

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

(+)-(1*S*,5*R*,6*R*)-6-[(*S*)-1-Hydroxy-2-(methoxymethyloxy)ethyl]-1-methyl-3trichloromethyl-2-aza-4,7-dioxabicyclo-[3.3.0]oct-2-en-8-one

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Received 9 October 2012; accepted 15 October 2012

Key indicators: single-crystal X-ray study; T = 90 K; mean σ (C–C) = 0.003 Å; R factor = 0.027; wR factor = 0.069; data-to-parameter ratio = 12.2.

In the title compound, $C_{11}H_{14}Cl_3NO_6$, the fused fivemembered oxazoline and tetrahydrofuran rings are essentially planar with maximum deviations of 0.069 (1) and 0.031 (1) Å, respectively, and make a dihedral angle of 64.23 (11)° with each other. In the crystal, molecules are linked by $O-H\cdots O$ and $C-H\cdots O$ hydrogen bonds, forming chains along the *b*axis direction. Further $C-H\cdots O$ hydrogen bonds are observed between the chains.

Related literature

For the synthesis, see: Oishi *et al.* (2012). For the isolation of sphingofungins, see: VanMiddlesworth, Giacobbe *et al.* (1992); VanMiddlesworth, Dufresne *et al.* (1992); Horn *et al.* (1992).



Experimental

Crystal data $C_{11}H_{14}Cl_3NO_6$ $M_r = 362.58$ Monoclinic, $P2_1$ a = 8.9311 (7) Å

b = 6.0283 (4) Å c = 13.8694 (10) Å $\beta = 99.699 (2)^{\circ}$ $V = 736.05 (9) \text{ Å}^{3}$ Z = 2Mo $K\alpha$ radiation $\mu = 0.65 \text{ mm}^{-1}$

Data collection

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.027$ $wR(F^2) = 0.069$ S = 1.342364 reflections 193 parameters 1 restraint

6662 measured reflections

2364 independent reflections 2291 reflections with $I > 2\sigma(I)$ $R_{int} = 0.028$ 227 H-atom parameters constrained $\Delta \rho_{max} = 0.37 \text{ e } \text{\AA}^{-3}$ $\Delta \rho_{min} = -0.22 \text{ e } \text{\AA}^{-3}$ Absolute structure: Flack (1983), 947 Friedel pairs

 $T = 90 \, {\rm K}$

Flack parameter: 0.01 (5)

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
O12−H12···O15 ⁱ	0.84	1.90	2.695 (2)	157
$C6-H6\cdots O9^n$	1.00	2.43	3.402 (3)	164
$C17 - H17C \cdot \cdot \cdot O9^{iii}$	0.98	2.53	3.320 (3)	137

Symmetry codes: (i) x, y - 1, z; (ii) x, y + 1, z; (iii) $-x + 1, y + \frac{1}{2}, -z + 1$.

Data collection: *APEX2* (Bruker, 2012); cell refinement: *SAINT* (Bruker, 2012); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *Mercury* (Macrae *et al.*, 2006); software used to prepare material for publication: *SHELXL97*.

We thank Professor S. Ohba, Professor N. Yoshioka and Dr C. Maeda (Keio University, Japan) for providing valuable advice.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: IS5207).

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 $0.50 \times 0.25 \times 0.16 \text{ mm}$

supporting information

Acta Cryst. (2012). E68, o3185 [doi:10.1107/S1600536812042912]

(+)-(1*S*,5*R*,6*R*)-6-[(*S*)-1-Hydroxy-2-(methoxymethyloxy)ethyl]-1-methyl-3-trichloromethyl-2-aza-4,7-dioxabicyclo[3.3.0]oct-2-en-8-one

Takeshi Oishi, Hiroki Oishi, Syun Tsuzaki, Takaaki Sato and Noritaka Chida

S1. Comment

Sphingofungins are natural antifungal agents isolated from *Aspergillus* and reported to be potent inhibitors of the biosynthesis of sphingolipids (VanMiddlesworth, Giacobbe *et al.*, 1992; VanMiddlesworth, Dufresne *et al.*, 1992; Horn *et al.*, 1992). The title compound (I), $C_{11}H_{14}Cl_3NO_6$, which has four contiguous stereogenic center including a tetrasubstituted carbon with nitrogen (Fig. 1), was provided in a synthetic study on the natural products sphingofungins from $_D$ -ribose (Oishi *et al.*, 2012). The absolute configurations were confirmed by the X-ray analysis as C1*S*, C5*R*, C6*R* and C10*S*. The crystal packing was stabilized by an intermolecular O—H…O hydrogen bond, forming molecular a chain along the *b* axis (Fig. 2). There are also C—H…O hydrogen bonds and intermolecular Cl…O short contacts, Cl19…O4 (*x*, *y* - 1, *z*) and Cl20…O7 (*x* + 1, *y*, *z*) being 3.070 (2) and 3.142 (2) Å, respectively.

