

# Synthesis and structure of tris(2-methyl-1*H*-imidazol-3-ium) 5-carboxybenzene-1,3-dicarboxylate 3,5-dicarboxybenzoate

Lina Maria Asprilla-Herrera,<sup>a</sup> Simone Techert<sup>b,c</sup> and Jose de Jesus Velazquez-Garcia<sup>c\*</sup>

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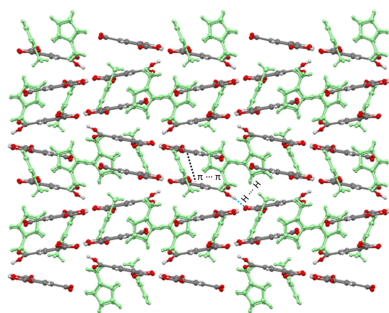
<sup>a</sup>Department of Chemistry, Faculty of Natural and Exact Sciences, Universidad del Valle, Calle 13 No. 100-00, 760042 Cali, Colombia, <sup>b</sup>Institut für Röntgenphysik, Georg-August-Universität Göttingen, Friedrich-Hund-Platz 1, 37077 Göttingen, Germany, and <sup>c</sup>Deutsches Elektronen-Synchrotron DESY, Notkestr. 85, 22607 Hamburg, Germany. \*Correspondence e-mail: jose.velazquez@desy.de

The structure of the title salt,  $3\text{C}_4\text{H}_7\text{N}_2^+ \cdot \text{C}_9\text{H}_5\text{O}_6^- \cdot \text{C}_9\text{H}_4\text{O}_6^{2-}$ , **1**, consists of three 2-methyl-imidazolium cations and both a single and a doubly deprotonated form of trimesic acid as anions. A detailed analysis of the bond lengths and angles reveals both differences and similarities between compound **1** and the previously reported 2-methyl-1*H*-imidazol-3-ium 3,5-dicarboxybenzoate structure [Baletska *et al.* (2023). *Acta Cryst. E* **79**, 1088–109], as well as the neutral counterpart of the ions. Examination of the crystal packing shows the formation of infinite chains by the anions, which, along with the cations, form zigzag planes parallel to the *ab* plane. The packing interactions are primarily driven by  $\pi$ – $\pi$  interactions and hydrogen bonding between anions.

## 1. Chemical context

Trimesic acid ( $\text{H}_3\text{btc}$ , or benzene-1,2,3-tricarboxylic acid) and 2-methylimidazole (2-mIm) are two well-known organic compounds with a wide range of applications. Trimesic acid, a planar and highly symmetrical trifunctional compound, has been used for self-assembled molecular monolayers and surface functionalization (Ha *et al.*, 2010; Lin *et al.*, 2023; Chen *et al.*, 2014; Korolkov *et al.*, 2012; MacLeod, 2019; Iancu *et al.*, 2013). Additionally,  $\text{H}_3\text{btc}$ , along with dendrimers based on it, has been employed in biomolecular delivery systems (Salamończyk, 2011; Mat Yusuf *et al.*, 2017; Emani *et al.*, 2023). On the other hand, 2-mIm, a nitrogen-containing heterocyclic organic compound, is widely used in the preparation of pharmaceuticals, photographic and photothermographic chemicals, dyes and pigments, agricultural chemicals, and in rubber production (Hachula *et al.*, 2010; Chan, 2004). Both  $\text{H}_3\text{btc}$  and 2-mIm are also well-known ligands in the syntheses of metal–organic frameworks (MOFs), such as HKUST-1 (Chui *et al.*, 1999), MIL-100 (Férey *et al.*, 2004), ZIF-8 (Park *et al.*, 2006), and ZIF-67 (Banerjee *et al.*, 2008), which have applications in gas adsorption, catalysis, and drug delivery, among others (Zhong *et al.*, 2018*a,b*; Zhao *et al.*, 2024; Huang *et al.*, 2011; Song *et al.*, 2024; Abdelhamid, 2021; Sun *et al.*, 2012).

In our previous studies, we synthesised hexaaquacobalt bis(2-methyl-1*H*-imidazol-3-ium) tetraaquabis(benzene-1,3,5-tricarboxylato- $\kappa\text{O}$ )cobalt (Velazquez-Garcia & Techert, 2022) and 2-methyl-1*H*-imidazol-3-ium 3,5-dicarboxybenzoate (Baletska *et al.*, 2023) using 2-mIm and  $\text{H}_3\text{btc}$  as organic compounds. In this work, we used the same organic compounds to synthesise the title compound, **1**.



**Table 1**

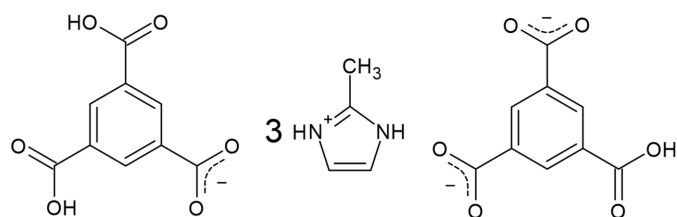
Selected bond lengths (Å), angles (°) and torsion angles (°) of the H<sub>2</sub>btc<sup>−</sup> anion in **1**.

C10–C11	1.392 (2)	C7–C12	1.391 (2)	C8–C9	1.389 (2)
C11–C12	1.393 (2)	C7–C8	1.394 (2)	C9–C10	1.394 (2)
C11–C21	1.499 (2)	C7–C22	1.492 (2)	C9–C20	1.519 (2)
O1–C21	1.214 (2)	O3–C20	1.247 (2)	O5–C22	1.218 (2)
O2–C21	1.303 (2)	O4–C20	1.258 (2)	O6–C22	1.318 (2)
C10–C11–C12	119.68 (15)	C7–C12–C11	119.86 (16)	O1–C21–O2	124.30 (16)
C9–C8–C7	120.68 (15)	C12–C7–C8	119.93 (15)	O3–C20–O4	126.76 (17)
C8–C9–C10	118.98 (16)	C11–C10–C9	120.84 (15)	O5–C22–O6	124.31 (15)
C10–C11–C21–O1	−4.4 (2)	C10–C9–C20–O4	−173.05 (15)	C10–C11–C12–C7	2.3 (2)
C12–C11–C21–O1	174.23 (16)	C8–C9–C20–O4	5.4 (2)	C12–C7–C8–C9	0.0 (2)
C10–C11–C21–O2	176.85 (15)	C12–C7–C22–O5	−177.56 (16)	C7–C8–C9–C10	1.5 (2)
C12–C11–C21–O2	−4.5 (2)	C8–C7–C22–O5	1.9 (2)	C8–C9–C10–C11	−1.2 (2)
C10–C9–C20–O3	6.0 (2)	C12–C7–C22–O6	1.2 (2)	C8–C7–C12–C11	−1.9 (2)
C8–C9–C20–O3	−175.60 (15)	C8–C7–C22–O6	−179.37 (15)	C9–C10–C11–C12	−0.7 (2)

**Table 2**

Selected bond lengths (Å), angles (°) and torsion angles (°) of the Hbtc<sup>2−</sup> anion in **1**.

C1–C6	1.393 (2)	C2–C3	1.39 (2)	C4–C5	1.392 (2)
C1–C2	1.398 (2)	C3–C4	1.391 (2)	C5–C6	1.388 (2)
C2–C19	1.504 (2)	C4–C18	1.486 (2)	C6–C17	1.510 (2)
O7–C17	1.255 (2)	O9–C18	1.214 (2)	O11–C19	1.2555 (19)
O8–C17	1.2650 (19)	O10–C18	1.338 (2)	O12–C19	1.263 (2)
C2–C3–C4	119.79 (15)	C6–C1–C2	120.38 (16)	O7–C17–O8	125.41 (15)
C6–C5–C4	120.39 (15)	C3–C2–C1	119.79 (14)	O9–C18–O10	123.24 (16)
C3–C4–C5	120.21 (16)	C5–C6–C1	119.38 (15)	O11–C19–O12	124.16 (15)
C1–C6–C17–O7	15.5 (2)	C3–C4–C18–O10	17.1 (2)	C1–C2–C3–C4	2.5 (2)
C5–C6–C17–O7	167.31 (14)	C5–C4–C18–O10	164.63 (14)	C2–C3–C4–C5	−0.5 (2)
C1–C6–C17–O8	163.84 (15)	C1–C2–C19–O11	−163.18 (15)	C2–C1–C6–C5	−0.4 (2)
C5–C6–C17–O8	−13.4 (2)	C3–C2–C19–O11	13.5 (2)	C3–C4–C5–C6	−1.5 (2)
C3–C4–C18–O9	−163.64 (16)	C1–C2–C19–O12	15.8 (2)	C4–C5–C6–C1	1.6 (2)
C5–C4–C18–O9	14.6 (2)	C3–C2–C19–O12	−167.44 (15)	C6–C1–C2–C3	−2.4 (2)



## 2. Structural commentary

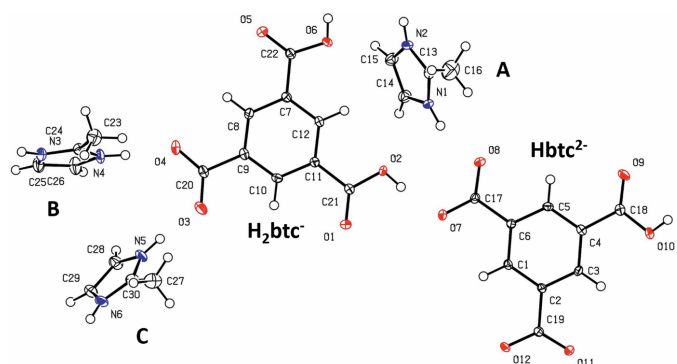
Compound **1** crystallizes with one H<sub>2</sub>btc<sup>−</sup>, one Hbtc<sup>2−</sup>, and three H<sub>2</sub>-mIm<sup>+</sup> ions in the asymmetric unit, space group *P*<sub>2</sub><sub>1</sub>/*n*. An ellipsoid plot illustrating these ionic species is shown in Fig. 1. For clarity, the three crystallographically independent cations are labelled as **A**, **B**, and **C** to facilitate their identification.

