

N-Acryloyl glycinamide

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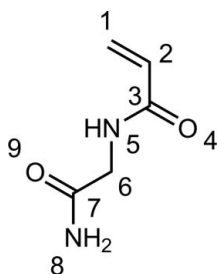
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 Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(C-C) = 0.002$ Å;
 R factor = 0.032; wR factor = 0.085; data-to-parameter ratio = 11.8.

The molecule of the title compound [systematic name: *N*-(carbamoylmethyl)prop-2-enamide], $C_5H_8N_2O_2$, which can be radically polymerized to polymers with thermoresponsive behavior in aqueous solution, consists of linked essentially planar acrylamide and amide segments [maximum deviations = 0.054 (1) and 0.009 (1) Å] with an angle of 81.36 (7)° between their mean planes. In the crystal, $N-H \cdots O$ hydrogen bonding leads to an infinite two-dimensional network along (100).

Related literature

For the first preparation of the title compound, see: Haas & Schuler (1964). For the properties of polymers of the title compound in aqueous solution, see: Haas *et al.* (1967, 1970*a,b,c,d*); Marstokk *et al.* (1998); Nagaoka *et al.* (2007); Ohnishi *et al.* (2007); Seuring & Agarwal (2010); Glatzel *et al.* (2010). For the structure of the related compound, 2-(2-acrylamidoacetamido)acetic acid monohydrate, see Gao *et al.* (2007).



Experimental

Crystal data

$C_5H_8N_2O_2$
 $M_r = 128.13$
 Monoclinic, $P2_1/c$
 $a = 15.938$ (2) Å
 $b = 4.8055$ (4) Å
 $c = 8.4920$ (12) Å
 $\beta = 98.109$ (11)°

$V = 643.91$ (14) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.10$ mm⁻¹
 $T = 100$ K
 $0.23 \times 0.19 \times 0.09$ mm

Data collection

Stoe IPDS 2T diffractometer
 Absorption correction: integration
 (*X-RED*; Stoe & Cie, 2006)
 $T_{\min} = 0.991$, $T_{\max} = 0.997$
 6001 measured reflections
 1362 independent reflections
 1065 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.049$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.032$
 $wR(F^2) = 0.085$
 $S = 0.97$
 1362 reflections
 115 parameters
 All H-atom parameters refined
 $\Delta\rho_{\max} = 0.18$ e Å⁻³
 $\Delta\rho_{\min} = -0.15$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$N5-H5 \cdots O4^i$	0.850 (19)	2.062 (19)	2.8946 (14)	166.3 (15)
$N8-H8B \cdots O9^{ii}$	0.881 (17)	2.081 (17)	2.9494 (14)	168.2 (15)
$N8-H8A \cdots O9^{iii}$	0.927 (18)	1.971 (18)	2.8855 (14)	168.6 (16)

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

Data collection: *X-AREA* (Stoe & Cie, 2006); cell refinement: *X-AREA*; data reduction: *X-AREA*; program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *DIAMOND* (Brandenburg, 2007); software used to prepare material for publication: *publCIF* (Westrip, 2010), *PLATON* (Spek, 2009) and *WinGX* (Farrugia, 1999).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SJ5179).

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supplementary materials

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Comment

N-acryloyl glycinamide was first synthesized by Haas and Schuler (1964). It can be polymerized radically to obtain polymers that exhibit thermoresponsive behavior in aqueous solution. The polymers show gelatin-like thermoreversible gelation (Haas & Schuler, 1964; Haas *et al.*, 1967, 1970*a*, 1970*b*, 1970*c*, 1970*d*; Marstokk *et al.*, 1998; Seuring & Agarwal, 2010; Glatzel *et al.*, 2010) and an upper critical solution temperature (Seuring & Agarwal, 2010; Ohnishi *et al.*, 2007; Nagaoka *et al.*, 2007) in water. These phenomena rely on intermolecular hydrogen bonding. Therefore, investigating the intermolecular hydrogen bonding between monomer units in the crystal may contribute to the understanding of interpolymer interactions.

The molecular structure of the title compound shows two planar parts with C1, C2, C3, O4, N5, C6 and C6, C7, N8, O9 in plane. The angle between these mean planes is 81.36 (7)°. In the packing the molecule forms three hydrogen bonds to three different neighbouring molecules. For details see Table 1. The intermolecular N5—H5···O4ⁱ contacts form an infinite chain in the (0 1 0) direction. Two of these chains are linked *via* N8—H8A···O9ⁱⁱⁱ and N8—H8B···O9ⁱⁱ interactions, respectively (symmetry codes: (i) $x, y + 1, z$; (ii) $-x + 1, y - 1/2, -z + 3/2$; (iii) $x, -y + 1/2, z + 1/2$). Herein the N8—H8A···O9ⁱⁱⁱ hydrogen bonds form a second chain with direction (0 0 1), and hydrogen bonded rings are generated. In conclusion, a two dimensional hydrogen bond network has been formed with the hydrophobic tails of the molecules as border planes.

