of open dimers (Yani *et al.*, 2012) present in saturated solutions. The presence or absence of glycine polymerization in coordination compounds such as these may be useful in understanding the association of glycine in saturated aqueous solutions during nucleation and the role of solvated metal ions in polymerizing amino acids in Earth's Hadean oceans (Kitadai *et al.*, 2011, 2016) or in extraterrestrial oceans elsewhere in our solar system (Kimura & Kitadai, 2015).

4. Database survey

A search of the Cambridge Structural Database (Groom *et al.*, 2016) identified the following directly relevant entries:

Glyc·*M*(II)SO₄ pentahydrates: 672589 (Mg); 857075 (Mg); 1451396 (Mg); 296329 (Co); 857073 (Co); 806684 (Zn); 857076 (Zn); 936400 (Zn).

Raman vibrational frequencies and mode assignments of α -glycine (*cf.*, Stenbäck, 1976: Rosado *et al.*, 1998: Yang *et al.*, 2008), Glyc·MgSO₄·5H₂O (Howard *et al.*, 2016) and the title compound.

Meaning of symbols: v =stretch; $\delta =$ deformation; $\rho =$ rock; $\omega =$ wag; $\Gamma =$ twist; (A) = asymmetric; (S) = symmetric.

	α -Glycine ^{<i>a</i>}	Glyc·MgSO ₄ ·5H ₂ O ^a	Glyc-ZnSO ₄ -5H ₂ O
Vibrational mode	180 s, 18 mW	1400 s, 18 mW	540 s, 18 mW
$\delta M^{2+} - O(?)$	-	208 236	203 220
$\delta \text{ CCN}^+ ho \text{ COO}^-$	356	361	382
$\delta(S) \operatorname{SO_4}^{2-}$ $\rho \operatorname{COO^-}$ $\omega \operatorname{COO^-}$	– 497 601	453 522 597	451 527 582 599
$\delta(A)$ SO ₄ ²⁻		623 645	626 644
δ COO ⁻ unknown	696 -	_ 794	_
$ \begin{array}{l} \nu C - C^{+} \\ \nu C - N \\ \nu C - O \end{array} $	893	890 905	890 906
$\rho CH_2 \nu(S) SO_4^{2-} \nu C-N$	922 - 1036	- 983.8 1020	- 983.2 1021
$\nu(A)$ SO ₄ ²⁻	-	1077 1100	1078 1101
$ ho \mathrm{NH}^{3+}$	1108 1140	1139	1141
ω CH ₂ Γ CH ₂	1325	1305 1328	1306 1327
$\nu(S) \operatorname{COO}^-$	1410	1395	1391
$\delta(S) \operatorname{CH}_2$	1441 1457	1434	1433
$\delta(A) \operatorname{NH}^{3+}$ $\delta(S) \operatorname{NH}^{3+}$	1502 1516 1569	1488	1488
$\nu C - C^+$ ωCH_2	1634	1597	1590
$\nu(A) \operatorname{COO}^{-} \\ \nu(S) \operatorname{CH}_{2} \\ \nu(A) \operatorname{CH}_{2} \\ \nu(S) \operatorname{NH}^{3+} $	1670 2972 3009 3143	1631 2997 3038 -	1614 2996 3037 -
ν(<i>S</i>) H ₂ O	_	3248	3204 3233
$\nu(A)$ H ₂ O	-	3384	3331 3405

^aHoward et al. (2016).

Glyc·*M*(II)SO₄ trihydrates: 989590 (Mg); 1451397 (Mg); 857074 (Co); 936396 (Fe); 243588 (Zn, orthorhombic); 936394 (Zn, monoclinic).

Glyc·M(II)SO₄ hexahydrates: 1285639 (Ni).

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Figure 7

Raman spectra of α -glycine (top) and Glyc·ZnSO₄·5H₂O (bottom). Selected vibrational modes are labelled and a complete quantitative listing is given in Table 4.

