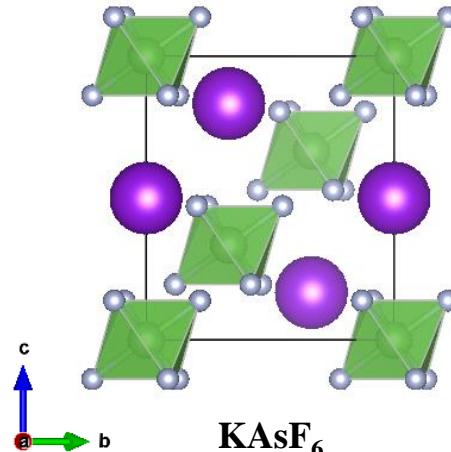


# Mathematics and Computer Science for Materials Innovation: Crystal Lattice Classifications

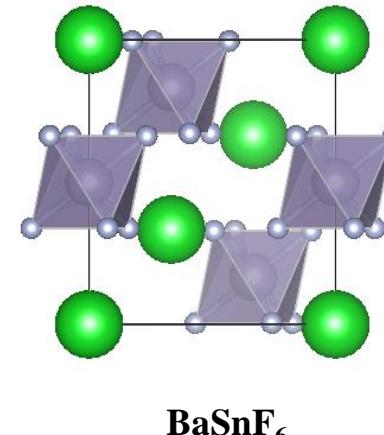
05.09 – 09.09 2022 Liverpool

gemma.delaflor@kit.edu

Comparison of structures by the Bilbao Crystallographic Server



?  
≈



# Content

## Crystal-structure descriptions

- Transformations between different crystal-structure descriptions
- Transformations to a lower symmetry space group
- Symmetrically equivalent descriptions

## Comparison between different structure descriptions

- Comparison of structures with the same/different composition
- Comparison of chiral structures

# Crystal structure description

- What type of information is necessary to describe a crystal structure?
  - Space Group
  - Lattice parameters
  - The number of independent atoms in the asymmetric unit
  - The atom type and the coordinates

141
6.6164 6.6164 6.0150 90 90 90
3
Zr 1 4a 0.000 0.750 0.125
Si 1 4b 0.000 0.750 0.625
O 1 16h 0.000 0.067 0.198

**BCS format**

# Standard setting

The settings of space groups hat coincide with the conventional space-group descriptions found in *Volume A of International Tables for Crystallography*. For space groups with more than one description, the following settings are chosen as standard:

- *Unique axis b* and *cell choice 1* for monoclinic space groups
- *Hexagonal axes* for rhombohedral space groups
- *Origin choice 2* (origin at  $\bar{1}$ ) for centrosymmetric space groups listed with two origin choices

**What can I do if my structures are described in a non-standard setting?**

# Example – Structure transformation

Transform the crystal structure of manganese trifluoride  $\text{MnF}_3$ , described in the space group  $I\bar{1}2/a1$  (No. 15), to its standard setting  $C\bar{1}2/c1$  taking into account that the transformation matrix  $(P, p) = -\mathbf{a} - \mathbf{c}, \mathbf{b}, \mathbf{a}; 0,0,0$

**Initial Setting:**  $I\bar{1}2/a$  (No. 15)

```

15
5.5017 5.0270 7.2619 90 92.814 90
3
Mn 1 4d 0.250000 0.250000 0.250000
F 1 8f 0.072200 -0.038000 0.305700
F 2 4e 0.250000 0.136000 0.000000
  
```

$$\xrightarrow{(P, p)} \quad -\mathbf{a} - \mathbf{c}, \mathbf{b}, \mathbf{a}$$

**Final Setting:**  $C\bar{1}2/c1$  (No. 15)

- 1) Transform the unit cell parameters
- 2) Transform of the atomic coordinates

# Example – Structure transformation

- 1) Transform the unit cell parameters

Calculate the metric tensor  $G$

**Initial Setting:** I12/a (No. 15)

$a$	$b$	$c$	$\alpha$	$\beta$	$\gamma$
5.5017	5.0270	7.2619	90	92.814	90

$$G = \begin{pmatrix} a^2 & ab \cos\gamma & ac \cos\beta \\ ab \cos\gamma & b^2 & bc \cos\alpha \\ ac \cos\beta & bc \cos\alpha & c^2 \end{pmatrix} \Rightarrow G = \begin{pmatrix} 30.268703 & 0 & -1.961435 \\ 0 & 25.270726 & 0 \\ -1.961435 & 0 & 52.735192 \end{pmatrix}$$

# Example – Structure transformation

- 1) Transform the unit cell parameters

Transform the unit cell parameters to the standard setting

**Initial Setting:**  $I\bar{1}2/a$  (No. 15)

$a$	$b$	$c$	$\alpha$	$\beta$	$\gamma$
5.5017	5.0270	7.2619	90	92.814	90

$$\mathbf{P} = \begin{pmatrix} \bar{1} & 0 & 1 \\ 0 & 1 & 0 \\ \bar{1} & 0 & 0 \end{pmatrix}$$

$$\mathbf{G}' = \mathbf{P}^T \cdot \mathbf{G} \cdot \mathbf{P} = \begin{pmatrix} \bar{1} & 0 & \bar{1} \\ 0 & 1 & 0 \\ 1 & 0 & 0 \end{pmatrix} \begin{pmatrix} 30.268703 & 0 & -1.961435 \\ 0 & 25.270726 & 0 \\ -1.961435 & 0 & 52.735192 \end{pmatrix} \begin{pmatrix} \bar{1} & 0 & 1 \\ 0 & 1 & 0 \\ \bar{1} & 0 & 0 \end{pmatrix}$$

$$\mathbf{G}' = \begin{pmatrix} 79.081024 & 0 & -28.307267 \\ 0 & 25.270726 & 0 \\ -28.307267 & 0 & 30.268703 \end{pmatrix}$$

**Final Setting:**  $C\bar{1}2/c1$  (No. 15)

$a$	$b$	$c$	$\alpha$	$\beta$	$\gamma$
8.8928	5.0270	5.5017	90	125.35	90

# Example – Structure transformation

2) Transform of the atomic coordinates  $\mathbf{x}' = \mathbf{P}^{-1}\mathbf{x}$

Mn	1	4d	0.250000	0.250000	0.250000
F	1	8f	0.072200	-0.038000	0.305700
F	2	4e	0.250000	0.136000	0.000000

$$\mathbf{P} = \begin{pmatrix} \bar{1} & 0 & 1 \\ 0 & 1 & 0 \\ \bar{1} & 0 & 0 \end{pmatrix} \Rightarrow \mathbf{P}^{-1} = \begin{pmatrix} 0 & 0 & \bar{1} \\ 0 & 1 & 0 \\ 1 & 0 & \bar{1} \end{pmatrix}$$

Mn       $x' = \begin{pmatrix} 0 & 0 & \bar{1} \\ 0 & 1 & 0 \\ 1 & 0 & \bar{1} \end{pmatrix} \begin{pmatrix} 0.25 \\ 0.25 \\ 0.25 \end{pmatrix} = \begin{pmatrix} -0.25 \\ 0.25 \\ 0 \end{pmatrix}$

F1       $x' = \begin{pmatrix} 0 & 0 & \bar{1} \\ 0 & 1 & 0 \\ 1 & 0 & \bar{1} \end{pmatrix} \begin{pmatrix} 0.0722 \\ -0.038 \\ 0.3057 \end{pmatrix} = \begin{pmatrix} -0.3057 \\ -0.038 \\ -0.2335 \end{pmatrix}$

F2       $x' = \begin{pmatrix} 0 & 0 & \bar{1} \\ 0 & 1 & 0 \\ 1 & 0 & \bar{1} \end{pmatrix} \begin{pmatrix} 0.25 \\ 0.136 \\ 0 \end{pmatrix} = \begin{pmatrix} 0 \\ 0.136 \\ 0.25 \end{pmatrix}$

**Final Setting:** C12/c1 (No. 15)

Mn	1	<span style="border: 1px solid blue; padding: 2px;"> </span>	-0.250000	0.250000	0.000000
F	1		-0.305700	-0.038000	-0.233500
F	2		0.000000	0.136000	0.250000

WP?

# Example – Structure transformation

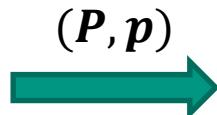
## Wyckoff Positions of Group C2/c (No. 15) [unique axis b]

Multiplicity	Wyckoff letter	Site symmetry	Coordinates	
			(0,0,0)	(1/2,1/2,0)
8	f	1	(x,y,z) (-x,y,-z+1/2) (-x,-y,-z) (x,-y,z+1/2)	
4	e	2	(0,y,1/4) (0,-y,3/4)	
4	d	-1	(1/4,1/4,1/2) (3/4,1/4,0)	
4	c	-1	(1/4,1/4,0) (3/4,1/4,1/2)	
4	b	-1	(0,1/2,0) (0,1/2,1/2)	
4	a	-1	(0,0,0) (0,0,1/2)	

Mn	1	4d	-0.250000	0.250000	0.000000
F	1	8f	-0.305700	0.038000	-0.233500
F	2	4e	0.000000	0.136000	0.250000

**Initial Setting:** I12/a (No. 15)

15  
 5.5017 5.0270 7.2619 90 92.814 90  
 3  
 Mn 1 4d 0.250000 0.250000 0.250000  
 F 1 8f 0.072200 -0.038000 0.305700  
 F 2 4e 0.250000 0.136000 0.000000



**Final Setting:** C12/c1 (No. 15)

15  
 8.8928 5.0270 5.5017 90 125.35 90  
 3  
 Mn 1 4d -0.250000 0.250000 0.000000  
 F 1 8f -0.305700 0.038000 -0.233500  
 F 2 4e 0.000000 0.136000 0.250000

# Crystal-structure descriptions

## Transformation between different structure descriptions

Structure Utilities	
CELLTRAN	Transform Unit Cells
STRAIN	Strain Tensor Calculation
WPASSIGN	Assignment of Wyckoff Positions
TRANSTRU	Transform structures.
SETSTRU	Alternative Settings for a given Crystal Structure
EQUIVSTRU	Equivalent Descriptions for a given Crystal Structure
CIF2STANDARD	Transforms a given CIF (in any setting) to that of standard setting
STRCONVERT	Convert & Edit Structure Data (supports the CIF, mCIF, VESTA, VASP formats -- with magnetic information where available)
VISUALIZE	Visualize structures using Jmol
COMPSTRU	Comparison of Crystal Structures with the same Symmetry
STRUCTURE RELATIONS	Evaluation of structure relationships [transformation matrix] between group-subgroup related phases
PSEUDOLATTICE	Pseudosymmetry of a lattice and compatible supergroups

# IT A setting structure description

**SETSTRU** <https://www.cryst.ehu.es/cryst/setSTRU.html>

Transform a structure to an alternative setting

**Transform to an alternative setting**

The program SETSTRU performs the transformations between crystal-structure descriptions referred to the so-called **IT A setting** of space groups.

