



Synthesis and Structural Characterization of $(\text{H}_2\text{dabco})[\text{Cu}^{\text{II}}\text{Br}_4]$

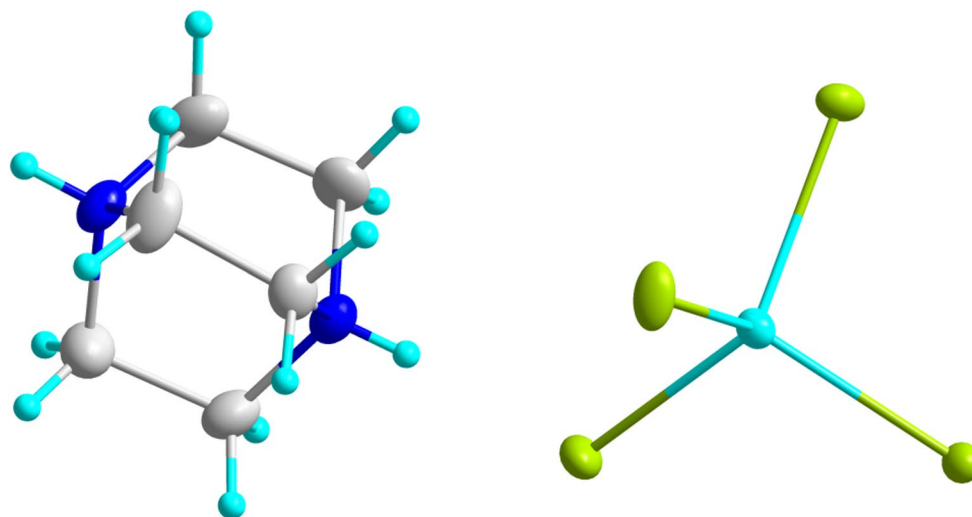
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Abstract

The $(\text{H}_2\text{dabco})[\text{Cu}^{\text{II}}\text{Br}_4]$ salt (dabco = 1,4-diazabicyclo[2.2.2]octane) was obtained and structurally characterized. The structure consists of isolated $\text{H}_2\text{dabco}^{2+}$ cations and pseudo-tetrahedral $\text{Cu}^{\text{II}}\text{Br}_4^{2-}$ anions linked via N–H...Br hydrogen bonds along the 001 direction. Because the β -angle is very close to 90° , the unit cell parameters were examined over a temperature range from 290 to 90 K. No signs of a phase transition were observed. Strong distortion of the tetrahedral geometry of the $\text{Cu}^{\text{II}}\text{Br}_4^{2-}$ anion was discussed. The Hirshfeld surface was constructed and analyzed to better understand the nature of weak interactions.

Graphical Abstract



Keywords Copper · Bromide · 1,4-diazabicyclo[2.2.2]octane

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Introduction

Hybrid organic–inorganic compounds comprising mono- or di-protonated 1,4-diazabicyclo[2.2.2]octane (dabco) moieties and cupro-halide anions are notable for both the number and diversity of cupro-halide moieties. The CCDC contains 160 entries for such compounds, from that 56 records contain tetracoordinated copper center [1]. Copper chloride derivatives with dabco alone form isolated, one-dimensional, layered [2] and three-dimensional [3] cuprochloride moieties. Up to now five cuprobromide derivatives of dabco [4–9] have been structurally studied, while including all dabco derivatives expands this list to 41 entries in the CCDC. These salts are of interest not only from an academic perspective but also because of their useful properties, particularly luminescence and reversible thermochromism [5, 6, 9–13]. In addition, effective photodegradation of tetracycline in the presence of dabco derivative cuprobromide and cupriodide has been reported [14].

In many cases, cupro-chlorides and cupro-bromides form isotypical compounds, such as $(\text{H}_2\text{dabco})[\text{Cu}_3\text{X}_5]$ ($\text{X}=\text{Cl}$ [2], Br [5]) or $(\text{H}_2\text{dabco})[\text{CuX}_4]\cdot\text{H}_2\text{O}$ ($\text{X}=\text{Cl}$ [15], Br [8]). However, a bromine analogue of $(\text{H}_2\text{dabco})[\text{CuCl}_4]$ [15, 16] has not been reported. To address this, $(\text{H}_2\text{dabco})[\text{CuBr}_4]$ was synthesized and characterized by single-crystal X-ray diffraction. The results are presented in this article.

