

N-(2-Aminophenyl)maleamic acid

Norma Francenia Santos-Sánchez,^a Raúl Salas-Coronado,^a Adrián Peña-Hueso^b and Angelina Flores-Parra^{b*}

^aInstituto de Agroindustrias, Universidad Tecnológica de la Mixteca, 69000 Huajuapan de León, Oaxaca, Mexico, and ^bDepartment of Chemistry, CINVESTAV-IPN, AP 14-740, 07000 México DF, Mexico
Correspondence e-mail: nsantos@mixteco.utm.mx

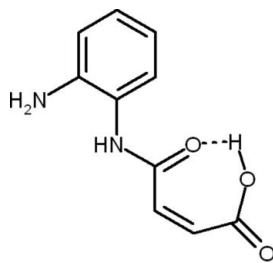
Received 10 August 2007; accepted 19 September 2007

Key indicators: single-crystal X-ray study; $T = 294\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.041; wR factor = 0.051; data-to-parameter ratio = 8.1.

The title compound [also called (*Z*)-3-(2-aminophenylcarbamoyl)propenoic acid], $C_{10}H_{10}N_2O_3$, is a non-planar amidated maleic acid derivative, with the mean planes through the 2-aminophenyl and maleamic acid groups inclined at an angle of $43.08(10)^\circ$. Symmetry-related molecules are linked by intermolecular $\text{N}-\text{H}\cdots\text{N}$ hydrogen bonds, forming centrosymmetric amine-amide dimers. The dimers are linked by $\text{N}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds and weak $\text{N}-\text{H}\cdots\pi$ and $\pi-\pi$ interactions [C-to-centroid distance $3.510(3)\text{ \AA}$] into a three-dimensional network.

Related literature

For related literature, see: Bernstein *et al.* (1995); Vaupel *et al.* (2006).



Experimental

Crystal data

$C_{10}H_{10}N_2O_3$
 $M_r = 206.2$
Monoclinic, $P2_1/c$
 $a = 7.3899(2)\text{ \AA}$
 $b = 12.2135(3)\text{ \AA}$
 $c = 12.2633(3)\text{ \AA}$
 $\beta = 119.726(1)^\circ$
 $V = 961.19(4)\text{ \AA}^3$
 $Z = 4$
Mo $K\alpha$ radiation
 $\mu = 0.11\text{ mm}^{-1}$
 $T = 294\text{ K}$
 $0.25 \times 0.25 \times 0.1\text{ mm}$

Data collection

Nonius KappaCCD area-detector diffractometer
Absorption correction: none
4773 measured reflections

2418 independent reflections
1155 reflections with $I > 2.5\sigma(I)$
 $R_{\text{int}} = 0.039$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.041$
 $wR(F^2) = 0.051$
 $S = 1.01$
1155 reflections
142 parameters
2 restraints

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.17\text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.17\text{ e \AA}^{-3}$

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O1—H1···O4	0.87	1.62	2.492 (2)	174
N4—H4···N6 ⁱ	0.87	2.17	3.009 (3)	163
N6—H61···O2 ⁱⁱ	0.878 (19)	2.125 (19)	2.969 (3)	161.0 (19)
C7—H7···O4 ⁱⁱⁱ	0.95	2.41	3.349 (3)	166
N6—H62···C9 ⁱⁱⁱ	0.85 (2)	2.77 (2)	3.496 (3)	144.0 (17)

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

Data collection: *COLLECT* (Nonius, 2001); cell refinement: *DENZO/SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *DENZO/SCALEPACK*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *CRYSTALS* (Betteridge *et al.*, 2003); molecular graphics: *CAMERON* (Watkin *et al.*, 1996); software used to prepare material for publication: *PLATON* (Spek, 2003).

The authors acknowledge Sonia Araceli Sánchez Ruiz, Department of Chemistry, CINVESTAV-IPN, for the acquisition of the MS, IR, ^1H and ^{13}C NMR spectra.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: ZL2058).

