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1,1'-Di(hydrazinocarbonylmethyl)-2,2'-biimidazole monohydrate and 1,1'-di[2-(hydrazinocarbonyl)ethyl]-2,2'-biimidazole

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The crystal structures of the title compounds, alternatively called 2,2'-(2,2'-biimidazole-1,1'-diyl)diacetohydrazide monohydrate, C₁₀H₁₄N₈O₂·H₂O, (I), and 3,3'-(2,2'-biimidazole-1,1'divl)dipropionohydrazide, C₁₂H₁₈N₈O₂, (II), respectively, have been determined. The molecules consist of half-molecule asymmetric units related by a twofold rotation in (I) and by a center of inversion in (II). The imidazole rings of both molecules crystallize in a nearly coplanar fashion [dihedral angles of 5.91 (3) and $0.0 (1)^{\circ}$ for (I) and (II), respectively]. Both planar hydrazinocarbonylalkyl substituents are essentially planar and assume the E orientation.

Comment

Compounds containing the 2,2'-biimidazole moiety have been the focus of several investigations due to their biological activity as cardiotonics (Matthews et al., 1990), antiprotozoals (Melloni et al., 1975), and enzyme active-site models (Overberger & Vorchheimer, 1963; Kirchner & Krebs, 1987). In addition to its physiological activity, the aromatic biheterocyclic structure has also been incorporated into a variety of polymer systems (Chi & Collier, 1988; Lui et al., 1988; Elmer & Collier, 1993; Lister & Collier, 1993; Barnett, 1997) in a series of attempts to imbue polymers with thermal stability, conductivity and metal-ion binding selectivity. Recently, a new class of 2,2'-biimidazole compounds, viz. the 1,1'-diester derivatives (Barnett, 1997; Barnett et al., 1996, 1997, 1999; Secondo et al., 1996), has been developed. Ready acylation of primary amines or hydrazine with these diesters proceeds with excellent yields. The hydrazide macromolecules 1,1'-di(hydrazinocarbonylmethyl)-2,2'-biimidazole monohydrate, (I), and 1,1'-di[2-(hydrazinocarbonyl)ethyl]-2,2'-biimidazole, (II),

illustrated in the Scheme below, were prepared as monomers for subsequent polymer syntheses.



As with similar unconjugated disubstituted biimidazole derivatives, the imidazole rings of (I) and (II) are coplanar, with r.m.s. deviations of 0.003 and 0.001 Å, respectively. The N1/C1/N2/C3/C2 planes and their associated symmetry partners, as shown in Figs. 1 and 2, are essentially coplanar, as they exhibit dihedral angles of 5.91 (3) and $0.0 (1)^{\circ}$ in (I) and (II), respectively. Previously reported derivatives (Barnett, 1997; Barnett et al., 1999; Secondo et al., 1996) have torsion angles along the ring-bridging C atoms of less than 1°, while the dihedral angle for the ring planes of unsubstituted 2,2'-biimidazole has been reported to be 4.6° (Cromer *et al.*, 1987). Thus, biimidazoles (I) and (II) represent the greatest disparity between imidazole-plane torsion angles for similar derivatives.





View of (I), showing the labeling of the non-H and selected H atoms. Displacement ellipsoids are shown at the 50% probability level and H atoms are drawn as small spheres of arbitrary radii.



Figure 2

View of (II), showing the labeling of the non-H atoms. Displacement ellipsoids are shown at 50% probability levels.

The N1/C4/C5 least-squares plane of (I) forms a dihedral angle of 76.7 (2)° with respect to the adjacent imidazole ring, while the same angle in (II) is 84.9 (1)° (*cf.* Figs. 1 and 2). Both dihedral angles are comparable to those observed in analogous compounds. Each hydrazinocarbonylalkyl (*R*) group adopts a zigzag conformation and is essentially planar, as the r.m.s. deviations for the non-H atoms (including N1) are 0.092 and 0.550 Å for (I) and (II), respectively. Fig. 1 shows the hydrogen bonding to the water molecules entrained in the crystal lattice of (I). The remaining bond lengths and angles do not differ significantly from 1,1'-disubstituted biimidazole structures reported previously. Selected distances and angles, together with hydrogen-bonding parameters, for (I) and (II) are given in Tables 1–4.