5. Vibrational spectroscopy

Laser-stimulated Raman spectra were measured using a portable B&WTek *i*-Raman Plus spectrometer equipped with a 532 nm laser ($P_{\text{max}} = 37 \text{ mW}$ at the probe tip) that records spectra over the range 171–4002 cm⁻¹ with an optimal resolution of 3 cm⁻¹. Measurements were carried out on powdered specimens of α -glycine and Glyc-ZnSO₄·5H₂O. Samples were measured in thin-walled glass vials using the BC100 fibre-optic coupled Raman probe; the total integration time and laser power for each sample is provided with the tabulated results (see supplementary material).

The Raman spectrum of Glyc·ZnSO₄·5H₂O (Fig. 6) is virtually identical with that of Glyc·MgSO₄·5H₂O reported in Howard *et al.* (2016) and is in excellent agreement with the spectrum shown in Tepavitcharova *et al.* (2012). Numerical data of the Raman spectrum are provided as an electronic supplement; peak positions and vibrational mode assignments are given in Table 4. The main differences between the two divalent-metal-substituted compounds include the blueshifting of octahedral deformation modes and blue-shifting of both symmetric and asymmetric COO⁻ stretching modes. A large blue-shift of $\nu(A)$ and $\nu(S)$ COO⁻ occurs when glycine coordinates to Mg²⁺ and the shift increases when glycine coordinates to Zn²⁺. Raman spectra of α -glycine and Glyc·ZnSO₄·5H₂O are shown in Fig. 7.

6. Synthesis and crystallization

Glyc·ZnSO₄·5H₂O was crystallized by evaporation at room temperature of an equimolar aqueous solution of α -glycine (Alfa Aesar A13816) and ZnSO₄·7H₂O (Sigma Aldrich Z4750) in deionized water (Alfa Aesar 36645). Unlike the MgSO₄-bearing analogue, Glyc·ZnSO₄·5H₂O forms large well-faceted crystals that are both amenable to morphological study and suitably large for single-crystal neutron diffraction analysis. Fig. 1 shows photographs of a representative crystal viewed along its a axis and series of drawings with indexed crystal faces.

7. Data collection and refinement

Crystal data, data collection and structure refinement details are summarized in Table 5. Data were collected from a pair of single crystals at a series of four discrete rotational positions about the vertical axis, each frame being counted for 5 h, equivalent to 800 µAhr of ISIS proton beam current per frame. The structure of $Glyc(d_5) \cdot MgSO_4 \cdot 5D_2O$ at 10 K reported by Howard et al. (2016) was used as a starting point for the refinement. A total of eleven peaks, with the largest $\sigma(F_{obs}-F_{calc})$ values were omitted from the refinement; such outliers are fairly common in SXD measurement when peaks occur close to the edges of detectors. A mild restraint on the U_{ii} parameters of the sulfur atom was imposed (SHELX ISOR command) in order to avoid a slightly non-positive-definite displacement ellipsoid. Since sulfur has the smallest neutron scattering cross section of any atom in the structure, and since it is both comparatively heavy and the temperature is very low, it is not surprising that – within errors – the effective U_{iso} parameter should refine to a small negative value.

Table 5Experimental details.	
Crystal data	
Chemical formula	$[Zn(H_2O)_6][Zn(C_2H_5NO_2)_2-(H_2O)_4](SO_4)_2$
M _r	653.20
Crystal system, space group	Triclinic, P1
Temperature (K)	10
a, b, c (Å)	5.9601 (15), 6.7670 (17), 13.112 (4)
α, β, γ (°)	84.955 (18), 83.25 (2), 83.042 (19)
$V(\dot{A}^3)$	519.8 (2)
Ζ	1
Radiation type	Neutron, $\lambda = 0.48-7.0$ A
$\mu (\text{mm}^{-1})$	$5.02 + 0.0182 * \lambda$
Crystal size (mm)	$4 \times 2.5 \times 1$
Data collection	
Diffractometer	SXD
Absorption correction	Numerical. The linear absorption coefficient is wavelength dependent and is calculated as: $\mu = 5.0165 + 0.0182 * \lambda \text{ [cm}^{-1}\text{] as}$ determined by Gaussian integration in <i>SXD2001</i> (Gutmann, 2005)
No. of measured, independent and	8296, 8296, 8296
observed $[I > 2\sigma(I)]$ reflections	
R _{int}	0.089
Refinement	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.089, 0.246, 1.09
No. of reflections	8296
No. of parameters	291
No. of restraints	12
H-atom treatment	All H-atom parameters refined
$\Delta \rho_{\rm max}, \Delta \rho_{\rm min} \ ({\rm e} \ {\rm A}^{-3})$	3.20, -3.47

