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Key indicators

Single-crystal X-ray study
T = 293 K
Mean $\sigma(C-C)$ = 0.004 Å
R factor = 0.054
wR factor = 0.105
Data-to-parameter ratio = 17.2

For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

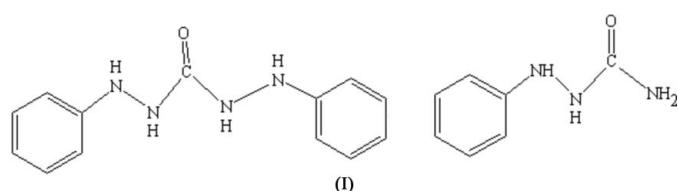
Diphenylcarbonohydrazide–phenylsemicarbazide (1/1)

In the title compound, $C_{13}H_{14}N_4O \cdot C_7H_9N_3O$, a phenylcarbonohydrazide molecule cocrystallizes with a phenylsemicarbazide molecule. In the crystal structure, extensive hydrogen-bonding and $\pi-\pi$ stacking interactions stabilize the structure.

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Comment

Diphenylcarbonohydrazide, an artificial electron-donor material, has various applications (Verma & Singh, 1995; Melis *et al.*, 1992; Prasad *et al.*, 1991; Sundari & Raghavendra, 1990; Mishra *et al.*, 1993), especially in the fields of biophysics, microbiology and analytical chemistry (El-Kabbany *et al.*, 1997). The structure of diphenylcarbonohydrazide ($C_{13}H_{14}N_4O$) has been determined previously (De Ranter *et al.*, 1979). A number of diphenylcarbonohydrazide derivatives have also been prepared (Jian *et al.*, 2003; Martynova *et al.*, 1985; Wang *et al.*, 2001; Hamuro *et al.*, 1999).



We present here the structure of the title compound, a cocrystal with a 1-phenylsemicarbazide, molecule 1, and a diphenylcarbonohydrazide, molecule 2 (Fig. 1 and Table 1). In molecule 2, ring 2 (C8–C13) is approximately parallel to ring 3 (C15–C20), with a dihedral angle of 12.64 (13)°. Ring 1, from molecule 1, makes dihedral angles of 49.53 (6) and 59.83 (6)°, respectively, with rings 2 and 3.

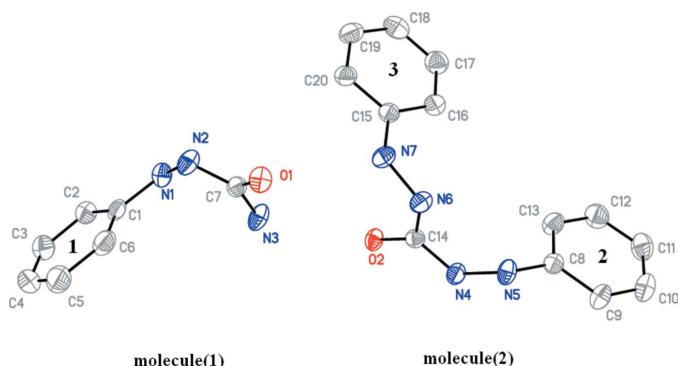


Figure 1

The asymmetric unit of (I), showing the atom-labeling scheme. Displacement ellipsoids are shown at the 50% probability level. All H atoms have been omitted for clarity.

In the crystal structure, molecules form a network structure through intermolecular N—H \cdots O hydrogen bonds (Fig. 2 and Table 2). Moreover, there are offset face-to-face π — π stacking interactions in the crystal structure, with centroid-centroid distances of 3.584–3.700 Å (Fig. 2). These interactions increase the stability of the structure.

Experimental

A mixture of $(\text{PhNNNH}_2)_2\text{C=O}$ (diphenylcarbonohydrazide, 0.0605 g, 0.25 mmol), $(\text{PhNNNH})(\text{NH}_2)\text{C=O}$ (phenylsemicarbazide, 0.0378 g, 0.25 mmol), $\text{Co}(\text{OAc})_2$ (0.0443 g, 0.25 mmol) and water (12 ml) was sealed in a 23 ml Teflon-lined stainless steel autoclave to approximately 60% of capacity. The resulting mixture was heated at a rate of *ca* 100 K h $^{-1}$ to 413 K and held at this temperature for 5 d. Subsequently, the autoclave was cooled at a rate of *ca* 3 K h $^{-1}$ to room temperature. The resulting colorless plates were filtered off, washed with water and dried at ambient temperature.

