

SYNTHESIS AND CRYSTAL CHARACTERIZATION OF BIS-(GUANIDINIUM) 3,3-DIMETHYLGLUTARATE HYDRATE

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ABSTRACT

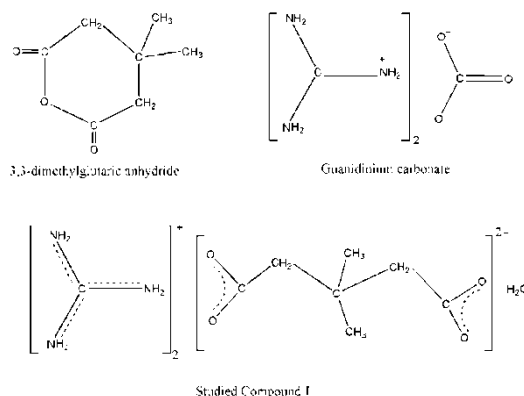
The salt $\{[\text{NH}_2\text{C}(\text{NH}_2)_2]_2\}(\text{C}_7\text{H}_{10}\text{O}_4)(\text{H}_2\text{O})(\text{I})$ was isolated from reaction between aqueous solutions of 3,3-dimethylglutaric anhydride and guanidinium carbonate. It crystallizes in the triclinic space group *P*-1 with $Z = 2$, $a = 8.7314(10)$ Å, $b = 9.1938(19)$ Å, $c = 10.9182(18)$ Å, $\alpha = 73.118(13)^\circ$, $\beta = 79.067(13)^\circ$, $\gamma = 76.353(13)^\circ$ and $V = 808.1(2)$ Å³. The asymmetric unit is comprised of a guanidinium cation, a 3,3-dimethylglutarate anion and a molecule of water. In the structure, guanidinium cations, 3,3-dimethylglutarate anions and water molecules are connected through expands N-H...O and O-H...O inter species hydrogen bonds leading to a supramolecular three dimensional structure.

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INTRODUCTION

Guanidinium with its six proton donors/acceptors may easily give rise to polydimensional topology self-assemblies. The possible propagation of multidirectional hydrogen bonding interactions makes guanidinium a useful candidate in crystal engineering. Several inorganic-organic hybrid materials with guanidinium have shown a great variety of networks (Halevas *et al.*, 2017; Thuery and Harrowfield, 2016; Han *et al.*, 2012; Galloway *et al.*, 2010; Moggach *et al.*, 2009). Videnova-Adrabinska and coworkers earlier reported some guanidinium dicarboxylates crystal structures among with guanidinium hydrogen glutarate, illustrating the prospective arrangement of guanidinium to grow two-dimensional hydrogen-bonded networks (Videnova-Adrabinska *et al.*, 2007). 3,3-dimethylglutarate and some related (glutarate substituted at the 3-position) have been used to synthesize and structurally characterize by single crystal crystallographic analysis, some compounds that exhibit diverse topologies (Nettleman *et al.*, 2010). An interesting structure with a two-fold interpenetration of layers has also been isolated and characterized (Contejean and La Duca, 2018). Diverse copper (II) compounds containing 3,3-dimethylglutarate or mixed

ligands such as 3,3-dimethylglutarate/2,2'-bipyridine were reported to be able to kill both liver and kidney cell lines at very low concentrations (Devereux *et al.*, 2006). The Dakar group has focused on glutarate and N,N,N',N'-tetramethylguanidinium, and has already published some articles (Boye *et al.*, 2007a, b; Ndiaye *et al.*, 2016a, b, 2017). Continuing our investigations on glutarate, especially on 3,3-dimethylglutarate and guanidinium, we investigated in this work the interactions between 3,3-dimethylglutaric anhydride and guanidinium carbonate which has afforded crystals of the salt **I**, namely bis (guanidinium) 3,3-dimethylglutarate hydrate (Scheme 1), whose crystal characterization is reported herein.



Scheme 1 Molecular representations of 3,3-dimethylglutaric anhydride and guanidinium carbonate reagents, and compound **I** related to this study.

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MATERIALS AND METHODS

Synthesis and crystallization

All chemicals were purchased from Aldrich Company and were used without any further purification. To 1.30 g of 3,3-dimethylglutaric anhydride, $C_7H_{10}O_3$ 99 % purity was added 25 mL hot water at 333 K. The mixture was first homogenized by stir then 25mL hot aqueous solution (333 K) of guanidinium carbonate, $\{[NH_2C(NH_2)_2]CO_3\}$ 98 % purity prepared by dissolving 1.66 g in 25 mL, was added slowly. The resulting clear solution was stirred 2h in the opened atmosphere. Submitted to a forced evaporation in an oven at 333 K, colorless prism like crystals of the salt **I** suitable for a X-ray crystallographic analysis were obtained after some days.