S2. Experimental

The title compound was obtained in a synthetic study of sphingofungins from _D-ribose (Oishi *et al.*, 2012), and recrystallized from ethyl acetate solution. $[a]^{27}_{D}$ +83.1 (*c* 0.345, CHCl₃); m.p. 409.7–411.2 K.

S3. Refinement

C-bound H atoms were positioned geometrically with C—H = 0.98–1.00 Å, and constrained to ride on their parent atoms with $U_{iso}(H) = 1.2U_{eq}(C)$ or $1.5U_{eq}(methyl C)$. The H atom of hydroxyl group (O12) was placed guided by difference maps, with O—H = 0.84 Å and with $U_{iso}(H) = 1.5U_{eq}(O)$. Three reflections (6 3 6, 6 3 7, 0 1 16) have been omitted in the final refinement.



Figure 1

The molecular structure of the title compound, with atom labels and 50% probability displacement ellipsoids for non-H atoms.



Figure 2

The crystal packing of the title compound, viewed down the *a* axis. The dashed lines indicate O—H…O hydrogen bonds.

(+)-(1*S*,5*R*,6*R*)-6-[(*S*)-1-Hydroxy-2-(methoxymethyloxy)ethyl]-1-methyl-3-trichloromethyl-2-aza-4,7-

dioxabicyclo[3.3.0]oct-2-en-8-one

Crystal data $C_{11}H_{14}Cl_3NO_6$ $M_r = 362.58$ Monoclinic, $P2_1$ a = 8.9311 (7) Å b = 6.0283 (4) Å c = 13.8694 (10) Å $\beta = 99.699$ (2)° V = 736.05 (9) Å³ Z = 2F(000) = 372

 $D_x = 1.636 \text{ Mg m}^{-3}$ Melting point: 409.7 K Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ Å}$ Cell parameters from 5778 reflections $\theta = 2.3-25.1^{\circ}$ $\mu = 0.65 \text{ mm}^{-1}$ T = 90 KPrism, colourless $0.50 \times 0.25 \times 0.16 \text{ mm}$ Data collection

Bruker D8 goniometer diffractometer Radiation source: fine-focus sealed tube Graphite monochromator Detector resolution: 10.4167 pixels mm ⁻¹ ω scans Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 2012) $T_{\min} = 0.738, T_{\max} = 0.904$	6662 measured reflections 2364 independent reflections 2291 reflections with $I > 2\sigma(I)$ $R_{int} = 0.028$ $\theta_{max} = 25.1^{\circ}, \theta_{min} = 2.3^{\circ}$ $h = -10 \rightarrow 10$ $k = -7 \rightarrow 6$ $l = -16 \rightarrow 16$
Refinement	
Refinement on F^2 Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.027$ $wR(F^2) = 0.069$ S = 1.34 2364 reflections 193 parameters 1 restraint Primary atom site location: structure-invariant direct methods Secondary atom site location: difference Fourier	Hydrogen site location: inferred from neighbouring sites H-atom parameters constrained $w = 1/[\sigma^2(F_o^2) + (0.0338P)^2]$ where $P = (F_o^2 + 2F_o^2)/3$ $(\Delta/\sigma)_{max} = 0.001$ $\Delta\rho_{max} = 0.37$ e Å ⁻³ $\Delta\rho_{min} = -0.22$ e Å ⁻³ Absolute structure: Flack (1983), 947 Friedel pairs Absolute structure parameter: 0.01 (5)

