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> data Na5Pr4(OH)[SiO4]4 publ requested journal Inorg.Chem. \_publ\_contact author name 'Hans-Conrad zur Loye' \_publ\_contact author address ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 publ contact author email zurloye@mailbox.sc.edu publ contact author phone +1-803-777-6916 publ contact author fax +1-803-777-8508 loop publ author name \_publ\_author\_address 'zur Loye, Hans-Conrad' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Latshaw, Allison M.' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Chance, W. Michael' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Morrison, Gregory' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Smith, Mark D.' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208

'Yeon, Jeongho' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 audit creation method SHELXL-2013 chemical name systematic ? ? chemical name common chemical melting point ? chemical formula moiety 'H Na5 O17 Pr4 Si4' chemical formula sum 'H Na5 O17 Pr4 Si4' chemical formula weight 1063.96 loop atom type symbol atom type description atom type scat dispersion real atom type scat dispersion imag atom type scat source O O 0.0106 0.0060 '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' Na Na 0.0362 0.0249 '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' Pr Pr -0.2180 2.8214 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4' \_space\_group\_crystal\_system tetragonal

\_space\_group\_IT\_number 82 \_space\_group\_name\_H-M\_alt 'I -4' \_space\_group\_name\_Hall 'I -4'

\_shelx\_space\_group\_comment

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The symmetry employed for this shelxl refinement is uniquely defined by the following loop, which should always be used as a source of symmetry information in preference to the above space-group names. They are only intended as comments.

loop\_ \_space\_group\_symop\_operation\_xyz 'x, y, z'

'-x, -y, z' 'y, -x, -z' '-y, x, -z' 'x+1/2, y+1/2, z+1/2' '-x+1/2, -y+1/2, z+1/2' 'y+1/2, -x+1/2, -z+1/2' '-y+1/2, x+1/2, -z+1/2'
_cell_length_a12.0588(3)_cell_length_b12.0588(3)_cell_length_c5.4668(2)_cell_angle_alpha90_cell_angle_beta90_cell_angle_gamma90_cell_volume794.95(5)_cell_formula_units_Z2_cell_measurement_temperature296(2)_cell_measurement_theta_min2.388_cell_measurement_theta_max33.996
<pre>_exptl_crystal_description cube _exptl_crystal_colour 'light green' _exptl_crystal_density_meas ? _exptl_crystal_density_method ? _exptl_crystal_density_diffrn 4.445 _exptl_crystal_F_000 968 _exptl_transmission_factor_min ? _exptl_transmission_factor_max ? _exptl_crystal_size_max 0.080 _exptl_crystal_size_min 0.060 _exptl_crystal_size_min 0.060 _exptl_absorpt_coefficient_mu 12.573 _shelx_estimated_absorpt_T_min 0.433 _shelx_estimated_absorpt_T_max 0.519 _exptl_absorpt_correction_type multi-scan _exptl_absorpt_correction_T_min 0.7211 _exptl_absorpt_correction_T_max 1.0000 _exptl_absorpt_process_details 'SADABS Version 2.05 (Bruker, 2001)'</pre>
_exptl_special_details ; ? ;

\_diffrn\_ambient\_temperature 296(2)

diffrn radiation wavelength 0.71073 diffrn radiation type MoK\a diffrn source diffrn measurement device type 'Bruker SMART APEX CCD diffractometer' diffrn measurement method 'omega and phi scans' diffrn detector area resol mean? diffrn reflns number 9092 0.0351 diffrn reflns av unetI/netI diffrn reflns av R equivalents 0.0374 diffrn reflns limit h min -20 diffrn reflns limit h max 18 diffrn reflns limit k min -17 diffrn reflns limit k max 19 diffrn reflns limit 1 min -9 diffrn reflns limit 1 max 8 diffrn reflns theta min 2.389 diffrn reflns theta max 36.276 diffrn reflns theta full 25.242 diffrn measured fraction theta max 0.999 diffrn measured fraction theta full 0.998 diffrn reflns Laue measured fraction max 0.999 diffrn reflns Laue measured fraction full 0.998 diffrn reflns point group measured fraction max 0.999 diffrn reflns point group measured fraction full 0.999 reflns number total 1852 reflns number gt 1691 reflns threshold expression 'I > 2 (I)reflns Friedel coverage 0.845 reflns Friedel fraction max 1.000 reflns Friedel fraction full 1.000

\_reflns\_special\_details

## ;

Reflections were merged by SHELXL according to the crystal class for the calculation of statistics and refinement.

\_reflns\_Friedel\_fraction is defined as the number of unique Friedel pairs measured divided by the number that would be possible theoretically, ignoring centric projections and systematic absences.

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_computing_data_collection	'SMART Version 5.625 (Bruker, 2001)'
_computing_cell_refinement	'SAINT-Plus Version 6.45 (Bruker, 2001)'
_computing_data_reduction	'SAINT-Plus Version 6.45 (Bruker, 2001)'
_computing_structure_solution	'SHELXS97 (Sheldrick, 1997)'

computing structure refinement 'SHELXL-2013 (Sheldrick, 2013)' computing molecular graphics various computing publication material various refine special details Refined as a 2-component inversion twin. refine ls structure factor coef Fsqd refine ls matrix type full refine ls weighting scheme calc refine ls weighting details  $w=1/[\sqrt{6^2}(6^2)+(0.0374P)^2+0.8588P]$ where  $P = (Fo^2 + 2Fc^2)/3$ atom sites solution primary ? atom sites solution secondary ? atom sites solution hydrogens refine ls hydrogen treatment undef refine ls extinction method none refine ls extinction coef refine ls abs structure details Refined as an inversion twin. refine ls abs structure Flack 0.48(5) chemical absolute configuration? refine ls number reflns 1852 refine ls number parameters 70 refine ls number restraints 0 refine ls R factor all 0.0379 refine ls R factor gt 0.0328 refine ls wR factor ref 0.0703 refine ls wR factor gt 0.0679 refine ls goodness of fit ref 1.039 refine ls restrained S all 1.039 refine ls shift/su max 0.000 refine ls shift/su mean 0.000 loop atom site label atom site type symbol atom site fract x atom site fract y atom site fract z

atom site U iso or equiv atom site adp type atom site occupancy atom site site symmetry order atom site calc flag atom site refinement flags posn atom site refinement flags adp atom site refinement flags occupancy atom site disorder assembly atom site disorder group Pr1 Pr 0.11806(2) 0.18538(2) 0.49185(8) 0.01392(10) Uani 1 1 d . . . . Si1 Si 0.24816(12) 0.10028(12) 0.0032(5) 0.0120(3) Uani 1 1 d . . . . Na1 Na 0.1054(2) 0.4110(2) 0.0020(9) 0.0236(5) Uani 1 1 d . . . . Na2A Na 0.0000 0.0000 0.101(3) 0.0239(18) Uiso 0.288(11) 2 d S . P . . Na2B Na 0.0000 0.0000 0.0000 0.0239(18) Uiso 0.424(11) 4 d S . P . . O1 O 0.0000 0.0000 0.5000 0.029(2) Uani 1 4 d S T P ... O2 O 0.2055(7) 0.0390(6) 0.2491(12) 0.0234(14) Uani 1 1 d . . . . O3 O 0.1068(3) 0.6173(3) 0.0044(15) 0.0215(9) Uani 1 1 d . . . . O4 O 0.2008(4) 0.2276(3) 0.0213(16) 0.0205(11) Uani 1 1 d . . . . . O5 O 0.3008(8) 0.4568(6) 0.2605(12) 0.0241(14) Uani 1 1 d . . . .

loop\_

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Na1 0.0188(11) 0.0212(11) 0.0309(15) -0.002(2) 0.003(2) 0.0001(9)
O1 0.014(2) 0.014(2) 0.059(8) 0.000 0.000 0.000
O2 0.040(5) 0.015(3) 0.015(3) 0.001(2) 0.011(3) -0.002(3)
O3 0.0168(18) 0.0147(17) 0.033(3) 0.001(3) -0.001(4) -0.0024(13)
O4 0.0149(17) 0.0103(16) 0.036(3) 0.000(3) 0.002(3) 0.0011(13)
O5 0.044(5) 0.015(3) 0.014(3) -0.001(2) 0.006(3) -0.001(3)
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\_geom\_special\_details

All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes. ;

loop \_geom\_bond\_atom site label 1 \_geom\_bond\_atom\_site\_label\_2 \_geom\_bond distance \_geom\_bond\_site\_symmetry 2 geom bond publ flag Pr1 O2 2.377(7) 4\_556 ? Pr1 O5 2.390(7) 7 455 ? Pr1 O4 2.429(4) 6 ? Pr1 O2 2.448(8) . ? Pr1 O5 2.460(8) 6 ? Pr1 O3 2.507(4) 7 455 ? Pr1 O1 2.6507(3) . ? Pr1 O4 2.806(8) . ? Pr1 Si1 3.0475(14) 6? Pr1 Si1 3.264(3) . ? Pr1 Si1 3.366(3) 1 556? Pr1 Na2A 3.403(12) . ? Si1 O5 1.607(7) 6 554 ? Si1 O2 1.618(7) . ? Si1 O3 1.624(4) 4 655 ? Si1 O4 1.641(4) . ? Sil Pr1 3.0475(14) 6 554 ? Sil Nal 3.164(3) 3? Si1 Na2B 3.2276(14) . ? Si1 Na1 3.251(5) 6? Si1 Na1 3.263(5) 6 554 ? Si1 Na2A 3.272(3) . ? Si1 Na2A 3.278(4) 3 ? Na1 O3 2.489(5) . ? Na1 O4 2.495(5) . ? Na1 O3 2.581(5) 2 565 ? Na1 O3 2.711(9) 7 455 ? Na1 O2 2.734(9) 6 554 ? Na1 O3 2.780(9) 7 454 ? Na1 O5 2.802(9) . ? Na1 Si1 3.164(3) 4? Nal Sil 3.251(5) 6 554? Na1 Si1 3.263(5) 6? Na1 Na1 3.327(5) 2 565 ? Na1 Pr1 3.532(2) 6 554 ? Na2A Na2B 0.554(18) . ? Na2A Na2A 1.11(4) 3? Na2A O1 2.179(18) . ?

Na2A O5 2.572(10) 7 455 ? Na2A O5 2.572(10) 8 545 ? Na2A O2 2.649(10) 2 ? Na2A O2 2.649(10) . ? Na2A Si1 3.272(3) 2 ? Na2A Si1 3.278(4) 3? Na2A Si1 3.278(4) 4 ? Na2A Pr1 3.403(12) 2 ? Na2B Na2A 0.554(18) 3 ? Na2B O1 2.73340(10) 1 554 ? Na2B O1 2.73340(10) . ? Na2B O5 2.785(9) 5 444 ? Na2B O5 2.785(9) 7 455 ? Na2B O5 2.785(9) 6 554 ? Na2B O5 2.785(9) 8 545 ? Na2B O2 2.867(8) 3 ? Na2B O2 2.867(8) 2 ? Na2B O2 2.867(8) 4 ? Na2B O2 2.867(8) . ? O1 Na2A 2.179(18) 3 556? O1 Pr1 2.6507(3) 2 ? O1 Pr1 2.6507(3) 4 556? O1 Pr1 2.6507(3) 3 556? O1 Na2B 2.73340(10) 1 556? O2 Pr1 2.377(7) 3 556 ? O2 Na1 2.734(9) 6? O3 Si1 1.624(4) 3 565 ? O3 Pr1 2.506(4) 8 ? O3 Na1 2.581(5) 2 565 ? O3 Na1 2.711(9) 8 ? O3 Na1 2.780(9) 8 554 ? O4 Pr1 2.429(4) 6 554 ? O5 Si1 1.607(7) 6 ? O5 Pr1 2.390(7) 8 ? O5 Pr1 2.460(8) 6 554 ? O5 Na2A 2.572(10) 7 ? O5 Na2B 2.785(9) 5 ? loop \_geom\_angle\_atom\_site label 1

