

1-Hydroxydiamantane

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Key indicators

Single-crystal X-ray study
 $T = 293\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$
 R factor = 0.049
 wR factor = 0.149
Data-to-parameter ratio = 17.9For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.The molecule of the title compound, $\text{C}_{14}\text{H}_{20}\text{O}$, possesses normal geometrical parameters. The compound crystallizes in the uncommon space group $P4_2/n$ and features discrete quartets of molecules linked by $\text{O}-\text{H}\cdots\text{O}$ bonds into a loop.

Comment

The title compound, (I) (Fig. 1), is a derivative of diamantane. Diamantane is a member of the diamondoid family: a hydrocarbon that has a carbon framework that is superimposable on the diamond lattice. Diamantane derivatives are potentially valuable molecules both in materials science and in pharmaceutical sciences. For example, (I) is an intermediate for the synthesis of a variety of other diamantane derivatives such as 1-bromodiamantane, 1-aminodiamantane and 1-diamantanecarboxylic acid.

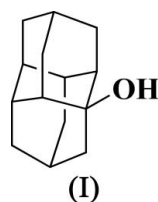
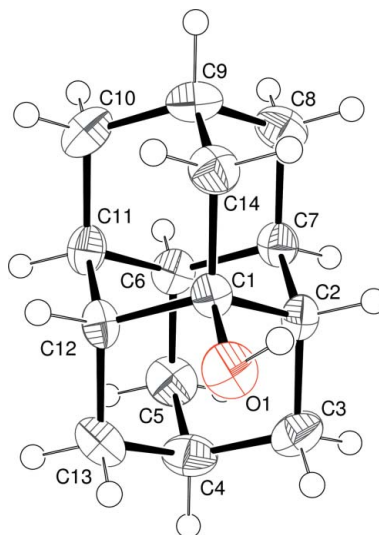
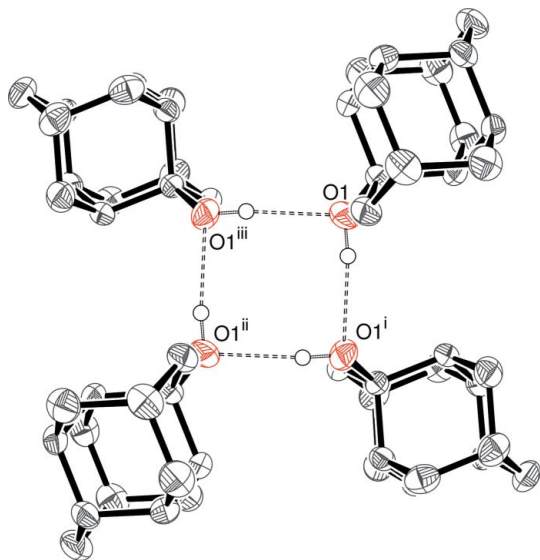
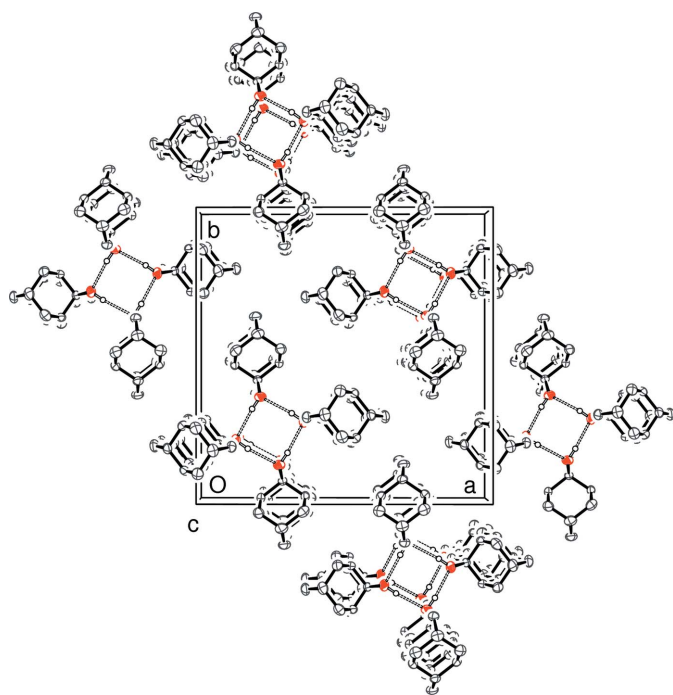
The geometrical parameters for (I) are normal. The crystal structure is stabilized by intermolecular $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds (Table 1), which result in the formation of discrete quartets (Figs. 2 and 3) of molecules.

Figure 1
View of (I), showing 50% displacement ellipsoids (arbitrary spheres for H atoms).


Figure 2

View of a quartet of molecules linked by hydrogen bonds (50% displacement ellipsoids) C-bound H atoms have been omitted for clarity and hydrogen bonds are shown as dashed lines. [Symmetry codes: (i) $\frac{1}{2} - y, x, \frac{1}{2} - z$; (ii) $\frac{1}{2} - x, \frac{1}{2} - y, z$; (iii) $y, \frac{1}{2} - x, \frac{1}{2} - z$.]


Figure 3

: The packing of (I). Drawing conventions as in Fig. 2.

Experimental

The title compound was prepared according to the procedure reported by Fokin *et al.* (2005) in 69.4% yield. Suitable crystals were recrystallized from acetone (m.p. 564–565 K). ^{13}C NMR (CDCl_3 ,

75 MHz): δ 70.72 (1C, C-1), 46.33 (1C, C-14), 43.31 (2C, C-2, C-12), 39.88 (2C, C-7, C-11), 37.92 (1C, C-5), 37.46 (2C, C-8, C-10), 36.66 (1C, C-6), 32.47 (2C, C-3, C-13), 30.37 (1C, C-9), 25.23 (1C, C-4). Analysis calculated for $\text{C}_{14}\text{H}_{20}\text{O}$: C 82.30, H 9.87%; found: C 82.34, H 9.80%.

Crystal data

$\text{C}_{14}\text{H}_{20}\text{O}$
 $M_r = 204.30$
 Tetragonal, $P4_2/n$
 $a = 16.572$ (2) Å
 $c = 7.7879$ (16) Å
 $V = 2138.8$ (6) Å³
 $Z = 8$

$D_x = 1.269$ Mg m⁻³
 Mo $K\alpha$ radiation
 $\mu = 0.08$ mm⁻¹
 $T = 293$ (2) K
 Prism, colorless
 $0.64 \times 0.22 \times 0.18$ mm

Data collection

Rigaku R-AXIS RAPID IP
 diffractometer
 ω scans
 Absorption correction: multi-scan
 (ABSCOR; Higashi, 1995)
 $T_{\min} = 0.952$, $T_{\max} = 0.986$

18440 measured reflections
 2441 independent reflections
 1899 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.038$
 $\theta_{\max} = 27.5^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.049$
 $wR(F^2) = 0.149$
 $S = 1.05$
 2441 reflections
 136 parameters

H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.1P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.33$ e Å⁻³
 $\Delta\rho_{\min} = -0.26$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O1}-\text{H1A}\cdots\text{O1}^i$	0.82	1.96	2.7581 (13)	166

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

All H atoms were initially located in a difference Fourier map. There were repositioned with ideal geometry ($\text{C}-\text{H} = 0.97\text{--}0.98$ Å and $\text{O}-\text{H} = 0.82$ Å) and refined as riding, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{carrier})$.

Data collection: *RAPID-AUTO* (Rigaku, 2001); cell refinement: *RAPID-AUTO*; data reduction: *RAPID-AUTO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *SHELXL97*.

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