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## Cyclohexanone at 150 K

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#### **Key indicators**

Single-crystal X-ray study T = 150 KMean  $\sigma(\text{C-C}) = 0.003 \text{ Å}$ R factor = 0.046wR factor = 0.119Data-to-parameter ratio = 12.1

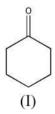
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

The structure of cyclohexanone,  $C_6H_{10}O$ , at 150 K is that of discrete molecules, with no strong intermolecular interactions.

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## Comment

Many of the esters and ketones used in the flavours and fragrances industry are liquid at room temperature, meaning that, in the past, crystalline derivatives have had to be prepared for X-ray analysis. As part of a programme to systematize *in situ* crystal growth from liquids, we have examined a range of commercially available chemicals. Low-molecular weight organic ketones are liquid at room temperature. The molecules of cyclohexanone, (I), exist in the crystal structure at 150 K as discrete entities, with no strong intermolecular interactions.



## **Experimental**

A 3 mm column of the title material, which is a liquid at room temperature, was sealed in a 0.3 mm Lindemann tube. The Lindemann tube was not precisely parallel to the  $\varphi$  axis. A single crystal of the compound was grown by keeping the compound under a cold nitrogen gas stream (Oxford Cryostream 600) at 180 K and slowly moving a small liquid zone, created by a micro-heating coil, up and down the sample. Once a suitable approximately single-crystal specimen had been obtained, the main data collection was carried out at 150 K. Because not all of the data were collected with the Lindemann tube perpendicular to the X-ray beam, the multi-scan corrections applied by DENZO/SCALEPACK (Otwinowski & Minor, 1997) also contain contributions due to changes in the illuminated volume of the cylindrical sample, which affects the value of  $T_{min}/T_{max}$ .

Crystal data

 $C_6H_{10}O$   $M_r = 98.14$ Orthorhombic,  $P2_12_12_1$  a = 5.3736 (2) Å b = 7.0394 (3) Å c = 15.1910 (7) Å V = 574.63 (4) Å<sup>3</sup> Z = 4 $D_x = 1.134$  Mg m<sup>-3</sup> Mo  $K\alpha$  radiation Cell parameters from 784 reflections  $\theta = 5{\text -}27^{\circ}$  $\mu = 0.08~\text{mm}^{-1}$ T = 150~KCylinder, colourless  $0.70 \times 0.30 \times 0.30~\text{mm}$ 

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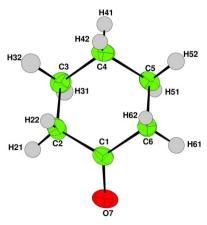


Figure 1
The title compound, with displacement ellipsoids drawn at the 50% probability level. H atoms are of arbitrary radii.

## Data collection

Nonius KappaCCD diffractometer  $\omega$  scans 593 reflections with  $I > 2\sigma(I)$  Absorption correction: multi-scan  $(DENZO/SCALEPACK; \ Otwinowski & Minor, 1997) h = -6 <math>\rightarrow 6$   $T_{min} = 0.74, T_{max} = 0.98$   $\mu = -9 \rightarrow 9$  9235 measured reflections  $\mu = 1.9 \rightarrow 19$ 

### Refinement

 $\begin{array}{lll} \mbox{Refinement on } F^2 & \mbox{H-atom parameters constrained} \\ R[F^2 > 2\sigma(F^2)] = 0.046 & \mbox{$w = 1/[\sigma^2(F) + 0.08 + 0.07P]$,} \\ wR(F^2) = 0.119 & \mbox{where } P = (\mbox{max}(F_o^2, 0) + 2F_c^2)/3 \\ S = 1.02 & (\Delta/\sigma)_{\mbox{max}} = 0.009 \\ 774 \mbox{ reflections} & \Delta\rho_{\mbox{max}} = 0.21 \mbox{ e Å}^{-3} \\ 64 \mbox{ parameters} & \Delta\rho_{\mbox{min}} = -0.17 \mbox{ e Å}^{-3} \\ \end{array}$ 

**Table 1**Selected geometric parameters (Å, °).

C1-C2	1.501 (2)	C3-C4	1.520 (3)
C1-C6	1.513 (2)	C4-C5	1.523 (3)
C1-O7	1.213 (2)	C5-C6	1.533 (2)
C2-C3	1.532 (3)		
C2-C1-C6	115.45 (14)	C2-C3-C4	111.63 (15)
C2-C1-O7	122.61 (15)	C3-C4-C5	110.85 (16)
C6 - C1 - O7	121.93 (15)	C4 - C5 - C6	111.04 (15)
C1 - C2 - C3	112.29 (15)	C5 - C6 - C1	111.65 (13)

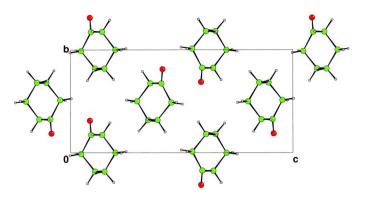


Figure 2 The crystal structure, viewed down the a axis.

All H atoms were located in a difference map and were repositioned geometrically. The H atoms were initially refined with soft restraints on the bond lengths and angles to regularize their geometry [C—H = 0.97–1.01 Å, and  $U_{\rm iso}({\rm H})$  = 1.2 $U_{\rm eq}({\rm C})$ ], after which they were refined with riding constraints. In the absence of significant anomalous scattering effects, Friedel pairs were merged.

Data collection: *COLLECT* (Nonius, 1997); cell refinement: *DENZO/SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *DENZO/SCALEPACK*; program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *CRYSTALS* (Betteridge *et al.*, 2003); molecular graphics: *CAMERON* (Watkin *et al.*, 1996); software used to prepare material for publication: *CRYSTALS*.

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