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## Structure Reports

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**(E)-N-(3,4-Dimethoxyphenethyl)-3-methoxybut-2-enamide**

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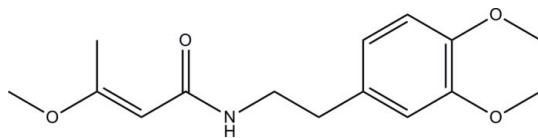
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Key indicators: single-crystal X-ray study;  $T = 173$  K; mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å;  $R$  factor = 0.059;  $wR$  factor = 0.133; data-to-parameter ratio = 14.3.

In the crystal of the title compound,  $\text{C}_{15}\text{H}_{21}\text{NO}_4$ , intermolecular  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bonds link molecules related by translation along the  $c$  axis into hydrogen-bonded chains.  $\text{C}-\text{H}\cdots\text{O}$  links are also present. The dihedral angle between benzene ring and enamide group is  $50.08$  ( $3^\circ$ )

## Related literature

For the applications of the title compound, see: Bernhard & Snieckus (1971); Ma *et al.* (2006). For bond-length data, see Allen *et al.* (1987).



## Experimental

## Crystal data

$\text{C}_{15}\text{H}_{21}\text{NO}_4$   
 $M_r = 279.33$

Monoclinic,  $P2_1/c$   
 $a = 12.509$  (3) Å

$b = 14.930$  (3) Å  
 $c = 8.2998$  (17) Å  
 $\beta = 107.59$  ( $3^\circ$ )  
 $V = 1477.5$  (5) Å<sup>3</sup>  
 $Z = 4$

Mo  $K\alpha$  radiation  
 $\mu = 0.09$  mm<sup>-1</sup>  
 $T = 173$  K  
 $0.30 \times 0.20 \times 0.15$  mm

## Data collection

Rigaku Mercury CCD/AFC diffractometer  
Absorption correction: multi-scan (*CrystalClear*; Rigaku, 2007)  
 $T_{\min} = 0.973$ ,  $T_{\max} = 0.987$

10754 measured reflections  
2591 independent reflections  
2435 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.049$

## Refinement

$R[F^2 > 2\sigma(F^2)] = 0.059$   
 $wR(F^2) = 0.133$   
 $S = 1.18$   
2591 reflections

181 parameters  
H-atom parameters constrained  
 $\Delta\rho_{\text{max}} = 0.16$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.18$  e Å<sup>-3</sup>

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1A}\cdots\text{O3}^i$	0.88	1.96	2.842 (2)	176
$\text{C15}-\text{H15A}\cdots\text{O1}^{ii}$	0.98	2.48	3.434 (3)	164

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

Data collection: *CrystalClear* (Rigaku, 2007); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HG2630).

## References

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## supporting information

*Acta Cryst.* (2010). E66, o432 [https://doi.org/10.1107/S1600536810001972]

**(*E*)-*N*-(3,4-Dimethoxyphenethyl)-3-methoxybut-2-enamide****Xiang Li****S1. Comment**

The title compound (*E*)-*N*-(3,4-dimethoxyphenethyl)-3-methoxybut-2-enamide was an important intermediate to the 3,4-dihydroisoquinoline and some other heterocyclic compounds (Bernhard & Snieckus, 1971; Ma *et al.*, 2006). In this paper, we use 3,4-dimethoxyphenethylamine and 3-methoxy-2-butenoyl chloride to synthesize the title compound and report its crystal structure here.

