

Supplementary Material

for the paper

Solvent-Free Methylthiomethylithium $[\text{LiCH}_2\text{SMe}]_\infty$: Solid State Structure and Thermal Decomposition.

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1. Thermolysis Details:

The capillaries were dried by repeated cycles of heating followed by application of a dynamic vacuum and subsequent storage of the capillary in vacuo overnight. Since the screw-cap of the vial was made of plastic, higher temperatures could not be applied; the vials were thus dried just by storing them at room temperature in a dynamic vacuum for four hours.

2. Structure Solution and Refinement Details:

A high resolution X-ray powder diffraction pattern of $[\text{LiCH}_2\text{SMe}]_\infty$ was recorded at room temperature on a laboratory powder diffractometer (D-8, Bruker, Cu-K α_1 radiation from primary Ge(111)-Johanson-type monochromator; V α ntag-1 position sensitive detector) in Debye-Scherrer geometry with the sample sealed in a lithium borate glass capillary of 0.5 mm diameter (Hilgenberg glass No. 50). Data were taken from 8.0 – 68.0° 2 Θ at a speed of 0.05°/minute. The step width was 0.017° 2 Θ . The sample was spun during measurement for better particle statistics.

Data reduction on the powder diffraction pattern was performed using GUF1.¹ Indexing with ITO² led to a primitive monoclinic unit cell for $[\text{LiCH}_2\text{SMe}]_\infty$. Application of the extinction rules revealed $P2_1/c$ as the most probable space group. The number of formula units per unit cell was calculated to be $Z = 4$ from volume increments. The peak profiles and precise lattice parameters for all powder patterns were determined by LeBail-type fits³ using the program GSAS⁴. The background was modeled manually using GUF1. The peak-profile was described by a pseudo-Voigt function in combination with a special function that accounts for the asymmetry due to axial divergence^{5, 6}. The powder pattern of $[\text{LiCH}_2\text{SMe}]_\infty$ exhibits severe anisotropic peak broadening caused by lattice strain. The phenomenological strain model of Stephens⁷ as implemented in GSAS was used to model the anisotropy of the FWHM. Nine parameters were refined for the monoclinic phase.

The crystal structure of $[\text{LiCH}_2\text{SMe}]_\infty$ was solved by global optimization in direct space using the DASH structure solution package⁸. The measured powder pattern at room temperature was subjected to a Pawley refinement⁹ in space group $P2_1/c$, in order to extract correlated integrated intensities from the pattern. Good fits to the data were obtained. An internal coordinate description of the CH₂SMe moiety was constructed using bond lengths and angles from ref[3]. The positions of the lithium cation as well as the position and orientation and torsion angles of the CH₂SMe molecule in the unit cell were postulated and the trial structure

was subjected to a global optimization. The structures giving the best fit to the data in space group $P2_1/c$ was validated by Rietveld refinement¹⁰ of the fractional coordinates obtained at the end of the simulated annealing run using the GSAS program. To stabilize the refinement of the CH₂SMe molecule, slack soft constraints for the bond lengths (C-S, C-H) and bond angles (H-C-S) were introduced.

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3. Rietveld Plot

