

Reema K. Thalji,† Mark L. McLaughlin,§ Steven F. Watkins and Frank R. Fronczek*

Department of Chemistry, Louisiana State University, Baton Rouge, LA 70803-1804, USA

† Current address: GlaxoSmithKline, 709 Swedeland Road, UW2810 PO Box 1539, King of Prussia, PA 19406-0939, USA

§ Current address: Department of Chemistry, University of South Florida, 4202 E. Fowler Ave., CHE 205A, Tampa, FL 33620-5250, USA, and Drug Discovery Program, H Lee Moffitt Cancer Center & Research Institute, 12902 Magnolia Drive, MRC 4East, Tampa, FL 33612, USA

Correspondence e-mail: ffroncz@lsu.edu

Key indicators

Single-crystal X-ray study

$T = 190\text{ K}$

Mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$

R factor = 0.056

wR factor = 0.150

Data-to-parameter ratio = 21.4

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

Di-*tert*-butyl 4-oxocyclohexane-1,1-dicarboxylate

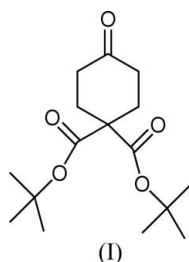
In the title compound, $\text{C}_{16}\text{H}_{26}\text{O}_5$, the cyclohexanone ring has a flattened chair conformation, with absolute values of the endocyclic torsion angles lying in the range $39.7(3)$ – $58.0(2)^\circ$.

Received 26 May 2006

Accepted 26 May 2006

Comment

The title compound, (I), was synthesized as a precursor in the synthesis of an α,α -disubstituted amino acid having the potential to stabilize the 3_{10} -helical secondary structure of a peptide in aqueous media. The crystal structure was determined in order to confirm its identity. Cooling the sample for data collection was limited to 190 K, as crystals were destroyed at lower temperatures.



The six-membered ring exists in a flattened chair conformation, as indicated by the torsion angles in Table 1. This flattening, which results from the presence of the ketone group, is similar to that seen in 4,4-dimethylcyclohexanone [Cambridge Structural Database (CSD; Version 5.27; Allen, 2002) refcode DMCYHX (Lichanot *et al.*, 1977)] and 4,4-

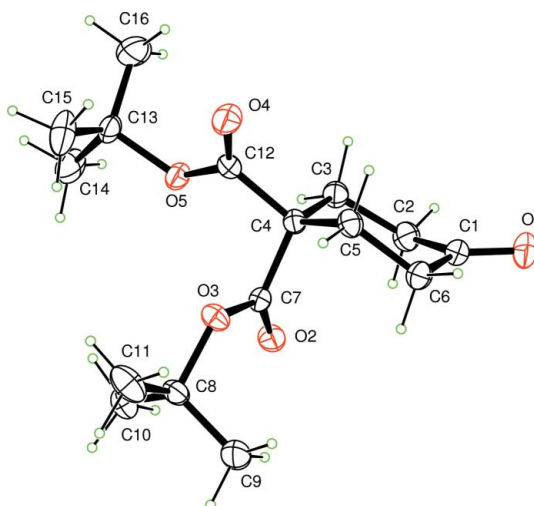


Figure 1
 View of (I), showing displacement ellipsoids drawn at the 30% probability level. H atoms are represented by spheres of arbitrary radius.

diphenylcyclohexanone (CSD refcode DPHCX10; Lambert *et al.*, 1969). The cyclohexanone ring in (I), however, is more distorted from local mirror symmetry than in either of those compounds. The mean difference of three torsion angle magnitudes across the local mirror is 7.9° in (I), 4.6° in DPHCX10, and zero for DMCYHX, which lies on a crystallographic mirror. The *tert*-butoxycarbonyl groups in (I) have conformations such that their carbonyl groups are approximately eclipsed with the C3–C4 and C4–C7 bonds (Table 1).

