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The *E* and *Z* isomers of 3-(benzoxazol-2-yl)prop-2-enoic acid

Jose G. Trujillo-Ferrara,^a Itzia I. Padilla-Martínez,^b Francisco J. Martínez-Martínez,^b Herbert Höpfl,^c Norberto Farfan-García^d and Efrén V. García-Báez^b*

^aSección de Graduados y Departamento de Bioquímica, Escuela Superior de Medicina, Instituto Politécnico Nacional, México, DF 11340, Mexico, ^bUnidad Profesional Interdisciplinaria de Biotecnología, Instituto Politécnico Nacional, Avenida Acueducto s/n, Barrio La Laguna Ticomán, México, DF 07340, Mexico, ^cCentro de Investigaciónes Químicas, Universidad Autónoma del Estado de Morelos, Cuernava Morelos, Mexico, and ^dCentro de Investigación y Estudios Avanzados del Instituto Politecnico Nacional, Avenida IPN 2508, México, DF 07330, Mexico Correspondence e-mail: vgarcia@acei.upibi.ipn.mx

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The carboxylic acid group and the double bond are coplanar in (E)-3-(benzoxazol-2-yl)prop-2-enoic acid, $C_{10}H_7NO_3$, whereas in isomeric (Z)-3-(benzoxazol-2-yl)prop-2-enoic acid, also $C_{10}H_7NO_3$, they are almost orthogonal. In both isomers, a strong $O-H\cdots N$ hydrogen bond, with the carboxylic acid group as a donor and the pyridine-like N atom as an acceptor, and weak $C-H\cdots O$ interactions contribute to the observed supramolecular structures, which are completed by $\pi-\pi$ stacking interactions between oxazole and benzenoid rings.

Comment

 γ -Aminobutyric acid (GABA) is the major inhibitory neurotransmitter in the central nervous system (CNS). GABA operates through three different classes of receptors, consisting of the ionotropic GABA_A and GABA_C receptors, and the G-protein coupled receptor GABA_B (Chebib & Johnston, 2000). Hypoactivity of the GABA neuronal function has been associated with pain and neurological disorders such as epilepsy, Huntington's chorea, anxiety and sleep disorders. In contrast, GABA-mediated hyperactivity has been suggested to be an important component of schizophrenic symptoms (Frølund *et al.*, 2002). In order to characterize these receptors pharmacologically, recent studies of the synthesis



and characterization of several GABA agonists (Chambers et al., 2003; Frølund et al. 1995) and antagonists (Frølund et al.,



Figure 1 The molecular structure of (I-E), showing displacement ellipsoids at the 30% probability level.

2002) have been undertaken. GABA is a conformationally flexible molecule, capable of adopting receptor subtype-specific conformations; in this context, we report the results of structural analyses of the E and Z geometric isomers of 3-(benzoxazol-2-yl)prop-2-enoic acid [(I-E) and (I-Z), respectively], which are analogues of GABA.

A one-pot synthesis was carried out by mixing equimolar quantities of maleic anhydride and o-aminophenol in tetrahydrofuran (THF). Careful regulation of the temperature to 323 K, using Ac_2O as catalyst, yielded isomer (I-Z); isomer (I-E) was obtained when the temperature was raised to 343 K and an excess of Ac_2O was used. It appears that (I-Z) is the kinetic product and (I-E) is the thermodynamic product, since the former is observed by thin-layer chromatography in the early stages of the reaction. The molecular structures of (I-E)and (I-Z) are shown in Figs. 1 and 2, respectively, while selected geometric parameters are given in Tables 1 and 3, respectively. The bond lengths and angles are very similar in the two isomers, as is the anti disposition of the double bond and the imine group. However, the carboxy and benzoxazole groups are on opposite sides in (I-E) and on the same side in (I-Z).