Experimental Section

All chemicals were of commercial origin: CuBr_2 (Sigma Aldrich, 99%), HBr (Sigma Aldrich, 48% solution in water), 1,4-diazabicyclo[2.2.2]octane (Alfa Aesar, 98%), ethanol (Carlo Erba, p.a) were used without further purification.

Synthesis

Crystals of the new compound $(\text{H}_2\text{dabco})[\text{CuBr}_4]$ were obtained by reacting 0.5–0.7 mmol of CuBr_2 with 0.1 mmol of dabco in a solution containing 18 ml of ethanol and 7 ml of 48% HBr , heated to 65–77 °C and stirred with a magnetic mixer for at least 5 h. Evaporation at elevated temperature produces a small amount of well-separated single crystals of compound **1** after about 3–4 days. Prolonged exposure to the mother liquor leads, after several days, to the transformation of crystals of compound **1** into the previously studied $(\text{H}_2\text{dabco})\text{CuBr}_4\cdot\text{H}_2\text{O}$.

X-Ray Structure Determination

Preliminary data collection and structure solution yielded a suspicious monoclinic unit cell with a beta angle extremely

close to 90 degrees. Due to concerns about the accuracy of the symmetry and to identify potential twinning, a multi-temperature experiment was conducted using $\text{CuK}\alpha$ radiation. Single-crystal X-ray data for compound **1** in the temperature range of 90–290 K were collected using graphite-monochromated $\text{CuK}\alpha$ radiation on a Gemini A diffractometer equipped with an Atlas CCD detector and a Cryostream 800 cooling device [17]. The data collection was performed up to 0.79 resolution limit, with 99.9% completeness and redundancy over 8. The data were processed using the CrysAlisPro program package. Despite the beta angle being very close to 90°, Rint values of 0.08 for the monoclinic setting and 0.54 for the orthorhombic setting (data measured at 150 K) clearly confirm the monoclinic symmetry. More than 90% reflections were indexed in the above monoclinic cell. Attempts to index the remaining fewer than 200 reflections as a second domain were unsuccessful. An analytical absorption correction was applied to the data set. The structure was solved using the dual-space algorithm in SHELXT [18], and refined using SHELXL-2014 [19], both implemented in the crystallographic software Olex [20]. Hydrogen atoms bound to carbon were placed in calculated positions (AFIX commands), with their thermal parameters set to 1.2Ueq of the corresponding carbon atoms. The positions of hydrogen atoms bound to nitrogen were located on difference Fourier maps and refined freely.

At the second stage, non-spherical atom refinement was performed using the NoSpherA2 procedure implemented in the Olex2 software [21]. Theoretical wavefunctions were calculated using ORCA 6.1 software [22], with the r2SCAN method and the def2-SVP basis set. Integration accuracy was set to Normal. Water was chosen as the solvation medium.

A multi-temperature experiment was performed with 20 K steps. The crystal was cooled at a rate of 30 K per hour. After reaching the desired temperature, a 15-min waiting period was used, then complete data were collected using the same strategy at each temperature.

The structural model at each temperature was examined using the PLATON/ADDSYM procedure. No indication of higher symmetry was observed. The CheckCIF report also does not contain any alerts regarding missed or higher symmetry. The PLATON/TwinRotMat procedure also did not indicate the presence of merohedral twinning. An analysis of geometric parameters CuBr_4 moieties from CCDC was performed by Mercury software [23]. Hirshfeld surface was built and analyzed using CrystalExplorer (version 21.5) software [24].