Experimental

Compound (I) was prepared by the method of Sanchez *et al.* (1985) from 2-methoxycarbonyl-4,4-di-*tert*-butoxycarbonylcyclohexanone. Crystals were grown by slow evaporation of a dimethyl sulfoxide solution.

Crystal data

$C_{16}H_{26}O_5$	$Z = 4$
$M_r = 298.37$	$D_x = 1.136 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
$a = 6.3412(3) \text{ \AA}$	$\mu = 0.08 \text{ mm}^{-1}$
$b = 16.186(2) \text{ \AA}$	$T = 190 \text{ K}$
$c = 17.211(2) \text{ \AA}$	Lath, colorless
$\beta = 99.096(6)^\circ$	$0.43 \times 0.28 \times 0.13 \text{ mm}$
$V = 1744.3(3) \text{ \AA}^3$	

Data collection

Enraf–Nonius CAD-4 diffractometer with an Oxford Cryosystems Cryostream cooler	2173 reflections with $I > 2\sigma(I)$
ω – 2θ scans	$R_{\text{int}} = 0.038$
Absorption correction: none	$\theta_{\text{max}} = 28.0^\circ$
6463 measured reflections	3 standard reflections
4198 independent reflections	frequency: 60 min
	intensity decay: none

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0524P)^2 + 0.2515P]$
$R[F^2 > 2\sigma(F^2)] = 0.056$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.150$	$(\Delta/\sigma)_{\text{max}} < 0.001$
$S = 1.01$	$\Delta\rho_{\text{max}} = 0.17 \text{ e \AA}^{-3}$
4198 reflections	$\Delta\rho_{\text{min}} = -0.21 \text{ e \AA}^{-3}$
196 parameters	
H-atom parameters constrained	

Table 1

Selected torsion angles ($^\circ$).

C1–C2–C3–C4	–43.1 (3)	C5–C6–C1–C2	–45.9 (3)
C2–C3–C4–C5	52.4 (2)	C6–C1–C2–C3	39.7 (3)
C3–C4–C5–C6	–58.0 (2)	C3–C4–C7–O2	8.8 (3)
C4–C5–C6–C1	55.1 (2)	C5–C4–C12–O4	12.9 (3)

All H atoms were placed in idealized positions (C–H = 0.98–0.99 Å) and refined as riding, with the constraint $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ or $1.5U_{\text{eq}}(\text{methyl C})$. A torsional parameter was refined for each methyl group.

Data collection: *CAD-4 Software* (Enraf–Nonius, 1994); cell refinement: *CAD-4 Software*; data reduction: *XCAD4* (Harms & Wocadlo, 1996); program(s) used to solve structure: *SHELXTL* (Bruker, 1998); program(s) used to refine structure: *SHELXTL*; molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *SHELXTL*.

The purchase of the diffractometer was made possible by a National Science Foundation chemical instrumentation grant, which we gratefully acknowledge. Improvements to the LSU X-ray Crystallography Facility were supported by Grant No. LEQSF(1996–97)-ENH-TR-10, administered by the Louisiana Board of Regents.

References

- Allen, F. H. (2002). *Acta Cryst.* **B58**, 380–388.
- Bruker (1998). *SHELXTL*. Version 5.10, Bruker AXS Inc., Madison, Wisconsin, USA.
- Enraf–Nonius (1994). *CAD4 Operations Manual*. Enraf–Nonius, Delft, The Netherlands.
- Farrugia, L. J. (1997). *J. Appl. Cryst.* **30**, 565.
- Harms, K. & Wocadlo, S. (1996). *XCAD4*. University of Marburg, Germany.
- Lambert, J. B., Carhart, R. E. & Corfield, P. W. R. (1969). *J. Am. Chem. Soc.* **91**, 3567–3571.
- Lichanot, A., Lectard, A., Metras, F., Gaultier, J. & Hauw, C. (1977). *Cryst. Struct. Commun.* **6**, 127–129.
- Sanchez, I., Ortega, A., Garcia, G., Larraza, M. & Flores, H. (1985). *Synth. Commun.* **15**, 141–149.