References

- Bernstein, J., Davis, R. E., Shimoni, L. & Chang, N.-L. (1995). *Angew. Chem. Int. Ed. Engl.* **34**, 1555–1573.
Betteridge, P. W., Carruthers, J. R., Cooper, R. I., Prout, K. & Watkin, D. J. (2003). *J. Appl. Cryst.* **36**, 1487.
Nonius (2001). *COLLECT*. Nonius BV, Delft, The Netherlands.
Otwinowski, Z. & Minor, W. (1997). *Methods in Enzymology*, Vol. 276, *Macromolecular Crystallography*, Part A, edited by C. W. Carter Jr & R. M. Sweet, pp. 307–326. New York: Academic Press.
Sheldrick, G. M. (1997). *SHELXS97*. University of Göttingen, Germany.
Spek, A. L. (2003). *J. Appl. Cryst.* **36**, 7–13.
Vaupel, S., Brutschy, B., Tarakeshwar, P. & Kim, K. S. (2006). *J. Am. Chem. Soc.* **128**, 5416–5426.
Watkin, D. J., Prout, C. K. & Pearce, L. J. (1996). *CAMERON*. Chemical Crystallography Laboratory, University of Oxford, England.

supplementary materials

Acta Cryst. (2007). E63, o4156 [doi:10.1107/S1600536807046132]

N-(2-Aminophenyl)maleamic acid

N. F. Santos-Sánchez, R. Salas-Coronado, A. Peña-Hueso and A. Flores-Parra

Comment

The title compound, (**I**), has been prepared by condensation from maleic anhydride and *o*-phenylenediamine as a intermediate for the synthesis of 1,2,3,4-tetrahydroquinoxalin-2-ones.

In the molecular structure of compound (**I**), the C2—C3 bond length of the maleamic moiety proves to have a clear double bond character and the bonds C1—C2 and C3—C4 are clearly single bonds, which confirms localized bonding, Table 1. The maleamic moiety is characterized by a rather short and almost linear O—H···O intramolecular hydrogen bond forming an *S*(7) ring (Bernstein *et al.*, 1995), Figure 1. The C—O distances in the carboxylic acid group are consistent with the position of the carboxyl H atom deduced from difference maps. The title compound is non-planar with the mean planes through the 2-aminophenyl and maleamic acid groups being inclined against each other at an angle of 43.08 (10)°.

The crystal packing for (**I**) shows that the molecules are linked by two close to linear N—H···N hydrogen bonds (see Table 2 for geometric parameters and symmetry codes) to form centrosymmetric $R^2_2(10)$ dimers (Figure 2). These hydrogen bonds probably are the reason for the non-planarity of compound (**I**).

Adjacent dimers of (**I**) are linked by N—H···O and C—H···O hydrogen bonds and also by weaker N—H···π contacts and π···π stacking interactions. Amino atom N6 acts as a hydrogen-bond donor, *via* H61, to atom O2 in the molecule at ($-x, y - 1/2, -z + 1/2$), while atom C7 at ($-x, y - 1/2, -z + 1/2$), in turn, acts as donor to O2 at ($x, y - 1, z$). In this manner, a *C*(10) chain is formed (motif a, Figure 3.), running along the *c* axis. The arrangement of N6—H61···O2ⁱⁱ, N6ⁱⁱ—H61ⁱⁱ···O2^{iv}, C7ⁱⁱⁱ—H7ⁱⁱⁱ···O4^{iv}, C7—H7···O4ⁱⁱⁱ interactions can be described by the graph-set notation $R^4_4(28)$. Aromatic carbon atom C7 acts as a hydrogen-bond donor, *via* H7, to O4 in the molecule at ($-x + 1, y - 1/2, -z + 3/2$), while C7 at ($-x + 1, y - 1/2, -z + 3/2$), in turn, acts as a donor to O4 at ($x, y - 1, z$). In this manner, a *C*(7) zigzag chain (motif b) is formed, running along the *b* axis, Figure 3. The geometry of the hydrogen bonding is given in Table 2. The amine hydrogen atom H62 is forming a N—H···π interaction with aromatic carbon C9 at ($-x + 1, y - 1/2, -z + 3/2$). These interactions are dominated by dispersion energies, Vaupel *et al.* (2006). Lastly, maleamic moieties do interact with the C5—C10 benzene ring (centroid Cg) at ($x - 1, -y + 3/2, z - 1/2$), the distance C1···Cg is 3.510 (3) Å and the gamma angle is 22.8°, Figure 4.