For the crystal structure of a related compound, 2-(2-acrylamidoacetamido)acetic acid monohydrate, see Gao *et al.* (2007).

Experimental

N-acryloyl glycinamide has been prepared according to the route of Haas and Schuler (1964). However, reagent ratios, workup and purification have been modified as follows.

In a 1 l three-necked round-bottom flask equipped with a mechanical stirrer glycinamide hydrochloride (23.11 g, 209 mmol) and potassium carbonate (56.7 g, 410 mmol) were dissolved in 125 ml of water. The solution was cooled in an ice bath and acryloyl chloride (16.65 ml, 205 mmol) dissolved in 250 ml of diethylether was added dropwise over 30 min with fast stirring (300 rpm). The suspension was further stirred at RT for 2 h. The diethylether was removed by rotary evaporation at 35 °C. The remaining aqueous phase was freeze dried. The crude brittle solid was extracted with acetone (6 times, 500 ml acetone, 40 °C, stirring for at least 15 min). Insoluble potassium salts were filtered off and the acetone was removed by rotary evaporation at 35 °C. 22.7 g (85%) of crude product were obtained. The crude product was dissolved in an eluent mixture of methanol and dichloromethane ($v/v = 1/4$, 600 ml) by heating to reflux once. The solution was filtered to remove polymeric impurities and purified by column chromatography ($d = 9$ cm, 900 g silica, porosity 60 Å, 0.063–0.2 mm mesh size, TLC: R_f (N-acryloyl glycinamide) = 0.40) to obtain 21.3 g (80%) of product which was recrystallized from 240 ml of a mixture of methanol and acetone ($v/v = 1/2$) to yield 15.3 g (57%) of colorless needle-like crystals.

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For obtaining crystals that are suitable for X-ray analysis a small fraction of purified N-acryloyl glycinamide was again recrystallized from a dilute solution in 2-propanol.

Refinement

All H atoms were located in a difference Fourier map and refined isotropically. The C—H bond distances vary from 0.937 (18) to 0.981 (18), the N—H bond lengths from 0.850 (19) to 0.927 (18) Å.

Figures

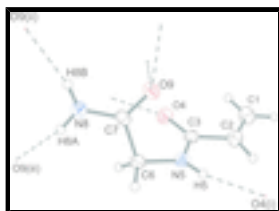


Fig. 1. View of the crystal structure of the title compound. The thermal ellipsoids are drawn at 50% probability level. Dashed lines indicate hydrogen bonding contacts.

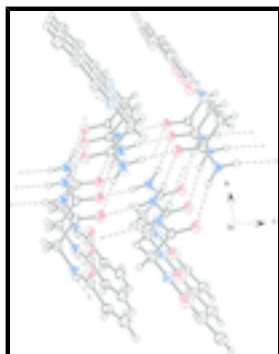


Fig. 2. Two dimensional hydrogen bond network with plane direction (1 0 0). Dashed lines indicate hydrogen bonds.

N-(Carbamoylmethyl)prop-2-enamide

Crystal data

$C_5H_8N_2O_2$

$M_r = 128.13$

Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

$a = 15.938$ (2) Å

$b = 4.8055$ (4) Å

$c = 8.4920$ (12) Å

$\beta = 98.109$ (11)°

$V = 643.91$ (14) Å³

$Z = 4$

$F(000) = 272$

$D_x = 1.322$ Mg m⁻³

Melting point: 143 K

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

Cell parameters from 7769 reflections

$\theta = 2.6$ – 27°

$\mu = 0.10$ mm⁻¹

$T = 100$ K

Plate, colourless

$0.23 \times 0.19 \times 0.09$ mm

Data collection

Stoe IPDS 2T
diffractometer

1362 independent reflections

Radiation source: fine-focus sealed tube
graphite
Detector resolution: 6.67 pixels mm⁻¹
rotation method scans
Absorption correction: integration
(*X-RED*; Stoe & Cie, 2006)
 $T_{\min} = 0.991$, $T_{\max} = 0.997$
6001 measured reflections

1065 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.049$
 $\theta_{\max} = 26.8^\circ$, $\theta_{\min} = 2.6^\circ$
 $h = -20 \rightarrow 17$
 $k = -6 \rightarrow 6$
 $l = -10 \rightarrow 10$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.032$
 $wR(F^2) = 0.085$
 $S = 0.97$
1362 reflections
115 parameters
0 restraints
Primary atom site location: structure-invariant direct methods