A summary of the crystallographic data and structure refinement is provided in Table 1. CCDC 2543564 (**1**) contains the supplementary crystallographic data for this paper. These data are available free of charge from the Cambridge

Table 1 Crystallographic data, details of data collection and structure refinement parameters

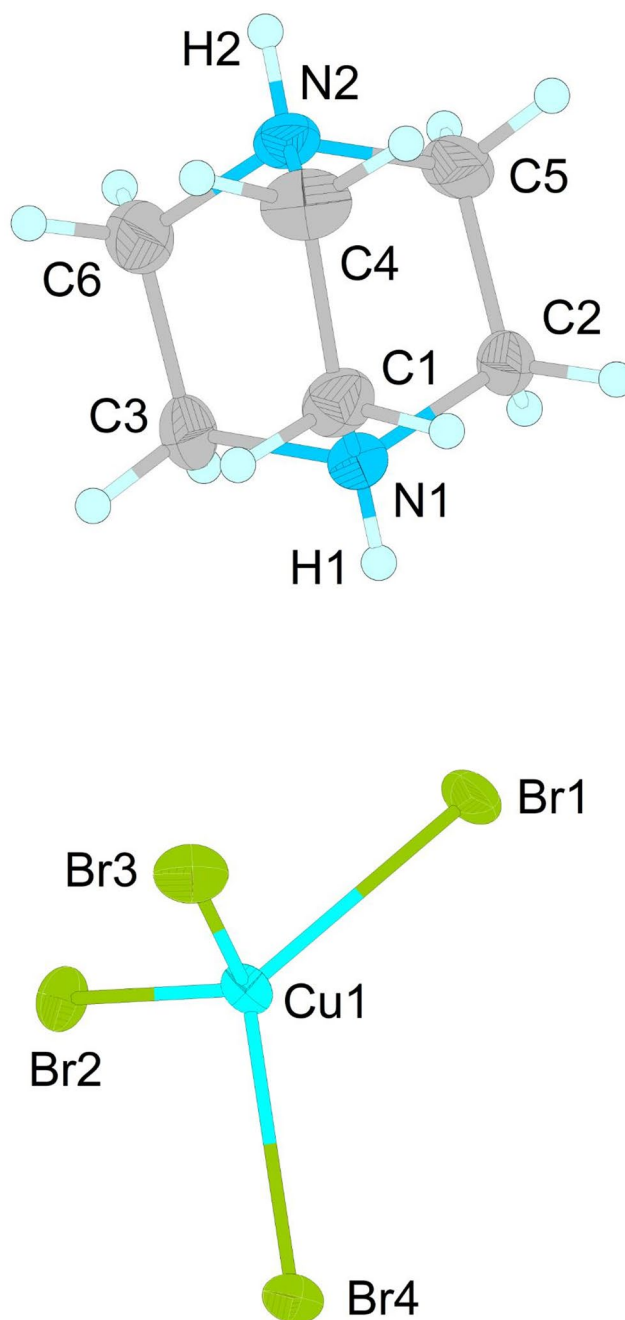
Formula	$C_6H_{14}Br_4CuN_2$	
M [g mol ⁻¹]	497.37	
T [K]	150	
Crystal system	monoclinic	
Space group	$P2_1/c$	
a [Å]	9.2820(2)	
b [Å]	7.07097(16)	
c [Å]	19.2884(5)	
β [°]	90.073(2)	
V [Å ³]	1265.95(5)	
Z	4	
F(000)	932	
ρ_{calc} [g cm ⁻³]	2.610	
Radiation, λ [Å]	CuK α , 1.54184	
μ [mm ⁻¹]	16.925	
Absorption correction	Analytical numeric absorption correction using a multifac- eted crystal model	
Crystal size, mm	0.50 × 0.37 × 0.20	
Crystal color and shape	Brown, block	
T _{max} , T _{min}	0.236, 0.039	
Rint	0.029	
Reflections all	23,758	
Independent	2642	
[I > 2 σ (I)]	2477	
	Independent atom model	NoSperA2
Goodness-of-fit on F ²	1.085	1.0651
Final R _i [I > 2 σ (I)]	0.0334	0.0324
Final R _i (all data)	0.0371	0.0362
wR ₂ [I > 2 σ (I)]	0.0851	0.0818
wR ₂ (all data)	0.0882	0.0848
Largest diff. peak and hole (e Å ⁻³)	1.201, -1.245	0.8804, -0.9774

Crystallographic Data Centre via <https://www.ccdc.cam.ac.uk/structures/>.