X-ray crystallography

A crystal of approximate dimensions $0.226 \times 0.192 \times 0.032$ mm was used for data collection. The X-ray crystallographic data were collected using a Nonius Kappa CCD diffractometer operating at $T = 293$ (2) K. Data were measured using φ and ω scans using MoK α radiation ($\lambda = 0.71073$ Å) using a collection strategy to obtain a hemisphere of unique data determined by COLLECT (Nonius, 2003). Cell parameters were determined and refined using Dirax (Duisenberg, 1992). Data were corrected for absorption correction using Eval CCD (Duisenberg *et al.*, 2003). The structure was solved using SHELXS (Sheldrick, 2015a) and the structure refined using least-squares minimization SHELXL (Sheldrick, 2015b).

Programs used for the representation of the molecular and crystal structures: Olex2 (Dolomanov *et al.*, 2009) and Mercury (Macrae *et al.*, 2008). The Crystallographic data and experimental details for structural analyses are summarized in Table 1. Selected bond lengths and angles are listed in Table 2. CCDC 1995963 contains the supplementary crystallographic data for this paper. These data can be obtained free of charge from the Cambridge Crystallographic Data Centre via www.ccdc.cam.ac.uk/data_request/cif, or by e-mailing data_request@ccdc.cam.ac.uk, or by contacting The Cambridge Crystallographic Data Centre, 12 Union Road, Cambridge CB2 1EZ, UK; fax: +44(0)1223-336033.

Table 1 Crystal data and structure refinement

Parameters	Compound I
Empirical formula	$C_7H_{10}O_4 \cdot 2(CH_6N_3) \cdot H_2O$
Formula weight	269.34
Temperature	293 (2) K
Wavelength	0.71073 Å
Crystal system	Triclinic
Space group	<i>P</i> -1
Unit cell dimensions	$a = 8.7314$ (10) Å $\alpha = 73.118$ (13)° $b = 9.1938$ (19) Å $\beta = 79.067$ (13)° $c = 10.9182$ (18) Å $\gamma = 76.353$ (13)°
Volume	808.1 (2) Å ³
Z	2
Calculated density	1.218 g cm ⁻³
Absorption coefficient	0.099 mm ⁻¹
F(000)	320
Crystal size	$0.226 \times 0.226 \times 0.226$ mm ³
Theta range for data collection	3.439–28.496°
Limiting indices	$-11 \leq h \leq 11$, $-12 \leq k \leq 12$, $-14 \leq l \leq 14$
Reflections	27442/4082

collected/unique	0.0368
R_{int}	Multi-scan
Absorption correction	Max. and min. transmission
	0.7457 and 0.7216
Refinement method	Full-matrix least-squares on F^2
Data/restraints/parameters	4082/0/189
Goodness-of-fit on F^2	1.065
Final R indices ($I > 2\sigma(I)$)	$R_1 = 0.0497$, $wR_2 = 0.1145$
R indices (all data)	$R_1 = 0.0813$, $wR_2 = 0.1018$
Largest diff. peak and hole	0.209 and -0.185e Å ⁻³

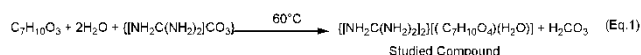
$$w = 1/[\sigma^2(F_o^2) + (0.0412P)^2 + 0.2475P] \text{ where } P = (F_o^2 + 2F_c^2)/3$$

Table 2 Selected bond lengths (Å) and angles (°)

Atom-Atom	Bond length	Atom-Atom	Bond length
O1—C3	1.2588 (17)	C7—O4	1.2446 (18)
O2—C3	1.2507 (17)	C7—O3	1.2636 (19)
C2—N6	1.322 (2)	N2—C1	1.316 (2)
C2—N4	1.3235 (19)	N1—C1	1.319 (2)
C2—N5	1.318 (2)	C1—N3	1.327 (2)
Atom-atom-atom	Angle value	Atom-atom-atom	Angle value
O1—C3—C4	118.26 (13)	O4—C7—C6	119.57 (14)
O2—C3—O1	122.63 (13)	O4—C7—O3	122.19 (14)
N6—C2—N4	119.20 (14)	N2—C1—N1	120.32 (14)
N5—C2—N6	120.28 (14)	N2—C1—N3	120.90 (15)
N5—C2—N4	120.52 (14)	N1—C1—N3	118.78 (15)

