

Ethyl 3-[(3,5-dimethylphenyl)amino-carbonyl]propanoate

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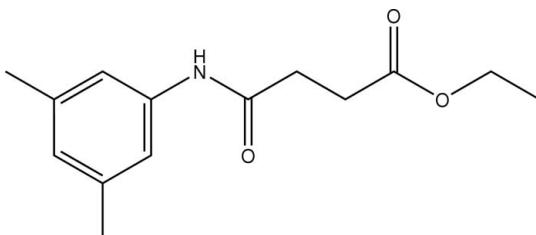
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Key indicators: single-crystal X-ray study; $T = 299\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; some non-H atoms missing; R factor = 0.058; wR factor = 0.169; data-to-parameter ratio = 11.4.

The non-H atoms in the title compound, $\text{C}_{14}\text{H}_{19}\text{NO}_3$, lie on a mirror plane. The amide O and ester carbonyl O atoms are *trans* to each other. Furthermore, the $\text{C}=\text{O}$ and $\text{O}-\text{CH}_2$ bonds of the ester group are *syn* with respect to each other. In the crystal, molecules are packed into centrosymmetric dimers through intermolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds.

Related literature

For related structures, see: Gowda *et al.* (2009a,b,c).



Experimental

Crystal data

$\text{C}_{14}\text{H}_{19}\text{NO}_3$

$M_r = 249.30$

Tetragonal, $I4/m$
 $a = 19.938(2)\text{ \AA}$
 $c = 7.0367(9)\text{ \AA}$
 $V = 2797.3(5)\text{ \AA}^3$
 $Z = 8$

$\text{Cu K}\alpha$ radiation
 $\mu = 0.67\text{ mm}^{-1}$
 $T = 299\text{ K}$
 $0.40 \times 0.28 \times 0.25\text{ mm}$

Data collection

Enraf–Nonius CAD-4 diffractometer
Absorption correction: none
4040 measured reflections
1367 independent reflections

1201 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.044$
3 standard reflections
frequency: 120 min
intensity decay: 1.0%

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.058$
 $wR(F^2) = 0.169$
 $S = 1.11$
1367 reflections
120 parameters

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.32\text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.32\text{ e \AA}^{-3}$

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

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N1—H1N \cdots O2 ⁱ	0.85 (3)	2.15 (3)	2.995 (3)	176 (3)

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

Data collection: *CAD-4-PC* (Enraf–Nonius, 1996); cell refinement: *CAD-4-PC*; data reduction: *REDU4* (Stoe & Cie, 1987); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *PLATON* (Spek, 2009); software used to prepare material for publication: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: TK2515).

References

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supporting information

Acta Cryst. (2009). E65, o2039 [doi:10.1107/S1600536809029511]

Ethyl 3-[(3,5-dimethylphenyl)aminocarbonyl]propanoate

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S1. Comment

As a part of studying the effect of ring and side chain substitutions on the structures of the substituted amides (Gowda *et al.*, 2009*a,b,c*), the crystal structure of ethyl *N*-(3,5-dimethylphenyl)succinamate (I) has been determined. The non-hydrogen atoms lie on a crystallographic mirror plane. The conformations of N—H and C=O bonds in the amide segment of the structure are *trans* to each other (Fig. 1). Likewise, the amide-O atom and ester carbonyl-O atoms are *trans* to each other. The C=O and O—CH₂ bonds of the ester group are in *syn* positions to each other, similar to that observed between the C=O and O—H bonds in the crystal structures of *N*-(2,6-dimethylphenyl)succinamic acid (Gowda *et al.*, 2009*b*) and *N*-(2-chlorophenyl)succinamic acid (Gowda *et al.*, 2009*a*).

