

Keywords: crystal structure; Ritter reaction; (\pm)-*exo*-*N*-isobornylacetamides; polymorph; hydrogen bonding

CCDC references: 1420761; 1420760

Supporting information: this article has supporting information at journals.iucr.org/e

Crystal structures of two (\pm)-*exo*-*N*-isobornylacetamides

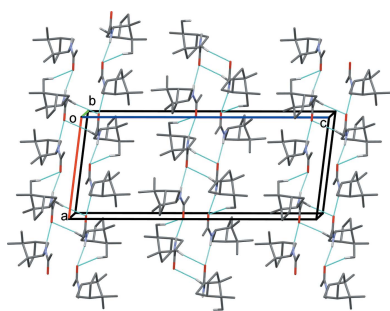
Dmitrijs Stepanovs,^{a,b*} Daniels Posevins^b and Maris Turks^{b*}

^aLatvian Institute of Organic Synthesis, Aizkraukles 21, Riga, LV-1006, Latvia, and ^bInstitute of Technology of Organic Chemistry, Faculty of Materials Science and Applied Chemistry, Riga Technical University, P. Valdena 3/7, Riga, LV-1048, Latvia. *Correspondence e-mail: d_stepanovs@osi.lv, maris_turks@ktf.rtu.lv

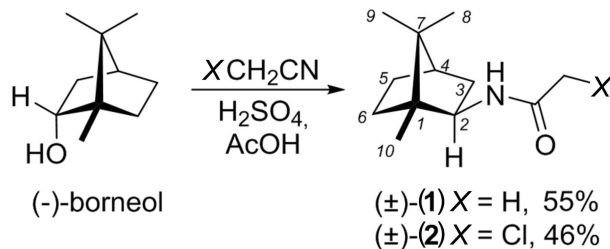
The title compounds consist of a bornane skeleton with attached acetamide, C₁₂H₂₁NO (\pm)-**(1)** {systematic name: (\pm)-*N*-[(1*RS*,2*RS*,4*RS*)-1,7,7-trimethylbicyclo[2.2.1]heptan-2-yl]acetamide}, and chloroacetamide, C₁₂H₂₀ClNO (\pm)-**(2)** {systematic name: (\pm)-2-chloro-*N*-[(1*RS*,2*RS*,4*RS*)-1,7,7-trimethylbicyclo[2.2.1]heptan-2-yl]acetamide}, functionalities to the 2-*exo*-position. The crystal structure of the first monoclinic polymorph of (\pm)-**(1)** has been reported previously [Ung *et al.* (2014). *Monatsh. Chem.* **145**, 983–992]. Compound (\pm)-**(1)** crystallizes in the space group *P*2₁/*n* with two independent molecules in the asymmetric unit, in contrast to the above-mentioned polymorph which crystallized in the space group *C*2/*c* with one molecule in the asymmetric unit. In the title compounds, the bicyclic bornane moieties have normal geometries. In the crystals of both compounds, molecules are linked by N—H \cdots O hydrogen bonds, reinforced by C—H \cdots O contacts, forming *trans*-amide chains propagating along the *a*-axis direction. In the case of compound (\pm)-**(1)**, neighbouring chains are linked by further C—H \cdots O contacts, forming double-chain ribbons along [100].

1. Chemical context

Isobornylamine-derived amides have recently been described as useful antimycobacterial agents (Stavrov *et al.*, 2014*a,b*). Promising biological activity profiles have been also discovered for other bornane derivatives such as 2-arylbornanes (Duclos *et al.*, 2008), camphor oximes (Schenone *et al.*, 2000), bornyl (3,4,5-trihydroxy)-cinnamate (Steinbrecher *et al.*, 2008) and others. There is no doubt that isobornylamine derivatives are chemically related to terpenoids camphor (Seebaluck *et al.*, 2015) and borneol (Horváthová *et al.*, 2012), which are well known for their biological activities. On the other hand, compounds containing the bornane skeleton are frequently used as chiral building blocks for various ligands, catalysts and chiral auxiliaries (Chelucci, 2006; Langlois & Kouklovsky, 2009; Ramón & Yus, 2007). In light of the aforementioned facts, there is a vast interest in developing new synthetic protocols for the synthesis of compounds of this class and in their structural studies. We have recently reported an application of the Ritter reaction (Jiang *et al.*, 2014) in the synthesis of amide-derivatized heterocycles (Turks *et al.*, 2012). Hence, we identified the possibility to obtain isobornylamine derived amides (\pm)-**(1)** and (\pm)-**(2)** from borneol in the direct Ritter reaction. When the optically active (–)-borneol was submitted to standard Ritter reaction conditions, the expected compounds were isolated in acceptable yields albeit in the racemic form. A similar type of racemization due to a 6,2-hydride shift was described in the Ritter reaction of (–)-bornyl acetate (Hanzawa *et al.*, 2012.). Previously, compounds (\pm)-**(1)** and (\pm)-**(2)** have been obtained as side products in a



cationic rearrangement of (–)-β-pinene in the presence of the corresponding nitriles (Ung *et al.*, 2014).



2. Structural commentary

The title compounds consist of a 1,7,7-trimethylbicyclo[2.2.1]-heptane (bornane or camphane) skeleton with attached acetamide [(±)-(1)] and chloroacetamide [(±)-(2)] functionalities in the 2-*exo*-position. The asymmetric unit of compound (±)-(1) (Fig. 1) contains two independent molecules having coincident geometry (r.m.s. deviation 0.057 Å). Compound (±)-(2) (Fig. 2) contains one molecule in the asymmetric unit. The bond lengths and angles in both compounds are close to those observed for the first monoclinic polymorph of compound (±)-(1) (Ung *et al.*, 2014).

3. Supramolecular features

In the crystals of both compounds, molecules are linked by N–H···O hydrogen bonds, reinforced by C–H···O contacts, forming *trans*-amide chains propagating along the *a* axis direction (Figs. 3 and 4 and Tables 1 and 2). In the case of compound (±)-(1), neighbouring chains are linked by further C–H···O contacts, forming ribbons along [100]; see Fig. 3 and Table 1.

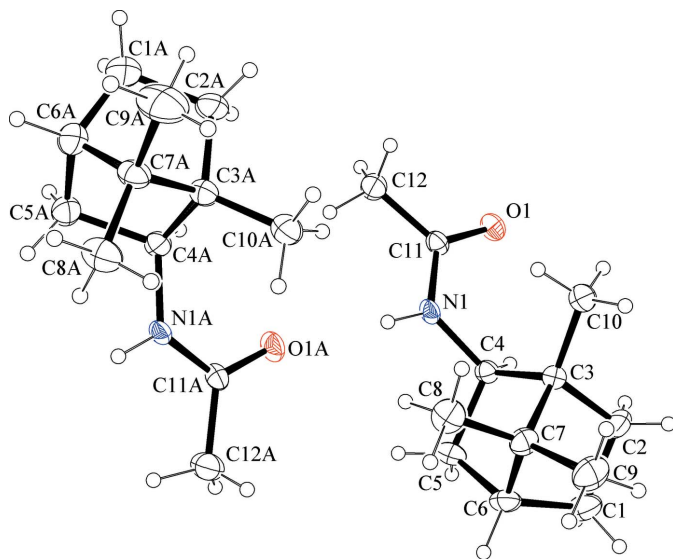


Figure 1
The molecular structure of the two independent molecules of compound (±)-(1), showing the atom labelling. Displacement ellipsoids are drawn at the 50% probability level.

Table 1
Hydrogen-bond geometry (Å, °) for (±)-(1).

