

Acta Crystallographica Section C

**Crystal Structure
Communications**

ISSN 0108-2701

Methylmalonamide and ethylmalonamide

Jun Sakamoto *et al.*

Electronic paper

This paper is published electronically. It meets the data-validation criteria for publication in Acta Crystallographica Section C. The submission has been checked by a Section C Co-editor though the text in the 'Comments' section is the responsibility of the authors.

© 2000 International Union of Crystallography • Printed in Great Britain – all rights reserved

Methylmalonamide and ethylmalonamide

Jun Sakamoto,^{a*} Takashi Nakagawa,^a Nobuko Kanehisa,^b Yasushi Kai^b and Masahiro Katsura^a^aDepartment of Nuclear Engineering, Graduate School of Engineering, Osaka University, Yamada-Oka 2-1, Suita, Osaka 565-0871, Japan, and ^bDepartment of Applied Chemistry, Graduate School of Engineering, Osaka University, Yamada-Oka 2-1, Suita, Osaka 565-0871, Japan

Correspondence e-mail: jsaka@nucl.eng.osaka-u.ac.jp

Received 24 July 2000

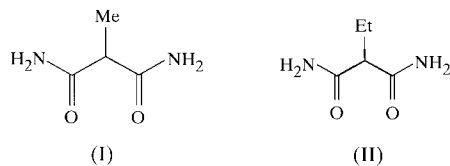
Accepted 2 August 2000

Data validation number: IUC0000207

In the title compounds, C₄H₈N₂O₂, (I), and C₅H₁₀N₂O₂, (II), respectively, the amide groups are rotated out of the central C—C—C plane by *ca* 76° in (I) and by 70–73° in (II). Compound (I) has crystallographic mirror symmetry perpendicular to the molecular plane.

Comment

The malonamides have a stronger extracting capacity than other diamides, which is due to the larger electron delocalization and the higher chelating abilities of malonamides (Siddall & Good, 1967). The title substituted malonamides, (I) and (II), were synthesized with the aim of studying their lipophilic suitability as extractants. The crystal structure analyses of (I) and (II) revealed different space groups, namely orthorhombic *Pnma* for (I) and monoclinic *P2₁/n* for (II). In the molecular structure of (I), the molecule is bisected



by a mirror plane passing through atoms H5, C3, C1 and H3. Bond lengths and angles are in the usual ranges. There are N—H...O intermolecular hydrogen bonds in (I), and N—H...O and C—H...O intermolecular hydrogen bonds in (II). In both title compounds, each amide O atom acts as an acceptor in two hydrogen bonds. These hydrogen bonds stabilize the intermolecular packing (Tables 1 and 2).

Experimental

A suspension of substituted diethyl malonate and NH₃ solution was refluxed for one week with vigorous stirring. Crystals of (I) and (II) were obtained by slow evaporation of an ethanol solution.

Compound (I)

Crystal data

C₄H₈N₂O₂
M_r = 116.12
 Orthorhombic, *Pnma*
a = 9.778 (2) Å
b = 10.879 (2) Å
c = 5.706 (2) Å
V = 607.0 (2) Å³
Z = 4
D_x = 1.271 Mg m⁻³

Mo *K*α radiation
 Cell parameters from 25 reflections
 θ = 13.8–14.9°
 μ = 0.102 mm⁻¹
T = 296.2 K
 Prismatic, colorless
 0.25 × 0.20 × 0.10 mm

Data collection

Rigaku AFC-5R diffractometer
 ω -2 θ scans
 Absorption correction: ψ scan (North *et al.*, 1968)
 T_{\min} = 0.980, T_{\max} = 0.997
 854 measured reflections
 735 independent reflections
 451 reflections with $I > 2\sigma(I)$

θ_{\max} = 27.51°
 h = -12 → 0
 k = 0 → 14
 l = 0 → 7
 3 standard reflections every 150 reflections
 intensity decay: 3.48%

Refinement

Refinement on F^2
 $R(F)$ = 0.0404
 $wR(F^2)$ = 0.1146
 S = 1.152
 735 reflections
 58 parameters

All H-atom parameters refined
 $w = 1/(\sigma^2(F_o^2) + \{0.050[\text{Max}(F_o^2, 0) + 2F_c^2]/3\}^2)$
 $(\Delta/\sigma)_{\max}$ = 0.0019
 $\Delta\rho_{\max}$ = 0.35 e Å⁻³
 $\Delta\rho_{\min}$ = -0.27 e Å⁻³