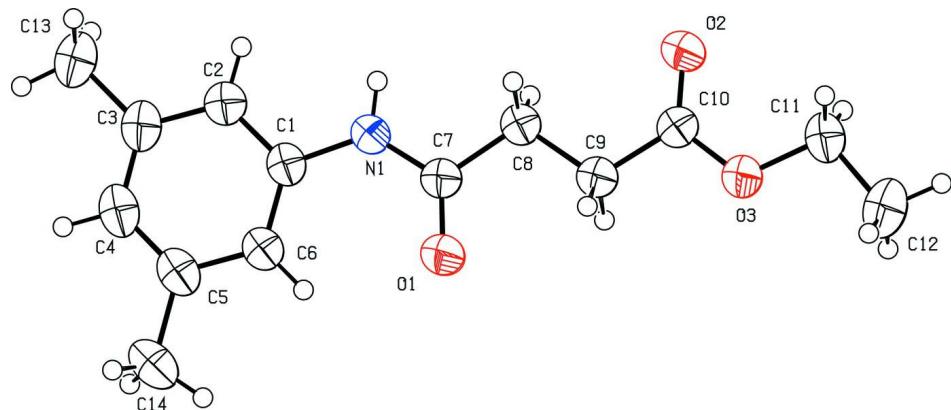
The presence of N—H···O hydrogen bonds connect the molecules into centrosymmetric dimers (Table 1).

S2. Experimental

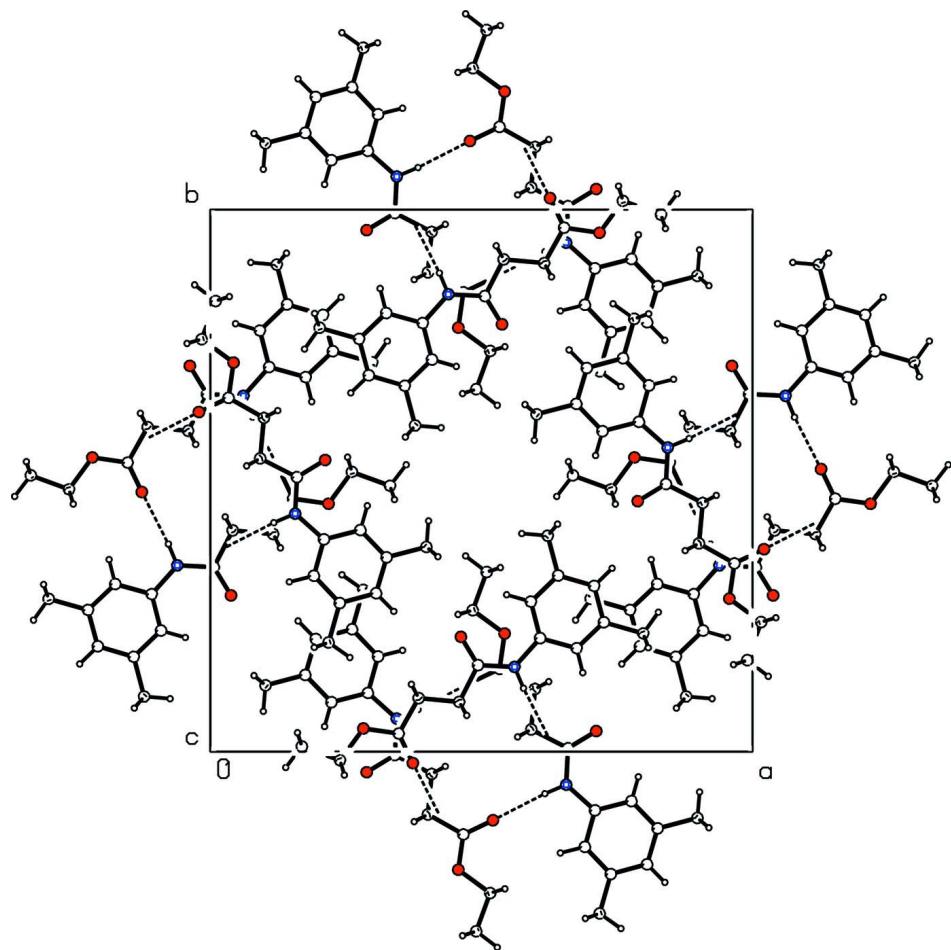
A solution of succinic anhydride (0.025 mole) in toluene (25 ml) was treated dropwise with a solution of 3,5-dimethylaniline (0.025 mole) also in toluene (20 ml) with constant stirring. The resulting mixture was stirred for about one hour and set aside for an additional hour at room temperature for the completion of reaction. The mixture was then treated with dilute hydrochloric acid to remove the unreacted 3,5-dimethylaniline. The resultant solid *N*-(3,5-dimethylphenyl)-succinamic acid was filtered under suction and washed thoroughly with water to remove the unreacted succinic anhydride and succinic acid. *N*-(3,5-Dimethylphenyl)succinamic acid was recrystallized into ethyl *N*-(3,5-dimethylphenyl)-succinamate (I) from hot ethanol. The rod like colourless single crystals of (I) were grown in hot ethanolic solution by slow evaporation.

