

Powder X-ray diffraction of levothyroxine sodium pentahydrate, $C_{15}H_{10}I_4NNaO_4(H_2O)_5$

J.A. Kaduk,¹ K. Zhong,² T.N. Blanton,^{2,a)} S. Gates,² and T.G. Fawcett² ¹Illinois Institute of Technology, 3101 S. Dearborn St., Chicago, Illinois 60616 ²International Centre for Diffraction Data, 12 Campus Blvd., Newtown Square, Pennsylvania 19073-3273

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The room-temperature crystal structure of levothyroxine sodium pentahydrate has been refined using synchrotron powder diffraction data. The compound crystallizes in space group P1 (#1) with a = 8.2489(4), b = 9.4868(5), c = 15.8298(6) Å, $\alpha = 84.1387(4)$, $\beta = 83.1560(3)$, $\gamma = 85.0482(3)$ deg, V = 1220.071(9) Å³, and Z = 2. Hydrogen atoms (missing from the previously-reported structure) were included. © 2015 International Centre for Diffraction Data. [doi:10.1017/S0885715615000676]

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Levothyroxine sodium pentahydrate (marketed as Synthroid) is a synthetic form of the thyroid hormone, thyroxine, which is used in the treatment of thyroid hormone deficiency. Commercial levothyroxine sodium pentahydrate crystallizes in the triclinic space group P1 (#1), with a = 8.2489(4) Å, b = 9.4868(5) Å, c = 15.8298(6) Å, $\alpha = 84.1387$

(4)°, $\beta = 83.1560(3)^\circ$, $\gamma = 85.0482(3)^\circ$, V = 1220.071(9) Å³, and Z = 2. A reduced cell search in the Cambridge Structural Database yielded a previously reported crystal structure (Allen, 2002), which did not include hydrogens (Katrusiak *et al.*, 2004). In this work, the sample was ordered from United States Pharmacopeia, and analyzed as received. The

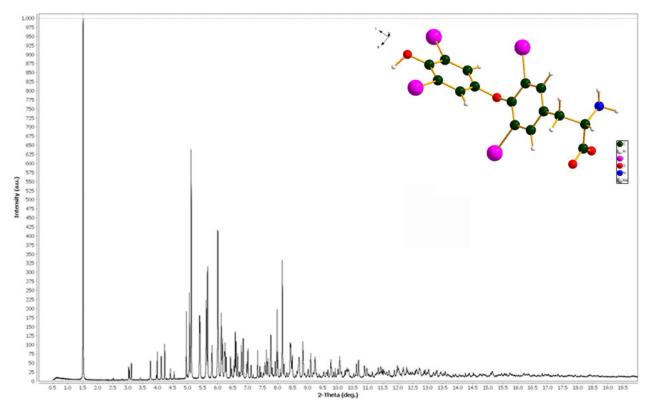


Figure 1. (Color online) Powder X-ray diffraction pattern of levothyroxine sodium pentahydrate.

^{a)}Author to whom correspondence should be addressed. Electronic mail: tblanton@icdd.com Data were submitted via Genie (http://www.icdd.com/ websubmission/launch.html), the ICDD[®] Web Submission Page.

room-temperature crystal structure was refined using synchrotron ($\lambda = 0.413\,891\,\text{\AA}$) powder diffraction data, density functional theory, and Rietveld refinement techniques. Hydrogen positions were included as part of the structure, and recalculated during the refinement. The diffraction data were collected on beamline 11-BM at the Advanced Photon Source, Argonne National Laboratory. Figure 1 shows the powder X-ray diffraction pattern of the compound.

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SUPPLEMENTARY MATERIALS

For supplementary material for this article, please visit http://dx.doi.org/10.1017/S0885715615000676

- Allen, F. H. (2002). "The Cambridge Structural Database: a quarter of a million crystal structures and rising," Acta Crystallogr. Sect. B: Struct. Sci. 58, 380–388. CSD Refcode QQQETG01.
- Katrusiak, A. and Katrusiak, A. (2004). "Thyroxine revisited," J. Pharm. Sci. 93(12), 3066–3075.