The first step consists in the input of the structure data. The data can be given using the form or it can be loaded from a CIF file. The necessary data for the structure consists in the number of its space group in the *International Tables for Crystallography*, Vol A, the lattice parameters (in Å and degrees), the number of the atoms in the asymmetric unit and the corresponding atomic positions.

Next, it is necessary to specify the initial and final settings of the structure descriptions among the listed ITA-settings of the structure's space group (e.g. to convert from **rhombohedral** to the **standard hexagonal** settings).

A detailed description of the structure with respect to the final setting of the space group is shown in the output.

Structure Data  
[In CIF format]

No file selected.  
HINT: [ The option for a given filename is preferential ]

15	5.5017	5.0270	7.2619	90	92.814	90
3	Mn	1	4d	0.250000	0.250000	0.250000
	F	1	8f	0.072200	-0.038000	0.305700
	F	2	4e	0.250000	0.136000	0.000000

Structure

**CIF file**

**BCS format**

To transform a structure described in a *IT A*-setting into another *IT A*-setting

# ITA setting structure description

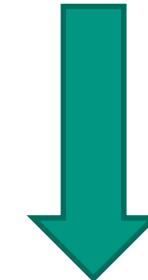
## Choose the initial and final space groups symbols

The standard setting (default) of the space group 15 is  $C12/c1$  [cell choice 1]

Initial	Final	Setting	P	$P^{-1}$
<input type="radio"/>	<input checked="" type="radio"/>	$C12/c1$ [cell choice 1]	<b>a,b,c</b>	<b>a,b,c</b>
<input type="radio"/>	<input type="radio"/>	$A12/a1$ [cell choice 1]	<b>c,-b,a</b>	<b>c,-b,a</b>
<input type="radio"/>	<input type="radio"/>	$A12/n1$ [cell choice 2]	<b>-a-c,b,a</b>	<b>c,b,-a-c</b>
<input type="radio"/>	<input type="radio"/>	$C12/n1$ [cell choice 2]	<b>a,-b,-a-c</b>	<b>a,-b,-a-c</b>
<input checked="" type="radio"/>	<input type="radio"/>	$I12/a1$ [cell choice 3]	<b>c,b,-a-c</b>	<b>-a-c,b,a</b>
<input type="radio"/>	<input type="radio"/>	$I12/c1$ [cell choice 3]	<b>-a-c,-b,c</b>	<b>-a-c,-b,c</b>
<input type="radio"/>	<input type="radio"/>	$A112/a$ [cell choice 1]	<b>c,a,b</b>	<b>b,c,a</b>
<input type="radio"/>	<input type="radio"/>	$B112/b$ [cell choice 1]	<b>a,c,-b</b>	<b>a,-c,b</b>
<input type="radio"/>	<input type="radio"/>	$B112/n$ [cell choice 2]	<b>a,-a-c,b</b>	<b>a,c,-a-b</b>
<input type="radio"/>	<input type="radio"/>	$A112/n$ [cell choice 2]	<b>-a-c,a,-b</b>	<b>b,-c,-a-b</b>
<input type="radio"/>	<input type="radio"/>	$I112/b$ [cell choice 3]	<b>-a-c,c,b</b>	<b>-a-b,c,b</b>
<input type="radio"/>	<input type="radio"/>	$I112/a$ [cell choice 3]	<b>c,-a-c,-b</b>	<b>-a-b,-c,a</b>
<input type="radio"/>	<input type="radio"/>	$B2/b11$ [cell choice 1]	<b>b,c,a</b>	<b>c,a,b</b>
<input type="radio"/>	<input type="radio"/>	$C2/c11$ [cell choice 1]	<b>-b,a,c</b>	<b>b,-a,c</b>
<input type="radio"/>	<input type="radio"/>	$C2/n11$ [cell choice 2]	<b>b,a,-a-c</b>	<b>b,a,-b-c</b>
<input type="radio"/>	<input type="radio"/>	$B2/n11$ [cell choice 2]	<b>-b,-a-c,a</b>	<b>c,-a,-b-c</b>
<input type="radio"/>	<input type="radio"/>	$I2/c11$ [cell choice 3]	<b>b,-a-c,c</b>	<b>-b-c,a,c</b>
<input type="radio"/>	<input type="radio"/>	$I2/b11$ [cell choice 3]	<b>-b,c,-a-c</b>	<b>-b-c,-a,b</b>

## Initial Setting: $I12/a$ (No. 15)

15  
 5.5017 5.0270 7.2619 90 92.814 90  
 3  
 Mn 1 4d 0.250000 0.250000 0.250000  
 F 1 8f 0.072200 -0.038000 0.305700  
 F 2 4e 0.250000 0.136000 0.000000



## Final Setting: $C12/c1$ (No. 15)

# ITA setting structure description

## Transformation to standard setting of space group 15

### Initial structure

**Initial Setting:** I12/a1 [ cell choice 3 ] (No. 15)

```

15
5.5017 5.0270 7.2619 90 92.814 90
3
Mn    1      -    0.250000    0.250000    0.250000
F     1      -    0.072200   -0.038000    0.305700
F     2      -    0.250000    0.136000    0.000000
  
```

### Final structure

**Final Setting:** C12/c1 [ cell choice 1 ] (No. 15)

```

15
8.8928 5.0270 5.5017 90.00 125.35 90.00
3
Mn    1      4d    -0.250000    0.250000    0.000000
F     1      8f    -0.305700   -0.038000   -0.233500
F     2      4e    0.000000    0.136000    0.250000
  
```

CIF File

Transformation matrix ( $P, p$ ): -a-c,b,a; 0,0,0

Matrix form:

$$(P, p) = \begin{bmatrix} 1 & -1 & 0 & 1 \\ 0 & 1 & 0 & 0 \\ -1 & 0 & 0 & 0 \end{bmatrix} \begin{bmatrix} 0 \\ 0 \\ 0 \end{bmatrix}$$

### Atoms Data:

The data shown in this table corresponds to the final setting which corresponds to the standard setting

AT.	WP	SS	Representative	Atomic orbit
Mn1	4d (1/4,1/4,1/2)	-1	(0.750000, 0.250000, 0.000000)	(0.750000, 0.250000, 0.500000) (0.250000, 0.250000, 0.750000) (0.250000, 0.750000, 0.000000) (0.750000, 0.750000, 0.500000)
F1	8f (x,y,z)	1	(0.694300, 0.962000, 0.766500)	(0.694300, 0.962000, 0.766500) (0.305700, 0.962000, 0.733500) (0.305700, 0.038000, 0.233500) (0.694300, 0.038000, 0.266500) (0.194300, 0.462000, 0.766500) (0.805700, 0.462000, 0.733500) (0.805700, 0.538000, 0.233500) (0.194300, 0.538000, 0.266500)
F2	4e (0,y,1/4)	2	(0.000000, 0.136000, 0.250000)	(0.000000, 0.136000, 0.250000) (0.000000, 0.864000, 0.750000) (0.500000, 0.636000, 0.250000) (0.500000, 0.364000, 0.750000)

This data is only calculated by the program if the final setting corresponds to the standard

# Example – Scheelite

- Scheelite ( $\text{CaWO}_4$ ) is a mineral that crystallizes in the space group  $I4_1/a$  (No. 88). In the *Inorganic Crystal Structure Database* the following two descriptions of  $\text{CaWO}_4$  can be found:

# (a) Origin choice 1

# ICSD: 15869

88

5.243 5.243 11.376 90 90 90

3

Ca 1 4b 0.0000 0.0000 0.5000

W 1 4a 0.0000 0.0000 0.0000

O 1 16f 0.2413 0.1511 0.0861

# (b) Origin choice 2

# ICSD: 15586

88

5.243 5.243 11.376 90 90 90

3

Ca 1 4b 0.0000 0.2500 0.6250

W 1 4a 0.0000 0.2500 0.1250

O 1 16f 0.1504 0.0085 0.2111

Compare the two structure descriptions

**Hint:** In order to compare the different data, the parameters of Structure (a) are to be transformed to 'origin at center 2/m', i.e. ORIGIN CHOICE 2

# Example – Scheelite

## # Structure 1 – origin 1

88  
 5.243 5.243 11.376 90 90 90

3  
 Ca 1 4b 0.0000 0.0000 0.5000  
 W 1 4a 0.0000 0.0000 0.0000  
 O 1 16f 0.2413 0.1511 0.0861

 SETSTRU

## # Structure 1 – origin 2

88  
 5.2430 5.2430 11.3760 90 90 90

3  
 Ca 1 4b 0.0000 -0.2500 0.375000  
 W 1 4a 0.0000 -0.2500 -0.125000  
 O 1 16f 0.2413 -0.0989 -0.038900

## # Structure 2 – origin 2

88  
 5.243 5.243 11.376 90 90 90

3  
 Ca 1 4b 0.0000 0.2500 0.6250  
 W 1 4a 0.0000 0.2500 0.1250  
 O 1 16f 0.1504 0.0085 0.2111

AT.	WP	SS	Representative	Atomic orbit
Ca1	4b (0,1/4,5/8)	-4..	(0.000000, 0.750000, 0.375000) (0.500000, 0.250000, 0.875000) (0.000000, 0.250000, 0.625000) (0.500000, 0.750000, 0.125000)	
W1	4a (0,1/4,1/8)	-4..	(0.000000, 0.750000, 0.875000) (0.500000, 0.250000, 0.375000) (0.000000, 0.250000, 0.125000) (0.500000, 0.750000, 0.625000)	
O1	16f (x,y,z)	1	(0.241300, 0.901100, 0.961100) (0.258700, 0.098900, 0.461100) (0.848900, 0.491300, 0.211100) (0.651100, 0.508700, 0.711100) (0.758700, 0.098900, 0.038900) (0.741300, 0.901100, 0.538900) (0.151100, 0.508700, 0.788900) (0.348900, 0.491300, 0.288900) (0.741300, 0.401100, 0.461100) (0.758700, 0.598900, 0.961100) (0.348900, 0.991300, 0.711100) (0.151100, 0.008700, 0.211100) (0.258700, 0.598900, 0.538900) (0.241300, 0.401100, 0.038900) (0.651100, 0.008700, 0.288900) (0.848900, 0.991300, 0.788900)	

# Structure transformation

**TRANSTRU**

<https://www.cryst.ehu.es/cryst/tranSTRU.html>

## Transform Structure

### Transform Structure

TRANSTRU can transform a structure in two ways:

- To a lower symmetry space group. The transformed structure is given in the low symmetry space group basis, taking care of all possible splittings of the Wyckoff positions.
- With an arbitrary matrix. The structure, including the cell parameters and the atoms in the unit cell, is transformed with an arbitrary matrix introduced by the user.