Table 1 presents selected bond distances and angles of the H<sub>2</sub>btc<sup>−</sup> ion, while Table 2 shows those for the Hbtc<sup>2−</sup> ion. The shortest bond in the H<sub>2</sub>btc<sup>−</sup> ion is between C21 and O1 at 1.214 (2) Å, while the longest is between C9 and C20 at 1.519 (2) Å. In the Hbtc<sup>2−</sup> ion, the shortest bond is C18–O9 at 1.214 (2) Å, and the longest is C6–C17 at 1.510 (2) Å.

The C–C and C–O bond lengths in the H<sub>2</sub>btc<sup>−</sup> ion range from 1.389 (2) to 1.519 (2) Å and 1.214 (2) to 1.318 (2) Å, respectively. For the Hbtc<sup>2−</sup> ion, the C–C bond lengths span 1.388 (2) to 1.510 (2) Å, while the C–O bonds range from 1.214 (2) to 1.338 (2) Å. These values are comparable to those

in the neutral H<sub>3</sub>btc molecule (Tothadi *et al.*, 2020), where the C–C bond lengths range from 1.381 (6) to 1.494 (9) Å, and C–O bonds range from 1.229 (5) to 1.303 (5) Å. They are also consistent with the bond lengths observed in the H<sub>2</sub>btc<sup>−</sup> anion reported in our previous work (Baletska *et al.*, 2023), and featuring ranges of 1.388 (2)–1.511 (2) Å for C–C bonds and 1.224 (2)–1.320 (2) Å for C–O bonds.

The C–C–C angles in H<sub>2</sub>btc<sup>−</sup> in **1** range from 118.9 (2) to 120.8 (2)°, while in the Hbtc<sup>2−</sup> ion, they fall between 119.4 (2) and 120.4 (2)°. These values are comparable to the corresponding angles in H<sub>3</sub>btc [119.0 (4)–121.1 (4)°] and H<sub>2</sub>btc<sup>−</sup> reported by Baletska *et al.* (2023) [118.9 (2)–121.4 (4)°]. The



**Figure 1**

Single-crystal X-ray structure of **1** with displacement ellipsoids drawn at the 50% probability level.

**Table 3**  
Selected bond lengths (Å), angles (°) and torsion angles (°) of the H2-mIm<sup>+</sup> cations in **1**.

A		B		C	
C13—C16	1.483 (3)	C23—C24	1.482 (3)	C27—C30	1.482 (3)
C14—C15	1.348 (2)	C25—C26	1.339 (3)	C28—C29	1.346 (3)
N1—C13	1.326 (2)	N3—C24	1.332 (2)	N5—C30	1.330 (2)
N1—C14	1.370 (2)	N3—C25	1.383 (2)	N5—C28	1.380 (2)
N2—C13	1.330 (2)	N4—C24	1.323 (2)	N6—C30	1.335 (2)
N2—C15	1.371 (2)	N4—C26	1.380 (2)	N6—C29	1.380 (2)
C13—N2—C15	109.13 (14)	C28—C29—N6	106.06 (17)	C24—N4—C26	108.48 (16)
C13—N1—C14	109.87 (15)	C29—C28—N5	107.12 (17)	C24—N3—C25	109.08 (15)
C14—C15—N2	107.24 (16)	C30—N5—C28	109.18 (16)	C25—C26—N4	107.85 (16)
C15—C14—N1	106.39 (15)	C30—N6—C29	108.92 (16)	C26—C25—N3	106.35 (17)
N1—C13—N2	107.36 (16)	N4—C24—N3	108.31 (16)	N5—C30—N6	107.91 (16)
N1—C14—C15—N2	0.1 (2)	N3—C25—C26—N4	0.0 (2)	N5—C28—C29—N6	−0.5 (2)
C13—N1—C14—C15	0.4 (2)	C24—N3—C25—C26	0.4 (2)	C30—N5—C28—C29	0.5 (2)
C14—N1—C13—N2	−0.8 (2)	C25—N3—C24—N4	0.7 (2)	C28—N5—C30—N6	−0.2 (2)
C13—N2—C15—C14	−0.5 (2)	C24—N4—C26—C25	−0.5 (2)	C30—N6—C29—C28	0.4 (2)
C15—N2—C13—N1	0.8 (2)	C26—N4—C24—N3	0.7 (2)	C29—N6—C30—N5	0.0 (2)
C14—N1—C13—C16	177.8 (2)	C26—N4—C24—C23	179.13 (18)	C28—N5—C30—C27	−179.94 (17)
C15—N2—C13—C16	−177.7 (2)	C25—N3—C24—C23	179.12 (18)	C29—N6—C30—C27	179.67 (18)

O—C—O angles in the H<sub>2</sub>btc<sup>−</sup> ion in complex **1** span 124.3 (2) to 126.8 (2)°, and in the Hbtc<sup>2−</sup> ion, they range from 123.2 (2) to 125.4 (2)°. These values are also consistent with those found in neutral H<sub>3</sub>btc [124.4 (4)–125.0 (4)°] and in H<sub>2</sub>btc<sup>−</sup> from [123.9 (2)–126.1 (2)°; Baletska *et al.*, 2023].

The main difference between the anions in **1**, the neutral H<sub>3</sub>btc molecule, and the H<sub>2</sub>btc<sup>−</sup> ion (Baletska *et al.*, 2023) lies in their torsion angles. In the H<sub>3</sub>btc molecule, the oxygen atoms are nearly coplanar with the aromatic ring, with torsion angles deviating from 0 or 180° by no more than 4.2 (4)°. H<sub>2</sub>btc<sup>−</sup> (Baletska *et al.*, 2023) shows a wider deviation range, from 4.2 (2) to 16.6 (2)°. In comparison, the H<sub>2</sub>btc<sup>−</sup> ion in **1** exhibits intermediate values, ranging from 0.6 (2) to 7.0 (2)°, whereas the Hbtc<sup>2−</sup> ion shows the largest torsion angles, ranging from 12.6 (2) to 17.1 (2)°.

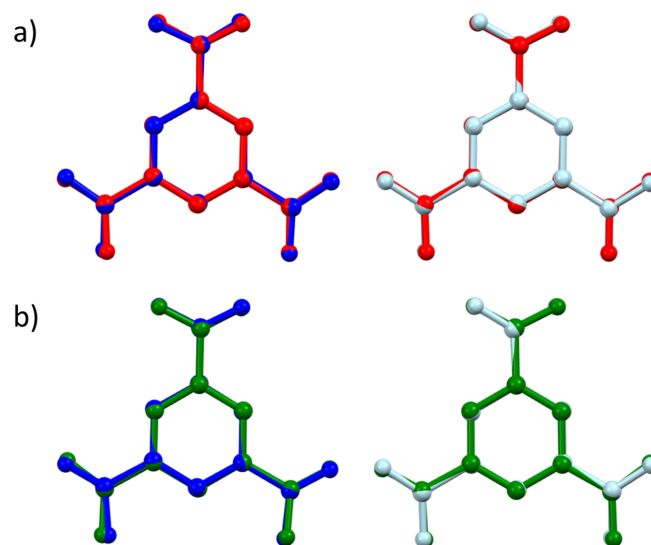
These differences are further emphasised through molecular overlays generated using *Mercury* software (Macrae *et al.*, 2020). The overlays (Fig. 2) show that the H<sub>2</sub>btc<sup>−</sup> ion in **1** resembles the neutral H<sub>3</sub>btc more closely (root-mean-square deviation, r.m.s.d. = 0.0683 Å; maximal deviation, max. d. = 0.1257 Å) than the H<sub>2</sub>btc<sup>−</sup> ion (r.m.s.d. = 0.1039 Å; max. d. = 0.2189 Å; Baletska *et al.*, 2023). On the other hand, the Hbtc<sup>2−</sup> ion in **1** shows a lower resemblance to H<sub>3</sub>btc (r.m.s.d. = 0.1856 Å; max. d. = 0.3985 Å) compared to the H<sub>2</sub>btc<sup>−</sup> ion (r.m.s.d. = 0.09 Å; max. d. = 0.2344 Å; Baletska *et al.*, 2023). Note that hydrogen atoms were excluded from the model during the overlay process.

Table 3 presents selected bond lengths, angles, and torsions for the H2-mIm<sup>+</sup> cations. The C—C bond distances fall in the range 1.339 (3)–1.483 (3) Å, while the C—N bonds vary from 1.323 (2) to 1.383 (2) Å. These values are comparable to the corresponding distances observed in the neutral 2-mIm<sup>+</sup> molecule reported by Hachuła *et al.* (2010) [C—C = 1.367 (1)–1.488 (1) Å, C—N = 1.329 (1)–1.385 (1) Å] and in the H2-mIm<sup>+</sup> ion reported by Baletska *et al.* (2023) [C—C = 1.345 (3)–1.481 (3) Å, C—N = 1.327 (2)–1.377 (2) Å].