The carboxy group in (I-*E*) is coplanar with the double bond, whereas in (I-*Z*) these groups are almost orthogonal $[O1B-C1-C2-C3 = -4.1 (3) \text{ and } 103.7 (3)^\circ, \text{ respectively}]$, in agreement with the conformation reported for other (*Z*)-propenoic acids (Stomberg *et al.*, 1995). This conformational difference between the isomers determines the observed crystal packing (see below). Isomer (I-*E*) forms





The molecular structure of (I-Z), showing displacement ellipsoids at the 30% probability level.

supramolecular sheets in the $(10\overline{2})$ plane through both strong (Steiner, 2002) and weak (Desiraju, 1996) hydrogen-bonding interactions. The hydrogen-bonding geometry for isomer (I-*E*) is listed in Table 2. The carboxylic acid donor and pyridine-like N-atom acceptor form a strong $O1A - H1A \cdots N3^{\prime i}$ interaction [symmetry code: (i) $x - 1, \frac{1}{2} - y, -\frac{1}{2} + z$], which is complemented by a weak $C5' - H5' \cdots O1B^{ii}$ interaction [symmetry code: (ii) $1 - x, y - \frac{1}{2}, \frac{1}{2} - z$] between an aromatic H-atom donor and a carbonyl O-atom acceptor (Fig. 3).

Centrosymmetric sheets pack via π - π stacking interactions (Hunter et al., 1991; Singh & Thornton, 1990) between oxazole and benzenoid rings along the c axis. The mean intercentroid and interplanar distances [3.707 (1) and 3.435 (1) Å, respectively] at the (1 - x, -y, -z) symmetry position indicate a face-to-face π - π stacking arrangement, whereas the distances [4.4487 (11) and 3.38 (2) A] at the (1 - x, -y, 1 - z)symmetry position indicate a parallel displaced π - π stacking arrangement. The former arrangement is overwhelmingly preferred in the stacking of electron-rich and electron-deficient aromatic rings, such as benzene and hexafluorobenzene, whose 1:1 complex has mean intercentroid and interplanar distances of 3.7 and 3.4 Å, respectively (Williams, 1993). The electron-deficient carbonyl C atom is located 3.36 (4) Å from the electron-rich double bond of the neighbouring molecule [symmetry code: (v) $x, \frac{1}{2} - y, \frac{1}{2} + z$], forming an angle of 68 (1)° with this double bond (Fig. 3). This distance is as short as the sum of the van der Waals radii of two C atoms (3.4 Å; Hunter et al., 1991), and the carbonyl group is parallel but slightly displaced relative to the double bond. Therefore, this interaction can be considered as a parallel-to-displaced olefin-type π - π stacking interaction (Kim *et al.*, 2000), which should be an important contributor to the slippage of the hydrogen-bonded supramolecular sheets of (I-E).

Molecules of (I-Z) are linked by a strong hydrogen-bonding interaction $[O1A - H1A \cdots N3'^{vi};$ symmetry code: (vi) $\frac{1}{2} + x$, $\frac{1}{2} + y$, z] between the carboxylic acid donor and the pyridinelike N-atom acceptor, and a weak interaction $[C3 - H3 \cdots O1B^{vii};$ symmetry code: (vii) $\frac{1}{2} - x$, $y - \frac{1}{2}, \frac{1}{2} - z$] involving a vinyl H-atom donor and the carbonyl O-atom acceptor (Fig. 4). As a consequence, a twisted eight-membered chain is



Figure 3

The supramolecular structure of (I-*E*), viewed along the *a* axis. Centrosymmetric sheets of molecules pack through face-to-face and parallel-displaced π - π stacking interactions. [Symmetry codes: (i) x - 1, $\frac{1}{2} - y, -\frac{1}{2} + z$; (ii) $1 - x, y - \frac{1}{2}, \frac{1}{2} - z$; (iii) 1 - x, -y, -z; (iv) 1 - x, -y, 1 - z; (v) $x, \frac{1}{2} - y, \frac{1}{2} + z$.]



