

Summer School on Mathematical Crystallography

3-7 June 2019, Nancy (France)

**International Union of Crystallography Commission on Mathematical
and Theoretical Crystallography**



C2MP



métropole
GrandNancy

CRYSTAL-STRUCTURE DESCRIPTIONS

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Karlsruhe Institute of Technology



CRYSTAL-STRUCTURE TOOLS

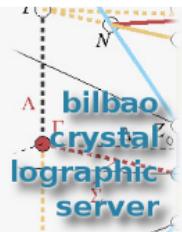
Crystal structure descriptions

Structure transformations

Equivalent structure descriptions

Comparison between different structure descriptions





Bilbao Crystallographic
Server
in forthcoming schools and
workshops

News:

- **New Article in Acta Cryst. A** 05/2019: Gallego et al. "Automatic calculation of symmetry-adapted tensors in magnetic and non-magnetic materials: a new tool of the Bilbao Crystallographic Server." *Acta Cryst.* (2019) A75, 438-447.
- **New Article in Nature** 03/2019: Vergniory et al. "A complete catalogue of high-quality topological materials" *Nature* (2019). 566, 480-485.
- **Updated versions of TENSOR and MTENSOR** 03/2019: The programs give the general expression of tensor properties for a given point group and magnetic point group, respectively..

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Space-group symmetry

Magnetic Symmetry and Applications

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

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

complete catalogue of high-quality topological materials"
Nature (2019). 566, 480-485.

- Updated versions of **TENSOR** and **MTENSOR** 03/2019: The programs give the general expression of tensor properties for a given point group and magnetic point group, respectively.

Raman and hyper-Raman scattering

Point-group symmetry

Plane-group symmetry

Double point and space groups

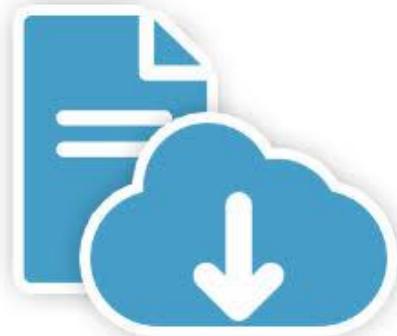
CRYSTAL-STRUCTURE TOOLS

You can access to the material of this session:

<http://www.cryst.ehu.es/resources/nancy2019/>

You need to download:

- CrystalStructureTools.txt



STRUCTURE TRANSFORMATION

Crystal structure descriptions

What do we need to describe a crystal structure?

Space Group (*IT*A number)

Lattice parameters

Number of independent atoms in the asymmetric unit

Atom type and number

The Wyckoff position

The coordinates x, y, z

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



ITA settings structure descriptions

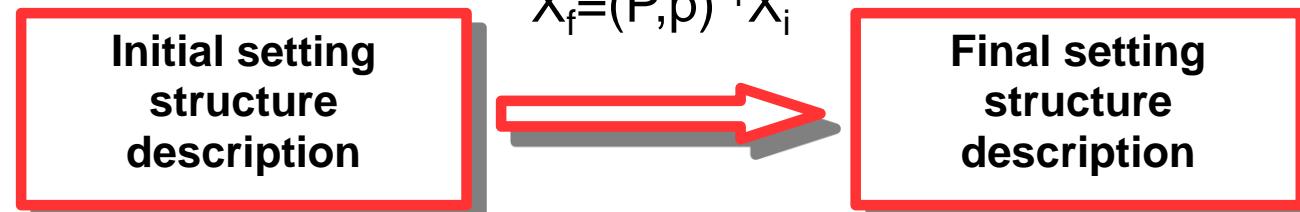
Inorganic Crystal Structure Database

CaPbO₃

		CIF	Export	Bonds	Pattern	Structure	Jmol
Title	Crystal structure and its role in electrical properties of the perovskite Ca Pb O ₃ synthesized at high pressure.						
Authors	Yamamoto, A.;Khasanova, N.R.;Izumi, F.;Wu, X.-J.;Kamiyama, T.;Torii, S.;Tajima, S.						
Reference	Chemistry of Materials (1999) 11 , 747-753 Link XRef SCOPUS SCIRUS Google						
Compound	Ca ₁ O ₃ Pb ₁ - Calcium plumbate [ABX3] [oP20] [d c2 b] []						
Cell	5.67102(4), 5.88752(4), 8.14954(6), 90., 90., 90. PBNM (62) $\text{V} = 272.1$						
Remarks	R=0.011000 : RVP NDP						
Atom (site) Oxid.		x, y, z, B, Occupancy					
Ca1	(4c)	2	0.9860(3)	0.0563(2)	0.25	0.84(3)	1
Pb1	(4b)	4	0.5	0	0	0.38(2)	1
O1	(4c)	-2	0.1200(3)	0.4452(2)	0.25	0.67(3)	1
O2	(8d)	-2	0.6907(2)	0.3051(2)	0.0613(2)	0.96(2)	1

→ Non-standard setting

$$(a, b, c)_n = (a, b, c)_s P$$



ITA number	Setting	P	P^{-1}
62	<i>P n m a</i>	a,b,c	a,b,c
62	<i>P m n b</i>	b,a,-c	b,a,-c
62	<i>P b n m</i>	c,a,b	b,c,a
62	<i>P c m n</i>	-c,b,a	c,b,-a
62	<i>P m c n</i>	b,c,a	c,a,b
62	<i>P n a m</i>	a,-c,b	a,c,-b

ITA settings structure descriptions

SETSTRU

<http://www.cryst.ehu.es/cryst/setstru.html>

CaPbO₃

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 **ITA 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]

Datei auswählen Keine ausgewählt

HINT: [The option for a given filename is preferential]

