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## Crystal Structure

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# Triaquamanganese sulfite revisited 

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Orthorhombic $\mathrm{Mn}\left(\mathrm{SO}_{3}\right)\left(\mathrm{H}_{2} \mathrm{O}\right)_{3}$ has been reinvestigated by single-crystal X-ray diffraction in two possible space groups, viz. $P 2_{1} 2_{1} 2_{1}$ (with all atoms in general positions) and Pnma (with the molecule bisected by a mirror plane). The results confirm the lower symmetry assigned in a previous singlecrystal neutron diffraction study. However, the refinement of the $P 2_{1} 2_{1} 2_{1}$ model requires the introduction of racemic twinning and soft positional and displacement restraints for the H atoms. The importance of a scrupulous report on symmetry absence violations as standard policy in crystallographic work is discussed.

## Comment

$\mathrm{Mn}\left(\mathrm{SO}_{3}\right)\left(\mathrm{H}_{2} \mathrm{O}\right)_{3}$ is known to crystallize in two polymorphic varieties, viz. a monoclinic form (Engelen \& Freiburg, 1979; Johansson \& Lindqvist, 1980) and an orthorhombic form (Baggio \& Baggio, 1976; Engelen, 1983; Basso et al., 1991; Gonschorek et al., 1996). Both structures present the same local coordination scheme $\left(\mathrm{SO}_{3}\right.$ bound to three different cationic units via its three O atoms) but with a different threedimensional connectivity (to be discussed below). While the structure of the monoclinic form is well established, that of the orthorhombic form still presents an apparent contradiction between the Pnma symmetry originally determined and the $P 2_{1} 2_{1} 2_{1}$ symmetry associated with an ordered distribution of its aqua H atoms.

X-ray studies could not distinguish definitively between the options 'Pnma + splitting' and ' $P 2_{1} 2_{1} 2_{1}+$ ordering'. As a result of 'real symmetry' or 'strong pseudosymmetry', both Pnma and $P 2_{1} 2_{1} 2_{1}$ treatments resulted in similar difference Fourier maps, from which it was almost impossible to differentiate genuine H atoms from eventual ghost images. A single-crystal neutron diffraction study (Gonschorek et al., 1996) determined $P 2_{1} 2_{1} 2_{1}$ as the correct space group, at least for the deuterated samples used. Unfortunately, this seemingly
accurate work ended up in a report impaired by a number of numerical inconsistencies, some of which affected relevant atomic coordinates involved in the 'hot' part of the structure, and rendered much of the reported geometry meaningless. ${ }^{1}$

We present here the results of the structure refinement of a good X-ray data set in the two controversial space groups, namely (i) $P 2_{1} 2_{1} 2_{1}$ and (ii) Pnma. Fig. 1 shows the corresponding ellipsoid plots and numbering schemes, while Tables 1 and 3 present relevant distances and angles. Some redundant values have been included in Table 3 in order to facilitate comparison with corresponding values in Table 1.

A simplified view of the structure as obtainable in a difference Fourier synthesis is shown in Fig. 2(a). The 'H-like' atoms drawn are the maxima clearly appearing in both centroand non-centrosymmetric difference Fourier maps; the two maps look equivalent except for the presence of a mirror in Pnma (Fig. 1b) versus a pseudo-mirror in $P 2_{1} 2_{1} 2_{1}$ (Fig. 1a).