<i>D</i> –H··· <i>A</i>	<i>D</i> –H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> –H··· <i>A</i>
N1–H1N···O1A ⁱ	0.85 (2)	2.06 (2)	2.900 (2)	170 (2)
N1A–H1AN···O1 ⁱⁱ	0.87 (2)	2.03 (2)	2.886 (2)	172 (2)
C8A–H8A1···O1 ⁱⁱ	0.98	2.57	3.524 (3)	165
C12–H12C···O1 ⁱⁱ	0.98	2.52	3.468 (3)	164

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

Table 2
Hydrogen-bond geometry (Å, °) for (±)-(2).

<i>D</i> –H··· <i>A</i>	<i>D</i> –H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> –H··· <i>A</i>
N1–H1···O1 ⁱ	0.79 (3)	2.21 (3)	2.983 (2)	168 (2)
C12–H12A···O1 ⁱ	0.97	2.36	3.238 (3)	151

Symmetry code: (i) $x + \frac{1}{2}, -y + \frac{3}{2}, z$.

4. Database survey

A search of the Cambridge Structural Database (Version 5.36; Groom & Allen, 2014) for substituted bornanes gave 1517 hits (excluding organometallics). 119 structures are substituted at the 2-position. Only two of these are amides, *viz.* the previously reported polymorph of (±)-(1) (LOPQEO: Ung *et al.*, 2014) and 2,2,2-triphenyl-*N*-(1,7,7-trimethylbicyclo[2.2.1]-hept-2-yl)acetamide (TOQWED: Prusinowska *et al.*, 2015).

5. Synthesis and crystallization

Compound (±)-(1): (–)-Borneol (463 mg, 3 mmol, 1 equiv.) was added to a stirred solution of acetonitrile (790 μL, 15 mmol, 5.0 equiv.) in glacial acetic acid (7.0 ml) and conc.

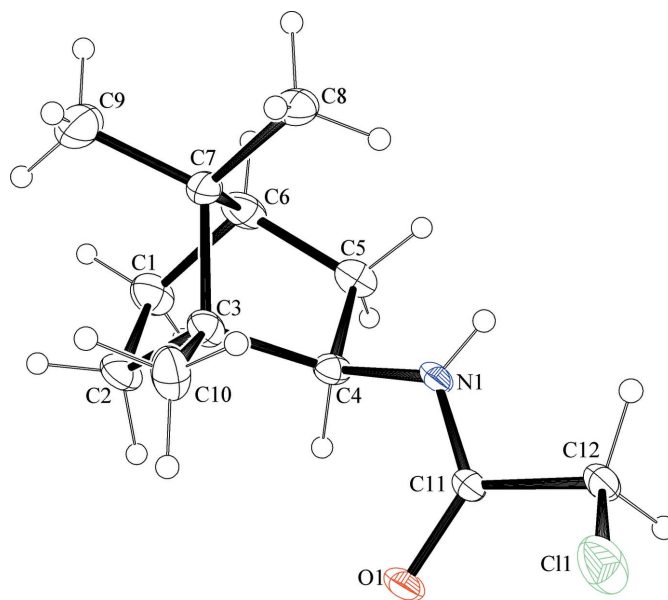


Figure 2
The molecular structure of compound (±)-(2), showing the atom labelling. Displacement ellipsoids are drawn at the 50% probability level.

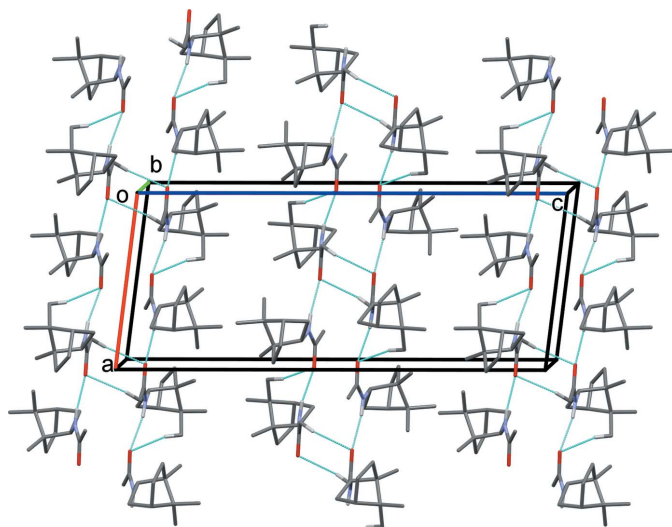


Figure 3
The crystal packing of compound (±)-(1), viewed along the *b* axis. Hydrogen bonds are shown as dashed lines (see Table 1 for details). For clarity, only H atoms involved in these interactions have been included.

H₂SO₄ (3.07 g, 30 mmol, 10.0 equiv.). The resulting reaction mixture was stirred at 343 K for 16 h (TLC control). The reaction mixture was cooled to 273 K and poured into a vigorously stirred 10% aqueous solution of NaOH (30–40 mL) at 273 K. Ethyl acetate (30 mL) was added and the phases were separated. The aqueous phase was extracted with ethyl acetate (3 × 20 mL). The combined organic phase was washed with brine, dried over anhydrous Na₂SO₄, filtered and evaporated under reduced pressure. The resulting residue was purified by silica gel column chromatography to provide (±)-(1) (yield: 319 mg, 55%). The NMR data of (±)-(1) correspond fully to those reported earlier (Ung *et al.*, 2014): ¹H NMR (300 MHz, CDCl₃) δ (p.p.m.): 5.44 (*br s*, 1H), 3.87 (*td*, *J* = 9.0, 5.2 Hz, 1H), 1.97–1.77 (*m*, 4H), 1.73–1.61 (*m*, 2H), 1.60–1.46 (*m*, 2H), 1.32–1.20 (*m*, 1H), 1.18–1.07 (*m*, 1H), 0.88 (*s*, 3H), 0.81 (*s*, 3H), 0.80 (*s*, 3H); ¹³C NMR (75.5 MHz, CDCl₃) δ (p.p.m.): 169.35, 56.81, 48.87, 47.14, 44.92, 39.15, 36.02, 27.06, 23.69, 20.38, 20.35, 11.77. GC–MS (C₁₂H₂₁NO): *t*_R = 5.92 min; *m/z*: calculated 195.2; found 195.1. (GC–MS method: column: HP5 (5% phenyl methyl siloxane), 30 m × 0.25 mm ID, 0.25 μm; column temp.: 323 K (hold for 2 min) to 583 K at 323 K min⁻¹ (hold at 583 K for 3 min); injector/detector: 523 K/503 K; carrier gas: helium at 1.0 mL min⁻¹, linear velocity; injection mode: splitless (solvent delay: 3 min); injection volume: 1 μL). X-ray quality single crystals were obtained by slow evaporation of a solution of (±)-(1) in hexanes/ethyl acetate (2:1).

Compound (±)-(2): (–)-Borneol (463 mg, 3 mmol, 1 equiv.) was added to a stirred solution of chloroacetonitrile (950 μL, 15 mmol, 5.0 equiv.) in glacial acetic acid (7.0 ml) and conc. H₂SO₄ (3.07 g, 30 mmol, 10.0 equiv.). The resulting reaction mixture was stirred at 343 K for 16 h (TLC control). The reaction mixture was cooled to 273 K and poured into a vigorously stirred 10% aqueous solution of NaOH (30–40 mL)