Structure

```
# Space ITA number
62
# Lattice parameters
5.67102 5.88752 8.14954 90. 90. 90.
# Number of independent atoms in the asymmetric unit
4
# [atom type] [number] [WP] [x] [y] [z]
Ca 1 4c 0.9860 0.0563 0.2500
Pb 1 4b 0.5000 0.0000 0.0000
O 1 4c 0.1200 0.4452 0.2500
O 2 8d 0.6907 0.3051 0.0613
```

CIF file

BCS format

Transform Structure

[Bilbao Crystallographic Server Main Menu]

ITA settings structure descriptions

SETSTRU

CaPbO_3

Choose the initial and final space groups symbols

The standard setting (default) of the space group 62 is *Pnma*

Initial	Final	Setting	\mathbf{P}	\mathbf{P}^{-1}
	●	<i>Pnma</i>	a,b,c	a,b,c
○	○	<i>Pmn b</i>	b,a,-c	b,a,-c
●	○	<i>Pbnm</i>	c,a,b	b,c,a
○	○	<i>Pcmn</i>	-c,b,a	c,b,-a
○	○	<i>Pmcn</i>	b,c,a	c,a,b
○	○	<i>Pnam</i>	a,-c,b	a,c,-b

Note:

1. The listed transformation matrices \mathbf{P} describe the transformation from standard to non-conventional setting:

$$(\mathbf{a}, \mathbf{b}, \mathbf{c})_n = (\mathbf{a}, \mathbf{b}, \mathbf{c})_s \mathbf{P}$$

2. The non-zero elements of the transformation matrices \mathbf{P} are listed by columns, i.e. $\mathbf{P} = -\mathbf{a}, -\mathbf{a}, -\mathbf{c}, -\mathbf{b}$ means:

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

Transform

Transformation to standard setting of space group 62

Initial structure

Initial Setting: *Pbnm* (62)

62
 5.67102 5.88752 8.14954 90. 90. 90.
 4
 Ca 1 - 0.986000 0.056300 0.250000
 Pb 1 - 0.500000 0.000000 0.000000
 O 1 - 0.120000 0.445200 0.250000
 O 2 - 0.690700 0.305100 0.061300

Final structure

Final Setting: *Pnma* (62)

62
 5.8875 8.1495 5.6710 90.00 90.00 90.00
 4
 Ca 1 4c 0.056300 0.250000 0.986000
 Pb 1 4b 0.000000 0.000000 0.500000
 O 1 4c 0.445200 0.250000 0.120000
 O 2 8d 0.305100 0.061300 0.690700

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

Transformation matrix (\mathbf{P}, p): b,c,a; 0,0,0

Matrix form:

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

ITA settings structure descriptions

SETSTRU

CaPbO3

Choose the initial and final space groups symbols

The standard setting (default) of the space group 62 is *Pnma*

Initial	Final	Setting	P	P ⁻¹
	<input checked="" type="radio"/>	<i>P n m a</i>	a,b,c	a,b,c
	<input type="radio"/>	<i>P m n b</i>	b,a,-c	b,a,-c
	<input checked="" type="radio"/>	<i>P b n m</i>	c,a,b	b,c,a
	<input type="radio"/>	<i>P c m n</i>	-c,b,a	c,b,-a
	<input type="radio"/>	<i>P m c n</i>	b,c,a	c,a,b
	<input type="radio"/>	<i>P n a m</i>	a,-c,b	a,c,-b

Note:

1. The listed transformation matrices **P** describe the transformation from standard to non-conventional setting:

$$(\mathbf{a}, \mathbf{b}, \mathbf{c})_n = (\mathbf{a}, \mathbf{b}, \mathbf{c})_s P$$

2. The non-zero elements of the transformation matrices **P** are listed by columns, i.e. **P = -a, -a-c, -b** means:

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

Atoms Data:

AT.	WP	SS	Representative	Atomic orbit
Ca1	4c (x,1/4,z)	.m.	(0.056300, 0.250000, 0.986000) (0.443700, 0.750000, 0.486000) (0.943700, 0.750000, 0.014000) (0.556300, 0.250000, 0.514000)	(0.056300, 0.250000, 0.986000) (0.443700, 0.750000, 0.486000) (0.943700, 0.750000, 0.014000) (0.556300, 0.250000, 0.514000)
Pb1	4b (0,0,1/2)	-1	(0.000000, 0.000000, 0.500000) (0.500000, 0.000000, 0.000000) (0.000000, 0.500000, 0.500000) (0.500000, 0.500000, 0.000000)	(0.000000, 0.000000, 0.500000) (0.500000, 0.000000, 0.000000) (0.000000, 0.500000, 0.500000) (0.500000, 0.500000, 0.000000)
O1	4c (x,1/4,z)	.m.	(0.445200, 0.250000, 0.120000) (0.054800, 0.750000, 0.620000) (0.554800, 0.750000, 0.880000) (0.945200, 0.250000, 0.380000)	(0.445200, 0.250000, 0.120000) (0.054800, 0.750000, 0.620000) (0.554800, 0.750000, 0.880000) (0.945200, 0.250000, 0.380000)
O2	8d (x,y,z)	1	(0.305100, 0.061300, 0.690700) (0.194900, 0.938700, 0.190700) (0.694900, 0.561300, 0.309300) (0.805100, 0.438700, 0.809300) (0.694900, 0.938700, 0.309300) (0.805100, 0.061300, 0.809300) (0.305100, 0.438700, 0.690700) (0.194900, 0.561300, 0.190700)	(0.305100, 0.061300, 0.690700) (0.194900, 0.938700, 0.190700) (0.694900, 0.561300, 0.309300) (0.805100, 0.438700, 0.809300) (0.694900, 0.938700, 0.309300) (0.805100, 0.061300, 0.809300) (0.305100, 0.438700, 0.690700) (0.194900, 0.561300, 0.190700)

Transform

Exercise 3.1



Compare the two structure descriptions and check if they belong to the same structure type

ZrSiO₄

Print 2 entries selected.