First, there is a clear distinction between atom $\mathrm{O} 1 W$ with its two well behaved H atoms and the remaining atoms $\mathrm{O} 2 W$ and O3W, each showing three plausible maxima in the Fourier maps. These 'proto-H atoms', in turn, fall into two categories, viz. (i) $\mathrm{H} 2 W A$ and $\mathrm{H} 3 W A$ (one for each aqua ligand), which do not present any possible 'steric collision' with other atoms in the structure and are in principle good candidates to be real H atoms, and (ii) other H atoms that do present steric hindrance problems [in $P 2_{1} 2_{1} 2_{1}$ : H2WB $\cdots \mathrm{H} 3 W C\left(2-x, \frac{1}{2}+y\right.$, $\left.\frac{1}{2}-z\right)=1.24 \AA$ and $\mathrm{H} 2 W C \cdots \mathrm{H} 3 W B(x, 1+y, z)=0.93 \AA$; in Pnma: H2WB $\cdots \mathrm{H} 2 W B(2-x, 2-y, 1-z)=1.34 \AA$ and $\left.\mathrm{H} 2 W C \cdots \mathrm{H} 2 W C\left(x, \frac{5}{2}-y, z\right)=1.04 \AA\right]$. In $P 2_{1} 2_{2} 2_{1}$, this problem could be overcome by the judicious choice of one H atom from each set ( $\mathrm{H} 2 W B$ and $\mathrm{H} 3 W C$, or $\mathrm{H} 2 W C$ and $\mathrm{H} 3 W B$ ) as the correct partners of $\mathrm{H} 2 W A$ and $\mathrm{H} 3 W A$. In Pnma, the existence of $\mathrm{H} 2 W B(C)$ would necessarily imply the existence of $\mathrm{H} 3 W B(C)$; in this case, only a split model is possible in the form of a $50: 50$ mixture of the two possible non-centrosymmetric solutions.

The $a b$ initio analysis of systematic absences (and their violations) favoured $P 2_{1} 2_{1} 2_{1}$ rather than Pnma (Table 5). The strongest violation occurs for the multiply determined 041 reflection with an intensity $I \simeq 60 \sigma$ in seven different measurements in the whole data set. Overall, the strong pseudosymmetry is apparent from the very low mean intensities of the offending reflections.

In order to test both hypotheses, three unrestrained refinements in the two possible space groups were performed, including two ordered models in $P 2_{1} 2_{1} 2_{1}$ (one for each possible choice of the controversial H atoms, viz. $\mathrm{H} 2 W A$, $\mathrm{H} 2 W B / \mathrm{H} 3 W A, \mathrm{H} 3 W C$ or $\mathrm{H} 2 W A, \mathrm{H} 2 W C / \mathrm{H} 3 W A, \mathrm{H} 3 W B)$ and a disordered structure in Pnma, with a fully occupied $\mathrm{H} 2 W A$ and two half-occupied $\mathrm{H} 2 W B$ and $\mathrm{H} 2 W C$ sites. The results of both $P 2_{1} 2_{1} 2_{1}$ refinements yielded significantly lower $R$ factors

[^0]( 0.0199 and 0.0215 versus 0.0238 ), though at the cost of nearly twice as many parameters and with the need of a (nearly equipopulated) racemic twin model ( $0.45 / 0.55$ ). In addition, the behaviour of the controversial H atoms during the completely free refinement was rather erratic and a smooth convergence could only be achieved when soft restraints were applied. In contrast, the Pnma model proved to be quite robust and, in spite of its slightly larger $R$ factor, provided a sensible and stable solution even when severely over-refined.

Our conclusion is that $\mathrm{Mn}\left(\mathrm{SO}_{3}\right)\left(\mathrm{H}_{2} \mathrm{O}\right)_{3}$ crystallizes in the non-centrosymmetric $P 2_{1} 2_{1} 2_{1}$ space group, as confirmed by the lower $R$ indices and the violations of the expected systematic absences in Pnma. In ordinary circumstances, without previous information or with only a medium-quality X-ray data set, it would be impossible to discriminate between the $P 2_{1} 2_{1} 2_{1}$ and Pnma space groups and the latter would be normally preferred. This result alerts us to the fact that it might sometimes be easy to overlook annoying violations of systematic absences.