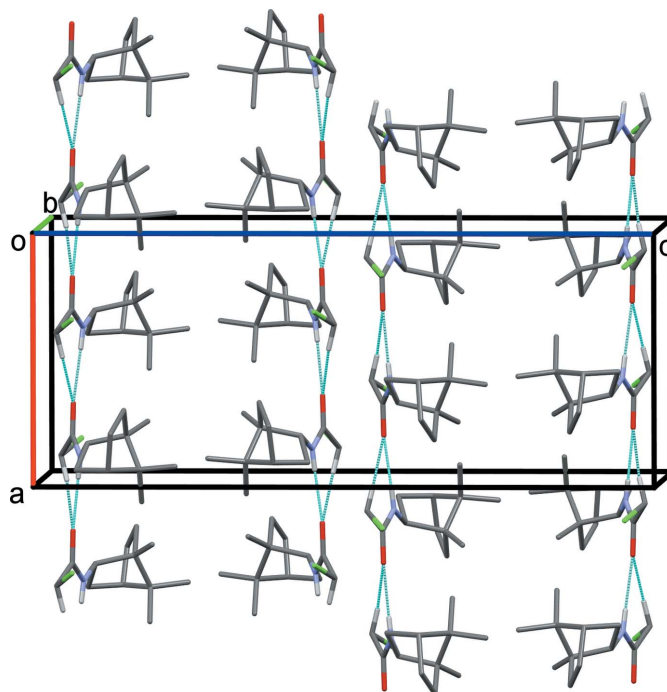


Figure 4
The crystal packing of compound (±)-(2), viewed along the *b* axis. Hydrogen bonds are shown as dashed lines (see Table 2 for details). For clarity, only H atoms involved in these interactions have been included.

at 273 K. Ethyl acetate (30 mL) was added and the phases were separated. The aqueous phase was extracted with ethyl acetate (3 × 20 mL). The combined organic phase was washed with brine, dried over anhydrous Na₂SO₄, filtered and evaporated under reduced pressure. The resulting residue was purified by silica gel column chromatography to provide (±)-(2) (yield: 318 mg, 46%). The NMR data of (±)-(2) fully correspond to those reported earlier (Ung *et al.*, 2014): ¹H NMR (300 MHz, CDCl₃) δ (p.p.m.): 6.63 (*br s*, 1H), 4.03 (*d*, *J* = 1.5 Hz, 2H), 3.88 (*td*, *J* = 9.1, 4.9 Hz, 1H), 1.87 (*dd*, *J* = 13.3, 9.1 Hz, 1H), 1.80–1.52 (*m*, 4H), 1.35–1.23 (*m*, 1H), 1.22–1.10 (*m*, 1H), 0.94 (*s*, 3H), 0.86–0.83 (*m*, 6H); ¹³C NMR (75.5 MHz, CDCl₃) δ (p.p.m.): 164.97, 57.12, 48.69, 47.21, 45.01, 43.00, 39.01, 35.95, 27.10, 20.33, 20.19, 11.82. GC–MS (C₁₂H₂₀³⁵ClNO): *t*_R = 6.21 min; *m/z*: calculated 229.1; found 229.1. (GC–MS method: *vide supra*). X-ray quality single crystals were obtained by slow evaporation of a solution of (±)-(2) in hexanes/ethyl acetate (2:1).

6. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 3. For both compounds, the H atom on the amino group were located in difference Fourier maps and freely refined, and the C-bound H atoms were positioned geometrically and refined as riding on their parent atoms: C–H = 0.93–0.97 Å with *U*_{iso}(H) = 1.5*U*_{eq}(C) for methyl H atoms and 1.2*U*_{eq}(C) for other H atoms. Reflection (0,1,1) whose intensity was affected by the beam-stop was removed from the final refinement of compound (±)-(1).

Table 3
Experimental details.

	(±)-(1)	(±)-(2)
Crystal data		
Chemical formula	C ₁₂ H ₂₁ NO	C ₁₂ H ₂₀ CINO
<i>M_r</i>	195.30	229.74
Crystal system, space group	Monoclinic, <i>P</i> 2 ₁ / <i>n</i>	Orthorhombic, <i>Pcab</i>
Temperature (K)	173	173
<i>a</i> , <i>b</i> , <i>c</i> (Å)	9.6820 (6), 10.6540 (3), 23.3676 (7)	9.6852 (2), 10.7589 (3), 23.7261 (8)
α , β , γ (°)	90, 97.184 (10), 90	90, 90, 90
<i>V</i> (Å ³)	2391.49 (19)	2472.31 (12)
<i>Z</i>	8	8
Radiation type	Mo <i>K</i> α	Mo <i>K</i> α
μ (mm ⁻¹)	0.07	0.29
Crystal size (mm)	0.18 × 0.12 × 0.09	0.35 × 0.10 × 0.09
Data collection		
Diffractometer	Nonius KappaCCD	Nonius KappaCCD
No. of measured, independent and observed [<i>I</i> > 2 σ (<i>I</i>)] reflections	7908, 4320, 2637	6757, 3611, 1854
<i>R</i> _{int}	0.056	0.097
(sin θ / λ) _{max} (Å ⁻¹)	0.600	0.704
Refinement		
<i>R</i> [<i>F</i> ² > 2 σ (<i>F</i> ²)], <i>wR</i> (<i>F</i> ²), <i>S</i>	0.065, 0.159, 1.05	0.073, 0.160, 1.02
No. of reflections	4320	3611
No. of parameters	269	143
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement	H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{\text{max}}$, $\Delta\rho_{\text{min}}$ (e Å ⁻³)	0.37, -0.20	0.51, -0.38

Computer programs: *KappaCCD Server Software* (Nonius, 1997), *DENZO* and *SCALEPACK* (Otwinowski & Minor, 1997), *SIR2011* (Burla *et al.*, 2012), *ORTEP-3 for Windows* (Farrugia, 2012), *Mercury* (Macrae *et al.*, 2008), *SHELXL97* (Sheldrick, 2008), *PLATON* (Spek, 2009) and *pubCIF* (Westrip, 2010).

Acknowledgements

This work was financed by the Latvian Council of Science (grant No 12.0291). DP thanks the JSC ‘Olainfarm’ for a scholarship. JSC ‘Grindeks’ is acknowledged for the donation of organic solvents.

References

- Burla, M. C., Caliandro, R., Camalli, M., Carrozzini, B., Cascarano, G. L., Giacovazzo, C., Mallamo, M., Mazzone, A., Polidori, G. & Spagna, R. (2012). *J. Appl. Cryst.* **45**, 357–361.
- Chelucci, G. (2006). *Chem. Soc. Rev.* **35**, 1230–1243.
- Duclos, R. I. Jr, Lu, D., Guo, J. & Makriyannis, A. (2008). *Tetrahedron Lett.* **49**, 5587–5589.
- Farrugia, L. J. (2012). *J. Appl. Cryst.* **45**, 849–854.
- Groom, C. R. & Allen, F. H. (2014). *Angew. Chem. Int. Ed.* **53**, 662–671.
- Hanzawa, Y., Kasashima, Y., Tomono, K., Mino, T., Sakamoto, M. & Fujita, T. (2012). *J. Oleo Sci.* **61**, 391–399.
- Horváthová, E., Kozics, K., Srančíková, A., Hunáková, L., Gálová, E., Ševčovičová, A. & Slameňová, D. (2012). *Mutagenesis*, **27**, 581–588.
- Jiang, D., He, T., Ma, L. & Wang, Z. (2014). *RSC Adv.* **4**, 64936–64946.
- Langlois, Y. & Kouklovsky, C. (2009). *Synlett*, pp. 3065–3081.
- Macrae, C. F., Bruno, I. J., Chisholm, J. A., Edgington, P. R., McCabe, P., Pidcock, E., Rodriguez-Monge, L., Taylor, R., van de Streek, J. & Wood, P. A. (2008). *J. Appl. Cryst.* **41**, 466–470.
- Nonius (1997). *KappaCCD Server Software*. Nonius BV, Delft, The Netherlands.
- Otwinowski, Z. & Minor, W. (1997). *Methods in Enzymology*, Vol. 276, *Macromolecular Crystallography*, Part A, edited by C. W. Carter Jr & R. M. Sweet, pp. 307–326. New York: Academic Press.
- Prusinowska, N., Bendzińska-Berus, W., Jelecki, M., Rychlewska, U. & Kwit, M. (2015). *Eur. J. Org. Chem.* **2015**, 738–749.
- Ramón, D. J. & Yus, M. (2007). *Synlett*, pp. 2309–2320.
- Schenone, S., Bruno, O., Ranise, A., Bondavalli, F., Filippelli, W., Falcone, G. & Rinaldi, B. (2000). *Farmaco*, **55**, 495–498.
- Seebaluck, R., Gurib-Fakim, A. & Mahomoodally, F. (2015). *J. Ethnopharmacol.* **159**, 137–157.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
- Spek, A. L. (2009). *Acta Cryst.* **D65**, 148–155.
- Stavrov, G., Philipova, I., Valcheva, V. & Momekov, G. (2014a). *Bioorg. Med. Chem. Lett.* **24**, 165–167.
- Stavrov, G., Valcheva, V., Philipova, I. & Doytchinova, I. (2014b). *J. Mol. Graph. Model.* **51**, 7–12.
- Steinbrecher, T., Hrenn, A., Dormann, K. L., Merfort, I. & Labahn, A. (2008). *Bioorg. Med. Chem.* **16**, 2385–2390.
- Turks, M., Strakova, I., Gorovojs, K., Belyakov, S., Piven, Y., Khlebnicova, T. S. & Lakhvich, F. (2012). *Tetrahedron*, **68**, 6131–6140.
- Ung, A. T., Williams, S. G., Angeloski, A., Ashmore, J., Kuzhiumpar-ambal, U., Bhadbhade, M. & Bishop, R. (2014). *Monatsh. Chem.* **145**, 983–992.
- Westrip, S. P. (2010). *J. Appl. Cryst.* **43**, 920–925.