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O5 Na2B O2 56.42(13) 6 554 . ? O5 Na2B O2 75.9(3) 8 545 . ? O2 Na2B O2 103.04(11) 3 . ? O2 Na2B O2 123.3(3) 2 . ? O2 Na2B O2 103.04(11) 4 . ? Na2A O1 Na2A 180.0 3 556 . ? Na2A O1 Pr1 90.963(10) 3 556.? Na2A O1 Pr1 89.037(10) . . ? Na2A O1 Pr1 90.963(10) 3 556 2 ? Na2A O1 Pr1 89.037(10) . 2 ? Pr1 O1 Pr1 178.074(19) . 2 ? Na2A O1 Pr1 89.037(10) 3 556 4 556 ? Na2A O1 Pr1 90.963(10) . 4 556? Pr1 O1 Pr1 90.0 . 4 556? Pr1 O1 Pr1 90.0 2 4 556 ? Na2A O1 Pr1 89.037(10) 3 556 3 556 ? Na2A O1 Pr1 90.963(10) . 3 556? Pr1 O1 Pr1 90.0 . 3 556? Pr1 O1 Pr1 90.0 2 3 556? Pr1 O1 Pr1 178.074(19) 4 556 3 556? Na2A O1 Na2B 180.0 . 1 556? Pr1 O1 Na2B 90.963(10) . 1 556? Pr1 O1 Na2B 90.963(10) 2 1 556 ? Pr1 O1 Na2B 89.037(10) 4 556 1 556? Pr1 O1 Na2B 89.037(10) 3 556 1 556 ? Na2A O1 Na2B 180.0 3 556 . ? Pr1 O1 Na2B 89.037(10) . . ? Pr1 O1 Na2B 89.037(10) 2 . ? Pr1 O1 Na2B 90.963(10) 4 556.? Pr1 O1 Na2B 90.963(10) 3 556.? Na2B O1 Na2B 180.0 1 556 . ? Si1 O2 Pr1 153.1(5) . 3 556? Si1 O2 Pr1 104.9(4) . . ? Pr1 O2 Pr1 102.0(2) 3 556.? Si1 O2 Na2A 97.2(5) . . ? Pr1 O2 Na2A 86.8(3) 3 556 . ? Pr1 O2 Na2A 83.7(4) . . ? Si1 O2 Na1 93.1(3) . 6? Pr1 O2 Na1 87.7(3) 3 556 6? Pr1 O2 Na1 85.8(2) . 6? Na2A O2 Na1 166.8(5) . 6 ? Si1 O2 Na2B 87.4(3) . . ? Pr1 O2 Na2B 93.7(2) 3 556.? Pr1 O2 Na2B 90.2(3) . . ? Na1 O2 Na2B 175.9(3) 6 . ? Si1 O3 Na1 176.4(3) 3 565 .?

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Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 zurloye@mailbox.sc.edu publ contact author email publ contact author phone +1-803-777-6916 \_publ\_contact\_author\_fax +1-803-777-8508 loop \_publ\_author\_name \_\_publ\_author\_address 'zur Loye, Hans-Conrad' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Latshaw, Allison M.' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Latshaw, Allison M.' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Chance, W. Michael' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Morrison, Gregory' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Smith, Mark D.' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208

'Yeon, Jeongho' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 audit creation method SHELXL-2013 chemical name systematic ? ? chemical name common chemical melting point ? chemical formula moiety 'H Na5 Nd4 O17 Si4' chemical formula sum 'H Na5 Nd4 O17 Si4' chemical formula weight 1077.28 loop atom type symbol atom type description atom type scat dispersion real atom type scat dispersion imag atom type scat source O O 0.0106 0.0060 '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' Na Na 0.0362 0.0249 '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' Nd Nd -0.1943 3.0179 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4' \_space\_group\_crystal\_system tetragonal

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\_shelx\_space\_group\_comment

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The symmetry employed for this shelxl refinement is uniquely defined by the following loop, which should always be used as a source of symmetry information in preference to the above space-group names. They are only intended as comments.

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'-V, X, -Z'		
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'-x+1/2, $-y+1/2$ , $z+1/2'$		
$\frac{1}{2}$ $\frac{1}{2}$ $\frac{1}{2}$ $\frac{1}{2}$ $\frac{1}{2}$ $\frac{1}{2}$		
-v+1/2, $x+1/2$ , $z+1/2$		
$y \in 1/2, X \in 1/2, Z \in 1/2$		
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volume /85.80(5)		
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\_reflns\_special\_details

## ;

Reflections were merged by SHELXL according to the crystal class for the calculation of statistics and refinement.

\_reflns\_Friedel\_fraction is defined as the number of unique Friedel pairs measured divided by the number that would be possible theoretically, ignoring centric projections and systematic absences.

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atom site U iso or equiv atom site adp type atom site occupancy atom site site symmetry order atom site calc flag atom site refinement flags posn atom site refinement flags adp atom site refinement flags occupancy atom site disorder assembly atom site disorder group Nd1 Nd 0.11766(2) 0.18506(2) 0.48281(4) 0.01047(6) Uani 1 1 d . . . . Si1 Si 0.24759(9) 0.10078(9) 0.0076(2) 0.00995(18) Uani 1 1 d . . . . Na1 Na 0.10534(15) 0.41080(16) 0.0063(5) 0.0221(4) Uani 1 1 d . . . . . Na2 Na 0.0000 0.0000 0.0714(14) 0.037(2) Uani 0.5 2 d S T P . . . O1 O 0.0000 0.0000 0.5000 0.0226(15) Uani 1 4 d S T P . . O2 O 0.2087(3) 0.0359(3) 0.2520(6) 0.0178(7) Uani 1 1 d . . . . O3 O 0.1066(2) 0.6165(2) 0.0125(7) 0.0173(6) Uani 1 1 d . . . . O4 O 0.2013(3) 0.2288(2) 0.0441(6) 0.0137(5) Uani 1 1 d . . . . O5 O 0.3063(3) 0.4536(3) 0.2653(6) 0.0177(6) Uani 1 1 d . . . .

loop\_

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O3 0.0137(12) 0.0117(11) 0.0266(16) 0.0002(13) -0.0027(14) -0.0015(9)
O4\ 0.0143(12)\ 0.0075(11)\ 0.0192(13)\ -0.0012(11)\ 0.0019(11)\ 0.0009(9)
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All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes. ;

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Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 zurloye@mailbox.sc.edu publ contact author email publ contact author phone +1-803-777-6916 \_publ\_contact\_author\_fax +1-803-777-8508 loop \_publ\_author\_name \_\_publ\_author\_address 'zur Loye, Hans-Conrad' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Latshaw, Allison M.' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Chance, W. Michael' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Morrison, Gregory' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Smith, Mark D.' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Yeon, Jeongho' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208

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\_reflns\_special\_details

Reflections were merged by SHELXL according to the crystal class for the calculation of statistics and refinement.

\_reflns\_Friedel\_fraction is defined as the number of unique Friedel pairs measured divided by the number that would be possible theoretically, ignoring centric projections and systematic absences.

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\_computing\_data\_collection'SMART Version 5.625 (Bruker, 2001)'\_computing\_cell\_refinement'SAINT-Plus Version 6.45 (Bruker, 2001)'\_computing\_data\_reduction'SAINT-Plus Version 6.45 (Bruker, 2001)'\_computing\_structure\_solution'SHELXS97 (Sheldrick, 1997)'\_computing\_molecular\_graphicsvariouscomputing\_publicationmaterial various

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Refined as a 2-component inversion twin.

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loop\_

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\_geom\_special\_details

All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

loop

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O2 Na1 2.692(7) 6? O3 Si1 1.636(5) 3 565 ? O3 Sm1 2.461(5) 8 ? O3 Na1 2.535(5) 2 565 ? O3 Na1 2.626(7) 8 ? O3 Na1 2.881(7) 8 554 ? O4 Sm1 2.400(5) 6 554 ? O5 Si1 1.619(6) 6 ? O5 Sm1 2.381(5) 8 ? O5 Sm1 2.424(6) 6 554 ? O5 Na2 2.491(7) 7 ? O5 Na2 2.870(9) 5 ? loop geom angle atom site label 1 geom angle atom site label 2 \_geom\_angle\_atom\_site\_label\_3 \_geom angle geom angle site symmetry 1 geom angle site symmetry 3 geom angle publ flag O2 Sm1 O5 73.62(18) 4 556 7 455 ? O2 Sm1 O4 122.0(2) 4 556 6 ? O5 Sm1 O4 145.29(19) 7 455 6 ? O2 Sm1 O2 138.9(3) 4 556 . ? O5 Sm1 O2 95.86(19) 7 455.? O4 Sm1 O2 89.45(19) 6 . ? O2 Sm1 O5 87.1(2) 4 556 6? O5 Sm1 O5 133.3(3) 7 455 6? O4 Sm1 O5 80.80(19) 6 6 ? O2 Sm1 O5 71.37(18).6? O2 Sm1 O3 82.1(2) 4 556 7 455 ? O5 Sm1 O3 88.3(2) 7 455 7 455 ? O4 Sm1 O3 66.07(16) 6 7 455 ? O2 Sm1 O3 138.27(18).7 455? O5 Sm1 O3 131.42(19) 67 455? O2 Sm1 O4 146.37(18) 4 556 . ? O5 Sm1 O4 78.57(17) 7 455 .? O4 Sm1 O4 73.85(11) 6.? O2 Sm1 O4 61.89(16) . . ? O5 Sm1 O4 126.18(18) 6 . ? O3 Sm1 O4 78.51(17) 7 455 .? O2 Sm1 O1 70.31(16) 4 556.? O5 Sm1 O1 67.06(16) 7 455 . ? O4 Sm1 O1 144.96(12) 6 . ? O2 Sm1 O1 69.05(14) . . ?

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631 Sumter Street Columbia, SC 29208 'Latshaw, Allison M.' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Chance, W. Michael' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Morrison, Gregory' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Smith, Mark D.' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Yeon, Jeongho' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 audit creation method SHELXL-2013 \_chemical\_name\_systematic , ? \_chemical\_name common ? \_chemical\_melting\_point ? chemical formula moiety ? \_chemical\_formula\_sum 'Eu4 H Na5 O17 Si4' chemical formula weight 1108.16

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\_reflns\_special\_details

Reflections were merged by SHELXL according to the crystal class for the calculation of statistics and refinement.

\_reflns\_Friedel\_fraction is defined as the number of unique Friedel pairs measured divided by the number that would be possible theoretically, ignoring centric projections and systematic absences.

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atom site site symmetry order

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atom site calc flag

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## \_geom\_special\_details

All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

loop

geom bond atom site label 1 geom bond atom site label 2 geom bond distance geom bond site symmetry 2 geom bond publ flag Eu1 O2 2.321(4) 4 556 ? Eu1 O5 2.373(4) 7 455 ? Eu1 O4 2.383(4) 6 ? Eu1 O2 2.408(4) . ? Eu1 O5 2.410(4) 6 ? Eu1 O3 2.456(3) 7 455 ? Eu1 O4 2.572(4) . ? Eu1 O1 2.5811(2) . ? Eu1 Si1 2.9877(13) 6? Eu1 Si1 3.1380(16) . ? Eu1 Na2 3.402(5) . ? Eu1 Si1 3.4382(17) 1 556? Si1 O5 1.625(4) 6 554 ? Si1 O2 1.628(4).? Si1 O3 1.643(4) 4 655 ? Si1 O4 1.653(4) . ? Sil Eul 2.9877(13) 6 554? Si1 Na1 3.111(3) 3?