Results and Discussion

NoSperA2 vs IAM Refinement

The effects the IAM neglects include chemical bonding phenomena, electron lone pairs, weak interactions, and atom or molecular polarization [25]. These effects can only be described by applying a non-spherical treatment of the electron density of atoms in a crystal structure. As it is clear from the Table 1, the main improvement after NoSperA2

**Fig. 1** Crystallographically independent part of crystal structure 1. Thermal ellipsoids are drawn at 50% probability**Table 2** Comparison of unit cell parameters (H₂dabco)[CuCl₄] (203 K, [16]) and (H₂dabco)[CuBr₄] (200 K)

	Sp.gr	a (Å)	b (Å)	c (Å)	β (°)
(H ₂ dabco) [CuCl ₄]	$P2_1/c$	8.997(2)	6.920(1)	18.804(4)	90.09(1)
(H ₂ dabco) [CuBr ₄]	$P2_1/c$	9.2946(10)	7.0703(5)	19.2632(17)	90.103(7)

refinement in the case of $(\text{H}_2\text{dabco})[\text{CuBr}_4]$ is noticeable decreasing of highest peaks of residual electron density.

Crystal Structure of 1

The compound $(\text{H}_2\text{dabco})[\text{CuBr}_4]$ (**1**), consisting of $(\text{H}_2\text{dabco})^{2+}$ di-cations and pseudo-tetrahedral CuBr_4^{2-} anions (Fig. 1), is isotopic with the chloride analogue $(\text{H}_2\text{dabco})[\text{CuCl}_4]$ (Table 2) [15, 16].

Three Cu–Br distances fall within a narrow range from 2.3619(6) to 2.3724(7) Å, while the Br4–Cu1 bond is slightly elongated at 2.4029(7) Å. In the closest related compound, $(\text{H}_2\text{dabco})[\text{CuBr}_4]\cdot\text{H}_2\text{O}$, the Cu–Br distances are very similar: 2.3625(6)–2.4112(6) Å. The Br–Cu–Br angles span a broader range from 97.00(3) to 135.79(3) degrees (96.83(2)–133.95(3) degrees in $(\text{H}_2\text{dabco})[\text{CuBr}_4]\cdot\text{H}_2\text{O}$). The peculiarities of the CuBr_4 geometry will be discussed later. Cations and anions are linked by N–H...Br hydrogen bonds. Each H(N) atom exhibits a different type of bonding: the H1 center forms an ordinary hydrogen bond, whereas the H2 atom participates in a bifurcated hydrogen bond with two bromide centers from the same anion (Table 3, Fig. 2).

Table 3 Hydrogen-bond geometry (Å, °) in the structure of compound **1**

$D\cdots H\cdots A$	$D\cdots H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1–H1...Br1	0.79 (6)	2.67 (6)	3.275 (4)	135 (5)
N2–H2...Br2 ⁱ	0.84 (6)	2.76 (6)	3.346 (3)	128 (4)
N2–H2...Br4 ⁱ	0.84 (6)	2.70 (6)	3.417 (4)	144 (4)
C1–H1B...Br2 ⁱⁱ	1.11	2.78	3.684(4)	138
C2–H2A...Br3 ⁱⁱⁱ	1.11	2.89	3.754(4)	135

Symmetry code: (i) $x, -y+1/2, z-1/2$; (ii) $-x+1, -y+1, -z+1$; (iii) $x, y-1, z$

These hydrogen bonds connect cations and anions into slightly wavy chains. Weak C–H...Br hydrogen bonds ($H\cdots Br$ 2.78–2.89 Å) further associate these chains.