CC=Collection Code: [AB2X4]=ANX Form: [cF56]=Pearson: [e d a]=Wyckoff Symbol: [Al2MgO4]=Structure Type:

Click the ANX, Pearson or Wyckoff Symbol to find structures with that symbol.

CC=45520 Details Bonds Pattern Structure Jmol

Title Redetermination of the oxygen parameters in zircon (Zr Si O₄).

Authors Krstanovic, I.R.

Reference Acta Crystallographica (1958) 11, 896-897
[Link](#) [XRef](#) [SCOPUS](#) [SCIRUS](#) [Google](#)

Compound Zr (Si O₄) - [Zircon] Zirconium silicate [**ABX4**] [**tI24**] [**h b a**] [**ZrSiO₄**]

Cell 6.6164(5), 6.6164, 6.0150(5), 90., 90., 90.
I41/AMDZ (141) V=263.32

Remarks R=0.070000 : PDC =01-073-6646 : PDF =6-266 : TYP =ZrSiO₄ :
 XDS MIN =Zircon :
 At least one temperature factor missing in the paper.
 hk0- and 0kl-data, crystals not metamict

Atom (site) Oxid. **x, y, z, B, Occupancy**

Zr1	(4a)	4	0 0.75	0.125	0 1
Si1	(4b)	4	0 0.75	0.625	0 1
O1	(16h)	-2	0 0.067(3)	0.198(3)	0 1

CC=31101 Details Bonds Pattern Structure Jmol

Title Die Kristallstruktur von Zirkon und die Kriterien fuer spezielle Lagen in tetragonalen Raumgruppen..

Authors Wyckoff, R.W.G.;Hendricks, S.B.

Reference Zeitschrift fuer Kristallographie, Kristallgeometrie, Kristallphysik, Kristallchemie (1927) 66, 73-102
[Link](#) [XRef](#) [SCOPUS](#) [SCIRUS](#) [Google](#)
 Also: Philosophical Magazine, Serie (1926) 1, 1151-1151

Compound Zr (Si O₄) - [Zircon] Zirconium silicate [**ABX4**] [**tI24**] [**h b a**] [**ZrSiO₄**]

Cell 6.61, 6.61, 5.98, 90., 90., 90.
I41/AMDS (141) V=261.28

Remarks COR MIN =Zircon : PDF =6-266 : TYP =ZrSiO₄ : XDS At least one temperature factor missing in the paper.
 No R value given in the paper.
 Revised data of 31084

Atom (site) Oxid. **x, y, z, B, Occupancy**

Zr1	(4a)	4	0 0	0	0 1
Si1	(4b)	4	0 0	0.5	0 1
O1	(16h)	-2	0 0.2(1)	0.34(2)	0 1

Origin choice 2

Origin choice 1



Exercise 3.1

Structure 1

Space group $I4_1/AMD$ (No. 141)
 $a=6.600 \text{ \AA}$ $c=5.88 \text{ \AA}$ Origin choice 1 at $-4m2$

Structure 2

Space group $I4_1/AMD$ (No. 141)
 $a=6.616 \text{ \AA}$ $c=6.015 \text{ \AA}$ Origin choice 2 at $2/m$

Compare the two structure descriptions and check if they belong to the same structure type

Use the tool of Bilbao Crystallographic Server: **SETSTRU**

Hint: In order to compare the different data, the parameters of Structure 1 are to be transformed to 'origin at center 2/m', i.e. ORIGIN CHOICE 2

Structure transformation

TRANSTRU

<http://www.cryst.ehu.es/cryst/transtru.html>

Transform Structure

Pb₃(VO₄)₂

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

[in CIF format]

Datei auswählen Keine ausgewählt

HINT: [The option for a given filename is preferential]

CIF file

```
166  
5.6739 5.6739 20.3412 90. 90. 120.  
6  
Pb 1 3a 0.0000 0.0000 0.0000  
Pb 2 6c 0.0000 0.0000 0.2073  
PV 1 6c 0.0000 0.0000 0.3924  
O 1 6c 0.0000 0.0000 0.3247  
O 2 18h 0.8433 0.1567 0.4299
```

High Symmetry Structure

BCS format

Transform structure to a subgroup basis

Transform structure with an arbitrary matrix

subgroup basis

Show

arbitrary transformation

Structure transformation

Description
R-3m (No. 166)

Pb₃(VO₄)₂

(P,p)

166	5.6748	5.6748	20.3784	90	90	120
5	Pb	1	3a	0.000000	0.000000	0.000000
	Pb	2	6c	0.000000	0.000000	0.207100
	PV	3	6c	0.000000	0.000000	0.388400
	O	4	6c	0.000000	0.000000	0.324000
	O	5	18h	0.842400	0.157600	0.430100

Structure

Low symmetry Space Group ITA number

14

Transformation Matrix:

In matrix form:

Description
P2₁/c (No. 14)

Validity (P,p)

WP splitting

Linear part

2/3	0	-2
1/3	1	1
1/2	0	0

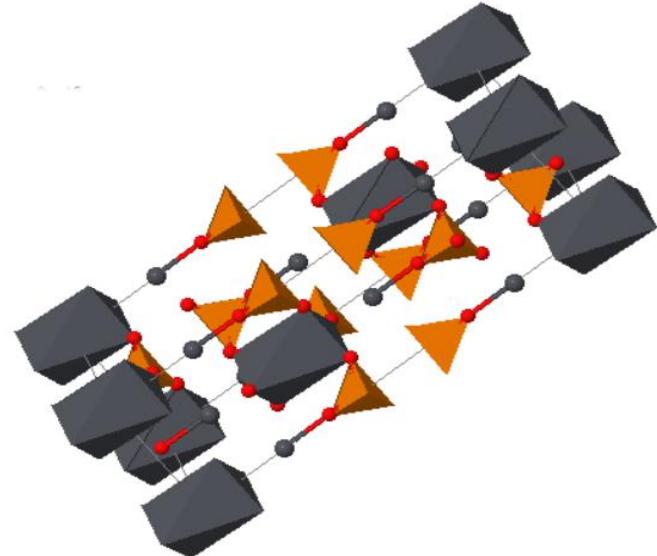
Origin Shift

0
0
0

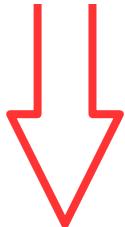
Structure transformation

High symmetry structure

```
166  
5.6739 5.6739 20.3412 90. 90. 120.  
5  
Pb     1      3a      0.000000      0.000000      0.000000  
Pb     2      6c      0.000000      0.000000      0.207300  
PV    1      6c      0.000000      0.000000      0.392400  
O     1      6c      0.000000      0.000000      0.324700  
O     2      18h     0.843300      0.156700      0.429900
```