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 $(Å^2)$

	x	У	Ζ	$U_{ m iso}$ */ $U_{ m eq}$
C1	0.7195 (2)	0.5236 (4)	0.43062 (15)	0.0152 (5)
N2	0.7869 (2)	0.3750 (3)	0.36335 (12)	0.0139 (4)
C3	0.8554 (2)	0.5035 (4)	0.31378 (15)	0.0134 (5)
O4	0.85179 (17)	0.7255 (3)	0.32714 (10)	0.0156 (3)
C5	0.7464 (2)	0.7590 (4)	0.39563 (15)	0.0148 (5)
H5	0.7928	0.8550	0.4516	0.018*
C6	0.5893 (2)	0.8481 (4)	0.34831 (15)	0.0142 (5)
H6	0.5652	0.9825	0.3851	0.017*
O7	0.48036 (16)	0.6720 (3)	0.36046 (10)	0.0162 (4)
C8	0.5464 (2)	0.4965 (4)	0.40903 (15)	0.0142 (5)
09	0.47385 (18)	0.3412 (3)	0.42994 (10)	0.0181 (4)
C10	0.5688 (3)	0.9041 (4)	0.24039 (15)	0.0161 (5)
H10	0.6464	1.0185	0.2316	0.019*
C11	0.4125 (3)	1.0049 (4)	0.20566 (15)	0.0165 (5)

H11A	0.3335	0.9027	0.2216	0.020*
H11B	0.4039	1.1462	0.2408	0.020*
O12	0.59596 (18)	0.7153 (3)	0.18539 (11)	0.0208 (4)
H12	0.5132	0.6520	0.1642	0.031*
O13	0.38660 (19)	1.0460 (3)	0.10212 (11)	0.0205 (4)
C14	0.4558 (3)	1.2396 (4)	0.07599 (16)	0.0192 (5)
H14A	0.4577	1.2365	0.0049	0.023*
H14B	0.5622	1.2437	0.1105	0.023*
015	0.38097 (18)	1.4326 (3)	0.09848 (11)	0.0195 (4)
C16	0.2416 (3)	1.4735 (5)	0.03432 (17)	0.0256 (6)
H16A	0.1663	1.3625	0.0456	0.038*
H16B	0.2582	1.4641	-0.0336	0.038*
H16C	0.2045	1.6220	0.0468	0.038*
C17	0.7852 (3)	0.4676 (4)	0.53576 (15)	0.0194 (5)
H17A	0.8962	0.4793	0.5454	0.029*
H17B	0.7564	0.3159	0.5503	0.029*
H17C	0.7455	0.5714	0.5796	0.029*
C18	0.9561 (2)	0.4317 (4)	0.24145 (15)	0.0155 (5)
C119	0.95353 (6)	0.14171 (10)	0.22820 (4)	0.02283 (16)
C120	1.14459 (6)	0.51691 (10)	0.28730 (4)	0.02415 (16)
Cl21	0.89393 (7)	0.55359 (12)	0.12613 (4)	0.03251 (19)

Atomic displacement parameters $(Å^2)$

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0208 (12)	0.0143 (12)	0.0113 (10)	0.0026 (11)	0.0052 (9)	-0.0013 (10)
N2	0.0157 (10)	0.0130 (10)	0.0135 (8)	0.0008 (8)	0.0040 (8)	0.0008 (8)
C3	0.0127 (11)	0.0159 (12)	0.0107 (10)	-0.0001 (10)	-0.0005 (8)	-0.0026 (10)
O4	0.0157 (8)	0.0130 (8)	0.0195 (8)	-0.0007 (7)	0.0071 (6)	0.0001 (7)
C5	0.0158 (11)	0.0172 (13)	0.0122 (10)	0.0006 (10)	0.0050 (8)	-0.0015 (11)
C6	0.0191 (12)	0.0100 (11)	0.0147 (10)	-0.0041 (10)	0.0063 (9)	-0.0026 (10)
07	0.0162 (8)	0.0161 (9)	0.0174 (8)	-0.0006 (7)	0.0061 (6)	0.0032 (7)
C8	0.0219 (12)	0.0125 (12)	0.0094 (10)	0.0023 (11)	0.0065 (9)	-0.0024 (10)
09	0.0228 (9)	0.0154 (9)	0.0174 (7)	-0.0033 (7)	0.0074 (7)	0.0001 (7)
C10	0.0195 (12)	0.0159 (12)	0.0143 (10)	-0.0009 (10)	0.0064 (9)	-0.0022 (10)
C11	0.0216 (11)	0.0175 (13)	0.0109 (10)	-0.0012 (10)	0.0041 (9)	0.0026 (10)
O12	0.0208 (9)	0.0239 (9)	0.0179 (8)	0.0014 (8)	0.0035 (7)	-0.0072 (8)
013	0.0294 (9)	0.0177 (9)	0.0135 (8)	-0.0011 (8)	0.0010 (7)	0.0023 (7)
C14	0.0231 (13)	0.0195 (12)	0.0164 (11)	0.0038 (11)	0.0071 (9)	0.0027 (11)
015	0.0219 (9)	0.0171 (8)	0.0194 (8)	0.0012 (7)	0.0038 (7)	0.0004 (8)
C16	0.0234 (13)	0.0262 (15)	0.0257 (12)	0.0023 (11)	-0.0005 (10)	-0.0014 (12)
C17	0.0190 (12)	0.0231 (14)	0.0168 (11)	0.0052 (10)	0.0048 (9)	0.0037 (11)
C18	0.0158 (12)	0.0159 (12)	0.0151 (11)	-0.0009 (10)	0.0032 (9)	-0.0003 (10)
Cl19	0.0254 (3)	0.0168 (3)	0.0290 (3)	-0.0034 (3)	0.0126 (3)	-0.0074 (3)
Cl20	0.0162 (3)	0.0259 (3)	0.0317 (3)	-0.0044 (3)	0.0079 (2)	-0.0094 (3)
Cl21	0.0345 (4)	0.0461 (5)	0.0202 (3)	0.0192 (3)	0.0139 (3)	0.0130 (3)
		. ,				