Computer programs: *SXD2001* (Gutmann, 2005), *SHELXT2014* (Sheldrick, 2015*a*; Gruene *et al.*, 2014), *SHELXL2014* (Sheldrick, 2015*b*; Gruene *et al.*, 2014), *DIAMOND* (Putz & Brandenburg, 2006) and *publCIF* (Westrip, 2010).

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supporting information

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Glycine zinc sulfate pentahydrate: redetermination at 10 K from time-of-flight neutron Laue diffraction

A. Dominic Fortes, Christopher M. Howard, Ian G. Wood and Matthias J. Gutmann

Computing details

Data collection: *SXD2001* (Gutmann, 2005); cell refinement: *SXD2001* (Gutmann, 2005); data reduction: *SXD2001* (Gutmann, 2005); program(s) used to solve structure: SHELXT2014 (Sheldrick, 2015a; Gruene *et al.*, 2014); program(s) used to refine structure: *SHELXL2014* (Sheldrick, 2015b; Gruene *et al.*, 2014); molecular graphics: *DIAMOND* (Putz & Brandenburg, 2006); software used to prepare material for publication: *publCIF* (Westrip, 2010).

Hexaaquazinc(II) tetraaquadiglycinezinc(II) bis(sulfate)

Crystal data

 $[Zn(H_2O)_6][Zn(C_2H_5NO_2)_2(H_2O)_4](SO_4)_2$ $M_r = 653.20$ Triclinic, $P\overline{1}$ a = 5.9601 (15) Å b = 6.7670 (17) Å c = 13.112 (4) Å $a = 84.955 (18)^{\circ}$ $\beta = 83.25 (2)^{\circ}$ $\gamma = 83.042 (19)^{\circ}$ $V = 519.8 (2) \text{ Å}^3$

Data collection

SXD

diffractometer Radiation source: ISIS neutron spallation source time–of–flight LAUE diffraction scans Absorption correction: numerical The linear absorption coefficient is wavelength dependent and is calculated as: $\mu = 5.0165 +$ $0.0182 * \lambda$ [cm⁻¹] as determined by Gaussian integration in *SXD2001* (Gutmann, 2005) *Refinement*

Refinement on F^2 Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.089$ $wR(F^2) = 0.246$ S = 1.098296 reflections 291 parameters 12 restraints Hydrogen site location: difference Fourier map Z = 1 F(000) = 336 $D_x = 2.086 \text{ Mg m}^{-3}$ Neutron radiation, $\lambda = 0.48-7.0 \text{ Å}$ Cell parameters from 550 reflections $\mu = 5.02 + 0.0182 * \lambda \text{ mm}^{-1}$ T = 10 KTabular, colourless $4 \times 2.5 \times 1 \text{ mm}$

8296 measured reflections 8296 independent reflections 8296 reflections with $I > 2\sigma(I)$ $R_{int} = 0.089$ $\theta_{max} = 87.4^{\circ}, \theta_{min} = 8.2^{\circ}$ $h = -15 \rightarrow 15$ $k = -18 \rightarrow 16$ $l = -28 \rightarrow 29$

All H-atom parameters refined $w = 1/[\sigma^2(F_o^2) + (0.1376P)^2 + 36.2519P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} < 0.001$ $\Delta\rho_{max} = 3.20 \text{ e} \text{ Å}^{-3}$ $\Delta\rho_{min} = -3.47 \text{ e} \text{ Å}^{-3}$ Extinction correction: SHELXL, $Fc^* = kFc[1+0.001xFc^2\lambda^3/sin(2\theta)]^{-1/4}$ Extinction coefficient: 0.0386 (18)