Experimental

To a solution of *o*-phenylenediamine (9.2 mmol) in THF (20 ml) is added a solution of maleic anhydride (0.91 g, 9.3 mmol) in THF (20 ml). The suspension was stirred for 12 h at room temperature. The mixture was filtered and the solvent distilled off. The residue was precipitated with a mixture of acetone/hexane to give a yellow solid (1.87 g, 98%), which was washed with ethyl ether and recrystallized from THF to give yellow crystals suitable for single-crystal X-ray diffraction (m.p. 412–413 K). R_F (4:6:0.1, hexane: AcOEt: formic acid) 0.23. ^1H NMR (300.13 MHz, DMSO-*d*₆): δ 9.85 (s, 1H, CO₂H), 8.04 (bs, 2H, NH₂), 7.14 (bd, *J* = 7.9 Hz, 1H, H10), 6.97 (td, *J* = 7.9, 1.2 Hz, 1H, H8), 6.74 (bd, *J* = 7.9 Hz, 1H, H7), 6.61 (d, *J* = 12.2

supplementary materials

Hz, 1H, H3), 6.56 (td, $J = 7.9, 1.0$ Hz, 1H, H9), 6.27 (d, $J = 12.2$ Hz, 1H, H2); ^{13}C NMR (75.47 MHz, DMSO- d_6): δ 167.5 (CO₂), 165.0 (C4), 143.9 (C5), 134.5 (C3), 130.6 (C2), 127.8 (C8), 127.0 (C10), 122.4 (C6), 116.8 (C9), 116.5 (C7). MS [EI, m/z (%)]: 206 (100), 160 (64), 147 (99), 119 (36). HRMS calcd for [C₁₀H₁₀N₂O₃ + H]⁺ 207.0764, found 207.0769. FT-IR (KBr) (cm⁻¹): 3388, 3308, 1706, 1308, 839.

Refinement

All non-hydrogen atoms were refined anisotropically, C—H and amide hydrogen were placed in calculated positions, with distances C—H = 0.95 Å, N—H = 0.87 Å and O—H = 0.87 Å and $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{C}, \text{N})$ or $1.5 U_{\text{eq}}(\text{O})$. The H atoms of the amine group were refined but the N—H distances were restrained to 0.87 (2) Å [$U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{N})$].

Figures

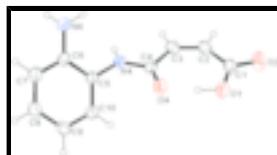


Fig. 1. A view of (I), showing 30% probability displacement ellipsoids (H atoms are drawn as small spheres of arbitrary radii).

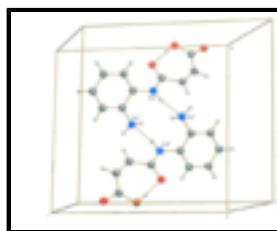


Fig. 2. Part of the crystal structure of (I), showing the formation of a hydrogen-bonded $R^2_2(10)$ dimer. Dashed lines indicate hydrogen bonds. Atoms marked with (i) are at the symmetry related position ($-x + 1, -y + 1, -z + 1$).



Fig. 3. The packing of (I), showing the $R^4_4(28)$ ring pattern. Dashed lines indicate hydrogen bonds. For the sake of clarity, H atoms not involved in the motifs shown have been omitted. [Symmetry codes: (ii) $-x, y - 1/2, -z + 1/2$; (iii) $-x + 1, y - 1/2, -z + 3/2$; (iv) $x, y - 1, z$.]