For the sake of completeness, a very brief description of the $P 2_{1} 2_{1} 2_{1}$ structure of $\operatorname{Mn}\left(\mathrm{SO}_{3}\right)\left(\mathrm{H}_{2} \mathrm{O}\right)_{3}$ follows. The basic structural unit consists of an $\mathrm{Mn}^{\mathrm{II}}$ cation octahedrally coordinated to three water molecules and atoms $\mathrm{O} 1, \mathrm{O} 2^{\mathrm{i}}$ and $\mathrm{O} 3^{\mathrm{ii}}$ from three symmetry-related sulfite ions (Fig. 1a). The anion is bonded through each of its O atoms to a different cation in a $\mu_{3}$-mode to generate a double chain running along $b$, where both the cations and the anions are surrounded by aqua ligands (Fig. 2b). These good hydrogen-bond donors find an adequate number of acceptors to build up an extremely complex hydrogen-bonding network. The interactions are either inter-chain (entries $1-3$ in Table 2) or intra-chain (entries 4-6) and connect each chain to six others (Fig. 3). This chain structure is the most conspicuous difference between the orthorhombic and monoclinic forms of $\mathrm{Mn}\left(\mathrm{SO}_{3}\right)\left(\mathrm{H}_{2} \mathrm{O}\right)_{3}$. In fact, the latter presents a completely analogous cation coordination scheme but with a slightly different interconnectivity between polyhedra (Table 4), which leads to a tightly woven three-dimensional covalent structure. These packing differ-

(a)

(b)

Figure 1
Displacement ellipsoid plots ( $50 \%$ probability level) of the Mn polyhedron in (a) the $P 2_{1} 2_{1} 2_{1}$ and (b) the Pnma model. Symmetry-independent atoms are drawn as filled ellipsoids. The symmetry codes are as given in Tables 1 and 3 .

(a)

(b)

Figure 2
Schematic diagrams presenting $(a)$ the possible H-atom positions and $(b)$ their correct assignment in $P 2_{1} 2_{2} 2_{1}$. Note the formation of chains along the horizontal $b$ direction.


Figure 3
The packing in $P 2_{1} 2_{1} 2_{1}$, viewed along the chain direction, showing a central chain (bold lines) connected to six different neighbours (thin lines). The inter-chain hydrogen-bonding interactions are indicated by broken lines.
ences, however, do not affect the compactness of the material, as the two structures differ in their crystal densities by less than $1 \%$.

## Experimental

The title compound was prepared as described by Baggio \& Baggio (1976) by allowing equimolar aqueous solutions of $\mathrm{MnSO}_{4}$ and $\mathrm{Na}_{2} \mathrm{SO}_{3}$ to diffuse slowly in a U-shaped tube through a much denser $\mathrm{Na}_{2} \mathrm{SO}_{4}$ solution, which acted both as a separation medium (avoiding initial mixing of the original solutions) and as a deterring agent (slowing down the diffusion rate). After a few weeks, colourless prisms adequate for X-ray diffraction analysis were observed at the glass walls.

## Determination in $\mathbf{P 2} \mathbf{1 2}_{1} \mathbf{2}_{1}$

## Crystal data

| $\mathrm{Mn}\left(\mathrm{SO}_{3}\right)\left(\mathrm{H}_{2} \mathrm{O}\right)_{3}$ | $Z=4$ |
| :--- | :--- |
| $M_{r}=189.05$ | $D_{x}=2.391 \mathrm{Mg} \mathrm{m}$ |
| Orthorhombic, $P_{2} 2_{1} 2_{1} 2_{1}$ | $\mathrm{Mo} \mathrm{K} \mathrm{\alpha} \mathrm{radiation}$ |
| $a=9.7577(10) \AA$ | $\mu=2.86 \mathrm{~mm}^{-1}$ |
| $b=5.6319(6) \AA$ | $T=295(2) \mathrm{K}$ |
| $c=9.5579(10) \AA$ | Prism, colourless |
| $V=525.25(10) \AA^{3}$ | $0.16 \times 0.10 \times 0.08 \mathrm{~mm}$ |
| Data collection |  |
| Bruker SMART CCD area-detector | 4391 measured reflections |
| $\quad$ diffractometer | 1190 independent reflections |
| $\varphi$ and $\omega$ scans | 1152 reflections with $I>2 \sigma(I)$ |
| Absorption correction: multi-scan | $R_{\text {int }}=0.021$ |
| $\quad(S A D A B S ;$ Sheldrick, 2001) | $\theta_{\text {max }}=27.9^{\circ}$ |
| $\quad T_{\text {min }}=0.658, T_{\text {max }}=0.804$ |  |

