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Titanyl sulfate dihydrate - template for preparation of 1D titania structures

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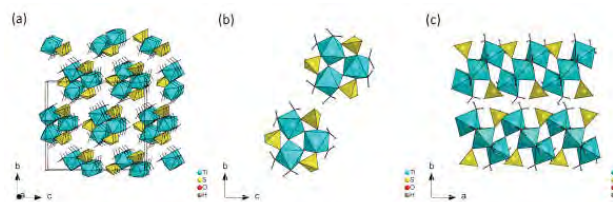
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Titanyl sulfate dihydrate (TSD, $\text{TiOSO}_4 \cdot 2\text{H}_2\text{O}$) can serve as a template for preparation of 1D titania structures. When immersed in aqueous ammonia at a temperature of about 0 °C the crystals of TSD provide solid residue composed of particles, whose shape and dimensions correspond perfectly to the rod-like particle morphology of the initial titanil sulfate with composition matching that of metatitanic acid. When annealed up to 1200 °C, the rod-like morphology of particles is retained, while the phase composition changes to anatase/rutile. To understand the mechanism of the phase transformation that is not accompanied with a change of particle shape, the structure of TSD had to be determined.

The TSD crystals form very thin platelets unsuitable for investigation by single-crystal x-ray diffraction. The structure was therefore solved using dynamical refinement of precession electron diffraction tomography data using PETS [1] and Jana2006 [2].

The structure is monoclinic Pn with lattice parameters $a = 5.54(2)$ Å, $b = 16.77(9)$ Å, $c = 18.74(10)$ Å, $\beta = 90.5(3)^\circ$. 54 independent non-hydrogen atoms were easily identified in the electrostatic potential distribution obtained ab initio by charge flipping. To keep charge balance such model requires 24 hydrogen atoms, whose positions were estimated based on the bond-valence calculations [3]. 17 hydrogens out of 24 could be matched to the maxima in the difference map calculated from combined refinement of 6 datasets. The remaining hydrogen positions were estimated based on the knowledge of water molecule geometry. The agreement factor R_{obs} decreased from 10.36 % to 9.89 % after including the hydrogens.

The TSD structure is composed of two symmetry independent helices of corner sharing TiO_6 octahedra extending along **a** (Figure 1). Each pseudotrigonal helix is held together by SO_4 tetrahedra that share corners with odd/even TiO_6 octahedra. Both hydrogens in each water molecule form hydrogen bonds toward oxygens of two different SO_4 tetrahedra. One H-bond points to the oxygen of SO_4 tetrahedra from the same helix, while the other H-bond points to the oxygen of SO_4 tetrahedra from the neighboring helix. Based on the structure a transformation mechanism could be proposed. During the reaction the sulfate anions are replaced with OH^- . The replacement of sulfate by OH^- anions in the helix does not cause a collapse of the helices as the newly bonded OH-groups repel one another. However, the interaction between the helices is weakened, resulting in polymer-like chains of TiO_6 octahedra characteristic for the amorphous metatitanic acid.



References:

- [1] L. Palatinus (2011). PETS - program for analysis of electron diffraction data. Prague: Institute of Physics of the AS CR. <http://pets.fzu.cz/>.
- [2] V. Petříček, M. Dušek, L. Palatinus, Crystallographic Computing System JANA2006: General features, Z. Krist. 229-5 (2014) 345–352.
- [3] I. D. Brown, The Chemical Bond in Inorganic Chemistry: The Bond Valence Model (Oxford Univ. Press, 2002).

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