Special details

Experimental. For peak integration a local UB matrix refined for each frame, using approximately 50 reflections from each of the 11 detectors. Hence _cell_measurement_reflns_used 550 For final cell dimensions a weighted average of all local cells was calculated Because of the nature of the experiment, it is not possible to give values of theta_min and theta_max for the cell determination. The same applies for the wavelength used for the experiment. The range of wavelengths used was 0.48–7.0 Angstroms, BUT the bulk of the diffraction information is obtained from wavelengths in the range 0.7–2.5 Angstroms. The data collection procedures on the SXD instrument used for the single-crystal neutron data collection are most recently summarized in the Appendix to the following paper Wilson, C.C. (1997). J. Mol. Struct. 405, 207–217

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

	x	У	Ζ	$U_{ m iso}$ */ $U_{ m eq}$
S1	0.0141 (6)	0.9128 (6)	0.1847 (4)	0.0008 (6)
O1	0.1809 (4)	1.0196 (3)	0.1159 (2)	0.0036 (4)
O2	-0.0249 (4)	0.7303 (3)	0.1375 (2)	0.0037 (4)
O3	-0.2027 (4)	1.0416 (3)	0.2010 (2)	0.0040 (4)
O4	0.1050 (4)	0.8522 (4)	0.2845 (2)	0.0038 (3)
Zn1	0.5000	0.0000	0.5000	0.0017 (4)
05	0.1954 (4)	0.1618 (4)	0.5392 (2)	0.0049 (4)
H5A	0.0913 (11)	0.1389 (11)	0.6009 (6)	0.0198 (11)
H5B	0.0998 (10)	0.2109 (9)	0.4826 (6)	0.0170 (10)
O6	0.4360 (4)	0.0464 (4)	0.3456 (2)	0.0043 (4)
H6A	0.5654 (10)	0.0305 (9)	0.2922 (5)	0.0162 (10)
H6B	0.3082 (10)	-0.0116 (10)	0.3229 (6)	0.0179 (11)
07	0.6512 (4)	0.2793 (4)	0.4768 (2)	0.0037 (3)
O8	0.9812 (4)	0.3195 (4)	0.3809 (2)	0.0060 (4)
N1	0.3991 (2)	0.5025 (2)	0.34432 (14)	0.0044 (2)
H1N	0.3634 (11)	0.5559 (12)	0.4165 (6)	0.0218 (13)
H2N	0.3159 (12)	0.6073 (10)	0.2963 (7)	0.0212 (12)
H3N	0.3294 (11)	0.3719 (9)	0.3435 (6)	0.0199 (12)
C1	0.7711 (3)	0.3465 (3)	0.39746 (19)	0.0029 (3)
C2	0.6484 (3)	0.4781 (3)	0.3163 (2)	0.0041 (3)
H2A	0.7111 (11)	0.6233 (9)	0.3086 (7)	0.0220 (13)
H2B	0.6871 (11)	0.4134 (12)	0.2420 (6)	0.0222 (13)
Zn2	0.5000	0.5000	0.0000	0.0008 (3)

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters $(Å^2)$

supporting information

09	0.2958 (4)	0.3897 (3)	0.1320 (2)	0.0045 (4)	
H9A	0.2434 (11)	0.2593 (8)	0.1279 (6)	0.0175 (11)	
H9B	0.1606 (10)	0.4824 (9)	0.1443 (6)	0.0190 (11)	
O10	0.7911 (4)	0.3443 (4)	0.0509(2)	0.0048 (4)	
H10A	0.7828 (12)	0.2322 (9)	0.1030 (6)	0.0180 (10)	
H10B	0.8989 (11)	0.2959 (10)	-0.0061 (6)	0.0197 (11)	
011	0.5434 (4)	0.7298 (4)	0.0881 (2)	0.0054 (4)	
H11A	0.6938 (9)	0.7484 (9)	0.1036 (6)	0.0182 (11)	
H11B	0.4422 (10)	0.8523 (8)	0.0915 (7)	0.0200 (12)	

