

## Crystal structure of [1,1':3',1''-terphenyl]-2',3,3''-tricarboxylic acid

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The asymmetric unit of the title compound,  $C_{21}H_{14}O_6$ , comprises two symmetrically independent molecules that form a locally centrosymmetric hydrogen-bonded dimer, with the planes of the corresponding carboxylic acid groups rotated by  $15.8(1)$  and  $17.5(1)^\circ$  relative to those of the adjacent benzene rings. The crystal as a whole, however, exhibits a noncentrosymmetric packing, described by the polar space group  $Pca2_1$ . The dimers form layers along the  $ab$  plane, being interconnected by hydrogen bonds involving the remaining carboxylic acid groups. The plane of the central carboxylic acid group forms dihedral angles of  $62.5(1)$  and  $63.0(1)^\circ$  with those of the adjacent benzene rings and functions as a hydrogen-bond donor and acceptor. As a donor, it interconnects adjacent layers, while as an acceptor it stabilizes the packing within the layers. The ‘distal’ carboxylic acid groups are nearly coplanar with the planes of the adjacent benzene rings, forming dihedral angles of  $1.8(1)$  and  $7.1(1)^\circ$ . These groups also form intra- and inter-layer hydrogen bonds, but with ‘reversed’ functionality, as compared with the central carboxylic acid groups.

**Keywords:** crystal structure; hydrogen bonding; meta-terphenyl.

**CCDC reference:** 1418223

### 1. Related literature

For a detailed discussion on local centers of symmetry in the space group  $Pca2_1$ , see: Marsh *et al.* (1998). For the synthesis of the starting material 3,3''-dimethyl-[1,1':3',1''-terphenyl]-2'-carboxylic acid, see: Du *et al.* (1986).

### 2. Experimental

#### 2.1. Crystal data

$C_{21}H_{14}O_6$   
 $M_r = 362.32$   
Orthorhombic,  $Pca2_1$   
 $a = 23.1735(9)\text{ \AA}$   
 $b = 7.2480(2)\text{ \AA}$   
 $c = 20.3320(8)\text{ \AA}$

$V = 3415.0(2)\text{ \AA}^3$   
 $Z = 8$   
Mo  $K\alpha$  radiation  
 $\mu = 0.10\text{ mm}^{-1}$   
 $T = 100\text{ K}$   
 $0.3 \times 0.05 \times 0.05\text{ mm}$

#### 2.2. Data collection

Bruker D8 VENTURE DUO diffractometer  
Absorption correction: multi-scan (*SADABS*; Bruker, 2012)  
 $T_{\min} = 0.695$ ,  $T_{\max} = 0.745$

46489 measured reflections  
6467 independent reflections  
5899 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.028$

#### 2.3. Refinement

$R[F^2 > 2\sigma(F^2)] = 0.037$   
 $wR(F^2) = 0.091$   
 $S = 1.05$   
6467 reflections  
511 parameters  
2 restraints

H atoms treated by a mixture of independent and constrained refinement  
 $\Delta\rho_{\max} = 0.27\text{ e \AA}^{-3}$   
 $\Delta\rho_{\min} = -0.17\text{ e \AA}^{-3}$

**Table 1**  
Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ ).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$O2-\text{H}2\cdots O3^{\text{i}}$	0.91 (5)	1.79 (5)	2.657 (3)	159 (4)
$O4-\text{H}4\cdots O1^{\text{ii}}$	0.90 (6)	1.84 (6)	2.689 (3)	156 (5)
$O2'\cdots H2'\cdots O3^{\text{iii}}$	0.87 (6)	1.76 (6)	2.628 (3)	171 (5)
$O4'\cdots H4'\cdots O1^{\text{iv}}$	0.84 (5)	1.90 (5)	2.717 (3)	166 (4)
$O6'\cdots H6'\cdots O5$	0.92 (4)	1.69 (4)	2.593 (3)	168 (4)
$O6\cdots H6\cdots O5'$	1.01 (3)	1.58 (3)	2.581 (3)	172 (6)

Symmetry codes: (i)  $-x + 1, -y + 2, z - \frac{1}{2}$ ; (ii)  $x, y - 1, z$ ; (iii)  $-x + \frac{3}{2}, y + 1, z + \frac{1}{2}$ ; (iv)  $x, y + 1, z$ .

Data collection: *APEX2* (Bruker, 2012); cell refinement: *SAINT* (Bruker, 2012); data reduction: *SAINT*; program(s) used to solve structure: *SHELXT* (Sheldrick, 2015a); program(s) used to refine structure: *SHELXL2014* (Sheldrick, 2015b); molecular graphics: *OLEX2* (Dolomanov *et al.*, 2009); software used to prepare material for publication: *OLEX2*.

### Acknowledgements

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## data reports

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Supporting information for this paper is available from the IUCr electronic archives (Reference: LD2133).

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Du, C. J. F., Hart, H. & Ng, K. K. D. (1986). *J. Org. Chem.* **51**, 3162–3165.  
Marsh, R. E., Schomaker, V. & Herbstein, F. H. (1998). *Acta Cryst.* **B54**, 921–924.  
Sheldrick, G. M. (2015a). *Acta Cryst. A* **71**, 3–8.  
Sheldrick, G. M. (2015b). *Acta Cryst. C* **71**, 3–8.