Only the [default choice](#) for the conventional setting of the space groups is used.

Structure Data Examinar... No se ha seleccionado ningún archivo.  
 [in CIF format] HINT: [The option for a given filename is preferential]

```
# Space Group ITA number
221
# Lattice parameters
5.0 5.0 5.0 90 90 90
# Number of independent atoms in the asymmetric unit
3
# [atom type] [number] [WP] [x] [y] [z]
Ba 1 1a 0.0 0.0 0
Ti 2 1b 0.5 0.5 0.5
O 3 3c 0.5 0.0 0.5
```

**CIF file**

**BCS format**

Transform structure to a subgroup basis

Transform structure with an arbitrary matrix

Show

To transform a structure described in standard setting to a lower symmetry space group or with an arbitrary matrix

# Structure transformation – arbitrary matrix

**Transform Structure**

TRANSTRU transforms the structure, including the cell parameters and the atoms in the unit cell, with an arbitrary matrix introduced by the user.

**Structure**

```

221
5.0 5.0 5.0 90 90 90
3
Ba    1      1a     0.000000   0.000000   0.000000
Ti    2      1b     0.500000   0.500000   0.500000
O     3      3c     0.500000   0.000000   0.500000
  
```

**Input structure**

**Transformation matrix:**

1	0	0
0	1	0
0	0	1

**Linear part**

**Origin Shift**

0
0
0

**Show**

**Transformation matrix ( $P, p$ )**



# Structure transformation

## TRANSTRU

<https://www.cryst.ehu.es/cryst/tranSTRU.html>

**Transform Structure**

TRANSTRU can transform a structure in two ways:

- To a lower symmetry space group. The transformed structure is given in the low symmetry space group basis, taking care of all possible splittings of the Wyckoff positions.
- With an arbitrary matrix. The structure, including the cell parameters and the atoms in the unit cell, is transformed with an arbitrary matrix introduced by the user.

Only the [default choice](#) for the conventional setting of the space groups is used.

**Transform Structure**

Structure Data [Examinar...](#) No se ha seleccionado ningún archivo.  
 [in CIF format] **HINT:** [ The option for a given filename is preferential ]

```
# Space Group ITA number
221
# Lattice parameters
5.0 5.0 5.0 90 90 90
# Number of independent atoms in the asymmetric unit
3
# [atom type] [number] [WP] [x] [y] [z]
Ba 1 1a 0.0 0.0 0
Ti 2 1b 0.5 0.5 0.5
O 3 3c 0.5 0.0 0.5
```

High Symmetry Structure

**CIF file**

**BCS format**

Transform structure to a subgroup basis

Transform structure with an arbitrary matrix

[Show](#)

To transform a structure described in standard setting to a lower symmetry space group or with an arbitrary matrix

# Structure transformation – lower symmetry

**Transform Structure**

TRANSTRU transforms the structure to the low symmetry space group basis, taking care of all possible splittings of the Wyckoff positions.

**Structure**

```

221
5.0 5.0 5.0 90 90 90
3
Ba    1      1a     0.000000   0.000000   0.000000
Ti    2      1b     0.500000   0.500000   0.500000
O     3      3c     0.500000   0.000000   0.500000
  
```

**Input structure**

Low symmetry Space Group ITA number:

Transformation Matrix:

In matrix form:

Linear part			Origin Shift
0	0	1	0
1	0	0	0
0	1	0	0

Show

**Transformation matrix ( $P, p$ )**

# Output – TRANSTRU

## Transform structure

Transformation matrix: b,c,a

## High symmetry structure

```
221
5.0 5.0 5.0 90 90 90
3
Ba    1      1a      0.000000      0.000000      0.000000
Ti    2      1b      0.500000      0.500000      0.500000
O     3      3c      0.500000      0.000000      0.500000
```

[Visualize this structure](#) [CIF File](#) [Cartesian Coordinates](#)

## Low symmetry structure

```
123
5.000000 5.000000 5.000000 90.000000 90.000000 90.000000
4
Ba    1      1a      0.000000      0.000000      0.000000
Ti    2      1d      0.500000      0.500000      0.500000
O     3      2e      0.000000      0.500000      0.500000
O    3_2    1c      0.500000      0.500000      0.000000
```

[Visualize this structure](#) [CIF File](#) [Cartesian Coordinates](#)

Space Group: 123

Lattice Parameters: 5 5 5 90 90 90

AT	#	WP	Coordinates		
Ba	1	1a	0	0	0
Ti	2	1d	1/2	1/2	1/2
O	3	2e	0	1/2	1/2
O	3_2	1c	1/2	1/2	0

Detailed information

### Atoms Data:

AT.	WP	SS	Representative	Atomic orbit
Ba1	1a (0,0,0)	4/mmm	(0.000000, 0.000000, 0.000000)	(0.000000, 0.000000, 0.000000)
Ti2	1d (1/2,1/2,1/2)	4/mmm	(0.500000, 0.500000, 0.500000)	(0.500000, 0.500000, 0.500000)
O3	2e (0,1/2,1/2)	mmm .	(0.000000, 0.500000, 0.500000)	(0.000000, 0.500000, 0.500000)
O3_2	1c (1/2,1/2,0)	4/mmm	(0.500000, 0.500000, 0.000000)	(0.500000, 0.500000, 0.000000)

Note: The data shown in this table are correct if the input structure is referred to standard setting.

# Structure transformation – CIF2Standard

**CIF2Standard** <https://www.cryst.ehu.es/cryst/cif2standard.html>

Transform a given structure (in CIF format) to its description in the standard setting

**CIF to Standard**

**CIF to Standard**

CIF2Standard transforms a given structure (in CIF format) to its description in the standard setting of its space group. Its original setting is determined by analyzing its symmetry operators listed in the input CIF file.

The **default choice** of the conventional setting of the space groups is used.

This tool uses a combination of the **IDENTIFY GROUP**, **TRANSTRU** and **STRCONVERT** tools of the Bilbao Crystallographic Server, but optionally **STRUCTURE TIDY** of the **PLATON package** (after Parthe & Gelato) can also be used by checking the corresponding option in the form.

**CIF file**

Structure Data  
[in CIF format]  No file selected.

*Do the conversion via STRUCTURE TIDY implementation of the PLATON package*

# Structure transformation – CIF2Standard

## CIF to Standard Setting

The submitted structure's space group number is identified through the symmetry operators as: #15.

It has been transformed to the standard setting C2/c via the transformation matrix:  $a+c+1/4, b+1/4, c+1/4$

15						
8.8588	5.0084	7.2411	90.00	38.29	90.00	
3						
Mn	1	4a	0.000000	0.000000	0.000000	
F	1	8f	0.678050	0.209720	-0.735150	
F	2	4e	0.000000	-0.117620	-0.250000	

Transform structure  
(standard setting)

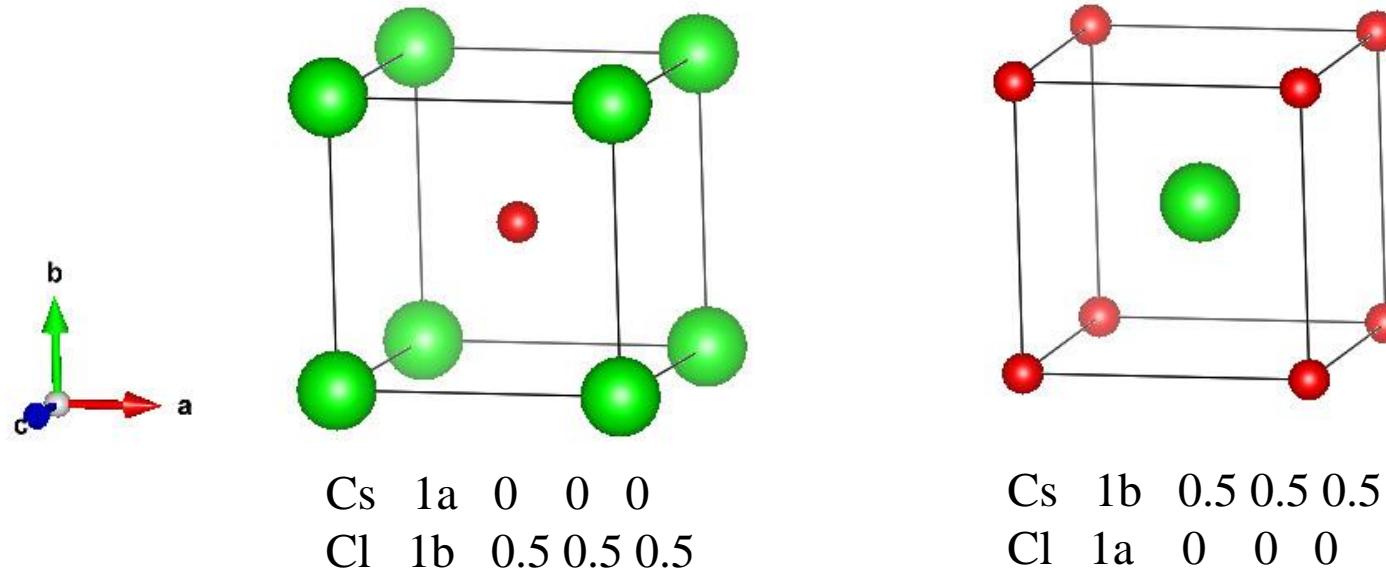
CIF file of the structure in standard setting: [cif2std\\_13517.cif](#)

Download CIF file  
(standard setting)

# Equivalent crystal structure descriptions

There are almost always several possible ways to describe the exactly same crystal structure

Space group  $Pm\bar{3}m$  (No. 221)



How many equivalent descriptions of the structure CsCl exist?