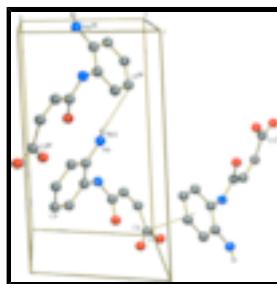


Fig. 4. Part of the crystal structure of (I), showing the formation of weak N—H···π and π···π interactions. For clarity, all H atoms not involved in the motifs shown have been omitted [Symmetry codes: (iii) $-x + 1, y - 1/2, -z + 3/2$; (v) $-1 + x, 3/2 - y, -1/2 + z$.]

(Z)-3-(2-aminophenylcarbamoyl)propenoic acid

Crystal data

C₁₀H₁₀N₂O₃

$F_{000} = 432$

$M_r = 206.2$

$D_x = 1.425 \text{ Mg m}^{-3}$

Monoclinic, $P2_1/c$

Mo $K\alpha$ radiation

$\lambda = 0.71073 \text{ \AA}$

Hall symbol: -P 2ybc	Cell parameters from 2535 reflections
$a = 7.3899 (2) \text{ \AA}$	$\theta = 1\text{--}29^\circ$
$b = 12.2135 (3) \text{ \AA}$	$\mu = 0.11 \text{ mm}^{-1}$
$c = 12.2633 (3) \text{ \AA}$	$T = 294 \text{ K}$
$\beta = 119.7260 (10)^\circ$	Prism, colourless
$V = 961.19 (4) \text{ \AA}^3$	$0.25 \times 0.25 \times 0.1 \text{ mm}$
$Z = 4$	

Data collection

Nonius KappaCCD area-detector diffractometer	1155 reflections with $I > 2.5\sigma(I)$
Monochromator: graphite	$R_{\text{int}} = 0.039$
Detector resolution: 9 pixels mm^{-1}	$\theta_{\text{max}} = 28.6^\circ$
$T = 291 \text{ K}$	$\theta_{\text{min}} = 2.5^\circ$
φ and ω scans	$h = -9 \rightarrow 9$
Absorption correction: none	$k = -16 \rightarrow 16$
4773 measured reflections	$l = -16 \rightarrow 16$
2418 independent reflections	

Refinement

Refinement on F	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.041$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.051$	Method = Modified Sheldrick $w = 1/[\sigma^2(F^2) + (0.04P)^2]$, where $P = (\max(F_o^2, 0) + 2F_c^2)/3$ (Sheldrick, 1997)
$S = 1.01$	$(\Delta/\sigma)_{\text{max}} = 0.0002$
1155 reflections	$\Delta\rho_{\text{max}} = 0.17 \text{ e \AA}^{-3}$
142 parameters	$\Delta\rho_{\text{min}} = -0.17 \text{ e \AA}^{-3}$
2 restraints	Extinction correction: none
Primary atom site location: structure-invariant direct methods	

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'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 $F^2 > 2\text{sigma}(F^2)$ 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.

supplementary materials

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	0.0694 (3)	0.9421 (2)	0.2238 (2)	0.0631
C2	0.1320 (3)	0.82792 (19)	0.21902 (19)	0.0626
C3	0.2464 (3)	0.75461 (18)	0.30799 (19)	0.0566
C4	0.3461 (3)	0.76698 (16)	0.44500 (18)	0.0479
C5	0.5634 (3)	0.67199 (14)	0.64461 (16)	0.0421
C6	0.5469 (3)	0.57449 (15)	0.69851 (16)	0.0449
C7	0.6578 (3)	0.56406 (18)	0.82783 (18)	0.057
C8	0.7808 (3)	0.64865 (18)	0.9029 (2)	0.0633
C9	0.7967 (3)	0.74467 (17)	0.84874 (19)	0.0575
C10	0.6885 (3)	0.75622 (15)	0.71998 (18)	0.0491
H1	0.2094	0.9452	0.392	0.1064*
H2	0.0782	0.8031	0.1371	0.0719*
H3	0.2663	0.6861	0.2818	0.0655*
H4	0.4646	0.6245	0.4705	0.0569*
H7	0.6456	0.4975	0.8645	0.0697*
H8	0.8545	0.6406	0.9915	0.0728*
H9	0.8829	0.8021	0.9008	0.0643*
H10	0.7023	0.8206	0.6838	0.0573*
H61	0.316 (3)	0.5050 (17)	0.5507 (16)	0.0671*
H62	0.409 (3)	0.4385 (16)	0.6653 (18)	0.0687*
N4	0.4533 (2)	0.67995 (12)	0.51109 (13)	0.0474
N6	0.4266 (3)	0.48726 (14)	0.62196 (17)	0.0557
O1	0.1358 (2)	0.99089 (13)	0.33127 (15)	0.0797
O2	-0.0414 (3)	0.99077 (16)	0.12712 (16)	0.0937
O4	0.3304 (2)	0.85106 (12)	0.49678 (12)	0.0711