Si1 Na2 3.1674(15) . ? Si1 Na2 3.1823(17) 3? Si1 Na1 3.240(3) 6? Sil Nal 3.267(3) 6 554? Si1 Eu1 3.4382(17) 1 554 ? Na1 O3 2.397(4) . ? Na1 O4 2.407(4) . ? Na1 O3 2.514(4) 2 565 ? Na1 O3 2.629(6) 7 455 ? Na1 O2 2.692(5) 6 554 ? Na1 O5 2.856(5) . ? Na1 O3 2.873(6) 7 454? Na1 Si1 3.111(3) 4? Na1 Si1 3.240(3) 6 554 ? Na1 Na1 3.242(4) 2 565 ? Na1 Si1 3.267(3) 6? Na1 Eu1 3.468(2) 6 554 ? Na2 Na2 0.800(16) 3? Na2 O1 2.341(8) . ? Na2 O5 2.462(5) 7 455 ? Na2 O5 2.462(5) 8 545 ? Na2 O2 2.671(6) 2 ? Na2 O2 2.671(6) . ? Na2 O5 2.846(6) 5 444 ? Na2 O5 2.846(6) 6 554 ? Na2 Si1 3.1674(15) 2 ? Na2 Si1 3.1823(17) 3 ? Na2 Si1 3.1823(17) 4? O1 Na2 2.341(8) 3 556? O1 Eu1 2.5811(2) 2 ? O1 Eu1 2.5811(2) 4 556 ? O1 Eu1 2.5811(2) 3 556? O2 Eu1 2.321(4) 3 556 ? O2 Na1 2.692(5) 6? O3 Si1 1.643(4) 3 565 ? O3 Eu1 2.456(3) 8 ? O3 Na1 2.515(4) 2 565 ? O3 Na1 2.629(6) 8 ? O3 Na1 2.873(6) 8 554 ? O4 Eu1 2.383(4) 6 554 ? O5 Si1 1.625(4) 6 ? O5 Eu1 2.373(4) 8 ? O5 Eu1 2.410(4) 6 554 ? O5 Na2 2.462(5) 7 ? O5 Na2 2.846(6) 5 ?

loop\_ \_geom\_angle\_atom\_site\_label\_1 \_geom\_angle\_atom\_site\_label\_2 \_geom\_angle\_atom\_site\_label\_3

\_geom\_angle atom site label 3 geom angle geom angle site symmetry 1 \_geom\_angle\_site\_symmetry 3 geom angle publ flag O2 Eu1 O5 73.98(14) 4 556 7 455 ? O2 Eu1 O4 121.45(15) 4 556 6? O5 Eu1 O4 146.29(14) 7 455 6? O2 Eu1 O2 138.22(18) 4 556.? O5 Eu1 O2 95.29(14) 7 455 .? O4 Eu1 O2 90.00(14) 6 . ? O2 Eu1 O5 86.04(15) 4 556 6? O5 Eu1 O5 132.61(18) 7 455 6? O4 Eu1 O5 80.58(14) 6 6 ? O2 Eu1 O5 71.77(14) . 6 ? O2 Eu1 O3 82.06(14) 4 556 7 455 ? O5 Eu1 O3 88.61(14) 7 455 7 455 ? O4 Eu1 O3 66.54(12) 6 7 455 ? O2 Eu1 O3 138.94(14). 7 455? O5 Eu1 O3 131.26(14) 6 7 455 ? O2 Eu1 O4 146.81(13) 4 556.? O5 Eu1 O4 78.90(13) 7 455 .? O4 Eu1 O4 74.20(8) 6 . ? O2 Eu1 O4 62.41(12) . . ? O5 Eu1 O4 126.85(13) 6 . ? O3 Eu1 O4 78.47(13) 7 455 . ? O2 Eu1 O1 70.00(11) 4 556.? O5 Eu1 O1 66.68(11) 7 455 . ? O4 Eu1 O1 144.52(9) 6 . ? O2 Eu1 O1 68.73(10) . . ? O5 Eu1 O1 66.16(10) 6 . ? O3 Eu1 O1 146.56(8) 7 455 . ? O4 Eu1 O1 115.85(8) . . ? O2 Eu1 Si1 105.94(12) 4 556 6? O5 Eu1 Si1 117.18(11) 7 455 6? O4 Eu1 Si1 33.51(9) 6 6 ? O2 Eu1 Si1 114.61(10).6? O5 Eu1 Si1 109.45(10) 6 6 ? O3 Eu1 Si1 33.35(9) 7 455 6? O4 Eu1 Si1 70.13(9).6? O1 Eu1 Si1 173.86(4) . 6? O2 Eu1 Si1 157.67(11) 4 556.? O5 Eu1 Si1 86.88(10) 7 455 .?

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Columbia, SC 29208 'Morrison, Gregory' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Smith, Mark D.' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Yeon, Jeongho' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 audit creation method SHELXL-97 chemical name systematic ? \_chemical\_name\_common ? ? chemical melting point chemical formula moiety 'H Na5 O17 Si4 Tb4' chemical formula sum 'H Na5 O17 Si4 Tb4' chemical formula weight 1136.00 loop atom type symbol atom type description atom type scat dispersion real atom type scat dispersion imag atom type scat source Na Na 0.0362 0.0249 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4' O O 0.0106 0.0060 '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' Tb Tb -0.1723 4.1537 '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 'I -4'

symmetry space group name Hall 'I -4' loop symmetry equiv pos as xyz 'x, y, z' 'y, -x, -z' '-x, -y, z' '-y, x, -z' 'x+1/2, y+1/2, z+1/2' y+1/2, -x+1/2, -z+1/2''-x+1/2, -y+1/2, z+1/2' '-y+1/2, x+1/2, -z+1/2' \_cell\_length a 11.6845(7)\_cell\_length b 11.6845(7)cell length c 5.4574(6) \_cell\_angle\_alpha 90.00 cell angle beta 90.00 \_cell\_angle gamma 90.00 \_cell\_volume 745.09(10) \_cell\_formula units Z 2 \_cell\_measurement temperature 296(2) \_cell\_measurement\_reflns used 4450 cell measurement theta min 2.465 cell measurement theta max 37.770 \_exptl\_crystal\_description 'square block' exptl crystal colour colorless exptl crystal size max 0.06 \_exptl\_crystal\_size\_mid 0.06 \_exptl\_crystal size min 0.04 exptl crystal density meas ? exptl crystal density diffrn 5.064 exptl crystal density method 'not measured' \_exptl\_crystal F 000 1016 exptl absorpt coefficient mu 19.319 exptl absorpt correction type multi-scan \_exptl\_absorpt\_correction\_T\_min 0.607237 \_exptl\_absorpt\_correction T max 1.000000 exptl absorpt process details 'SADABS Version 2.05 (Bruker, 2001)' \_exptl\_special\_details

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diffrn ambient temperature 296(2)diffrn radiation wavelength 0.71073 diffrn radiation type MoK\a diffrn radiation source 'fine-focus sealed tube' diffrn radiation monochromator graphite diffrn measurement device type 'Bruker SMART APEX CCD diffractometer' diffrn measurement method 'omega and phi scans' diffrn detector area resol mean? diffrn standards number diffrn standards interval count? diffrn standards interval time ? diffrn standards decay % ? diffrn reflns number 10635 diffrn reflns av R equivalents 0.0467 diffrn reflns av sigmal/netI 0.0439 diffrn reflns limit h min -20 diffrn reflns limit h max 20 diffrn reflns limit k min -20 diffrn reflns limit k max 20 diffrn reflns limit 1 min -8 9 diffrn reflns limit 1 max diffrn reflns theta min 2.47 diffrn reflns theta max 38.57 reflns number total 2112 reflns number gt 2009 reflns threshold expression >2sigma(I) computing data collection 'SMART Version 5 625 (Bruker 2001)'

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_computing_publication_material 'SHELXTL Version 6.14 (Bruker, 2000)'			

\_refine\_special\_details

Crystal refined as inversion twin based on value of Flack parameter after anisotropic refinement of all located atoms. The Flack parameter was included in the refinement as the twin fraction. The final value is 0.507(17), indicating a nearly perfect two-component inversion twin.

The compound is assumed to be a hydroxide to satisfy charge balance requirements (crystal electroneutrality). The hydroxy proton could not be located in the X-ray data. It is assumed to be bonded to the non-silicate oxygen atom O(5).

Refinement of F<sup>2</sup> against ALL reflections. The weighted R-factor wR and goodness of fit S are based on F<sup>2</sup>, conventional R-factors R are based on F, with F set to zero for negative F<sup>2</sup>. The threshold expression of F<sup>2</sup> > 2sigma(F<sup>2</sup>) is used only for calculating R-factors(gt) etc. and is not relevant to the choice of reflections for refinement. R-factors based on F<sup>2</sup> 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 calc refine ls weighting details 'calc w=1/[ $s^2(Fo^2)+(0.0258P)^2+0.0000P$ ] where P=(Fo^2+2Fc^2)/3' atom sites solution primary direct atom sites solution secondary difmap atom sites solution hydrogens ? refine ls hydrogen treatment ? refine ls extinction method none refine ls extinction coef ? refine ls abs structure details 'Flack H D (1983), Acta Cryst. A39, 876-881' refine ls abs structure Flack ? refine ls number reflns 2112 refine ls number parameters 72 refine ls number restraints 0 refine ls R factor all 0.0334 refine ls R factor gt 0.0300 refine ls wR factor ref 0.0551 refine ls wR factor gt 0.0537 refine ls goodness of fit ref 1.030 refine ls restrained S all 1.030 refine ls shift/su max 0.000 refine ls shift/su mean 0.000 loop atom site label atom site type symbol atom site fract x atom site fract y atom site fract z atom site U iso or equiv atom site adp type

\_atom\_site\_occupancy

atom site symmetry multiplicity

\_atom\_site\_calc\_flag

\_atom\_site\_refinement\_flags \_atom\_site\_disorder\_assembly \_atom\_site\_disorder\_group Tb1 Tb 0.116610(15) 0.316098(15) 0.27318(3) 0.00900(5) Uani 1 1 d . . . Si1 Si 0.25264(10) 0.10251(9) 0.2386(3) 0.0109(2) Uani 1 1 d . . . Na1 Na 0.39557(16) 0.41057(16) 0.2416(5) 0.0210(4) Uani 1 1 d . . . Na2 Na 0.0000 0.5000 0.6759(12) 0.0272(15) Uani 0.50 2 d SP . . O1 O 0.1136(2) 0.1083(2) 0.2641(7) 0.0134(6) Uani 1 1 d . . . O2 O 0.3117(3) 0.0488(3) 0.4831(6) 0.0132(6) Uani 1 1 d . . . O3 O 0.2975(3) 0.2345(3) 0.1869(6) 0.0128(6) Uani 1 1 d . . . O4 O 0.2930(3) 0.0336(3) -0.0058(6) 0.0137(6) Uani 1 1 d . . . O5 O 0.0000 0.5000 0.2500 0.0150(13) Uani 1 4 d S . .

loop\_

atom site aniso label atom site aniso U 11 atom site aniso U 22 atom site aniso U 33 atom site aniso U 23 atom site aniso U 13 atom site aniso U 12 Tb1 0.00777(8) 0.00847(8) 0.01076(8) 0.00022(6) -0.00039(6) -0.00060(6) Si1 0.0119(4) 0.0101(4) 0.0107(6) 0.0010(5) -0.0001(5) 0.0008(4) Na1 0.0163(8) 0.0154(8) 0.0312(12) -0.0016(10) 0.0008(9) -0.0005(6) Na2 0.027(4) 0.023(3) 0.031(4) 0.000 0.000 0.004(3) O1 0.0113(11) 0.0089(11) 0.0199(17) -0.0008(13) -0.0020(13) -0.0004(9) O2 0.0210(16) 0.0097(14) 0.0088(14) 0.0010(11) -0.0056(12) 0.0014(13)  $O3\ 0.0139(13)\ 0.0104(12)\ 0.0141(15)\ -0.0006(11)\ 0.0019(12)\ 0.0016(10)$  $O4\ 0.0216(17)\ 0.0099(14)\ 0.0094(14)\ -0.0010(11)\ 0.0038(12)\ 0.0005(12)$ O5 0.0101(14) 0.0101(14) 0.025(4) 0.000 0.000 0.000

\_geom\_special\_details

All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

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loop\_ \_geom\_bond\_atom\_site\_label\_1 \_geom\_bond\_atom\_site\_label\_2 \_geom\_bond\_distance \_geom\_bond\_site\_symmetry\_2