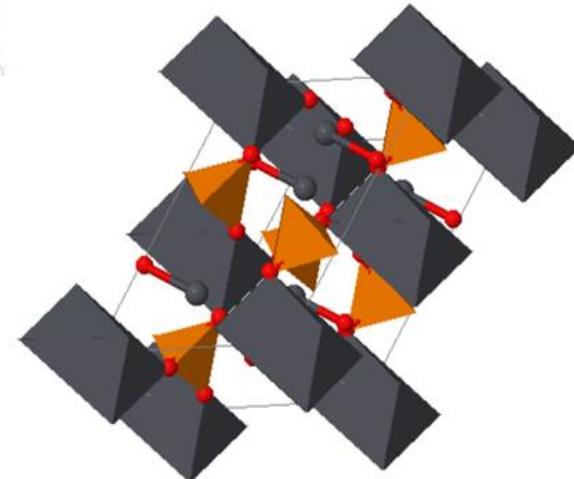


$$(P,p) = 2/3\mathbf{a} + 1/3\mathbf{b} + 1/3\mathbf{c}, \mathbf{b}, -2\mathbf{a}-\mathbf{b}$$



Low symmetry structure

```
014  
7.530264 5.673900 9.827483 90.000000 115.786674 90.000000  
7  
Pb     1      2a      0.000000      0.000000      0.000000  
Pb     2      4e      0.621900      0.000000      0.207300  
PV    1      4e      0.177200      0.000000      0.392400  
O     1      4e      0.974100      0.000000      0.324700  
O     2      4e      0.289700      0.735050      0.008250  
O    2_2     4e      0.289700      0.500000      0.773200  
O    2_3     4e      0.710300      0.764950      0.491750
```



Exercise 3.1 (cont.)



Apply the program **TRANSTRU** in order to check if the two structure descriptions belong to the same structure type

Structure 1

Space group $I4_1/amd$ (No. 141)
 $a=6.600 \text{ \AA}$ $c=5.88 \text{ \AA}$ Origin choice 1 at $-4m2$

Structure 2

Space group $I4_1/amd$ (No. 141)
 $a=6.616 \text{ \AA}$ $c=6.015 \text{ \AA}$ Origin choice 2 at $2/m$

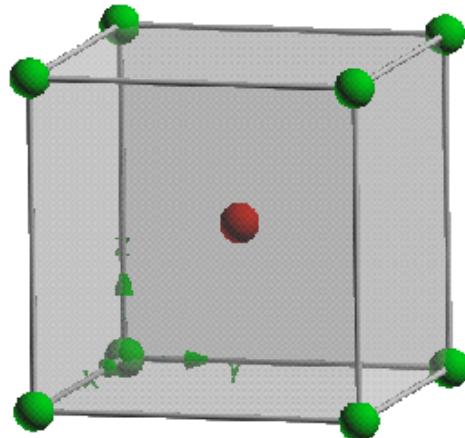
Coordinate transformation

Origin choice 2 $\xrightarrow{\hspace{1cm}}$ Origin choice 1

$$\mathbf{p}=0, 1/4, -1/8$$

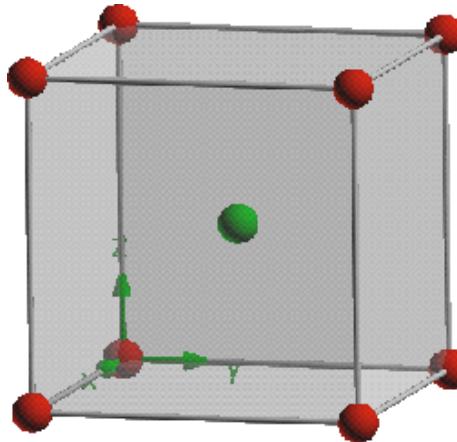
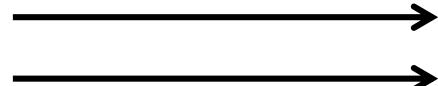
EQUIVALENT STRUCTURE DESCRIPTIONS

Equivalent structures descriptions



CsCl
Pm-3m (No. 221)

1a (0,0,0)
1b (1/2,1/2,1/2)



1b (1/2,1/2,1/2)
1a (0,0,0)

How to find all possible equivalent descriptions of a crystal structure?

$$\text{Number of equivalent descriptions} = |N(G)|/|G|$$

Index of the group in its Euclidean normalizer

Equivalent structures descriptions

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

Equivalent Descriptions of Crystal Structures

CsCl

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.

Structure Data

[in CIF format]

Examinar...

No se ha seleccionado ningún archivo.