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0556 (12)	0.0661 (15)	0.0542 (14)	-0.0072 (11)	0.0170 (11)	0.0160 (12)
C2	0.0636 (12)	0.0744 (16)	0.0390 (12)	-0.0143 (12)	0.0172 (10)	-0.0011 (11)
C3	0.0666 (13)	0.0534 (13)	0.0449 (12)	-0.0047 (11)	0.0240 (11)	-0.0063 (10)
C4	0.0549 (11)	0.0432 (11)	0.0430 (11)	-0.0034 (9)	0.0222 (9)	-0.0036 (9)
C5	0.0471 (10)	0.0398 (10)	0.0399 (10)	0.0036 (8)	0.0218 (9)	-0.0006 (8)
C6	0.0513 (10)	0.0395 (10)	0.0478 (11)	0.0024 (9)	0.0276 (9)	-0.0007 (8)
C7	0.0703 (13)	0.0511 (12)	0.0524 (12)	0.0047 (11)	0.0327 (11)	0.0069 (10)
C8	0.0742 (14)	0.0642 (15)	0.0421 (11)	0.0112 (12)	0.0217 (10)	0.0035 (11)
C9	0.0591 (12)	0.0517 (13)	0.0479 (13)	0.0028 (10)	0.0160 (10)	-0.0073 (10)
C10	0.0542 (11)	0.0399 (11)	0.0499 (12)	-0.0002 (9)	0.0233 (10)	-0.0017 (9)
N4	0.0635 (10)	0.0367 (9)	0.0416 (9)	0.0011 (8)	0.0257 (8)	-0.0037 (7)
N6	0.0686 (11)	0.0415 (10)	0.0590 (12)	-0.0063 (9)	0.0331 (10)	-0.0004 (8)
O1	0.0946 (12)	0.0566 (10)	0.0640 (11)	0.0115 (8)	0.0210 (9)	0.0103 (8)
O2	0.0916 (12)	0.0978 (14)	0.0665 (11)	0.0065 (10)	0.0199 (10)	0.0353 (10)
O4	0.0936 (11)	0.0542 (9)	0.0460 (9)	0.0215 (8)	0.0197 (8)	-0.0042 (7)