Table 1

Hydrogen-bonding geometry (Å, °) for (I).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N1—H1...O1 ⁱ	0.91 (2)	1.97 (2)	2.8445 (18)	162 (2)
N1—H2...O1 ⁱⁱ	0.97 (2)	1.987 (19)	2.943 (2)	169.4 (17)

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

Compound (II)

Crystal data

C₅H₁₀N₂O₂
M_r = 130.15
 Monoclinic, *P2₁/n*
a = 6.570 (1) Å
b = 9.502 (1) Å
c = 11.032 (1) Å
 β = 96.69 (1)°
V = 684.1 (2) Å³
Z = 4

D_x = 1.264 Mg m⁻³
 Mo *K*α radiation
 Cell parameters from 25 reflections
 θ = 12.8–13.5°
 μ = 0.098 mm⁻¹
T = 296.2 K
 Prismatic, colorless
 0.20 × 0.20 × 0.20 mm

Data collection

Rigaku AFC-5R diffractometer
 ω -2 θ scans
 Absorption correction: ψ scan (North *et al.*, 1968)
 T_{\min} = 0.974, T_{\max} = 0.998
 1830 measured reflections
 1589 independent reflections
 1077 reflections with $I > 2\sigma(I)$

R_{int} = 0.029
 θ_{\max} = 27.57°
 h = 0 → 8
 k = 0 → 12
 l = -14 → 14
 3 standard reflections every 150 reflections
 intensity decay: 0.50%

Refinement

Refinement on F^2
 $R(F)$ = 0.051
 $wR(F^2)$ = 0.120
 S = 1.131
 1589 reflections
 121 parameters

All H-atom parameters refined
 $w = 1/(\sigma^2(F_o^2) + \{0.048[\text{Max}(F_o^2, 0) + 2F_c^2]/3\}^2)$
 $(\Delta/\sigma)_{\max}$ = 0.0033
 $\Delta\rho_{\max}$ = 0.60 e Å⁻³
 $\Delta\rho_{\min}$ = -0.27 e Å⁻³

Table 2

Hydrogen-bonding geometry (Å, °) for (II).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N2—H4...O2 ⁱ	0.85 (3)	2.08 (3)	2.906 (2)	164 (2)
N2—H3...O1 ⁱⁱ	0.90 (2)	2.10 (2)	2.996 (2)	174.5 (18)
N1—H1...O1 ⁱ	0.90 (3)	1.98 (3)	2.859 (2)	165 (2)
N1—H2...O2 ⁱⁱⁱ	0.85 (2)	2.14 (2)	2.982 (2)	167.1 (19)
C1—H5...O2 ⁱ	0.967 (19)	2.59 (2)	3.441 (2)	147.5 (17)

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

All the H atoms were located from the difference Fourier map and refined isotropically. In (I), the C—H range is 0.95 (3)–0.96 (3) Å and the N—H range is 0.90 (2)–0.96 (2) Å. In (II), the C—H range is 0.95 (3)–1.00 (3) Å and the N—H range is 0.85 (3)–0.91 (2) Å.

For both compounds, data collection: *MSC/AFC Diffractometer Control Software* (Molecular Structure Corporation, 1990); cell refinement: *MSC/AFC Diffractometer Control Software*; data reduc-

tion: *TEXSAN* (Molecular Structure Corporation & Rigaku Corporation, 1999); program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1993); program(s) used to refine structure: *TEXSAN*; software used to prepare material for publication: *TEXSAN*.

References

- Altomare, A., Cascarano, M., Giacovazzo, C. & Guagliardi, A. (1993). *J. Appl. Cryst.* **26**, 343–350.
- Molecular Structure Corporation (1990). *MSC/AFC Diffractometer Control Software*. MSC, 3200 Research Forest Drive, The Woodlands, TX 77381, USA.
- Molecular Structure Corporation & Rigaku Corporation (1999). *TEXSAN*. Version 1.10. MSC, 3200 Research Forest Drive, The Woodlands, TX 77381, USA, and Rigaku, 3–9–12 Akishima, Tokyo, Japan.
- North, A. C. T., Phillips, D. C. & Mathews, F. S. (1968). *Acta Cryst.* **A24**, 351–359.
- Siddall, T. H. & Good, M. L. (1967). *J. Inorg. Nucl. Chem.* **29**, 149.