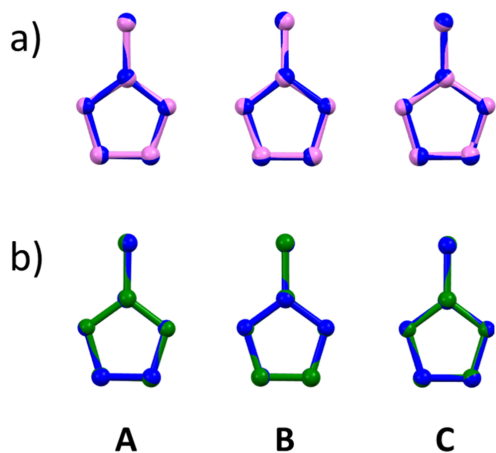
Imidazole derivatives often exhibit an asymmetry in the two endocyclic N—C bonds (Hachuła *et al.*, 2010). However, this

asymmetry is minimal in all three cations of **1**, with differences between the two N—C bond lengths of 0.001 (3), 0.003 (3), and 0.0 (3) Å for cations **A**, **B**, and **C**, respectively. These values are comparable with the asymmetry found in the H2-mIm<sup>+</sup> ion [0.008 (3) Å; Baletska *et al.*, 2023] and are significantly smaller than that reported for the neutral molecule [0.022 (1) Å]. This increased symmetry supports the idea that protonation of the imidazole reduces the disparity between the two endocyclic N—C bonds.

Protonation to an H2-mIm<sup>+</sup> ion also leads to a more symmetrical heterocyclic ring. In the H2-mIm<sup>+</sup> ion (Baletska *et al.*, 2023), this increased symmetry is observed in the C—C—N and N—C—N angles of the heterocyclic ring, which closely approach the ideal pentagon angle of 108°, with a maximum deviation of 1.6 (2)°. In contrast, the neutral 2-mIm



**Figure 2**  
Overlay plot comparing the H<sub>2</sub>btc<sup>−</sup> (dark blue) and Hbtc<sup>2−</sup> (light blue) ions in **1** with (a) H<sub>3</sub>btc (red; Tothadi *et al.* 2020) and (b) H<sub>2</sub>btc<sup>−</sup> (green; Baletska *et al.*, 2023). Hydrogen atoms are omitted for clarity.



**Figure 3**  
 Overlay plot comparing the three H<sub>2</sub>-mIm<sup>+</sup> ions (dark blue - **A**, **B** and **C**) in **1** with (a) 2-mIm (pink; Hachula et al., 2010) and (b) H<sub>2</sub>-mIm<sup>+</sup> ion (green; Baletska et al., 2023). Hydrogen atoms are omitted for clarity.

molecule shows a larger deviation of 3.4 (1)°. In compound **1**, the maximum deviations from the ideal angles of a pentagon are 1.9 (2), 1.9 (2), and 1.7 (2)° for cations **A**, **B**, and **C**, respectively. These values confirm that the protonated imidazole exhibits a more symmetrical ring structure than its neutral counterpart.

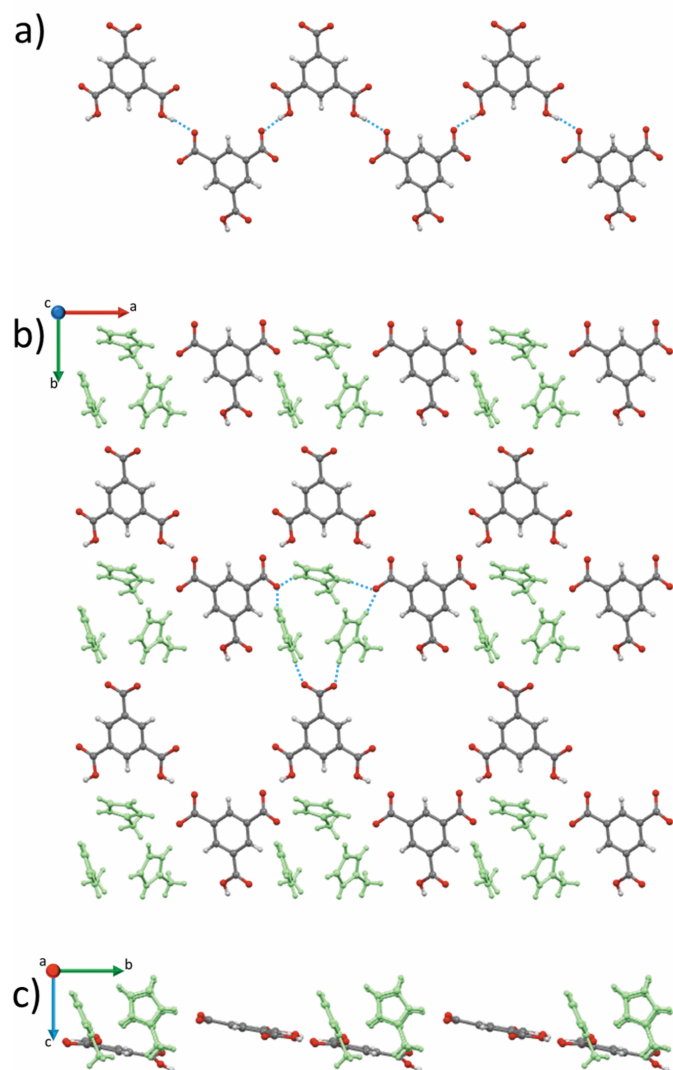
An analysis of the torsion angles in all cations in compound **1** reveals that the methyl group in cation **A** is less coplanar to the ring than in other cations. This is evident from the maximum deviation from 180° of the C–N–C–C<sup>Me</sup> torsion angles (where C<sup>Me</sup> represents the carbon from the methyl group). Cation **A** shows a deviation of 2.3 (2)°, while cations **B** and **C** exhibit smaller deviations of 0.9 (2) and 0.3 (2)°, respectively. The deviation in cation **A** is also larger than that observed in the neutral molecule [0.7 (1)°] and the H<sub>2</sub>-mIm<sup>+</sup> ion [0.5 (2)°; Baletska et al., 2023]. The root-mean-squared deviation (r. m. s. d.) and maximal deviation (max. d.) values, calculated by *Mercury* software for the molecular overlays of the three H<sub>2</sub>-mIm<sup>+</sup> cations in **1** with the H<sub>2</sub>-mIm<sup>+</sup> cation (Baletska et al., 2023) and the neutral molecule (Fig. 3), show a greater similarity between the protonated forms compared to the neutral molecule. The r. m. s. d. and max. d. values for the cations of **1** and the protonated H<sub>2</sub>-mIm<sup>+</sup> (Baletska et al., 2023) range from 0.0067 to 0.0140 Å and 0.0092 to 0.0201 Å, respectively, indicating a close resemblance. On the other hand, the values for the neutral molecule are notably higher, ranging from 0.0269 to 0.0297 Å (r.m.s.d.) and 0.0402 to 0.0474 Å (max. d.). In all cases, hydrogen atoms were omitted from the model during the overlay process.

### 3. Supramolecular features

The primary intermolecular interaction contributing to the crystal packing includes hydrogen bonds between all ions, along with  $\pi$ – $\pi$  stacking between anions. Table 4 provides a summary of the hydrogen bonds found within the compound. As shown in Fig. 4a, infinite chains are formed along the *a* axis through hydrogen bonding between H<sub>2</sub>btc<sup>−</sup> and Hbtc<sup>2−</sup>

anions. These chains are further linked, *via* hydrogen bonding, with all of the cations, forming zigzag planes parallel to the *ab* plane (Fig. 4b,c). Each plane interacts with two types of neighbouring planes: one with a parallel zigzag pattern, interacting *via*  $\pi$ – $\pi$  stacking between H<sub>2</sub>btc<sup>−</sup> and Hbtc<sup>2−</sup> ions [centroid-to-centroid distance of 3.5663 (12) Å, perpendicular distance between planes  $\sim$ 3.3 Å and offset of 1.249 Å], and another arranged in an antiparallel configuration, with the zigzag pattern running in the opposite direction. This antiparallel plane interacts *via* hydrogen bonding between Hbtc<sup>2−</sup> ions (Fig. 5). Note that the spaces observed in the planes in Fig. 4b are filled by counter-ions from the adjacent planes with a parallel zigzag pattern, ensuring no voids within compound **1**.

A graph-set analysis (Etter et al., 1990; Bernstein et al., 1995) allows a more detailed examination of the inter-



**Figure 4**  
 (a) View down the *c* axis showing an infinite chain of H<sub>2</sub>btc<sup>−</sup>–Hbtc<sup>2−</sup> anions running along the *a* axis. A plane formed by the H<sub>2</sub>-mIm<sup>+</sup> ions (green) and the H<sub>2</sub>btc<sup>−</sup>–Hbtc<sup>2−</sup> chains, view down (b) the *c* axis and (c) the *a* axis.



