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## Structure Reports

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## 4-(Chloromethyl)benzophenone

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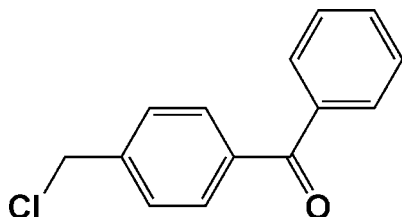
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Key indicators: single-crystal X-ray study;  $T = 293$  K; mean  $\sigma(\text{C}-\text{C}) = 0.002$  Å;  $R$  factor = 0.040;  $wR$  factor = 0.103; data-to-parameter ratio = 18.5.

In the title compound {systematic name: [4-(chloromethyl)phenyl](phenyl)methanone},  $\text{C}_{14}\text{H}_{11}\text{ClO}$ , the two benzene rings form a dihedral angle of  $58.10(8)^\circ$ . The crystal packing is stabilized by intermolecular  $\text{C}-\text{H}\cdots\pi$  interactions which link the molecules into a ribbon along the  $a$  axis.

## Related literature

For related literature, see: Green & Green (1989); James & Ridgefield (1980); Mejiritski *et al.* (1997); Xu *et al.* (2007).



## Experimental

## Crystal data

$\text{C}_{14}\text{H}_{11}\text{ClO}$   
 $M_r = 230.68$   
 Monoclinic,  $P2_1/n$   
 $a = 7.5227(5)$  Å

$b = 11.4535(7)$  Å  
 $c = 13.5932(7)$  Å  
 $\beta = 102.694(4)^\circ$   
 $V = 1142.58(12)$  Å<sup>3</sup>

$Z = 4$   
 Mo  $K\alpha$  radiation  
 $\mu = 0.31$  mm<sup>-1</sup>

$T = 293(2)$  K  
 $0.50 \times 0.23 \times 0.15$  mm

## Data collection

Bruker SMART CCD area-detector diffractometer  
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  
 $T_{\min} = 0.861$ ,  $T_{\max} = 0.955$

8204 measured reflections  
 2678 independent reflections  
 1936 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.024$

## Refinement

$R[F^2 > 2\sigma(F^2)] = 0.040$   
 $wR(F^2) = 0.103$   
 $S = 1.04$   
 2678 reflections

145 parameters  
 H-atom parameters constrained  
 $\Delta\rho_{\max} = 0.17$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.29$  e Å<sup>-3</sup>

Table 1

Hydrogen-bond geometry (Å, °).

Cg1 and Cg2 are the centroids of the C2–C7 and C9–C14 rings, respectively.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{C4}-\text{H4}\cdots\text{Cg2}^{\text{i}}$	0.93	2.85	3.6347 (17)	143
$\text{C10}-\text{H10}\cdots\text{Cg1}^{\text{ii}}$	0.93	2.94	3.6676 (17)	137

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

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.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CI2397).

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