# Equivalent crystal structure descriptions

- For all space groups, except  $Im\bar{3}m$  (No. 229) and  $Ia\bar{3}d$  (No. 230), one can choose several different sets of atomic coordinates describing the same structure in the same space-group setting.
- The number of equivalent descriptions can be calculated:

$$[i] = \frac{|N_\varepsilon(G)|}{|G|}$$

$N_\varepsilon(G)$  represents the Euclidean normalizer of the space group G

- By definition,  $i$  cosets result in the coset decomposition of  $N_\varepsilon(G)$  with respect to  $G$ . The cosets generate the different equivalent descriptions of a given structure

# Example – CsCl

- CsCl crystallizes in space group  $Pm\bar{3}m$  (No. 221)

**Euclidean normalizer (general metric) of the Group  $Pm\bar{3}m$  (No. 221)**

**Euclidean normalizer of  $Pm\bar{3}m$  ( $a,b,c$ ):  $Im\bar{3}m$  ( $a,b,c$ )**

Index of  $Pm\bar{3}m$  in  $Im\bar{3}m$  ( $a,b,c$ ): 2 with  $i_L=2$  and  $i_P=1$

Additional generators of  $Im\bar{3}m$  ( $a,b,c$ ) with respect to  $Pm\bar{3}m$

$x+1/2, y+1/2, z+1/2$	$\begin{pmatrix} 1 & 0 & 0 & 1/2 \\ 0 & 1 & 0 & 1/2 \\ 0 & 0 & 1 & 1/2 \end{pmatrix}$	$t(1/2, 1/2, 1/2)$
-----------------------	---------------------------------------------------------------------------------------	--------------------

The cosets representatives of the Euclidean normalizer  $Im\bar{3}m$  ( $a,b,c$ ) with respect to  $Pm\bar{3}m$

- As  $[i] = 2 \Rightarrow$  there are two possible sets of coordinates
- One set of coordinates is obtained from the other one according to the additional generators of  $N_\varepsilon(G)$

# Example – CsCl

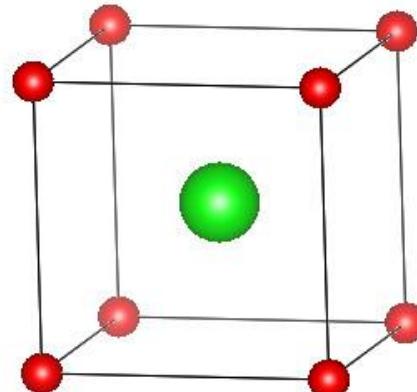
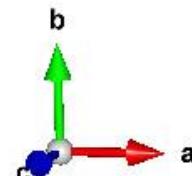
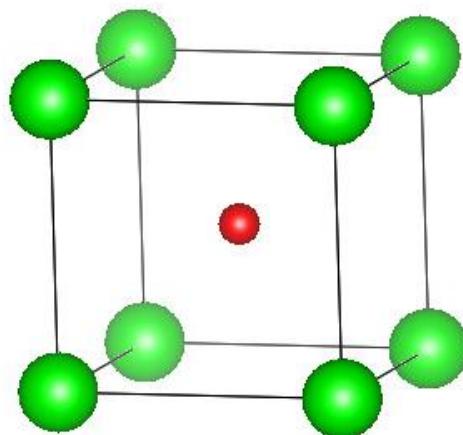
- CsCl crystallizes in space group  $Pm\bar{3}m$  (No. 221)

Cs	1a	0	0	0
Cl	1b	0.5	0.5	0.5

$t(1/2,1/2,1/2)$



Cs	1b	0.5	0.5	0.5
Cl	1a	0	0	0



# Equivalent crystal structure descriptions

**EQUIVSTRU** <https://www.cryst.ehu.es/cryst/equivSTRU.html>

**Equivalent Structures**

Given a space group ITA number, the cell parameters (separated with spaces) and the atom positions, the program EQUIVSTRU transforms the corresponding structure with the elements of the euclidean normalizer of the space group. All the transformed structures are equivalent symmetry descriptions of the given initial structure. The atom positions are identified generating the Wyckoff sets.

Only the default choice for the conventional setting of

Structure Data  No se ha seleccionado ningún archivo.  
 [in CIF format] **HINT:** [The option for a given filename is preferential]

```
#Exercise 2.4.2a(CsCl)
# Space Group ITA number
221
# Lattice parameters
4.12599 4.12599 4.12599 90.0 90.0 90.0
# Number of independent atoms in the asymmetric unit
2
# [atom type] [number] [WP] [x] [y] [z]
Cl    1    1a    0.000000 0.000000 0.000000
Cs    1    1b    0.500000 0.500000 0.500000

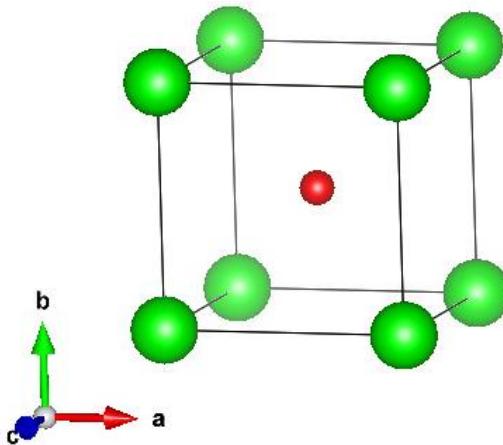
```

Structure

**CIF file**

**BCS format**

# EQUIVSTRU – Output



- Equivalent description 1 (original input structure)

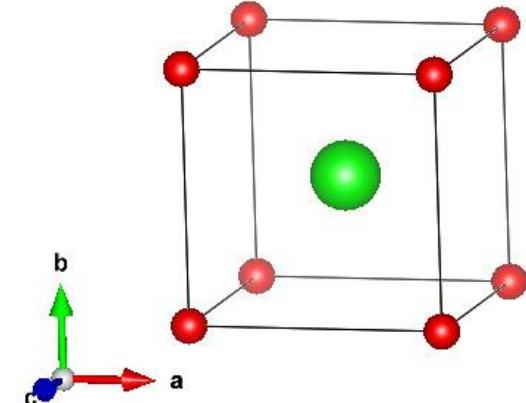
**Normalizer coset representative: x,y,z**

AT.	WP	SS	Representative	Atomic orbit
Cl1	1a (0,0,0)	m-3m	(0.000000, 0.000000, 0.000000)	(0.000000, 0.000000, 0.000000)
Cs1	1b (1/2,1/2,1/2)	m-3m	(0.500000, 0.500000, 0.500000)	(0.500000, 0.500000, 0.500000)

- Equivalent description 2

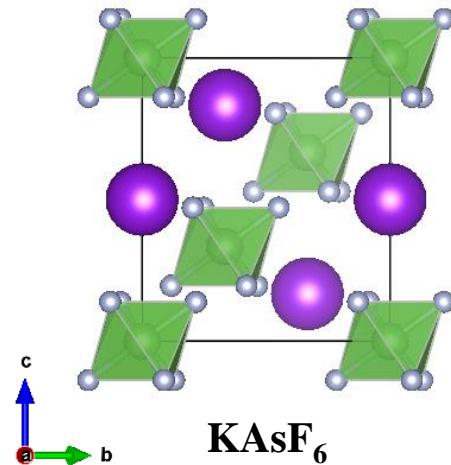
**Normalizer coset representative: x+1/2,y+1/2,z+1/2**

AT.	WP	SS	Representative	Atomic orbit
Cl1	1b (1/2,1/2,1/2)	m-3m	(0.500000, 0.500000, 0.500000)	(0.500000, 0.500000, 0.500000)
Cs1	1a (0,0,0)	m-3m	(0.000000, 0.000000, 0.000000)	(0.000000, 0.000000, 0.000000)

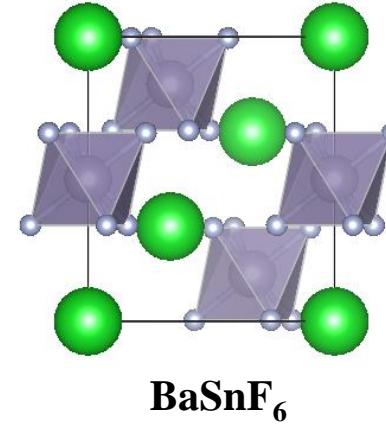


# Example – similarity

Evaluate the similarity between these two structures. Try to find analogous coordinate descriptions.



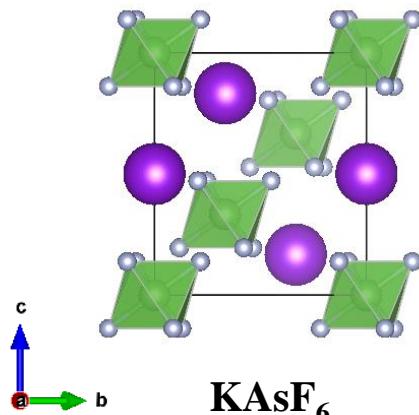
```
#ICSD: 59413
148
7.348 7.348 7.274 90. 90. 120.
3
K 1 3b 0.333333 0.666670 0.166670
As 1 3a 0.000000 0.000000 0.000000
F 1 18f 0.129200 0.216500 0.138100
```



```
#ICSD: 33788
148
7.4279 7.4279 7.418 90. 90. 120.
3
Ba 1 3a 0.000000 0.000000 0.000000
Sn 1 3b 0.000000 0.000000 0.500000
F 1 18f 0.258600 0.826200 0.004700
```

$R\bar{3}$  (No. 148)  $\Rightarrow$  4 equivalent descriptions:  $x, y, z$ ;  $x, y, z + 1/2$ ;  $y, x, -z$ ;  $y, x, -z + 1/2$

# Example – similarity



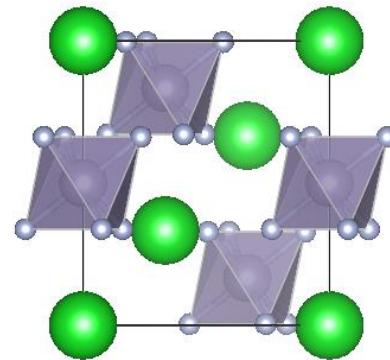
#ICSD: 59413

148

7.348 7.348 7.274 90. 90. 120.

