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#### Key indicators

Single-crystal X-ray study  
 $T = 123\text{ K}$   
Mean  $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$   
 $R$  factor = 0.036  
 $wR$  factor = 0.087  
Data-to-parameter ratio = 10.2

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

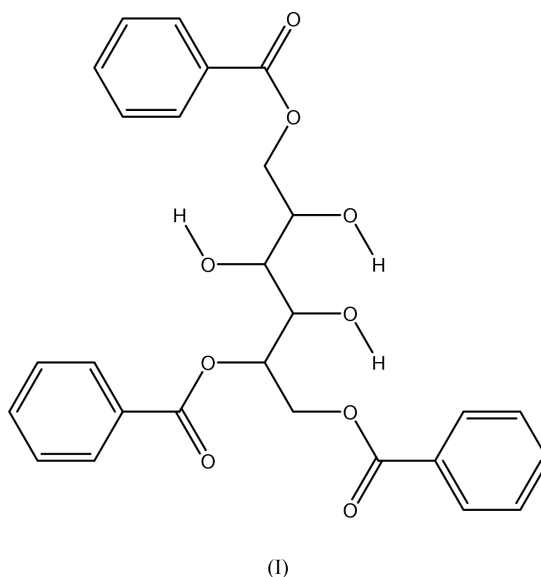
## D-Mannitol-1,2,6-tribenzoate

D-Mannitol-1,2,6-tribenzoate,  $\text{C}_{27}\text{H}_{26}\text{O}_9$ , was obtained as a side product from the reaction of D-mannitol with benzoyl chloride in hot pyridine. The major product of the reaction was D-mannitol-1,6-dibenzoate.

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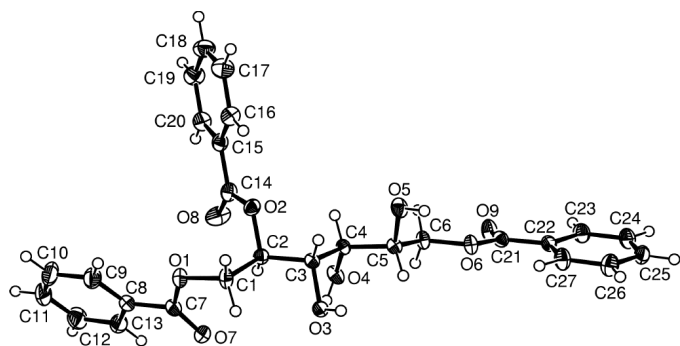
#### Comment

The application of boronic acids as labile protective agents in the selective functionalization of polyols is currently one of the focus areas of our research (Bhaskar *et al.*, 2001). The title compound, (I), was obtained from a control experiment, which was performed in order to determine the yield and selectivity of the reaction of D-mannitol with benzoyl chloride in the absence of a boronic acid. It is interesting to note that the third acylation of D-mannitol occurs at 2-OH, rather than at 3-OH.



#### Experimental

D-Mannitol (20.03 g, 0.11 mol) was treated with benzoyl chloride (12.30 ml, 0.11 mol) in hot pyridine (75 ml), following the procedure of Morpain & Tisserand (1979). After the reaction mixture was cooled, a white precipitate was obtained, which was recrystallized from hot methanol to afford pure D-mannitol-1,6-dibenzoate (4.29 g, 20%) (Bhaskar *et al.*, 2001). The addition of water to the recrystallization filtrate, followed by prolonged cooling, produced a crop of colourless needles that were collected by filtration, washed with diethyl ether and dried in air. Spectroscopic and X-ray crystallographic characterization of the needles revealed the second product to be D-mannitol-1,2,6-tribenzoate (0.31 g, 2%); m.p. 438–441 K [literature 439–440 K (Hockett & Fletcher, 1944)].



**Figure 1**  
View of (I) (50% probability displacement ellipsoids).

#### Crystal data

$C_{27}H_{26}O_9$   
 $M_r = 494.48$   
Orthorhombic,  $P2_12_12_1$   
 $a = 5.572$  (1) Å  
 $b = 15.530$  (1) Å  
 $c = 27.221$  (2) Å  
 $V = 2355.5$  (5) Å<sup>3</sup>  
 $Z = 4$   
 $D_x = 1.394$  Mg m<sup>-3</sup>

Mo  $K\alpha$  radiation  
Cell parameters from 35667  
reflections  
 $\theta = 2.6$ – $28.3^\circ$   
 $\mu = 0.11$  mm<sup>-1</sup>  
 $T = 123$  (2) K  
Acicular, colourless  
 $0.30 \times 0.22 \times 0.18$  mm

#### Data collection

KappaCCD diffractometer  
CCD ( $\varphi$  and  $\omega$ ) scans  
26379 measured reflections  
3347 independent reflections  
2894 reflections with  $I > 2\sigma(I)$

$R_{int} = 0.042$   
 $\theta_{max} = 28.3^\circ$   
 $h = -7 \rightarrow 7$   
 $k = -20 \rightarrow 17$   
 $l = -36 \rightarrow 36$

#### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.036$   
 $wR(F^2) = 0.087$   
 $S = 1.08$   
3347 reflections  
328 parameters  
H atoms treated by a mixture of  
independent and constrained  
refinement

$w = 1/[\sigma^2(F_o^2) + (0.0433P)^2 + 0.3834P]$   
where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{max} = 0.008$   
 $\Delta\rho_{max} = 0.24$  e Å<sup>-3</sup>  
 $\Delta\rho_{min} = -0.17$  e Å<sup>-3</sup>

**Table 1**

Hydrogen-bonding geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$O3-H1 \cdots O7^i$	0.84	2.15	2.9641 (19)	163
$O4-H2A \cdots O3^{ii}$	0.84	2.15	2.9164 (18)	152
$O4-H2A \cdots O3$	0.84	2.43	2.8636 (18)	113
$O5-H3A \cdots O9^{iii}$	0.84	1.94	2.7668 (19)	169

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

As the absolute structure cannot be reliably determined for this light-atom study, all Friedel pairs (2368) have been merged. The H atoms were included in the riding-model approximation. The torsion angles about the C–O bonds of the hydroxyl groups have been refined.

Data collection: *COLLECT* (Nonius, 1997–2000); cell refinement: *HKL* and *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *HKL*, *DENZO* (Otwinowski & Minor, 1997) and *SCALEPACK*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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