# supporting information

*Acta Cryst.* (2015). E71, o667–o668 [https://doi.org/10.1107/S2056989015015029]

## Crystal structure of [1,1':3',1''-terphenyl]-2',3,3''-tricarboxylic acid

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### S1. Experimental

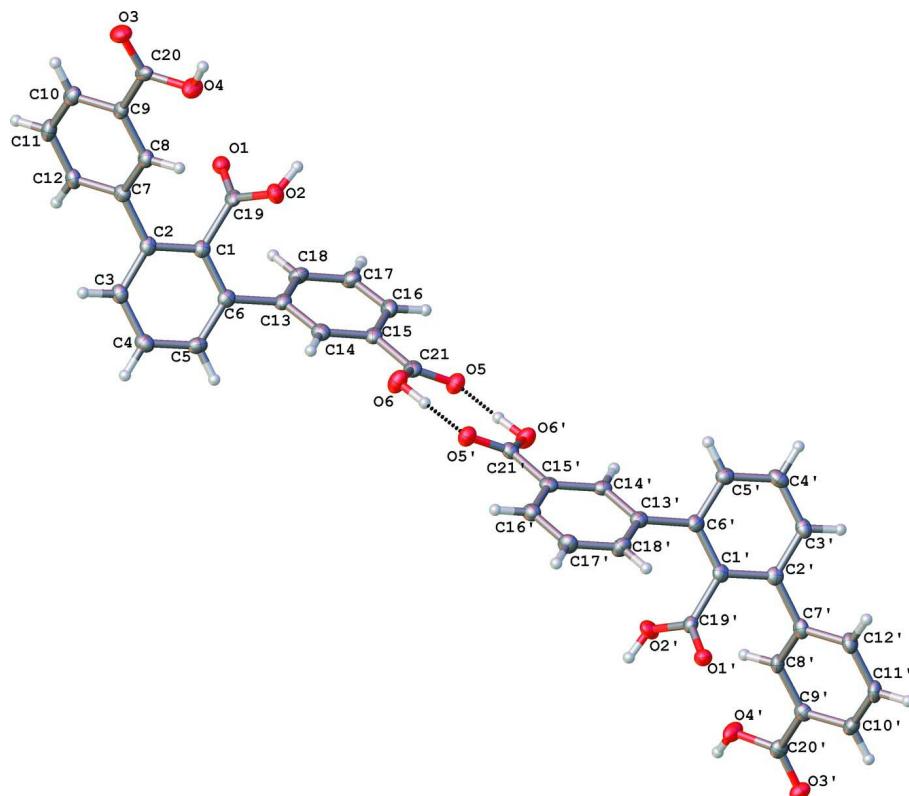
#### S1.1. Synthesis and crystallization

3,3''-dimethyl-[1,1':3',1''-terphenyl]-2'-carboxylic acid (0.432 g, 1.4 mmol) synthesized according to Du *et al.* (1986). The starting material was dissolved in 15 ml pyridine and brought to reflux. KMnO<sub>4</sub> (0.50 g, 3.1 mmol) in 1 ml of water was added, and allowed to react for 2 hours. Subsequently, 3 more additions of KMnO<sub>4</sub> (0.25 g, 1.6 mmol) in 1.5 ml of water were added every 2 hours for a total of 4 additions. After six hours, 10 ml of water was added and the reaction was refluxed overnight. The reaction mixture was filtered while hot to remove solid MnO<sub>2</sub>. The filtrate was then concentrated under reduced pressure, and treated by 12M HCl. The resulting white precipitate was then filtered and purified via silica gel column chromatograph (50/50 hexanes/ethyl acetate with 0.5% acetic acid) to give of the title compound. (0.366 g Yield 70%) <sup>1</sup>H NMR (DMSO, 400 MHz):  $\delta$  7.46 (d, 2H, *J* = 8 Hz), 7.58 (t, 2H, *J* = 8 Hz), 7.61 (t, 1H, *J* = 7.2 Hz), 7.68 (d, 2H, *J* = 8 Hz), 7.97 (d, 2H, *J* = 8 Hz), 8.01 (s, 2H), 13.01 (s, O-H). <sup>13</sup>C NMR (DMSO, 100 MHz):  $\delta$  128.4, 128.7, 129.1, 129.2, 129.3, 130.8, 132.7, 133.9, 138.0, 140.39, 167.0, 169.7. Single crystals suitable for X-ray diffraction were obtained by vapor diffusion of hexane into an ethyl acetate solution of the title compound.

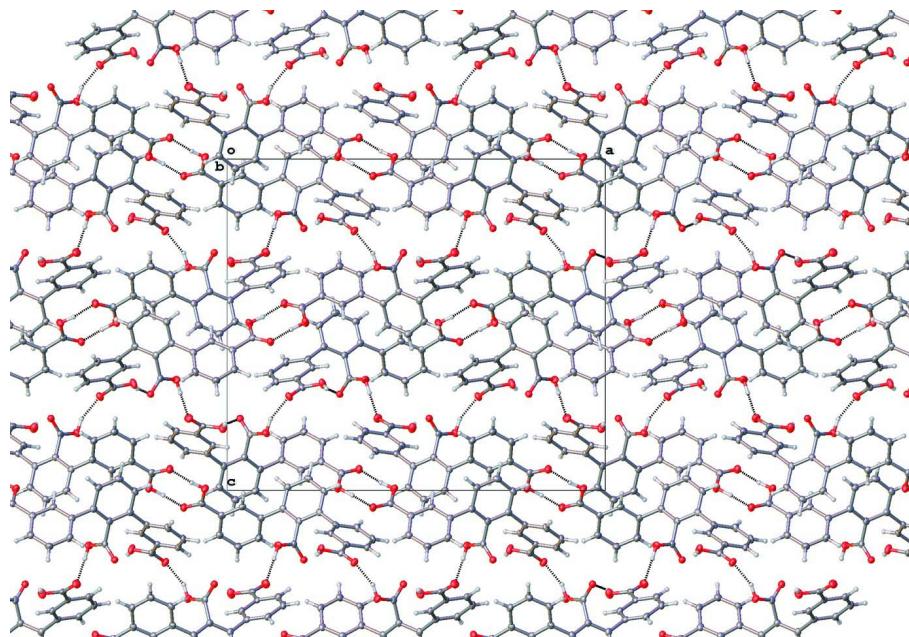
#### S1.2. Refinement

All H atoms were located in difference Fourier maps but finally their positions were determined geometrically, except for the carboxy H atoms that were refined with isotropic thermal parameters. The O6—H6 bond length in the carboxylic acid dimer was restrained at the distance from the corresponding residual electron density peak to the oxygen (0.996 (2) Å). This was done due to unreasonable lengthening (> 1.4 Å) of the H—O bond during the refinement. All other H atoms were refined using a riding model with fixed isotropic displacement parameters [Uiso(H) = 1.2Ueq(C) for the C(H) groups]. Additional crystal data, data collection and structure refinement details are summarized in Table 1.