**Table 4**  
Hydrogen-bond geometry (Å, °).

	Graph-set descriptor	type	D—H	H···A	D···A	D—H···A
N1—H1A···O8 <sup>v</sup>	D(2)	<i>d</i>	0.86 (2)	1.911 (18)	2.737 (2)	160.8 (7)
O2—H2···O7 <sup>i</sup>	D(2)	<i>a</i>	0.96 (3)	1.57 (2)	2.5222 (19)	170.7 (17)
N2—H2A···O11 <sup>iv</sup>	D(2)	<i>e</i>	0.88 (3)	1.93 (2)	2.806 (2)	172.5 (13)
N3—H3A···O11	D(2)	<i>f</i>	0.935 (19)	1.874 (19)	2.778 (2)	162.1 (18)
N4—H4···O4 <sup>vi</sup>	D(2)	<i>g</i>	1.01 (2)	1.59 (2)	2.593 (2)	172.6 (9)
N5—H5A···O3 <sup>viii</sup>	D(2)	<i>h</i>	1.01 (2)	1.69 (2)	2.655 (2)	159.9 (5)
O6—H6···O12 <sup>ii</sup>	D(2)	<i>b</i>	0.93 (3)	1.69 (2)	2.6189 (19)	171.7 (16)
N6—H6A···O8	D(2)	<i>i</i>	0.921 (17)	1.886 (19)	2.800 (2)	170.9 (13)
O10—H10A···O12 <sup>iii</sup>	C(8)	<i>c</i>	0.93 (3)	1.71 (3)	2.6156 (18)	162 (2)
C14—H14···O1 <sup>v</sup>			0.95	2.52	3.098 (2)	119
C15—H15···O10			0.95	2.46	3.280 (2)	144
C15—H15···O5 <sup>iv</sup>			0.95	2.38	3.038 (2)	126
C25—H25···O5			0.95	2.55	3.292 (2)	135
C27—H27B···O9			0.98	2.41	3.380 (3)	168
C28—H28···O9 <sup>vii</sup>			0.95	2.39	2.990 (2)	121
C29—H29···O1			0.95	2.33	3.108 (2)	138

(i)  $1 - x, 1 - y, 1 - z$ ; (ii)  $2 - x, 1 - y, 1 - z$ ; (iii)  $\frac{3}{2} - x, \frac{1}{2} + y, \frac{1}{2} - z$ ; (iv)  $-\frac{1}{2} + x, \frac{3}{2} - y, -\frac{1}{2} + z$ ; (v)  $\frac{1}{2} + x, \frac{3}{2} - y, -\frac{1}{2} + z$ ; (vi)  $2 - x, 2 - y, 1 - z$ ; (vii)  $1 - x, 2 - y, 1 - z$ .

molecular interaction patterns within **1**. The analysis reveals that **1** contains nine motifs at the first-level graph set, including eight discrete D(2) motifs and one chain motif C(8), labelled as type *c* in Table 4. The second-level graph set (Table 5) reveals a complex network of intermolecular interactions within **1**, featuring various patterns: C<sub>2</sub><sup>2</sup>(16) >*a*<*b*, C<sub>2</sub><sup>2</sup>(12) >*d*<*e*, several D<sub>3</sub><sup>3</sup> such as >*a*>*c*<*a*, >*d*>*c*<*d*, >*e*>*c*<*e*, >*f*>*c*<*f*, >*i*>*c*<*i* and many D<sub>2</sub><sup>2</sup>, for example >*a*<*d*, >*a*<*e* and >*a*<*f*. A different pattern, rather than discrete and chain, appears in the third order graph set with formation of the rings R<sub>5</sub><sup>6</sup>(42) >*a*>*c*(*b*)<*a*<*c*<*b* (Fig. 6a) and R<sub>3</sub><sup>6</sup>(36) >*c*<*d*>*e*<*c*<*d*>*e* (Fig. 6b).

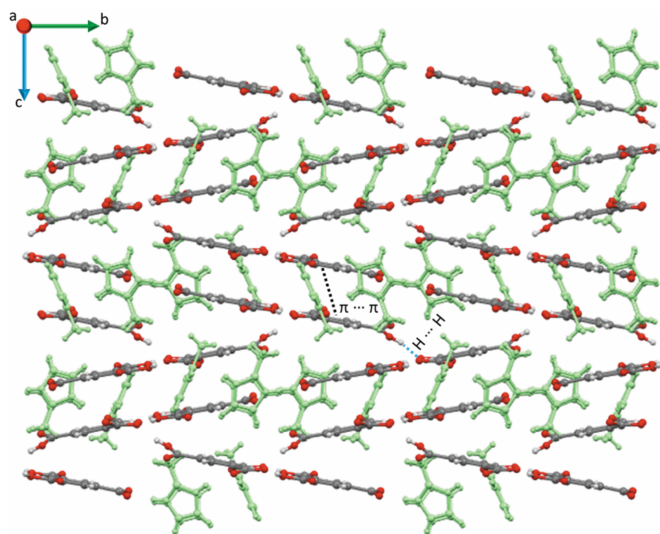
#### 4. Database survey

No reported structures of the title compound were found in the Cambridge Structural Database (CSD version 5.45, update of November 2023; Groom *et al.*, 2016). The closest to **1** is the previously mentioned structure reported under the refcode LODSUW (Baletska *et al.*, 2023).

Among the various reported structures containing the H2-mIm<sup>+</sup> cation, we highlight those with the following refcodes: BEZGEU (Dhanabal *et al.*, 2013), BOTTEK, BOTTIO, BOTTOU (Meng *et al.*, 2009), BOTTEK01, BOTTIO01, BOTTOU01, VURBUG, VURCAN, VURFAQ (Callear *et al.*,

