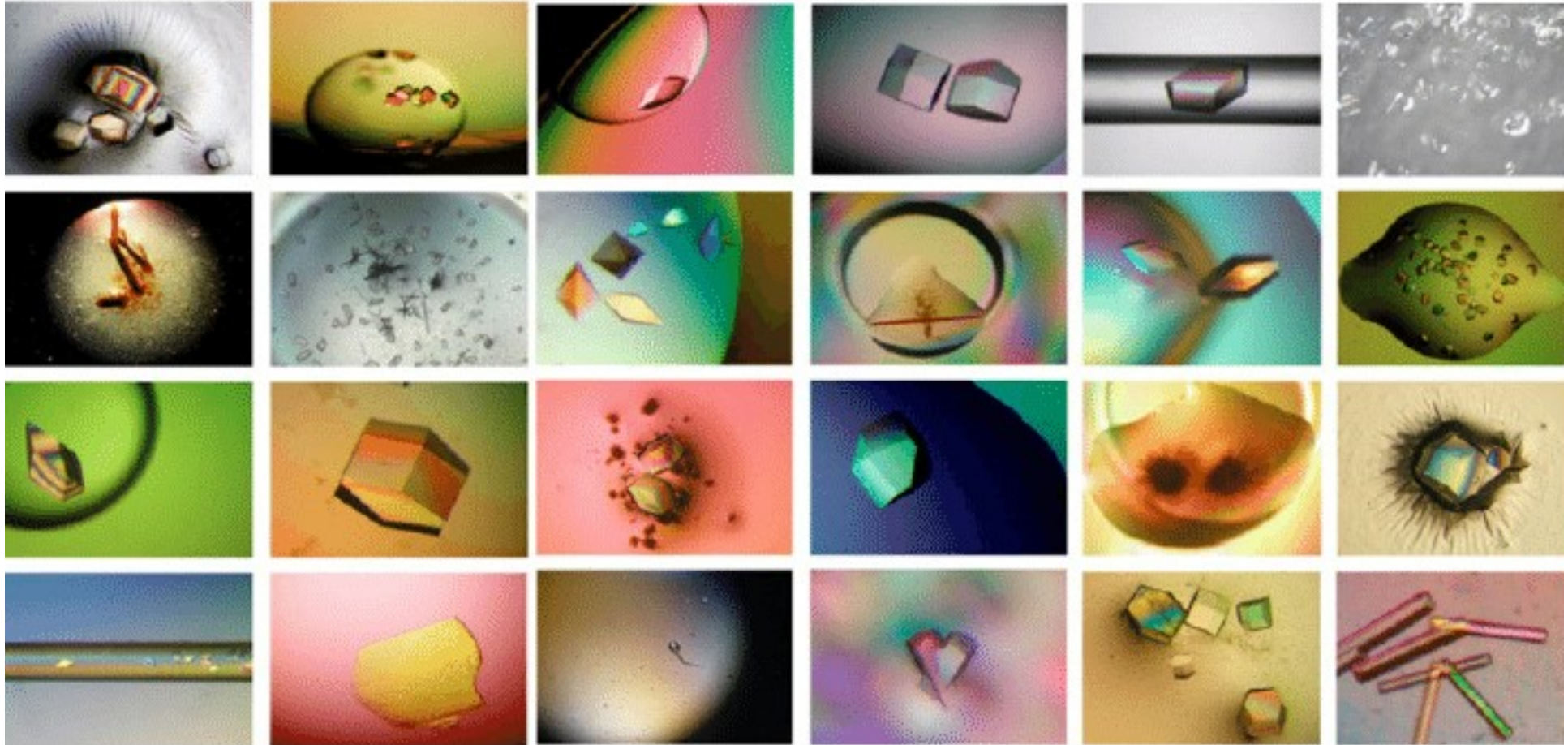


# Crystal geometry



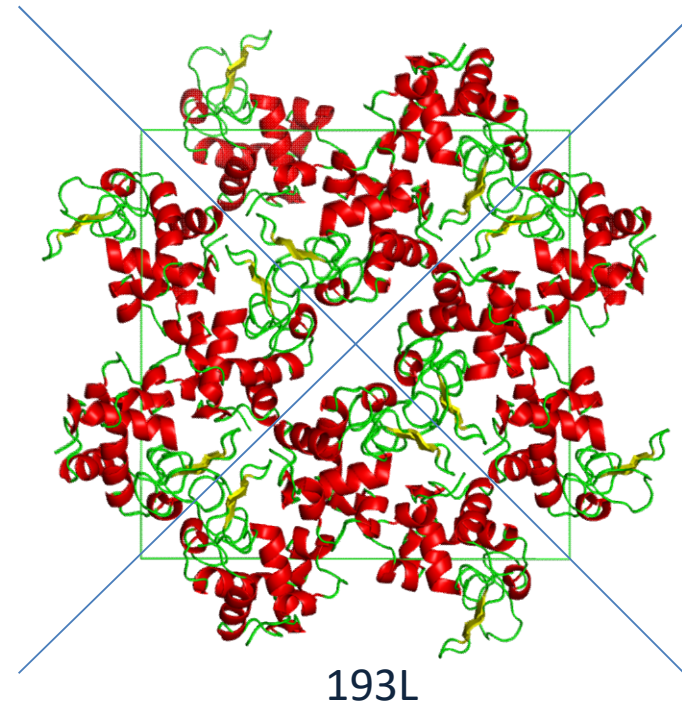
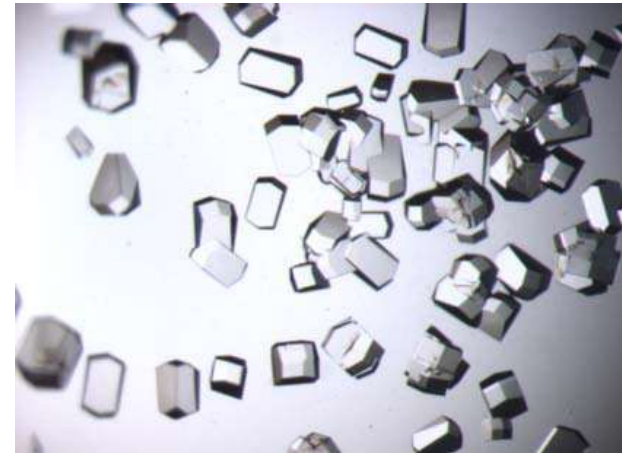
# Sharp edges and plane faces

Regular crystal habit depends on the internal long-range order of the crystal structure.

Diffraction properties of crystals depend on their internal order: X-ray interaction with the ordered molecules forming the crystal.