## S2. Results and discussion

**Figure 1**

The asymmetric unit of the title compound, with displacement ellipsoids drawn at 50% probability level. Hydrogen atoms presented by spheres of an arbitrary radius. Intra-dimer hydrogen bonds are represented by dotted lines.



**Figure 2**

Packing view along *b* axis. Hydrogen bonds are represented by dotted lines.

**[1,1':3',1''-Terphenyl]-2',3,3''-tricarboxylic acid***Crystal data*

$C_{21}H_{14}O_6$   
 $M_r = 362.32$   
Orthorhombic,  $Pca2_1$   
 $a = 23.1735$  (9) Å  
 $b = 7.2480$  (2) Å  
 $c = 20.3320$  (8) Å  
 $V = 3415.0$  (2) Å<sup>3</sup>  
 $Z = 8$   
 $F(000) = 1504$

$D_x = 1.409$  Mg m<sup>-3</sup>  
Mo  $K\alpha$  radiation,  $\lambda = 0.71073$  Å  
Cell parameters from 9141 reflections  
 $\theta = 3.1\text{--}25.7^\circ$   
 $\mu = 0.10$  mm<sup>-1</sup>  
 $T = 100$  K  
Needle, clear  
 $0.3 \times 0.05 \times 0.05$  mm

*Data collection*

Bruker D8 VENTURE DUO  
diffractometer  
Radiation source: sealed tube, fine-focus  
TRIUMPH graphite monochromator  
Detector resolution: 10.5 pixels mm<sup>-1</sup>  
 $\omega$  and  $\varphi$  scans  
Absorption correction: multi-scan  
(SADABS; Bruker, 2012)  
 $T_{\min} = 0.695$ ,  $T_{\max} = 0.745$

46489 measured reflections  
6467 independent reflections  
5899 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.028$   
 $\theta_{\max} = 25.7^\circ$ ,  $\theta_{\min} = 3.0^\circ$   
 $h = -28 \rightarrow 28$   
 $k = -7 \rightarrow 8$   
 $l = -24 \rightarrow 24$

*Refinement*

Refinement on  $F^2$   
Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.037$   
 $wR(F^2) = 0.091$   
 $S = 1.05$   
6467 reflections  
511 parameters  
2 restraints

Primary atom site location: structure-invariant  
direct methods  
Hydrogen site location: mixed  
H atoms treated by a mixture of independent  
and constrained refinement  
 $w = 1/[\sigma^2(F_o^2) + (0.0518P)^2 + 1.0855P]$   
where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.27$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.17$  e Å<sup>-3</sup>

*Special details*

**Experimental.** SADABS-2012/1 (Bruker, 2012) was used for absorption correction.  $wR2(\text{int})$  was 0.0509 before and 0.0455 after correction. The Ratio of minimum to maximum transmission is 0.9326. The  $\lambda/2$  correction factor is 0.0015.

**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.

*Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å<sup>2</sup>)*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.96578 (9)	0.7615 (3)	0.28358 (11)	0.0228 (5)
O2	0.89715 (9)	0.5814 (3)	0.32756 (11)	0.0244 (5)
H2	0.8852 (17)	0.572 (5)	0.285 (2)	0.042 (11)*

