# organic papers

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# Andrei S. Batsanov,\* Alistair J. Reid and Neil Cameron

Department of Chemistry, University of Durham, South Road, Durham DH1 3LE, England

Correspondence e-mail: a.s.batsanov@durham.ac.uk

#### **Key indicators**

Single-crystal X-ray study T = 100 KMean  $\sigma$ (C–C) = 0.001 Å R factor = 0.039 wR factor = 0.114 Data-to-parameter ratio = 28.8

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

# 2-*tert*-Butoxy-1-phenyl-1-(2,2,6,6-tetramethyl-piperidin-1-yloxy)ethane

The title compound,  $C_{21}H_{35}NO_2$ , contains a piperidine ring in a chair conformation, with a pyramidal N atom and a single (exocyclic) N-O bond in an equatorial orientation.

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# Comment

Nitroxide-mediated polymerization (NMP) has emerged in recent years as a successful controlled or 'living' radical polymerization technique which can be used to prepare polymers of target molecular weight, narrow polydispersity and complex architecture (block and graft copolymers, star polymers, etc.) (Matyjaszewski, 2003). Successful NMP requires the use of a monomolecular initiator, identified as an alkoxyamine, which is derived from a nitroxide. This alkoxyamine should be prepared separately and then added in a known concentration to the monomer to be polymerized. Synthetic routes to alkoxyamines include the trapping of alkyl radicals by free nitroxides at moderate temperatures (Braslau et al., 1997; Miura et al., 1998) and a catalytic route involving Mn-salen complexes [H2salen is bis(salicylidene)ethylenediamine; Dao et al., 1998]. The title compound, (I), has been synthesized in the course of these studies (Cameron et al., 2000).



The molecular structure of (I) (Fig. 1) is similar in its main features to the other *N*-oxy-2,2,6,6-tetramethylpiperidinyl derivatives reported by Ermert & Vasella (1993), Jahn *et al.* (2001, 2002) and Leitich *et al.* (2002). The piperidine ring adopts a chair conformation with the N–O1 bond in an equatorial orientation. The N–O1 distance is typical for a single bond (Allen *et al.*, 1987), while the N atom has pyramidal geometry, corresponding to  $sp^3$  hybridization.

# **Experimental**

A solution of 2,2,6,6-tetramethylpiperidinyloxy (TEMPO; 0.187 g, 1.20 mmol) and di-*tert*-butylperoxalate (0.148 g, 0.63 mmol) in styrene (5 ml) was prepared. The mixture was degassed by three freeze/pump/thaw cycles and backflushed with argon after the final cycle. The resulting solution was sealed, then heated overnight at 313 K. After cooling, the solution was dried *in vacuo* to constant

© 2005 International Union of Crystallography Printed in Great Britain – all rights reserved mass. The crude solid product obtained was then purified by flash chromatography on silica gel, eluting with a mixture of 40–60 petroleum ether/ethyl acetate (90:10  $\nu/\nu$ ). After removal of the solvent, a white crystalline solid was obtained in 79% yield. The characterization data of the isolated product were in agreement with those given in the literature (Bon *et al.*, 1999). <sup>1</sup>H NMR (300 MHz):  $\delta$  1.04 (*s*, 9H, CH<sub>3</sub> × 3), 0.58, 1.02, 1.19, 1.37 (*br s*, 3H, CH<sub>3</sub>), 0.9–1.7 (*br m*, 6H, CH<sub>2</sub> × 3), 3.41 (*m*, 1H, H1), 3.89 (*m*, 1H, H171), 4.74 (*m*, 1H, H172), 7.18–7.35 (*m*, 5H, Ph) p.p.m. (for the H-atom numbering, see Fig. 1).