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

C1—C2	1.479 (3)	C6—N6	1.407 (2)
C1—O1	1.301 (3)	C7—C8	1.383 (3)
C1—O2	1.211 (2)	C7—H7	0.954
C2—C3	1.339 (3)	C8—C9	1.381 (3)
C2—H2	0.929	C8—H8	0.948
C3—C4	1.470 (3)	C9—C10	1.379 (3)
C3—H3	0.933	C9—H9	0.949
C4—N4	1.333 (2)	C10—H10	0.932
C4—O4	1.243 (2)	H1—O1	0.872
C5—C6	1.396 (3)	H4—N4	0.869
C5—C10	1.386 (2)	H61—N6	0.879 (15)
C5—N4	1.425 (2)	H62—N6	0.849 (15)
C6—C7	1.383 (3)		
C2—C1—O1	120.3 (2)	C6—C7—H7	118.5
C2—C1—O2	119.8 (2)	C8—C7—H7	120.5
O1—C1—O2	119.9 (2)	C7—C8—C9	119.90 (19)
C1—C2—C3	133.0 (2)	C7—C8—H8	120
C1—C2—H2	112.1	C9—C8—H8	120.1
C3—C2—H2	115	C8—C9—C10	119.90 (19)
C2—C3—C4	128.2 (2)	C8—C9—H9	119.5
C2—C3—H3	117.6	C10—C9—H9	120.6
C4—C3—H3	114.2	C5—C10—C9	120.29 (17)
C3—C4—N4	115.02 (17)	C5—C10—H10	120.1
C3—C4—O4	123.22 (18)	C9—C10—H10	119.6
N4—C4—O4	121.74 (17)	C5—N4—C4	125.55 (15)
C6—C5—C10	120.22 (17)	C5—N4—H4	116.3
C6—C5—N4	117.91 (16)	C4—N4—H4	118
C10—C5—N4	121.85 (16)	C6—N6—H61	116.5 (14)
C5—C6—C7	118.70 (17)	C6—N6—H62	110.8 (15)
C5—C6—N6	120.31 (17)	H61—N6—H62	116.1 (19)
C7—C6—N6	120.95 (18)	H1—O1—C1	109.5
C6—C7—C8	120.98 (19)		

Hydrogen-bond geometry (\AA , $^\circ$)

$D\cdots H$	$D—H$	$H\cdots A$	$D\cdots A$	$D—H\cdots A$
O1—H1 \cdots O4	0.87	1.62	2.492 (2)	174
N4—H4 \cdots N6 ⁱ	0.87	2.17	3.009 (3)	163
N6—H61 \cdots O2 ⁱⁱ	0.878 (19)	2.125 (19)	2.969 (3)	161.0 (19)
C7—H7 \cdots O4 ⁱⁱⁱ	0.95	2.41	3.349 (3)	166
N6—H62 \cdots C9 ⁱⁱⁱ	0.85 (2)	2.77 (2)	3.496 (3)	144.0 (17)