supporting information

Acta Cryst. (2015). E71, 1117-1120 [doi:10.1107/S2056989015015984]

Crystal structures of two (\pm)-*exo*-*N*-isobornylacetamides

Dmitrijs Stepanovs, Daniels Posevins and Maris Turks

Computing details

For both compounds, data collection: *KappaCCD Server Software* (Nonius, 1997); cell refinement: *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *DENZO* and *SCALEPACK* (Otwinowski & Minor, 1997); program(s) used to solve structure: *SIR2011* (Burla *et al.*, 2012); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012) and *Mercury* (Macrae *et al.*, 2008); software used to prepare material for publication: *SHELXL97* (Sheldrick, 2008), *PLATON* (Spek, 2009) and *publCIF* (Westrip, 2010).

(1) (\pm)-*N*-[(1*RS*,2*RS*,4*RS*)-1,7,7-Trimethylbicyclo[2.2.1]heptan-2-yl]acetamide

Crystal data

$C_{12}H_{21}NO$

$M_r = 195.30$

Monoclinic, $P2_1/n$

Hall symbol: -P 2yn

$a = 9.6820$ (6) Å

$b = 10.6540$ (3) Å

$c = 23.3676$ (7) Å

$\beta = 97.184$ (10)°

$V = 2391.49$ (19) Å³

$Z = 8$

$F(000) = 864$

$D_x = 1.085$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 23028 reflections

$\theta = 1.0$ – 30.0 °

$\mu = 0.07$ mm⁻¹

$T = 173$ K

Plate, colourless

$0.18 \times 0.12 \times 0.09$ mm

Data collection

Nonius KappaCCD

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scan

7908 measured reflections

4320 independent reflections

2637 reflections with $I > 2\sigma(I)$

$R_{int} = 0.056$

$\theta_{max} = 25.3$ °, $\theta_{min} = 2.2$ °

$h = -11$ → 11

$k = -12$ → 11

$l = -28$ → 27

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.065$

$wR(F^2) = 0.159$

$S = 1.05$

4320 reflections

269 parameters

0 restraints

Primary atom site location: structure-invariant

direct methods

Secondary atom site location: difference Fourier map

Hydrogen site location: inferred from neighbouring sites

H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0684P)^2 + 0.5097P]$

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

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

$\Delta\rho_{max} = 0.37$ e Å⁻³

$\Delta\rho_{min} = -0.20$ e Å⁻³

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R-factor wR and goodness of fit S are based on F^2 , conventional R-factors R are based on F, with F set to zero for negative F^2 . The threshold expression of $F^2 > 2\sigma(F^2)$ is used only for calculating R-factors(gt) etc. and is not relevant to the choice of reflections for refinement. R-factors based on F^2 are statistically about twice as large as those based on F, and R-factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	−0.02258 (14)	0.12729 (16)	0.06523 (7)	0.0422 (5)
N1	0.18985 (18)	0.21667 (19)	0.07437 (8)	0.0296 (5)
H1N	0.276 (2)	0.204 (2)	0.0728 (9)	0.036 (7)*
C1	0.1084 (3)	0.6083 (3)	0.09760 (13)	0.0543 (8)
H1A	0.0660	0.6280	0.0579	0.065*
H1B	0.1171	0.6865	0.1206	0.065*
C2	0.0227 (2)	0.5082 (2)	0.12569 (12)	0.0474 (7)
H2A	−0.0068	0.5399	0.1621	0.057*
H2B	−0.0610	0.4837	0.0993	0.057*
C3	0.1236 (2)	0.3969 (2)	0.13719 (10)	0.0346 (6)
C4	0.1376 (2)	0.3440 (2)	0.07640 (10)	0.0311 (6)
H4	0.0440	0.3475	0.0529	0.037*
C5	0.2338 (2)	0.4415 (2)	0.05134 (11)	0.0402 (6)
H5A	0.3248	0.4039	0.0460	0.048*
H5B	0.1898	0.4752	0.0140	0.048*
C6	0.2501 (2)	0.5434 (2)	0.09772 (11)	0.0436 (7)
H6	0.3292	0.6024	0.0947	0.052*
C7	0.2638 (2)	0.4669 (2)	0.15414 (10)	0.0385 (6)
C8	0.3926 (2)	0.3821 (3)	0.16342 (12)	0.0478 (7)
H8A	0.3995	0.3333	0.1283	0.072*
H8B	0.3844	0.3248	0.1956	0.072*
H8C	0.4762	0.4339	0.1722	0.072*
C9	0.2675 (3)	0.5485 (3)	0.20856 (12)	0.0583 (8)
H9A	0.3534	0.5981	0.2135	0.087*
H9B	0.2645	0.4945	0.2423	0.087*
H9C	0.1870	0.6050	0.2046	0.087*
C10	0.0805 (3)	0.3019 (3)	0.17942 (12)	0.0481 (7)
H10D	0.0789	0.3419	0.2171	0.072*
H10E	0.1471	0.2323	0.1831	0.072*
H10F	−0.0126	0.2700	0.1654	0.072*
C11	0.1054 (2)	0.1169 (2)	0.06847 (9)	0.0318 (6)
C12	0.1737 (2)	−0.0091 (2)	0.06613 (12)	0.0422 (7)
H12A	0.1673	−0.0544	0.1022	0.063*
H12B	0.2719	0.0022	0.0610	0.063*
H12C	0.1267	−0.0574	0.0337	0.063*