Two peculiarities of the crystal structure attracted our attention. First, the monoclinic angle, which is very close to 90°, prompted us to investigate this structure at different temperatures to identify any possible phase transition. Surprisingly, the monoclinic symmetry remains unchanged over a rather wide temperature range—from room temperature to 90 K (Table 4). The data at each temperature were processed and carefully examined for the absence of reflections splitting, amount of non-indexed reflections, Rint

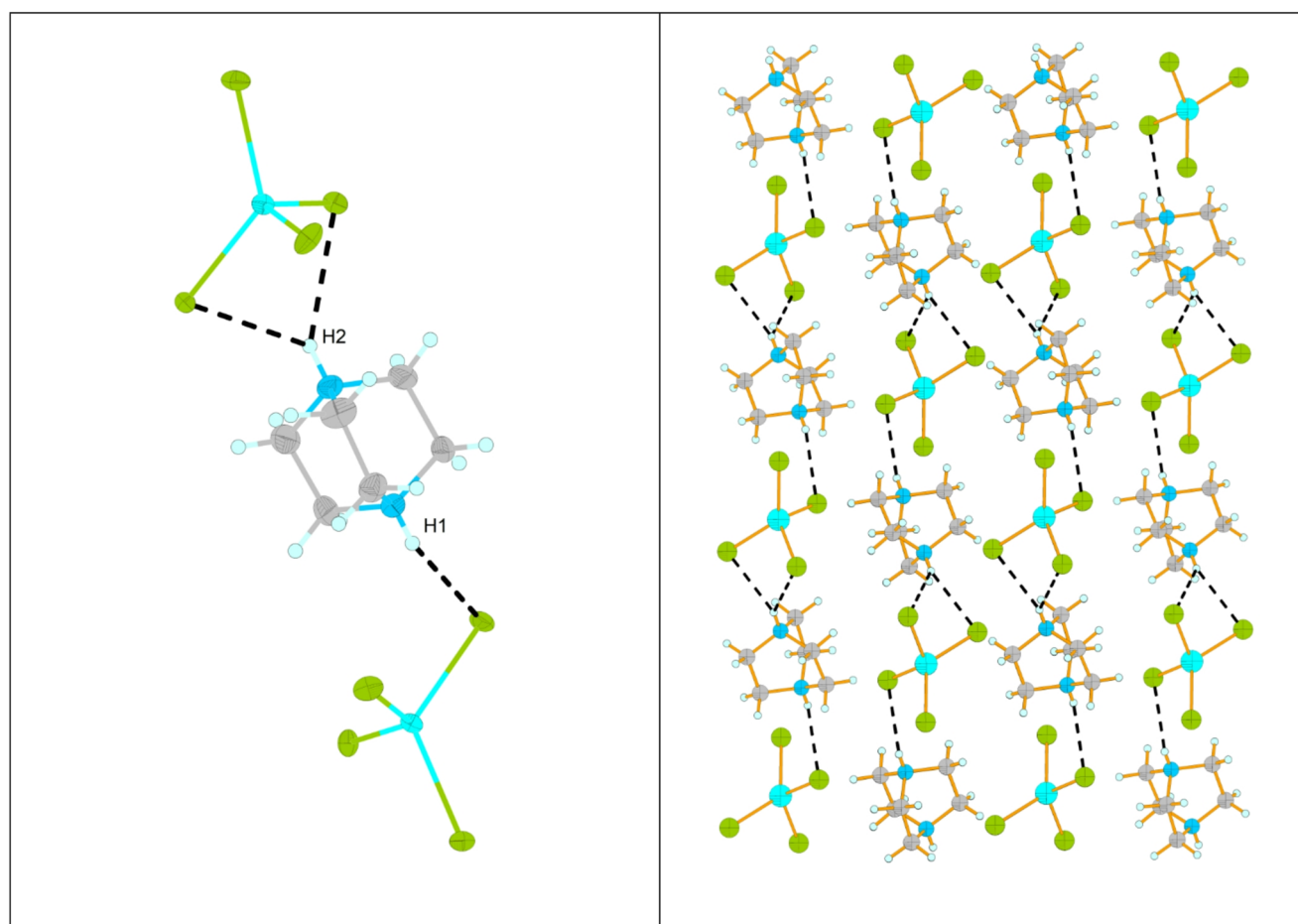
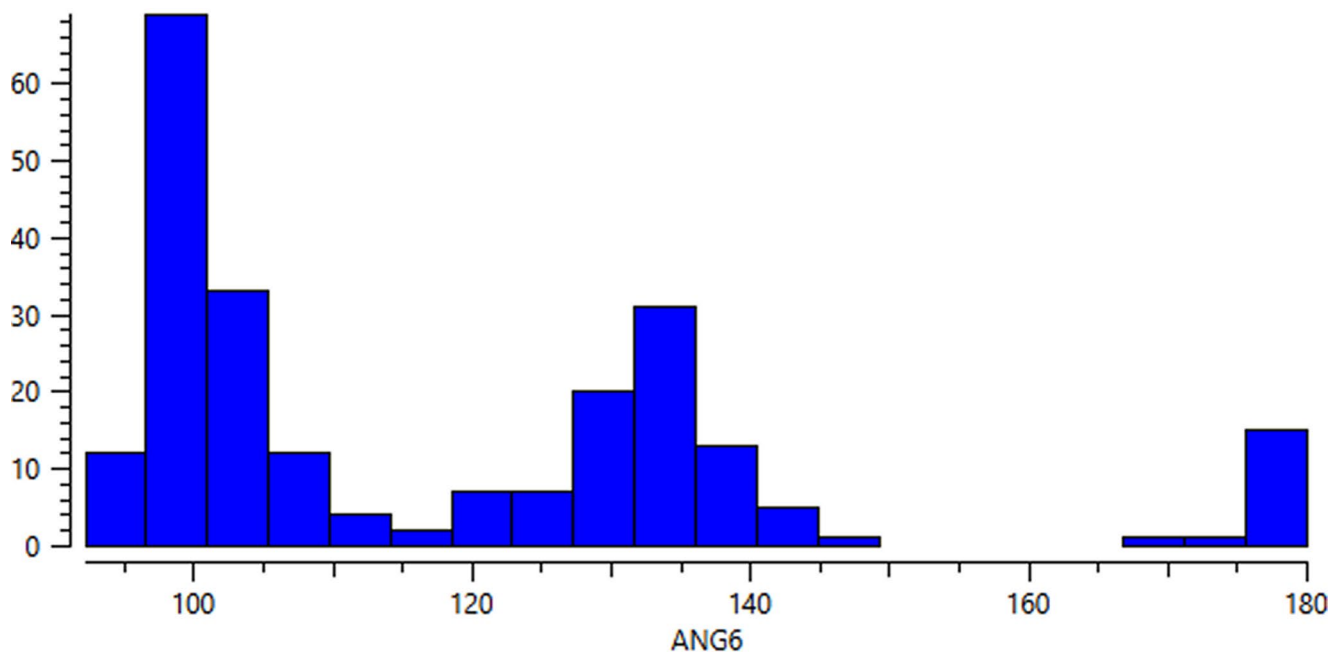