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

supplementary materials

Fig. 1

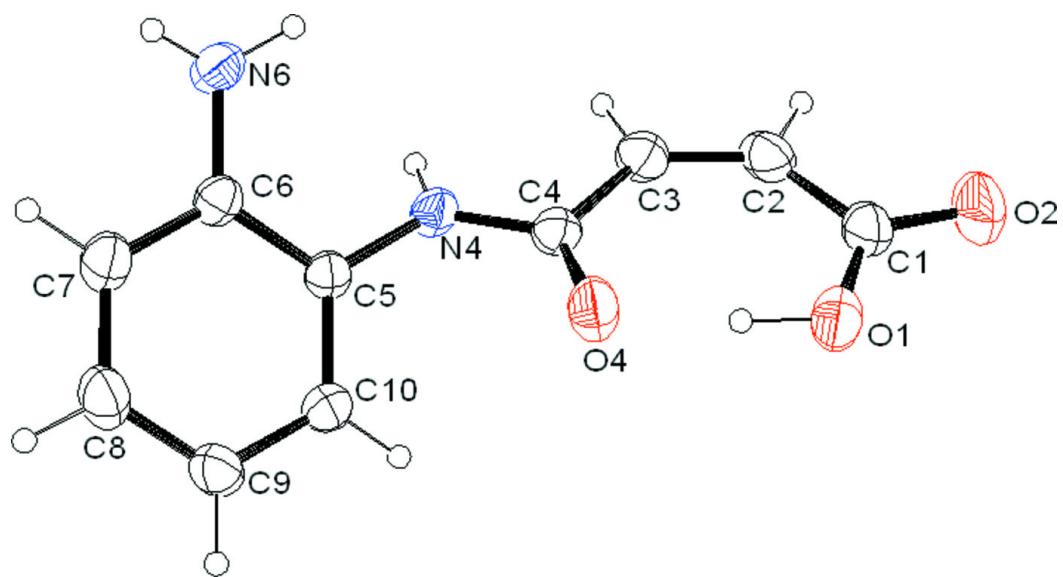
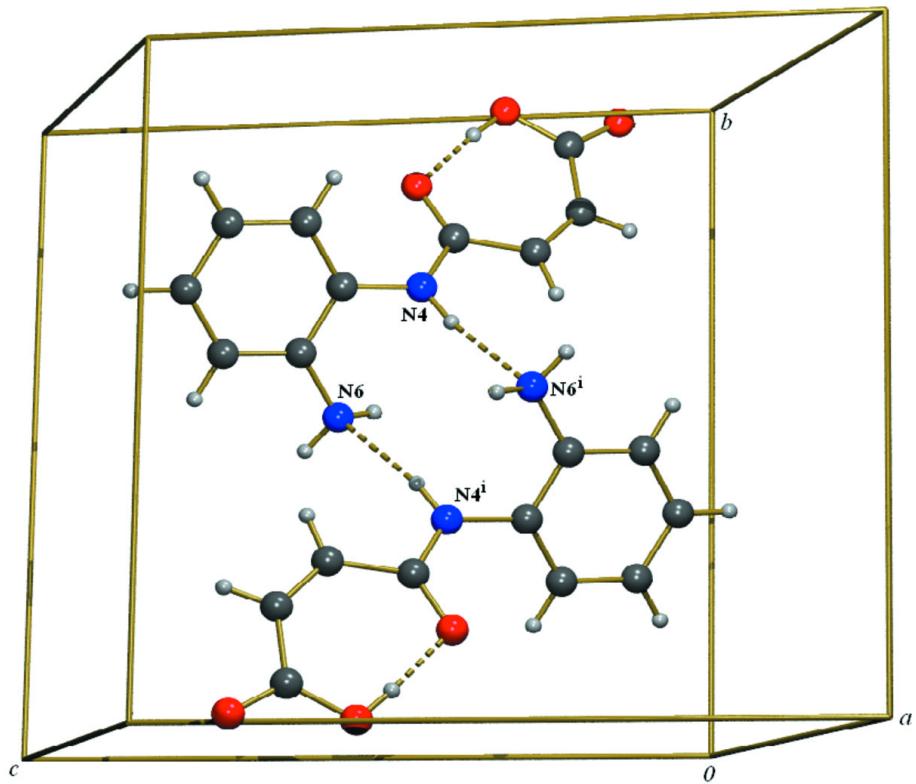


Fig. 2



supplementary materials

Fig. 3

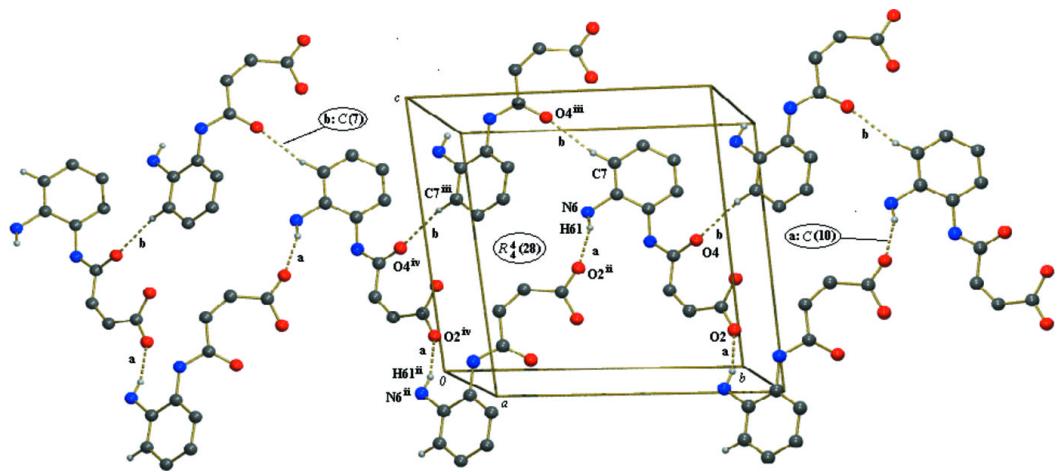


Fig. 4