O3	1.10167 (9)	0.0388 (3)	0.28278 (11)	0.0303 (5)
O4	1.01069 (10)	0.0942 (3)	0.31188 (12)	0.0306 (5)
H4	1.006 (2)	-0.025 (8)	0.301 (3)	0.085 (18)*
O5	0.65152 (9)	0.9096 (3)	0.43665 (11)	0.0258 (5)
O6	0.70682 (9)	0.7133 (3)	0.49305 (12)	0.0285 (5)
C1	0.95397 (12)	0.7560 (4)	0.40061 (15)	0.0188 (6)
C2	1.00836 (13)	0.7148 (4)	0.42736 (15)	0.0200 (6)
C3	1.02024 (14)	0.7721 (4)	0.49135 (16)	0.0214 (7)
H3	1.0572	0.7495	0.5098	0.026*
C4	0.97881 (14)	0.8613 (4)	0.52817 (16)	0.0235 (7)
H4A	0.9872	0.8968	0.5721	0.028*
C5	0.92531 (13)	0.8995 (4)	0.50187 (15)	0.0228 (7)
H5	0.8969	0.9586	0.5282	0.027*
C6	0.91246 (13)	0.8522 (4)	0.43706 (15)	0.0197 (6)
C7	1.05206 (13)	0.6053 (4)	0.38977 (15)	0.0188 (6)
C8	1.03776 (13)	0.4318 (4)	0.36459 (15)	0.0201 (7)
H8	0.9999	0.3841	0.3704	0.024*
C9	1.07890 (13)	0.3293 (4)	0.33113 (14)	0.0178 (6)
C10	1.13447 (14)	0.3978 (4)	0.32208 (16)	0.0239 (7)
H10	1.1625	0.3272	0.2991	0.029*
C11	1.14841 (13)	0.5694 (4)	0.34687 (16)	0.0251 (7)
H11	1.1861	0.6178	0.3404	0.030*
C12	1.10790 (13)	0.6712 (4)	0.38102 (15)	0.0221 (6)
H12	1.1183	0.7878	0.3988	0.026*
C13	0.85570 (13)	0.9045 (4)	0.40774 (15)	0.0203 (6)
C14	0.80514 (13)	0.8518 (4)	0.43914 (15)	0.0212 (7)
H14	0.8072	0.7839	0.4790	0.025*
C15	0.75168 (13)	0.8969 (4)	0.41302 (16)	0.0205 (6)
C16	0.74787 (13)	0.9963 (4)	0.35433 (16)	0.0226 (7)
H16	0.7113	1.0245	0.3357	0.027*
C17	0.79797 (13)	1.0528 (4)	0.32379 (16)	0.0243 (7)
H17	0.7959	1.1237	0.2846	0.029*
C18	0.85140 (13)	1.0069 (4)	0.34989 (16)	0.0233 (7)
H18	0.8855	1.0459	0.3280	0.028*
C19	0.94049 (13)	0.7010 (4)	0.33133 (15)	0.0199 (6)
C20	1.06555 (13)	0.1399 (4)	0.30611 (14)	0.0209 (6)
C21	0.69880 (13)	0.8403 (4)	0.44841 (16)	0.0220 (6)
O1'	0.29732 (9)	0.7389 (3)	0.70997 (10)	0.0213 (5)
O2'	0.36530 (10)	0.9209 (3)	0.66648 (12)	0.0265 (5)
H2'	0.376 (2)	0.949 (7)	0.706 (3)	0.070 (15)*
O3'	0.16368 (9)	1.4521 (3)	0.71744 (11)	0.0247 (5)
O4'	0.25240 (10)	1.4042 (3)	0.67837 (12)	0.0282 (5)
H4'	0.2621 (18)	1.507 (7)	0.693 (2)	0.054 (13)*
O5'	0.61435 (9)	0.6032 (3)	0.55199 (11)	0.0247 (5)
O6'	0.55852 (10)	0.7980 (3)	0.49534 (11)	0.0271 (5)
C1'	0.31076 (12)	0.7424 (4)	0.59342 (15)	0.0191 (6)
C2'	0.25613 (13)	0.7813 (4)	0.56681 (15)	0.0193 (6)
C3'	0.24393 (13)	0.7237 (4)	0.50299 (16)	0.0216 (7)

H3'	0.2067	0.7449	0.4850	0.026*
C4'	0.28570 (14)	0.6353 (4)	0.46537 (16)	0.0245 (7)
H4'A	0.2771	0.5986	0.4216	0.029*
C5'	0.33996 (13)	0.6003 (4)	0.49149 (16)	0.0214 (6)
H5'	0.3685	0.5418	0.4652	0.026*
C6'	0.35278 (13)	0.6504 (4)	0.55605 (15)	0.0198 (6)
C7'	0.21240 (13)	0.8890 (4)	0.60456 (16)	0.0209 (7)
C8'	0.22618 (12)	1.0626 (4)	0.63010 (14)	0.0183 (6)
H8'	0.2640	1.1105	0.6247	0.022*
C9'	0.18463 (13)	1.1658 (4)	0.66341 (15)	0.0198 (6)
C10'	0.12932 (13)	1.0961 (4)	0.67205 (15)	0.0212 (6)
H10'	0.1011	1.1663	0.6949	0.025*
C11'	0.11546 (13)	0.9241 (4)	0.64717 (17)	0.0242 (7)
H11'	0.0777	0.8760	0.6532	0.029*
C12'	0.15643 (13)	0.8211 (4)	0.61342 (16)	0.0228 (7)
H12'	0.1464	0.7036	0.5962	0.027*
C13'	0.41042 (13)	0.6027 (4)	0.58405 (15)	0.0198 (6)
C14'	0.46007 (13)	0.6550 (4)	0.55043 (15)	0.0192 (6)
H14'	0.4569	0.7180	0.5096	0.023*
C15'	0.51443 (13)	0.6154 (4)	0.57626 (15)	0.0199 (6)
C16'	0.51981 (13)	0.5223 (4)	0.63543 (15)	0.0231 (7)
H16'	0.5569	0.4950	0.6528	0.028*
C17'	0.47048 (13)	0.4694 (4)	0.66912 (16)	0.0252 (7)
H17'	0.4738	0.4067	0.7100	0.030*
C18'	0.41620 (13)	0.5075 (4)	0.64352 (16)	0.0252 (7)
H18'	0.3827	0.4685	0.6666	0.030*
C19'	0.32344 (12)	0.7990 (4)	0.66247 (15)	0.0183 (6)
C20'	0.19858 (13)	1.3529 (4)	0.68902 (15)	0.0210 (6)
C21'	0.56704 (13)	0.6716 (4)	0.53989 (14)	0.0197 (6)
H6'	0.5939 (17)	0.824 (6)	0.477 (2)	0.045 (12)*
H6	0.6693 (16)	0.681 (8)	0.515 (3)	0.098 (19)*

Atomic displacement parameters ( $\text{\AA}^2$ )