RESULTS

Compound **I** was isolated from reaction between two reagents, an anhydride and a carbonate salt, in aqueous solution. 3,3-dimethylglutaric anhydride, $C_7H_{10}O_4$ was preliminary dissolved in water at 333 K giving a clear solution. To this clear aqueous solution was added an equimolar aqueous solution of guanidinium carbonate, $\{[NH_2C(NH_2)_2]CO_3\}$. The resulting clear solution was stirred 2h. Colorless single crystals grew (Eq. 1) from the limpid solution and were characterized as **I**, $\{[NH_2C(NH_2)_2]_2\}[(C_7H_{10}O_4)(H_2O)]$.



DISCUSSION

Salt **I** crystallizes in the triclinic space group *P*-1. Its asymmetric unit depicted in Fig. 1 is comprised of two independent guanidinium cations, one 3,3-dimethylglutarate anion and a water molecule linked through strong H-bonds. The geometric parameters within the 3,3-dimethylglutarate are in accordance with those already reported in the literature (Arıcı *et al.*, 2019; Yeşilel & Wriedt, 2019; Nettleman *et al.*, 2010). The C–O bond lengths from 1.2446 (18) Å to 1.2636 (19) Å indicate a π delocalization, the slight difference is due to the involvement fashion in hydrogen bonding interactions. Angle and bond distance values for the guanidinium cation well corroborate those found in known containing guanidinium compounds (Videnova-Adrabsinska *et al.*, 2007; Peng *et al.*, 2010). The guanidinium cations C–N distances, whose C–N_{amine} and C=N_{imine} cannot be differentiated, as well as the sum of the angles at C1 and C2 carbon atoms of 360° indicate a π delocalization and a perfect planar arrangement, respectively. In a supramolecular point of view, in the crystal of **I** the 3,3-dimethylglutarates are organized into dimers *via* O–H \cdots O hydrogen bonds involving two water molecules. The hydrogen bonded dimers are strongly linked into chains by the cations (N4, N5, N6) through N–H \cdots O and N–H \cdots O_{water}. The chains are interconnected through N4–H4D \cdots O2 and N6–H6D \cdots O2

hydrogen bonds giving rise to layers represented in Fig. 2. The layers are then linked *via* the remaining cations (N1, N2, N3) and the carboxylate O atoms through N–H···O hydrogen bonds (see Table 3). Each 3,3-dimethylglutarate anion is linked to six cations and to two water molecules through N–H···O and O–H···O hydrogen bonds. The carboxylate oxygen atoms are each one involved in three H-bonds with two cations (O2 and O4) or with two cations and one molecule of water (O1 and O3), leading thus to tetrahedral environments at these O atoms. This hydrogen bonding interconnections fashion explains the slight difference of C–O bonds. The two longer C–O bonds involve two oxygen atoms (O1 and O3) that support two N–H···O and one stronger O–H···O hydrogen bonds while the shorter ones (O2 and O4) support three N–H···O hydrogen bonds. Each water molecule interacts with two O carboxylate atoms of two 3,3-dimethylglutarates and one guanidinium, through two O–H···O and one N–H···O hydrogen bonding interactions. Thus, the geometry around the O water is pyramidal. The inter species interactions exhibit five types of self-assemblies describing a 6-membered, a 8-membered and a 12-membered hydrogen bonded macrocycles involving both cations and anions, and a 8-membered and a 12-membered hydrogen bonded rings involving cations, anions and water molecules. Two different 8-membered and one 12-membered hydrogen bonded self-assemblies enable the junction between hydrogen bonded dimers, which grow into chains. The 12-membered hydrogen bonded macrocycle as well as the 6-membered ring comes from the joining chains. The rings encountered within the chains are fused, those allowing the sheets to grow as well. The extensive hydrogen bonding interactions between cations, anions and water molecules give rise to a three-dimensional supramolecular structure depicted in Fig. 3.

Table 3 Hydrogen-bond geometry (Å, °) [Symmetry codes: (i) $-x+1, -y+1, -z-3$; (ii) $-x+1, -y, -z-2$; (iii) $x+1, y-1, z$; (iv) $-x+1, -y+1, -z-2$; (v) $x, y, z+1$; (vi) $-x, -y+1, -z-2$.]