Secondary atom site location: difference Fourier map
Hydrogen site location: difference Fourier map
All H-atom parameters refined
 $w = 1/[\sigma^2(F_o^2) + (0.0548P)^2]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.18 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.15 \text{ e } \text{\AA}^{-3}$
Extinction correction: *SHELXL97* (Sheldrick, 2008),
 $F_c^* = kFc[1 + 0.001 \times Fc^2 \lambda^3 / \sin(2\theta)]^{-1/4}$
Extinction coefficient: 0.032 (7)

Special details

Experimental. DSC (rate of heating = 10 K min⁻¹): $T_m = 143 \text{ }^\circ\text{C}$. IR (ATR): $\nu = 3380$ (m, NH), 3312 (s, NH), 3187 (m, NH), 1652 (*vs*, C=O), 1621 (*vs*, C=O), 1551 (*vs*, NH) cm⁻¹. ¹H NMR (300 MHz, D₂O): $\delta = 3.93$ (s, 2H, N-CH₂-CONH₂), 5.77 [dd, J(doublet 1) = 2.0 Hz, J(doublet 2) = 9.5 Hz, 1H, H_{olef.}], 6.20 [dd, J(doublet 1) = 2.0 Hz, J(doublet 2) = 17.1 Hz, 1H, H_{olef.}], 6.29 [dd, J(doublet 1) = 9.5 Hz, J(doublet 2) = 17.2 Hz, 1H, H_{olef.}]. ¹³C NMR (75 MHz, D₂O): $\delta = 42.7$ (-N-CH₂-), 128.8 (C_{olef.}), 130.0 (C_{olef.}), 169.6 (-CO-), 174.8 (CO-). Flame emission spectroscopy: potassium content 5 p.p.m.

Geometry. All s.u.'s (except the s.u. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell s.u.'s are taken into account individually in the estimation of s.u.'s in distances, angles and torsion angles; correlations between s.u.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell s.u.'s is used for estimating s.u.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > 2\sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.91738 (9)	0.4644 (3)	0.66968 (16)	0.0334 (3)
C2	0.87077 (8)	0.6065 (3)	0.75755 (15)	0.0269 (3)

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C3	0.80276 (7)	0.4732 (3)	0.83438 (13)	0.0218 (3)
C6	0.68733 (7)	0.5421 (3)	0.98560 (13)	0.0229 (3)
C7	0.61296 (7)	0.4355 (3)	0.86952 (13)	0.0209 (3)
N5	0.75513 (6)	0.6473 (2)	0.90665 (12)	0.0221 (3)
N8	0.56460 (7)	0.2477 (2)	0.92691 (12)	0.0271 (3)
O4	0.79110 (6)	0.21891 (18)	0.83202 (10)	0.0280 (2)
O9	0.59828 (5)	0.52628 (19)	0.73164 (9)	0.0248 (2)
H1A	0.9069 (10)	0.273 (4)	0.6562 (19)	0.038 (4)*
H1B	0.9623 (10)	0.553 (4)	0.620 (2)	0.041 (4)*
H2	0.8770 (10)	0.802 (4)	0.7774 (18)	0.036 (4)*
H5	0.7628 (10)	0.822 (4)	0.8996 (19)	0.034 (4)*
H6A	0.7081 (8)	0.395 (3)	1.0551 (17)	0.023 (3)*
H6B	0.6661 (9)	0.697 (3)	1.0471 (17)	0.025 (3)*
H8A	0.5799 (10)	0.181 (4)	1.029 (2)	0.042 (4)*
H8B	0.5176 (11)	0.194 (4)	0.867 (2)	0.040 (4)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0316 (7)	0.0359 (8)	0.0336 (7)	0.0022 (6)	0.0078 (5)	0.0017 (6)
C2	0.0260 (6)	0.0247 (7)	0.0300 (6)	-0.0003 (5)	0.0039 (5)	0.0015 (5)
C3	0.0246 (6)	0.0198 (6)	0.0201 (5)	0.0000 (5)	0.0001 (4)	0.0001 (4)
C6	0.0261 (6)	0.0237 (6)	0.0189 (6)	-0.0002 (5)	0.0028 (4)	-0.0010 (5)
C7	0.0231 (6)	0.0211 (6)	0.0193 (5)	0.0030 (5)	0.0058 (4)	-0.0008 (4)
N5	0.0244 (5)	0.0177 (6)	0.0246 (5)	-0.0012 (4)	0.0040 (4)	-0.0010 (4)
N8	0.0286 (5)	0.0318 (7)	0.0204 (5)	-0.0076 (4)	0.0020 (4)	0.0030 (4)
O4	0.0361 (5)	0.0180 (5)	0.0304 (5)	-0.0008 (4)	0.0065 (4)	-0.0011 (4)
O9	0.0273 (4)	0.0291 (5)	0.0178 (4)	0.0006 (4)	0.0028 (3)	0.0025 (3)

Geometric parameters (\AA , $^\circ$)