S3. Refinement

The H atoms of the NH group, of C11 and C12 were located in a difference map and their position refined [N—H = 0.85 (3) Å, C—H = 0.98 (4)–1.03 (3) Å]. The other H atoms were positioned with idealized geometry using a riding model [C—H = 0.93–0.97 Å]. All H atoms were refined with isotropic displacement parameters (set to 1.2 times of the U_{eq} of the parent atom).

**Figure 1**

Molecular structure of (I), showing the atom labelling and the displacement ellipsoids are at the 50% probability level. The H atoms are represented as small spheres of arbitrary radii.

**Figure 2**

Molecular packing of (I) with hydrogen bonds shown as dashed lines.

Ethyl 3-[(3,5-dimethylphenyl)aminocarbonyl]propanoate*Crystal data*

$C_{14}H_{19}NO_3$
 $M_r = 249.30$
Tetragonal, $I4/m$
Hall symbol: -I 4
 $a = 19.938 (2)$ Å
 $c = 7.0367 (9)$ Å
 $V = 2797.3 (5)$ Å³
 $Z = 8$
 $F(000) = 1072$

$D_x = 1.184$ Mg m⁻³
Cu $K\alpha$ radiation, $\lambda = 1.54180$ Å
Cell parameters from 25 reflections
 $\theta = 4.4\text{--}20.5^\circ$
 $\mu = 0.67$ mm⁻¹
 $T = 299$ K
Rod, colourless
 $0.40 \times 0.28 \times 0.25$ mm

Data collection

Enraf–Nonius CAD-4
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 $\omega/2\theta$ scans
4040 measured reflections
1367 independent reflections
1201 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.044$
 $\theta_{\text{max}} = 67.0^\circ, \theta_{\text{min}} = 3.1^\circ$
 $h = -16 \rightarrow 23$
 $k = -16 \rightarrow 23$
 $l = -3 \rightarrow 8$
3 standard reflections every 120 min
intensity decay: 1.0%

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.058$
 $wR(F^2) = 0.169$
 $S = 1.11$
1367 reflections
120 parameters
0 restraints
Primary atom site location: structure-invariant
direct methods
Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites
H atoms treated by a mixture of independent
and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.1022P)^2 + 1.0554P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.32$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.32$ e Å⁻³
Extinction correction: *SHELXL97* (Sheldrick,
2008), $F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$
Extinction coefficient: 0.0012 (3)

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 $F^2 > \sigma(F^2)$ 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 (Å²)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
C1	0.60839 (11)	0.20945 (11)	0.0000	0.0465 (6)	
C2	0.67561 (12)	0.19085 (12)	0.0000	0.0513 (6)	
H2	0.6869	0.1456	0.0000	0.062*	

C3	0.72584 (12)	0.23863 (13)	0.0000	0.0532 (6)	
C4	0.70778 (13)	0.30566 (14)	0.0000	0.0579 (7)	
H4	0.7411	0.3383	0.0000	0.070*	
C5	0.64153 (13)	0.32504 (12)	0.0000	0.0562 (7)	
C6	0.59123 (12)	0.27666 (12)	0.0000	0.0522 (6)	
H6	0.5464	0.2894	0.0000	0.063*	
C7	0.49362 (12)	0.15957 (11)	0.0000	0.0506 (6)	
C8	0.45900 (11)	0.09187 (12)	0.0000	0.0543 (7)	
H8A	0.4728	0.0668	-0.1115	0.065*	0.50
H8B	0.4728	0.0668	0.1115	0.065*	0.50
C9	0.38416 (12)	0.09873 (11)	0.0000	0.0515 (6)	
H9A	0.3708	0.1242	0.1113	0.062*	0.50
H9B	0.3708	0.1242	-0.1113	0.062*	0.50
C10	0.34741 (11)	0.03335 (12)	0.0000	0.0491 (6)	
C11	0.23912 (14)	-0.01607 (15)	0.0000	0.0666 (8)	
H11	0.2470 (10)	-0.0432 (11)	0.118 (3)	0.080*	
C12	0.16867 (16)	0.0096 (2)	0.0000	0.0849 (10)	
H12A	0.139 (2)	-0.029 (2)	0.0000	0.102*	
H12B	0.1605 (12)	0.0349 (13)	0.126 (4)	0.102*	
C13	0.79850 (13)	0.21804 (16)	0.0000	0.0652 (7)	
H13A	0.8077	0.1916	-0.1110	0.078*	0.50
H13B	0.8078	0.1920	0.1118	0.078*	0.50
H13C	0.8263	0.2573	-0.0007	0.078*	
C14	0.62237 (18)	0.39835 (14)	0.0000	0.0829 (10)	
H14A	0.6403	0.4196	-0.1113	0.099*	0.50
H14B	0.6402	0.4196	0.1115	0.099*	0.50
H14C	0.5744	0.4024	-0.0003	0.099*	
N1	0.56095 (10)	0.15635 (10)	0.0000	0.0519 (6)	
H1N	0.5776 (15)	0.1170 (16)	0.0000	0.062*	
O1	0.46202 (9)	0.21173 (9)	0.0000	0.0821 (8)	
O2	0.37347 (9)	-0.02118 (8)	0.0000	0.0701 (7)	
O3	0.28217 (8)	0.04272 (8)	0.0000	0.0614 (6)	