Figure 4

The supramolecular structure of (I-Z), viewed along the [230] and [230] directions. [Symmetry codes: (vi) $\frac{1}{2} + x$, $\frac{1}{2} + y$, z; (vii) $\frac{1}{2} - x$, $y - \frac{1}{2}$, $\frac{1}{2} - z$; (viii) -x, 1 - y, -z.]

formed, whose topological motif is described by the C(8)[C(7)C(5)] graph-set descriptor (Bernstein *et al.*, 1995). This interaction is allowed because of the almost perpendicular conformation between the carboxy and double-bond groups. The hydrogen-bonding geometry for (I-Z) is listed in Table 4.

Centrosymmetric molecules of (I-Z) also form π - π stacks, viz. two almost perpendicular sets of columns developing in the [230] and [230] directions (Fig. 4). The mean intercentroid and interplanar distances between the oxazole [3.473 (1) and 3.376 (3) Å; symmetry code: -x, 1 - y, -z] and benzenoid rings [3.563 (1) and 3.30 (1) Å; symmetry code: $\frac{1}{2} - x$, $\frac{1}{2} - y$, -z] are also in the range expected for a face-to-face geometry (Hunter *et al.*, 1991; Singh & Thornton, 1990), with less slippage than observed for (I-*E*).

Experimental

The title propenoic acid derivatives (I-Z) and (I-E) were obtained by the reaction between equimolar quantities (1.0 mmol) of maleic anhydride and 2-aminophenol dissolved in THF. Ac₂O (0.1 mmol) was added as catalyst, and after heating at 323-328 K for 2 h, isomer (I-Z) was obtained. Under the same conditions, but heating at 343 K for 3 h with acetic anhydride (1.5 mol), isomer (I-E) was obtained. In both cases, THF was removed by evaporation and the resulting solid was treated with aqueous HCl (5%) until precipitation was complete. After washing three times with deionized water and drying, the products were recrystallized from ethanol solutions to give crystals suitable for X-ray analysis in approximately 70% yield. For (I-E), m.p. 500–502 K; IR (KBr, cm⁻¹): ν 1710 (COO), 1527 (C=N); ¹H NMR: δ 13.2 (*b*, 1H, OH), 7.84 (*d*, 1H, ³*J* = 8.6 Hz, H4'), 7.77 (*d*, 1H, ${}^{3}J = 8.6$ Hz, H7'), 7.52 (dd, 1H, ${}^{3}J = 8.6$ Hz, H6'), 7.45 (d, 1H, ${}^{3}J =$ 15.8 Hz, H3), 7.44 (*dd*, 1H, ³*J* = 8.6 Hz, H5'), 6.92 (*d*, 1H, ³*J* = 15.8 Hz, H2); ¹³C NMR: δ 166.0 (C1), 159.9 (C2'), 150.1 (C8'), 141.4 (C9'), 129.3 (C2), 128.2 (C3), 126.9 (C6'), 125.2 (C5'), 120.5 (C4'), 111.1 (C7'). For (I-Z), m.p. 396–397 K; IR (KBr, cm⁻¹): v 1707 (COO), 1527 (C=N); ¹H NMR: δ 13.2 (b, 1H, OH), 7.8 (dd, 1H, ³J = 8.5 Hz, ⁴J = 1.2 Hz, H4'), 7.7 (dd, 1H, ${}^{3}J = 8.0$ Hz, ${}^{4}J = 1.3$ Hz, H7'), 7.4 (ddd, 1H, ${}^{3}J = 8.6 \text{ Hz}, {}^{4}J = 1.3 \text{ Hz}, \text{H6}'), 7.5 (ddd, 1\text{H}, {}^{3}J = 8.6 \text{ Hz}, {}^{4}J = 1.3 \text{ Hz},$ H5'), 6.88 (d, 1H, ${}^{3}J$ = 12.0 Hz, H3), 6.66 (d, 1H, ${}^{3}J$ = 12.0 Hz, H2); ${}^{13}C$ NMR: § 167.0 (C1), 159.6 (C2'), 149.8 (C8'), 140.1 (C9'), 132.3 (C2), 120.0 (C3), 126.3 (C6'), 125.1 (C5'), 120.1 (C4'), 111.0 (C7').