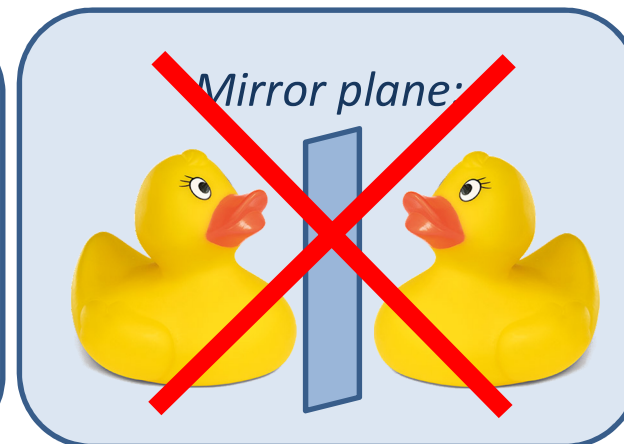
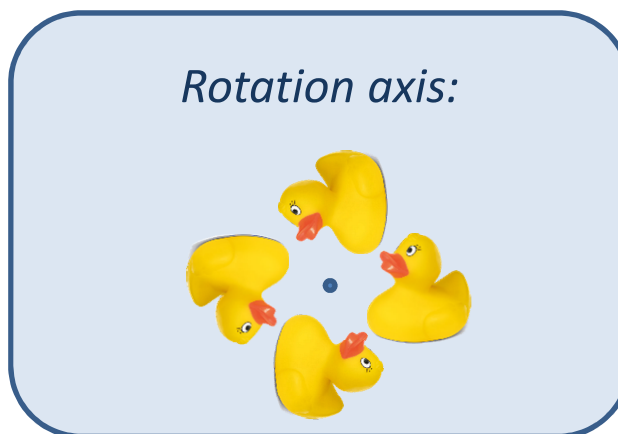
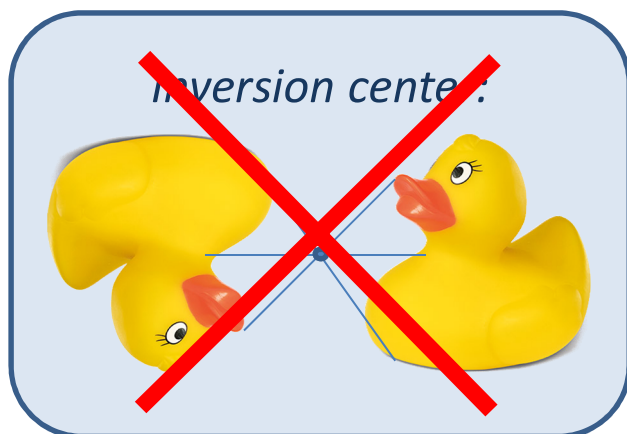
Combination of the requirements for:  
**long-range order**  
and  
**symmetry elements** in the crystal  
=  
230 **space groups** for crystals,  
But only **65** allowed for protein crystals

Understanding crystal order is important to maximize the information that can be extracted from crystallographic data.



# Point group symmetry elements

Point group symmetry elements are invariant points of transformations able to relate a point  $(x,y,z)$  to another point  $(x',y',z')$ .



During each of these transformations, a generic point of coordinates  $(x,y,z)$  is transformed into a symmetric point, with coordinates  $(x',y',z')$ , by a matrix operator:

*Inversion center,  $i$ :*

$$\mathbf{x}' = i \mathbf{x}$$

$$i = \begin{bmatrix} -1 & 0 & 0 \\ 0 & -1 & 0 \\ 0 & 0 & -1 \end{bmatrix}$$

*Rotation axis, e.g. 4 along z:*

$$\mathbf{x}' = A \mathbf{x}$$

$$A = \begin{bmatrix} 0 & -1 & 0 \\ 1 & 0 & 0 \\ 0 & 0 & 1 \end{bmatrix}$$

*Mirror plane, e.g. in yz:*

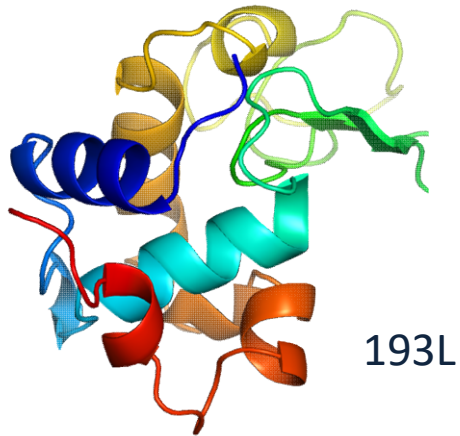
$$\mathbf{x}' = m \mathbf{x}$$

$$m = \begin{bmatrix} -1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{bmatrix}$$

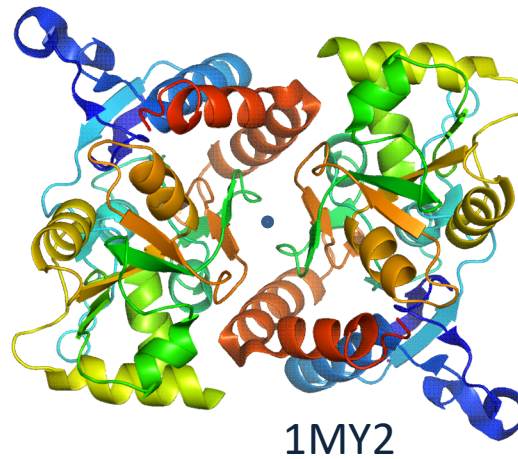
# Rotation axes

A rotation operation  $n$  (or of order  $n$ ) describes a rotation of each point of  $360^\circ/n$  around the rotation axis:

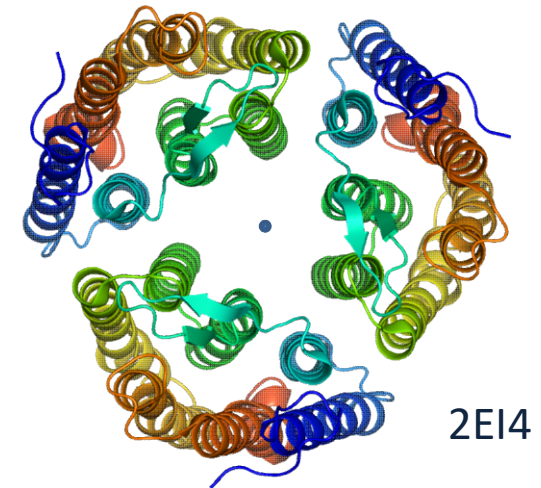
**1** Rotation of  $360^\circ$   
(identity!)