C1—C2	1.3160 (19)	C6—C7	1.5198 (16)
C1—H1A	0.937 (18)	C6—H6A	0.950 (15)
C1—H1B	0.981 (18)	C6—H6B	0.994 (15)
C2—C3	1.4867 (17)	C7—O9	1.2405 (13)
C2—H2	0.956 (18)	C7—N8	1.3235 (16)
C3—O4	1.2360 (15)	N5—H5	0.850 (19)
C3—N5	1.3356 (16)	N8—H8A	0.927 (18)
C6—N5	1.4412 (15)	N8—H8B	0.881 (17)
C2—C1—H1A	118.2 (10)	N5—C6—H6B	108.4 (8)
C2—C1—H1B	121.6 (10)	C7—C6—H6B	107.4 (8)
H1A—C1—H1B	120.2 (15)	H6A—C6—H6B	110.1 (11)
C1—C2—C3	122.01 (13)	O9—C7—N8	123.05 (11)
C1—C2—H2	123.8 (10)	O9—C7—C6	121.30 (11)
C3—C2—H2	114.2 (10)	N8—C7—C6	115.62 (10)
O4—C3—N5	122.19 (11)	C3—N5—C6	120.42 (11)
O4—C3—C2	122.42 (11)	C3—N5—H5	119.2 (11)
N5—C3—C2	115.39 (11)	C6—N5—H5	120.2 (11)

N5—C6—C7	112.57 (9)	C7—N8—H8A	119.6 (11)
N5—C6—H6A	109.3 (8)	C7—N8—H8B	118.6 (11)
C7—C6—H6A	108.9 (8)	H8A—N8—H8B	121.8 (15)
C1—C2—C3—O4	6.58 (19)	O4—C3—N5—C6	-0.03 (16)
C1—C2—C3—N5	-173.59 (12)	C2—C3—N5—C6	-179.87 (10)
N5—C6—C7—O9	-26.19 (17)	C7—C6—N5—C3	-70.81 (14)
N5—C6—C7—N8	155.59 (11)		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
N5—H5 \cdots O4 ⁱ	0.850 (19)	2.062 (19)	2.8946 (14)	166.3 (15)
N8—H8B \cdots O9 ⁱⁱ	0.881 (17)	2.081 (17)	2.9494 (14)	168.2 (15)
N8—H8A \cdots O9 ⁱⁱⁱ	0.927 (18)	1.971 (18)	2.8855 (14)	168.6 (16)

Symmetry codes: (i) $x, y+1, z$; (ii) $-x+1, y-1/2, -z+3/2$; (iii) $x, -y+1/2, z+1/2$.

Fig. 1

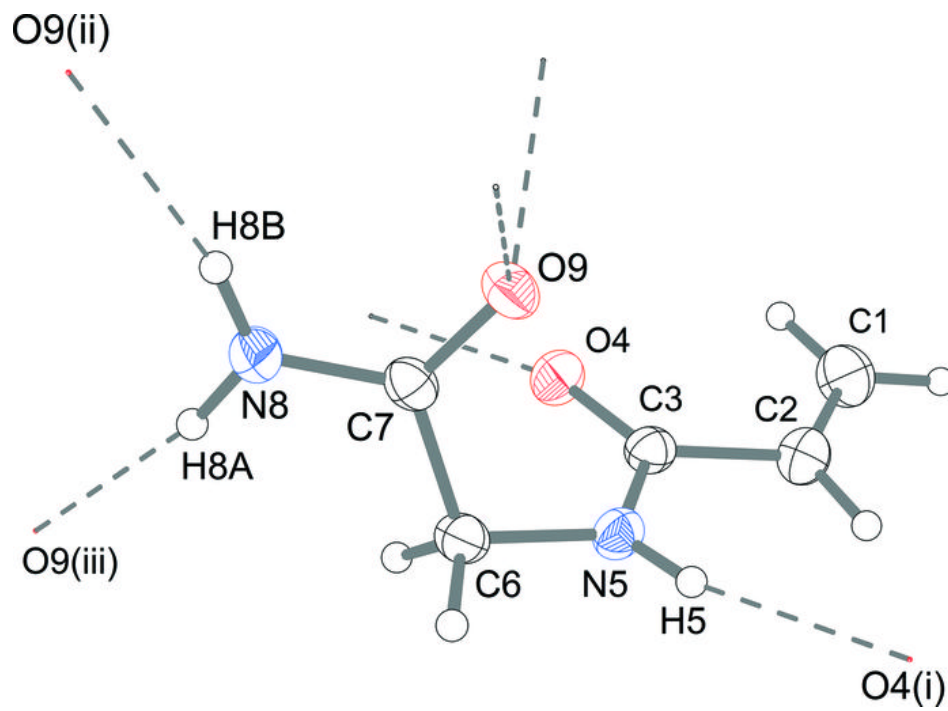


Fig. 2