**Table 5**  
Second- and third-level graph sets.

Second-level		Third-level	
C <sub>2</sub> <sup>2</sup> (16)	> <i>a</i> < <i>b</i>	C <sub>2</sub> <sup>3</sup> (18)	> <i>a</i> > <i>c</i> < <i>b</i>
D <sub>3</sub> <sup>3</sup> (17)	> <i>a</i> > <i>c</i> < <i>a</i>	C <sub>2</sub> <sup>3</sup> (24)	> <i>a</i> < <i>c</i> < <i>b</i>
D <sub>2</sub> <sup>2</sup> (5)	> <i>a</i> < <i>d</i>	R <sub>5</sub> <sup>6</sup> (42)	> <i>a</i> > <i>c</i> ( <i>b</i> )< <i>a</i> < <i>c</i> < <i>b</i>
D <sub>2</sub> <sup>2</sup> (9)	> <i>a</i> < <i>e</i>	D <sub>3</sub> <sup>3</sup> (17)	> <i>a</i> < <i>c</i> < <i>d</i>
D <sub>2</sub> <sup>2</sup> (9)	> <i>a</i> < <i>f</i>	D <sub>3</sub> <sup>3</sup> (17)	> <i>a</i> > <i>c</i> < <i>d</i>
D <sub>2</sub> <sup>2</sup> (10)	> <i>g</i> > <i>a</i>	D <sub>3</sub> <sup>3</sup> (13)	> <i>a</i> > <i>c</i> < <i>e</i>
D <sub>2</sub> <sup>2</sup> (10)	> <i>h</i> > <i>a</i>	D <sub>3</sub> <sup>3</sup> (17)	> <i>a</i> < <i>c</i> < <i>e</i>
D <sub>2</sub> <sup>2</sup> (5)	> <i>a</i> < <i>i</i>	D <sub>3</sub> <sup>3</sup> (13)	> <i>a</i> > <i>c</i> < <i>f</i>
D <sub>2</sub> <sup>3</sup> (11)	> <i>b</i> > <i>c</i> < <i>b</i>	D <sub>3</sub> <sup>3</sup> (17)	> <i>a</i> < <i>c</i> < <i>f</i>
D <sub>2</sub> <sup>2</sup> (9)	> <i>b</i> < <i>d</i>	D <sub>3</sub> <sup>3</sup> (17)	> <i>a</i> < <i>c</i> < <i>i</i>
D <sub>2</sub> <sup>2</sup> (5)	> <i>b</i> < <i>e</i>	D <sub>3</sub> <sup>3</sup> (17)	> <i>a</i> > <i>c</i> < <i>i</i>
D <sub>2</sub> <sup>2</sup> (5)	> <i>b</i> < <i>f</i>	D <sub>3</sub> <sup>3</sup> (13)	> <i>d</i> < <i>a</i> < <i>g</i>
D <sub>2</sub> <sup>2</sup> (10)	> <i>g</i> > <i>b</i>	D <sub>3</sub> <sup>3</sup> (13)	> <i>d</i> < <i>a</i> < <i>h</i>
D <sub>2</sub> <sup>2</sup> (10)	> <i>h</i> > <i>b</i>	D <sub>3</sub> <sup>3</sup> (17)	> <i>e</i> < <i>a</i> < <i>g</i>
D <sub>2</sub> <sup>2</sup> (9)	> <i>b</i> < <i>i</i>	D <sub>3</sub> <sup>3</sup> (17)	> <i>e</i> < <i>a</i> < <i>h</i>
D <sub>3</sub> <sup>3</sup> (17)	> <i>d</i> > <i>c</i> < <i>d</i>	C <sub>3</sub> <sup>3</sup> (20)	> <i>a</i> < <i>f</i> > <i>g</i>
D <sub>3</sub> <sup>3</sup> (13)	> <i>e</i> > <i>c</i> < <i>e</i>	D <sub>3</sub> <sup>3</sup> (17)	> <i>f</i> < <i>a</i> < <i>h</i>
D <sub>3</sub> <sup>3</sup> (13)	> <i>f</i> > <i>c</i> < <i>f</i>	D <sub>3</sub> <sup>3</sup> (13)	> <i>g</i> > <i>a</i> < <i>i</i>
D <sub>3</sub> <sup>3</sup> (17)	> <i>i</i> > <i>c</i> < <i>i</i>	C <sub>3</sub> <sup>3</sup> (16)	> <i>a</i> ( <i>i</i> ) <i>h</i>
C <sub>2</sub> <sup>2</sup> (12)	> <i>d</i> < <i>e</i>	D <sub>2</sub> <sup>2</sup> (11)	> <i>b</i> < <i>c</i> < <i>d</i>
D <sub>2</sub> <sup>2</sup> (9)	> <i>d</i> < <i>f</i>	D <sub>3</sub> <sup>3</sup> (17)	> <i>b</i> > <i>c</i> < <i>d</i>
D <sub>1</sub> <sup>2</sup> (3)	> <i>d</i> < <i>i</i>	D <sub>2</sub> <sup>2</sup> (11)	> <i>b</i> < <i>c</i> < <i>e</i>
D <sub>1</sub> <sup>2</sup> (3)	> <i>e</i> < <i>f</i>	D <sub>3</sub> <sup>3</sup> (13)	> <i>b</i> < <i>c</i> < <i>e</i>
D <sub>2</sub> <sup>2</sup> (9)	> <i>e</i> < <i>i</i>	D <sub>2</sub> <sup>2</sup> (11)	> <i>b</i> < <i>c</i> < <i>f</i>
D <sub>2</sub> <sup>2</sup> (7)	< <i>f</i> > <i>g</i>	D <sub>3</sub> <sup>3</sup> (13)	> <i>b</i> > <i>c</i> < <i>f</i>
D <sub>2</sub> <sup>2</sup> (9)	> <i>f</i> < <i>i</i>	D <sub>2</sub> <sup>2</sup> (11)	> <i>b</i> < <i>c</i> < <i>i</i>
D <sub>2</sub> <sup>2</sup> (5)	> <i>g</i> < <i>h</i>	D <sub>3</sub> <sup>3</sup> (17)	> <i>b</i> > <i>c</i> < <i>i</i>
D <sub>2</sub> <sup>2</sup> (7)	< <i>h</i> > <i>i</i>	D <sub>3</sub> <sup>3</sup> (17)	> <i>d</i> < <i>b</i> < <i>g</i>
		D <sub>3</sub> <sup>3</sup> (17)	> <i>e</i> < <i>c</i> < <i>f</i>
		D <sub>3</sub> <sup>3</sup> (13)	> <i>e</i> < <i>b</i> < <i>g</i>
		D <sub>3</sub> <sup>3</sup> (16)	> <i>b</i> < <i>f</i> > <i>g</i>
		D <sub>3</sub> <sup>3</sup> (13)	> <i>f</i> < <i>b</i> < <i>h</i>
		D <sub>3</sub> <sup>3</sup> (17)	> <i>g</i> > <i>b</i> < <i>i</i>
		D <sub>3</sub> <sup>3</sup> (20)	> <i>b</i> ( <i>i</i> ) <i>h</i>
		C <sub>3</sub> <sup>3</sup> (16)	> <i>c</i> < <i>e</i> > <i>d</i>
		C <sub>3</sub> <sup>3</sup> (20)	> <i>c</i> < <i>d</i> > <i>e</i>
		R <sub>6</sub> <sup>6</sup> (36)	> <i>c</i> < <i>d</i> > <i>e</i> < <i>c</i> < <i>d</i> > <i>e</i>
		D <sub>3</sub> <sup>3</sup> (13)	> <i>d</i> > <i>c</i> < <i>f</i>
		D <sub>3</sub> <sup>3</sup> (17)	> <i>d</i> < <i>c</i> < <i>f</i>
		D <sub>3</sub> <sup>3</sup> (17)	> <i>d</i> < <i>c</i> < <i>i</i>
		D <sub>3</sub> <sup>3</sup> (17)	> <i>d</i> > <i>c</i> < <i>i</i>
		D <sub>3</sub> <sup>3</sup> (13)	> <i>e</i> < <i>c</i> < <i>f</i>
		D <sub>3</sub> <sup>3</sup> (13)	> <i>e</i> > <i>c</i> < <i>f</i>
		D <sub>3</sub> <sup>3</sup> (17)	> <i>e</i> < <i>c</i> < <i>i</i>
		D <sub>3</sub> <sup>3</sup> (13)	> <i>f</i> < <i>c</i> < <i>i</i>
		D <sub>3</sub> <sup>3</sup> (17)	> <i>f</i> > <i>c</i> < <i>i</i>
		D <sub>3</sub> <sup>3</sup> (14)	> <i>d</i> < <i>f</i> > <i>g</i>
		D <sub>2</sub> <sup>2</sup> (8)	> <i>d</i> ( <i>i</i> ) <i>h</i>
		D <sub>2</sub> <sup>2</sup> (8)	> <i>e</i> < <i>f</i> > <i>g</i>
		D <sub>3</sub> <sup>3</sup> (14)	> <i>e</i> ( <i>i</i> ) <i>h</i>
		D <sub>3</sub> <sup>3</sup> (10)	> <i>h</i> < <i>g</i> > <i>f</i>
		D <sub>3</sub> <sup>3</sup> (14)	> <i>i</i> < <i>f</i> > <i>g</i>
		D <sub>3</sub> <sup>3</sup> (14)	> <i>f</i> ( <i>i</i> ) <i>h</i>
		D <sub>3</sub> <sup>3</sup> (10)	> <i>g</i> < <i>h</i> > <i>i</i>



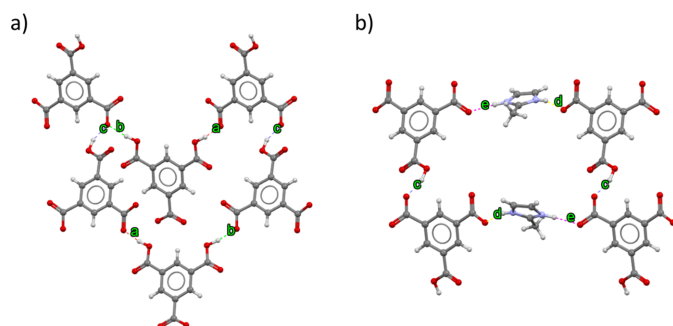
**Figure 5**  
Crystal packing in compound **1** viewed down the *a* axis showing the  $\pi$ - $\pi$  interactions and hydrogen bonding connecting the  $2H\text{-mim}^+-H\text{btc}^-$  planes that run parallel to the *ab* plane. The  $H_2\text{-mIm}^+$  ions are highlighted in green.

2010), DAMGIL (Hinokimoto *et al.*, 2021), DOWVUI (Shi *et al.*, 2014), FAMFIL, FAMFOR, FAMFUX (Zhang & Zhang, 2017), FETDAK (Aakeröy *et al.*, 2005), and HILSOL (Qu, 2007).

Organic compounds containing both  $H_2\text{btc}^-$  and  $H\text{btc}^{2-}$  were found with the refcodes: RAVPOV (Arunachalam *et al.*, 2012), SADKUE (Fan *et al.*, 2003), and TUBBAT (Melendez *et al.*, 1996). Some compounds with low resemblance to the title compound were reported under the refcodes CUMQUX (Basu *et al.*, 2009), HICSUJ (Lie *et al.*, 2013), ILELAO (Li & Li, 2016), JOCBAH (Falek *et al.*, 2019), LUBGUM, LUBHAT, LUBHEX, LUBHIB, LUBHOH, LUBHUN, LUBJAV (Singh *et al.*, 2015), SUHRAR (Rajkumar *et al.*, 2020), YOCSIT (Habib & Janiak, 2008), and WOGBED (Sosa-Rivadeneira *et al.*, 2024).

## 5. Synthesis and crystallization

To obtain the title compound, 800  $\mu\text{l}$  of an ethanolic solution of 2-mlm (1.57 *M*) was diluted in 20 ml of ethanol, followed by



**Figure 6**  
View along the *c* axis showing the formation of hydrogen-bonded ring patterns with the graph-set descriptors: (a)  $R_2^2(42)$  and (b)  $R_6^6(36)$ .

**Table 6**  
Experimental details.

Crystal data	
Chemical formula	$3C_4H_7N_2^+ \cdot C_9H_4O_6^{2-} \cdot C_9H_5O_6^-$
$M_r$	666.60
Crystal system, space group	Monoclinic, $P2_1/n$
Temperature (K)	100
<i>a</i> , <i>b</i> , <i>c</i> (Å)	14.172 (3), 15.902 (3), 14.644 (3)
$\beta$ (°)	110.46 (3)
<i>V</i> (Å <sup>3</sup> )	3092.0 (12)
<i>Z</i>	4
Radiation type	Mo <i>K</i> $\alpha$
$\mu$ (mm <sup>-1</sup> )	0.11
Crystal size (mm)	0.08 × 0.07 × 0.05
Data collection	
Diffractometer	Bruker P4
Absorption correction	Multi-scan ( <i>SADABS</i> ; Krause <i>et al.</i> , 2015)
$T_{\min}$ , $T_{\max}$	0.695, 0.746
No. of measured, independent and observed [ $I > 2\sigma(I)$ ] reflections	36740, 7127, 5287
$R_{\text{int}}$	0.052
$(\sin \theta/\lambda)_{\text{max}}$ (Å <sup>-1</sup> )	0.651
Refinement	
$R[F^2 > 2\sigma(F^2)]$ , $wR(F^2)$ , <i>S</i>	0.045, 0.119, 1.05
No. of reflections	7127
No. of parameters	457
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{\text{max}}$ , $\Delta\rho_{\text{min}}$ (e Å <sup>-3</sup> )	0.35, -0.27

Computer programs: *APEX2* and *SAINT* (Bruker, 2016), *SHELXT2018/2* (Sheldrick, 2015a), *SHELXL2018/3* (Sheldrick, 2015b) and *OLEX2* (Dolomanov *et al.*, 2009).

the addition of 1 ml of an ethanolic solution of  $H_3\text{btc}$  (0.12 *M*). The mixture was shaken gently, but no visible changes were observed after 5 min. Crystals of **1** were obtained after 24 h.