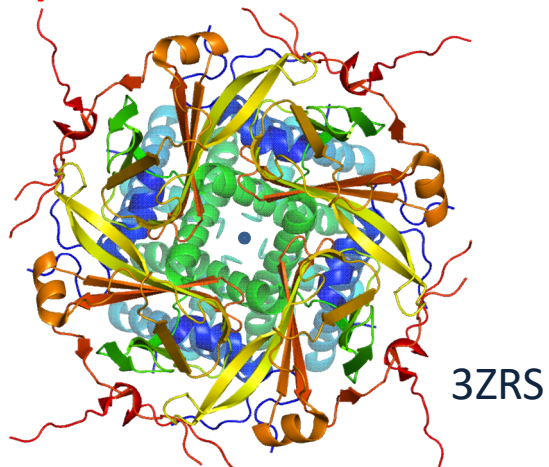
**2** Rotation of  $180^\circ$



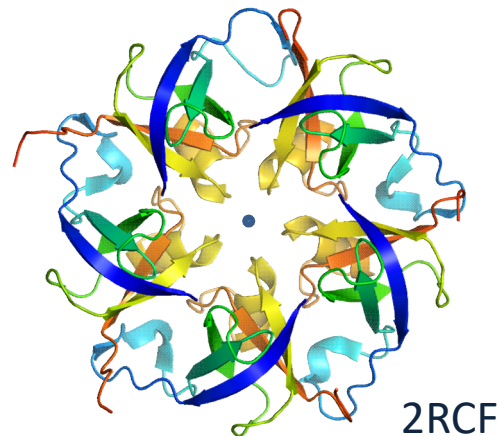
**3** Rotation of  $120^\circ$



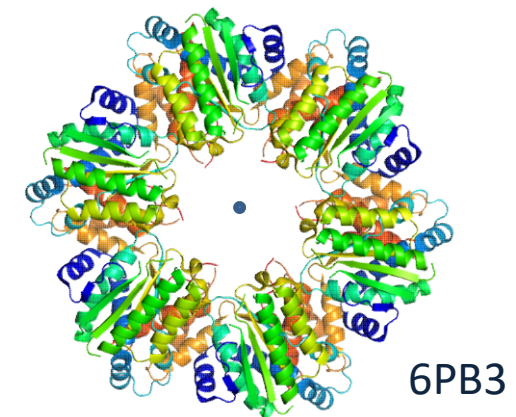
**4** Rotation of  $90^\circ$



**5** Rotation of  $72^\circ$

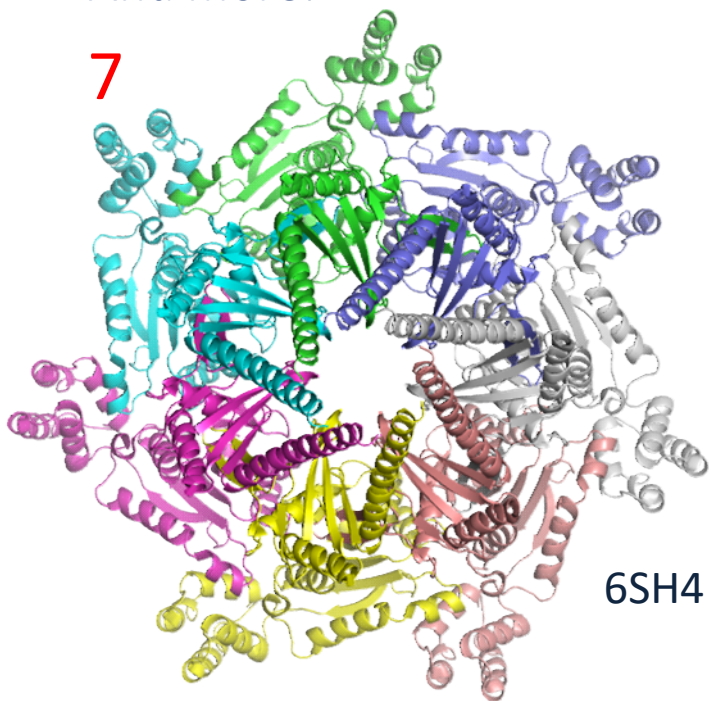


**6** Rotation of  $60^\circ$

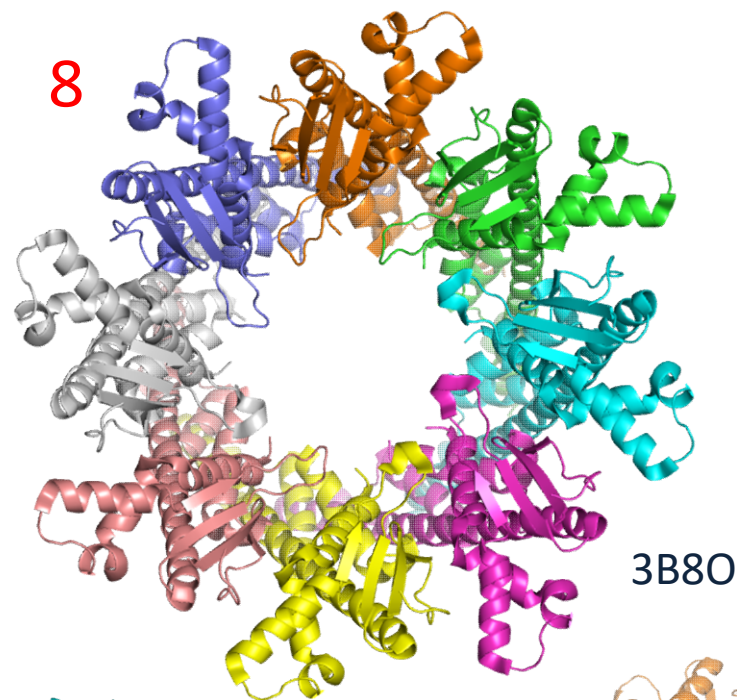


And more:

7

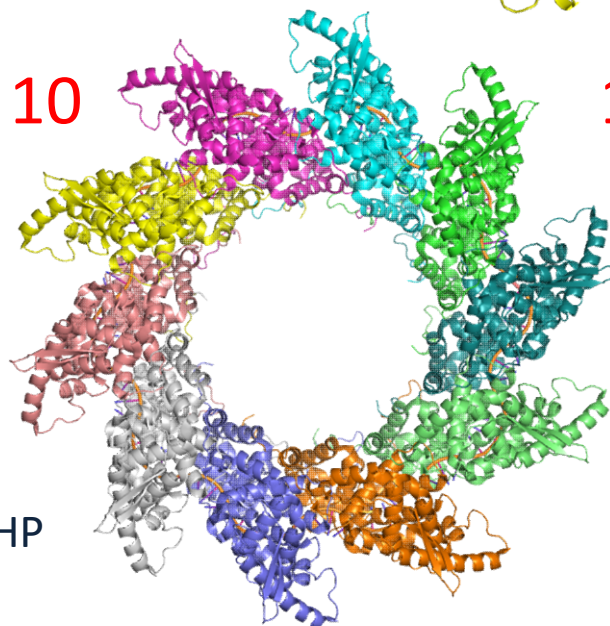


8



When symmetry elements are present, they involve more than one protein chain: proteins are **ASYMMETRIC!**

10



12



# Translational periodicity: lattice and unit cell

Crystals are formed by repeated units along the three directions, but to simplify the problem we can start to describe lattice in the 2D case

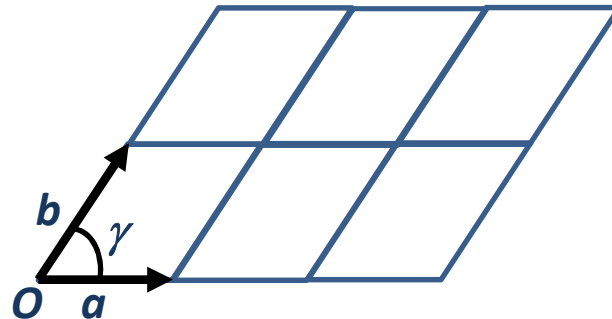
*How to cover a floor with tiles??*



Due to the requirement for **translational periodicity**, lattices can have only the following rotational symmetries:

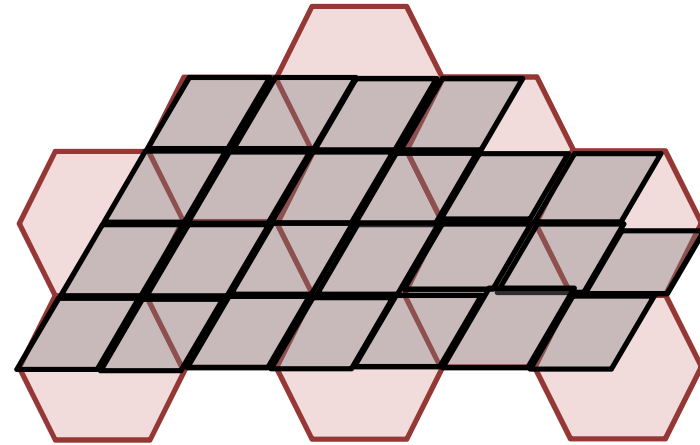
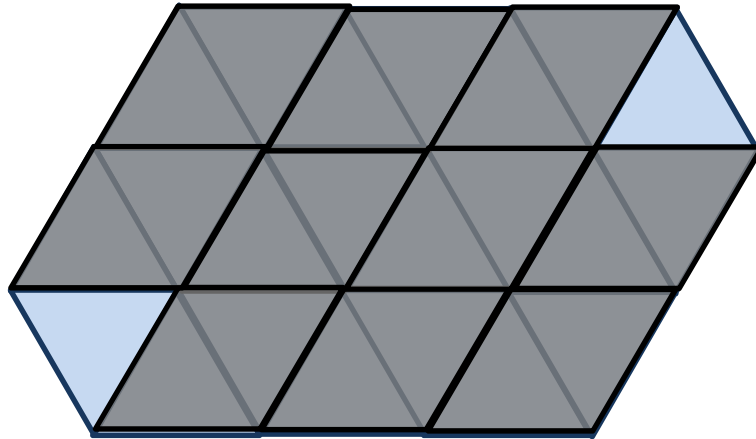
**1, 2, 3, 4 and 6**

In 2D, the lattice is formed by a repeated unit, translated in 2 directions:

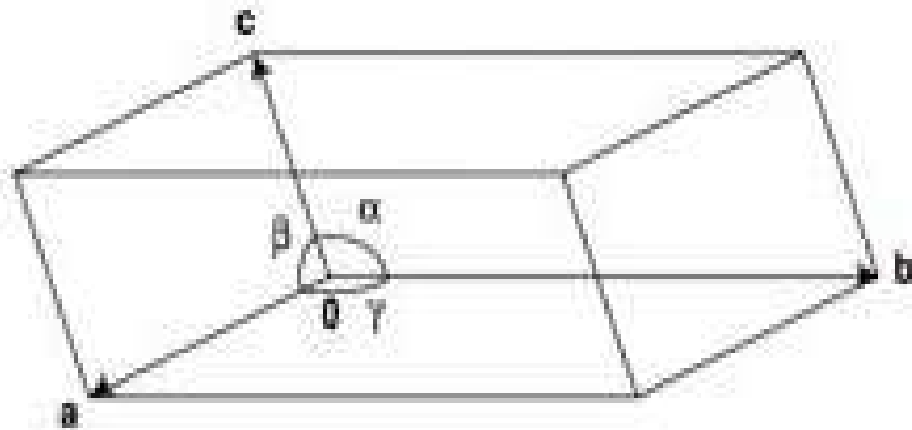


The repetitive unit is called **unit cell**. The lattice is defined by an **origin** of the translations and the **unit cell parameters** (or dimensions):  
(in 2D)  $a$ ,  $b$  and  $\gamma$

Even when symmetry elements include a 3-fold axis or a 6-fold axis, unit cell can be described as a parallelogram:



In 3D:



Unit cell parameters:  
 $a, b, c$  and  $\alpha, \beta, \gamma$

# Crystal families and crystal systems



Crystal family	Crystal system	Unit cell dimensions	Minimal symmetry elements
Triclinic	Triclinic	$a, b, c, \alpha, \beta, \gamma$	none
Monoclinic	Monoclinic	$a, b, c, \beta$ ( $\alpha, \gamma=90^\circ$ )	2
Orthorhombic	Orthorhombic	$a, b, c$ ( $\alpha=\beta=\gamma=90^\circ$ )	three perpendicular 2 axes
Tetragonal	Tetragonal	$a, c$ ( $b=a, \alpha=\beta=\gamma=90^\circ$ )	4
Hexagonal	Trigonal	$a, c$ ( $b=a, \alpha=\beta=90^\circ, \gamma=120^\circ$ )	3
	Hexagonal		6
Cubic	Cubic	$a$ ( $b=a, c=a, \alpha=\beta=\gamma=90^\circ$ )	four 3 axes along diagonal directions



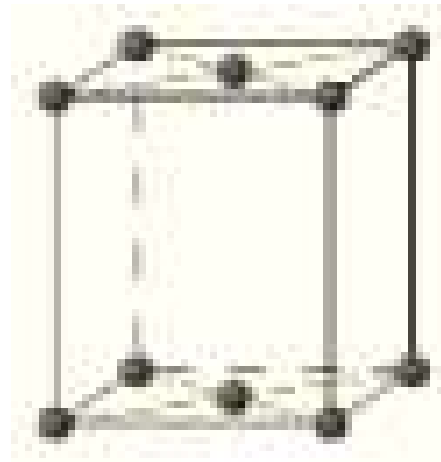
# Primitive and centered lattices

**Primitive** lattices have *nodal points (nodes)* at the vertices of the unit cell.

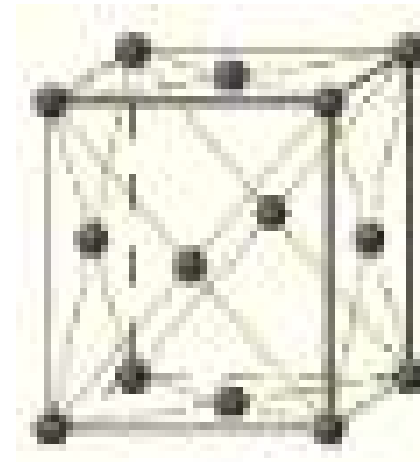
But further translations are possible in the unit cell, forming **centered lattices**, with additional nodes:



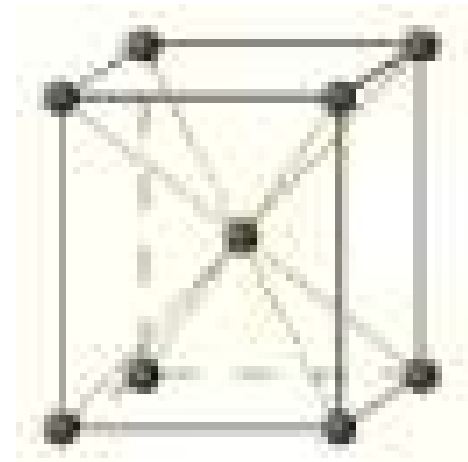
*Primitive  
lattice*



*Base centered*



*Face centered*



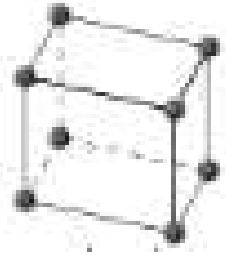
*Body  
centered  
lattice*

*Face(s) centered  
lattices*

A primitive lattice can **always** describe the structure, but in presence of specific elements of symmetry it can be useful to describe the lattice as centered.