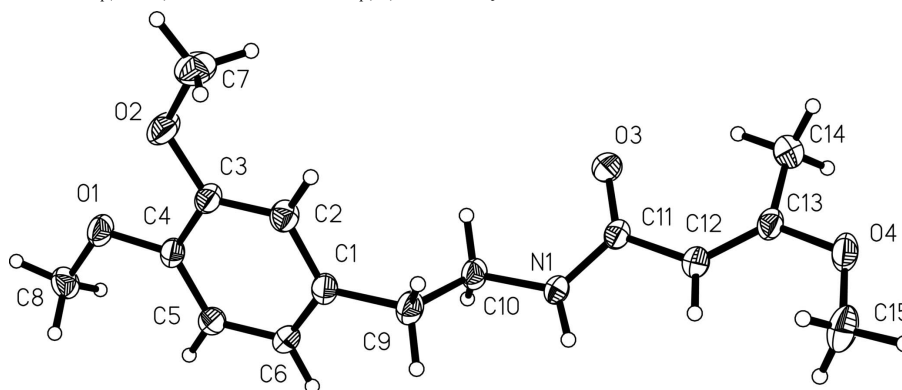
The title compound  $C_{15}H_{21}NO_4$  (Fig. 1), all bond lengths in the molecule are normal (Allen *et al.*, 1987). The intermolecular N—H $\cdots$ O hydrogen bonds [ $N\cdots O$  2.842 (2) Å] link the molecules related by translation along *c* axis into hydrogen-bonded chains.

**S2. Experimental**

3,4-dimethoxyphenethylamine (20 mmol) was solved in  $CH_2Cl_2$ ,  $Et_3N$  (30 mmol) was added, then 3-methoxy-2-butenoyl chloride (20 mmol) was added during 30 min at 273 K, after react 2 h at room temperature, the solution was washed with water, the organic layer was separated, dried with  $Na_2SO_4$ , evaporated to obtain the primary product, the pure product was isolated by recrystallization from ethyl acetate. (4.74 g, 84.9%). Single crystals suitable for X-ray measurements were obtained by recrystallization from ethyl acetate at room temperature.

**S3. Refinement**

H atoms were positioned geometrically and refined using a riding model, with C—H = 0.93–0.97 Å and N—H = 0.86 Å; with  $U_{iso}(H) = 1.2$  times  $U_{eq}(C, N)$  and 1.5 times  $U_{eq}(C)$  for methyl H atoms.

**Figure 1**

The molecular structure of (I), with atom labels and 40% probability displacement ellipsoids for non-H atoms.

*(E)*-*N*-(3,4-Dimethoxyphenethyl)-3-methoxybut-2-enamide*Crystal data*

$C_{15}H_{21}NO_4$	$F(000) = 600$
$M_r = 279.33$	$D_x = 1.256 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
Hall symbol: -P 2ybc	Cell parameters from 4588 reflections
$a = 12.509 (3) \text{ \AA}$	$\theta = 1.4\text{--}27.5^\circ$
$b = 14.930 (3) \text{ \AA}$	$\mu = 0.09 \text{ mm}^{-1}$
$c = 8.2998 (17) \text{ \AA}$	$T = 173 \text{ K}$
$\beta = 107.59 (3)^\circ$	Rod, colorless
$V = 1477.5 (5) \text{ \AA}^3$	$0.30 \times 0.20 \times 0.15 \text{ mm}$
$Z = 4$	

*Data collection*

Rigaku Mercury CCD/AFC diffractometer	10754 measured reflections
Radiation source: Sealed Tube	2591 independent reflections
Graphite Monochromator monochromator	2435 reflections with $I > 2\sigma(I)$
$\varphi$ and $\omega$ scans	$R_{\text{int}} = 0.049$
Absorption correction: multi-scan ( <i>CrystalClear</i> ; Rigaku, 2007)	$\theta_{\text{max}} = 25.0^\circ$ , $\theta_{\text{min}} = 2.7^\circ$
$T_{\text{min}} = 0.973$ , $T_{\text{max}} = 0.987$	$h = -14 \rightarrow 14$
	$k = -17 \rightarrow 17$
	$l = -9 \rightarrow 8$

*Refinement*

Refinement on $F^2$	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.059$	H-atom parameters constrained
$wR(F^2) = 0.133$	$w = 1/[\sigma^2(F_o^2) + (0.045P)^2 + 0.6856P]$
$S = 1.18$	where $P = (F_o^2 + 2F_c^2)/3$
2591 reflections	$(\Delta/\sigma)_{\text{max}} < 0.001$
181 parameters	$\Delta\rho_{\text{max}} = 0.16 \text{ e \AA}^{-3}$
0 restraints	$\Delta\rho_{\text{min}} = -0.18 \text{ e \AA}^{-3}$
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  $F^2 > \sigma(F^2)$  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 ( $\text{\AA}^2$ )*