 $\theta_{\rm max}=27.5^\circ$

 $h = -12 \rightarrow 12$ $k = -12 \rightarrow 12$

 $l = -23 \rightarrow 23$

Isomer (I-E)

Crystal data

C₁₀H₇NO₃ $M_r = 189.17$ Monoclinic, $P2_1/c$ a = 5.882(1) Å b = 19.694 (1) Åc = 7.4910(1) Å $\beta = 91.715 \ (10)^{\circ}$ $V = 867.37 (15) \text{ Å}^3$ Z = 4

Data collection

Enraf-Nonius CAD-4 diffractometer $\omega/2\theta$ scans 2082 measured reflections 2082 independent reflections 1554 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.011$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_a^2) + (0.058P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.037$	+ 0.105P]
$wR(F^2) = 0.124$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.15	$(\Delta/\sigma)_{\rm max} = 0.001$
2082 reflections	$\Delta \rho_{\rm max} = 0.19 \ {\rm e} \ {\rm \AA}^{-3}$
128 parameters	$\Delta \rho_{\rm min} = -0.18 \text{ e} \text{ \AA}^{-3}$
H-atom parameters constrained	

 $D_x = 1.449 \text{ Mg m}^{-3}$

Cell parameters from 24

 $0.50 \times 0.50 \times 0.40 \text{ mm}$

3 standard reflections

every 200 reflections

intensity decay: 2.5%

Mo $K\alpha$ radiation

reflections $\theta = 10 - 11^{\circ}$

 $\mu = 0.11 \text{ mm}^{-1}$

T = 293 (2) K

Block, yellow

 $\theta_{\rm max} = 28.0^{\circ}$

 $h = -7 \rightarrow 7$

 $k = 0 \rightarrow 25$

 $l = 0 \rightarrow 9$

Table 1

Selected geometric parameters (Å, $^{\circ}$) for isomer (I-E).

1.3588 (17) 1.3800 (16) 1.324 (2) 104.23 (10) 104.76 (11)	O1B-C1 N3'-C2' N3'-C9' N3'-C2'-C3 O1'-C2'-C3	1.1978 (19) 1.2987 (17) 1.3962 (18) 127.00 (12) 114.92 (12)
1.3800 (16) 1.324 (2) 104.23 (10) 104 76 (11)	N3' - C2' N3' - C9' N3' - C2' - C3 O1' - C2' - C3	1.2987 (17) 1.3962 (18) 127.00 (12) 114.02 (12)
1.324 (2) 104.23 (10) 104.76 (11)	N3' - C9' N3' - C2' - C3 O1' - C2' - N2'	1.3962 (18) 127.00 (12) 114.02 (12)
104.23 (10) 104.76 (11)	N3' - C2' - C3	127.00 (12)
104.23 (10) 104.76 (11)	N3' - C2' - C3	127.00 (12)
104 76 (11)	O1' C2' N2'	11402(12)
101.70(11)	01 - 02 - N5	114.92 (12)
111.12 (12)	O1′-C8′-C9′	107.77 (12)
124.61 (16)	N3′-C9′-C8′	108.32 (11)
124.25 (14)	N3'-C9'-C4'	131.53 (12)
118.08 (11)		. ,
2.5 (2)	N3' - C2' - C3 - C2	-178.33 (14)
	111.12 (12) 124.61 (16) 124.25 (14) 118.08 (11) 2.5 (2)	$\begin{array}{cccc} 111.12 & 01' - C8' - C9' \\ 124.61 & 16) & N3' - C9' - C8' \\ 124.25 & (14) & N3' - C9' - C4' \\ 118.08 & (11) \end{array}$

Table 2

Hydrogen-bonding geometry (Å, $^{\circ}$) for isomer (I-*E*).