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.020$
$w R\left(F^{2}\right)=0.052$
$S=0.97$
1190 reflections
99 parameters
All H -atom parameters refined

$$
\begin{aligned}
& w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.033 P)^{2}\right. \\
& +0.26 P \text { ] } \\
& \text { where } P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}^{2}\right) / 3 \\
& (\Delta / \sigma)_{\max }<0.001 \text { 。 } \\
& \Delta \rho_{\text {max }}=0.28 \mathrm{e}^{-3} \\
& \Delta \rho_{\min }=-0.29 \mathrm{e}^{-3} \\
& \text { Absolute structure: Flack (1983), } \\
& 424 \text { Friedel pairs } \\
& \text { Flack parameter: } 0.45 \text { (4) }
\end{aligned}
$$

Table 1
Selected geometric parameters ( $\left({ }^{\circ},^{\circ}\right.$ ) for (I) in $P 2_{1} 2_{1} 2_{1}$.

| $\mathrm{Mn}-\mathrm{O} 1$ | 2.1673 (13) | $\mathrm{Mn}-\mathrm{O} 3 \mathrm{~W}$ | 2.242 (2) |
| :---: | :---: | :---: | :---: |
| $\mathrm{Mn}-\mathrm{O} 2{ }^{\text {i }}$ | 2.162 (2) | $\mathrm{S}-\mathrm{O} 1$ | 1.5202 (13) |
| $\mathrm{Mn}-\mathrm{O}^{\text {ii }}$ | 2.185 (2) | $\mathrm{S}-\mathrm{O} 2$ | 1.522 (2) |
| $\mathrm{Mn}-\mathrm{O} 1{ }^{\text {W }}$ | 2.1647 (14) | $\mathrm{S}-\mathrm{O} 3$ | 1.533 (2) |
| $\mathrm{Mn}-\mathrm{O} 2 \mathrm{~W}$ | 2.236 (2) |  |  |
| $\mathrm{O} 1^{\mathrm{i}}-\mathrm{Mn}-\mathrm{O} 2$ | 96.40 (10) | $\mathrm{O} 3{ }^{\text {ii }}-\mathrm{Mn}-\mathrm{O} 1 W$ | 85.64 (11) |
| $\mathrm{O} 1-\mathrm{Mn}-\mathrm{O}^{\text {ii }}$ | 94.40 (10) | $\mathrm{O} 3^{\text {ii }}-\mathrm{Mn}-\mathrm{O} 2 \mathrm{~W}$ | 93.30 (9) |
| $\mathrm{O} 1-\mathrm{Mn}-\mathrm{O} 1 W$ | 176.08 (8) | $\mathrm{O} 3{ }^{\text {ii }}-\mathrm{Mn}-\mathrm{O} 3 W$ | 171.50 (9) |
| $\mathrm{O} 1-\mathrm{Mn}-\mathrm{O} 2 W$ | 89.43 (10) | $\mathrm{O} 1 W-\mathrm{Mn}-\mathrm{O} 2 W$ | 86.66 (11) |
| $\mathrm{O} 1-\mathrm{Mn}-\mathrm{O} 3 W$ | 90.19 (10) | $\mathrm{O} 1 W-\mathrm{Mn}-\mathrm{O} 3 W$ | 89.28 (10) |
| $\mathrm{O} 2{ }^{\text {i }}-\mathrm{Mn}-\mathrm{O} 3^{\text {ii }}$ | 95.58 (6) | $\mathrm{O} 2 W-\mathrm{Mn}-\mathrm{O} 3 W$ | 79.58 (5) |
| $\mathrm{O} 2{ }^{\text {i }}-\mathrm{Mn}-\mathrm{O} 1 W$ | 87.49 (11) | $\mathrm{O} 1-\mathrm{S}-\mathrm{O} 2$ | 104.65 (14) |
| $\mathrm{O} 2{ }^{\text {i }}-\mathrm{Mn}-\mathrm{O} 2 W$ | 168.96 (9) | $\mathrm{O} 1-\mathrm{S}-\mathrm{O} 3$ | 104.21 (13) |
| $\mathrm{O} 2{ }^{\text {i }}-\mathrm{Mn}-\mathrm{O} 3 W$ | 91.00 (9) | $\mathrm{O} 2-\mathrm{S}-\mathrm{O} 3$ | 104.36 (8) |