	$x$	$y$	$z$	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	1.17428 (12)	0.48028 (10)	0.66017 (19)	0.0356 (4)
O2	1.16017 (13)	0.30958 (10)	0.6857 (2)	0.0446 (4)
O3	0.57963 (13)	0.16546 (10)	0.56367 (18)	0.0352 (4)
O4	0.36035 (13)	0.06101 (10)	0.8340 (2)	0.0417 (4)

N1	0.62813 (14)	0.26957 (12)	0.7715 (2)	0.0308 (4)
H1A	0.6128	0.2920	0.8601	0.037*
C1	0.91080 (17)	0.37551 (15)	0.8108 (3)	0.0303 (5)
C2	0.99168 (17)	0.32078 (14)	0.7740 (3)	0.0320 (5)
H2A	0.9871	0.2576	0.7838	0.038*
C3	1.07767 (17)	0.35758 (14)	0.7239 (3)	0.0307 (5)
C4	1.08529 (16)	0.45083 (14)	0.7088 (2)	0.0285 (5)
C5	1.00533 (18)	0.50451 (14)	0.7431 (3)	0.0321 (5)
H5A	1.0093	0.5677	0.7322	0.038*
C6	0.91870 (18)	0.46703 (15)	0.7936 (3)	0.0331 (5)
H6A	0.8642	0.5050	0.8167	0.040*
C7	1.1604 (2)	0.21485 (16)	0.7102 (3)	0.0484 (6)
H7A	1.2231	0.1881	0.6793	0.073*
H7B	1.0896	0.1895	0.6390	0.073*
H7C	1.1688	0.2019	0.8292	0.073*
C8	1.1861 (2)	0.57474 (15)	0.6482 (3)	0.0379 (5)
H8A	1.2524	0.5878	0.6126	0.057*
H8B	1.1948	0.6022	0.7587	0.057*
H8C	1.1192	0.5992	0.5650	0.057*
C9	0.81715 (17)	0.33412 (16)	0.8642 (3)	0.0342 (5)
H9A	0.7954	0.3757	0.9418	0.041*
H9B	0.8445	0.2781	0.9271	0.041*
C10	0.71449 (17)	0.31312 (15)	0.7155 (3)	0.0316 (5)
H10A	0.6843	0.3693	0.6557	0.038*
H10B	0.7364	0.2735	0.6351	0.038*
C11	0.57000 (16)	0.19712 (14)	0.6969 (3)	0.0283 (5)
C12	0.49832 (17)	0.15920 (14)	0.7913 (3)	0.0304 (5)
H12A	0.5082	0.1820	0.9017	0.036*
C13	0.42041 (17)	0.09551 (14)	0.7360 (3)	0.0321 (5)
C14	0.3818 (2)	0.05392 (18)	0.5650 (3)	0.0500 (7)
H14A	0.4248	0.0788	0.4944	0.075*
H14B	0.3019	0.0667	0.5129	0.075*
H14C	0.3933	-0.0110	0.5751	0.075*
C15	0.3837 (2)	0.09322 (19)	1.0043 (3)	0.0500 (7)
H15A	0.3349	0.0627	1.0594	0.075*
H15B	0.3700	0.1579	1.0026	0.075*
H15C	0.4623	0.0811	1.0669	0.075*

Atomic displacement parameters ( $\text{\AA}^2$ )

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
O1	0.0300 (8)	0.0331 (8)	0.0494 (9)	-0.0036 (6)	0.0206 (7)	0.0028 (7)
O2	0.0372 (9)	0.0326 (9)	0.0727 (12)	0.0074 (7)	0.0298 (8)	0.0031 (8)
O3	0.0398 (9)	0.0370 (9)	0.0334 (9)	-0.0040 (7)	0.0180 (7)	-0.0029 (7)
O4	0.0338 (9)	0.0372 (9)	0.0607 (11)	-0.0059 (7)	0.0242 (8)	0.0051 (8)
N1	0.0318 (10)	0.0340 (10)	0.0330 (9)	-0.0074 (8)	0.0193 (8)	-0.0034 (8)
C1	0.0255 (11)	0.0361 (12)	0.0296 (11)	-0.0041 (9)	0.0087 (8)	-0.0002 (9)
C2	0.0300 (11)	0.0301 (12)	0.0365 (12)	0.0005 (9)	0.0109 (9)	0.0042 (9)