Geometric parameters (Å, °)

C1—N2	1.491 (3)	C11—H11A	0.9900
C1—C17	1.515 (3)	C11—H11B	0.9900
C1—C5	1.532 (3)	O12—H12	0.8400
C1—C8	1.533 (3)	O13—C14	1.396 (3)
N2—C3	1.260 (3)	C14—O15	1.403 (3)
C3—O4	1.352 (3)	C14—H14A	0.9900
C3—C18	1.519 (3)	C14—H14B	0.9900
O4—C5	1.459 (3)	O15—C16	1.424 (3)
C5—C6	1.542 (3)	C16—H16A	0.9800
С5—Н5	1.0000	C16—H16B	0.9800
C6—O7	1.468 (3)	C16—H16C	0.9800
C6—C10	1.515 (3)	C17—H17A	0.9800
С6—Н6	1.0000	C17—H17B	0.9800
O7—C8	1.337 (3)	C17—H17C	0.9800
C8—O9	1.202 (3)	C18—C119	1.757 (3)
C10—O12	1.414 (3)	C18—Cl21	1.763 (2)
C10-C11	1.524 (3)	C18—Cl20	1.773 (2)
C10—H10	1.0000	Cl19—O4 ⁱ	3.0697 (17)
C11—O13	1.437 (2)	Cl20—O7 ⁱⁱ	3.1419 (16)
			()
N2—C1—C17	109.67 (18)	O13—C11—H11A	109.3
N2—C1—C5	104.89 (16)	C10-C11-H11A	109.3
C17—C1—C5	117.1 (2)	O13—C11—H11B	109.3
N2—C1—C8	108.29 (17)	C10-C11-H11B	109.3
C17—C1—C8	112.22 (18)	H11A—C11—H11B	107.9
C5—C1—C8	104.10 (18)	C10—O12—H12	109.5
C3—N2—C1	104.83 (19)	C14—O13—C11	113.55 (17)
N2—C3—O4	120.6 (2)	O13—C14—O15	112.79 (17)
N2—C3—C18	125.5 (2)	O13—C14—H14A	109.0
O4—C3—C18	113.8 (2)	O15—C14—H14A	109.0
C3—O4—C5	105.01 (17)	O13—C14—H14B	109.0
O4—C5—C1	103.34 (17)	O15—C14—H14B	109.0
O4—C5—C6	114.30 (16)	H14A—C14—H14B	107.8
C1—C5—C6	106.11 (18)	C14—O15—C16	113.79 (18)
O4—C5—H5	110.9	O15—C16—H16A	109.5
C1—C5—H5	110.9	O15—C16—H16B	109.5
С6—С5—Н5	110.9	H16A—C16—H16B	109.5
O7—C6—C10	107.59 (17)	O15—C16—H16C	109.5
O7—C6—C5	105.81 (18)	H16A—C16—H16C	109.5
C10—C6—C5	116.62 (18)	H16B—C16—H16C	109.5
O7—C6—H6	108.9	C1—C17—H17A	109.5
С10—С6—Н6	108.9	C1—C17—H17B	109.5
С5—С6—Н6	108.9	H17A—C17—H17B	109.5
C8—O7—C6	112.65 (16)	C1—C17—H17C	109.5
O9—C8—O7	121.9 (2)	H17A—C17—H17C	109.5
O9—C8—C1	127.1 (2)	H17B—C17—H17C	109.5