Atomic displacement parameters $(Å^2)$

	U^{11}	<i>U</i> ²²	U^{33}	U^{12}	U^{13}	U ²³
S1	0.0006 (10)	0.0009 (11)	0.0010 (18)	0.0001 (8)	-0.0003 (9)	0.0000 (11)
01	0.0027 (6)	0.0037 (7)	0.0038 (10)	-0.0006 (5)	0.0016 (5)	0.0001 (7)
O2	0.0029 (6)	0.0028 (6)	0.0058 (11)	-0.0007 (5)	-0.0007 (6)	-0.0012 (7)
03	0.0017 (6)	0.0040 (7)	0.0053 (11)	0.0018 (5)	0.0010 (6)	0.0000 (7)
04	0.0039 (6)	0.0054 (7)	0.0018 (10)	0.0001 (5)	-0.0015 (6)	0.0011 (7)
Zn1	0.0015 (4)	0.0017 (4)	0.0017 (5)	-0.0001 (3)	-0.0001 (3)	0.0000 (3)
05	0.0036 (7)	0.0071 (8)	0.0032 (11)	0.0014 (5)	-0.0002 (6)	0.0011 (7)
H5A	0.015 (2)	0.028 (3)	0.014 (3)	-0.0027 (18)	0.0046 (18)	0.002 (2)
H5B	0.015 (2)	0.019 (2)	0.016 (3)	0.0017 (15)	-0.0065 (17)	0.004 (2)
06	0.0041 (7)	0.0064 (7)	0.0021 (11)	-0.0005 (5)	-0.0004 (6)	0.0002 (7)
H6A	0.0142 (18)	0.021 (2)	0.013 (3)	-0.0020 (15)	0.0034 (16)	-0.005 (2)
H6B	0.017 (2)	0.023 (2)	0.017 (3)	-0.0076 (17)	-0.0052 (18)	-0.005 (2)
07	0.0042 (6)	0.0041 (7)	0.0026 (10)	-0.0011 (5)	-0.0004 (6)	0.0014 (7)
08	0.0028 (7)	0.0091 (9)	0.0050 (12)	0.0008 (5)	0.0001 (6)	0.0022 (8)
N1	0.0030 (4)	0.0057 (5)	0.0041 (7)	0.0012 (3)	-0.0013 (4)	0.0004 (5)
H1N	0.017 (2)	0.033 (3)	0.015 (3)	0.002 (2)	0.0004 (19)	-0.009 (3)
H2N	0.019 (2)	0.020 (2)	0.025 (4)	0.0015 (18)	-0.011 (2)	0.006 (2)
H3N	0.018 (2)	0.015 (2)	0.027 (4)	-0.0066 (16)	-0.001 (2)	0.001 (2)
C1	0.0020 (5)	0.0034 (6)	0.0032 (9)	0.0000 (4)	-0.0011 (5)	0.0007 (6)
C2	0.0042 (6)	0.0047 (6)	0.0030 (9)	-0.0002 (4)	-0.0004 (5)	0.0017 (6)
H2A	0.020 (2)	0.013 (2)	0.033 (4)	-0.0078 (17)	-0.005 (2)	0.007 (2)
H2B	0.018 (2)	0.035 (3)	0.013 (3)	0.007 (2)	-0.0031 (19)	-0.006 (3)
Zn2	0.0008 (4)	0.0008 (4)	0.0009 (5)	-0.0001 (3)	-0.0001 (3)	0.0001 (3)
09	0.0045 (7)	0.0035 (7)	0.0050 (11)	-0.0005 (5)	0.0005 (6)	0.0004 (7)
H9A	0.021 (2)	0.0115 (17)	0.021 (3)	-0.0064 (15)	-0.001 (2)	-0.001 (2)
H9B	0.0135 (19)	0.016 (2)	0.024 (4)	0.0064 (14)	0.0009 (18)	-0.001 (2)
O10	0.0039 (7)	0.0047 (7)	0.0050 (11)	0.0019 (5)	-0.0012 (6)	0.0010 (7)
H10A	0.024 (3)	0.015 (2)	0.013 (3)	0.0011 (17)	0.0002 (19)	0.0054 (19)
H10B	0.020 (2)	0.020 (2)	0.016 (3)	0.0065 (17)	0.0040 (19)	-0.002 (2)
011	0.0033 (6)	0.0048 (7)	0.0086 (12)	0.0002 (5)	-0.0015 (6)	-0.0033 (8)
H11A	0.0091 (16)	0.022 (2)	0.025 (3)	-0.0030 (15)	-0.0057 (17)	-0.004 (2)
H11B	0.017 (2)	0.0118 (18)	0.031 (4)	0.0059 (14)	-0.003 (2)	-0.006 (2)