HINT: [The option for a given filename is preferential]

Structure

```
#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
```

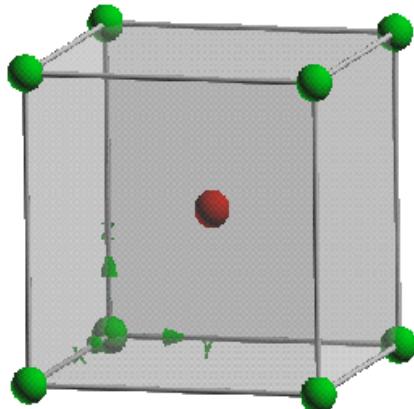
CIF file

BCS format

Only the default choice for
the conventional setting of

Show

Equivalent structures descriptions



- Equivalent description 1 (original input structure)

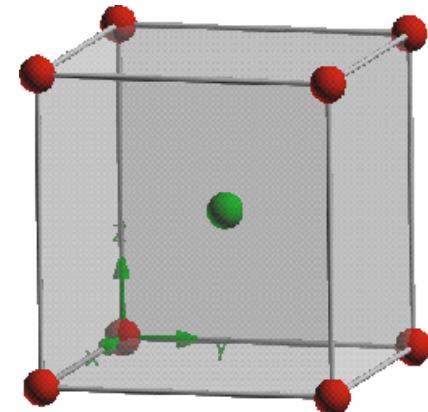
Normalizer coset representative: x,y,z

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)



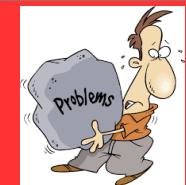
Exercise 3.2(b)

How many equivalent sets of coordinates can be used to describe the structure $\text{P}(\text{C}_6\text{C}_5)_4[\text{MoNCl}_4]$? What are their coordinates?

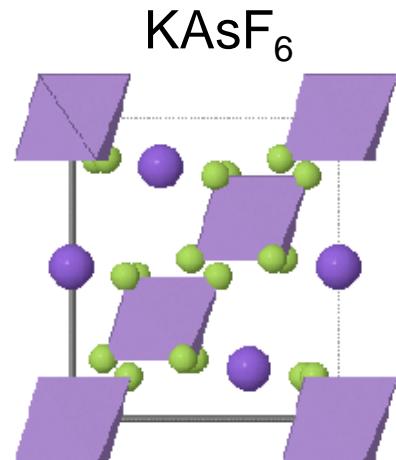
Atom	Wyckoff position	Coordinate	triplets	
			x	y
P	2b	0.25	0.75	0
Mo	2c	0.25	0.25	0.121
N	2c	0.25	0.25	-0.093
C1	8g	0.362	0.760	0.141
C2	8g	0.437	0.836	0.117
Cl	8g	0.400	0.347	0.191



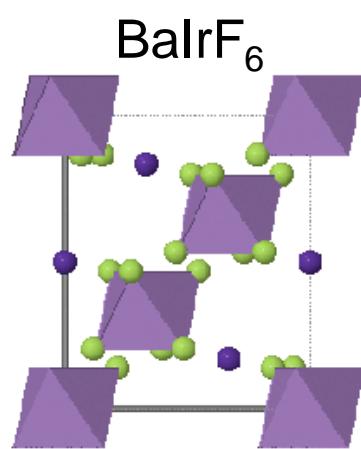
Exercise 3.3



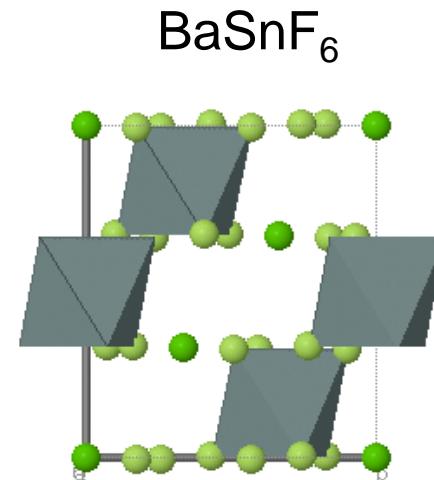
Do the following three structures belong to the same structure type?
Try to find analogous coordinate descriptions for all three crystal structures



148
7.3480 7.3480 7.2740 90.00 90.00 120.00
3
K 1 3b 0.333333 0.666666 0.166666
As 1 3a 0 0 0
F 1 18f 0.1292 0.2165 0.1381



148
7.3965 7.3965 7.2826 90.00 90.00 120.00
3
Ba 1 3b 0.333333 0.666666 0.166666
Ir 1 3a 0 0 0
F 1 18f 0.0729 0.2325 0.1640



148
7.4279 7.4279 7.4180 90.00 90.00 120.00
3
Ba 1 3a 0 0 0
Sn 1 3b 0 0 0.5
F 1 18f 0.2586 0.8262 0.0047

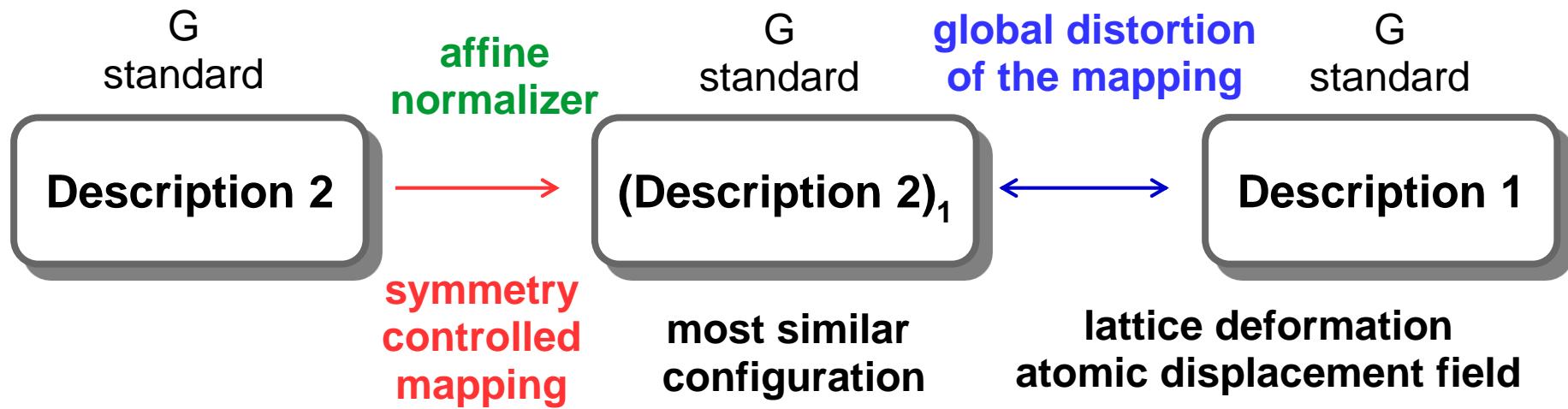
R-3 (No. 148) → 4 equivalent descriptions: x,y,z; x,y,z+1/2; y,x,-z; y,x,-z+1/2