O7—C8—C1	111.04 (19)	C3—C18—C119	110.75 (17)
O12—C10—C11	112.46 (18)	C3—C18—Cl21	110.46 (16)
O12—C10—C6	110.51 (19)	Cl19—C18—Cl21	108.87 (12)
C11—C10—C6	110.68 (17)	C3—C18—Cl20	108.21 (15)
O12—C10—H10	107.7	Cl19—C18—Cl20	108.69 (13)
C11—C10—H10	107.7	Cl21—C18—Cl20	109.84 (12)
C6—C10—H10	107.7	C18-C119-O4 ⁱ	139.97 (8)
O13—C11—C10	111.67 (17)	C18—Cl20—O7 ⁱⁱ	177.84 (8)
C17—C1—N2—C3	-118.3 (2)	C17—C1—C8—O9	-47.7 (3)
C5—C1—N2—C3	8.3 (2)	C5-C1-C8-09	-175.3 (2)
C8—C1—N2—C3	119.0 (2)	N2-C1-C8-O7	-105.6 (2)
C1—N2—C3—O4	-1.7 (3)	C17—C1—C8—O7	133.2 (2)
C1—N2—C3—C18	173.61 (19)	C5-C1-C8-O7	5.6 (2)
N2—C3—O4—C5	-6.0 (3)	O7—C6—C10—O12	-59.9 (2)
C18—C3—O4—C5	178.18 (16)	C5-C6-C10-O12	58.7 (3)
C3—O4—C5—C1	10.3 (2)	O7—C6—C10—C11	65.4 (2)
C3—O4—C5—C6	-104.5 (2)	C5-C6-C10-C11	-176.0 (2)
N2-C1-C5-O4	-11.3 (2)	O12—C10—C11—O13	-52.1 (3)
C17—C1—C5—O4	110.49 (19)	C6-C10-C11-O13	-176.20 (19)
C8—C1—C5—O4	-125.02 (17)	C10-C11-O13-C14	-79.6 (2)
N2-C1-C5-C6	109.24 (18)	C11—O13—C14—O15	-74.1 (2)
C17—C1—C5—C6	-128.9 (2)	O13-C14-O15-C16	-74.3 (2)
C8—C1—C5—C6	-4.4 (2)	N2-C3-C18-C119	4.6 (3)
O4—C5—C6—O7	115.31 (19)	O4—C3—C18—C119	-179.78 (15)
C1C5C6O7	2.1 (2)	N2-C3-C18-Cl21	125.3 (2)
O4C5C10	-4.2 (3)	O4—C3—C18—Cl21	-59.1 (2)
C1C5C10	-117.4 (2)	N2-C3-C18-Cl20	-114.4 (2)
C10—C6—O7—C8	126.83 (19)	O4—C3—C18—Cl20	61.2 (2)
C5—C6—O7—C8	1.5 (2)	C3-C18-C119-O4 ⁱ	-12.9 (2)
C6—O7—C8—O9	176.28 (19)	Cl21—C18—Cl19—O4 ⁱ	-134.58 (9)
C6—O7—C8—C1	-4.6 (2)	Cl20—C18—Cl19—O4 ⁱ	105.81 (13)
N2-C1-C8-09	73.5 (3)		

Symmetry codes: (i) *x*, *y*–1, *z*; (ii) *x*+1, *y*, *z*.

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H··· A
012—H12…O15 ⁱ	0.84	1.90	2.695 (2)	157
C6—H6····O9 ⁱⁱⁱ	1.00	2.43	3.402 (3)	164
C17—H17 <i>C</i> ···O9 ^{iv}	0.98	2.53	3.320 (3)	137

Symmetry codes: (i) *x*, *y*-1, *z*; (iii) *x*, *y*+1, *z*; (iv) -*x*+1, *y*+1/2, -*z*+1.