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
O1	0.0208 (11)	0.0242 (11)	0.0235 (11)	0.0006 (8)	-0.0009 (9)	-0.0026 (9)
O2	0.0228 (12)	0.0250 (11)	0.0255 (13)	-0.0031 (9)	-0.0048 (10)	-0.0020 (9)
O3	0.0304 (12)	0.0302 (12)	0.0304 (13)	0.0075 (10)	0.0024 (10)	-0.0070 (10)
O4	0.0275 (13)	0.0232 (12)	0.0411 (14)	-0.0028 (10)	0.0041 (10)	-0.0084 (10)
O5	0.0209 (11)	0.0256 (11)	0.0309 (12)	0.0030 (9)	0.0016 (9)	0.0029 (10)
O6	0.0221 (12)	0.0292 (12)	0.0341 (13)	0.0023 (10)	0.0037 (11)	0.0100 (11)
C1	0.0199 (15)	0.0150 (14)	0.0215 (15)	-0.0009 (11)	-0.0003 (12)	0.0006 (11)
C2	0.0192 (15)	0.0149 (13)	0.0259 (16)	-0.0022 (11)	-0.0009 (12)	0.0019 (12)
C3	0.0224 (16)	0.0195 (15)	0.0225 (17)	-0.0004 (12)	-0.0040 (14)	0.0022 (13)
C4	0.0287 (18)	0.0191 (15)	0.0228 (16)	0.0006 (13)	-0.0033 (13)	-0.0010 (12)
C5	0.0252 (17)	0.0213 (15)	0.0218 (16)	-0.0007 (12)	0.0047 (13)	-0.0016 (13)
C6	0.0207 (15)	0.0149 (14)	0.0236 (15)	-0.0017 (12)	0.0019 (13)	0.0017 (12)
C7	0.0217 (16)	0.0163 (14)	0.0183 (15)	0.0021 (12)	-0.0029 (12)	0.0022 (12)

C8	0.0181 (15)	0.0199 (15)	0.0222 (16)	0.0020 (12)	-0.0025 (12)	0.0011 (12)
C9	0.0166 (15)	0.0189 (14)	0.0178 (14)	0.0028 (11)	-0.0020 (12)	0.0042 (11)
C10	0.0211 (17)	0.0269 (16)	0.0235 (16)	0.0045 (13)	-0.0009 (13)	0.0041 (13)
C11	0.0166 (16)	0.0287 (17)	0.0299 (18)	-0.0010 (12)	-0.0020 (13)	0.0061 (14)
C12	0.0203 (16)	0.0222 (15)	0.0237 (16)	-0.0014 (12)	-0.0056 (13)	0.0025 (12)
C13	0.0201 (15)	0.0173 (14)	0.0234 (16)	0.0029 (12)	0.0012 (13)	-0.0033 (12)
C14	0.0255 (17)	0.0171 (14)	0.0211 (16)	0.0024 (12)	-0.0009 (13)	-0.0009 (12)
C15	0.0206 (15)	0.0158 (13)	0.0250 (16)	0.0026 (12)	-0.0001 (12)	-0.0050 (12)
C16	0.0206 (15)	0.0211 (15)	0.0262 (17)	0.0044 (13)	-0.0033 (13)	-0.0026 (13)
C17	0.0252 (17)	0.0252 (15)	0.0225 (17)	0.0048 (13)	0.0026 (13)	-0.0007 (13)
C18	0.0238 (16)	0.0227 (15)	0.0234 (17)	0.0021 (14)	0.0065 (12)	0.0007 (13)
C19	0.0173 (15)	0.0159 (14)	0.0264 (16)	0.0051 (12)	-0.0016 (13)	-0.0044 (12)
C20	0.0226 (17)	0.0248 (15)	0.0153 (14)	0.0081 (13)	-0.0009 (12)	0.0002 (12)
C21	0.0227 (17)	0.0170 (14)	0.0261 (16)	0.0037 (12)	-0.0006 (13)	-0.0049 (12)
O1'	0.0212 (11)	0.0221 (11)	0.0207 (11)	-0.0001 (8)	0.0011 (9)	-0.0013 (9)
O2'	0.0268 (12)	0.0283 (11)	0.0244 (12)	-0.0108 (9)	-0.0009 (10)	-0.0065 (10)
O3'	0.0237 (11)	0.0205 (11)	0.0301 (12)	0.0016 (9)	0.0070 (10)	-0.0003 (9)
O4'	0.0217 (11)	0.0201 (11)	0.0428 (15)	-0.0052 (9)	0.0083 (10)	-0.0080 (11)
O5'	0.0205 (12)	0.0258 (11)	0.0277 (12)	0.0002 (9)	0.0022 (9)	0.0023 (10)
O6'	0.0232 (12)	0.0300 (12)	0.0280 (12)	0.0029 (9)	0.0047 (10)	0.0076 (10)
C1'	0.0191 (15)	0.0141 (13)	0.0241 (16)	-0.0039 (11)	-0.0004 (13)	0.0006 (12)
C2'	0.0209 (16)	0.0136 (13)	0.0233 (16)	-0.0028 (11)	-0.0002 (12)	0.0020 (12)
C3'	0.0221 (16)	0.0162 (14)	0.0264 (17)	-0.0011 (12)	-0.0064 (13)	0.0002 (12)
C4'	0.0330 (18)	0.0204 (16)	0.0201 (15)	-0.0046 (13)	-0.0039 (13)	-0.0020 (12)
C5'	0.0235 (17)	0.0177 (14)	0.0229 (15)	-0.0012 (12)	0.0020 (13)	-0.0020 (12)
C6'	0.0204 (16)	0.0157 (13)	0.0234 (15)	-0.0037 (11)	0.0000 (12)	-0.0017 (12)
C7'	0.0201 (16)	0.0197 (14)	0.0227 (16)	-0.0001 (12)	-0.0060 (13)	0.0049 (12)
C8'	0.0140 (15)	0.0198 (15)	0.0212 (16)	-0.0025 (11)	-0.0027 (12)	0.0031 (12)
C9'	0.0228 (15)	0.0182 (14)	0.0185 (14)	-0.0001 (12)	-0.0024 (13)	0.0029 (12)
C10'	0.0171 (15)	0.0237 (15)	0.0227 (15)	0.0025 (12)	0.0018 (13)	0.0025 (12)
C11'	0.0151 (16)	0.0265 (16)	0.0309 (18)	-0.0040 (12)	-0.0045 (13)	0.0029 (14)
C12'	0.0203 (16)	0.0181 (14)	0.0301 (17)	-0.0029 (12)	-0.0057 (13)	-0.0005 (12)
C13'	0.0217 (15)	0.0152 (13)	0.0226 (16)	-0.0012 (12)	0.0011 (12)	-0.0024 (12)
C14'	0.0240 (16)	0.0155 (13)	0.0180 (14)	0.0003 (11)	0.0029 (12)	-0.0023 (12)
C15'	0.0245 (16)	0.0146 (14)	0.0205 (15)	0.0011 (12)	0.0003 (12)	-0.0041 (12)
C16'	0.0241 (16)	0.0230 (15)	0.0221 (16)	0.0028 (13)	-0.0014 (12)	-0.0017 (12)
C17'	0.0275 (17)	0.0275 (16)	0.0207 (16)	0.0011 (13)	-0.0002 (13)	0.0040 (14)
C18'	0.0226 (16)	0.0256 (16)	0.0272 (18)	0.0000 (14)	0.0057 (13)	0.0026 (14)
C19'	0.0151 (14)	0.0174 (13)	0.0223 (15)	0.0017 (11)	-0.0002 (13)	-0.0004 (12)
C20'	0.0211 (16)	0.0202 (14)	0.0217 (15)	-0.0021 (13)	0.0026 (13)	0.0052 (12)
C21'	0.0224 (16)	0.0167 (13)	0.0198 (15)	-0.0017 (12)	-0.0019 (12)	-0.0023 (12)