Crystal data

$\text{C}_{13}\text{H}_{14}\text{N}_4\text{O}\cdot\text{C}_7\text{H}_9\text{N}_3\text{O}$	$Z = 4$
$M_f = 393.45$	$D_x = 1.308 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/n$	$\text{Mo K}\alpha$ radiation
$a = 14.0097 (9) \text{ \AA}$	$\mu = 0.09 \text{ mm}^{-1}$
$b = 5.8841 (3) \text{ \AA}$	$T = 293 (2) \text{ K}$
$c = 24.3124 (19) \text{ \AA}$	Plate, colorless
$\beta = 94.273 (3)^\circ$	$0.32 \times 0.17 \times 0.07 \text{ mm}$
$V = 1998.6 (2) \text{ \AA}^3$	

Data collection

Rigaku Weissenberg IP diffractometer	18299 measured reflections
ω scans	4516 independent reflections
Absorption correction: multi-scan (<i>TEXSAN</i> ; Molecular Structure Corporation, 1998)	1815 reflections with $I > 2\sigma(I)$
$R_{\text{int}} = 0.059$	$R_{\text{int}} = 0.059$
$T_{\min} = 0.836$, $T_{\max} = 0.994$	$\theta_{\max} = 27.5^\circ$

Refinement

Refinement on F^2	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.054$	$w = 1/[\sigma^2(F_o^2) + (0.0229P)^2]$
$wR(F^2) = 0.105$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 0.97$	$(\Delta/\sigma)_{\text{max}} < 0.001$
4516 reflections	$\Delta\rho_{\text{max}} = 0.35 \text{ e \AA}^{-3}$
262 parameters	$\Delta\rho_{\text{min}} = -0.38 \text{ e \AA}^{-3}$

Table 1
Selected geometric parameters (Å, °).

O1—C7	1.248 (3)	N4—N5	1.406 (3)
O2—C14	1.257 (3)	N5—C8	1.421 (3)
N1—N2	1.391 (3)	N6—C14	1.345 (3)
N1—C1	1.419 (3)	N6—N7	1.398 (3)
N2—C7	1.351 (3)	N7—C15	1.408 (3)
N3—C7	1.341 (3)	C1—C6	1.372 (4)
N4—C14	1.337 (3)	C15—C20	1.397 (3)
N2—N1—C1	116.9 (2)	O1—C7—N2	121.2 (2)
C7—N2—N1	121.67 (19)	N3—C7—N2	116.7 (2)
C14—N4—N5	121.88 (19)	C13—C8—C9	118.9 (2)
N4—N5—C8	116.27 (19)	C13—C8—N5	121.6 (2)
C14—N6—N7	122.8 (2)	C9—C8—N5	119.6 (2)
N6—N7—C15	117.85 (18)	O2—C14—N4	120.5 (2)
C6—C1—C2	120.0 (3)	O2—C14—N6	122.7 (2)
C6—C1—N1	122.3 (3)	N4—C14—N6	116.8 (2)
C2—C1—N1	117.7 (2)	C16—C15—C20	119.2 (3)
O1—C7—N3	122.1 (2)	C20—C15—N7	117.8 (2)

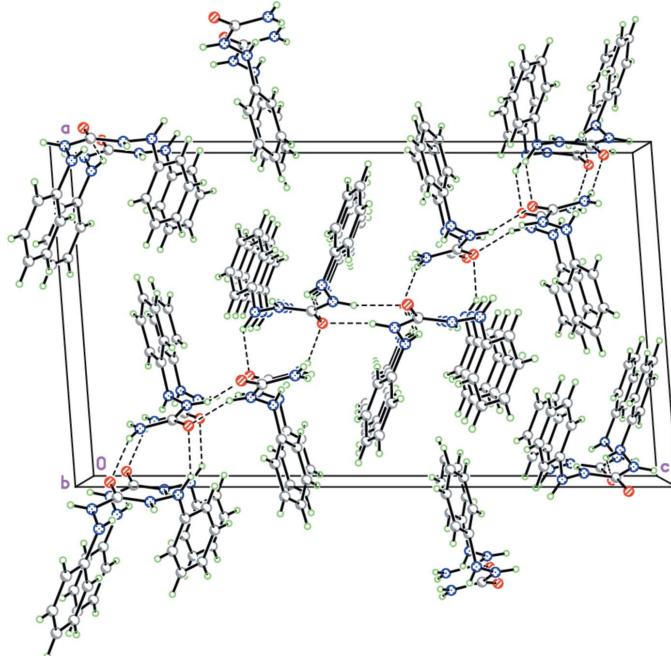


Figure 2

Packing diagram of (I), showing the π — π stacking along the b axis. Hydrogen bonds are drawn as dashed lines.

Table 2
Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N3—H3B \cdots O2	0.86	2.15	2.952 (2)	154
N7—H7A \cdots O1	0.86	2.35	2.973 (3)	129
N1—H1A \cdots O1 ⁱ	0.86	2.61	3.098 (3)	117
N2—H2B \cdots O1 ⁱⁱ	0.86	2.11	2.899 (2)	153
N4—H4B \cdots O2 ⁱⁱⁱ	0.86	2.10	2.946 (2)	168
N5—H5B \cdots O2 ^{iv}	0.86	2.48	3.079 (3)	128

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

All H atoms were positioned geometrically and allowed to ride on their parent C or N atoms, with aromatic C—H = 0.93 Å and N—H = 0.86 Å, and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$.

Data collection: *TEXSAN* (Molecular Structure Corporation, 1998); cell refinement: *TEXSAN*; data reduction: *TEXSAN*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL/PC* (Sheldrick, 1993); software used to prepare material for publication: *SHELXL97*.

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