Fig. 2 Hydrogen bonds scheme (left) and packing (view along 010) of crystal structure **1**

Table 4 Unit cell parameters of compound **1** at different temperatures

T, K	<i>a</i> [Å]	<i>b</i> [Å]	<i>c</i> [Å]	β [°]	<i>V</i> [Å ³]
240	9.3184(10)	7.0829(9)	19.284(2)	90.087(9)	1272.8(3)
220	9.3057(10)	7.0761(9)	19.275(2)	90.096(9)	1269.2(3)
200	9.2946(10)	7.0703(5)	19.2632(17)	90.103(7)	1265.71(18)
180	9.2856(10)	7.0644(9)	19.262(2)	90.093(9)	1263.6(3)
160	9.2749(9)	7.0557(8)	19.255(2)	90.092(9)	1260.1(2)
140	9.2654(9)	7.0480(8)	19.249(2)	90.093(8)	1257.0(2)
120	9.2624(11)	7.0401(9)	19.262(2)	90.047(10)	1256.0(3)
100	9.2474(9)	7.0347(8)	19.244(2)	90.099(8)	1251.9(2)
90	9.2464(9)	7.0333(8)	19.248(2)	90.095(8)	1251.7(2)

**Fig. 3** Distribution of Br–Cu–Br angles in known CuBr₄ moieties

values, R- and GOOF values. Decreasing the temperature results in a monotonic contraction of each axis of the unit cell, while the beta angle remains unchanged within standard uncertainty. The linear expansion coefficients in the 100 and 010 directions are similar ($5.17 \times 10^{-5} \text{ K}^{-1}$ and $4.69 \times 10^{-5} \text{ K}^{-1}$, respectively), whereas the value for the 001 direction, $1.25 \times 10^{-5} \text{ K}^{-1}$, is noticeably smaller. This difference is likely related to the shortest Br...Br distances of 3.88 Å (slightly longer than the sum of bromine van der Waals radii), which roughly coincide with the 001 direction. Thus, the thermal contraction of compound **1** is anisotropic.