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0444 (12)	0.0435 (12)	0.0516 (12)	-0.0050 (9)	0.000	0.000
C2	0.0475 (13)	0.0481 (13)	0.0583 (13)	-0.0009 (9)	0.000	0.000
C3	0.0465 (12)	0.0632 (15)	0.0498 (12)	-0.0103 (10)	0.000	0.000
C4	0.0567 (15)	0.0593 (15)	0.0578 (14)	-0.0177 (11)	0.000	0.000
C5	0.0621 (15)	0.0464 (13)	0.0601 (14)	-0.0105 (10)	0.000	0.000
C6	0.0494 (13)	0.0426 (12)	0.0645 (14)	-0.0036 (9)	0.000	0.000
C7	0.0433 (12)	0.0411 (12)	0.0673 (15)	0.0014 (9)	0.000	0.000
C8	0.0432 (13)	0.0422 (13)	0.0775 (16)	-0.0015 (9)	0.000	0.000
C9	0.0447 (13)	0.0425 (12)	0.0673 (15)	0.0006 (9)	0.000	0.000
C10	0.0430 (12)	0.0449 (12)	0.0594 (14)	0.0013 (9)	0.000	0.000
C11	0.0497 (14)	0.0559 (15)	0.094 (2)	-0.0124 (11)	0.000	0.000
C12	0.0490 (16)	0.094 (3)	0.112 (3)	-0.0109 (15)	0.000	0.000

C13	0.0460 (14)	0.0815 (19)	0.0681 (16)	-0.0094 (12)	0.000	0.000
C14	0.083 (2)	0.0443 (15)	0.121 (3)	-0.0118 (13)	0.000	0.000
N1	0.0424 (11)	0.0367 (10)	0.0768 (14)	-0.0013 (8)	0.000	0.000
O1	0.0481 (10)	0.0416 (10)	0.157 (2)	0.0021 (7)	0.000	0.000
O2	0.0514 (10)	0.0397 (10)	0.1193 (18)	0.0015 (7)	0.000	0.000
O3	0.0420 (9)	0.0458 (9)	0.0965 (14)	-0.0017 (7)	0.000	0.000

Geometric parameters (\AA , $\text{^{\circ}}$)