3

K 1 3b 0.333333 0.666670 0.166670  
 As 1 3a 0.000000 0.000000 0.000000  
 F 1 18f 0.129200 0.216500 0.138100



**BaSnF<sub>6</sub>**

#ICSD: 33788

148

7.4279 7.4279 7.418 90. 90. 120.

3

Ba 1 3a 0.000000 0.000000 0.000000  
 Sn 1 3b 0.000000 0.000000 0.500000  
 F 1 18f 0.258600 0.826200 0.004700



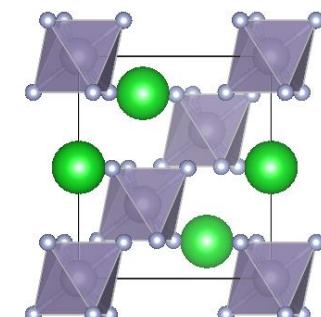
*x, y, z+1/2*

148

7.4279 7.4279 7.4180 90.00 90.00 120.00

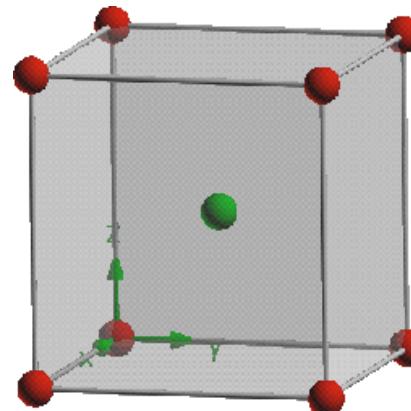
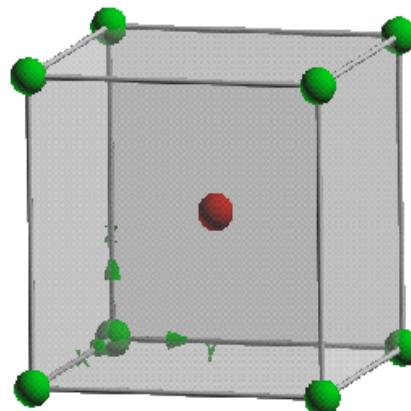
3

Ba 1 3b 0.333333 0.666667 0.166667  
 Sn 1 3a 0.000000 0.000000 0.000000  
 F 1 18f 0.159533 0.234267 0.161967



# Comparison of structures

- Comparison of crystal structures is convenient to:
  - cross-check different experimental and/or theoretical structural models of the same phase coming from different sources
  - identify different phases with the same symmetry
  - classify structures into structure types
- The existence of various equivalent structure descriptions makes the comparison of different structural models a non-trivial task in general.



Are these two structures  
of Cs Cl similar?

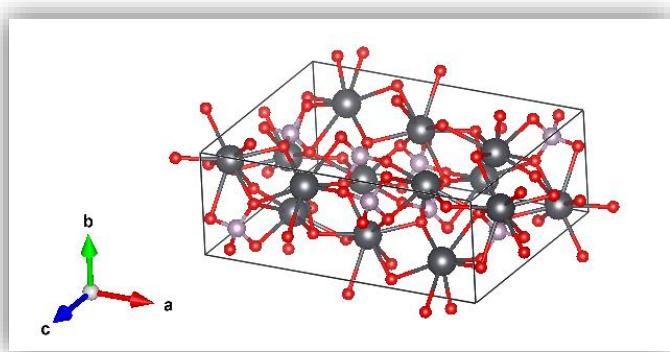
Yes

# Similarity between two crystal structures

## #Structure 1

```

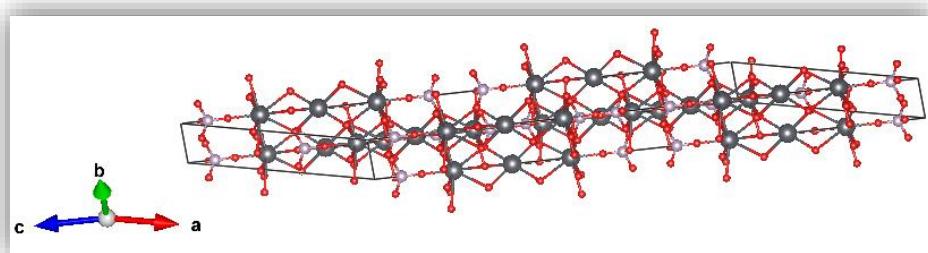
15
13.800 5.691 9.420 90.0 102.3 90.0
7
Pb    1      4e      0.000000      0.291000      0.250000
Pb    2      8f      0.317000      0.309000      0.352000
P     1      8f      0.599000      0.241000      0.447000
O     1      8f      0.643000      0.030000      0.392000
O     2      8f      0.634000      0.464000      0.374000
O     3      8f      0.642000      0.280000      0.612000
O     4      8f      0.491000      0.222000      0.420000
  
```



## #Structure 2

```

15
13.967 5.560 40.778 90.0 166.713 90.0
7
Pb    1      4e      0.000000      0.000000      0.750000
Pb    2      8f      0.000000      0.000000      0.856300
P     1      8f      0.000000      0.000000      0.951100
O     1      8f      0.000000      0.000000      0.914500
O     2      8f      0.271500      0.728500      0.888500
O     3      8f      0.957000      0.500000      0.117000
O     4      8f      0.728500      0.271500      0.611500
  
```



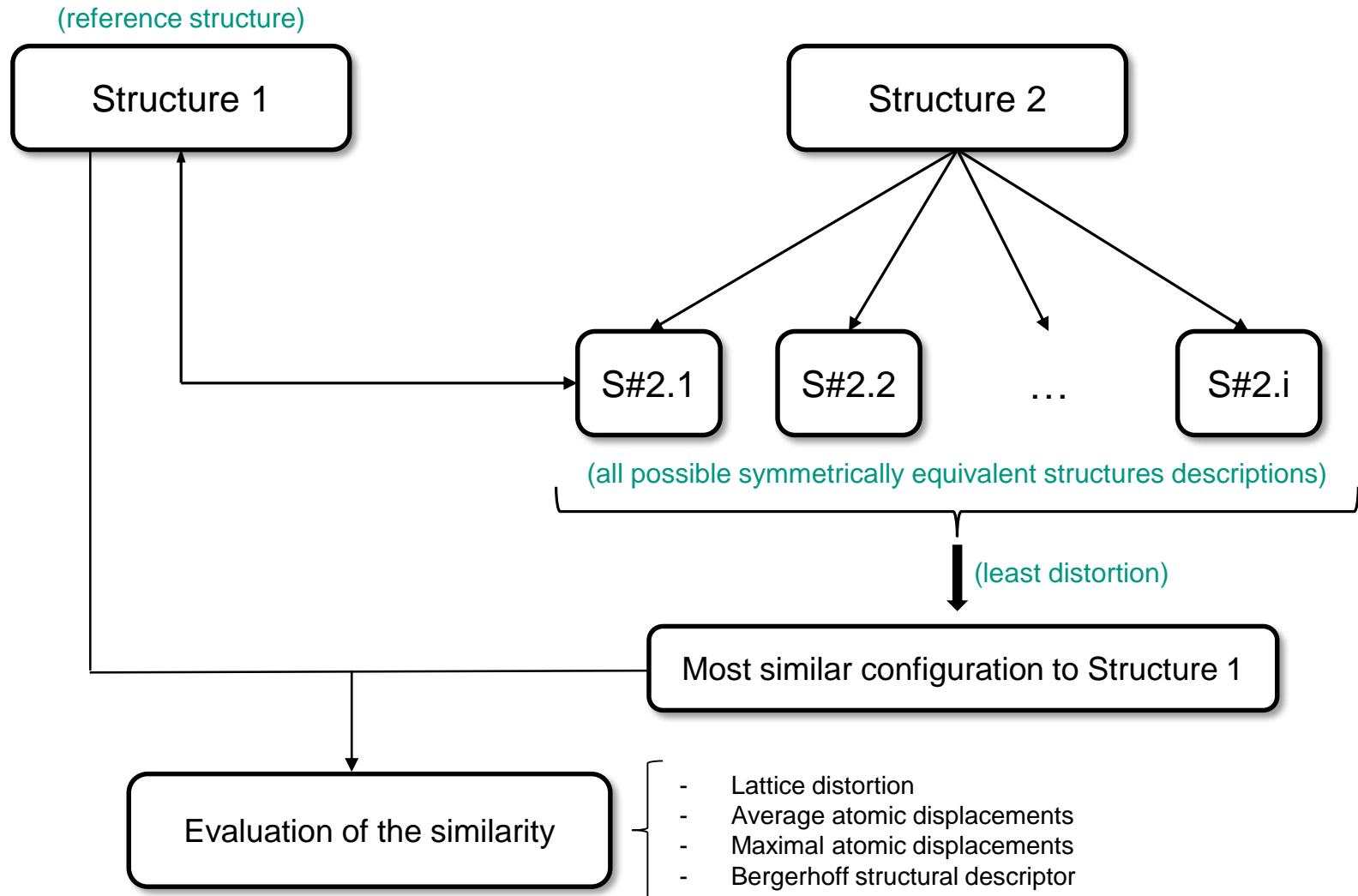
Are these two structures similar?

# Crystal Structure Relationships

## Structure Utilities

<b>CELLTRAN</b>	Transform Unit Cells
<b>STRAIN</b>	Strain Tensor Calculation
<b>WPASSIGN</b>	Assignment of Wyckoff Positions
<b>TRANSTRU</b>	Transform structures.
<b>SETSTRU</b>	Alternative Settings for a given Crystal Structure
<b>EQUIVSTRU</b>	Equivalent Descriptions for a given Crystal Structure
<b>CIF2STANDARD</b>	Transforms a given CIF (in any setting) to that of standard setting
<b>STRCONVERT</b>	Convert & Edit Structure Data (supports the CIF, mCIF, VESTA, VASP formats -- with magnetic information where available)
<b>VISUALIZE</b>	Visualize structures using Jmol
<b>COMPSTRU</b>	Comparison of Crystal Structures with the same Symmetry
<b>STRUCTURE RELATIONS</b>	Evaluation of structure relationships [transformation matrix] between group-subgroup related phases
<b>PSEUDOLATTICE</b>	Pseudosymmetry of a lattice and compatible supergroups

# The program COMPSTRU



# The program COMPSTRU

- How to measure the **similarity** between two descriptions ?