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O2 Na2 Si1 84.84(8) 2 566 5 455 ?
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O2 Na2 Si1 30.41(8) 2 566 2 566 ?
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O4 Na2 Si1 81.05(8) 7 2 566 ?
O2 Na2 Si1 78.59(10) 7 2 566 ?
O2 Na2 Si1 91.31(12) 5 455 2 566?
Si1 Na2 Si1 89.08(3) 7 2 566 ?
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O2 Na2 Si1 30.41(8) 4 556 4 556?
O2 Na2 Si1 165.4(2) 2 566 4 556?
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O2 Na2 Si1 78.59(10) 5\_455 4\_556? Si1 Na2 Si1 89.08(3) 7 4 556 ? Si1 Na2 Si1 89.08(3) 5 455 4 556? Si1 Na2 Si1 163.0(2) 2 566 4 556? Si1 O1 Na1 173.4(2) . 6 455 ? Sil Ol Tbl 91.66(12)..? Na1 O1 Tb1 91.93(11) 6 455 .? Si1 O1 Na1 94.02(13) . 8 545 ? Na1 O1 Na1 82.41(11) 6 455 8 545? Tb1 O1 Na1 174.32(14) . 8 545 ? Sil Ol Nal 97.04(16).7? Na1 O1 Na1 88.31(14) 6 455 7 ? Tb1 O1 Na1 93.71(12).7? Na1 O1 Na1 85.64(13) 8 545 7 ? Si1 O1 Na1 87.08(16) . 7 554 ? Na1 O1 Na1 87.01(13) 6 455 7 554? Tb1 O1 Na1 95.61(12).7 554? Na1 O1 Na1 84.65(12) 8 545 7 554 ? Na1 O1 Na1 169.70(14) 7 7 554 ? Si1 O2 Tb1 142.3(2) . 2 556 ? Si1 O2 Tb1 115.96(17).7? Tb1 O2 Tb1 99.27(12) 2 556 7 ? Si1 O2 Na2 100.5(2) . 2 556 ? Tb1 O2 Na2 89.39(14) 2 556 2 556 ? Tb1 O2 Na2 94.28(16) 7 2 556 ? Si1 O2 Na2 85.51(18) . 5 544 ? Tb1 O2 Na2 98.80(13) 2 556 5 544 ? Tb1 O2 Na2 104.72(15) 7 5 544 ? Si1 O2 Na1 88.91(16) . 7 ? Tb1 O2 Na1 82.80(12) 2 556 7 ? Tb1 O2 Na1 81.92(11) 7 7 ? Na2 O2 Na1 170.6(2) 2 556 7 ? Na2 O2 Na1 172.73(17) 5 544 7 ? Si1 O3 Tb1 93.36(14) . . ? Si1 O3 Na1 160.1(2) . . ? Tb1 O3 Na1 93.25(12) . . ? Si1 O3 Tb1 93.44(14) . 7 554 ? Tb1 O3 Tb1 128.55(14).7 554? Na1 O3 Tb1 97.05(13) . 7 554 ? Si1 O4 Tb1 158.8(2) . 2 ? Si1 O4 Tb1 100.30(16) . 7 554 ? Tb1 O4 Tb1 100.94(13) 2 7 554 ? Si1 O4 Na2 91.8(2) . 5 544 ? Tb1 O4 Na2 91.00(15) 2 5 544 ? Tb1 O4 Na2 83.89(14) 7 554 5 544 ? Si1 O4 Na1 93.60(17) . 7 554 ?

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631 Sumter Street Columbia, SC 29208 'Latshaw, Allison M.' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Chance, W. Michael' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Morrison, Gregory' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Smith, Mark D.' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Yeon, Jeongho' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 audit creation method SHELXL-97 \_chemical\_name\_systematic

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computing cell refinement 'SAINT-Plus Version 6.45 (Bruker, 2001)'

computing data reduction 'SAINT-Plus Version 6.45 (Bruker, 2001)'

computing structure solution 'SHELXS-97 (Sheldrick, 1990)'

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computing molecular graphics various

computing publication material ?

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Refinement of F<sup>2</sup> against ALL reflections. The weighted R-factor wR and goodness of fit S are based on F<sup>2</sup>, conventional R-factors R are based on F, with F set to zero for negative  $F^2$ . The threshold expression of  $F^2 > 2$ sigma( $F^2$ ) is used only for calculating R-factors(gt) etc. and is not relevant to the choice of reflections for refinement. R-factors based on F<sup>2</sup> are statistically about twice as large as those based on F, and Rfactors based on ALL data will be even larger.

;

refine ls structure factor coef Fsqd

refine ls matrix type full \_refine\_ls\_weighting\_scheme calc

\_refine\_ls\_weighting\_details

 $\overline{\text{calc w=1/[}} = 1/[\sqrt{6^2}) + (0.0362P)^2 + 0.4530P]$  where P=(Fo^2^+2Fc^2)/3'

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\_atom\_site\_aniso\_U\_12

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All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

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O3 Na1 2.826(5) 8\_554 ? O4 Dy1 2.356(4) 7\_554 ? O5 Si1 1.626(4) 7 ? O5 Dy1 2.331(4) 8 ? O5 Dy1 2.372(4) 7\_554 ? O5 Na2 2.451(5) 6 ? O5 Na2 2.782(7) 5 ?

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'Morrison, Gregory' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 ; 'Smith, Mark D.' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208

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Refinement of F<sup>2</sup> against ALL reflections. The weighted R-factor wR and goodness of fit S are based on F<sup>2</sup>, conventional R-factors R are based on F, with F set to zero for negative F<sup>2</sup>. The threshold expression of  $F^2^> 2 \text{sigma}(F^2^>)$  is used only for calculating R-factors(gt) etc. and is not relevant to the choice of reflections for refinement. R-factors based on F<sup>2</sup> are statistically about twice as large as those based on F, and Rfactors based on ALL data will be even larger.

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All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

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O2 Ho1 O5 71.76(14) . 7 ?
O2 Ho1 O3 82.12(15) 4 556 6 455 ?
O5 Ho1 O3 88.73(14) 6 455 6 455 ?
O4 Ho1 O3 67.47(12) 7 6 455 ?
O2 Ho1 O3 140.29(13) . 6 455 ?
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O2 Ho1 O1 69.31(11) 4 556 . ?
O5 Ho1 O1 66.47(10) 6 455.?
O4 Ho1 O1 143.45(9) 7 . ?
O2 Ho1 O1 68.11(10) . . ?
O5 Ho1 O1 65.94(9) 7 . ?
O3 Ho1 O1 146.06(8) 6 455.?
O4 Ho1 O1 117.18(8) . . ?
O2 Ho1 Si1 106.29(12) 4 5567?
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O5 Ho1 Si1 117.80(10) 6 455 7 ? O4 Ho1 Si1 34.08(9) 7 7 ? O2 Ho1 Si1 115.42(10) . 7 ? O5 Ho1 Si1 109.16(10) 77? O3 Ho1 Si1 33.87(9) 6 455 7 ? O4 Ho1 Si1 69.31(9) . 7 ? O1 Ho1 Si1 173.30(3).7? O2 Ho1 Si1 157.66(11) 4 556.? O5 Ho1 Si1 87.13(10) 6 455.? O4 Ho1 Si1 81.72(10) 7 . ? O2 Ho1 Si1 31.59(10) . . ? O5 Ho1 Si1 99.74(10) 7 . ? O3 Ho1 Si1 109.72(11) 6 455 .? O4 Ho1 Si1 32.44(8) . . ? O1 Ho1 Si1 92.53(2) . . ? Sil Hol Sil 92.85(4) 7 . ? O2 Ho1 Na2 100.58(14) 4 556.? O5 Ho1 Na2 46.21(11) 6 455 .? O4 Ho1 Na2 138.27(12) 7 . ? O2 Ho1 Na2 50.68(11) . . ? O5 Ho1 Na2 99.16(13) 7 . ? O3 Ho1 Na2 129.74(14) 6 455 .? O4 Ho1 Na2 73.61(15) ...? O1 Ho1 Na2 44.54(13) . . ? Sil Hol Na2 142.16(13) 7 . ? Sil Hol Na2 57.19(10) . . ? O2 Ho1 Si1 79.59(11) 4 556 1 556? O5 Ho1 Si1 147.93(10) 6 455 1 556? O4 Ho1 Si1 62.70(10) 7 1 556? O2 Ho1 Si1 91.93(11) . 1 556? O5 Ho1 Si1 25.52(9) 7 1 556 ? O3 Ho1 Si1 105.38(11) 6 455 1 556? O4 Ho1 Si1 130.88(9) . 1 556 ? O1 Ho1 Si1 87.56(2) . 1 556? Si1 Ho1 Si1 86.63(4) 7 1 556 ? Si1 Ho1 Si1 113.63(4) . 1 556? Na2 Ho1 Si1 124.61(10) . 1 556? O5 Si1 O2 111.0(2) 7 554.? O5 Si1 O3 111.3(2) 7 554 4 655 ? O2 Si1 O3 112.3(2) . 4 655 ? O5 Si1 O4 112.0(2) 7 554.? O2 Si1 O4 103.4(2) . . ? O3 Si1 O4 106.4(2) 4 655 . ? O5 Si1 Ho1 119.50(16) 7 554 7 554 ? O2 Si1 Ho1 129.10(16) . 7 554? O3 Si1 Ho1 54.62(12) 4 655 7 554?

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Si1 Na1 Na1 78.95(8) 4 3 565 ? O4 Na1 Si1 79.27(12) . 7 554 ? O3 Na1 Si1 92.97(14). 7 554? O3 Na1 Si1 123.20(15) 3 565 7 554? O3 Na1 Si1 144.19(11) 6 455 7 554 ? O2 Na1 Si1 30.73(9) 7 554 7 554 ? O3 Na1 Si1 30.90(8) 6 454 7 554? O5 Na1 Si1 87.85(10) . 7 554 ? Si1 Na1 Si1 111.71(8) 4 7 554? Na1 Na1 Si1 116.35(6) 3 565 7 554? O4 Na1 Si1 65.58(12).7? O3 Na1 Si1 91.23(13) . 7 ? O3 Na1 Si1 120.72(15) 3 565 7 ? O3 Na1 Si1 30.17(9) 6 455 7 ? O2 Na1 Si1 89.79(11) 7 554 7 ? O3 Na1 Si1 145.00(11) 6 4547? O5 Na1 Si1 30.02(9). 7? Si1 Na1 Si1 114.36(9) 47? Na1 Na1 Si1 115.89(6) 3 565 7? Si1 Na1 Si1 114.64(7) 7 554 7 ? O4 Na1 Ho1 43.44(10) . 7 554 ? O3 Na1 Ho1 107.64(11) . 7 554 ? O3 Na1 Ho1 154.81(11) 3 565 7 554? O3 Na1 Ho1 89.08(10) 6 455 7 554? O2 Na1 Ho1 43.45(9) 7 554 7 554 ? O3 Na1 Ho1 83.97(9) 6 454 7 554 ? O5 Na1 Ho1 43.23(8) . 7 554 ? Si1 Na1 Ho1 122.56(7) 4 7 554 ? Na1 Na1 Ho1 158.19(11) 3 565 7 554? Si1 Na1 Ho1 55.41(4) 7 554 7 554 ? Sil Nal Hol 61.31(5) 7 7 554? Na2 Na2 O1 180.0 2 . ? Na2 Na2 O5 112.8(2) 2 6 455 ? O1 Na2 O5 67.2(2) . 6 455 ? Na2 Na2 O5 112.8(2) 2 8 545 ? O1 Na2 O5 67.2(2) . 8 545 ? O5 Na2 O5 134.5(5) 6 455 8 545 ? Na2 Na2 O2 113.7(2) 2 3 ? O1 Na2 O2 66.3(2) . 3 ? O5 Na2 O2 76.22(19) 6 455 3 ? O5 Na2 O2 85.8(2) 8 545 3 ? Na2 Na2 O2 113.7(2) 2 . ? O1 Na2 O2 66.3(2) . . ? O5 Na2 O2 85.8(2) 6 455 . ? O5 Na2 O2 76.22(19) 8 545 . ? O2 Na2 O2 132.5(5) 3 . ?

