

## 3-(2-Chlorophenyl)-N-(2-cyano-4-methyl-2-pentyl)-5-methylisoxazole-4-carboxamide

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## Key indicators

Single-crystal X-ray study  
 $T = 294\text{ K}$   
Mean  $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$   
 $R$  factor = 0.045  
 $wR$  factor = 0.150  
Data-to-parameter ratio = 17.1For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The title compound,  $\text{C}_{18}\text{H}_{20}\text{ClN}_3\text{O}_2$ , is a potent new herbicide. X-ray analysis reveals that the benzene ring is twisted away from the isoxazole ring by  $67.7(1)^\circ$ . The crystal packing is stabilized by intermolecular  $\text{N}-\text{H}\cdots\text{N}$  and  $\text{C}-\text{H}\cdots\text{O}$  hydrogen bonds.

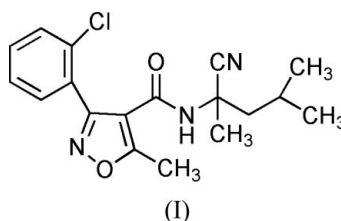
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## Comment

Isoxazole derivatives represent one of the most active classes of compounds, possessing a wide spectrum of biological activity. They are widely used in agrochemicals and pharmaceuticals (He *et al.*, 2000). Similarly, aminoacetonitrile derivatives exhibit various biological activities, *e.g.* insecticidal (Andoh *et al.*, 1999; Ducray, Goebel & Bouvier, 2003; Ducray, Goebel & Fruechtel, 2003), fungicidal (Walker & Baker 1979; Kim *et al.*, 2004) and herbicidal (Berliner & Richter, 1971). In view of these properties and in a continuation of our interest in the chemistry of isoxazoles, we have attempted to synthesize a series of 3-(2-chlorophenyl)-5-methylisoxazole-4-carboxamide derivatives, some of which have comparatively high herbicidal activity. The crystal structure determination of the title compound, (I), was undertaken to investigate the relationship between structure and herbicidal activity.



The molecular structure of (I) is shown in Fig. 1. The X-ray analysis reveals that the benzene ring is twisted away from the isoxazole ring by  $67.7(1)^\circ$ . The dihedral angle between the isoxazole ring and the carboxamide  $\text{N}2/\text{O}2/\text{C}11/\text{C}12$  plane is  $30.2(1)^\circ$ . The  $\text{C}11=\text{O}2$  distance of  $1.220(2)\text{ \AA}$  is comparable to the mean value of the  $\text{C}=\text{O}$  distance [ $1.231(12)\text{ \AA}$ ; Zhong *et al.*, 2004] in amides. In the crystal structure, adjacent molecules are linked by weak  $\text{N}-\text{H}\cdots\text{N}$  and  $\text{C}-\text{H}\cdots\text{O}$  hydrogen bonds (Table 2).

## Experimental

Compound (I) was prepared according to the reported procedure of Liu *et al.* (2004). Single crystals suitable for X-ray diffraction were obtained by recrystallization from a mixture of ethyl acetate and petroleum ether (1:5 *v/v*).

Crystal data

C<sub>18</sub>H<sub>20</sub>ClN<sub>3</sub>O<sub>2</sub>  
*M<sub>r</sub>* = 345.82  
 Monoclinic, *P*2<sub>1</sub>/*n*  
*a* = 12.340 (2) Å  
*b* = 8.7775 (14) Å  
*c* = 17.678 (3) Å  
 β = 104.861 (3)°  
*V* = 1850.7 (5) Å<sup>3</sup>  
*Z* = 4

*D<sub>x</sub>* = 1.241 Mg m<sup>-3</sup>  
 Mo Kα radiation  
 Cell parameters from 2447 reflections  
 θ = 2.6–22.7°  
 μ = 0.22 mm<sup>-1</sup>  
*T* = 294 (2) K  
 Prism, colorless  
 0.26 × 0.22 × 0.20 mm