# Crystal families + centering = 14 Bravais lattices

## Triclinic lattice

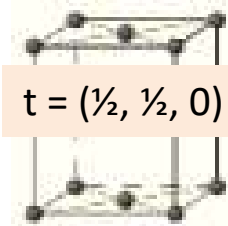


Primitive, *P*

## Orthorhombic lattice

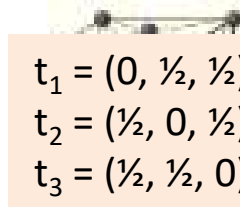


Primitive, *P*



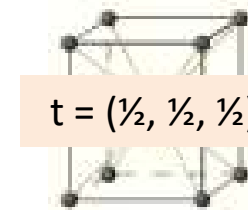
$$t = (\frac{1}{2}, \frac{1}{2}, 0)$$

Base centered, *C*



$$\begin{aligned} t_1 &= (0, \frac{1}{2}, \frac{1}{2}) \\ t_2 &= (\frac{1}{2}, 0, \frac{1}{2}) \\ t_3 &= (\frac{1}{2}, \frac{1}{2}, 0) \end{aligned}$$

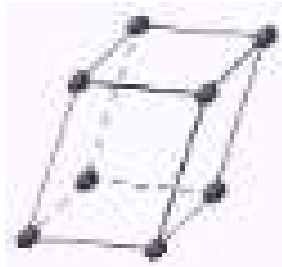
Face centered, *F*



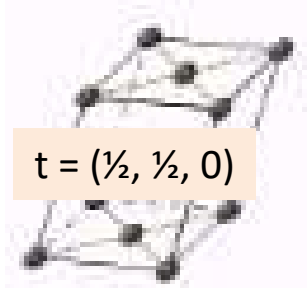
$$t = (\frac{1}{2}, \frac{1}{2}, \frac{1}{2})$$

Body centered, *I*

## Monoclinic lattice



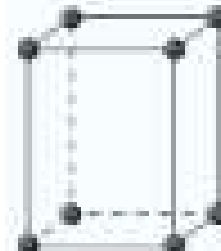
Primitive, *P*



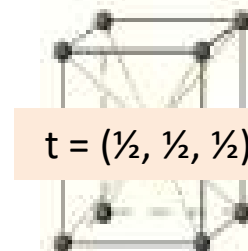
$$t = (\frac{1}{2}, \frac{1}{2}, 0)$$

Base centered, *C*

## Tetragonal lattice



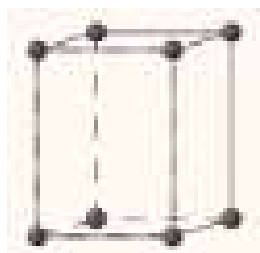
Primitive, *P*



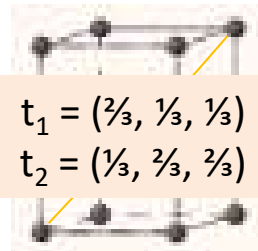
$$t = (\frac{1}{2}, \frac{1}{2}, \frac{1}{2})$$

Body centered, *I*

## Hexagonal lattice



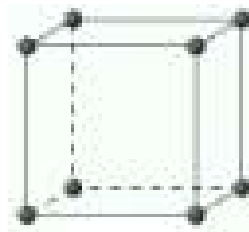
Primitive, *P*



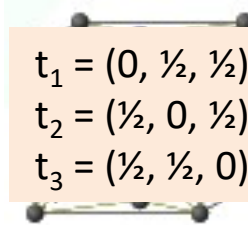
$$\begin{aligned} t_1 &= (\frac{2}{3}, \frac{1}{3}, \frac{1}{3}) \\ t_2 &= (\frac{1}{3}, \frac{2}{3}, \frac{2}{3}) \end{aligned}$$

Rhombohedral, *R*

## Cubic lattice

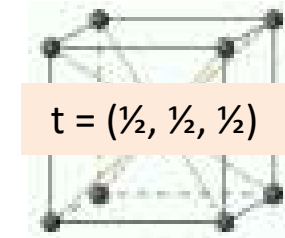


Primitive, *P*



$$\begin{aligned} t_1 &= (0, \frac{1}{2}, \frac{1}{2}) \\ t_2 &= (\frac{1}{2}, 0, \frac{1}{2}) \\ t_3 &= (\frac{1}{2}, \frac{1}{2}, 0) \end{aligned}$$