**degree of lattice distortion**

$$S = \frac{1}{3} \sqrt{\sum_i \eta_i^2}$$

$\eta_i$ -eigenvalues of the Lagrangian strain tensor

**average atomic displacements**

$$d_{av} = \frac{1}{n} \sum_i m_i u_i$$

$u_i$  atomic displacements

**maximal atomic displacements**

maximal displacements of the paired atoms

# The program COMPSTRU

- How to measure the **similarity** between two descriptions ?

**structural  
descriptor**

$$\Delta = [2^{1/2} \Delta(c) + 1] \Delta(d) - 1$$



$$\Delta(c) = \frac{\sum m[(x_1 - x_2)^2 + (y_1 - y_2)^2 + (z_1 - z_2)^2]^{1/2}}{\sum m}$$

weighted mean difference  
between atomic coordinates

$$\Delta(d) = \frac{[(b_1/a_1)(c_1/a_1)]}{[(b_2/a_2)(c_2/a_2)]}$$

relation between  
axial ratios

Bergerhoff *et al.*, Acta Cryst B55 (1998)

# Input – COMPSTRU

## COMPSTRU:

<https://www.cryst.ehu.es/cryst/compstru.html>

## Input:

Two crystal structures described  
in the standard setting

- Isopointal structures with the same/different composition
- Chiral structures

Structure Data  
[in CIF format]

Examinar... No se ha seleccionado ningún archivo.  
HINT: [The option for a given filename is preferential]

```
15
13.800 5.691 9.420 90.0 102.3 90.0
7
Pb 1 4e 0.0000 0.2910 0.2500
Pb 2 8f 0.3170 0.3090 0.3520
P 1 8f 0.5990 0.2410 0.4470
O 1 8f 0.6430 0.0300 0.3920
O 2 8f 0.6340 0.4640 0.3740
O 3 8f 0.6420 0.2800 0.6120
O 4 8f 0.4910 0.2220 0.4200
```

CIF file

Structure Data  
[in CIF format]

Examinar... No se ha seleccionado ningún archivo.  
HINT: [The option for a given filename is preferential]

```
15
13.967 5.560 40.778 90.0 166.713 90.0
7
Pb 1 4e 0.0000 0.0000 0.7500
Pb 2 8f 0.0000 0.0000 0.8563
P 1 8f 0.0000 0.0000 0.9511
O 1 8f 0.0000 0.0000 0.9145
O 2 8f 0.2715 0.7285 0.8885
O 3 8f 0.9570 0.5000 0.1170
O 4 8f 0.7285 0.2715 0.6115
```

BCS format

Structure Data  
[in CIF format]

Examinar... No se ha seleccionado ningún archivo.  
HINT: [The option for a given filename is preferential]

```
15
13.967 5.560 40.778 90.0 166.713 90.0
7
Pb 1 4e 0.0000 0.0000 0.7500
Pb 2 8f 0.0000 0.0000 0.8563
P 1 8f 0.0000 0.0000 0.9511
O 1 8f 0.0000 0.0000 0.9145
O 2 8f 0.2715 0.7285 0.8885
O 3 8f 0.9570 0.5000 0.1170
O 4 8f 0.7285 0.2715 0.6115
```

CIF file

Enter the maximum distance allowed between the paired atoms:  Å

Enter the allowed tolerance (a b c α β γ):

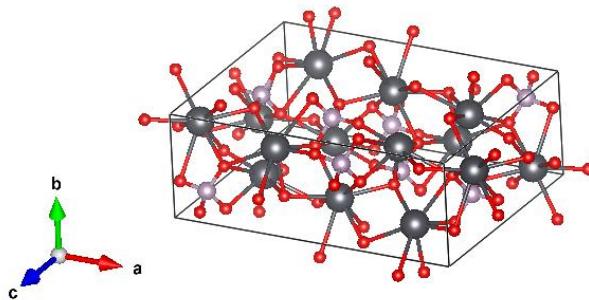
BCS format

Tolerance

# Are these two structures similar?

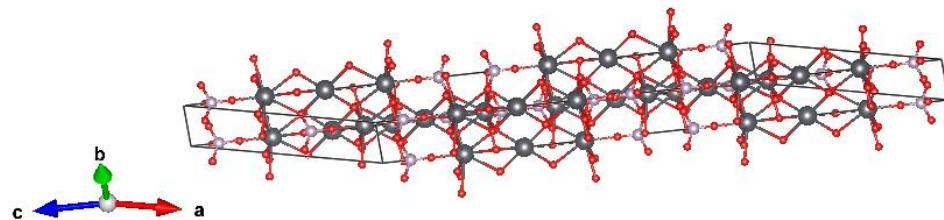
**Pb<sub>3</sub>(PO<sub>4</sub>)<sub>2</sub>**

#Structure 1



15  
 13.800 5.691 9.420 90.0 102.3 90.0  
 7  
 Pb 1 4e 0.0000 0.2910 0.2500  
 Pb 2 8f 0.3170 0.3090 0.3520  
 P 1 8f 0.5990 0.2410 0.4470  
 O 1 8f 0.6430 0.0300 0.3920  
 O 2 8f 0.6340 0.4640 0.3740  
 O 3 8f 0.6420 0.2800 0.6120  
 O 4 8f 0.4910 0.2220 0.4200

#Structure 2



15  
 13.967 5.560 40.778 90.0 166.713  
 90.0  
 7  
 Pb 1 4e 0.0000 0.0000 0.7500  
 Pb 2 8f 0.0000 0.0000 0.8563  
 P 1 8f 0.0000 0.0000 0.9511  
 O 1 8f 0.0000 0.0000 0.9145  
 O 2 8f 0.2715 0.7285 0.8885  
 O 3 8f 0.9570 0.5000 0.1170  
 O 4 8f 0.7285 0.2715 0.6115

# COMPSTRU

Pb<sub>3</sub>(PO<sub>4</sub>)<sub>2</sub>

## Comparison of crystal structures of the same symmetry C2/c ( No. 15 ) [ unique axis b ]

### Structure #1

```

15
13.800 5.691 9.420 90.0 102.3 90.0
7
Pb    1      4e      0.000000      0.291000      0.250000
Pb    2      8f      0.317000      0.309000      0.352000
P     1      8f      0.599000      0.241000      0.447000
O     1      8f      0.643000      0.030000      0.392000
O     2      8f      0.634000      0.464000      0.374000
O     3      8f      0.642000      0.280000      0.612000
O     4      8f      0.491000      0.222000      0.420000

```

### Structure #2

```

15
13.967 5.560 40.778 90.0 166.713 90.0
7
Pb    1      4e      0.000000      0.000000      0.000000      0.750000
Pb    2      8f      0.000000      0.000000      0.000000      0.856300
P     1      8f      0.000000      0.000000      0.000000      0.951100
O     1      8f      0.000000      0.000000      0.000000      0.914500
O     2      8f      0.271500      0.728500      0.728500      0.888500
O     3      8f      0.957000      0.500000      0.500000      0.117000
O     4      8f      0.728500      0.271500      0.271500      0.611500

```

The next step is to select a transformation that best matches the lattice parameters of the structures for the chosen tolerance (a b c  $\alpha$   $\beta$   $\gamma$ ) = (.5 .5 .5 5 5 5):

Select	Transformation (P,p)	Cell parameters of the Structure #1 Cell parameters of the Structure #2 applying the transformation matrix	Strain
<input checked="" type="radio"/>	-a,-b,3a+c	13.8000 5.6910 9.4200 90.0000 102.3000 90.0000 13.9670 5.5600 9.6301 90.0000 103.2951 90.0000	0.0116

Continue

## Structure #1

15  
13.800 5.691 9.420 90.0 102.3 90.0

7

Pb	1	4e	0.000000	0.291000	0.250000
Pb	2	8f	0.317000	0.309000	0.352000
P	1	8f	0.599000	0.241000	0.447000
O	1	8f	0.643000	0.030000	0.392000
O	2	8f	0.634000	0.464000	0.374000
O	3	8f	0.642000	0.280000	0.612000
O	4	8f	0.491000	0.222000	0.420000

WP	Atom	Atomic Displacements				
		u <sub>x</sub>	u <sub>y</sub>	u <sub>z</sub>	u	
4e	(0,y,1/4)	Pb1	0.0000	-0.0410	0.0000	0.2333
8f	(x,y,z)	Pb2	0.0019	-0.0590	0.0043	0.3386
8f	(x,y,z)	P1	0.0043	0.0090	0.0041	0.0816
8f	(x,y,z)	O1	0.0010	-0.0085	-0.0035	0.0617
8f	(x,y,z)	O2	0.0100	0.0145	0.0145	0.1910
8f	(x,y,z)	O3	0.0020	-0.0300	0.0050	0.1777
8f	(x,y,z)	O4	0.0025	0.0280	-0.0055	0.1733