Na2 Na2 O5 54.34(19) 2 5 444 ? O1 Na2 O5 125.66(19) . 5 444 ? O5 Na2 O5 103.04(10) 6 455 5 444 ? O5 Na2 O5 103.04(10) 8 545 5 444 ? O2 Na2 O5 59.63(12) 3 5 444 ? O2 Na2 O5 167.1(4) . 5 444 ? Na2 Na2 O5 54.34(19) 2 7 554 ? O1 Na2 O5 125.66(19).7 554? O5 Na2 O5 103.04(10) 6 455 7 554 ? O5 Na2 O5 103.04(10) 8 545 7 554 ? O2 Na2 O5 167.1(4) 3 7 554 ? O2 Na2 O5 59.63(12) . 7 554 ? O5 Na2 O5 108.7(4) 5 444 7 554 ? Na2 Na2 O2 54.29(18) 2 2 ? O1 Na2 O2 125.71(18) . 2 ? O5 Na2 O2 166.2(4) 6 455 2 ? O5 Na2 O2 58.71(14) 8 545 2 ? O2 Na2 O2 103.59(10) 3 2 ? O2 Na2 O2 103.59(10) . 2 ? O5 Na2 O2 66.12(19) 5 444 2 ? O5 Na2 O2 74.0(2) 7 554 2 ? Na2 Na2 O2 54.29(18) 2 4 ? O1 Na2 O2 125.71(18) . 4 ? O5 Na2 O2 58.71(14) 6 455 4 ? O5 Na2 O2 166.2(4) 8 545 4 ? O2 Na2 O2 103.59(10) 3 4 ? O2 Na2 O2 103.59(10) . 4 ? O5 Na2 O2 74.0(2) 5 444 4 ? O5 Na2 O2 66.12(19) 7 554 4 ? O2 Na2 O2 108.6(4) 2 4 ? Na2 Na2 Si1 84.91(19) 2.? O1 Na2 Si1 95.09(19) . . ? O5 Na2 Si1 84.54(10) 6 455 . ? O5 Na2 Si1 99.42(11) 8 545 .? O2 Na2 Si1 157.2(3) 3 . ? O2 Na2 Si1 31.77(9) . . ? O5 Na2 Si1 138.5(4) 5 444 . ? O5 Na2 Si1 31.47(9) 7 554.? O2 Na2 Si1 98.04(17) 2.? O2 Na2 Si1 75.91(13) 4 . ? Na2 Na2 Si1 84.91(19) 2 3 ? O1 Na2 Si1 95.09(19) . 3? O5 Na2 Si1 99.42(11) 6 455 3 ? O5 Na2 Si1 84.54(10) 8 545 3? O2 Na2 Si1 31.77(9) 3 3 ? O2 Na2 Si1 157.2(3).3?

O5 Na2 Si1 31.47(9) 5 444 3 ? O5 Na2 Si1 138.5(4) 7 554 3 ? O2 Na2 Si1 75.91(13) 2 3 ? O2 Na2 Si1 98.04(17) 4 3 ? Si1 Na2 Si1 169.8(4) . 3 ? Na2 O1 Na2 180.000(1) . 2 556? Na2 O1 Ho1 87.033(5) . . ? Na2 O1 Ho1 92.967(5) 2 556 . ? Na2 O1 Ho1 87.034(5).3? Na2 O1 Ho1 92.966(5) 2 556 3 ? Ho1 O1 Ho1 174.067(10) . 3 ? Na2 O1 Ho1 92.967(5) . 2 556? Na2 O1 Ho1 87.033(5) 2 556 2 556 ? Ho1 O1 Ho1 90.2 . 2 556 ? Ho1 O1 Ho1 90.2 3 2 556? Na2 O1 Ho1 92.967(5) . 4 556? Na2 O1 Ho1 87.033(5) 2 556 4 556 ? Ho1 O1 Ho1 90.153(1) . 4 556? Ho1 O1 Ho1 90.153(1) 3 4 556 ? Ho1 O1 Ho1 174.067(9) 2 556 4 556 ? Si1 O2 Ho1 158.8(2) . 2 556 ? Si1 O2 Ho1 99.41(18) . . ? Ho1 O2 Ho1 101.72(16) 2 556 . ? Si1 O2 Na2 90.4(3) . . ? Ho1 O2 Na2 93.04(19) 2 556 . ? Ho1 O2 Na2 85.39(18) . . ? Sil O2 Nal 92.57(19).7? Ho1 O2 Na1 87.39(15) 2 556 7 ? Ho1 O2 Na1 85.22(13) . 7 ? Na2 O2 Na1 170.5(2) . 7 ? Si1 O2 Na2 79.8(2) . 2 ? Ho1 O2 Na2 101.05(18) 2 556 2 ? Ho1 O2 Na2 92.47(17) . 2 ? Na2 O2 Na2 12.0(4) . 2 ? Na1 O2 Na2 171.54(19) 7 2 ? Si1 O3 Na1 173.3(3) 2 565 . ? Si1 O3 Ho1 91.50(15) 2 565 8 ? Na1 O3 Ho1 92.39(13) . 8 ? Si1 O3 Na1 93.68(16) 2 565 3 565 ? Na1 O3 Na1 82.46(13) . 3 565 ? Ho1 O3 Na1 174.81(17) 8 3 565 ? Si1 O3 Na1 96.43(19) 2 565 8 ? Na1 O3 Na1 88.80(17) . 8 ? Ho1 O3 Na1 93.57(14) 8 8 ? Na1 O3 Na1 85.78(16) 3 565 8? Si1 O3 Na1 87.20(19) 2 565 8 554?

Na1 O3 Na1 86.93(16) . 8\_554 ? Ho1 O3 Na1 96.11(15) 8 8 554 ? Na1 O3 Na1 84.24(15) 3 565 8 554 ? Na1 O3 Na1 169.58(17) 8 8 554 ? Si1 O4 Na1 159.1(3) . . ? Si1 O4 Ho1 93.16(17).7 554? Na1 O4 Ho1 93.28(13) . 7 554 ? Si1 O4 Ho1 93.16(17) . . ? Na1 O4 Ho1 98.20(15) . . ? Ho1 O4 Ho1 128.82(17) 7 554 .? Si1 O5 Ho1 142.0(2) 7 8 ? Si1 O5 Ho1 115.74(19) 7 7 554 ? Ho1 O5 Ho1 100.01(15) 8 7 554 ? Si1 O5 Na2 98.0(3) 7 6 ? Ho1 O5 Na2 90.42(19) 8 6 ? Ho1 O5 Na2 95.7(2) 7 554 6? Si1 O5 Na2 85.6(2) 7 5 ? Ho1 O5 Na2 98.16(17) 8 5 ? Ho1 O5 Na2 104.37(18) 7 554 5 ? Na2 O5 Na2 12.9(4) 6 5 ? Sil O5 Nal 89.58(18) 7.? Ho1 O5 Na1 82.92(13) 8.? Ho1 O5 Na1 81.76(13) 7 554.? Na2 O5 Na1 172.3(3) 6 . ? Na2 O5 Na1 173.4(2) 5 . ?

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\_diffrn\_reflns\_theta\_full 36.38

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\_refine\_diff\_density\_max 2.348

\_refine\_diff\_density\_min -1.398

refine diff density rms 0.339

#===END

data\_Na5Er4(OH)[SiO4]4 \_publ\_requested\_journal Inorg.Chem. \_publ\_contact\_author\_name 'Hans-Conrad zur Loye' \_publ\_contact\_author\_address

;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 publ contact author email zurloye@mailbox.sc.edu \_publ\_contact\_author\_phone +1-803-777-6916 publ contact author fax +1-803-777-8508 loop publ author name \_publ\_author address 'zur Loye, Hans-Conrad' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Latshaw, Allison M.' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Chance, W. Michael' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Morrison, Gregory' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Smith, Mark D.' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Yeon, Jeongho' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208

audit creation method SHELXL-2013 chemical name systematic ? chemical name common ? \_chemical melting point ? chemical formula moiety 'Er4 H Na5 O17 Si4' chemical formula sum 'Er4 H Na5 O17 Si4' chemical formula weight 1169.36 loop atom type symbol atom type description atom type scat dispersion real \_atom\_type\_scat\_dispersion imag \_atom\_type scat source O O 0.0106 0.0060 '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' Na Na 0.0362 0.0249 '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' Er Er -0.2586 4.9576 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4' space group crystal system tetragonal 82

\_space\_group\_erystar\_system tetragona \_space\_group\_IT\_number 82 \_space\_group\_name\_H-M\_alt 'I -4' \_space\_group\_name\_Hall 'I -4'

\_shelx\_space\_group\_comment

The symmetry employed for this shelxl refinement is uniquely defined by the following loop, which should always be used as a source of symmetry information in preference to the above space-group names. They are only intended as comments.

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loop\_ \_space\_group\_symop\_operation\_xyz 'x, y, z' '-x, -y, z' 'y, -x, -z' '-y, x, -z' 'x+1/2, y+1/2, z+1/2' '-x+1/2, -y+1/2, z+1/2'

'y+1/2, -x+1/2, -z+1/2' '-y+1/2, x+1/2, -z+1/2' \_cell\_length a 11.5731(3) cell length b 11.5731(3)cell length c 5.4107(2) \_cell\_angle alpha 90 cell angle beta 90 \_cell\_angle gamma 90 cell volume 724.69(5) cell formula units Z 2 \_cell\_measurement temperature 296(2)cell measurement reflns used 4436 \_cell\_measurement theta min 2.489 cell measurement theta max 33.100 \_exptl\_crystal description plate exptl crystal colour pink exptl crystal density meas \_exptl\_crystal\_density method ? exptl crystal density diffrn 5.359 \_exptl\_crystal F 000 1040 exptl transmission factor min ? exptl transmission factor max ? \_exptl\_crystal\_size max 0.060 exptl crystal size mid 0.050 exptl crystal size min 0.050 exptl absorpt coefficient mu 23.506 shelx estimated absorpt T min 0.333 shelx estimated absorpt T max 0.386 \_exptl\_absorpt\_correction type multi-scan exptl absorpt correction T min 0.5819 exptl absorpt correction T max 1.0000 exptl absorpt process details 'SADABS Version 2.05 (Bruker, 2001)' \_exptl\_special details ? diffrn ambient temperature 296(2)diffrn radiation wavelength 0.71073 diffrn radiation type MoK\a diffrn source diffrn measurement device type 'Bruker SMART APEX CCD diffractometer' diffrn measurement method 'omega and phi scans'

diffrn detector area resol mean? diffrn reflns number 10726 diffrn reflns av unetI/netI 0.0286 diffrn reflns av R equivalents 0.0369 diffrn reflns limit h min -19 diffrn reflns limit h max 19 diffrn reflns limit k min -19 diffrn reflns limit k max 17 diffrn reflns limit 1 min -8 diffrn reflns limit 1 max 8 diffrn reflns theta min 2.489 diffrn reflns theta max 36.281 diffrn reflns theta full 25.242 diffrn measured fraction theta max 1.000 diffrn measured fraction theta full 0.997 diffrn reflns Laue measured fraction max 1.000 diffrn reflns Laue measured fraction full 0.997 diffrn reflns point group measured fraction max 1.000 diffrn reflns point group measured fraction full 0.998 reflns number total 1652 reflns number gt 1611 reflns threshold expression 'I > 2 (I)'reflns Friedel coverage 0.841 reflns Friedel fraction max 1.000 reflns Friedel fraction full 1.000

\_reflns\_special\_details

Reflections were merged by SHELXL according to the crystal class for the calculation of statistics and refinement.

\_reflns\_Friedel\_fraction is defined as the number of unique Friedel pairs measured divided by the number that would be possible theoretically, ignoring centric projections and systematic absences.