Mo *Kα* radiation Cell parameters from 902 reflections

7229 independent reflections

 $w = 1/[\sigma^2(F_o^2) + (0.0633P)^2]$ 

where  $P = (F_o^2 + 2F_c^2)/3$ 

+ 0.7171P]

 $\Delta \rho_{\rm max} = 0.52 \text{ e} \text{ Å}^{-3}$ 

 $\Delta \rho_{\rm min} = -0.17 \text{ e} \text{ Å}^{-3}$ 

 $(\Delta/\sigma)_{\rm max} = 0.001$ 

6298 reflections with  $I > 2\sigma(I)$ 

 $\theta = 12.1-24.3^{\circ}$   $\mu = 0.07 \text{ mm}^{-1}$  T = 100 (2) KBlock, colourless  $1.00 \times 0.66 \times 0.14 \text{ mm}$ 

 $R_{\rm int} = 0.020$  $\theta_{\rm max} = 32.5^{\circ}$ 

 $h = -23 \rightarrow 23$ 

 $\begin{array}{l} k=-16 \rightarrow 16 \\ l=-34 \rightarrow 35 \end{array}$ 

#### Crystal data

$C_{21}H_{35}NO_2$
$M_r = 333.50$
Orthorhombic, Pbca
a = 15.532 (4)  Å
<i>b</i> = 11.119 (3) Å
c = 23.247 (5)  Å
$V = 4014.8 (17) \text{ Å}^3$
Z = 8
$D_{\rm r} = 1.104 {\rm Mg m}^{-3}$

### Data collection

Bruker SMART 6000 CCD area-
detector diffractometer
$\omega$ scans
Absorption correction: multi-scan
(SADABS; Bruker, 2001)
$T_{\min} = 0.579, T_{\max} = 1.000$
60196 measured reflections

#### Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.039$   $wR(F^2) = 0.114$  S = 1.067229 reflections 251 parameters H atoms treated by a mixture of independent and constrained refinement

#### Table 1

Selected	geometric	parameters	(A,	°).
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O1-C1	1.4422 (8)	C2-C3	1.5399 (11)
O1-N	1.4595 (7)	C3-C4	1.5198 (14)
O2-C17	1.4158 (8)	C4-C5	1.5153 (13)
O2-C18	1.4446 (8)	C5-C6	1.5312 (10)
N-C6	1.4987 (9)	C6-C10	1.5280 (10)
N-C2	1.5033 (9)	C6-C9	1.5353 (11)
C1-C11	1.5168 (9)	C18-C20	1.5187 (10)
C1-C17	1.5228 (9)	C18-C21	1.5200 (11)
C2-C8	1.5354 (11)	C18-C19	1.5238 (11)
C2-C7	1.5377 (12)		
C1-O1-N	112.33 (5)	C5-C4-C3	108.36 (7)
C17-O2-C18	116.92 (5)	C4-C5-C6	112.78 (6)
O1-N-C6	106.17 (5)	C10-C6-C5	107.62 (6)
O1-N-C2	107.19 (5)	O2-C17-C1	108.48 (5)
C6-N-C2	117.35 (5)	O2-C18-C20	103.38 (6)
O1-C1-C11	114.45 (5)	O2-C18-C21	111.07 (6)
O1-C1-C17	105.37 (5)	C20-C18-C21	110.35 (6)
C11-C1-C17	110.04 (5)	O2-C18-C19	111.02 (6)
N-C2-C3	107.44 (6)	C20-C18-C19	110.12 (7)
C4-C3-C2	114.33 (7)	C21-C18-C19	110.68 (7)



#### Figure 1

The molecular structure of (I), showing atomic displacement ellipsoids at the 50% probability level.

Methyl groups were treated as threefold symmetrical bodies rotating around C–C bonds, with a refined common  $U_{iso}$  for the three H atoms. Other H atoms were treated as riding on the corresponding C atoms, with refined  $U_{iso}$  values. C–H distances are 0.95–1.00 Å.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SMART*; data reduction: *SAINT* (Bruker, 2001); program(s) used to solve structure: *SHELXTL* (Bruker, 2001); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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