Symmetry codes: (i) $-x+1, y-\frac{1}{2},-z+\frac{1}{2}$; (ii) $-x+1, y+\frac{1}{2},-z+\frac{1}{2}$.
Table 2
Hydrogen-bond geometry ( $\AA,^{\circ}$ ) for (I) in $P 2_{1} 2_{1} 2_{1}$.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :---: | :---: | :---: | :---: | :---: |
| $\mathrm{O} 1 W-\mathrm{H} 1 W A \cdots \mathrm{O} 1^{\text {iii }}$ | 0.79 (3) | 2.44 (3) | 3.227 (4) | 172 (5) |
| $\mathrm{O} 2 W-\mathrm{H} 2 W A \cdots \mathrm{O}^{\text {iv }}$ | 0.79 (3) | 1.90 (4) | 2.686 (5) | 171 (6) |
| $\mathrm{O} 3 W-\mathrm{H} 3 W A \cdots \mathrm{O} 2^{\mathrm{iv}}$ | 0.80 (3) | 1.92 (3) | 2.711 (4) | 171 (3) |
| $\mathrm{O} 1 W-\mathrm{H} 1 W B \cdots \mathrm{O} 1^{\text {v }}$ | 0.77 (3) | 2.42 (3) | 3.134 (4) | 155 (5) |
| $\mathrm{O} 2 W-\mathrm{H} 2 W B \cdots \mathrm{O} 3 W^{\text {vi }}$ | 0.79 (2) | 2.04 (3) | 2.830 (2) | 176 (3) |
| $\mathrm{O} 3 W-\mathrm{H} 3 W C \cdots \mathrm{O} 2 W^{\text {vii }}$ | 0.81 (3) | 1.99 (3) | 2.766 (2) | 158 (3) |

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

## Determination in Pnma

## Crystal data

$\mathrm{Mn}\left(\mathrm{SO}_{3}\right)\left(\mathrm{H}_{2} \mathrm{O}\right)_{3}$
$M_{r}=189.05$
Orthorhombic, Pnma
$a=9.7577$ (10) $\AA$
$b=5.6319$ (6) A
$c=9.5579$ (10) $\AA$
$V=525.25(10) \AA^{3}$
Data collection
Bruker SMART CCD area-detector diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan (SADABS; Sheldrick, 2001) $T_{\text {min }}=0.658, T_{\text {max }}=0.804$
$Z=4$
$D_{x}=2.391 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation
$\mu=2.86 \mathrm{~mm}^{-1}$
$T=295$ (2) K
Prism, colourless
$0.16 \times 0.10 \times 0.08 \mathrm{~mm}$