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\_computing\_data\_collection'SMART Version 5.625 (Bruker, 2001)'\_computing\_cell\_refinement'SAINT-Plus Version 6.45 (Bruker, 2001)'\_computing\_data\_reduction'SAINT-Plus Version 6.45 (Bruker, 2001)'\_computing\_structure\_solution'SHELXS97 (Sheldrick, 1997)'\_computing\_molecular\_graphicsvarious\_computing\_publication\_material'SHELXTL Version 6.14 (Bruker, 2000)'

\_refine\_special\_details

Refined as a 2-component inversion twin. refine ls structure factor coef Fsqd refine ls matrix type full refine ls weighting scheme calc refine ls weighting details  $w=1/[\sqrt{6^2}(6^2)+(0.0395P)^2+3.8969P]$ where  $P = (Fo^{2} + 2Fc^{2})/3$ atom sites solution primary ? atom sites solution secondary ? atom sites solution hydrogens refine ls hydrogen treatment undef refine ls extinction method none refine ls extinction coef refine ls abs structure details Refined as an inversion twin. refine ls abs structure Flack 0.44(3) chemical absolute configuration? refine ls number reflns 1652 refine ls number parameters 69 refine ls number restraints 0 refine ls R factor all 0.0292 refine ls R factor gt 0.0279 refine ls wR factor ref 0.0660 refine ls wR factor gt 0.0655 refine ls goodness of fit ref 1.066 refine ls restrained S all 1.066 refine ls shift/su max 0.000 refine ls shift/su mean 0.000 loop atom site label atom site type symbol atom site fract x atom site fract y atom site fract z atom site U iso or equiv \_atom\_site\_adp type atom site occupancy atom site site symmetry order atom site calc flag

\_atom\_site\_refinement\_flags\_posn \_atom\_site\_refinement\_flags\_adp \_atom\_site\_refinement\_flags\_occupancy \_atom\_site\_disorder\_assembly \_atom\_site\_disorder\_group Er1 Er 0.11786(3) 0.18444(3) 0.47556(6) 0.01096(9) Uani 1 1 d . . . . . Si1 Si 0.2457(2) 0.1024(2) 0.0111(6) 0.0166(4) Uani 1 1 d . . . . . Na1 Na 0.1043(3) 0.4110(3) 0.0044(9) 0.0215(8) Uani 1 1 d . . . . . Na2 Na 0.0000 0.0000 0.055(3) 0.033(3) Uiso 0.5 2 d S . P . . O1 O 0.0000 0.0000 0.5000 0.031(4) Uani 1 4 d S T P . . O2 O 0.2026(6) 0.0319(5) 0.2571(13) 0.0174(12) Uani 1 1 d . . . . . O3 O 0.1086(5) 0.6124(5) 0.0097(13) 0.0162(11) Uani 1 1 d . . . . O4 O 0.2006(5) 0.2374(5) 0.0649(13) 0.0157(11) Uani 1 1 d . . . . . O5 O 0.3128(6) 0.4518(5) 0.2608(12) 0.0153(11) Uani 1 1 d . . . .

loop\_

\_atom\_site\_aniso\_label \_atom\_site\_aniso\_U\_11 \_atom\_site\_aniso\_U\_22 \_atom\_site\_aniso\_U\_33 \_atom\_site\_aniso\_U\_23 \_atom\_site\_aniso\_U\_13 \_atom\_site\_aniso\_U\_12 Er1 0.00886(13) 0.01120(14) 0.01282(14) 0.00113(11) 0.00076(10) 0.00226(10) Si1 0.0180(9) 0.0142(9) 0.0176(12) -0.0002(10) 0.0003(9) -0.0010(7) Na1 0.0181(15) 0.0130(13) 0.033(2) 0.0014(17) 0.0023(16) -0.0001(11) O1 0.023(4) 0.023(4) 0.047(12) 0.000 0.000 0.000 O2 0.025(3) 0.011(3) 0.016(3) 0.002(2) 0.004(2) 0.000(2) O3 0.010(2) 0.016(2) 0.022(3) -0.001(2) -0.002(2) -0.0006(17) O4 0.015(2) 0.012(2) 0.020(3) 0.003(2) 0.002(2) -0.0006(19) O5 0.021(3) 0.012(3) 0.013(3) 0.000(2) 0.001(2) 0.000(2)

\_geom\_special\_details

All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

loop

\_geom\_bond\_atom\_site\_label\_1 \_geom\_bond\_atom\_site\_label\_2 \_geom\_bond\_distance

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geom bond publ flag
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Er1 O3 2.398(5) 7 455 ?
Er1 O4 2.496(7).?
Er1 O1 2.5366(3) . ?
Er1 Si1 2.935(2) 6?
Er1 Si1 3.067(3).?
Er1 Si1 3.390(3) 1 556?
Er1 Na1 3.404(3) 6?
Si1 O2 1.639(7) . ?
Sil O5 1.639(7) 6 554?
Si1 O3 1.647(6) 4 655 ?
Si1 O4 1.673(6) . ?
Si1 Er1 2.935(2) 6 554 ?
Si1 Na1 3.064(4) 3 ?
Si1 Na2 3.090(2) . ?
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Sil Nal 3.188(5) 6?
Sil Nal 3.249(5) 6 554?
Si1 Er1 3.389(3) 1 554 ?
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'Smith, Mark D.' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Yeon, Jeongho' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 audit creation method SHELXL-2013 chemical name systematic ? ? chemical name common ? chemical melting point chemical formula moiety 'H Na5 O17 Si4 Tm4' chemical formula sum 'H Na5 O17 Si4 Tm4' chemical formula weight 1175.03 loop atom type symbol atom type description atom type scat dispersion real atom type scat dispersion imag atom type scat source O O 0.0106 0.0060 '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' Na Na 0.0362 0.0249 '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' Tm Tm -0.3139 5.2483 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4' space group crystal system tetragonal space group IT number 82 \_space\_group name H-M alt 'I -4' 'I -4' space group name Hall loop \_space\_group\_symop\_operation xyz 'x, y, z' '-x, -y, z'

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'v+1/2, -x+1/2, -z+1/2'				
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\_reflns\_special\_details

Reflections were merged by SHELXL according to the crystal class for the calculation of statistics and refinement.

\_reflns\_Friedel\_fraction is defined as the number of unique Friedel pairs measured divided by the number that would be possible theoretically, ignoring centric projections and systematic absences.

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\_computing\_molecular\_graphics various \_computing\_publication\_material various

\_refine\_special details Refined as a 2-component inversion twin. refine ls structure factor coef Fsqd \_refine\_ls\_matrix\_type full refine ls weighting scheme calc refine ls weighting details  $w=1/[\sqrt{c^2}(Fo^2)+(0.0330P)^2]$ where  $P = (Fo^2 + 2Fc^2)/3$ atom sites solution primary ? atom sites solution secondary ? atom sites solution hydrogens \_refine\_ls\_hydrogen\_treatment undef refine ls extinction method none refine ls extinction coef refine ls abs structure details Refined as an inversion twin. refine ls abs structure Flack 0.54(3) chemical absolute configuration? refine ls number reflns 1735 refine ls number parameters 72 refine ls number restraints 0 \_refine\_ls\_R\_factor\_all 0.0320 refine ls R factor gt 0.0291 refine ls wR factor ref 0.0603 refine ls wR factor gt 0.0593 \_refine\_ls\_goodness of fit ref 1.062 refine ls restrained S all 1.062 refine ls shift/su max 0.000 refine ls shift/su mean 0.000 loop\_ atom site label atom site type symbol \_atom\_site\_fract x atom site fract y atom site fract z atom site U iso or equiv

atom site adp type atom site occupancy atom site site symmetry order atom site calc flag atom site refinement flags posn atom site refinement flags adp atom site refinement flags occupancy atom site disorder assembly atom site disorder group Tm1 Tm 0.11651(2) 0.18374(3) 0.47653(6) 0.01052(8) Uani 1 1 d . . . . Si1 Si 0.24583(17) 0.10295(17) 0.0115(5) 0.0117(4) Uani 1 1 d . . . . . Na1 Na 0.1052(3) 0.4104(3) 0.0060(8) 0.0196(7) Uani 1 1 d . . . . . Na2 Na 0.0000 0.0000 0.060(2) 0.031(4) Uani 0.5 2 d S T P . . O1 O 0.0000 0.0000 0.5000 0.026(3) Uani 1 4 d S T P . . O2 O 0.2044(6) 0.0328(5) 0.2586(12) 0.0148(11) Uani 1 1 d . . . . O3 O 0.1091(4) 0.6131(4) 0.0116(13) 0.0149(11) Uani 1 1 d . . . . O4 O 0.2019(5) 0.2362(4) 0.0662(12) 0.0124(10) Uani 1 1 d . . . . O5 O 0.3131(6) 0.4512(5) 0.2633(11) 0.0140(11) Uani 1 1 d . . . .

## loop\_

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atom site aniso U 22
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Si1 0.0120(8) 0.0100(8) 0.0131(10) -0.0001(9) -0.0005(9) -0.0008(6)
Na1 0.0168(14) 0.0143(13) 0.028(2) 0.0024(16) 0.0025(15) 0.0016(10)
Na2 0.020(7) 0.028(7) 0.044(12) 0.000 0.000 -0.005(7)
O1 0.021(3) 0.021(3) 0.035(9) 0.000 0.000 0.000
O2\ 0.021(3)\ 0.012(3)\ 0.012(3)\ 0.003(2)\ 0.002(2)\ 0.000(2)
O3\ 0.011(2)\ 0.010(2)\ 0.023(3)\ -0.003(2)\ 0.001(2)\ -0.0006(16)
O4 0.010(2) 0.009(2) 0.017(3) 0.003(2) 0.001(2) 0.0006(16)
O5\ 0.022(3)\ 0.010(3)\ 0.011(3)\ 0.000(2)\ 0.002(2)\ 0.000(2)
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\_geom\_special\_details

All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

loop geom bond atom site label 1 \_geom\_bond atom site label 2 geom bond distance geom bond site symmetry 2 \_geom\_bond\_publ\_flag Tm1 O2 2.250(6) 4 556 ? Tm1 O5 2.303(6) 7\_455 ? Tm1 O2 2.331(6) . ? Tm1 O4 2.338(5) 6 ? Tm1 O5 2.339(6) 6 ? Tm1 O3 2.390(5) 7 455 ? Tm1 O4 2.497(6) . ? Tm1 O1 2.5112(3) . ? Tm1 Si1 2.933(2) 6? Tm1 Si1 3.063(3).? Tm1 Na2 3.366(9) . ? Tm1 Si1 3.379(3) 1 556? Sil O5 1.626(6) 6 554 ? Sil O2 1.631(7) . ? Si1 O3 1.633(6) 4 655 ? Si1 O4 1.644(6) . ? Sil Tml 2.933(2) 6 554? Si1 Na1 3.061(4) 3? Si1 Na2 3.084(2) . ? Si1 Na2 3.097(3) 3? Si1 Na1 3.176(5) 6? Sil Nal 3.227(5) 6 554? Si1 Tm1 3.379(3) 1 554? Na1 O4 2.320(6) . ? Na1 O3 2.336(6) . ? Na1 O3 2.485(6) 2 565 ? Na1 O3 2.614(8) 7 455 ? Na1 O2 2.651(7) 6 554 ? Na1 O3 2.803(8) 7 454 ? Na1 O5 2.809(7) . ? Na1 Si1 3.061(4) 4? Na1 Si1 3.176(5) 6 554 ? Na1 Na1 3.185(6) 2 565 ? Na1 Si1 3.227(5) 6? Na1 Tm1 3.390(3) 6 554 ? Na2 Na2 0.65(3) 3 ? Na2 O1 2.371(13) . ? Na2 O5 2.421(8) 7 455 ? Na2 O5 2.421(8) 8 545 ?

