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Key indicators

Single-crystal X-ray study T = 296 KMean $\sigma(\text{C-C}) = 0.004 \text{ Å}$ Some non-H atoms missing Disorder in main residue R factor = 0.049 wR factor = 0.162 Data-to-parameter ratio = 14.1

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

Benzyltributylammonium 4-hydroxynaphthalene-1-sulfonate

The title compound, $C_{19}H_{34}N^+ \cdot C_{10}H_7O_4S^-$, is a charge-control agent used in electrophotography. The anions form chains along the *b* axis through $O-H \cdot \cdot \cdot O$ hydrogen bonding.

Comment

The title compound, (I), is an ammonium salt used widely as a charge-control agent (CCA) of the positive type for toners in electrophotography (Tanaka, 1995). CCAs are usually added to toners to create a desired charge level and polarity (Nash *et al.*, 2001). However, the charge-control mechanism of CCA is not fully understood at the moment. We have, therefore, determined the title crystal structure as a step to elucidating the mechanism.



Fig. 1 shows the asymmetric unit of (I). The ions have no crystallographically imposed symmetry. Fig. 2 shows a hydrogen-bonded chain along the *b* axis, formed by $O-H\cdots O$ hydrogen bonding (Table 1).

Experimental

Compound (I) was obtained from Orient Chemical Industries Ltd, and was recrystallized from a methanol solution. After 48 h, a number of colorless crystals were obtained in the form of blocks.

Crystal data

 $\begin{array}{l} C_{19}H_{34}N^{+}\cdot C_{10}H_7O_4S^{-}\\ M_r = 499.70\\ \text{Monoclinic, } P_{2_1}/n\\ a = 14.3810 \ (11) \text{ Å}\\ b = 9.8124 \ (7) \text{ Å}\\ c = 19.7757 \ (15) \text{ Å}\\ \beta = 92.560 \ (5)^{\circ} \end{array}$

 $V = 2787.8 \text{ (4) } \text{\AA}^3$ Z = 4Cu K\alpha radiation $\mu = 1.29 \text{ mm}^{-1}$ T = 296.1 K $0.20 \times 0.20 \times 0.20 \text{ mm}$

Data collection

Rigaku R-AXIS RAPID diffractometer Absorption correction: multi-scan (ABSCOR; Higashi, 1995) $T_{min} = 0.745, T_{max} = 0.772$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.049$ $wR(F^2) = 0.162$ S = 1.194902 reflections 24226 measured reflections 4902 independent reflections 3490 reflections with $F^2 > 2\sigma(F^2)$ $R_{\rm int} = 0.036$

347 parameters H-atom parameters constrained
$$\begin{split} &\Delta \rho_{max} = 0.45 \text{ e } \text{\AA}^{-3} \\ &\Delta \rho_{min} = -0.24 \text{ e } \text{\AA}^{-3} \end{split}$$

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The asymmetric unit of (I), showing 30% probability displacement ellipsoids and only the major disorder components.

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - H \cdots A$
$O4-H4\cdots O1^i$	0.82	1.85	2.657 (2)	169
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Symmetry code: (i) $-x + \frac{1}{2}, y + \frac{1}{2}, -z + \frac{1}{2}$.

C4, C5 and C6 were found to be disordered over two sites each. The site occupancies for C4A/C4B are 0.742 (7):0.258 (7), whereas those for for C5A/C5B and C6A/C6B are 0.460 (6):0.540 (6). These atoms were isotropically refined. All H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with C–H = 0.93 Å (aromatic), 0.96 Å (methyl) or 0.97 Å (methylene), and O–H = 0.82 Å; $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm parent atom})$.

Data collection: *PROCESS-AUTO* (Rigaku, 1998); cell refinement: *PROCESS-AUTO*; data reduction: *CrystalStructure* (Rigaku/ MSC, 2006); program(s) used to solve structure: *SIR2004* (Burla *et al.*,



Figure 2

The formation of a hydrogen-bonded (dashed lines) chain.

2003); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPIII* (Burnett & Johnson, 1996); software used to prepare material for publication: *CrystalStructure*.

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