## Evaluation of the structure similarity

S	d <sub>max.</sub> (Å)	d <sub>av.</sub> (Å)	Δ
0.0116	0.3386	0.1755	0.066

## Structure #2

Pb<sub>3</sub>(PO<sub>4</sub>)<sub>2</sub>

15  
13.967 5.560 40.778 90.0 166.713 90.0

7

Pb	1	4e	0.000000	0.000000	0.750000
Pb	2	8f	0.000000	0.000000	0.856300
P	1	8f	0.000000	0.000000	0.951100
O	1	8f	0.000000	0.000000	0.914500
O	2	8f	0.271500	0.728500	0.888500
O	3	8f	0.957000	0.500000	0.117000
O	4	8f	0.728500	0.271500	0.611500

$$(P, p) = -a, -b, 3a + c; 1/4, 1/4, 0$$

Description of Structure #2 in the most similar configuration to Structure #1

015  
13.967000 5.560000 9.630055 90.000000 103.295059 90.000000

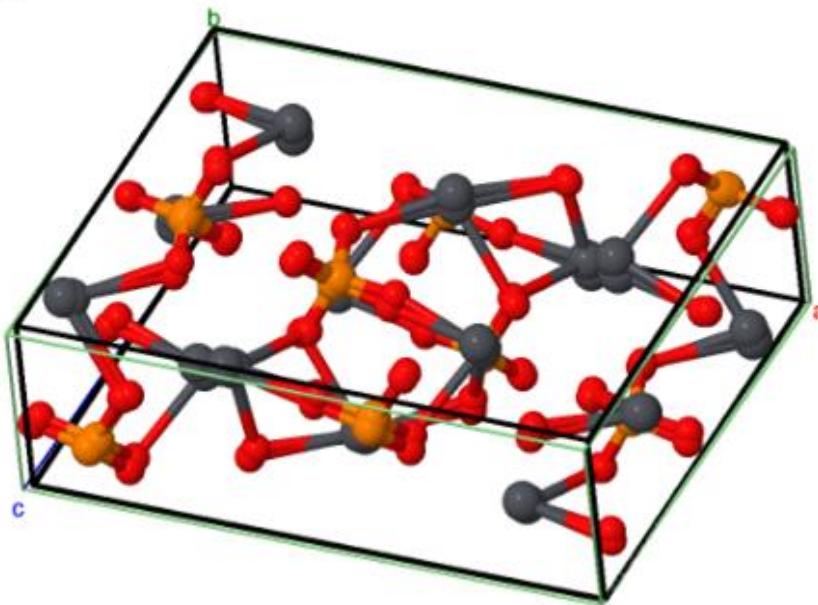
7

Pb	1	4e	0.500000	0.250000	0.750000
Pb	2	8f	0.818900	0.250000	0.856300
P	1	8f	0.103300	0.250000	0.951100
O	1	8f	0.993500	0.250000	0.914500
O	2	8f	0.644000	0.521500	0.888500
O	3	8f	0.644000	0.750000	0.117000
O	4	8f	0.356000	0.978500	0.611500

# Input – COMPSTRU

Pb<sub>3</sub>(PO<sub>4</sub>)<sub>2</sub>

```
-C 2yc [C 1 2/c 1] #15
a=13.967Å
b=5.560Å
c=9.630Å
α=90.000°
β=103.295°
γ=90.000°
```



Structure #1

Structure #2

Structure #2 (most similar)

opaque

ball&stick  stick  cross

opaque

ball&stick  stick  cross

opaque

ball&stick  stick  cross

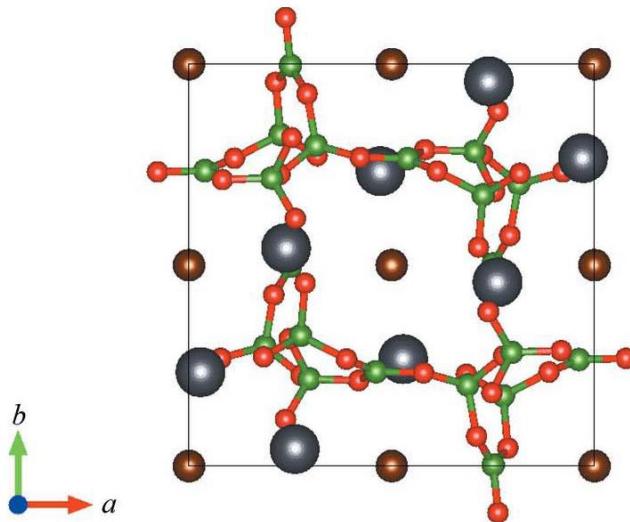
Show Distances

cutoff: 0.5

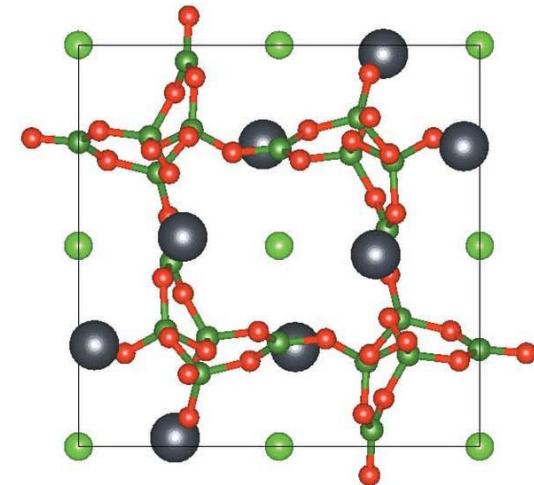
Save PNG+Jmol

# Example: Structures with different composition

Are these two  
structures equivalent?



Belokoneva *et al.*, 2003



Egorova *et al.*, 2008

Comparison of isopointal structures with different composition

# Example: Structures with different composition

Comparison of crystal structures of the same symmetry *Pnn2* ( No. 34 )

**Structure #1**

```

34
11.524 11.431 6.5399 90. 90. 90.
18
Pb   1    4c    0.260700    0.042500    0.000000
Pb   2    4c    0.028000    0.233700    0.663000
Br   1    2a    0.000000    0.000000    0.893000
Br   2    2b    0.000000    0.500000    0.626000
O    1    4c    0.247000    0.317000    0.590000
O    2    4c    0.209000    0.426000    0.179000
O    3    4c    0.276000    0.226000    0.253000
O    4    4c    0.078000    0.268000    0.241000
O    5    4c    0.286000    0.455000    0.853000
O    6    4c    0.388000    0.266000    0.842000
O    7    4c    0.184000    0.272000    0.939000
O    8    4c    0.418000    0.213000    0.500000
O    9    4c    0.238000    0.116000    0.570000
B    1    4c    0.275000    0.325000    0.800000
B    2    4c    0.187000    0.298000    0.160000
B    3    4c    0.296000    0.217000    0.480000
B    4    4c    0.461000    0.235000    0.700000
B    5    4c    0.250000    0.497000    0.030000
  
```

**Structure #2**

```

34
11.3810 11.3840 6.56335 90. 90. 90.
18
Pb   1    4c    0.039700    0.252300    0.000100
Pb   2    4c    0.240500    0.021900    0.332600
Cl   1    2a    0.000000    0.000000    0.121800
Cl   2    2b    0.500000    0.000000    0.381300
O    1    4c    0.318400    0.238700    0.418600
O    2    4c    0.427100    0.212600    0.826900
O    3    4c    0.225600    0.271700    0.753700
O    4    4c    0.274800    0.072000    0.752600
O    5    4c    0.448100    0.286300    0.160800
O    6    4c    0.260500    0.380800    0.162500
O    7    4c    0.273900    0.180300    0.069800
O    8    4c    0.210400    0.419200    0.506500
O    9    4c    0.115700    0.232100    0.438700
B    1    4c    0.322100    0.271200    0.203400
B    2    4c    0.301500    0.181600    0.854700
B    3    4c    0.217300    0.288900    0.534400
B    4    4c    0.230700    0.458800    0.308800
B    5    4c    0.498800    0.258300    0.976800
  
```

Select	Transformation (P,p)	Cell parameters of the Structure #1 Cell parameters of the Structure #2 applying the transformation matrix	Strain
<input type="radio"/>	a,b,c	11.5240 11.4310 6.5399 90.0000 90.0000 90.0000 11.3810 11.3840 6.5633 90.0000 90.0000 90.0000	0.0046
<input checked="" type="radio"/>	-b,-a,-c	11.5240 11.4310 6.5399 90.0000 90.0000 90.0000 11.3840 11.3810 6.5633 90.0000 90.0000 90.0000	0.0045

All equivalent unit-cell parameters are calculated and compared with the unit-cell parameters of structure 1

# Example: Structures with different composition

## Comparison of crystal structures of the same symmetry *Pnn2* ( No. 34 )

**Structure 1**

```

34
11.3810 11.3840 6.56335 90. 90. 90.
18
Pb   1   4c    0.039700 0.252300 0.000100
Pb   2   4c    0.240500 0.021900 0.332600
Cl   1   2a    0.000000 0.000000 0.121800
Cl   2   2b    0.500000 0.000000 0.381300
O    1   4c    0.318400 0.238700 0.418600
O    2   4c    0.427100 0.212600 0.826900
O    3   4c    0.225600 0.271700 0.753700
O    4   4c    0.274800 0.072000 0.752600
O    5   4c    0.448100 0.286300 0.160800
O    6   4c    0.260500 0.380800 0.162500
O    7   4c    0.273900 0.180300 0.069800
O    8   4c    0.210400 0.419200 0.506500
O    9   4c    0.115700 0.232100 0.438700
B    1   4c    0.322100 0.271200 0.203400
B    2   4c    0.301500 0.181600 0.854700
B    3   4c    0.217300 0.288900 0.534400
B    4   4c    0.230700 0.458800 0.308800
B    5   4c    0.498800 0.258300 0.976800
  
```

**Structure 2**

```

34
11.524 11.431 6.5399 90. 90. 90.
18
Pb   1   4c    0.260700 0.042500 0.000000
Pb   2   4c    0.028000 0.233700 0.663000
Br   1   2a    0.000000 0.000000 0.893000
Br   2   2b    0.000000 0.500000 0.626000
O    1   4c    0.247000 0.317000 0.590000
O    2   4c    0.209000 0.426000 0.179000
O    3   4c    0.276000 0.226000 0.253000
O    4   4c    0.078000 0.268000 0.241000
O    5   4c    0.286000 0.455000 0.853000
O    6   4c    0.388000 0.266000 0.842000
O    7   4c    0.184000 0.272000 0.939000
O    8   4c    0.418000 0.213000 0.500000
O    9   4c    0.238000 0.116000 0.570000
B    1   4c    0.275000 0.325000 0.800000
B    2   4c    0.187000 0.298000 0.160000
B    3   4c    0.296000 0.217000 0.480000
B    4   4c    0.461000 0.235000 0.700000
B    5   4c    0.250000 0.497000 0.030000
  