Geometric parameters ( $\text{\AA}$ ,  $^\circ$ )

O1—C19	1.216 (4)	O1'—C19'	1.220 (4)
O2—H2	0.91 (5)	O2'—H2'	0.87 (6)
O2—C19	1.329 (4)	O2'—C19'	1.314 (4)
O3—C20	1.209 (4)	O3'—C20'	1.227 (4)

O4—H4	0.90 (6)	O4'—H4'	0.84 (5)
O4—C20	1.319 (4)	O4'—C20'	1.319 (4)
O5—C21	1.229 (4)	O5'—C21'	1.228 (4)
O6—C21	1.306 (4)	O6'—C21'	1.303 (4)
O6—H6	1.01 (3)	O6'—H6'	0.92 (4)
C1—C2	1.405 (4)	C1'—C2'	1.405 (4)
C1—C6	1.400 (4)	C1'—C6'	1.404 (4)
C1—C19	1.497 (4)	C1'—C19'	1.492 (4)
C2—C3	1.393 (5)	C2'—C3'	1.392 (4)
C2—C7	1.496 (4)	C2'—C7'	1.491 (4)
C3—H3	0.9500	C3'—H3'	0.9500
C3—C4	1.378 (4)	C3'—C4'	1.390 (4)
C4—H4A	0.9500	C4'—H4'A	0.9500
C4—C5	1.378 (4)	C4'—C5'	1.388 (4)
C5—H5	0.9500	C5'—H5'	0.9500
C5—C6	1.394 (5)	C5'—C6'	1.394 (4)
C6—C13	1.493 (4)	C6'—C13'	1.493 (4)
C7—C8	1.398 (4)	C7'—C8'	1.398 (4)
C7—C12	1.391 (4)	C7'—C12'	1.399 (4)
C8—H8	0.9500	C8'—H8'	0.9500
C8—C9	1.387 (4)	C8'—C9'	1.395 (4)
C9—C10	1.393 (4)	C9'—C10'	1.389 (4)
C9—C20	1.496 (4)	C9'—C20'	1.488 (4)
C10—H10	0.9500	C10'—H10'	0.9500
C10—C11	1.380 (5)	C10'—C11'	1.384 (4)
C11—H11	0.9500	C11'—H11'	0.9500
C11—C12	1.381 (5)	C11'—C12'	1.389 (5)
C12—H12	0.9500	C12'—H12'	0.9500
C13—C14	1.388 (4)	C13'—C14'	1.391 (4)
C13—C18	1.394 (5)	C13'—C18'	1.399 (4)
C14—H14	0.9500	C14'—H14'	0.9500
C14—C15	1.387 (4)	C14'—C15'	1.394 (4)
C15—C16	1.397 (5)	C15'—C16'	1.385 (4)
C15—C21	1.479 (5)	C15'—C21'	1.483 (4)
C16—H16	0.9500	C16'—H16'	0.9500
C16—C17	1.379 (5)	C16'—C17'	1.387 (4)
C17—H17	0.9500	C17'—H17'	0.9500
C17—C18	1.387 (4)	C17'—C18'	1.389 (4)
C18—H18	0.9500	C18'—H18'	0.9500
C19—O2—H2	110 (3)	C19'—O2'—H2'	115 (3)
C20—O4—H4	109 (3)	C20'—O4'—H4'	117 (3)
C21—O6—H6	111 (3)	C21'—O6'—H6'	107 (3)
C2—C1—C19	119.7 (3)	C2'—C1'—C19'	119.0 (3)
C6—C1—C2	121.1 (3)	C6'—C1'—C2'	120.8 (3)
C6—C1—C19	119.2 (3)	C6'—C1'—C19'	120.2 (3)
C1—C2—C7	121.5 (3)	C1'—C2'—C7'	121.3 (3)
C3—C2—C1	118.4 (3)	C3'—C2'—C1'	118.8 (3)