<i>D</i> –H··· <i>A</i>	<i>D</i> –H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> –H··· <i>A</i>
N6–H6C···O4 ⁱ	0.86	2.08	2.9217 (18)	167
N6–H6D···O2	0.86	2.22	2.9664 (18)	146
N4–H4C···O1 ⁱⁱ	0.86	2.07	2.9173 (19)	167
N4–H4D···O2	0.86	2.10	2.8789 (19)	151
N5–H5A···O3 ⁱ	0.86	2.01	2.8528 (18)	165
N5–H5B···O5 ⁱⁱⁱ	0.86	2.20	2.9919 (19)	153
N2–H2A···O2	0.86	2.18	3.0216 (18)	166
N2–H2B···O5 ^{iv}	0.86	2.23	3.075 (2)	167
N1–H1A···O1	0.86	1.96	2.8238 (17)	178
N1–H1B···O4 ^v	0.86	2.18	2.9070 (18)	142
N3–H3A···O3 ^{iv}	0.86	2.18	2.977 (2)	155
N3–H3B···O4 ^v	0.86	2.21	2.9246 (19)	141
O5–H5C···O1 ^{vi}	0.85 (3)	1.94 (3)	2.7623 (17)	160 (2)
O5–H5D···O3	0.85 (2)	1.83 (2)	2.6689 (18)	168 (2)

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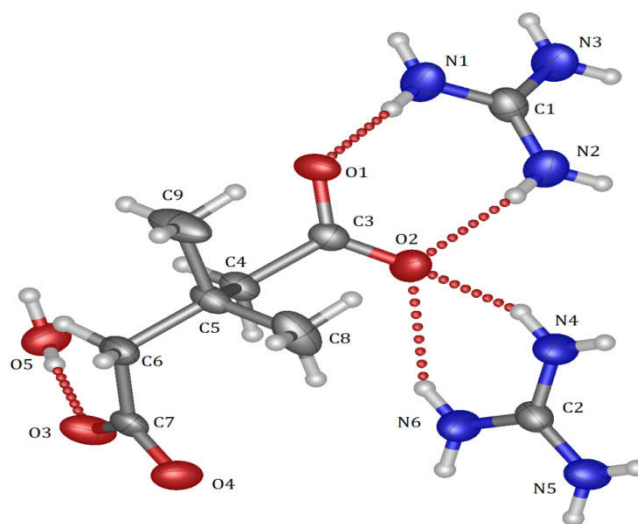


Fig 1 The molecular structure of salt I showing the atom labelling. Displacement ellipsoids are drawn at the 50% probability level.

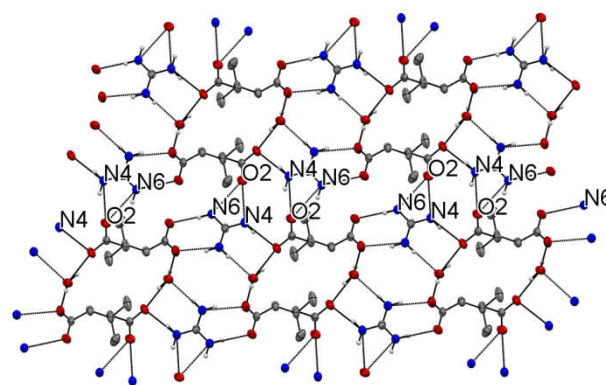


Fig 2 Partial packing diagram of 1 showing the hydrogen-bonded chains connected into sheets parallel to (0 1 1). Only hydrogen atoms involved in the interactions are shown. Symmetry identifiers are those found in Table 3.

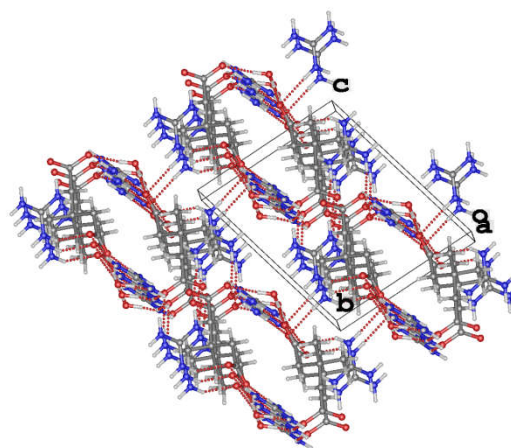


Fig 3 The partial crystal packing of salt I. Displacement ellipsoids are drawn at the 50% probability level.

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