Another unusual feature observed in compound **1** and related structures is the pronounced tetrahedral distortion of the CuX₄ coordination polyhedron. The τ_4 parameter value of 0.76 indicates a significantly distorted tetrahedral environment (a value of 1.00 corresponds to a perfect tetrahedral geometry, while 0 corresponds to a perfect square-planar geometry) [26]. This value changes imperceptibly as

the temperature decreases. Corresponding value observed in (H₂dabco)[CuBr₄]·H₂O salt is 0.68. Previously described [H₂dabco][CuCl₄] and [H₂dabco][CuCl₄]·H₂O salts also contain pseudo-tetrahedral CuCl₄²⁻ anions with τ_4 parameter values of 0.74 and 0.66 respectively [15]. It should be noted that all known (H₂dabco)²⁺ cupro(II) halides exhibit similar tetrahedral distortion of the CuX₄ anion. The CCDC contains 215 entries with CuBr₄ moieties [1]. A histogram of Br–Cu–Br angles (Fig. 3) in these anions shows a wide distribution, with two statistical maxima at 100 and 133 degrees respectively.

Alternative parameter for description of tetrahedral unit distortion is a value of dihedral angle between Br1–Cu–Br2 and Br3–Cu–Br4 planes. This parameter, being 90° for ideal tetrahedral CuBr₄ geometry, decreases with tetrahedron flattening. In the case of compound **1** respective value is 72.3°. In earlier described [H₂dabco][CuCl₄] and [H₂dabco][CuCl₄]·H₂O salts these parameters are 67.83 and 61.61

Fig. 4 Hirshfeld surface for (H₂dabco)[CuBr₄]₂ fragment mapped with *d* norm over the range -0.65 to 1.7, as well as fingerprint plots for molecules resolved into H...H and H...Br interactions

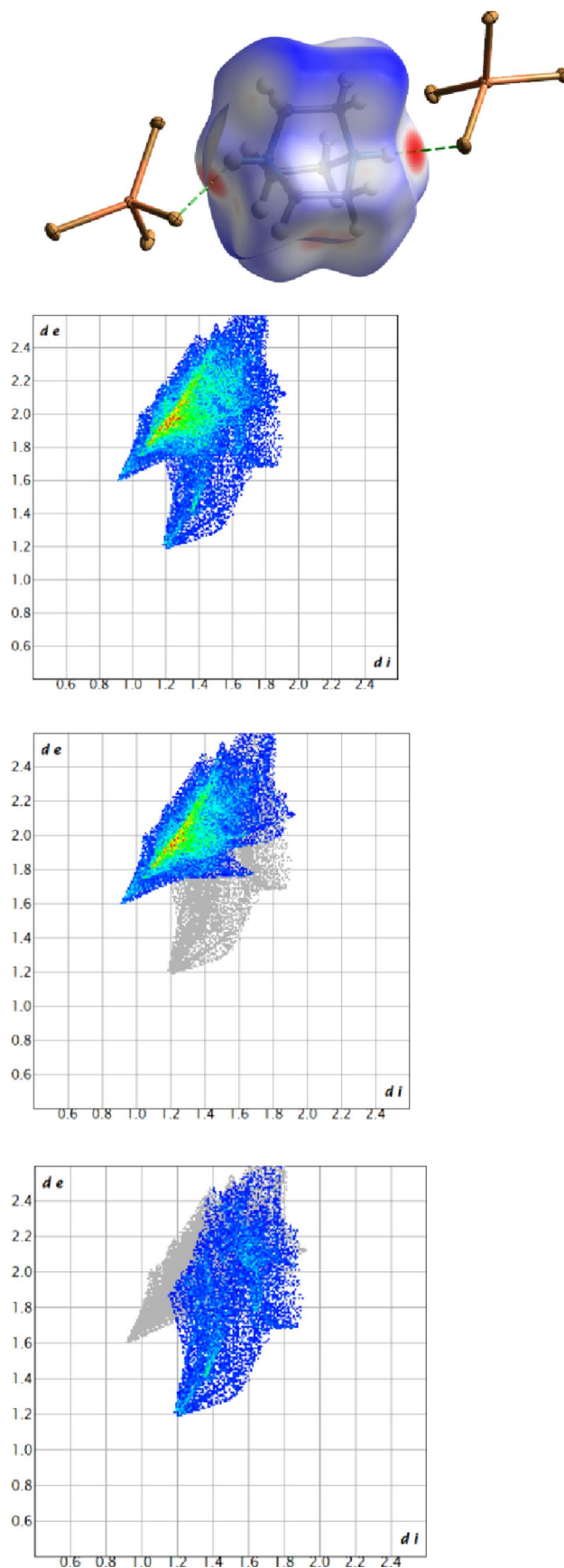
respectively. Another parameter of tetrahedral geometry distortion is the Bauer index. The angular $\Delta\theta$ parameter, calculated using the equation $\Delta\theta = \sqrt{\frac{1}{6} \sum_{i=1}^6 (i - 109.47)^2}$ (θ_i —