## 6. Refinement

Crystal data, data collection, and structure refinement details are summarized in Table 6. The positions of hydrogen atoms were refined with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$  for CH. Hydrogen atoms bonded to nitrogen atoms (N-H) and oxygen atoms (O-H) were treated with free refinement of bond distances and isotropic displacement parameters ( $U_{\text{iso}}$ ).

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## supporting information

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## Synthesis and structure of tris(2-methyl-1*H*-imidazol-3-ium) 5-carboxybenzene-1,3-dicarboxylate 3,5-dicarboxybenzoate

Lina Maria Asprilla-Herrera, Simone Techert and Jose de Jesus Velazquez-Garcia

### Computing details

#### Tris(2-methyl-1*H*-imidazol-3-ium) 5-carboxybenzene-1,3-dicarboxylate 3,5-dicarboxybenzoate

##### Crystal data



$M_r = 666.60$

Monoclinic,  $P2_1/n$

$a = 14.172$  (3) Å

$b = 15.902$  (3) Å

$c = 14.644$  (3) Å

$\beta = 110.46$  (3)°

$V = 3092.0$  (12) Å<sup>3</sup>

$Z = 4$

$F(000) = 1392$

$D_x = 1.432$  Mg m<sup>-3</sup>

Mo  $K\alpha$  radiation,  $\lambda = 0.71073$  Å

Cell parameters from 5589 reflections

$\theta = 2.5$ – $26.9$ °

$\mu = 0.11$  mm<sup>-1</sup>

$T = 100$  K

Irregular, clear light colourless

$0.08 \times 0.07 \times 0.05$  mm

##### Data collection

Bruker P4

diffractometer

$\omega$  scans

Absorption correction: multi-scan  
(SADABS; Krause *et al.*, 2015)

$T_{\min} = 0.695$ ,  $T_{\max} = 0.746$

36740 measured reflections

7127 independent reflections

5287 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.052$

$\theta_{\max} = 27.6$ °,  $\theta_{\min} = 2.0$ °

$h = -18$ → $18$

$k = -17$ → $20$

$l = -19$ → $19$

Standard reflections: not measured; every not measured reflections

intensity decay: not measured

##### Refinement

Refinement on  $F^2$

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.045$

$wR(F^2) = 0.119$

$S = 1.05$

7127 reflections

457 parameters

0 restraints

Primary atom site location: dual

Hydrogen site location: inferred from neighbouring sites

H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0496P)^2 + 1.5554P]$

where  $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} < 0.001$

$\Delta\rho_{\max} = 0.35$  e Å<sup>-3</sup>

$\Delta\rho_{\min} = -0.27$  e Å<sup>-3</sup>



*Special details*

**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.

**Refinement.** Hydrogen atoms bonded to nitrogen and oxygen were refined with free isotropic displacement parameters and bond lengths (AFIX 44/148)

*Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.53578 (9)	0.64000 (8)	0.61849 (11)	0.0259 (3)
O2	0.61872 (9)	0.52011 (8)	0.62429 (10)	0.0220 (3)
H2	0.5602 (19)	0.4963 (7)	0.6328 (19)	0.053 (8)*
O3	0.68318 (10)	0.90396 (8)	0.54264 (10)	0.0241 (3)
O4	0.83140 (9)	0.89827 (8)	0.52006 (10)	0.0234 (3)
O5	1.02817 (8)	0.63301 (8)	0.58809 (9)	0.0180 (3)
O6	0.95050 (9)	0.51574 (8)	0.60965 (10)	0.0190 (3)
H6	1.0140 (19)	0.4921 (7)	0.6229 (19)	0.051 (7)*
C7	0.86039 (11)	0.64249 (11)	0.58751 (11)	0.0121 (3)
C8	0.85460 (12)	0.72890 (11)	0.57047 (11)	0.0129 (3)
H8	0.910200	0.757596	0.562900	0.015*
C9	0.76831 (12)	0.77347 (11)	0.56445 (11)	0.0128 (3)
C10	0.68832 (11)	0.73080 (11)	0.57804 (11)	0.0124 (3)
H10	0.629504	0.760943	0.575260	0.015*
C11	0.69353 (11)	0.64458 (11)	0.59565 (11)	0.0119 (3)
C12	0.77922 (11)	0.60010 (11)	0.59865 (11)	0.0121 (3)
H12	0.782241	0.540944	0.608288	0.015*
C20	0.76001 (12)	0.86677 (11)	0.54084 (12)	0.0152 (4)
C21	0.60758 (12)	0.60123 (11)	0.61338 (12)	0.0140 (3)
C22	0.95475 (12)	0.59710 (11)	0.59439 (12)	0.0137 (3)
O7	0.52944 (8)	0.55672 (8)	0.35759 (9)	0.0170 (3)
O8	0.45915 (8)	0.67838 (8)	0.37705 (9)	0.0168 (3)
O9	0.60022 (9)	0.94414 (8)	0.30521 (9)	0.0191 (3)
O10	0.73044 (8)	0.93889 (8)	0.25158 (9)	0.0168 (3)
H10A	0.7044 (12)	0.9897 (17)	0.2214 (19)	0.054 (8)*
O11	0.95291 (8)	0.68773 (7)	0.35827 (8)	0.0146 (3)
O12	0.87752 (8)	0.56353 (7)	0.34820 (9)	0.0168 (3)
C1	0.69841 (11)	0.64122 (11)	0.34545 (11)	0.0110 (3)
H1	0.704556	0.582377	0.357238	0.013*
C2	0.77796 (11)	0.68585 (10)	0.33282 (11)	0.0105 (3)
C3	0.76750 (11)	0.77131 (10)	0.31188 (11)	0.0114 (3)
H3	0.820330	0.801448	0.300719	0.014*
C4	0.67944 (12)	0.81258 (11)	0.30732 (11)	0.0121 (3)
C5	0.60164 (11)	0.76843 (11)	0.32292 (11)	0.0120 (3)
H5	0.542357	0.797249	0.321299	0.014*
C6	0.61016 (11)	0.68255 (10)	0.34084 (11)	0.0108 (3)
C17	0.52617 (11)	0.63554 (11)	0.35966 (11)	0.0120 (3)

C18	0.66564 (12)	0.90441 (11)	0.28876 (12)	0.0140 (3)
C19	0.87669 (11)	0.64296 (11)	0.34702 (11)	0.0116 (3)
N1	0.81021 (10)	0.86622 (9)	-0.05350 (11)	0.0158 (3)
H1A	0.8634 (15)	0.86270 (13)	-0.0687 (4)	0.036 (6)*
N2	0.65710 (10)	0.85634 (9)	-0.06081 (11)	0.0169 (3)
H2A	0.5919 (19)	0.8450 (3)	-0.0820 (6)	0.047 (7)*
C13	0.71886 (12)	0.84187 (12)	-0.10972 (13)	0.0180 (4)
C14	0.80713 (13)	0.89782 (12)	0.03242 (13)	0.0191 (4)
H14	0.862098	0.919778	0.085000	0.023*
C15	0.71066 (13)	0.89158 (12)	0.02759 (13)	0.0202 (4)
H15	0.684631	0.908466	0.076363	0.024*
C16	0.69127 (17)	0.80710 (16)	-0.20962 (16)	0.0405 (6)
H16A	0.652114	0.848892	-0.256907	0.061*
H16B	0.650842	0.756108	-0.215027	0.061*
H16C	0.752612	0.793341	-0.222948	0.061*
N3	1.04278 (11)	0.82926 (10)	0.46433 (11)	0.0190 (3)
H3A	1.0001 (13)	0.7868 (13)	0.4287 (11)	0.054 (8)*
N4	1.12129 (11)	0.94728 (10)	0.50099 (11)	0.0194 (3)
H4	1.1453 (6)	1.0059 (16)	0.49572 (18)	0.051 (7)*
C23	1.00047 (15)	0.93620 (14)	0.33006 (14)	0.0280 (5)
H23A	0.928518	0.941715	0.319353	0.042*
H23B	1.009719	0.896478	0.282717	0.042*
H23C	1.027755	0.991192	0.321944	0.042*
C24	1.05406 (13)	0.90477 (11)	0.43016 (13)	0.0177 (4)
C25	1.10444 (14)	0.82389 (12)	0.56105 (14)	0.0231 (4)
H25	1.111071	0.777215	0.603334	0.028*
C26	1.15295 (15)	0.89749 (12)	0.58330 (14)	0.0253 (4)
H26	1.200691	0.912727	0.644847	0.030*
N5	0.38559 (11)	0.94219 (10)	0.51325 (11)	0.0194 (3)
H5A	0.3614 (7)	1.0020 (16)	0.50764 (19)	0.054 (8)*
N6	0.43032 (11)	0.82074 (10)	0.47674 (12)	0.0203 (3)
H6A	0.4446 (4)	0.7773 (13)	0.4420 (10)	0.049 (7)*
C27	0.37791 (15)	0.92675 (15)	0.34081 (14)	0.0313 (5)
H27A	0.343417	0.882455	0.294699	0.047*
H27B	0.441827	0.940409	0.332440	0.047*
H27C	0.335313	0.977045	0.328545	0.047*
C28	0.41150 (14)	0.89349 (12)	0.59641 (14)	0.0223 (4)
H28	0.410189	0.910375	0.658189	0.027*
C29	0.43899 (13)	0.81759 (12)	0.57354 (13)	0.0213 (4)
H29	0.460320	0.770711	0.616009	0.026*
C30	0.39781 (13)	0.89704 (12)	0.44175 (13)	0.0195 (4)