COMPARISON OF STRUCTURES

Comparison of structures

The program measures **the similarity between two structures** with the same or different compositions:

- same space-group (or space groups that form an enantiomorph pair)
- same sequence of the occupied Wyckoff positions
- the same total number of atoms in the unit cells



Comparison of structures

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

degree of lattice distortion

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

η_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

Comparison of structures

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

Comparison of structures

COMPSTRU

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

Structure Data
[in CIF format]

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

CIF file

Pb₃(PO₄)₂

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
```

BCS format

Structure Data
[in CIF format]

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

CIF file

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
```

BCS format

Enter the maximum distance allowed between the paired atoms: Å

Enter the allowed tolerance (a b c α β γ): .5 .5 .5 5 5 5

Standard setting

Tolerance

Comparison of 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

```



WP	Atom	Atomic Displacements				
		u_x	u_y	u_z	$ 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	$d_{\max.} (\text{\AA})$	$d_{\text{av.}} (\text{\AA})$	Δ
0.0116	0.3386	0.1755	0.066

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

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

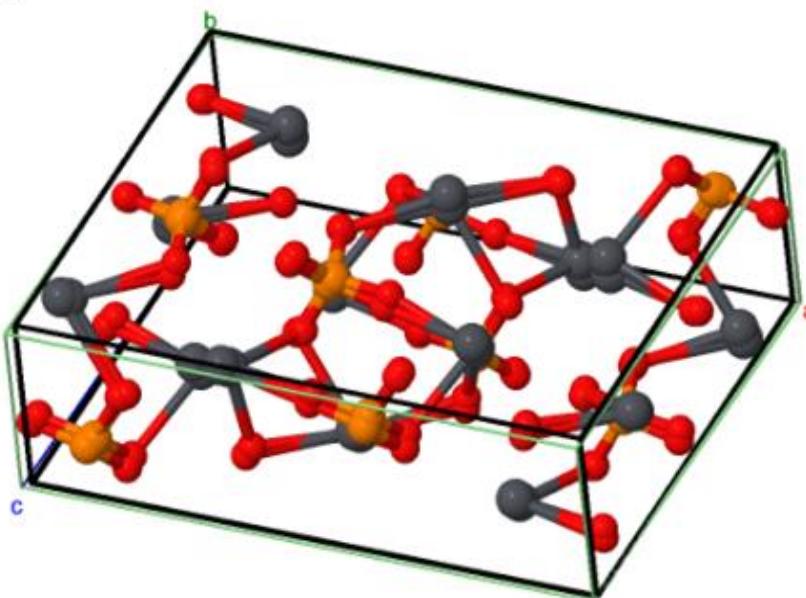
```

Comparison of structures

Visualization of the comparison

Pb₃(PO₄)₂

-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)

Compare Structures

Compare Lattices

Atomic Displacements

Structure 1: opaque
 ball&stick stick cross

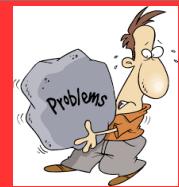
Structure 2: opaque
 ball&stick stick cross

Show Distances
cutoff: 0.5

Save PNG+Jmol

NEW

Exercise 3.4



In ICSD can be found several structure data sets of $\epsilon\text{-Fe}_2\text{O}_3$, all of them of symmetry Pna2₁(No.33). Compare the following two descriptions and check if they belong to the same structure type.

ICSD for WWW

Details of the selected entries

2 entries selected.

CC=Collection Code: [AB2X4]=ANX Form: [cF56]=Pearson: [e d a]=Wyckoff Symbol: [Al2MgO4]=Structure Type:
*****Click the ANX, Pearson or Wyckoff Symbol to find structures with that symbol***,**