Extensive hydrogen-bonding networks exist in (I) and (II), as both compounds take advantage of the presence of various amine and amide H atoms and a carbonyl O atom. These are supplemented in (I) by the presence of a water of hydration. In the crystal, an intra-asymmetric unit hydrogen bond exists between atoms H4D and O2 (dashed lines in Fig. 1), while inter-asymmetric unit hydrogen bonds are present between H2A and N4, and H3A and O1, as well as between H4C and N2 (cf. Table 2). Thus, the presence of water in the crystal is an important contributor to the overall packing of this biimidazole derivative. In contrast, no strong intramolecular hydrogen bond is present in (II), although a number of intermolecular bonds, similar to those in (I), do exist (cf. Table 4). While both compounds were crystallized from water, only (I) draws on the hydrogen-bonding potential of the water in crystal formation.

Experimental

The title compounds were prepared by reacting 1,1'-di(methoxycarbonylmethyl)-2,2'-biimidazole (0.25 g, 0.898 mmol) or 1,1'-di[2-(ethoxycarbonyl)ethyl]-2,2'-biimidazole (0.25 g, 0.748 mmol) with 64% hydrazine (10 ml, 200 mmol) in a 25 ml round-bottomed flask with stirring. The mixtures were stirred for a minimum of 3 d, after which time the white powdery products were filtered off, washed with water, and crystallized by slow cooling from hot aqueous solutions. Yields were ~60 and 78% for (I) and (II), respectively. The crystal of (I) was sealed with mother liquor in a capillary to minimize possible gain or loss of water.

Table 1	
Selected geometric parameters (Å, $^{\circ}$) for (I).	

O1-C5	1.222 (4)	N3-N4	1.410 (4)
N1-C4	1.455 (3)	C1-C1 ⁱ	1.453 (5)
N3-C5	1.331 (4)	C4-C5	1.521 (4)
<i></i>		o., o., va	
C1 - N1 - C4	129.2 (2)	O1 - C5 - N3	124.0 (3)
C2-N1-C4	123.7 (3)	O1 - C5 - C4	122.4 (2)
C5-N3-N4	123.6 (3)	N3-C5-C4	113.5 (3)
N2-C1-C1 ⁱ	125.3 (3)	H4D - O2 - H2A	102
N1-C1-C1 ⁱ	123.6 (3)	$H4D - O2 - H2A^{ii}$	122
N1-C4-C5	110.8 (2)		

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

Table 2

Hydrogen-bonding geometry (Å, °) for (I).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$\begin{array}{c} 02 - H2A \cdots N4^{i} \\ N3 - H3A \cdots 01^{ii} \\ N4 - H4C \cdots N2^{iii} \\ N4 - H4D \cdots 02 \end{array}$	0.850 (5) 0.90 0.90 0.90	2.27 (2) 1.93 2.41 2.30	3.080 (3) 2.830 (3) 3.260 (4) 3.165 (4)	161 (4) 176 157 161

Symmetry codes: (i) $\frac{1}{2} + y$, 1 - x, $z - \frac{1}{4}$; (ii) 1 - y, $x - \frac{1}{2}$, $\frac{1}{4} + z$; (iii) $\frac{3}{2} - y$, x, $z - \frac{1}{4}$.

Compound (I)

Crystal data

C₁₀H₁₄N₈O₂·H₂O $M_r = 296.29$ Tetragonal, I4₁ a = 9.7451 (3) Å c = 14.1764 (6) Å V = 1346.30 (8) Å³ Z = 4 $D_x = 1.462$ Mg m⁻³ Mo Kα radiation

Data collection

Bruker P4 diffractometer $\theta/2\theta$ scans 939 measured reflections 620 independent reflections 562 reflections with $I > 2\sigma(I)$ $R_{int} = 0.015$ $\theta_{max} = 25.0^{\circ}$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.032$ $wR(F^2) = 0.077$ S = 1.14620 reflections 99 parameters H-atom parameters constrained

Compound (II)

Crystal data

 $C_{12}H_{18}N_8O_2$ $M_r = 306.33$ Monoclinic, $P2_1/c$ a = 10.0505 (5) Å b = 4.9326 (2) Å c = 15.4663 (7) Å $\beta = 106.904$ (3)° V = 733.61 (6) Å³ Z = 2

Data collection

Bruker P4 diffractometer $\theta/2\theta$ scans 1899 measured reflections 1287 independent reflections 1078 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.026$ $\theta_{\text{max}} = 25.0^{\circ}$ Cell parameters from 100 reflections $\theta = 6.4-19.7^{\circ}$ $\mu = 0.11 \text{ mm}^{-1}$ T = 293 (2) K Block cut from larger crystal, colorless $0.59 \times 0.50 \times 0.45 \text{ mm}$