C3—C2—C7	120.1 (3)	C3'—C2'—C7'	119.9 (3)
C2—C3—H3	119.7	C2'—C3'—H3'	119.7
C4—C3—C2	120.6 (3)	C4'—C3'—C2'	120.7 (3)
C4—C3—H3	119.7	C4'—C3'—H3'	119.7
C3—C4—H4A	119.7	C3'—C4'—H4'A	119.9
C5—C4—C3	120.7 (3)	C5'—C4'—C3'	120.3 (3)
C5—C4—H4A	119.7	C5'—C4'—H4'A	119.9
C4—C5—H5	119.7	C4'—C5'—H5'	119.8
C4—C5—C6	120.6 (3)	C4'—C5'—C6'	120.4 (3)
C6—C5—H5	119.7	C6'—C5'—H5'	119.8
C1—C6—C13	121.3 (3)	C1'—C6'—C13'	121.6 (3)
C5—C6—C1	118.4 (3)	C5'—C6'—C1'	119.1 (3)
C5—C6—C13	120.2 (3)	C5'—C6'—C13'	119.3 (3)
C8—C7—C2	120.2 (3)	C8'—C7'—C2'	120.5 (3)
C12—C7—C2	120.9 (3)	C8'—C7'—C12'	118.7 (3)
C12—C7—C8	118.9 (3)	C12'—C7'—C2'	120.8 (3)
C7—C8—H8	120.0	C7'—C8'—H8'	119.8
C9—C8—C7	119.9 (3)	C9'—C8'—C7'	120.4 (3)
C9—C8—H8	120.0	C9'—C8'—H8'	119.8
C8—C9—C10	120.6 (3)	C8'—C9'—C20'	120.6 (3)
C8—C9—C20	121.1 (3)	C10'—C9'—C8'	120.2 (3)
C10—C9—C20	118.3 (3)	C10'—C9'—C20'	119.2 (3)
C9—C10—H10	120.3	C9'—C10'—H10'	120.1
C11—C10—C9	119.3 (3)	C11'—C10'—C9'	119.7 (3)
C11—C10—H10	120.3	C11'—C10'—H10'	120.1
C10—C11—H11	119.8	C10'—C11'—H11'	119.8
C10—C11—C12	120.4 (3)	C10'—C11'—C12'	120.4 (3)
C12—C11—H11	119.8	C12'—C11'—H11'	119.8
C7—C12—H12	119.5	C7'—C12'—H12'	119.7
C11—C12—C7	120.9 (3)	C11'—C12'—C7'	120.6 (3)
C11—C12—H12	119.5	C11'—C12'—H12'	119.7
C14—C13—C6	119.3 (3)	C14'—C13'—C6'	119.3 (3)
C14—C13—C18	118.3 (3)	C14'—C13'—C18'	118.7 (3)
C18—C13—C6	122.3 (3)	C18'—C13'—C6'	122.0 (3)
C13—C14—H14	119.6	C13'—C14'—H14'	119.8
C15—C14—C13	120.9 (3)	C13'—C14'—C15'	120.4 (3)
C15—C14—H14	119.6	C15'—C14'—H14'	119.8
C14—C15—C16	120.3 (3)	C14'—C15'—C21'	119.9 (3)
C14—C15—C21	119.2 (3)	C16'—C15'—C14'	120.6 (3)
C16—C15—C21	120.4 (3)	C16'—C15'—C21'	119.5 (3)
C15—C16—H16	120.5	C15'—C16'—H16'	120.3
C17—C16—C15	119.0 (3)	C15'—C16'—C17'	119.3 (3)
C17—C16—H16	120.5	C17'—C16'—H16'	120.3
C16—C17—H17	119.7	C16'—C17'—H17'	119.8
C16—C17—C18	120.5 (3)	C16'—C17'—C18'	120.4 (3)
C18—C17—H17	119.7	C18'—C17'—H17'	119.8
C13—C18—H18	119.5	C13'—C18'—H18'	119.7
C17—C18—C13	120.9 (3)	C17'—C18'—C13'	120.6 (3)