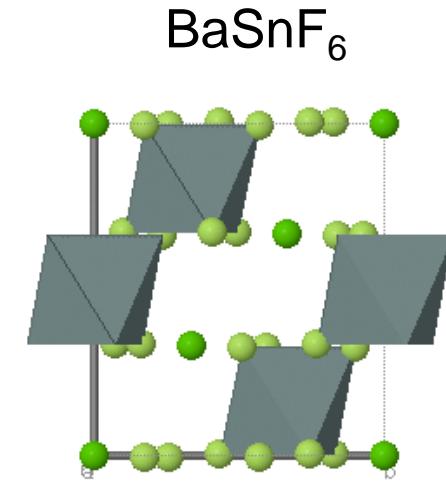
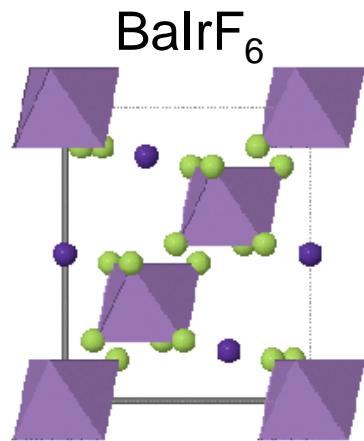
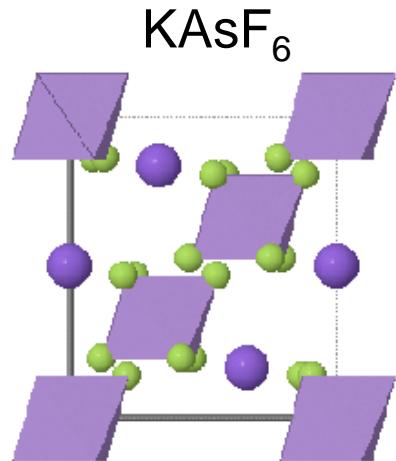
CC=173024	Details	Bonds	Pattern	Structure	Jmol
Title	High- and low-temperature crystal and magnetic structure of epsilon-Fe ₂ O ₃ and their correlation to its magnetic properties.				
Authors	Gich, M.;Frontera, C.;Roig, A.;Taboada, E.;Molins, E.;Rechenberg, H.R.;Ardisson, J.D.;Macedo, W.A.A.;Ritter, C.;Hardy, V.;Sort, J.;Skumryev, V.;Nogues, J.				
Reference	Chemistry of Materials (2007) 18 , 3889-3897 Link XRef SCOPUS SCIRUS Google				
Compound	Fe ₂ O ₃ - Iron(III) oxide - epsilon [A2X3] [oP40] [a10] [AlFeO3]				
Cell	5.0885(5), 8.7802(14), 9.4709(13), 90., 90., 90. PNA21 (33) V=423.14				
Remarks	R=0.013300 : NPD RVP SNP TEM =200 : TYP =AlFeO3 : XDP MAG At least one temperature factor missing in the paper.				
Atom (site) Oxid.	x, y, z, B, Occupancy				
O1 (4a)	-2	0.978(2)	0.3282(15)	0.4314(11)	0 1
O2 (4a)	-2	0.515(2)	0.4907(17)	0.4187(16)	0 1
O3 (4a)	-2	0.650(3)	0.9979(13)	0.1883(9)	0 1
O4 (4a)	-2	0.160(3)	0.1637(15)	0.1956(7)	0 1
O5 (4a)	-2	0.841(3)	0.1680(15)	0.6669(7)	0 1
O6 (4a)	-2	0.527(2)	0.1637(19)	0.9362(9)	0 1
Fe1 (4a)	3	0.1928(11)	0.1505(6)	0.5807(3)	0 1
Fe2 (4a)	3	0.6826(6)	0.0291(3)	0.7897(5)	0 1
Fe3 (4a)	3	0.1858(10)	0.1519(6)	0	0 1
Fe4 (4a)	3	0.8104(7)	0.1580(4)	0.3071(3)	0 1

CC=415250	Details	Bonds	Pattern	Structure	Jmol	
Title	Synthesis and structural analysis of epsilon-(Fe ₂ O ₃).					
Authors	Kelm, K.;Mader, W.					
Reference	Zeitschrift fuer Anorganische und Allgemeine Chemie (2005) 631 , 2383-2389 Link XRef SCOPUS SCIRUS Google					
Compound	Fe ₂ O ₃ - Diliron(III) oxide - epsilon [A2X3] [oP40] [a10] [AlFeO3]					
Cell	5.0715(2), 8.7359(4), 9.4178(4), 90, 90, 90 PNA21 (33) V=417.24					
Remarks	R=0.039000 : TYP =AlFeO3 : XDP RVP					
Atom (site) Oxid.	x, y, z, B, Occupancy					
Fe1 (4a)	3	0.6768(9)	0.8427(5)	0.0000000	0.050(2)	1.0000000
Fe2 (4a)	3	0.204(1)	0.3509(8)	0.7726(9)	0.063(3)	1.0000000
Fe3 (4a)	3	0.807(1)	0.6605(8)	0.693(1)	0.059(2)	1.0000000
Fe4 (4a)	3	0.6852(9)	0.4634(5)	0.983(2)	0.046(1)	1.0000000
O1 (4a)	-2	0.337(2)	0.853(2)	0.887(1)	0.0063326	1.0000000
O2 (4a)	-2	0.019(3)	0.474(2)	0.610(2)	0.0063326	1.0000000
O3 (4a)	-2	0.453(3)	0.677(2)	0.651(2)	0.0063326	1.0000000
O4 (4a)	-2	0.527(3)	0.669(2)	0.100(1)	0.0063326	1.0000000
O5 (4a)	-2	0.868(3)	0.334(2)	0.863(1)	0.0063326	1.0000000
O6 (4a)	-2	0.335(3)	0.513(1)	0.891(1)	0.0063326	1.0000000

Exercise 3.3 (cont)



Do these compounds belong to the **same structure type** ?



148
7.3480 7.3480 7.2740 90.00 90.00 120.00
3
K 1 3b 0.333333 0.666666 0.166666
As 1 3a 0 0 0
F 1 18f 0.1292 0.2165 0.1381

148
7.3965 7.3965 7.2826 90.00 90.00 120.00
3
Ba 1 3b 0.333333 0.666666 0.166666
Ir 1 3a 0 0 0
F 1 18f 0.0729 0.2325 0.1640

148
7.4279 7.4279 7.4180 90.00 90.00 120.00
3
Sn 1 3b 0 0 0.5
Ba 1 3a 0 0 0
F 1 18f 0.2586 0.8262 0.0047