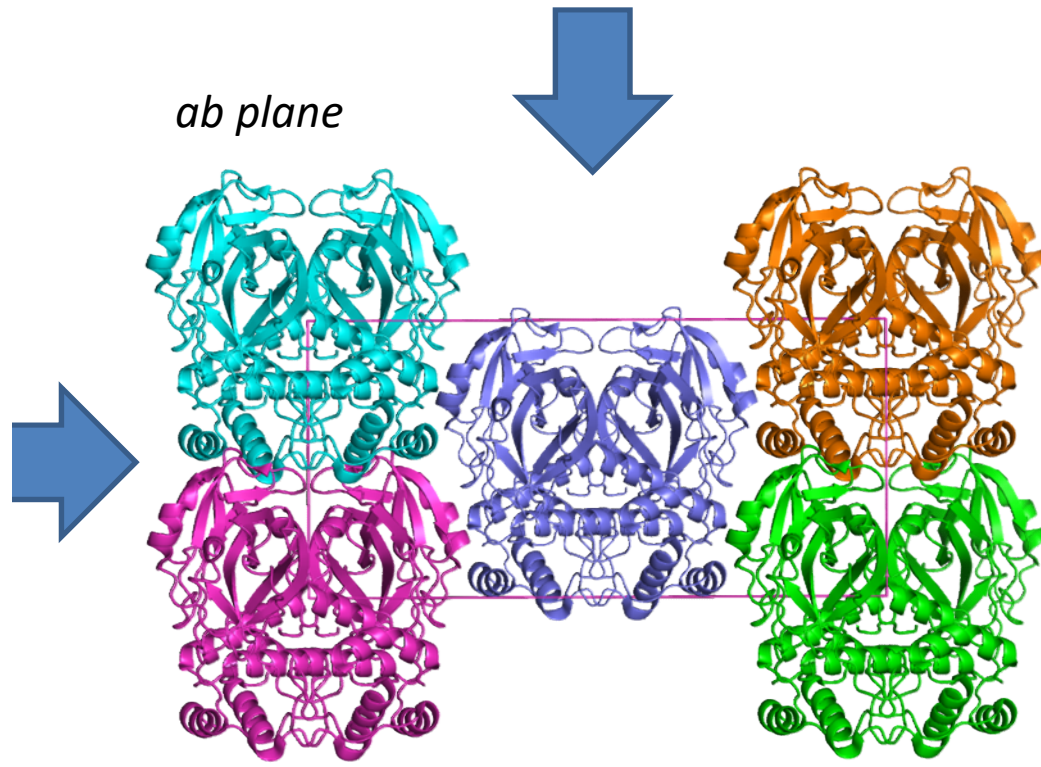
Face centered, *F*



$$t = (\frac{1}{2}, \frac{1}{2}, \frac{1}{2})$$

Body centered, *I*

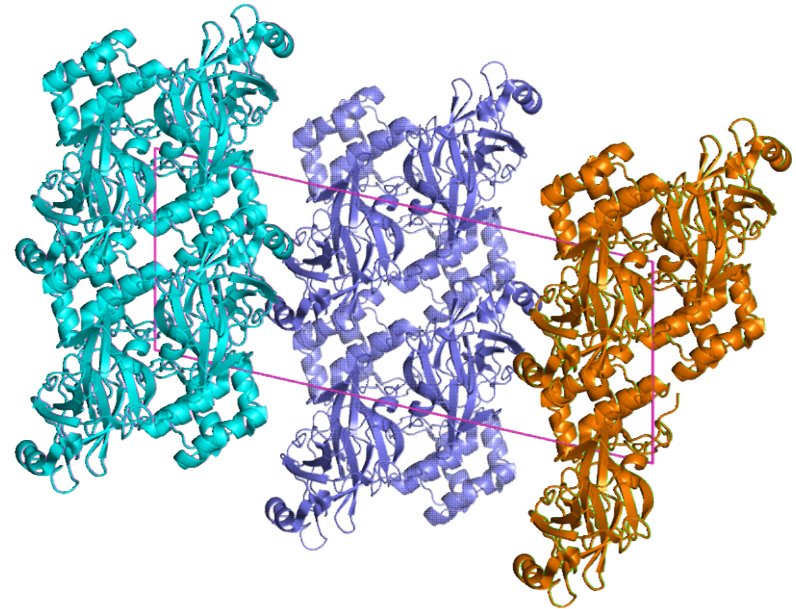
Case in point: monoclinic C2 lattice



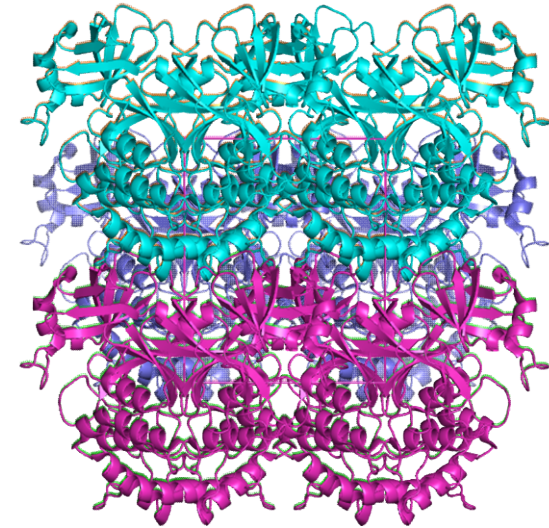
Monoclinic:  
 $a, b, c$   
 $\alpha = \gamma = 90^\circ, \beta$

5RE9

*ac plane*



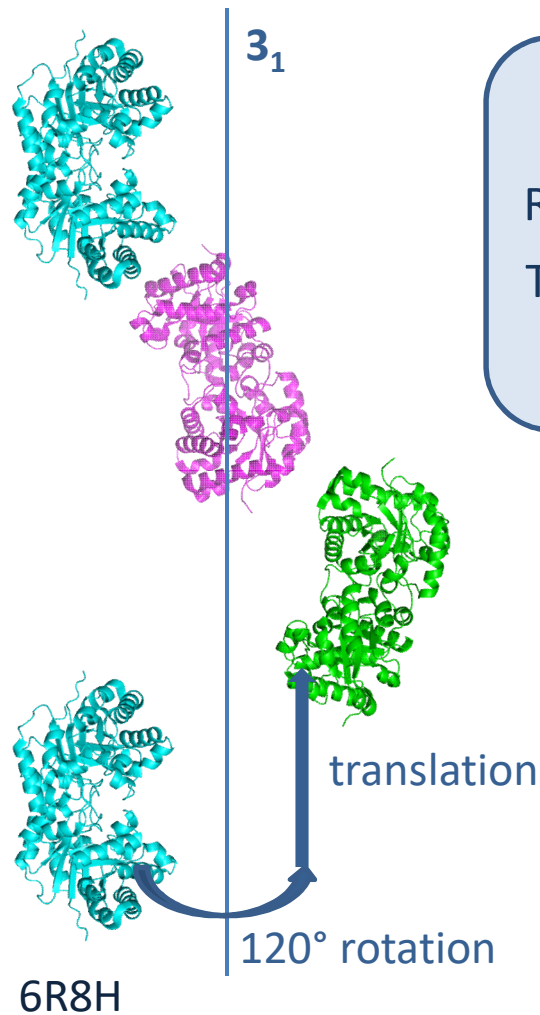
*bc plane*



# Rotation + translation = screw axis

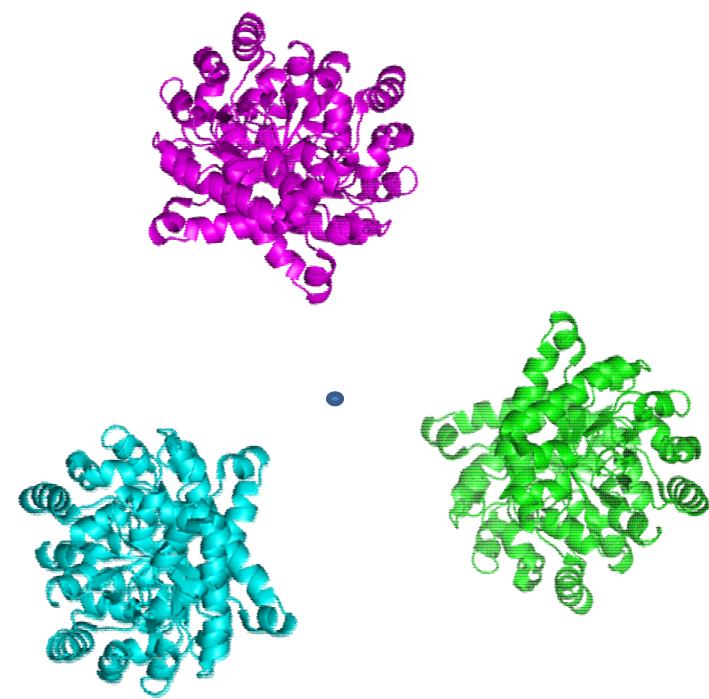
Besides the point group symmetry elements, additional symmetry elements are possible in crystal cells:

- **screw axis** = rotation + translation
- ~~glide planes = mirror + translation~~