C17—C18—H18	119.5	C17'—C18'—H18'	119.7
O1—C19—O2	123.6 (3)	O1'—C19'—O2'	123.8 (3)
O1—C19—C1	123.7 (3)	O1'—C19'—C1'	123.3 (3)
O2—C19—C1	112.7 (3)	O2'—C19'—C1'	112.9 (3)
O3—C20—O4	123.3 (3)	O3'—C20'—O4'	122.4 (3)
O3—C20—C9	123.1 (3)	O3'—C20'—C9'	123.8 (3)
O4—C20—C9	113.6 (2)	O4'—C20'—C9'	113.9 (3)
O5—C21—O6	123.4 (3)	O5'—C21'—O6'	123.9 (3)
O5—C21—C15	122.1 (3)	O5'—C21'—C15'	121.5 (3)
O6—C21—C15	114.6 (3)	O6'—C21'—C15'	114.5 (3)
C1—C2—C3—C4	2.2 (4)	C1'—C2'—C3'—C4'	-2.3 (4)
C1—C2—C7—C8	-54.7 (4)	C1'—C2'—C7'—C8'	54.2 (4)
C1—C2—C7—C12	127.4 (3)	C1'—C2'—C7'—C12'	-127.7 (3)
C1—C6—C13—C14	125.4 (3)	C1'—C6'—C13'—C14'	-127.7 (3)
C1—C6—C13—C18	-55.3 (4)	C1'—C6'—C13'—C18'	52.0 (4)
C2—C1—C6—C5	-2.8 (4)	C2'—C1'—C6'—C5'	1.0 (4)
C2—C1—C6—C13	176.9 (3)	C2'—C1'—C6'—C13'	-178.1 (3)
C2—C1—C19—O1	-63.6 (4)	C2'—C1'—C19'—O1'	62.6 (4)
C2—C1—C19—O2	117.5 (3)	C2'—C1'—C19'—O2'	-117.0 (3)
C2—C3—C4—C5	-1.6 (5)	C2'—C3'—C4'—C5'	1.2 (4)
C2—C7—C8—C9	-178.5 (3)	C2'—C7'—C8'—C9'	177.8 (3)
C2—C7—C12—C11	179.4 (3)	C2'—C7'—C12'—C11'	-178.3 (3)
C3—C2—C7—C8	122.8 (3)	C3'—C2'—C7'—C8'	-123.3 (3)
C3—C2—C7—C12	-55.2 (4)	C3'—C2'—C7'—C12'	54.8 (4)
C3—C4—C5—C6	-1.4 (5)	C3'—C4'—C5'—C6'	1.1 (4)
C4—C5—C6—C1	3.5 (4)	C4'—C5'—C6'—C1'	-2.2 (4)
C4—C5—C6—C13	-176.2 (3)	C4'—C5'—C6'—C13'	176.9 (3)
C5—C6—C13—C14	-54.9 (4)	C5'—C6'—C13'—C14'	53.2 (4)
C5—C6—C13—C18	124.4 (3)	C5'—C6'—C13'—C18'	-127.1 (3)
C6—C1—C2—C3	0.0 (4)	C6'—C1'—C2'—C3'	1.2 (4)
C6—C1—C2—C7	177.4 (3)	C6'—C1'—C2'—C7'	-176.3 (3)
C6—C1—C19—O1	114.7 (3)	C6'—C1'—C19'—O1'	-116.9 (3)
C6—C1—C19—O2	-64.1 (3)	C6'—C1'—C19'—O2'	63.6 (4)
C6—C13—C14—C15	-179.5 (3)	C6'—C13'—C14'—C15'	178.7 (3)
C6—C13—C18—C17	179.8 (3)	C6'—C13'—C18'—C17'	-178.3 (3)
C7—C2—C3—C4	-175.3 (3)	C7'—C2'—C3'—C4'	175.2 (3)
C7—C8—C9—C10	-0.2 (4)	C7'—C8'—C9'—C10'	0.6 (4)
C7—C8—C9—C20	177.4 (3)	C7'—C8'—C9'—C20'	-178.2 (3)
C8—C7—C12—C11	1.4 (4)	C8'—C7'—C12'—C11'	-0.2 (5)
C8—C9—C10—C11	0.1 (4)	C8'—C9'—C10'—C11'	-0.3 (4)
C8—C9—C20—O3	-172.2 (3)	C8'—C9'—C20'—O3'	178.4 (3)
C8—C9—C20—O4	7.9 (4)	C8'—C9'—C20'—O4'	-1.5 (4)
C9—C10—C11—C12	0.8 (5)	C9'—C10'—C11'—C12'	-0.3 (5)
C10—C9—C20—O3	5.5 (4)	C10'—C9'—C20'—O3'	-0.5 (5)
C10—C9—C20—O4	-174.4 (3)	C10'—C9'—C20'—O4'	179.6 (3)
C10—C11—C12—C7	-1.5 (5)	C10'—C11'—C12'—C7'	0.5 (5)
C12—C7—C8—C9	-0.5 (4)	C12'—C7'—C8'—C9'	-0.3 (4)

C13—C14—C15—C16	0.1 (4)	C13'—C14'—C15'—C16'	0.5 (4)
C13—C14—C15—C21	-179.2 (3)	C13'—C14'—C15'—C21'	179.9 (3)
C14—C13—C18—C17	-0.9 (4)	C14'—C13'—C18'—C17'	1.4 (4)
C14—C15—C16—C17	-1.7 (4)	C14'—C15'—C16'—C17'	-0.4 (4)
C14—C15—C21—O5	163.2 (3)	C14'—C15'—C21'—O5'	-162.4 (3)
C14—C15—C21—O6	-16.1 (4)	C14'—C15'—C21'—O6'	18.1 (4)
C15—C16—C17—C18	1.9 (5)	C15'—C16'—C17'—C18'	0.7 (5)
C16—C15—C21—O5	-16.1 (4)	C16'—C15'—C21'—O5'	17.0 (4)
C16—C15—C21—O6	164.5 (3)	C16'—C15'—C21'—O6'	-162.6 (3)
C16—C17—C18—C13	-0.6 (5)	C16'—C17'—C18'—C13'	-1.2 (5)
C18—C13—C14—C15	1.1 (4)	C18'—C13'—C14'—C15'	-1.0 (4)
C19—C1—C2—C3	178.3 (3)	C19'—C1'—C2'—C3'	-178.2 (3)
C19—C1—C2—C7	-4.2 (4)	C19'—C1'—C2'—C7'	4.3 (4)
C19—C1—C6—C5	178.9 (3)	C19'—C1'—C6'—C5'	-179.5 (3)
C19—C1—C6—C13	-1.4 (4)	C19'—C1'—C6'—C13'	1.3 (4)
C20—C9—C10—C11	-177.6 (3)	C20'—C9'—C10'—C11'	178.6 (3)
C21—C15—C16—C17	177.7 (3)	C21'—C15'—C16'—C17'	-179.7 (3)

*Hydrogen-bond geometry (Å, °)*

D—H···A	D—H	H···A	D···A	D—H···A
O2—H2···O3 <sup>i</sup>	0.91 (5)	1.79 (5)	2.657 (3)	159 (4)
O4—H4···O1 <sup>ii</sup>	0.90 (6)	1.84 (6)	2.689 (3)	156 (5)
O2'—H2'···O3 <sup>iii</sup>	0.87 (6)	1.76 (6)	2.628 (3)	171 (5)
O4'—H4'···O1 <sup>iv</sup>	0.84 (5)	1.90 (5)	2.717 (3)	166 (4)
O6'—H6'···O5	0.92 (4)	1.69 (4)	2.593 (3)	168 (4)
O6—H6···O5'	1.01 (3)	1.58 (3)	2.581 (3)	172 (6)

Symmetry codes: (i)  $-x+1, -y+2, z-1/2$ ; (ii)  $x, y-1, z$ ; (iii)  $-x+3/2, y+1, z+1/2$ ; (iv)  $x, y+1, z$ .