4138 measured reflections 670 independent reflections 661 reflections with $I>2 \sigma(I)$ $R_{\text {int }}=0.022$
$\theta_{\text {max }}=27.9^{\circ}$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.024$
$w R\left(F^{2}\right)=0.068$
$S=1.35$
670 reflections
59 parameters
All H -atom parameters refined

$$
\begin{aligned}
& w=1 /[ \sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.0226 P)^{2} \\
&+0.7961 P] \\
& \text { where } P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}^{2}\right) / 3 \\
&(\Delta / \sigma)_{\max }<0.001 \\
& \Delta \rho_{\max }=0.30 \mathrm{e} \AA^{-3} \\
& \Delta \rho_{\min }=-0.33 \mathrm{e} \AA^{-3}
\end{aligned}
$$

Table 3
Selected geometric parameters $\left(\AA,^{\circ}\right)$ for (I) in Pnma.

| $\mathrm{Mn}-\mathrm{O} 1$ | $2.166(3)$ | $\mathrm{Mn}-\mathrm{O} 2 W$ | $2.2396(19)$ |
| :--- | :---: | :--- | :---: |
| $\mathrm{Mn}-\mathrm{O} 2^{\mathrm{i}}$ | $2.1716(18)$ | $\mathrm{S}-\mathrm{O} 1$ | $1.519(3)$ |
| $\mathrm{Mn}-\mathrm{O} 1 W$ | $2.167(3)$ | $\mathrm{S}-\mathrm{O} 2$ | $1.5289(18)$ |
|  |  |  |  |
|  |  |  |  |
| $\mathrm{O} 1-\mathrm{Mn}-\mathrm{O} 2^{\mathrm{i}}$ | $95.34(7)$ | $\mathrm{O} 2^{\mathrm{ii}}-\mathrm{Mn}-\mathrm{O} 2 W$ | $92.15(7)$ |
| $\mathrm{O} 1-\mathrm{Mn}-\mathrm{O} 2 W$ | $89.82(8)$ | $\mathrm{O} 1 W-\mathrm{Mn}-\mathrm{O} 2 W$ | $87.99(9)$ |
| $\mathrm{O} 2^{\mathrm{i}}-\mathrm{Mn}-\mathrm{O} 2^{2 i}$ | $95.39(11)$ | $\mathrm{O} 2 W-\mathrm{Mn}-\mathrm{O} 2 W^{\text {iii }}$ | $79.78(10)$ |
| $\mathrm{O}^{\mathrm{i}}-\mathrm{Mn}-\mathrm{O} 1 W$ | $86.58(8)$ | $\mathrm{O} 1-\mathrm{S}-\mathrm{O} 2$ | $104.38(10)$ |
| $\mathrm{O} 2^{\mathrm{i}}-\mathrm{Mn}-\mathrm{O} 2 W$ | $170.43(8)$ | $\mathrm{O} 2^{\mathrm{iii}}-\mathrm{S}-\mathrm{O} 2$ | $104.63(15)$ |
| Symmetry codes: | (i) $-x+1, y-\frac{1}{2},-z+1 ;$ | (ii) $-x+1,-y+2,-z+1 ; \quad$ (iii) $x$, |  |
| $-y+\frac{3}{2}, z$. |  |  |  |

Table 4
Hydrogen-bond geometry ( $\AA,^{\circ}$ ) for (I) in Pnma.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | H $\cdots$ A | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :---: | :---: | :---: | :---: | :---: |
| $\mathrm{O} 1 W-\mathrm{H} 1 W \cdots \mathrm{O} 2^{\text {iv }}$ | 0.72 (4) | 2.47 (4) | 3.062 (3) | 141 (4) |
| $\mathrm{O} 2 W-\mathrm{H} 2 W A \cdots \mathrm{O} 2^{\text {v }}$ | 0.80 (4) | 1.91 (4) | 2.698 (3) | 167 (4) |
| $\mathrm{O} 2 W-\mathrm{H} 2 W B \cdots \mathrm{O} 2 W^{\text {vi }}$ | 0.73 (6) | 2.10 (6) | 2.834 (4) | 178 (7) |
| $\mathrm{O} 2 W-\mathrm{H} 2 W C \cdots \mathrm{O} 2 W^{\text {vii }}$ | 0.93 (6) | 1.89 (6) | 2.759 (4) | 154 (5) |

Symmetry codes: (i) $-x+1, y-\frac{1}{2},-z+1$; (ii) $-x+1,-y+2,-z+1$; (iii) $x,-y+\frac{3}{2}, z$; (iv) $-x+\frac{3}{2},-y+2, z+\frac{1}{2}$; (v) $x+\frac{1}{2}, y,-z+\frac{1}{2}$; (vi) $-x+2,-y+2,-z+1$; (vii) $x,-y+\frac{5}{2}, z$.