O1A	0.52606 (14)	-0.16700 (16)	-0.05366 (7)	0.0413 (5)
N1A	0.31245 (18)	-0.08247 (18)	-0.07554 (8)	0.0297 (5)
H1AN	0.223 (2)	-0.092 (2)	-0.0758 (9)	0.035 (6)*
C1A	0.3751 (3)	0.2963 (3)	-0.12778 (15)	0.0615 (8)
H1A1	0.4035	0.3332	-0.0892	0.074*
H1A2	0.3675	0.3640	-0.1570	0.074*
C2A	0.4775 (2)	0.1949 (2)	-0.14177 (13)	0.0523 (8)
H2A1	0.5552	0.1864	-0.1102	0.063*
H2A2	0.5161	0.2145	-0.1780	0.063*
C3A	0.3881 (2)	0.0737 (2)	-0.14821 (11)	0.0396 (6)
C4A	0.3603 (2)	0.0434 (2)	-0.08641 (10)	0.0343 (6)
H4A	0.4474	0.0600	-0.0597	0.041*
C5A	0.2472 (3)	0.1450 (3)	-0.07462 (12)	0.0463 (7)
H5A1	0.1567	0.1053	-0.0702	0.056*
H5A2	0.2786	0.1956	-0.0399	0.056*
C6A	0.2383 (3)	0.2240 (3)	-0.12928 (14)	0.0568 (8)
H6A	0.1537	0.2786	-0.1354	0.068*
C7A	0.2459 (2)	0.1260 (3)	-0.17734 (11)	0.0418 (7)
C8A	0.1271 (2)	0.0305 (3)	-0.18521 (12)	0.0530 (8)
H8A1	0.1113	-0.0029	-0.1475	0.079*
H8A2	0.1518	-0.0383	-0.2098	0.079*
H8A3	0.0421	0.0715	-0.2033	0.079*
C9A	0.2545 (3)	0.1849 (4)	-0.23710 (14)	0.0819 (11)
H9A1	0.1685	0.2313	-0.2495	0.123*
H9A2	0.2664	0.1184	-0.2650	0.123*
H9A3	0.3340	0.2424	-0.2347	0.123*
C10A	0.4535 (3)	-0.0307 (3)	-0.17793 (12)	0.0485 (7)
H10A	0.3955	-0.1061	-0.1779	0.073*
H10B	0.5462	-0.0483	-0.1576	0.073*
H10C	0.4618	-0.0063	-0.2178	0.073*
C11A	0.3980 (2)	-0.1770 (2)	-0.05835 (9)	0.0326 (6)
C12A	0.3303 (2)	-0.2983 (3)	-0.04552 (14)	0.0577 (8)
H12D	0.3754	-0.3321	-0.0089	0.087*
H12E	0.3397	-0.3585	-0.0765	0.087*
H12F	0.2314	-0.2838	-0.0428	0.087*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0203 (8)	0.0470 (12)	0.0600 (12)	-0.0052 (7)	0.0079 (7)	-0.0016 (9)
N1	0.0170 (9)	0.0367 (13)	0.0359 (12)	-0.0036 (9)	0.0059 (7)	-0.0039 (10)
C1	0.0598 (17)	0.0345 (17)	0.067 (2)	0.0022 (14)	0.0014 (13)	0.0058 (15)
C2	0.0429 (14)	0.0350 (16)	0.0646 (18)	0.0079 (12)	0.0082 (12)	-0.0045 (14)
C3	0.0343 (12)	0.0313 (15)	0.0397 (15)	-0.0006 (10)	0.0107 (10)	0.0004 (12)
C4	0.0247 (11)	0.0325 (15)	0.0356 (14)	-0.0007 (10)	0.0023 (9)	0.0018 (11)
C5	0.0426 (13)	0.0387 (16)	0.0396 (15)	-0.0083 (12)	0.0064 (10)	0.0103 (13)
C6	0.0446 (14)	0.0321 (16)	0.0540 (17)	-0.0119 (12)	0.0059 (11)	0.0012 (14)
C7	0.0433 (14)	0.0356 (16)	0.0362 (15)	-0.0025 (11)	0.0028 (10)	-0.0070 (13)

C8	0.0377 (13)	0.0527 (19)	0.0494 (17)	-0.0031 (12)	-0.0083 (11)	-0.0073 (14)
C9	0.0698 (18)	0.049 (2)	0.0551 (19)	-0.0046 (15)	0.0030 (14)	-0.0151 (16)
C10	0.0614 (16)	0.0407 (18)	0.0467 (17)	-0.0006 (13)	0.0242 (13)	0.0000 (14)
C11	0.0270 (12)	0.0370 (16)	0.0320 (14)	-0.0064 (11)	0.0064 (9)	-0.0031 (11)
C12	0.0338 (13)	0.0377 (17)	0.0558 (17)	-0.0037 (11)	0.0081 (11)	-0.0057 (13)
O1A	0.0206 (8)	0.0514 (12)	0.0522 (11)	0.0030 (7)	0.0057 (7)	0.0043 (9)
N1A	0.0180 (10)	0.0332 (13)	0.0384 (12)	-0.0016 (9)	0.0056 (7)	0.0044 (9)
C1A	0.0579 (17)	0.045 (2)	0.084 (2)	-0.0083 (14)	0.0174 (15)	-0.0010 (17)
C2A	0.0414 (14)	0.0380 (17)	0.078 (2)	-0.0099 (12)	0.0102 (13)	0.0029 (15)
C3A	0.0366 (13)	0.0381 (16)	0.0450 (16)	-0.0065 (11)	0.0084 (10)	-0.0007 (13)
C4A	0.0249 (11)	0.0332 (15)	0.0437 (15)	-0.0044 (10)	0.0001 (9)	-0.0020 (12)
C5A	0.0415 (14)	0.0390 (17)	0.0590 (18)	-0.0028 (12)	0.0089 (12)	-0.0067 (14)
C6A	0.0408 (15)	0.0461 (19)	0.085 (2)	0.0059 (13)	0.0128 (13)	0.0081 (18)
C7A	0.0384 (13)	0.0450 (17)	0.0410 (15)	-0.0032 (12)	0.0016 (10)	0.0135 (14)
C8A	0.0400 (14)	0.064 (2)	0.0507 (17)	-0.0095 (14)	-0.0097 (11)	0.0142 (16)
C9A	0.079 (2)	0.091 (3)	0.072 (2)	-0.009 (2)	-0.0030 (17)	0.040 (2)
C10A	0.0528 (15)	0.0471 (19)	0.0481 (16)	-0.0036 (13)	0.0164 (12)	0.0013 (14)
C11A	0.0262 (12)	0.0401 (16)	0.0324 (14)	-0.0002 (11)	0.0065 (9)	0.0024 (12)
C12A	0.0390 (14)	0.0480 (19)	0.086 (2)	0.0008 (13)	0.0093 (13)	0.0232 (17)

Geometric parameters (Å, °)

O1—C11	1.237 (2)	O1A—C11A	1.236 (2)
N1—C11	1.338 (3)	N1A—C11A	1.334 (3)
N1—C4	1.451 (3)	N1A—C4A	1.452 (3)
N1—H1N	0.85 (2)	N1A—H1AN	0.87 (2)
C1—C6	1.536 (3)	C1A—C6A	1.529 (4)
C1—C2	1.547 (4)	C1A—C2A	1.529 (4)
C1—H1A	0.9900	C1A—H1A1	0.9900
C1—H1B	0.9900	C1A—H1A2	0.9900
C2—C3	1.539 (3)	C2A—C3A	1.552 (3)
C2—H2A	0.9900	C2A—H2A1	0.9900
C2—H2B	0.9900	C2A—H2A2	0.9900
C3—C10	1.508 (3)	C3A—C10A	1.494 (3)
C3—C4	1.550 (3)	C3A—C4A	1.536 (3)
C3—C7	1.556 (3)	C3A—C7A	1.560 (3)
C4—C5	1.557 (3)	C4A—C5A	1.588 (3)
C4—H4	1.0000	C4A—H4A	1.0000
C5—C6	1.528 (4)	C5A—C6A	1.523 (4)
C5—H5A	0.9900	C5A—H5A1	0.9900
C5—H5B	0.9900	C5A—H5A2	0.9900
C6—C7	1.541 (4)	C6A—C7A	1.542 (4)
C6—H6	1.0000	C6A—H6A	1.0000
C7—C8	1.534 (3)	C7A—C8A	1.530 (3)
C7—C9	1.537 (4)	C7A—C9A	1.543 (4)
C8—H8A	0.9800	C8A—H8A1	0.9800
C8—H8B	0.9800	C8A—H8A2	0.9800
C8—H8C	0.9800	C8A—H8A3	0.9800

