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#### **Key indicators**

Single-crystal X-ray study T = 150 KMean  $\sigma$ (C–C) = 0.017 Å Disorder in solvent or counterion R factor = 0.040 wR factor = 0.100 Data-to-parameter ratio = 7.7

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. In the title compound,  $C_{32}H_{46}O_{19}\cdot 0.7H_2O$ , all hydroxy groups on the parent compound, trehalose, have been substituted and some disordered bound water is located in the weakly diffracting crystal.

2,2',3,3',4,4'-Hexaacetato-6,6'-bis(isobutanoyl)-a,a-

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# Comment

trehalose 0.7-hydrate

Trehalose is a naturally occurring non-reducing disaccharide which is effective in preserving the structural and functional integrity of membranes and proteins from the effects of dehydration (Crowe *et al.*, 1988). Polymorphic, amorphous and crystalline forms of trehalose have been studied (Sussich *et al.*, 1998), and the common crystalline form is the dihydrate (Brown *et al.*, 1972; Taga *et al.*, 1972). The ability of trehalose to readily form hydrates may be related to its support of an-hydrobiosis (life without water). We have undertaken a study of substituted trehalose molecules in an attempt to understand the enhanced stabilizing potential of trehalose. In the title compound, (I), all eight hydroxy groups have been substituted, hence removing the possibility of hydrogen-bonding networks with water through these groups.



However, an ill defined water molecule (occupation factor close to 0.7) was found to be incorporated into the crystal lattice. Short distances here are  $O1' \cdots O5$  3.097 (15) Å and  $O1' \cdots O6'(-1 + x, y, z)$  2.965 (13) Å, but uncertainty in location of water H atoms precludes a reliable description of the hydrogen bonding. The substitution at C1,C1' is  $\alpha,\alpha$ , with C1-O1 = 1.453 (11) Å, C1'-O1 = 1.435 (11) Å and C1-O1-C1' = 115.5 (8)°. The absolute configuration is *R* at the C atoms C1, C2, C4 and C5 (also at C1', C2', C4' and C5') and *S* at C3 and C3'. The two six-membered rings adopt chair conformations, with puckering parameters (Cremer & Pople, 1975) calculated with *PLATON* (Spek, 1998) of *Q* = 0.56 (1) Å,  $\theta = 9$  (1)°,  $\varphi = 94$  (7)° (primed atoms) and *Q* = 0.57 (1) Å,  $\theta = 3$  (1)° and  $\varphi = 322$  (21)° (unprimed atoms).

# **Experimental**

The title compound, (I), was prepared *via* the reaction mixture sequence of trehalose and  $Ph_3CCl$  followed by addition of  $CH_3COCl$  and pyridine, then aqueous hydrobromide was added and finally a

© 2001 International Union of Crystallography Printed in Great Britain – all rights reserved mixture of CH3CH(CH3)COCl and pyridine was prepared and added. Purification was achieved through a combination of column chromatography and recrystallization. Crystals for X-ray work were obtained by slow evaporation from 95% ethanol.

Mo  $K\alpha$  radiation

reflections

 $\mu = 0.11 \text{ mm}^{-1}$ 

T = 150 (2) K

Lozenge, colourless  $0.21 \times 0.10 \times 0.10$  mm

 $\theta = 2.0-25.0^{\circ}$ 

Cell parameters from 250

### Crystal data

 $C_{32}H_{46}O_{19} \cdot 0.7H_2O$  $M_r = 747.30$ Orthorhombic, P212121 a = 8.857 (3) Åb = 17.813 (11) Å c = 24.085 (3) Å  $V = 3800 (3) \text{ Å}^3$ Z = 4 $D_x = 1.306 \text{ Mg m}^{-3}$ 

### Data collection

Delft Instruments FAST diffract-	$R_{\rm int} = 0.311$
ometer	$\theta_{\rm max} = 25.0^{\circ}$
$\varphi$ and $\omega$ scans to fill Ewald sphere	$h = -10 \rightarrow 8$
17 076 measured reflections	$k = -20 \rightarrow 21$
3541 independent reflections	$l = -27 \rightarrow 27$
565 reflections with $I > 2\sigma(I)$	

### Refinement

Refinement on $F^2$	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.040$	$w = 1/[\sigma^2(F_o^2)]$
$wR(F^2) = 0.100$	$(\Delta/\sigma)_{\rm max} < 0.001$
S = 0.39	$\Delta \rho_{\rm max} = 0.19 \ {\rm e} \ {\rm A}^{-3}$
3541 reflections	$\Delta \rho_{\rm min} = -0.17 \text{ e} \text{ \AA}^{-3}$
460 parameters	Absolute structure: none

Detailed data collection procedures are described by Darr et al. (1993). The number of intensities with  $F^2 > 2\sigma F^2$  was only 16% of the total (565 compared with a total of 3541 unique reflections). Structure solution and refinement with such a weak data led to results that were below the standard normally expected, but are adequate for a qualitative assessment of the structure. With SIR92 (Altomare et al., 1994), the number of atoms in the unit cell was lowered to C100H168O76 from C128H192O80 to achieve suitable phasing. All non-H atoms were refined using the DELU command which applies rigid bond restraints was used. The H atoms were initially placed in calculated positions and thereafter allowed to ride on their attached atoms  $[U_{iso} = 1.2U_{eq}(C)]$ . Water H atoms were not observed on a final difference map but were calculated with CALC-OH (Nardelli, 1999) and the coordinates fixed. Friedel pairs were merged. The absolute configuration is known for trehalose.

Data collection: MADNES (Plugrath & Messerschmidt, 1989); cell refinement: MADNES; data reduction: ABSMAD (Karaulov, 1992); program(s) used to refine structure: SIR92 (Altomare et al., 1994); program(s) used to solve structure: SHELX97 (Sheldrick, 1997);





The atomic arrangement in the title molecule. Displacement ellipsoids are shown at the 50% probability level.

molecular graphics: ORTEP-3 (Farrugia, 1997); software used to prepare material for publication: ORTEP-3 (Farrugia, 1997).

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