```

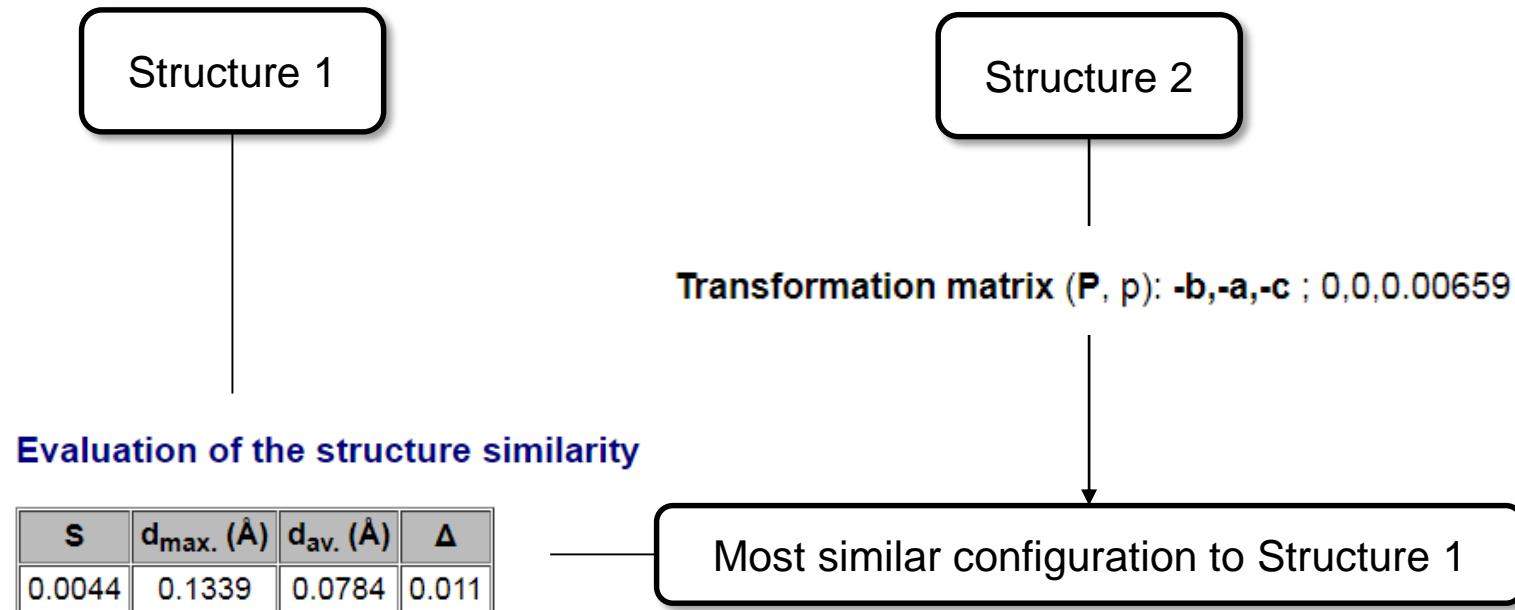
The similarity of the two structures will be evaluated taking into account the following correspondence between the atomic species:

Structure #1	Structure #2
Cl	Br
O	O
Pb	Pb
B	B

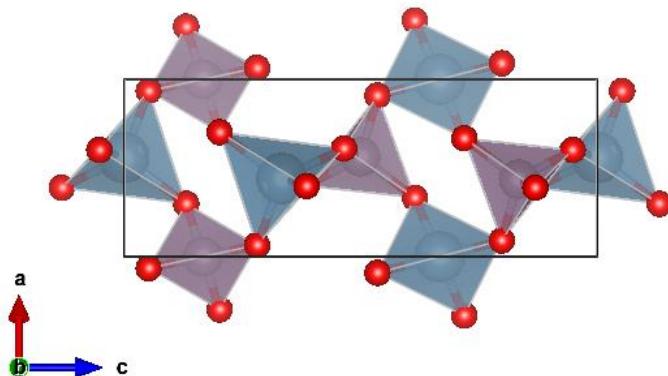
Do you agree with the proposed correspondence scheme?

**DIFFERENT  
COMPOSITION**

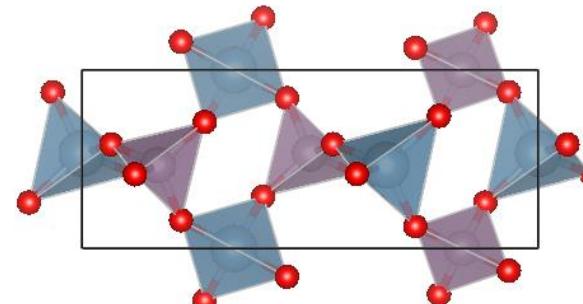
# Example: Structures with different composition



# Example: Chiral Structures



**P3<sub>1</sub>21 (No. 152)**



**P3<sub>2</sub>21 (No. 154)**

#ICSD: 158620

152

4.5191 4.5191 10.471 90. 90. 120.

4

Al	1	3a	0.441200	0.000000	0.333333
P	1	3b	0.437300	0.000000	0.833300
O	1	6c	0.398200	0.332600	0.385500
O	2	6c	0.389200	0.297900	0.868700

#ICSD: 50100

154

4.9438 4.9438 10.9498 90. 90. 120.

4

Al	1	3a	0.466460	0.000000	0.666667
P	1	3b	0.466900	0.000000	0.166700
O	1	6c	0.416400	0.291900	0.602540
O	2	6c	0.415500	0.257400	0.116180

# Example: Chiral Structures

## Structure #1

```

152
4.5191 4.5191 10.471 90. 90. 120.
4
Al    1      3a     0.441200   0.000000   0.333333
P     1      3b     0.437300   0.000000   0.833300
O     1      6c     0.398200   0.332600   0.385500
O     2      6c     0.389200   0.297900   0.868700
  
```

## Atom pairings and distances

Atom Mappings					
WP	Atom	Coordinates in S <sub>1</sub>		Atom	Coordinates in S <sub>2</sub>
3a (x,0,1/3)	Al1	(0.441200,0.000000,0.333333)		Al1	(0.466460,0.000000,0.333333)
3b (x,0,5/6)	P1	(0.437300,0.000000,0.833300)		P1	(0.466900,0.000000,0.833300)
6c (x,y,z)	O1	(0.398200,0.332600,0.385500)		O1	(0.416400,0.291900,0.397460)
6c (x,y,z)	O2	(0.389200,0.297900,0.868700)		O2	(0.415500,0.257400,0.883820)

WP	Atom	Atomic Displacements			
		u <sub>x</sub>	u <sub>y</sub>	u <sub>z</sub>	u
3a (x,0,1/3)	Al1	0.0253	0.0000	0.0000	0.1142
3b (x,0,5/6)	P1	0.0296	0.0000	0.0000	0.1338
6c (x,y,z)	O1	0.0182	-0.0407	0.0120	0.2672
6c (x,y,z)	O2	0.0263	-0.0405	0.0151	0.3073

## Structure #2

```

154
4.94388 4.94388 10.9498 90. 90. 120.
4
Al    1      3a     0.466460   0.000000   0.666667
P     1      3b     0.466900   0.000000   0.166700
O     1      6c     0.416400   0.291900   0.602540
O     2      6c     0.415500   0.257400   0.116180
  
```

Transformation matrix (P, p): a,b,-c ; 0,0,0



The enantiomorphic description of Structure #2 in the most similar configuration to Structure #1

```

152
4.943800 4.943800 10.949800 90.000000 90.000000 120.000000
4
Al    1      3a     0.466460   0.000000   0.333333
P     1      3b     0.466900   0.000000   0.833300
O     1      6c     0.416400   0.291900   0.397460
O     2      6c     0.415500   0.257400   0.883820
  
```

S	d <sub>max.</sub> (Å)	d <sub>av.</sub> (Å)	Δ
0.0413	0.3073	0.2328	0.107

# Isopointal and isoconfigurational structures

- Two structures are defined as *isopointal* if:
  - (1) they have the same space-group type or belong to a pair of enantiomorphic space groups, and
  - (2) the atomic positions are the same in both structures
  
- Two structures are defined as *isoconfigurational* (or belonging to the same structure type) if
  - (1) they are isopointal, and
  - (2) for all corresponding Wyckoff positions, both, the crystallographic configurations (crystallographic orbits) and their geometric interrelationships, are similar.

Lima-de Faria *et al.* *Acta Cryst.* (1990), A41, 1

# Structure types - COMPSTRU

**ABX<sub>6</sub> family R-3 (148); WP sequence: fba; Pearson: hR8**

KCrF <sub>6</sub>	LiNbF <sub>6</sub>	VNbF <sub>6</sub>	HgRhF <sub>6</sub>	MgPbF <sub>6</sub>	InAsF <sub>6</sub>
RbCrF <sub>6</sub>	LiRuF <sub>6</sub>	CoZrF <sub>6</sub>	NiRhF <sub>6</sub>	ZnPbF <sub>6</sub>	CsNbF <sub>6</sub>
KAsF <sub>6</sub>	LiRhF <sub>6</sub>	PdPtF <sub>6</sub>	CaCrF <sub>6</sub>	NiPbF <sub>6</sub>	HgCrF <sub>6</sub>
RuAsF <sub>6</sub>	LiTaF <sub>6</sub>	FeNbF <sub>6</sub>	MgCrF <sub>6</sub>	MgPdF <sub>6</sub>	CoSnF <sub>6</sub>
CsAsF <sub>6</sub>	LiOsF <sub>6</sub>	CaSnF <sub>6</sub>	CdCrF <sub>6</sub>	CaPdF <sub>6</sub>	CsNbF <sub>6</sub>
RbSbF <sub>6</sub>	LilrF <sub>6</sub>	FeZrF <sub>6</sub>	MnSnF <sub>6</sub>	ZnPdF <sub>6</sub>	MnPtF <sub>6</sub>
BaSnF <sub>6</sub>	LiPtF <sub>6</sub>	CuZrF <sub>6</sub>	FeSnF <sub>6</sub>	CdPdF <sub>6</sub>	CdRhF <sub>6</sub>
CsBrF <sub>6</sub>	LiAuF <sub>6</sub>	CaPtF <sub>6</sub>	ZnSnF <sub>6</sub>	LiSbF <sub>6</sub>	NaBiF <sub>6</sub>
CsSbF <sub>6</sub>	NiPtF <sub>6</sub>	ZnPtF <sub>6</sub>	NiSnF <sub>6</sub>	BalrF <sub>6</sub>	TlAsF <sub>6</sub>
CsBiF <sub>6</sub>	CdPtF <sub>6</sub>	CoPtF <sub>6</sub>	CuSnF <sub>6</sub>	RbBiF <sub>6</sub>	
CsUF <sub>6</sub>	LiPF <sub>6</sub>	MgRhF <sub>6</sub>	CdSnF <sub>6</sub>	KRhF <sub>6</sub>	
KOsF <sub>6</sub>	LiAsF <sub>6</sub>	CaRhF <sub>6</sub>	CdTlF <sub>6</sub>	CsReF <sub>6</sub>	
NaCrF <sub>6</sub>	PdZrF <sub>6</sub>	ZnRhF <sub>6</sub>	LiBiF <sub>6</sub>	KPF <sub>6</sub>	