Na2 O2 2.614(9) . ? Na2 O2 2.615(9) 2 ? Na2 O5 2.744(10) 5 444 ? Na2 O5 2.744(10) 6 554 ? Na2 O2 2.942(10) 3 ? Na2 O2 2.942(10) 4 ? Na2 Si1 3.084(2) 2 ? O1 Na2 2.371(13) 3 556? O1 Tm1 2.5112(3) 2 ? O1 Tm1 2.5112(3) 4 556? O1 Tm1 2.5112(3) 3 556? O2 Tm1 2.250(6) 3 556 ? O2 Na1 2.651(7) 6 ? O2 Na2 2.942(10) 3 ? O3 Si1 1.633(6) 3 565 ? O3 Tm1 2.390(5) 8 ? O3 Na1 2.485(6) 2 565 ? O3 Na1 2.614(8) 8 ? O3 Na1 2.803(8) 8 554 ? O4 Tm1 2.338(5) 6 554 ? O5 Si1 1.626(6) 6 ? O5 Tm1 2.303(6) 8 ? O5 Tm1 2.339(6) 6 554 ? O5 Na2 2.421(8) 7 ? O5 Na2 2.744(10) 5 ?

loop

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Tm1 O2 Na2 100.9(3) 3 556 3 ? Tm1 O2 Na2 92.5(2) . 3 ? Na1 O2 Na2 171.6(3) 6 3 ? Si1 O3 Na1 173.7(4) 3 565 .? Si1 O3 Tm1 91.6(2) 3 565 8 ? Na1 O3 Tm1 92.06(19) . 8 ? Si1 O3 Na1 93.7(2) 3 565 2 565 ? Na1 O3 Na1 82.64(19) . 2 565 ? Tm1 O3 Na1 174.6(3) 8 2 565 ? Si1 O3 Na1 96.1(3) 3 565 8 ? Na1 O3 Na1 88.7(2) . 8 ? Tm1 O3 Na1 93.4(2) 8 8 ? Na1 O3 Na1 85.6(2) 2 565 8 ? Si1 O3 Na1 87.3(3) 3 565 8 554? Na1 O3 Na1 87.3(2) . 8 554 ? Tm1 O3 Na1 96.2(2) 8 8 554 ? Na1 O3 Na1 84.5(2) 2 565 8 554? Na1 O3 Na1 169.7(2) 8 8 554 ? Si1 O4 Na1 158.7(4) . . ? Si1 O4 Tm1 93.2(2) . 6 554 ? Na1 O4 Tm1 93.4(2) . 6 554 ? Si1 O4 Tm1 93.1(3) . . ? Na1 O4 Tm1 98.3(2) . . ? Tm1 O4 Tm1 129.2(3) 6 554.? Si1 O5 Tm1 141.9(4) 6 8 ? Si1 O5 Tm1 115.7(3) 6 6 554 ? Tm1 O5 Tm1 100.0(2) 8 6 554 ? Si1 O5 Na2 97.9(4) 6 7 ? Tm1 O5 Na2 90.8(3) 8 7 ? Tm1 O5 Na2 96.1(3) 6 554 7 ? Si1 O5 Na2 85.7(3) 6 5 ? Tm1 O5 Na2 98.4(2) 8 5 ? Tm1 O5 Na2 104.5(3) 6 554 5 ? Si1 O5 Na1 89.2(3) 6 . ? Tm1 O5 Na1 82.8(2) 8 . ? Tm1 O5 Na1 81.83(19) 6 554.? Na2 O5 Na1 172.8(4) 7 . ? Na2 O5 Na1 173.2(3) 5 . ?

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\_publ\_requested\_journal Inorg.Chem. publ contact author name 'Hans-Conrad zur Loye' \_publ\_contact\_author\_address ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 \_publ\_contact\_author\_email zurloye@mailbox.sc.edu publ contact author phone +1-803-777-6916 publ contact author fax +1-803-777-8508loop publ author name \_publ\_author\_address 'zur Loye, Hans-Conrad' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Latshaw, Allison M.' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Chance, W. Michael' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Morrison, Gregory' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Smith, Mark D.' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Yeon, Jeongho' ;University of South Carolina

Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 audit creation method SHELXL-2013 \_chemical\_name\_systematic ? ? chemical name common chemical melting point 9 chemical formula moiety 'H Na5 O17 Si4 Yb4' chemical formula sum 'H Na5 O17 Si4 Yb4' chemical formula weight 1192.48 loop atom type symbol atom type description atom type scat dispersion real atom type scat dispersion imag atom type scat source O O 0.0106 0.0060 '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' Na Na 0.0362 0.0249 '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' Yb Yb -0.3850 5.5486 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4' tetragonal space group crystal system

\_space\_group\_IT\_number 82 \_space\_group\_name\_H-M\_alt 'I -4' \_space\_group\_name\_Hall 'I -4'

\_shelx\_space\_group\_comment

;

The symmetry employed for this shelxl refinement is uniquely defined by the following loop, which should always be used as a source of symmetry information in preference to the above space-group names. They are only intended as comments.

;

loop\_ \_space\_group\_symop\_operation\_xyz 'x, y, z' '-x, -y, z' 'y, -x, -z'

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'-x+1/2, -y+1/2, z+1/2'
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Friedel pairs measured divided by the number that would be possible theoretically, ignoring centric projections and systematic absences.

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computing publication material various refine special details Refined as a 2-component inversion twin. \_refine\_ls\_structure\_factor coef Fsqd refine ls matrix type full refine ls weighting scheme calc refine ls weighting details  $w=1/[\sqrt{0.0448P}^{-1.002P}]$ where  $P = (Fo^{2^{+2}Fc^{2^{-2}}})/3$ \_atom\_sites\_solution primary ? atom sites solution secondary ? atom sites solution hydrogens refine ls hydrogen treatment undef refine ls extinction method none refine ls extinction coef refine ls abs structure details Refined as an inversion twin. refine ls abs structure Flack 0.53(3) chemical absolute configuration? refine ls number reflns 1750 refine ls number parameters 69 refine ls number restraints 0 refine ls R factor all 0.0327 refine ls R factor gt 0.0309 refine ls wR factor ref 0.0710 refine ls wR factor gt 0.0703 refine ls goodness of fit ref 1.061 \_refine\_ls\_restrained S all 1.061 refine ls shift/su max 0.000 \_refine\_ls\_shift/su mean 0.000 loop atom site label atom site type symbol atom site fract x atom site fract y atom site fract z

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loop\_

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\_geom\_special\_details

All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

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Yb1 O1 Yb1 174.124(16) . 2 ?
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Yb1 O1 Na2 92.938(8) 3 556 3 ?
Yb1 O1 Na2 92.938(8) 4 556 3 ?
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Na2 O1 Na2 180.0 1 556 3 ? Si1 O2 Yb1 158.1(4) . 3 556? Si1 O2 Yb1 99.3(3) . . ? Yb1 O2 Yb1 102.4(3) 3 556 . ? Si1 O2 Na2 89.7(4) . . ? Yb1 O2 Na2 94.5(3) 3 556 . ? Yb1 O2 Na2 86.8(3) . . ? Si1 O2 Na1 91.8(3) . 6? Yb1 O2 Na1 87.1(3) 3 556 6 ? Yb1 O2 Na1 84.9(2) . 6? Na2 O2 Na1 171.7(3) . 6? Si1 O2 Na2 80.8(4) . 3 ? Yb1 O2 Na2 101.1(3) 3 556 3 ? Yb1 O2 Na2 92.6(3) . 3 ? Na1 O2 Na2 171.8(3) 6 3 ? Sil O3 Nal 174.1(4) 3 565.? Si1 O3 Yb1 91.3(3) 3 565 8 ? Na1 O3 Yb1 92.6(2) . 8 ? Si1 O3 Na1 93.6(3) 3 565 2 565 ? Na1 O3 Na1 82.5(2) . 2 565 ? Yb1 O3 Na1 175.0(3) 8 2 565 ? Si1 O3 Na1 95.5(3) 3 565 8? Na1 O3 Na1 88.6(3) . 8 ? Yb1 O3 Na1 93.1(2) 8 8 ? Na1 O3 Na1 85.5(3) 2 565 8 ? Si1 O3 Na1 87.9(3) 3 565 8 554 ? Na1 O3 Na1 87.3(3) . 8 554 ? Yb1 O3 Na1 96.6(2) 8 8 554 ? Na1 O3 Na1 84.5(2) 2 565 8 554 ? Na1 O3 Na1 169.6(3) 8 8 554 ? Si1 O4 Na1 158.8(4) . . ? Sil O4 Yb1 92.8(3) . 6 554 ? Na1 O4 Yb1 93.5(2) . 6 554 ? Sil O4 Yb1 93.0(3) . . ? Na1 O4 Yb1 98.8(3) . . ? Yb1 O4 Yb1 128.7(3) 6 554 . ? Si1 O5 Yb1 141.0(4) 6 8 ? Si1 O5 Yb1 115.8(3) 6 6 554 ? Yb1 O5 Yb1 100.8(2) 8 6 554 ? Si1 O5 Na2 96.5(4) 6 7 ? Yb1 O5 Na2 91.5(3) 8 7 ? Yb1 O5 Na2 97.3(3) 6 554 7 ? Si1 O5 Na2 86.3(4) 6 5 ? Yb1 O5 Na2 97.7(3) 8 5 ? Yb1 O5 Na2 104.3(3) 6 554 5 ? Sil O5 Nal 89.3(3) 6 . ?

Yb1 O5 Na1 82.7(2) 8 . ? Yb1 O5 Na1 81.8(2) 6 554 . ? Na2 O5 Na1 173.8(4) 7 . ? Na2 O5 Na1 173.6(4) 5 . ? refine diff density max 3.098 refine diff density min -1.765 refine diff density rms 0.380 #===END data Na5Y4(OH)[SiO4]4 publ requested journal Inorg.Chem. \_publ\_contact\_author name 'Hans-Conrad zur Loye' \_publ\_contact author address ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 \_publ\_contact\_author\_email zurloye@mailbox.sc.edu \_publ\_contact\_author phone +1-803-777-6916 \_publ\_contact\_author fax +1-803-777-8508 loop \_publ\_author name \_\_\_\_\_publ\_author\_address 'zur Loye, Hans-Conrad' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Latshaw, Allison M.' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Chance, W. Michael' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Morrison, Gregory' ;University of South Carolina

Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Smith, Mark D.' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 'Yeon, Jeongho' ;University of South Carolina Department of Chemistry and Biochemistry 631 Sumter Street Columbia, SC 29208 audit creation method SHELXL-2013 chemical name systematic ? ? chemical name common ? chemical melting point \_chemical\_formula\_moiety ? chemical formula sum 'H Na5 O17 Si4 Y4' chemical formula weight 855.96 loop atom type symbol atom type description atom type scat dispersion real atom type scat dispersion imag atom type scat source O O 0.0106 0.0060 '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' Na Na 0.0362 0.0249 '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' Y Y -2.7962 3.5667 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4' \_space\_group\_crystal\_system tetragonal space group IT number 82 \_space\_group\_name H-M alt 'I -4' space group name Hall 'I -4'

\_shelx\_space\_group\_comment

, The symmetry employed for this shelxl refinement is uniquely defined by the following loop, which should always be used as a source of symmetry information in preference to the above space-group names. They are only intended as comments.