C3	0.0244 (10)	0.0324 (12)	0.0369 (12)	0.0027 (9)	0.0116 (9)	0.0016 (9)
C4	0.0230 (10)	0.0332 (11)	0.0301 (11)	-0.0028 (8)	0.0092 (8)	0.0003 (9)
C5	0.0319 (11)	0.0270 (11)	0.0399 (12)	-0.0020 (9)	0.0147 (9)	-0.0024 (9)
C6	0.0291 (11)	0.0347 (12)	0.0401 (12)	0.0000 (9)	0.0173 (9)	-0.0043 (9)
C7	0.0537 (16)	0.0342 (13)	0.0615 (16)	0.0135 (11)	0.0237 (13)	0.0057 (12)
C8	0.0401 (13)	0.0358 (13)	0.0407 (13)	-0.0122 (10)	0.0165 (10)	-0.0027 (10)
C9	0.0303 (12)	0.0414 (13)	0.0341 (12)	-0.0049 (10)	0.0144 (9)	0.0021 (10)
C10	0.0304 (11)	0.0358 (12)	0.0320 (12)	-0.0059 (9)	0.0147 (9)	0.0019 (9)
C11	0.0240 (10)	0.0298 (11)	0.0331 (12)	0.0011 (9)	0.0115 (9)	0.0029 (9)
C12	0.0294 (11)	0.0322 (12)	0.0318 (11)	-0.0025 (9)	0.0125 (9)	0.0018 (9)
C13	0.0244 (10)	0.0278 (11)	0.0476 (13)	0.0024 (9)	0.0164 (9)	0.0042 (9)
C14	0.0434 (14)	0.0491 (15)	0.0623 (17)	-0.0151 (12)	0.0235 (12)	-0.0199 (13)
C15	0.0445 (15)	0.0637 (17)	0.0492 (15)	-0.0082 (12)	0.0252 (12)	0.0135 (13)

*Geometric parameters (Å, °)*

O1—C4	1.366 (2)	C7—H7B	0.9800
O1—C8	1.425 (3)	C7—H7C	0.9800
O2—C3	1.370 (2)	C8—H8A	0.9800
O2—C7	1.429 (3)	C8—H8B	0.9800
O3—C11	1.241 (2)	C8—H8C	0.9800
O4—C13	1.364 (2)	C9—C10	1.521 (3)
O4—C15	1.436 (3)	C9—H9A	0.9900
N1—C11	1.345 (3)	C9—H9B	0.9900
N1—C10	1.453 (2)	C10—H10A	0.9900
N1—H1A	0.8800	C10—H10B	0.9900
C1—C6	1.380 (3)	C11—C12	1.471 (3)
C1—C2	1.404 (3)	C12—C13	1.339 (3)
C1—C9	1.505 (3)	C12—H12A	0.9500
C2—C3	1.379 (3)	C13—C14	1.490 (3)
C2—H2A	0.9500	C14—H14A	0.9800
C3—C4	1.404 (3)	C14—H14B	0.9800
C4—C5	1.377 (3)	C14—H14C	0.9800
C5—C6	1.392 (3)	C15—H15A	0.9800
C5—H5A	0.9500	C15—H15B	0.9800
C6—H6A	0.9500	C15—H15C	0.9800
C7—H7A	0.9800		
C4—O1—C8	116.74 (16)	H8B—C8—H8C	109.5
C3—O2—C7	117.00 (18)	C1—C9—C10	112.77 (17)
C13—O4—C15	118.37 (18)	C1—C9—H9A	109.0
C11—N1—C10	124.27 (17)	C10—C9—H9A	109.0
C11—N1—H1A	117.9	C1—C9—H9B	109.0
C10—N1—H1A	117.9	C10—C9—H9B	109.0
C6—C1—C2	118.36 (19)	H9A—C9—H9B	107.8
C6—C1—C9	121.55 (19)	N1—C10—C9	111.08 (17)
C2—C1—C9	120.1 (2)	N1—C10—H10A	109.4
C3—C2—C1	120.8 (2)	C9—C10—H10A	109.4