Atomic displacement parameters ( $\text{\AA}^2$ )

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
O1	0.0182 (6)	0.0143 (7)	0.0524 (9)	0.0038 (5)	0.0212 (6)	0.0053 (6)
O2	0.0198 (6)	0.0087 (7)	0.0457 (8)	-0.0005 (5)	0.0219 (6)	0.0028 (6)
O3	0.0297 (7)	0.0136 (7)	0.0334 (7)	0.0068 (5)	0.0167 (6)	0.0054 (6)

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O4	0.0255 (6)	0.0112 (7)	0.0367 (8)	-0.0038 (5)	0.0149 (6)	0.0028 (6)
O5	0.0141 (5)	0.0155 (7)	0.0271 (7)	0.0002 (5)	0.0105 (5)	0.0017 (5)
O6	0.0120 (5)	0.0089 (7)	0.0361 (7)	0.0021 (5)	0.0083 (5)	0.0016 (5)
C7	0.0131 (7)	0.0112 (9)	0.0118 (7)	0.0003 (6)	0.0039 (6)	-0.0007 (6)
C8	0.0134 (7)	0.0120 (9)	0.0137 (8)	-0.0024 (6)	0.0054 (6)	-0.0006 (7)
C9	0.0156 (7)	0.0103 (9)	0.0117 (7)	-0.0005 (6)	0.0037 (6)	-0.0003 (6)
C10	0.0117 (7)	0.0121 (9)	0.0132 (7)	0.0019 (6)	0.0042 (6)	-0.0017 (6)
C11	0.0124 (7)	0.0110 (9)	0.0127 (7)	0.0003 (6)	0.0048 (6)	0.0008 (6)
C12	0.0141 (7)	0.0088 (9)	0.0133 (7)	0.0001 (6)	0.0046 (6)	0.0001 (6)
C20	0.0198 (8)	0.0107 (9)	0.0156 (8)	-0.0004 (7)	0.0066 (6)	-0.0006 (7)
C21	0.0144 (7)	0.0108 (9)	0.0176 (8)	0.0004 (6)	0.0066 (6)	-0.0004 (7)
C22	0.0145 (7)	0.0114 (9)	0.0153 (8)	0.0001 (6)	0.0051 (6)	-0.0008 (7)
O7	0.0164 (6)	0.0090 (7)	0.0297 (7)	-0.0022 (5)	0.0131 (5)	-0.0004 (5)
O8	0.0149 (5)	0.0150 (7)	0.0255 (6)	0.0011 (5)	0.0134 (5)	0.0003 (5)
O9	0.0242 (6)	0.0121 (7)	0.0263 (7)	0.0059 (5)	0.0155 (5)	0.0020 (5)
O10	0.0162 (6)	0.0102 (7)	0.0252 (7)	0.0003 (5)	0.0086 (5)	0.0046 (5)
O11	0.0092 (5)	0.0118 (6)	0.0223 (6)	-0.0014 (4)	0.0050 (5)	-0.0006 (5)
O12	0.0129 (5)	0.0075 (6)	0.0304 (7)	0.0010 (5)	0.0080 (5)	-0.0024 (5)
C1	0.0130 (7)	0.0089 (9)	0.0112 (7)	-0.0005 (6)	0.0045 (6)	-0.0009 (6)
C2	0.0112 (7)	0.0086 (9)	0.0118 (7)	-0.0002 (6)	0.0041 (6)	-0.0020 (6)
C3	0.0105 (7)	0.0111 (9)	0.0131 (7)	-0.0027 (6)	0.0047 (6)	-0.0006 (6)
C4	0.0143 (7)	0.0101 (9)	0.0119 (7)	0.0012 (6)	0.0044 (6)	0.0004 (6)
C5	0.0114 (7)	0.0127 (9)	0.0130 (7)	0.0021 (6)	0.0058 (6)	-0.0001 (6)
C6	0.0117 (7)	0.0101 (9)	0.0118 (7)	-0.0004 (6)	0.0055 (6)	-0.0008 (6)
C17	0.0132 (7)	0.0126 (9)	0.0115 (7)	-0.0011 (6)	0.0060 (6)	0.0010 (6)
C18	0.0151 (7)	0.0120 (9)	0.0145 (8)	-0.0007 (6)	0.0046 (6)	0.0000 (7)
C19	0.0121 (7)	0.0108 (9)	0.0122 (7)	0.0000 (6)	0.0046 (6)	-0.0010 (6)
N1	0.0130 (6)	0.0137 (8)	0.0243 (8)	0.0002 (6)	0.0111 (6)	0.0004 (6)
N2	0.0105 (6)	0.0143 (8)	0.0263 (8)	-0.0011 (6)	0.0071 (6)	0.0015 (6)
C13	0.0175 (8)	0.0144 (10)	0.0233 (9)	0.0017 (7)	0.0088 (7)	-0.0002 (7)
C14	0.0162 (8)	0.0208 (10)	0.0194 (9)	0.0005 (7)	0.0051 (7)	-0.0016 (7)
C15	0.0200 (8)	0.0229 (11)	0.0210 (9)	0.0012 (7)	0.0114 (7)	-0.0002 (8)
C16	0.0375 (12)	0.0498 (16)	0.0311 (12)	0.0013 (11)	0.0079 (9)	-0.0155 (11)
N3	0.0185 (7)	0.0134 (8)	0.0254 (8)	-0.0034 (6)	0.0081 (6)	-0.0021 (6)
N4	0.0237 (7)	0.0116 (8)	0.0232 (8)	-0.0034 (6)	0.0087 (6)	-0.0001 (6)
C23	0.0291 (10)	0.0294 (12)	0.0227 (10)	-0.0011 (9)	0.0055 (8)	0.0049 (8)
C24	0.0187 (8)	0.0143 (10)	0.0225 (9)	-0.0010 (7)	0.0102 (7)	-0.0013 (7)
C25	0.0250 (9)	0.0177 (10)	0.0246 (9)	-0.0021 (8)	0.0061 (7)	0.0048 (8)
C26	0.0306 (10)	0.0203 (11)	0.0203 (9)	-0.0054 (8)	0.0030 (8)	0.0027 (8)
N5	0.0235 (7)	0.0125 (8)	0.0245 (8)	0.0036 (6)	0.0113 (6)	-0.0008 (6)
N6	0.0194 (7)	0.0152 (9)	0.0276 (8)	0.0025 (6)	0.0096 (6)	-0.0050 (7)
C27	0.0305 (10)	0.0392 (14)	0.0256 (10)	0.0070 (9)	0.0115 (8)	0.0062 (9)
C28	0.0266 (9)	0.0191 (11)	0.0226 (9)	0.0031 (8)	0.0105 (7)	-0.0009 (8)
C29	0.0222 (8)	0.0171 (10)	0.0247 (9)	0.0045 (7)	0.0083 (7)	0.0019 (8)
C30	0.0165 (8)	0.0196 (10)	0.0234 (9)	0.0021 (7)	0.0082 (7)	-0.0009 (8)

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*Geometric parameters (Å, °)*