**Screw axis  $N_m$**   
Rotation of  $360^\circ/N$   
Translation of  $m/N$   
of the unit cell

**Screw axis  $3_1$**   
Rotation of  $360^\circ/3$   
Translation of  $1/3$  of the unit cell

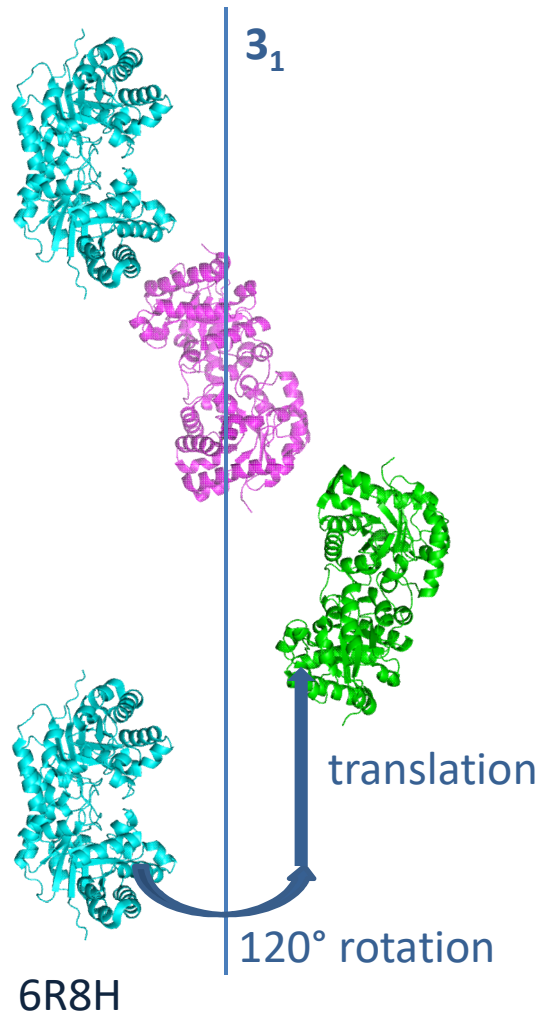


# Rotation + translation = screw axis

Screw axis  $3_1$

Rotation of  $360^\circ/3$

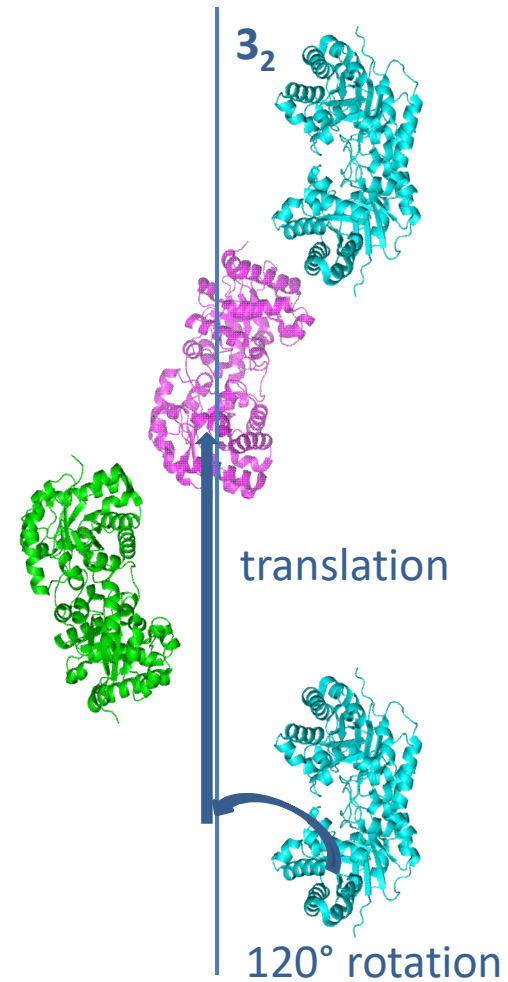
Translation of  $1/3$  of the unit cell



Screw axis  $3_2$






















Rotation of  $360^\circ/3$

Translation of  $2/3$  of the unit cell



# Symmetry operations

Symmetry operations are represented with specific figures when perpendicular to the plane, or arrows when parallel to the plane of the figure:

	2-fold axis		
	$2_1$ screw axis		
	3-fold axis		
	$3_1$ screw axis	} Rototranslations in opposite directions!	 Rotation axes
	$3_2$ screw axis		
	 4-fold axis		
	 $4_1$ screw axis		
	 $4_2$ screw axis		
	 $4_3$ screw axis		
	6-fold axis		
	$6_1$ screw axis	} Rototranslations in opposite directions!	 Rototranslation axes
	$6_2$ screw axis		
	$6_3$ screw axis		
	$6_4$ screw axis		
	$6_5$ screw axis		

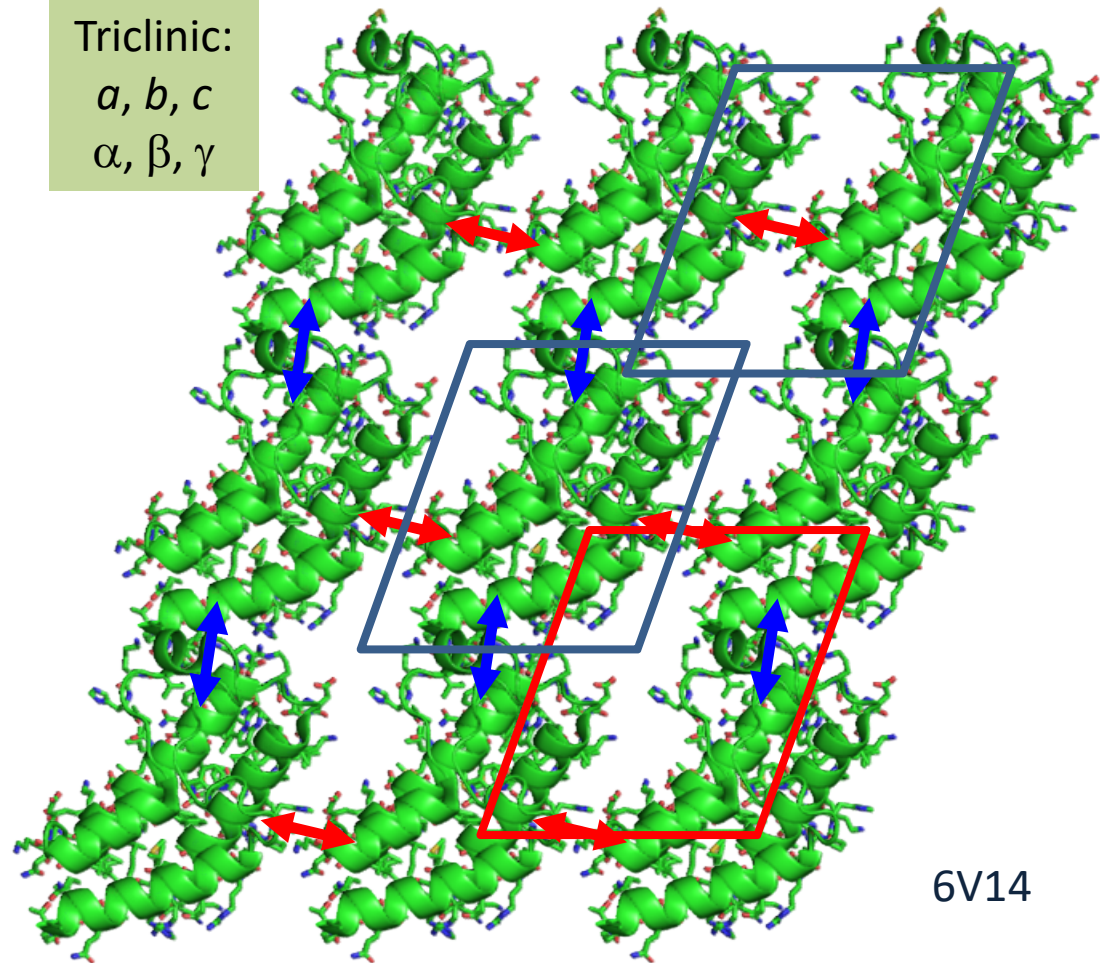
# Intermolecular contacts and crystal packing

Crystals are held together by intermolecular contacts, that determine their packing:

**Crystal packing** must be evaluated:

- 1) to evaluate the effect of crystal contacts on protein conformation
- 2) to obtain biologically active unit (for symmetry related oligomers)

Triclinic:  
 $a, b, c$   
 $\alpha, \beta, \gamma$



Different **origin** choices are possible (while unit cell dimensions are the same!) for a lattice with no additional symmetry.