C3—C2—H2A	119.6	N1—C10—H10B	109.4
C1—C2—H2A	119.6	C9—C10—H10B	109.4
O2—C3—C2	124.9 (2)	H10A—C10—H10B	108.0
O2—C3—C4	114.96 (18)	O3—C11—N1	122.13 (18)
C2—C3—C4	120.15 (19)	O3—C11—C12	124.53 (19)
O1—C4—C5	125.55 (19)	N1—C11—C12	113.31 (18)
O1—C4—C3	115.39 (18)	C13—C12—C11	126.0 (2)
C5—C4—C3	119.06 (19)	C13—C12—H12A	117.0
C4—C5—C6	120.6 (2)	C11—C12—H12A	117.0
C4—C5—H5A	119.7	C12—C13—O4	122.6 (2)
C6—C5—H5A	119.7	C12—C13—C14	128.0 (2)
C1—C6—C5	121.0 (2)	O4—C13—C14	109.43 (19)
C1—C6—H6A	119.5	C13—C14—H14A	109.5
C5—C6—H6A	119.5	C13—C14—H14B	109.5
O2—C7—H7A	109.5	H14A—C14—H14B	109.5
O2—C7—H7B	109.5	C13—C14—H14C	109.5
H7A—C7—H7B	109.5	H14A—C14—H14C	109.5
O2—C7—H7C	109.5	H14B—C14—H14C	109.5
H7A—C7—H7C	109.5	O4—C15—H15A	109.5
H7B—C7—H7C	109.5	O4—C15—H15B	109.5
O1—C8—H8A	109.5	H15A—C15—H15B	109.5
O1—C8—H8B	109.5	O4—C15—H15C	109.5
H8A—C8—H8B	109.5	H15A—C15—H15C	109.5
O1—C8—H8C	109.5	H15B—C15—H15C	109.5
H8A—C8—H8C	109.5		
C6—C1—C2—C3	0.7 (3)	C9—C1—C6—C5	-179.60 (19)
C9—C1—C2—C3	179.61 (19)	C4—C5—C6—C1	0.1 (3)
C7—O2—C3—C2	-4.2 (3)	C6—C1—C9—C10	89.1 (3)
C7—O2—C3—C4	175.8 (2)	C2—C1—C9—C10	-89.8 (2)
C1—C2—C3—O2	179.9 (2)	C11—N1—C10—C9	-134.8 (2)
C1—C2—C3—C4	-0.1 (3)	C1—C9—C10—N1	177.36 (18)
C8—O1—C4—C5	1.6 (3)	C10—N1—C11—O3	-5.9 (3)
C8—O1—C4—C3	-178.34 (18)	C10—N1—C11—C12	172.39 (18)
O2—C3—C4—O1	-0.7 (3)	O3—C11—C12—C13	-11.6 (3)
C2—C3—C4—O1	179.31 (18)	N1—C11—C12—C13	170.1 (2)
O2—C3—C4—C5	179.38 (19)	C11—C12—C13—O4	177.17 (19)
C2—C3—C4—C5	-0.6 (3)	C11—C12—C13—C14	-5.3 (4)
O1—C4—C5—C6	-179.30 (19)	C15—O4—C13—C12	-1.4 (3)
C3—C4—C5—C6	0.6 (3)	C15—O4—C13—C14	-179.3 (2)
C2—C1—C6—C5	-0.7 (3)		

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

<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>
N1—H1A $\cdots$ O3 <sup>i</sup>	0.88	1.96	2.842 (2)	176

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C15—H15A···O1 <sup>ii</sup>	0.98	2.48	3.434 (3)	164
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Symmetry codes: (i)  $x, -y+1/2, z+1/2$ ; (ii)  $x-1, -y+1/2, z+1/2$ .