C9—H9A	0.9800	C9A—H9A1	0.9800
C9—H9B	0.9800	C9A—H9A2	0.9800
C9—H9C	0.9800	C9A—H9A3	0.9800
C10—H10D	0.9800	C10A—H10A	0.9800
C10—H10E	0.9800	C10A—H10B	0.9800
C10—H10F	0.9800	C10A—H10C	0.9800
C11—C12	1.500 (3)	C11A—C12A	1.497 (3)
C12—H12A	0.9800	C12A—H12D	0.9800
C12—H12B	0.9800	C12A—H12E	0.9800
C12—H12C	0.9800	C12A—H12F	0.9800
C11—N1—C4	122.38 (18)	C11A—N1A—C4A	123.40 (18)
C11—N1—H1N	117.6 (16)	C11A—N1A—H1AN	119.6 (16)
C4—N1—H1N	119.6 (16)	C4A—N1A—H1AN	116.4 (16)
C6—C1—C2	102.5 (2)	C6A—C1A—C2A	102.8 (2)
C6—C1—H1A	111.3	C6A—C1A—H1A1	111.2
C2—C1—H1A	111.3	C2A—C1A—H1A1	111.2
C6—C1—H1B	111.3	C6A—C1A—H1A2	111.2
C2—C1—H1B	111.3	C2A—C1A—H1A2	111.2
H1A—C1—H1B	109.2	H1A1—C1A—H1A2	109.1
C3—C2—C1	104.02 (18)	C1A—C2A—C3A	103.87 (19)
C3—C2—H2A	111.0	C1A—C2A—H2A1	111.0
C1—C2—H2A	111.0	C3A—C2A—H2A1	111.0
C3—C2—H2B	111.0	C1A—C2A—H2A2	111.0
C1—C2—H2B	111.0	C3A—C2A—H2A2	111.0
H2A—C2—H2B	109.0	H2A1—C2A—H2A2	109.0
C10—C3—C2	114.18 (18)	C10A—C3A—C4A	114.5 (2)
C10—C3—C4	114.7 (2)	C10A—C3A—C2A	113.63 (19)
C2—C3—C4	104.27 (19)	C4A—C3A—C2A	104.2 (2)
C10—C3—C7	117.3 (2)	C10A—C3A—C7A	117.7 (2)
C2—C3—C7	100.96 (19)	C4A—C3A—C7A	103.72 (17)
C4—C3—C7	103.54 (16)	C2A—C3A—C7A	101.3 (2)
N1—C4—C3	116.15 (19)	N1A—C4A—C3A	117.1 (2)
N1—C4—C5	112.55 (17)	N1A—C4A—C5A	110.91 (17)
C3—C4—C5	103.13 (18)	C3A—C4A—C5A	103.01 (19)
N1—C4—H4	108.2	N1A—C4A—H4A	108.5
C3—C4—H4	108.2	C3A—C4A—H4A	108.5
C5—C4—H4	108.2	C5A—C4A—H4A	108.5
C6—C5—C4	102.74 (18)	C6A—C5A—C4A	101.5 (2)
C6—C5—H5A	111.2	C6A—C5A—H5A1	111.5
C4—C5—H5A	111.2	C4A—C5A—H5A1	111.5
C6—C5—H5B	111.2	C6A—C5A—H5A2	111.5
C4—C5—H5B	111.2	C4A—C5A—H5A2	111.5
H5A—C5—H5B	109.1	H5A1—C5A—H5A2	109.3
C5—C6—C1	107.8 (2)	C5A—C6A—C1A	107.5 (2)
C5—C6—C7	102.8 (2)	C5A—C6A—C7A	103.5 (2)
C1—C6—C7	102.75 (19)	C1A—C6A—C7A	103.8 (2)
C5—C6—H6	114.1	C5A—C6A—H6A	113.7

C1—C6—H6	114.1	C1A—C6A—H6A	113.7
C7—C6—H6	114.1	C7A—C6A—H6A	113.7
C8—C7—C9	106.4 (2)	C8A—C7A—C6A	115.7 (2)
C8—C7—C6	114.5 (2)	C8A—C7A—C9A	106.6 (2)
C9—C7—C6	113.5 (2)	C6A—C7A—C9A	113.4 (3)
C8—C7—C3	114.9 (2)	C8A—C7A—C3A	115.1 (2)
C9—C7—C3	114.28 (19)	C6A—C7A—C3A	92.51 (19)
C6—C7—C3	93.26 (18)	C9A—C7A—C3A	113.3 (2)
C7—C8—H8A	109.5	C7A—C8A—H8A1	109.5
C7—C8—H8B	109.5	C7A—C8A—H8A2	109.5
H8A—C8—H8B	109.5	H8A1—C8A—H8A2	109.5
C7—C8—H8C	109.5	C7A—C8A—H8A3	109.5
H8A—C8—H8C	109.5	H8A1—C8A—H8A3	109.5
H8B—C8—H8C	109.5	H8A2—C8A—H8A3	109.5
C7—C9—H9A	109.5	C7A—C9A—H9A1	109.5
C7—C9—H9B	109.5	C7A—C9A—H9A2	109.5
H9A—C9—H9B	109.5	H9A1—C9A—H9A2	109.5
C7—C9—H9C	109.5	C7A—C9A—H9A3	109.5
H9A—C9—H9C	109.5	H9A1—C9A—H9A3	109.5
H9B—C9—H9C	109.5	H9A2—C9A—H9A3	109.5
C3—C10—H10D	109.5	C3A—C10A—H10A	109.5
C3—C10—H10E	109.5	C3A—C10A—H10B	109.5
H10D—C10—H10E	109.5	H10A—C10A—H10B	109.5
C3—C10—H10F	109.5	C3A—C10A—H10C	109.5
H10D—C10—H10F	109.5	H10A—C10A—H10C	109.5
H10E—C10—H10F	109.5	H10B—C10A—H10C	109.5
O1—C11—N1	122.0 (2)	O1A—C11A—N1A	122.7 (2)
O1—C11—C12	121.4 (2)	O1A—C11A—C12A	121.1 (2)
N1—C11—C12	116.64 (18)	N1A—C11A—C12A	116.19 (19)
C11—C12—H12A	109.5	C11A—C12A—H12D	109.5
C11—C12—H12B	109.5	C11A—C12A—H12E	109.5
H12A—C12—H12B	109.5	H12D—C12A—H12E	109.5
C11—C12—H12C	109.5	C11A—C12A—H12F	109.5
H12A—C12—H12C	109.5	H12D—C12A—H12F	109.5
H12B—C12—H12C	109.5	H12E—C12A—H12F	109.5
C6—C1—C2—C3	-1.4 (3)	C6A—C1A—C2A—C3A	-1.5 (3)
C1—C2—C3—C10	163.3 (2)	C1A—C2A—C3A—C10A	163.7 (2)
C1—C2—C3—C4	-70.7 (2)	C1A—C2A—C3A—C4A	-71.1 (2)
C1—C2—C3—C7	36.5 (2)	C1A—C2A—C3A—C7A	36.4 (3)
C11—N1—C4—C3	91.9 (2)	C11A—N1A—C4A—C3A	91.2 (3)
C11—N1—C4—C5	-149.5 (2)	C11A—N1A—C4A—C5A	-151.0 (2)
C10—C3—C4—N1	-35.4 (3)	C10A—C3A—C4A—N1A	-38.7 (3)
C2—C3—C4—N1	-161.05 (18)	C2A—C3A—C4A—N1A	-163.36 (18)
C7—C3—C4—N1	93.7 (2)	C7A—C3A—C4A—N1A	91.0 (2)
C10—C3—C4—C5	-159.00 (19)	C10A—C3A—C4A—C5A	-160.7 (2)
C2—C3—C4—C5	75.4 (2)	C2A—C3A—C4A—C5A	74.6 (2)
C7—C3—C4—C5	-29.9 (2)	C7A—C3A—C4A—C5A	-31.1 (2)

