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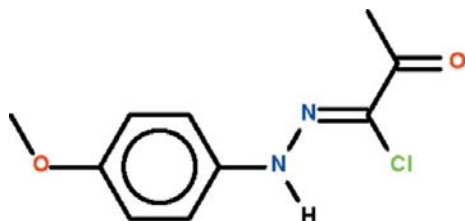
## 1-Chloro-1-[(4-methoxyphenyl)-hydrazinylidene]propan-2-one

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Key indicators: single-crystal X-ray study;  $T = 100$  K; mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å;  $R$  factor = 0.038;  $wR$  factor = 0.102; data-to-parameter ratio = 14.7.The non-H atoms of the title compound,  $\text{C}_{10}\text{H}_{11}\text{ClN}_2\text{O}_2$ , lie nearly on a plane (r.m.s. deviation = 0.150 Å), and the  $\text{C}=\text{N}$  double bond has a  $Z$  configuration. In the crystal, adjacent molecules are linked by an  $\text{N}-\text{H}\cdots\text{O}_{\text{carbonyl}}$  hydrogen bond, forming a chain running along [201].

## Related literature

For the synthesis, see: Benincori *et al.* (1990); Sayed *et al.* (2002). For background to the title compound, see: Asiri *et al.* (2010).

## Experimental

## Crystal data

 $\text{C}_{10}\text{H}_{11}\text{ClN}_2\text{O}_2$   
 $M_r = 226.66$ Monoclinic,  $P2_1/c$   
 $a = 5.8873$  (3) Å $b = 25.0467$  (10) Å  
 $c = 7.3041$  (3) Å  
 $\beta = 99.016$  (4)°  
 $V = 1063.74$  (8) Å<sup>3</sup>  
 $Z = 4$ Cu  $K\alpha$  radiation  
 $\mu = 3.05$  mm<sup>-1</sup>  
 $T = 100$  K  
 $0.35 \times 0.10 \times 0.05$  mm

## Data collection

Agilent SuperNova Dual  
diffractometer with an Atlas  
detector  
Absorption correction: multi-scan  
(*CrysAlis PRO*; Agilent, 2010)  
 $T_{\text{min}} = 0.415$ ,  $T_{\text{max}} = 0.863$ 3802 measured reflections  
2090 independent reflections  
1776 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.026$ 

## Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.038$   
 $wR(F^2) = 0.102$   
 $S = 1.05$   
2090 reflections  
142 parametersH atoms treated by a mixture of  
independent and constrained  
refinement  
 $\Delta\rho_{\text{max}} = 0.25$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.30$  e Å<sup>-3</sup>

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N2}-\text{H2}\cdots\text{O1}^i$	0.87 (3)	2.22 (3)	3.021 (2)	153 (2)

Symmetry code: (i)  $x + 1, -y + \frac{1}{2}, z + \frac{1}{2}$ Data collection: *CrysAlis PRO* (Agilent, 2010); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *publCIF* (Westrip, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: XU5259).

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