Data collection

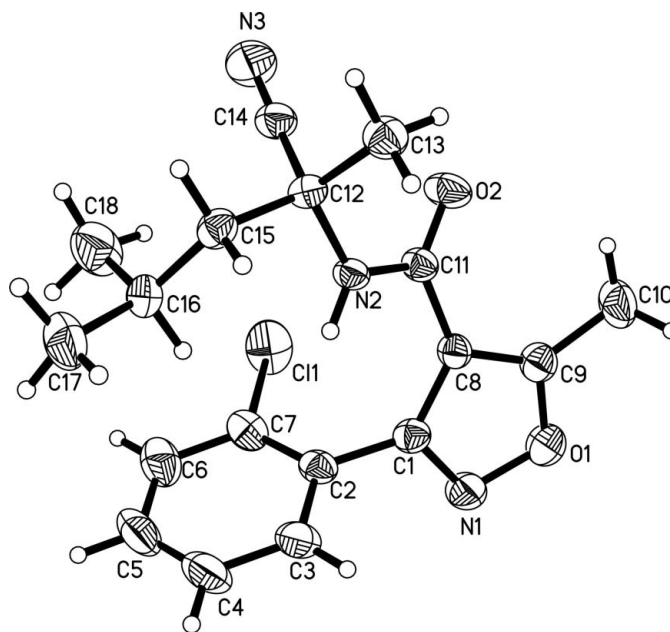
Bruker SMART CCD area-detector diffractometer  
 φ and ω scans  
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  
*T<sub>min</sub>* = 0.943, *T<sub>max</sub>* = 0.957  
 10142 measured reflections

3784 independent reflections  
 2123 reflections with *I* > 2σ(*I*)  
*R<sub>int</sub>* = 0.032  
 θ<sub>max</sub> = 26.5°  
*h* = -15 → 15  
*k* = -8 → 10  
*l* = -19 → 22

Refinement

Refinement on *F*<sup>2</sup>  
*R* [*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.045  
*wR* (*F*<sup>2</sup>) = 0.150  
*S* = 0.94  
 3784 reflections  
 221 parameters  
 H-atom parameters constrained

*w* = 1/[σ<sup>2</sup>(*F<sub>o</sub>*<sup>2</sup>) + (0.07*P*)<sup>2</sup> + 0.7394*P*]  
 where *P* = (*F<sub>o</sub>*<sup>2</sup> + 2*F<sub>c</sub>*<sup>2</sup>)/3  
 (Δ/σ)<sub>max</sub> = 0.014  
 Δρ<sub>max</sub> = 0.25 e Å<sup>-3</sup>  
 Δρ<sub>min</sub> = -0.20 e Å<sup>-3</sup>



**Figure 1**  
 The structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at 40% probability level.

**Table 1**

Selected geometric parameters (Å, °).

O1–C9	1.344 (3)	N1–C1	1.311 (3)
O1–N1	1.407 (3)	N2–C11	1.348 (3)
O2–C11	1.220 (2)	N2–C12	1.463 (3)
O2–C11–C8	122.2 (2)	N2–C11–C8	114.98 (18)
C8–C1–C2–C7	67.4 (4)	C1–C8–C11–N2	29.4 (4)
N1–C1–C2–C3	67.2 (3)	C11–N2–C12–C15	174.5 (2)
C12–N2–C11–O2	-4.8 (4)	C12–C15–C16–C17	170.8 (2)
C9–C8–C11–O2	29.9 (4)		

**Table 2**

Hydrogen-bond geometry (Å, °).

<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–H2...N1 <sup>i</sup>	0.86	2.63	3.187 (3)	124
C10–H10B...O2	0.96	2.53	3.156 (4)	123
C10–H10C...O2 <sup>ii</sup>	0.96	2.59	3.407 (4)	143
C13–H13C...O2	0.96	2.54	3.110 (3)	118
C16–H16A...N2	0.98	2.60	2.963 (3)	102

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

The H atoms were placed in calculated positions and allowed to ride on their parent atoms, with an N–H distance of 0.86 Å and C–

H distances in the range 0.93–0.98 Å, and with *U*<sub>iso</sub>(H) = 1.2*U*<sub>eq</sub>(C/N) and 1.5*U*<sub>eq</sub>(methyl C).

Data collection: SMART (Bruker, 1998); cell refinement: SAINT (Bruker, 1999); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 1999); software used to prepare material for publication: SHELXTL.

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