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loop\_

space group symop operation xyz 'x, y, z' '-x, -y, z' 'y, -x, -z' '-y, x, -z' 'x+1/2, y+1/2, z+1/2' '-x+1/2, -y+1/2, z+1/2' y+1/2, -x+1/2, -z+1/2''-y+1/2, x+1/2, -z+1/2' cell length a 11.6025(4) \_cell\_length b 11.6025(4) cell length c 5.4264(5) \_cell\_angle\_alpha 90 cell angle beta 90 cell angle gamma 90 \_cell\_volume 730.49(8) cell formula units Z 2 cell measurement temperature 296(2)cell measurement reflns used 681 cell measurement theta min 2.482 cell measurement theta max 26.484 exptl crystal description 'square plate' exptl crystal colour colorless exptl crystal density meas ? \_exptl\_crystal density method ? \_exptl\_crystal density diffrn 3.891 exptl crystal F 000 808 \_exptl\_transmission\_factor min ? exptl transmission factor max ? exptl crystal size max 0.050 \_exptl\_crystal size mid 0.040

\_exptl\_crystal\_size\_min 0.020 \_exptl\_absorpt\_coefficient\_mu 16.310 \_shelx\_estimated\_absorpt\_T\_min 0.496 \_shelx\_estimated\_absorpt\_T\_max 0.736 \_exptl\_absorpt\_correction\_type\_multi-scan

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exptl absorpt correction T max 1.0000
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exptl special details
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diffrn radiation wavelength
                              0.71073
diffrn radiation type
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diffrn source
diffrn measurement device type 'Bruker SMART APEX CCD diffractometer'
diffrn measurement method
                                'omega and phi scans'
diffrn detector area resol mean?
diffrn reflns number
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diffrn reflns av unetI/netI
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diffrn reflns av R equivalents 0.0921
diffrn reflns limit h min
                             -17
diffrn reflns limit h max
                              17
diffrn reflns limit k min
                             -17
diffrn reflns limit k max
                             17
 diffrn reflns limit 1 min
                             -8
diffrn reflns limit 1 max
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diffrn reflns theta min
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diffrn reflns theta max
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diffrn reflns theta full
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diffrn measured fraction theta full 0.997
diffrn reflns Laue measured fraction max 1.000
diffrn reflns Laue measured fraction full 0.997
diffrn reflns point group measured fraction max 1.000
diffrn reflns point group measured fraction full 0.998
reflns number total
                           1365
reflns number gt
                           1207
reflns threshold expression
                              'I > 2 (I)'
reflns Friedel coverage
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reflns Friedel fraction max
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reflns Friedel fraction full
                             1.000
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\_reflns\_special\_details

Reflections were merged by SHELXL according to the crystal class for the calculation of statistics and refinement.
\_reflns\_Friedel\_fraction is defined as the number of unique Friedel pairs measured divided by the number that would be possible theoretically, ignoring centric projections and systematic absences.

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\_computing\_data\_collection'SMART Version 5.625 (Bruker, 2001)'\_computing\_cell\_refinement'SAINT-Plus Version 6.45 (Bruker, 2001)'\_computing\_data\_reduction'SAINT-Plus Version 6.45 (Bruker, 2001)'\_computing\_structure\_solution'SHELXS97 (Sheldrick, 1997)'\_computing\_molecular\_graphicsvariouscomputing\_publication'SHELXTL Version 6.14 (Bruker, 2000)'

\_refine\_special\_details

Refined as a 2-component inversion twin. refine ls structure factor coef Fsqd refine ls matrix type full refine ls weighting scheme calc refine ls weighting details  $w=1/[\sqrt{c^2(Fo^2)}+(0.0549P)^2]$ where  $P = (Fo^2 + 2Fc^2)/3$ atom sites solution primary atom sites solution secondary ? atom sites solution hydrogens refine ls hydrogen treatment undef refine ls extinction method none refine ls extinction coef \_refine\_ls\_abs\_structure details Refined as an inversion twin. refine ls abs structure Flack 0.51(2)chemical absolute configuration? refine ls number reflns 1365 refine ls number parameters 72 refine\_ls\_number\_restraints 0 refine ls R factor all 0.0598 refine ls R factor gt 0.0476 refine ls wR factor ref 0.1049 refine ls wR factor gt 0.0998 refine ls goodness of fit ref 1.024

\_refine\_ls\_restrained\_S\_all 1.024 \_refine\_ls\_shift/su\_max 0.000 \_refine\_ls\_shift/su\_mean 0.000

loop\_

atom site label atom site type symbol atom site fract x atom site fract y atom site fract z atom site U iso or equiv atom site adp type atom site occupancy atom site site symmetry order atom site calc flag atom site refinement flags posn atom site refinement flags adp atom site refinement flags occupancy \_atom\_site\_disorder assembly atom site disorder group Y1 Y 0.11718(6) 0.18419(6) 0.47609(15) 0.0106(2) Uani 1 1 d . . . . Si1 Si 0.24579(19) 0.10245(19) 0.0109(6) 0.0148(5) Uani 1 1 d . . . . Na1 Na 0.1044(3) 0.4111(3) 0.0051(10) 0.0219(8) Uani 1 1 d . . . . . Na2 Na 0.0000 0.0000 0.062(2) 0.034(5) Uani 0.5 2 d S T P . . O1 O 0.0000 0.0000 0.5000 0.026(3) Uani 1 4 d S T P . . O2 O 0.2039(7) 0.0321(6) 0.2589(13) 0.0171(15) Uani 1 1 d . . . . O3 O 0.1088(5) 0.6128(5) 0.0118(15) 0.0155(12) Uani 1 1 d . . . . O4 O 0.2017(5) 0.2365(5) 0.0651(13) 0.0150(13) Uani 1 1 d . . . . O5 O 0.3117(7) 0.4515(6) 0.2644(12) 0.0140(13) Uani 1 1 d . . . . loop\_ atom site aniso label atom site aniso U 11 atom site aniso U 22 atom site aniso U 33 atom site aniso U 23 atom site aniso U 13 atom site aniso U 12 Y1 0.0077(3) 0.0094(3) 0.0146(3) 0.0006(3) 0.0005(3) 0.0017(3) Si1 0.0126(10) 0.0124(10) 0.0193(13) 0.0007(12) -0.0002(12) -0.0013(8) Na1 0.0163(16) 0.0137(15) 0.036(2) 0.001(2) -0.001(2) 0.0000(12) Na2 0.035(10) 0.020(9) 0.046(14) 0.000 0.000 -0.003(9) O1 0.021(4) 0.021(4) 0.035(9) 0.000 0.000 0.000  $O2\ 0.025(4)\ 0.010(3)\ 0.016(3)\ 0.002(3)\ 0.004(3)\ 0.001(3)$  $O3\ 0.009(2)\ 0.012(3)\ 0.026(4)\ 0.002(3)\ -0.004(3)\ 0.0002(19)$  $O4\ 0.013(3)\ 0.011(3)\ 0.021(4)\ 0.001(3)\ 0.001(3)\ -0.002(2)$ 

## O5 0.021(4) 0.007(3) 0.014(3) -0.001(2) 0.002(3) -0.002(3)

\_geom\_special\_details

All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

loop

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geom bond atom site label 1 geom bond atom site label 2 geom bond distance geom bond site symmetry 2 geom bond publ flag Y1 O2 2.262(7) 4 556 ? Y1 O5 2.324(7) 7 455 ? Y1 O4 2.344(6) 6 ? Y1 O2 2.349(7).? Y1 O5 2.367(7) 6 ? Y1 O3 2.403(5) 7 455 ? Y1 O4 2.511(7) . ? Y1 O1 2.5363(8) . ? Y1 Si1 2.948(2) 6? Y1 Si1 3.082(3).? Y1 Na2 3.387(9).? Y1 Si1 3.398(3) 1 556? Sil O5 1.620(7) 6 554? Si1 O3 1.646(6) 4 655 ? Si1 O2 1.648(8) . ? Si1 O4 1.664(7).? Si1 Y1 2.948(2) 6 554 ? Si1 Na1 3.073(4) 3? Si1 Na2 3.102(3) . ? Si1 Na2 3.115(3) 3? Si1 Na1 3.200(6) 6? Sil Nal 3.252(6) 6 554? Si1 Y1 3.398(3) 1 554 ? Na1 O3 2.342(6) . ? Na1 O4 2.342(7) . ? Na1 O3 2.490(6) 2 565 ? Na1 O3 2.633(9) 7 455 ? Na1 O2 2.677(9) 6 554 ?

Na1 O3 2.816(9) 7 454 ? Nal O5 2.826(8).? Na1 Si1 3.073(4) 4 ? Na1 Na1 3.182(7) 2 565 ? Na1 Si1 3.200(6) 6 554? Na1 Si1 3.252(6) 6? Na1 Y1 3.418(3) 6 554 ? Na2 Na2 0.67(3) 3 ? Na2 O1 2.378(13) . ? Na2 O5 2.445(9) 7 455 ? Na2 O5 2.445(9) 8 545 ? Na2 O2 2.623(10) . ? Na2 O2 2.623(10) 2 ? Na2 O5 2.774(11) 5 444 ? Na2 O5 2.774(11) 6 554 ? Na2 O2 2.960(11) 3 ? Na2 O2 2.960(11) 4 ? Na2 Si1 3.102(3) 2 ? O1 Na2 2.378(13) 3 556? O1 Y1 2.5363(8) 2 ? O1 Y1 2.5363(8) 3 556 ? O1 Y1 2.5363(8) 4 556? O2 Y1 2.262(7) 3 556 ? O2 Na1 2.677(9) 6 ? O2 Na2 2.960(11) 3 ? O3 Si1 1.647(6) 3 565 ? O3 Y1 2.403(5) 8 ? O3 Na1 2.490(6) 2 565 ? O3 Na1 2.633(9) 8 ? O3 Na1 2.816(9) 8 554 ? O4 Y1 2.344(6) 6 554 ? O5 Si1 1.620(7) 6 ? O5 Y1 2.324(7) 8 ? O5 Y1 2.367(7) 6 554 ? O5 Na2 2.445(9) 7 ? O5 Na2 2.774(11) 5? loop \_geom\_angle atom site label 1

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O2 Si1 Y1 128.9(3) . 6 554 ?
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Na1 Si1 Y1 124.44(14) 3 . ?
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Y1 Si1 Na2 66.4(2) . . ? O5 Si1 Na2 51.0(4) 6 554 3? O3 Si1 Na2 156.9(3) 4 655 3 ? O2 Si1 Na2 69.2(4) . 3 ? O4 Si1 Na2 95.6(2) . 3 ? Y1 Si1 Na2 143.72(11) 6 554 3? Na1 Si1 Na2 105.65(9) 3 3 ? Y1 Si1 Na2 77.2(2) . 3 ? Na2 Si1 Na2 12.4(5) . 3 ? O5 Si1 Na1 153.7(3) 6 554 6? O3 Si1 Na1 61.6(3) 4 655 6? O2 Si1 Na1 56.7(3).6? O4 Si1 Na1 93.7(3) . 6? Y1 Si1 Na1 78.57(9) 6 554 6? Na1 Si1 Na1 69.29(16) 3 6 ? Y1 Si1 Na1 65.89(10).6? Na2 Si1 Na1 113.9(3).6? Na2 Si1 Na1 125.8(3) 3 6 ? O5 Si1 Na1 60.3(3) 6 554 6 554 ? O3 Si1 Na1 53.6(3) 4 655 6 554 ? O2 Si1 Na1 145.4(3) . 6 554 ? O4 Si1 Na1 111.0(3) . 6 554 ? Y1 Si1 Na1 72.44(9) 6 554 6 554 ? Na1 Si1 Na1 66.73(15) 3 6 554? Y1 Si1 Na1 164.75(10) . 6 554 ? Na2 Si1 Na1 123.2(3) . 6 554 ? Na2 Si1 Na1 111.4(3) 3 6 554 ? Na1 Si1 Na1 114.49(12) 6 6 554 ? O5 Si1 Y1 38.9(2) 6 554 1 554 ? O3 Si1 Y1 111.2(3) 4 655 1 554 ? O2 Si1 Y1 134.9(3) . 1 554 ? O4 Si1 Y1 75.8(3) . 1 554 ? Y1 Si1 Y1 87.01(7) 6 554 1 554 ? Na1 Si1 Y1 117.90(14) 3 1 554 ? Y1 Si1 Y1 113.64(7) . 1 554 ? Na2 Si1 Y1 77.3(2) . 1 554 ? Na2 Si1 Y1 66.2(2) 3 1 554 ? Na1 Si1 Y1 165.47(10) 6 1 554 ? Na1 Si1 Y1 61.81(10) 6 554 1 554 ? O3 Na1 O4 148.4(3) . . ? O3 Na1 O3 97.6(2) . 2 565 ? O4 Na1 O3 112.4(2) . 2 565 ? O3 Na1 O3 94.1(3) . 7 455 ? O4 Na1 O3 76.6(3) . 7 455 ? O3 Na1 O3 90.7(3) 2 565 7 455 ? O3 Na1 O2 75.1(2) . 6 554 ?

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O3 Na1 O5 78.9(2) . . ?
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