C1—C6	1.383 (3)	C9—H9A	0.9700
C1—C2	1.391 (3)	C9—H9B	0.9700
C1—N1	1.420 (3)	C10—O2	1.205 (3)
C2—C3	1.382 (3)	C10—O3	1.314 (3)
C2—H2	0.9300	C11—H11 ⁱ	1.00 (2)
C3—C4	1.384 (4)	C11—O3	1.453 (3)
C3—C13	1.506 (4)	C11—C12	1.495 (4)
C4—C5	1.376 (4)	C11—H11	1.00 (2)
C4—H4	0.9300	C12—H12B ⁱ	1.03 (3)
C5—C6	1.392 (3)	C12—H12A	0.98 (4)
C5—C14	1.511 (4)	C12—H12B	1.03 (3)
C6—H6	0.9300	C13—H13A	0.9600
C7—O1	1.216 (3)	C13—H13B	0.9600
C7—N1	1.344 (3)	C13—H13C	0.9600
C7—C8	1.516 (3)	C14—H14A	0.9600
C8—C9	1.498 (3)	C14—H14B	0.9600
C8—H8A	0.9700	C14—H14C	0.9600
C8—H8B	0.9700	N1—H1N	0.85 (3)
C9—C10	1.495 (3)		
C6—C1—C2	119.8 (2)	H9A—C9—H9B	107.6
C6—C1—N1	123.9 (2)	O2—C10—O3	123.7 (2)
C2—C1—N1	116.3 (2)	O2—C10—C9	125.1 (2)
C3—C2—C1	121.0 (2)	O3—C10—C9	111.17 (19)
C3—C2—H2	119.5	H11 ⁱ —C11—O3	110.1 (12)
C1—C2—H2	119.5	H11 ⁱ —C11—C12	109.4 (12)
C2—C3—C4	118.5 (2)	O3—C11—C12	106.2 (2)
C2—C3—C13	120.6 (3)	H11 ⁱ —C11—H11	112 (3)
C4—C3—C13	120.9 (2)	O3—C11—H11	110.1 (12)
C5—C4—C3	121.4 (2)	C12—C11—H11	109.4 (12)
C5—C4—H4	119.3	H12B ⁱ —C12—C11	108.4 (14)
C3—C4—H4	119.3	H12B ⁱ —C12—H12A	106.9 (16)
C4—C5—C6	119.8 (2)	C11—C12—H12A	107 (2)
C4—C5—C14	121.0 (2)	H12B ⁱ —C12—H12B	118 (3)
C6—C5—C14	119.2 (3)	C11—C12—H12B	108.4 (14)
C1—C6—C5	119.6 (2)	H12A—C12—H12B	106.9 (16)
C1—C6—H6	120.2	C3—C13—H13A	109.5
C5—C6—H6	120.2	C3—C13—H13B	109.5
O1—C7—N1	123.9 (2)	H13A—C13—H13B	109.5

O1—C7—C8	121.7 (2)	C3—C13—H13C	109.5
N1—C7—C8	114.3 (2)	H13A—C13—H13C	109.5
C9—C8—C7	111.8 (2)	H13B—C13—H13C	109.5
C9—C8—H8A	109.2	C5—C14—H14A	109.5
C7—C8—H8A	109.2	C5—C14—H14B	109.5
C9—C8—H8B	109.2	H14A—C14—H14B	109.5
C7—C8—H8B	109.2	C5—C14—H14C	109.5
H8A—C8—H8B	107.9	H14A—C14—H14C	109.5
C10—C9—C8	114.1 (2)	H14B—C14—H14C	109.5
C10—C9—H9A	108.7	C7—N1—C1	129.0 (2)
C8—C9—H9A	108.7	C7—N1—H1N	116 (2)
C10—C9—H9B	108.7	C1—N1—H1N	115 (2)
C8—C9—H9B	108.7	C10—O3—C11	118.0 (2)
C6—C1—C2—C3	0.0	C7—C8—C9—C10	180.0
N1—C1—C2—C3	180.0	C8—C9—C10—O2	0.0
C1—C2—C3—C4	0.0	C8—C9—C10—O3	180.0
C1—C2—C3—C13	180.0	H11 ⁱ —C11—C12—H12B ⁱ	-54 (2)
C2—C3—C4—C5	0.0	O3—C11—C12—H12B ⁱ	64.8 (16)
C13—C3—C4—C5	180.0	O1—C7—N1—C1	0.0
C3—C4—C5—C6	0.0	C8—C7—N1—C1	180.0
C3—C4—C5—C14	180.0	C6—C1—N1—C7	0.0
C2—C1—C6—C5	0.0	C2—C1—N1—C7	180.0
N1—C1—C6—C5	180.0	O2—C10—O3—C11	0.0
C4—C5—C6—C1	0.0	C9—C10—O3—C11	180.0
C14—C5—C6—C1	180.0	H11 ⁱ —C11—O3—C10	-61.7 (13)
O1—C7—C8—C9	0.0	C12—C11—O3—C10	180.0
N1—C7—C8—C9	180.0		

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

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

$D\cdots H$	$D—H$	$H\cdots A$	$D\cdots A$	$D—H\cdots A$
N1—H1N ⁱⁱ —O2 ⁱⁱ	0.85 (3)	2.15 (3)	2.995 (3)	176 (3)

Symmetry code: (ii) $-x+1, -y, -z$.