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Supplementary Information

(1) Crystallographic information file for Sr[Be(OH)<sub>4</sub>]

(2) Figure S1 Electron micrographs of Sr[Be(OH)<sub>4</sub>] crystals

(3) Profile fit to NPD data and crystallographic model with hydrogen positions alone refined against these data; extracted O-H distances from this model.

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# Cif for Sr[Be(OH)<sub>4</sub>]<sub>2</sub>

data 2009btl0118i41acd SHELXL-97 audit creation method \_chemical\_name\_systematic ; ? ; \_chemical\_name\_common 'Strontium Tetrahydroxyberyllate' \_chemical\_melting\_point 'None' \_chemical\_formula\_moiety '40, 4H, Sr, Be' \_chemical\_formula\_sum 'Be H4 O4 Sr' chemical formula weight 164.66 loop atom type symbol \_atom\_type\_description atom type scat dispersion real \_atom\_type\_scat\_dispersion imag \_atom\_type\_scat\_source 'Be' 'Be' 0.0005 0.0002 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4' '0' '0' 0.0106 0.0060 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4' 'Ge' 'Ge' 0.1547 1.8001 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4' 'Sr' 'Sr' -1.5307 3.2498 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4' 'Si' 'Si' 0.0817 0.0704 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4' 'C' 'C' 0.0033 0.0016 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4' 'H' 'H' 0.0000 0.0000 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4' symmetry cell setting tetragonal \_symmetry\_space\_group\_name\_H-M I4(1)/acd loop \_symmetry\_equiv\_pos\_as\_xyz 'x, y, z' '-x+1/2, -y, z+1/2' '-y+1/4, x+3/4, z+1/4' 'y+1/4, -x+1/4, z+3/4' '-x+1/2, y, -z' 'x, -y, -z+1/2' 'y+1/4, x+3/4, -z+3/4' '-y+1/4, -x+1/4, -z+1/4' 'x+1/2, y+1/2, z+1/2'

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diffrn radiation type
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computing cell refinement
                                   ?
                                   ?
computing data reduction
computing structure solution
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computing structure refinement
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computing molecular graphics
                                   ?
computing publication material
                                   ?
refine special details
;
 Refinement of F<sup>2</sup> against ALL reflections. The weighted R-factor wR
and
 goodness of fit S are based on F^2\, conventional R-factors R are based
 on F, with F set to zero for negative F^2. The threshold expression of
 F^2 > 2 \setminus s(F^2) is used only for calculating R-factors(gt) etc. and is
 not relevant to the choice of reflections for refinement. R-factors
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 on \texttt{F}^2^{\star} are statistically about twice as large as those based on F, and
R-
 factors based on ALL data will be even larger.
;
```

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refine ls structure factor coef Fsqd
refine ls matrix type
                                   full
refine_ls_weighting_scheme
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refine ls weighting details
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                                   direct
atom sites solution secondary
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H2 H 0.260(6) 0.397(3) 0.069(3) 0.048(19) Uiso 1 1 d D . .
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04 0.0199(14) 0.0232(18) 0.0233(13) -0.0055(14) -0.0009(13) -0.0002(15)
Be 0.020(4) 0.014(4) 0.014(3) 0.000 0.002(3) 0.000
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into account individually in the estimation of s.u.'s in distances,
angles
and torsion angles; correlations between s.u.'s in cell parameters are
only
used when they are defined by crystal symmetry. An approximate
(isotropic)
treatment of cell s.u.'s is used for estimating s.u.'s involving l.s.
planes.
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O1 Sr2 2.586(3) 11 544 ?
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04 Sr2 2.626(3) . ?
Be O1 1.625(5) 14_454 ?
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Sr2 O1 2.586(3) 3 545 ?
Sr2 O1 2.586(3) 12 454 ?
Sr2 O1 2.586(3) 5 ?
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O4 Sr1 Be 118.35(15) . 20 656 ?
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Figure S1 Electron micrograph images of Sr[Be(OH)4] crystals



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### 3. NPD data analysis



Profile fit to NPD collected from  $SrBe(OH)_4$  at 250 K on D20, ILL Grenoble for 2 hours. Crystallographic model was that obtained at 298 K from SXD data (see cif above) with refinement of hydrogen atomic positions and ADPs only – see below. Red crosses are experimental data, green continuous line the calculated profile and the magneta lower continuous line the difference. Tick marks show reflection positions.

Refined hydrogen positions – crystallographic coordinates and esds. Labelling as in cif above.

H1 0.6018(9) 0.4680(10) 0.0117(7) H2 0.2493(18) 0.4154(8) 0.0714(5)

Extracted bond lengths

O1_H1	0.948(9)
O4_H2	0.969(8)