Table 5
Statistics of violations to (eventual) systematic absences.

|  | Along $a$ | Along $b$ | Along $c$ |
| :--- | :--- | :--- | :--- |
| SO | $b c n 2_{1}$ | $a c n 2_{1}$ |  |
| $N($ tot $)$ | 14013513310 | 2392382336 | $a b n 2_{1}$ |
| $N(\sigma)$ | 8195361 | 2162092111 | 1441481388 |
| $\langle I\rangle$ | 145.3153 .03 .90 .6 | 362.6398 .7402 .11 .1 | 5799820 |
| $\langle I / \sigma\rangle$ | 115.917 .42 .51 .5 | 37.037 .938 .61 .2 | 3.520 .4201 .70 .5 |

Notes: SO: eventual symmetry operations defining conditions among $h k l$ indices; $N($ tot $)$ : total number of reflections to which the former condition applies; $N(\sigma)$ : subset of $N$ (tot) with $I>3 \sigma(I)$ [number of violations to a $3 \sigma(I)$ level; $\langle I\rangle$ : mean intensity of the group; $\langle I / \sigma\rangle$ : mean intensity (in units of $\sigma$ ).

Owing to the real or strong pseudosymmetry presented by non-H atoms in both possible space groups, three plausible candidates for the two H atoms corresponding to each of the two controversial aqua molecules ( $\mathrm{O} 2 W$ and $\mathrm{O} 3 W$ ) clearly appeared in the difference

Fourier map (see Fig. 2a). In $P 2_{1} 2_{1} 2_{1}$, the correct set was chosen through a combination of steric arguments and least-squares refinement, which was extremely sensitive to the correct choice $(R / w R$ factors: $0.0199 / 0.0520$ and $0.0215 / 0.0579$ ). The model with lower $R$ effectively seems to correspond to the set reported by Gonschorek et al. (1996). However, and in spite of the extremely accurate data set ( $R_{\text {int }}=0.020$ ), refinement of the H atoms in the non-centrosymmetric space group was not possible without the use of restraints $[\mathrm{O}-\mathrm{H}=$ $0.80(5) \AA$ and $\left.U_{\text {iso }}(\mathrm{H})=0.06(4) \AA^{2}\right]$. The refinement also required an allowance for racemic twinning ( $0.45 / 0.55$ ). In the Pnma refinement, the split H -atom positions and their $U_{\text {iso }}$ values converged quite smoothly to reasonable values without the need for any external restraint.

For both determinations, data collection: SMART-NT (Bruker, 2001); cell refinement: SAINT-NT (Bruker, 2000); data reduction: SAINT-NT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL-NT (Bruker, 2000); software used to prepare material for publication: SHELXTL$N T$ and PLATON (Spek, 2005).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: BC3010). Services for accessing these data are described at the back of the journal.

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[^0]:    ${ }^{\mathbf{1}}$ The coordinates for atoms O3 and D2 as reported by Gonschorek et al. (1996) are clearly in error, as confirmed by their calculated distances to the Mn cation (not presented in the paper), viz. $\mathrm{Mn}-\mathrm{O} 3=1.84 \AA$ (expected $c a$ $2.20 \AA$ ) and $\mathrm{Mn} \cdots \mathrm{D} 2=1.58 \AA$ (expected $2.65-2.85 \AA$ ). In fact, only a few of the reported bonds and angles can be derived precisely from the published coordinates.