	2	$D=\prod \cdots A$
1.89 2.46	2.6995 (16) 3.372 (2)	167 165
	1.89 2.46	1.89 2.6995 (16) 2.46 3.372 (2)

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

Isomer (I-Z)

Crystal data

C ₁₀ H ₇ NO ₃	$D_x = 1.414 \text{ Mg m}^{-3}$
$M_r = 189.17$	Mo $K\alpha$ radiation
Monoclinic, $C2/c$	Cell parameters from 1010
a = 9.8191 (19) Å	reflections
b = 9.905 (2) Å	$\theta = 1.6-32.0^{\circ}$
c = 18.320 (4) Å	$\mu = 0.11 \text{ mm}^{-1}$
$\beta = 94.037 \ (4)^{\circ}$	T = 293 (2) K
V = 1777.4 (6) Å ³	Block, yellow
Z = 8	$0.36 \times 0.21 \times 0.19 \text{ mm}$

Bruker SMART area-detector
diffractometer
φ and ω scans
9736 measured reflections
2013 independent reflections
1559 reflections with $I > 2\sigma(I)$
$R_{\rm int} = 0.035$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.045P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.053$	+ 1.031P]
$wR(F^2) = 0.127$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.05	$(\Delta/\sigma)_{\rm max} < 0.001$
2013 reflections	$\Delta \rho_{\rm max} = 0.23 \ {\rm e} \ {\rm \AA}^{-3}$
128 parameters	$\Delta \rho_{\rm min} = -0.18 \text{ e} \text{ Å}^{-3}$
H-atom parameters constrained	

Table 3

Selected geometric parameters (Å, $^{\circ}$) for isomer (I-Z).

O1'-C2'	1.359 (2)	O1B-C1	1.200 (2)
O1'-C8'	1.382 (2)	N3' - C2'	1.298 (2)
O1A - C1	1.305 (2)	N3′-C9′	1.399 (3)
C2'-O1'-C8'	104.51 (13)	O1'-C2'-C3	119.65 (15)
C2'-N3'-C9'	105.13 (15)	N3' - C2' - C3	125.74 (17)
O1A-C1-C2	113.76 (16)	O1'-C2'-N3'	114.59 (16)
O1B-C1-C2	121.59 (18)	O1′-C8′-C7′	128.29 (17)
O1A-C1-O1B	124.57 (18)	N3'-C9'-C4'	131.94 (18)
	70 5 (2)	01/ 02/ 02 02	2.2 (2)
01A - C1 - C2 - C3	-79.5 (3)	O1' - C2' - C3 - C2	-3.2(3)
O1 <i>B</i> -C1-C2-C3	103.7 (3)	N3' - C2' - C3 - C2	178.51 (19)

Table 4

Hydrogen-bonding geometry (Å, $^{\circ}$) for isomer (I-Z).

$D - H \cdot \cdot \cdot A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$O1A - H1A \cdots N3'^{vi}$	0.82	1.90	2.715 (2)	172
$C3-H3\cdots O1B^{vii}$	0.93	2.43	3.310 (3)	159

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

All H atoms were refined as riding on their parent atoms using SHELXL97 (Sheldrick, 1997) defaults [O-H = 0.82 Å, C-H =0.93 Å and $U_{iso}(H) = 1.2U_{eq}(C,O)$].

For (I-E), data collection: CAD-4 EXPRESS (Enraf-Nonius, 1992); cell refinement: CAD-4 EXPRESS; data reduction: JANA98 (Vaclav, 1998); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: WinGX (Farrugia, 1999); software used to prepare material for publication: SHELXL97.

For (I-Z), data collection: SMART (Bruker, 2000); cell refinement: SAINT (Bruker, 2000); data reduction: SAINT; program(s) used to solve structure: SHELXS97; program(s) used to refine structure: SHELXL97; molecular graphics: SHELXTL (Bruker, 2000); software used to prepare material for publication: SHELXL97 and WinGX.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: BM1572). Services for accessing these data are described at the back of the journal.

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