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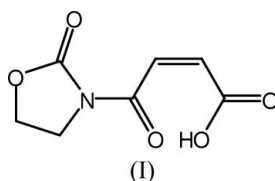
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## Key indicators

Single-crystal X-ray study  
 $T = 173$  K  
Mean  $\sigma(C-C) = 0.002$  Å  
 $R$  factor = 0.036  
 $wR$  factor = 0.109  
Data-to-parameter ratio = 14.1For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.**(Z)-3-(2-Oxo-1,3-oxazolidin-3-ylcarbonyl)-prop-2-enoic acid**In the title compound,  $C_7H_7NO_5$ , intermolecular  $O-H \cdots O$  hydrogen bonds connect two symmetry-related molecules, forming a pseudo-dimer.Received 3 January 2007  
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## Comment

In the past, furazolidone [*N*-(5-nitro-2-furfurylideneamino)-2-oxazolidinone] was once used as a veterinary drug for its obvious growth-promotion effect and good antibacterial activity. Since 1995, the use of furazolidone has been prohibited in food-animal production in many countries because of its carcinogenicity and mutagenicity (European Commission Regulation, 1995; Cooper *et al.*, 2004). However, as a result of the low costs of furazolidone and its treatment benefits, it has been proven by detecting the metabolites AOZ (3-amino-2-oxazolidinone) in animal-origin food that illegal abuse of furazolidone is still active (European Commission, 2005; Cooper *et al.*, 2004).



Among analytical methods for detecting AOZ residue, immunoassay is an efficient screening technique for the monitoring of illegal furazolidone. However, immunoassay formats reported are mainly based on the detection of the AOZ derivative NPAOZ (3-(2-nitrobenzylideneamino)oxazolidin-2-one), which must undergo a troublesome derivati-

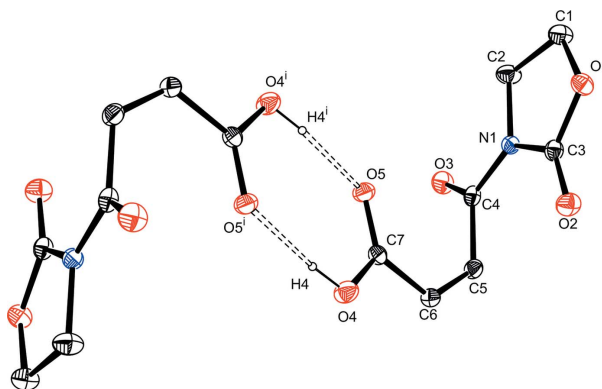


Figure 1

The structure of the dimer. Displacement ellipsoids are drawn at the 30% probability level.  $O-H \cdots O$  hydrogen bonds are shown as dashed lines. H atoms not involved in hydrogen bonding have been omitted for clarity. [Symmetry code: (i)  $1 - x, 1 - y, 1 - z$ .]

zation step (Cooper *et al.*, 2004; Franek *et al.*, 2006). As part of our investigation of immunoassay methods for directly detecting AOZ itself, we report here the synthesis and crystal structure of a new hapten, (I), for producing a specific antibody of AOZ.

The title compound is a non-aromatic AOZ analogue with a free carboxylic acid group, as shown in Fig. 1. The three C—O distances [C3—O2 = 1.198 (2), C4—O3 = 1.212 (2) and C7—O5 = 1.224 (2) Å] fall within the normal double-bond range; the C5—C6 bond length is 1.321 (2) Å, showing double-bond character. The bond distances within the O1/C1/C2/N1/C3 five-membered ring vary from 1.343 (2) to 1.506 (2) Å, indicating a non-aromatic character, which is also consistent with the NMR experiment. O—H...O hydrogen bonds (Table 1) link two adjacent molecules, forming a centrosymmetric dimer (Fig. 1).

## Experimental

To a solution of oxazolidinone (1.74 g, 0.02 mol), lithium chloride (0.84 g, 0.02 mol), and triethylamine (2.02 g, 0.02 mol) in dry THF (20 ml) was added maleic anhydride (2.35 g, 0.024 mol) at 253 K. The mixture was allowed to warm to 263 K and stirred overnight. THF was removed *in vacuo*. The residue was treated with saturated NaHCO<sub>3</sub> and adjusted to pH = 2 with concentrated hydrochloric acid. The mixture was then extracted with EtOAc, and the organic layer was washed with saturated NaCl and subsequently dried with Na<sub>2</sub>SO<sub>4</sub>. The solvents were removed under reduced pressure. The resulting solid was recrystallized from dichloromethane to give (I) in 92% yield. Crystals suitable for X-ray analysis were grown from an acetone–dichloromethane (1:2) solution.

### Crystal data

C <sub>7</sub> H <sub>7</sub> NO <sub>5</sub>	Z = 4
M <sub>r</sub> = 185.14	D <sub>x</sub> = 1.598 Mg m <sup>-3</sup>
Monoclinic, P2 <sub>1</sub> /c	Mo Kα radiation
a = 8.4248 (16) Å	μ = 0.14 mm <sup>-1</sup>
b = 5.0585 (9) Å	T = 173 (2) K
c = 18.060 (3) Å	Block, colourless
β = 91.128 (3)°	0.48 × 0.46 × 0.35 mm
V = 769.5 (2) Å <sup>3</sup>	

### Data collection

Bruker SMART diffractometer	4303 measured reflections
φ and ω scans	1679 independent reflections
Absorption correction: multi-scan (SADABS; Bruker, 2001)	1325 reflections with I > 2σ(I)
T <sub>min</sub> = 0.935, T <sub>max</sub> = 0.953	R <sub>int</sub> = 0.023
	θ <sub>max</sub> = 27.1°

### Refinement

Refinement on F <sup>2</sup>	w = 1/[σ <sup>2</sup> (F <sub>o</sub> <sup>2</sup> ) + (0.0618P) <sup>2</sup> + 0.1606P]
R[F <sup>2</sup> > 2σ(F <sup>2</sup> )] = 0.037	where P = (F <sub>o</sub> <sup>2</sup> + 2F <sub>c</sub> <sup>2</sup> )/3
wR(F <sup>2</sup> ) = 0.109	(Δ/σ) <sub>max</sub> < 0.001
S = 1.08	Δρ <sub>max</sub> = 0.30 e Å <sup>-3</sup>
1679 reflections	Δρ <sub>min</sub> = -0.20 e Å <sup>-3</sup>
119 parameters	
H-atom parameters constrained	

**Table 1**

Hydrogen-bond geometry (Å, °).

D—H...A	D—H	H...A	D...A	D—H...A
O4—H4...O5 <sup>i</sup>	0.82	1.87	2.6817 (16)	168

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

All H atoms were positioned geometrically and refined as riding, with C—H = 0.93 (aromatic) or 0.97 Å (methylene) and O—H = 0.82 Å, and with U<sub>iso</sub>(H) = 1.2U<sub>eq</sub>(C, O).

Data collection: SMART (Bruker, 2001); cell refinement: SAINT (Bruker, 2001); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Siemens, 1994); program(s) used to refine structure: SHELXTL; molecular graphics: ORTEPIII (Burnett & Johnson, 1996) and ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: SHELXTL.

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