Comparison of structures

STUDY OF THE FAMILY ABX₃

R-3 (148); WP sequence: fc2; Pearson: hR10

	AA1	AA2	AA3	AA4	AA5	AA6	AA7	AA8	AA9	AB1	AB2	AB3	AB4	AB5	AB6	AB7	AC1	AC2	AC3	AD1	AD2	AE1	AE2	B1	C1		
AA1	0	0,0	0,1	0,1	0,1	0,1	0,1	0,2	0,1	0,1	0,2	0,0	0,2	0,3	0,1	0,2	0,2	0,2	0,2	0,2	1,3	1,2	0,5	0,7			
AA2	0,0	0	0,1	0,1	0,1	0,1	0,1	0,2	0,1	0,1	0,2	0,0	0,2	0,3	0,1	0,2	0,2	0,2	0,2	0,2	1,3	1,2	0,5	0,7			
AA3	0,1	0,1	0	0,2	0,2	0,2	0,1	0,1	0,3	0,0	0,1	0,1	0,1	0,1	0,1	0,1	0,1	0,1	0,1	0,1	0,1	1,4	1,3	0,4	0,7		
AA4	0,1	0,1	0,2	0	0,1	0,0	0,1	0,1	0,2	0,1	0,2	0,2	0,1	0,3	0,2	0,2	0,2	0,2	0,3	0,2	0,3	1,3	1,2	0,4	0,7		
AA5	0,1	0,1	0,2	0,1	0	0,1	0,1	0,2	0,2	0,1	0,1	0,2	0,1	0,2	0,3	0,1	0,2	0,2	0,2	0,2	0,2	1,2	1,2	0,5	0,7		
AA6	0,1	0,1	0,2	0,0	0,1	0	0,1	0,2	0,2	0,1	0,2	0,2	0,1	0,3	0,2	0,2	0,2	0,2	0,3	0,2	0,3	1,2	1,2	0,4	0,8		
AA7	0,1	0,1	0,1	0,1	0,1	0	0	0,1	0,3	0,1	0,2	0,2	0,1	0,2	0,2	0,2	0,2	0,2	0,3	0,2	0,3	1,3	1,3	0,4	0,7		
AA8	0,1	0,1	0,1	0,1	0,2	0,2	0,1	0	0,3	0,1	0,1	0,1	0,1	0,2	0,2	0,1	0,1	0,1	0,2	0,2	0,2	1,4	1,3	0,4	0,7		
AA9	0,2	0,2	0,3	0,2	0,2	0,2	0,3	0,3	0	0,3	0,3	0,4	0,2	0,4	0,3	0,3	0,3	0,4	0,3	0,4	0,4	1,5	1,4	0,6	0,7		
AB1	0,1	0,1	0,0	0,1	0,1	0,1	0,1	0,3	0		0,1	0,1	0,1	0,1	0,1	0,1	0,1	0,2	0,1	0,2	0,2	2,0	1,9	0,5	0,7		
AB2	0,1	0,1	0,1	0,2	0,1	0,2	0,2	0,1	0,3	0,1		0	0,1	0,1	0,1	0,2	0,0	0,1	0,1	0,2	0,2	0,2	1,3	1,3	0,5	0,7	
AB3	0,2	0,2	0,1	0,2	0,2	0,2	0,2	0,1	0,4	0,1		0,1	0	0,2	0,1	0,2	0,2	0,0	0,1	0,1	0,1	0,1	1,4	1,3	0,5	0,7	
AB4	0,0	0,0	0,1	0,1	0,1	0,1	0,1	0,2	0,2	0,1		0,1	0,1	0	0,2	0,2	0,0	0,1	0,2	0,2	0,2	0,2	1,3	1,2	0,5	0,7	
AB5	0,2	0,2	0,1	0,3	0,2	0,3	0,2	0,2	0,4	0,1		0,1	0,1	0,2	0	0,2	0,1	0,1	0,1	0,1	0,1	1,5	1,5	0,4	0,7		
AB6	0,3	0,3	0,1	0,2	0,3	0,2	0,2	0,2	0,3	0,1		0,2	0,2	0,2	0,2	0	0,3	0,2	0,1	0,2	0,1	0,2	1,9	1,8	0,5	0,7	
AB7	0,1	0,1	0,1	0,2	0,1	0,2	0,2	0,1	0,3	0,1		0,0	0,2	0,0	0,1	0,3	0	0,2	0,2	0,2	0,2	0,2	1,3	1,2	0,4	0,7	
AC1	0,2	0,2	0,1	0,2	0,2	0,2	0,2	0,1	0,3	0,1		0,1	0,0	0,1	0,1	0,2	0,2	0		0,1	0,1	0,1	0,1	1,5	1,9	0,4	0,7
AC2	0,2	0,2	0,1	0,2	0,2	0,2	0,2	0,1	0,3	0,1		0,1	0,1	0,2	0,1	0,1	0,2	0,1		0	0,1	0,1	0,1	1,4	1,3	0,3	0,7
AC3	0,2	0,2	0,1	0,3	0,2	0,3	0,3	0,2	0,4	0,2		0,2	0,2	0,1	0,2	0,1	0,2	0,1		0,1	0,1	0,0	0,1	1,6	1,5	0,4	0,7
AD1	0,2	0,2	0,1	0,2	0,2	0,2	0,2	0,2	0,3	0,1		0,2	0,1	0,2	0,1	0,1	0,2	0,1		0,1	0,1	0,1	0,1	2,2	2,0	0,4	0,7
AD2	0,2	0,2	0,1	0,3	0,2	0,3	0,3	0,2	0,4	0,2		0,2	0,2	0,1	0,2	0,1	0,2	0,1		0,1	0,1	0,1	0	2,2	2,0	0,4	0,7
AE1	1,3	1,3	1,4	1,3	1,2	1,2	1,3	1,4	1,5	2,0		1,3	1,4	1,3	1,5	1,9	1,3	1,5	1,4	1,6	2,2	2,2	0	0,1	1,9	1,3	
AE2	1,2	1,2	1,3	1,2	1,2	1,2	1,3	1,3	1,4	1,9		1,3	1,3	1,2	1,5	1,8	1,2	1,9	1,3	1,5	2,0	2,0	0,1	0	1,8	1,2	
B1	0,5	0,5	0,4	0,4	0,5	0,4	0,4	0,6	0,5	0,5		0,5	0,5	0,4	0,5	0,4	0,4	0,3	0,4	0,4	0,4	0,4	1,9	1,8	0	0,5	
C1	0,7	0,7	0,7	0,7	0,7	0,8	0,7	0,7	0,7	0,7		0,7	0,7	0,7	0,7	0,7	0,7	0,7	0,7	0,7	0,7	0,7	1,3	1,2	0,5	0	



Bergerhoff
(structure descriptor)



Bilbao Server
(global distortion)