all six Br—Cu—Br angles within CuBr₄ moiety) of 12.02 (0 for ideal tetrahedral geometry, 43.7 for square-planar geometry) indicates also strong flattening of CuBr₄ polyhedron. A highly flattened geometry was observed, for example, in the crystal structure of [C₅H₇N₂]₂[CuBr₄] [27]. It was assumed that strong hydrogen bonds involving the bromine centers decrease the charge of the bromide ions and favor the flattened geometry of the CuBr₄²⁻ anion [28]. Another (indirect) confirmation of this assumption is the regular tetrahedral geometry of the CuBr₄²⁻ anion in [(C₂H₅)₄N]₂[CuBr₄] salt, where the possibility of hydrogen bond formation is absent [29]. As shown above, in both monohydrate salts (H₂dabco)[CuCl₄]·H₂O and (H₂dabco)[CuBr₄]·H₂O with more opportunities for hydrogen bond formation the τ_4 parameter values are noticeably lower than those in (H₂dabco)[CuCl₄] and (H₂dabco)[CuBr₄], respectively. On the other hand, the Jahn–Teller effect was cited as the reason for the distorted anion geometry in (DAP-H₂)[CuBr₄] (DAP=hexahydrodiazepine (C₅H₁₄N₂)) which contains discrete CuBr₄ tetrahedra [30]. Probably, the presence of hydrogen bonds has stronger influence on coordination polyhedron distortion than Jahn–Teller effect.

To analyze intermolecular interactions, the Hirshfeld surface was constructed for the [H₂dabco][CuBr₄]₂ fragment in **1**. Whole di-cation and two counter-ions were chosen to show complete picture of interactions around [H₂dabco] moiety. The most significant N...Br interactions between cations and anions appear as red areas on the Hirshfeld surface plot (Fig. 4). Fingerprint plots were generated to show the intermolecular surface bond distances. The main contributions to the surface area are from H...Br (65.5%) and H...H (30.3%) contacts.

Conclusions

The new (H₂dabco)[CuBr₄] salt was synthesized and structurally studied. Although the beta angle is very close to 90°, no signs of a phase transition were observed in the temperature range from 290 to 90 K. Similar to other H₂dabco cupro-halides, the Cu^{II}Br₄²⁻ anion exhibits a well-pronounced tetrahedral distortion.



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Author Contributions Conceptualisation—E.G., synthetic work—O.P., X-ray diffraction experiment —E.G., manuscript preparation—E.G. and O.P.

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Data Availability Crystallographic data are available free of charge from the Cambridge Crystallographic Data Centre via <https://www.cdc.cam.ac.uk/structures/>.

Declarations

Competing interests The authors declare no competing interests.

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