N1—C4—C5—C6	-131.5 (2)	N1A—C4A—C5A—C6A	-131.0 (2)
C3—C4—C5—C6	-5.6 (2)	C3A—C4A—C5A—C6A	-4.9 (2)
C4—C5—C6—C1	-68.3 (2)	C4A—C5A—C6A—C1A	-69.5 (3)
C4—C5—C6—C7	39.7 (2)	C4A—C5A—C6A—C7A	39.9 (2)
C2—C1—C6—C5	73.3 (2)	C2A—C1A—C6A—C5A	74.5 (3)
C2—C1—C6—C7	-34.8 (3)	C2A—C1A—C6A—C7A	-34.7 (3)
C5—C6—C7—C8	63.2 (2)	C5A—C6A—C7A—C8A	62.4 (3)
C1—C6—C7—C8	175.1 (2)	C1A—C6A—C7A—C8A	174.6 (2)
C5—C6—C7—C9	-174.4 (2)	C5A—C6A—C7A—C9A	-173.9 (2)
C1—C6—C7—C9	-62.5 (3)	C1A—C6A—C7A—C9A	-61.7 (3)
C5—C6—C7—C3	-56.1 (2)	C5A—C6A—C7A—C3A	-57.1 (2)
C1—C6—C7—C3	55.8 (2)	C1A—C6A—C7A—C3A	55.1 (2)
C10—C3—C7—C8	60.6 (3)	C10A—C3A—C7A—C8A	60.7 (3)
C2—C3—C7—C8	-174.6 (2)	C4A—C3A—C7A—C8A	-66.9 (3)
C4—C3—C7—C8	-66.9 (2)	C2A—C3A—C7A—C8A	-174.8 (2)
C10—C3—C7—C9	-62.8 (3)	C10A—C3A—C7A—C6A	-179.3 (2)
C2—C3—C7—C9	62.0 (3)	C4A—C3A—C7A—C6A	53.1 (2)
C4—C3—C7—C9	169.7 (2)	C2A—C3A—C7A—C6A	-54.7 (2)
C10—C3—C7—C6	179.6 (2)	C10A—C3A—C7A—C9A	-62.4 (3)
C2—C3—C7—C6	-55.7 (2)	C4A—C3A—C7A—C9A	170.0 (2)
C4—C3—C7—C6	52.0 (2)	C2A—C3A—C7A—C9A	62.1 (3)
C4—N1—C11—O1	-1.1 (3)	C4A—N1A—C11A—O1A	-4.6 (3)
C4—N1—C11—C12	179.1 (2)	C4A—N1A—C11A—C12A	176.2 (2)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N1—H1N...O1A ⁱ	0.85 (2)	2.06 (2)	2.900 (2)	170 (2)
N1A—H1AN...O1 ⁱⁱ	0.87 (2)	2.03 (2)	2.886 (2)	172 (2)
C8A—H8A1...O1 ⁱⁱ	0.98	2.57	3.524 (3)	165
C12—H12C...O1 ⁱⁱ	0.98	2.52	3.468 (3)	164

Symmetry codes: (i) $-x+1, -y, -z$; (ii) $-x, -y, -z$.(2) (\pm)-2-Chloro-*N*-[(1*RS*,2*RS*,4*RS*)-1,7,7-trimethylbicyclo[2.2.1]heptan-2-yl]acetamide

Crystal data

C₁₂H₂₀ClNO*M_r* = 229.74Orthorhombic, *Pcab*

Hall symbol: -P 2bc 2ac

a = 9.6852 (2) Å*b* = 10.7589 (3) Å*c* = 23.7261 (8) Å*V* = 2472.31 (12) Å³*Z* = 8*F*(000) = 992*D_x* = 1.234 Mg m⁻³Mo *K*α radiation, λ = 0.71073 Å

Cell parameters from 24915 reflections

θ = 1.0–30.0°

μ = 0.28 mm⁻¹*T* = 173 K

Plate, colorless

0.35 × 0.10 × 0.09 mm

Data collection

Nonius KappaCCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scan

6757 measured reflections

3611 independent reflections

1854 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.097$

$\theta_{\text{max}} = 30.0^\circ$, $\theta_{\text{min}} = 2.6^\circ$

$h = -13 \rightarrow 13$

$k = -15 \rightarrow 15$

$l = -33 \rightarrow 33$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.073$

$wR(F^2) = 0.160$

$S = 1.02$

3611 reflections

143 parameters

0 restraints

Primary atom site location: structure-invariant

direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites

H atoms treated by a mixture of independent
and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0497P)^2 + 1.7639P]$

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

$(\Delta/\sigma)_{\text{max}} < 0.001$

$\Delta\rho_{\text{max}} = 0.51 \text{ e } \text{\AA}^{-3}$

$\Delta\rho_{\text{min}} = -0.38 \text{ e } \text{\AA}^{-3}$

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Cl1	0.83924 (7)	1.01572 (7)	0.43765 (4)	0.0460 (3)
O1	0.68852 (15)	0.76668 (17)	0.44335 (8)	0.0326 (5)
N1	0.8914 (2)	0.6671 (2)	0.42922 (10)	0.0241 (5)
H1	0.972 (3)	0.673 (2)	0.4329 (11)	0.026 (8)*
C1	0.7790 (3)	0.3051 (3)	0.36991 (15)	0.0436 (8)
H1A	0.7757	0.2399	0.3417	0.052*
H1B	0.7467	0.2723	0.4056	0.052*
C2	0.6933 (3)	0.4183 (3)	0.35161 (14)	0.0377 (7)
H2A	0.6194	0.4345	0.3782	0.045*
H2B	0.6539	0.4055	0.3145	0.045*
C3	0.7980 (2)	0.5264 (2)	0.35096 (12)	0.0292 (6)
C4	0.8313 (2)	0.5475 (2)	0.41407 (11)	0.0260 (6)
H4	0.7456	0.5373	0.4355	0.031*
C5	0.9296 (3)	0.4360 (2)	0.42903 (13)	0.0332 (7)
H5A	1.0224	0.4647	0.4373	0.040*
H5B	0.8950	0.3891	0.4609	0.040*

C6	0.9261 (3)	0.3591 (3)	0.37543 (15)	0.0409 (8)
H6	0.9999	0.2970	0.3727	0.049*
C7	0.9285 (3)	0.4580 (3)	0.32796 (12)	0.0321 (7)
C8	1.0598 (3)	0.5389 (3)	0.32675 (14)	0.0435 (8)
H8A	1.1362	0.4900	0.3136	0.065*
H8B	1.0792	0.5688	0.3640	0.065*
H8C	1.0458	0.6082	0.3019	0.065*
C9	0.9126 (4)	0.4040 (4)	0.26906 (16)	0.0626 (11)
H9A	0.9126	0.4701	0.2419	0.094*
H9B	0.8271	0.3592	0.2667	0.094*
H9C	0.9880	0.3486	0.2614	0.094*
C10	0.7496 (3)	0.6398 (3)	0.31989 (14)	0.0427 (8)
H10A	0.7357	0.6197	0.2809	0.064*
H10B	0.8179	0.7042	0.3230	0.064*
H10C	0.6642	0.6682	0.3359	0.064*
C11	0.8159 (2)	0.7647 (2)	0.44464 (11)	0.0239 (6)
C12	0.8982 (2)	0.8735 (2)	0.46654 (14)	0.0324 (7)
H12A	0.9947	0.8620	0.4571	0.039*
H12B	0.8906	0.8767	0.5073	0.039*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C11	0.0364 (4)	0.0305 (4)	0.0711 (6)	−0.0062 (3)	−0.0118 (4)	0.0006 (4)
O1	0.0136 (8)	0.0345 (10)	0.0499 (13)	0.0001 (7)	−0.0002 (8)	−0.0073 (10)
N1	0.0120 (10)	0.0280 (11)	0.0323 (14)	−0.0021 (8)	−0.0006 (9)	−0.0028 (10)
C1	0.0349 (15)	0.0323 (15)	0.063 (2)	−0.0059 (12)	−0.0009 (15)	−0.0040 (16)
C2	0.0213 (13)	0.0361 (16)	0.056 (2)	−0.0064 (11)	0.0011 (12)	−0.0065 (15)
C3	0.0238 (12)	0.0276 (14)	0.0363 (17)	−0.0023 (10)	−0.0014 (11)	−0.0005 (13)
C4	0.0203 (12)	0.0248 (13)	0.0329 (16)	−0.0026 (10)	0.0057 (11)	0.0000 (11)
C5	0.0273 (13)	0.0285 (13)	0.0439 (18)	0.0001 (11)	0.0000 (12)	0.0083 (14)
C6	0.0261 (14)	0.0277 (15)	0.069 (2)	0.0058 (11)	−0.0011 (14)	−0.0065 (15)
C7	0.0247 (13)	0.0378 (16)	0.0338 (17)	−0.0059 (11)	0.0055 (11)	−0.0111 (14)
C8	0.0324 (15)	0.054 (2)	0.0442 (19)	−0.0137 (14)	0.0148 (14)	−0.0130 (16)
C9	0.0484 (19)	0.078 (3)	0.062 (3)	−0.0176 (18)	0.0151 (17)	−0.035 (2)
C10	0.0449 (17)	0.0419 (17)	0.0412 (19)	0.0003 (15)	−0.0156 (14)	−0.0008 (15)
C11	0.0161 (11)	0.0268 (13)	0.0289 (15)	−0.0006 (9)	−0.0008 (10)	−0.0028 (12)
C12	0.0180 (12)	0.0304 (14)	0.0488 (19)	0.0023 (10)	−0.0066 (12)	−0.0108 (14)