Geometric parameters (Å, °)

<u>S1—O3</u>	1.473 (4)	N1—H1N	1.033 (7)
S101	1.474 (5)	N1—C2	1.481 (2)
S1—O4	1.480 (6)	C1—C2	1.523 (4)
S1—O2	1.484 (4)	C2—H2A	1.085 (6)
Zn1—O5	2.039 (2)	C2—H2B	1.091 (7)
Zn1—O5 ⁱ	2.039 (2)	Zn2—O10	2.067 (3)
Zn1—O6	2.093 (3)	Zn2—O10 ⁱⁱ	2.067 (3)
Zn1—O6 ⁱ	2.094 (3)	Zn2—O11 ⁱⁱ	2.075 (2)
Zn1—O7 ⁱ	2.173 (2)	Zn2—O11	2.075 (2)
Zn1—O7	2.173 (2)	Zn2—09	2.129 (3)
O5—H5A	0.973 (7)	Zn2—O9 ⁱⁱ	2.129 (3)
O5—H5B	0.997 (7)	O9—H9B	0.966 (6)
O6—H6A	0.981 (6)	O9—H9A	0.979 (5)
O6—H6B	0.985 (6)	O10—H10A	0.977 (8)
O7—C1	1.272 (4)	O10—H10B	0.979 (6)
O8—C1	1.240 (3)	O11—H11B	0.966 (6)
N1—H3N	1.022 (6)	O11—H11A	0.966 (6)
N1—H2N	1.027 (8)		
03—\$1—01	110.2 (3)	O8—C1—O7	126.0 (3)
O3—S1—O4	110.1 (3)	O8—C1—C2	116.2 (2)
01—S1—O4	109.5 (3)	O7—C1—C2	117.7 (2)
O3—S1—O2	109.3 (3)	H2A—C2—H2B	107.9 (7)
01—S1—O2	109.3 (3)	H2A—C2—N1	109.5 (4)
O4—S1—O2	108.4 (3)	H2B—C2—N1	109.5 (4)
$O5$ — $Zn1$ — $O5^i$	180.0	H2A—C2—C1	108.5 (5)
O5—Zn1—O6	88.59 (11)	H2B—C2—C1	109.7 (4)
05 ⁱ —Zn1—O6	91.41 (11)	O10—Zn2—O10 ⁱⁱ	180.0
O5—Zn1—O6 ⁱ	91.41 (11)	O10-Zn2-O11 ⁱⁱ	91.57 (10)
$O5^{i}$ —Zn1— $O6^{i}$	88.59 (11)	O10 ⁱⁱ —Zn2—O11 ⁱⁱ	88.43 (10)
06—Zn1—O6 ⁱ	180.0	O10—Zn2—O11	88.43 (10)
O5—Zn1—O7 ⁱ	92.01 (9)	O10 ⁱⁱ —Zn2—O11	91.57 (10)
$O5^{i}$ —Zn1— $O7^{i}$	87.99 (9)	O11 ⁱⁱ —Zn2—O11	180.0
06—Zn1—O7 ⁱ	93.38 (10)	O10—Zn2—O9	91.52 (11)
$O6^{i}$ —Zn1— $O7^{i}$	86.62 (10)	O10 ⁱⁱ —Zn2—O9	88.48 (11)
05—Zn1—07	87.99 (9)	O11 ⁱⁱ —Zn2—O9	94.16 (10)
05 ⁱ —Zn1—07	92.01 (9)	O11—Zn2—O9	85.84 (10)
06—Zn1—07	86.62 (10)	O10—Zn2—O9 ⁱⁱ	88.48 (11)
06 ⁱ —Zn1—O7	93.38 (10)	O10 ⁱⁱ —Zn2—O9 ⁱⁱ	91.52 (11)
O7 ⁱ —Zn1—O7	180.0	O11 ⁱⁱ —Zn2—O9 ⁱⁱ	85.84 (10)
H5A—O5—H5B	106.7 (6)	O11—Zn2—O9 ⁱⁱ	94.16 (10)
H6A—O6—H6B	108.1 (6)	O9—Zn2—O9 ⁱⁱ	180.00 (12)
C1—07—Zn1	128.26 (18)	H9B—O9—H9A	106.2 (6)
H3N—N1—H2N	107.6 (6)	H10A—O10—H10B	106.1 (6)