O1—C21	1.214 (2)	N1—H1A	0.86 (2)
O2—H2	0.96 (3)	N1—C13	1.326 (2)
O2—C21	1.303 (2)	N1—C14	1.370 (2)
O3—C20	1.247 (2)	N2—H2A	0.89 (3)
O4—C20	1.258 (2)	N2—C13	1.330 (2)
O5—C22	1.218 (2)	N2—C15	1.371 (2)
O6—H6	0.93 (3)	C13—C16	1.483 (3)
O6—C22	1.318 (2)	C14—H14	0.9500
C7—C8	1.394 (2)	C14—C15	1.348 (2)
C7—C12	1.391 (2)	C15—H15	0.9500
C7—C22	1.492 (2)	C16—H16A	0.9800
C8—H8	0.9500	C16—H16B	0.9800
C8—C9	1.389 (2)	C16—H16C	0.9800
C9—C10	1.394 (2)	N3—H3A	0.93 (3)
C9—C20	1.519 (2)	N3—C24	1.332 (2)
C10—H10	0.9500	N3—C25	1.383 (2)
C10—C11	1.392 (2)	N4—H4	1.00 (3)
C11—C12	1.393 (2)	N4—C24	1.323 (2)
C11—C21	1.499 (2)	N4—C26	1.380 (2)
C12—H12	0.9500	C23—H23A	0.9800
O7—C17	1.255 (2)	C23—H23B	0.9800
O8—C17	1.2650 (19)	C23—H23C	0.9800
O9—C18	1.214 (2)	C23—C24	1.482 (3)
O10—H10A	0.93 (3)	C25—H25	0.9500
O10—C18	1.338 (2)	C25—C26	1.339 (3)
O11—C19	1.2555 (19)	C26—H26	0.9500
O12—C19	1.263 (2)	N5—H5A	1.01 (3)
C1—H1	0.9500	N5—C28	1.380 (2)
C1—C2	1.398 (2)	N5—C30	1.330 (2)
C1—C6	1.393 (2)	N6—H6A	0.92 (3)
C2—C3	1.390 (2)	N6—C29	1.380 (2)
C2—C19	1.504 (2)	N6—C30	1.335 (2)
C3—H3	0.9500	C27—H27A	0.9800
C3—C4	1.391 (2)	C27—H27B	0.9800
C4—C5	1.392 (2)	C27—H27C	0.9800
C4—C18	1.486 (2)	C27—C30	1.482 (3)
C5—H5	0.9500	C28—H28	0.9500
C5—C6	1.388 (2)	C28—C29	1.346 (3)
C6—C17	1.510 (2)	C29—H29	0.9500
C21—O2—H2	109.5	C13—N2—H2A	125.4
C22—O6—H6	109.5	C13—N2—C15	109.13 (14)
C8—C7—C22	118.97 (14)	C15—N2—H2A	125.4
C12—C7—C8	119.95 (15)	N1—C13—N2	107.36 (16)
C12—C7—C22	121.07 (15)	N1—C13—C16	125.80 (17)
C7—C8—H8	119.7	N2—C13—C16	126.82 (17)



C9—C8—C7	120.64 (15)	N1—C14—H14	126.8
C9—C8—H8	119.7	C15—C14—N1	106.39 (15)
C8—C9—C10	118.98 (16)	C15—C14—H14	126.8
C8—C9—C20	120.62 (15)	N2—C15—H15	126.4
C10—C9—C20	120.39 (15)	C14—C15—N2	107.24 (16)
C9—C10—H10	119.6	C14—C15—H15	126.4
C11—C10—C9	120.84 (15)	C13—C16—H16A	109.5
C11—C10—H10	119.6	C13—C16—H16B	109.5
C10—C11—C12	119.68 (15)	C13—C16—H16C	109.5
C10—C11—C21	119.43 (14)	H16A—C16—H16B	109.5
C12—C11—C21	120.88 (15)	H16A—C16—H16C	109.5
C7—C12—C11	119.86 (16)	H16B—C16—H16C	109.5
C7—C12—H12	120.1	C24—N3—H3A	125.5
C11—C12—H12	120.1	C24—N3—C25	108.95 (15)
O3—C20—O4	126.72 (17)	C25—N3—H3A	125.5
O3—C20—C9	117.21 (15)	C24—N4—H4	125.8
O4—C20—C9	116.06 (15)	C24—N4—C26	108.48 (16)
O1—C21—O2	124.30 (16)	C26—N4—H4	125.8
O1—C21—C11	121.75 (16)	H23A—C23—H23B	109.5
O2—C21—C11	113.94 (14)	H23A—C23—H23C	109.5
O5—C22—O6	124.31 (15)	H23B—C23—H23C	109.5
O5—C22—C7	122.51 (16)	C24—C23—H23A	109.5
O6—C22—C7	113.17 (14)	C24—C23—H23B	109.5
C18—O10—H10A	109.5	C24—C23—H23C	109.5
C2—C1—H1	119.8	N3—C24—C23	126.04 (17)
C6—C1—H1	119.8	N4—C24—N3	108.31 (16)
C6—C1—C2	120.38 (16)	N4—C24—C23	125.65 (17)
C1—C2—C19	120.38 (15)	N3—C25—H25	126.8
C3—C2—C1	119.79 (14)	C26—C25—N3	106.40 (17)
C3—C2—C19	119.76 (14)	C26—C25—H25	126.8
C2—C3—H3	120.1	N4—C26—H26	126.1
C2—C3—C4	119.79 (15)	C25—C26—N4	107.85 (16)
C4—C3—H3	120.1	C25—C26—H26	126.1
C3—C4—C5	120.21 (16)	C28—N5—H5A	125.5
C3—C4—C18	122.08 (15)	C30—N5—H5A	125.5
C5—C4—C18	117.69 (14)	C30—N5—C28	108.92 (16)
C4—C5—H5	119.8	C29—N6—H6A	125.6
C6—C5—C4	120.39 (15)	C30—N6—H6A	125.6
C6—C5—H5	119.8	C30—N6—C29	108.78 (16)
C1—C6—C17	120.59 (15)	H27A—C27—H27B	109.5
C5—C6—C1	119.38 (15)	H27A—C27—H27C	109.5
C5—C6—C17	119.97 (14)	H27B—C27—H27C	109.5
O7—C17—O8	125.41 (15)	C30—C27—H27A	109.5
O7—C17—C6	116.86 (14)	C30—C27—H27B	109.5
O8—C17—C6	117.73 (15)	C30—C27—H27C	109.5
O9—C18—O10	123.24 (16)	N5—C28—H28	126.4
O9—C18—C4	122.42 (15)	C29—C28—N5	107.12 (17)
O10—C18—C4	114.33 (14)	C29—C28—H28	126.4

O11—C19—O12	124.16 (14)	N6—C29—H29	126.5
O11—C19—C2	118.50 (15)	C28—C29—N6	107.06 (17)
O12—C19—C2	117.33 (14)	C28—C29—H29	126.5
C13—N1—H1A	125.1	N5—C30—N6	108.11 (16)
C13—N1—C14	109.87 (15)	N5—C30—C27	125.55 (18)
C14—N1—H1A	125.1	N6—C30—C27	126.33 (17)
C7—C8—C9—C10	1.5 (2)	C3—C4—C5—C6	-1.5 (2)
C7—C8—C9—C20	-176.91 (14)	C3—C4—C18—O9	-163.64 (16)
C8—C7—C12—C11	-1.9 (2)	C3—C4—C18—O10	17.1 (2)
C8—C7—C22—O5	1.9 (2)	C4—C5—C6—C1	1.6 (2)
C8—C7—C22—O6	-179.37 (14)	C4—C5—C6—C17	178.78 (14)
C8—C9—C10—C11	-1.2 (2)	C5—C4—C18—O9	14.6 (2)
C8—C9—C20—O3	-175.60 (15)	C5—C4—C18—O10	-164.63 (14)
C8—C9—C20—O4	5.4 (2)	C5—C6—C17—O7	167.31 (14)
C9—C10—C11—C12	-0.7 (2)	C5—C6—C17—O8	-13.4 (2)
C9—C10—C11—C21	177.89 (14)	C6—C1—C2—C3	-2.4 (2)
C10—C9—C20—O3	6.0 (2)	C6—C1—C2—C19	174.27 (14)
C10—C9—C20—O4	-173.05 (15)	C18—C4—C5—C6	-179.87 (14)
C10—C11—C12—C7	2.3 (2)	C19—C2—C3—C4	-174.28 (14)
C10—C11—C21—O1	-4.4 (2)	N1—C14—C15—N2	0.1 (2)
C10—C11—C21—O2	176.85 (15)	C13—N1—C14—C15	0.4 (2)
C12—C7—C8—C9	0.0 (2)	C13—N2—C15—C14	-0.5 (2)
C12—C7—C22—O5	-177.56 (16)	C14—N1—C13—N2	-0.8 (2)
C12—C7—C22—O6	1.2 (2)	C14—N1—C13—C16	177.78 (19)
C12—C11—C21—O1	174.23 (16)	C15—N2—C13—N1	0.8 (2)
C12—C11—C21—O2	-4.5 (2)	C15—N2—C13—C16	-177.7 (2)
C20—C9—C10—C11	177.28 (14)	N3—C25—C26—N4	0.0 (2)
C21—C11—C12—C7	-176.33 (14)	C24—N3—C25—C26	-0.4 (2)
C22—C7—C8—C9	-179.44 (14)	C24—N4—C26—C25	0.5 (2)
C22—C7—C12—C11	177.51 (14)	C25—N3—C24—N4	0.7 (2)
C1—C2—C3—C4	2.5 (2)	C25—N3—C24—C23	-179.16 (18)
C1—C2—C19—O11	-163.18 (15)	C26—N4—C24—N3	-0.7 (2)
C1—C2—C19—O12	15.8 (2)	C26—N4—C24—C23	179.13 (18)
C1—C6—C17—O7	-15.5 (2)	N5—C28—C29—N6	0.4 (2)
C1—C6—C17—O8	163.84 (14)	C28—N5—C30—N6	0.3 (2)
C2—C1—C6—C5	0.4 (2)	C28—N5—C30—C27	179.94 (17)
C2—C1—C6—C17	-176.77 (14)	C29—N6—C30—N5	0.0 (2)
C2—C3—C4—C5	-0.5 (2)	C29—N6—C30—C27	-179.67 (18)
C2—C3—C4—C18	177.76 (14)	C30—N5—C28—C29	-0.5 (2)
C3—C2—C19—O11	13.5 (2)	C30—N6—C29—C28	-0.2 (2)
C3—C2—C19—O12	-167.44 (15)		