 $h = -11 \rightarrow 1$ $k = -1 \rightarrow 11$ $l = -16 \rightarrow 1$ 3 standard reflections every 100 reflections intensity decay: average in $\sigma(I)$'s of 1.7%

$$\begin{split} &w = 1/[\sigma^2(F_o^{-2}) + (0.0380P)^2 \\ &+ 0.5103P] \\ &where \ P = (F_o^2 + 2F_c^{-2})/3 \\ (\Delta/\sigma)_{\rm max} < 0.001 \\ \Delta\rho_{\rm max} = 0.14 \ {\rm e} \ {\rm \mathring{A}}^{-3} \\ \Delta\rho_{\rm min} = -0.16 \ {\rm e} \ {\rm \mathring{A}}^{-3} \end{split}$$

 $D_x = 1.387 \text{ Mg m}^{-3}$ Mo K α radiation Cell parameters from 100 reflections $\theta = 7.7-19.7^{\circ}$ $\mu = 0.10 \text{ mm}^{-1}$ T = 293 (2) K Rectangular block, colorless $0.50 \times 0.36 \times 0.30 \text{ mm}$

 $h = -11 \rightarrow 1$ $k = -5 \rightarrow 1$ $l = -17 \rightarrow 18$ 3 standard reflections every 100 reflections intensity decay: average in $\sigma(I)$'s of 1.7%

Table 3					
Selected	geometric	parameters	(Å, °`) for (II).

O1-C6	1.233 (2)	C1-C1 ⁱ	1.458 (3)
N1-C4	1.464 (2)	C4-C5	1.518 (3)
N3-C6	1.328 (2)	C5-C6	1.509 (2)
N3-N4	1.417 (2)		
C1-N1-C4	129.90 (15)	N1-C4-C5	112.03 (14)
C2-N1-C4	123.6 (2)	C6-C5-C4	112.07 (15)
C6-N3-N4	123.37 (15)	O1-C6-N3	122.2 (2)
$N2-C1-C1^{i}$	125.1 (2)	O1-C6-C5	121.3 (2)
$N1-C1-C1^{i}$	123.9 (2)	N3-C6-C5	116.58 (15)

Symmetry code: (i) 1 - x, 1 - y, 2 - z.

Table 4

Hydrogen-bonding geometry (Å, $^\circ)$ for (II).

$D - H \cdots A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$N4-H4C\cdots N2^{i}$	0.90	2.47	3.367 (2)	174
$N4-H4D\cdots N4^{ii}$	0.90	2.40	3.213 (2)	151
N3-H3A···O1 ⁱⁱⁱ	0.90	2.03	2.923 (2)	176
$C3-H3\cdots O1^{iv}$	0.96	2.56	3.441 (2)	152
$C4-H4A\cdots N2^{v}$	0.96	2.32	2.960 (2)	123
$C4-H4B\cdots O1$	0.96	2.55	2.900 (2)	102
$C5-H5B\cdotsO1^{vi}$	0.96	2.54	3.471 (2)	162

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

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0496P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.040$	+ 0.2482P]
$wR(F^2) = 0.101$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.08	$(\Delta/\sigma)_{\rm max} < 0.001$
1283 reflections	$\Delta \rho_{\rm max} = 0.18 \ {\rm e} \ {\rm \AA}^{-3}$
100 parameters	$\Delta \rho_{\rm min} = -0.19 \text{ e } \text{\AA}^{-3}$
H atoms refined by a mixture of constrained and independent refinement	

For (I), the water H atom, H2A, was first located in a difference map, then refined to produce an O2–H2A distance of 0.85 Å and an H2A–O2–H2A' angle of 105°. For both structures, all other H atoms were refined as riding, with C–H bond lengths constrained to 0.96 Å and N–H to 0.90 Å.

For both compounds, data collection: P3/P4-PC Diffractometer Program (Siemens, 1991); cell refinement: P3/P4-PC Diffractometer Program; data reduction: XDISK (Siemens, 1991); program(s) used to solve structure: SHELXS86 (Sheldrick, 1990a); program(s) used to refine structure: SHELXL93 (Sheldrick, 1993); molecular graphics: SHELXTL/PC (Sheldrick, 1990b); software used to prepare material for publication: SHELXTL/PC and SHELXL93.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: SK1568). Services for accessing these data are described at the back of the journal.

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