Geometric parameters (Å, °)

C11—C12	1.771 (3)	C5—H5A	0.9700
O1—C11	1.235 (3)	C5—H5B	0.9700
N1—C11	1.331 (3)	C6—C7	1.550 (4)
N1—C4	1.457 (3)	C6—H6	0.9800
N1—H1	0.78 (3)	C7—C9	1.522 (4)
C1—C2	1.536 (4)	C7—C8	1.541 (4)
C1—C6	1.544 (4)	C8—H8A	0.9600

C1—H1A	0.9700	C8—H8B	0.9600
C1—H1B	0.9700	C8—H8C	0.9600
C2—C3	1.543 (4)	C9—H9A	0.9600
C2—H2A	0.9700	C9—H9B	0.9600
C2—H2B	0.9700	C9—H9C	0.9600
C3—C10	1.501 (4)	C10—H10A	0.9600
C3—C4	1.548 (4)	C10—H10B	0.9600
C3—C7	1.561 (4)	C10—H10C	0.9600
C4—C5	1.573 (4)	C11—C12	1.508 (3)
C4—H4	0.9800	C12—H12A	0.9700
C5—C6	1.517 (4)	C12—H12B	0.9700
C11—N1—C4	123.0 (2)	C5—C6—H6	114.2
C11—N1—H1	117 (2)	C1—C6—H6	114.2
C4—N1—H1	119 (2)	C7—C6—H6	114.2
C2—C1—C6	102.9 (2)	C9—C7—C8	106.4 (2)
C2—C1—H1A	111.2	C9—C7—C6	113.8 (3)
C6—C1—H1A	111.2	C8—C7—C6	114.4 (2)
C2—C1—H1B	111.2	C9—C7—C3	114.8 (2)
C6—C1—H1B	111.2	C8—C7—C3	114.1 (2)
H1A—C1—H1B	109.1	C6—C7—C3	93.3 (2)
C1—C2—C3	104.2 (2)	C7—C8—H8A	109.5
C1—C2—H2A	110.9	C7—C8—H8B	109.5
C3—C2—H2A	110.9	H8A—C8—H8B	109.5
C1—C2—H2B	110.9	C7—C8—H8C	109.5
C3—C2—H2B	110.9	H8A—C8—H8C	109.5
H2A—C2—H2B	108.9	H8B—C8—H8C	109.5
C10—C3—C2	114.3 (2)	C7—C9—H9A	109.5
C10—C3—C4	114.9 (2)	C7—C9—H9B	109.5
C2—C3—C4	103.7 (2)	H9A—C9—H9B	109.5
C10—C3—C7	117.7 (2)	C7—C9—H9C	109.5
C2—C3—C7	100.4 (2)	H9A—C9—H9C	109.5
C4—C3—C7	103.8 (2)	H9B—C9—H9C	109.5
N1—C4—C3	116.8 (2)	C3—C10—H10A	109.5
N1—C4—C5	112.1 (2)	C3—C10—H10B	109.5
C3—C4—C5	103.4 (2)	H10A—C10—H10B	109.5
N1—C4—H4	108.0	C3—C10—H10C	109.5
C3—C4—H4	108.0	H10A—C10—H10C	109.5
C5—C4—H4	108.0	H10B—C10—H10C	109.5
C6—C5—C4	102.3 (2)	O1—C11—N1	123.8 (2)
C6—C5—H5A	111.3	O1—C11—C12	121.5 (2)
C4—C5—H5A	111.3	N1—C11—C12	114.63 (19)
C6—C5—H5B	111.3	C11—C12—C11	111.55 (18)
C4—C5—H5B	111.3	C11—C12—H12A	109.3
H5A—C5—H5B	109.2	C11—C12—H12A	109.3
C5—C6—C1	107.3 (2)	C11—C12—H12B	109.3
C5—C6—C7	103.6 (2)	C11—C12—H12B	109.3
C1—C6—C7	102.1 (2)	H12A—C12—H12B	108.0

C6—C1—C2—C3	-2.3 (3)	C1—C6—C7—C9	-63.4 (3)
C1—C2—C3—C10	164.5 (3)	C5—C6—C7—C8	62.6 (3)
C1—C2—C3—C4	-69.6 (3)	C1—C6—C7—C8	174.0 (2)
C1—C2—C3—C7	37.5 (3)	C5—C6—C7—C3	-55.8 (2)
C11—N1—C4—C3	91.6 (3)	C1—C6—C7—C3	55.6 (2)
C11—N1—C4—C5	-149.3 (2)	C10—C3—C7—C9	-62.8 (3)
C10—C3—C4—N1	-35.2 (3)	C2—C3—C7—C9	61.9 (3)
C2—C3—C4—N1	-160.7 (2)	C4—C3—C7—C9	168.9 (2)
C7—C3—C4—N1	94.7 (2)	C10—C3—C7—C8	60.3 (3)
C10—C3—C4—C5	-158.8 (2)	C2—C3—C7—C8	-175.0 (3)
C2—C3—C4—C5	75.6 (2)	C4—C3—C7—C8	-67.9 (3)
C7—C3—C4—C5	-28.9 (2)	C10—C3—C7—C6	179.0 (2)
N1—C4—C5—C6	-132.9 (2)	C2—C3—C7—C6	-56.3 (2)
C3—C4—C5—C6	-6.2 (2)	C4—C3—C7—C6	50.8 (2)
C4—C5—C6—C1	-67.7 (3)	C4—N1—C11—O1	-6.0 (4)
C4—C5—C6—C7	39.8 (2)	C4—N1—C11—C12	171.7 (2)
C2—C1—C6—C5	74.5 (3)	O1—C11—C12—C11	-47.2 (3)
C2—C1—C6—C7	-34.1 (3)	N1—C11—C12—C11	135.1 (2)
C5—C6—C7—C9	-174.8 (2)		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N1—H1...O1 ⁱ	0.79 (3)	2.21 (3)	2.983 (2)	168 (2)
C12—H12 <i>A</i> ...O1 ⁱ	0.97	2.36	3.238 (3)	151

Symmetry code: (i) $x+1/2, -y+3/2, z$.