supporting information

H3N—N1—H1N	109.8 (7)	H11B—O11—H11A	111.4 (6)
H2N—N1—H1N	105.0 (6)		

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

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D····A	<i>D</i> —H… <i>A</i>
O5—H5 <i>A</i> …O4 ⁱⁱⁱ	0.973 (7)	1.793 (7)	2.755 (4)	169.0 (7)
O5—H5 <i>B</i> ···O8 ^{iv}	0.997 (7)	1.656 (8)	2.642 (4)	168.9 (6)
O6—H6 <i>A</i> ···O3 ^v	0.981 (6)	1.722 (6)	2.696 (3)	170.8 (5)
O6—H6B…O4 ^{vi}	0.985 (6)	1.751 (5)	2.729 (3)	171.8 (7)
O9—H9A···O1 ^{vi}	0.979 (5)	1.732 (5)	2.707 (3)	173.8 (6)
О9—H9 <i>B</i> …О2	0.966 (6)	1.895 (6)	2.811 (3)	157.2 (6)
O10—H10 <i>A</i> ···O3 ^v	0.977 (8)	1.740 (8)	2.713 (4)	173.0 (7)
O10—H10 <i>B</i> ····O2 ⁱⁱ	0.979 (6)	1.811 (7)	2.745 (4)	158.5 (7)
O11—H11A····O2 ^{vii}	0.966 (6)	1.772 (6)	2.726 (3)	168.6 (6)
O11—H11 <i>B</i> …O1	0.966 (6)	1.824 (6)	2.750 (3)	159.5 (7)
C2—H2A····O2 ^{vii}	1.085 (6)	2.682 (9)	3.351 (4)	119.4 (6)
C2—H2A····O5 ^{viii}	1.085 (6)	2.716 (8)	3.489 (3)	127.9 (6)
C2—H2B…O10	1.091 (7)	2.579 (8)	3.649 (4)	166.7 (7)
N1—H1 <i>N</i> ···O7 ^{viii}	1.033 (7)	1.853 (7)	2.848 (3)	160.8 (7)
N1—H2 <i>N</i> ···O4	1.027 (8)	1.961 (7)	2.877 (3)	147.0 (7)
N1—H3 <i>N</i> ···O6	1.022 (6)	2.216 (7)	3.066 (3)	139.5 (5)

Symmetry codes: (ii) -*x*+1, -*y*+1, -*z*; (iii) -*x*, -*y*+1, -*z*+1; (iv) *x*-1, *y*, *z*; (v) *x*+1, *y*-1, *z*; (vi) *x*, *y*-1, *z*; (vii) *x*+1, *y*, *z*; (viii) -*x*+1, -*y*+1, -*z*+1.