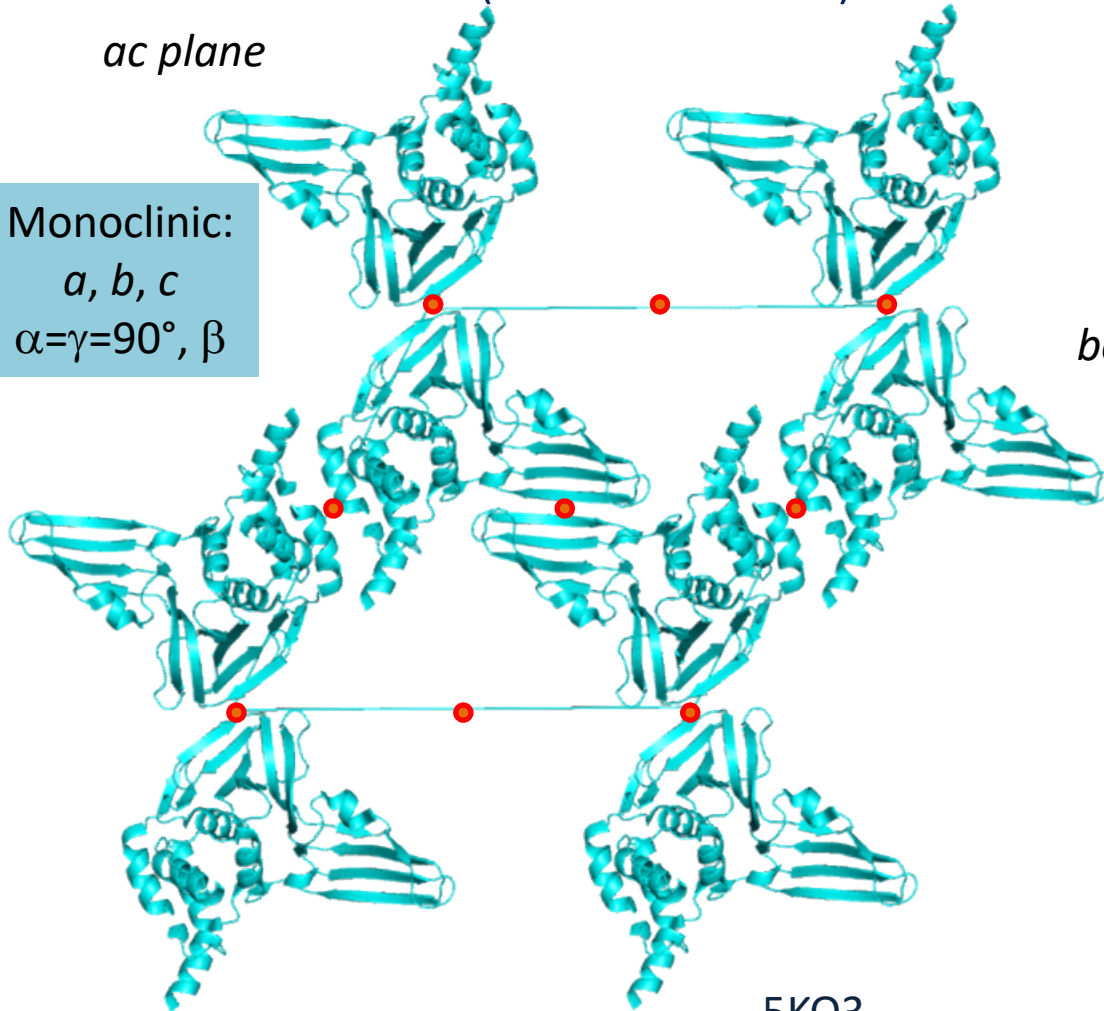
# Choice of the origin

The choice of the origin depends on the symmetry of the crystal lattice.

Case in point: lattice with 2-fold axis along  $b$   
(monoclinic lattice)

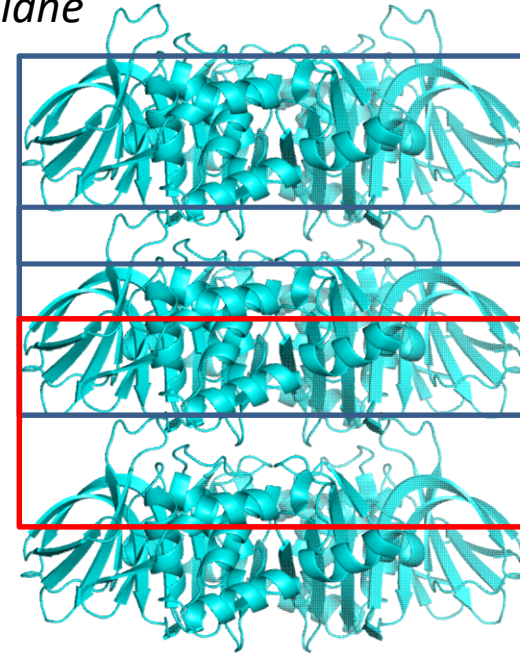
$ac$  plane

Monoclinic:  
 $a, b, c$   
 $\alpha=\gamma=90^\circ, \beta$

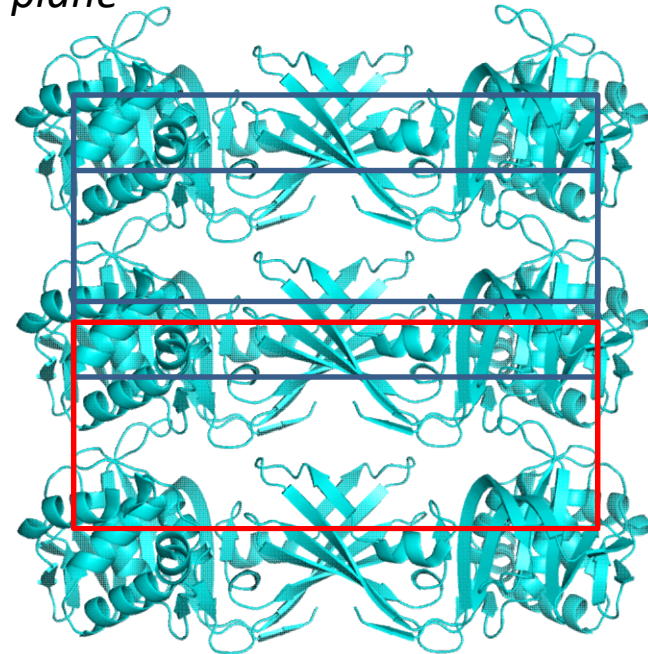


5KO3

$ab$  plane



$bc$  plane

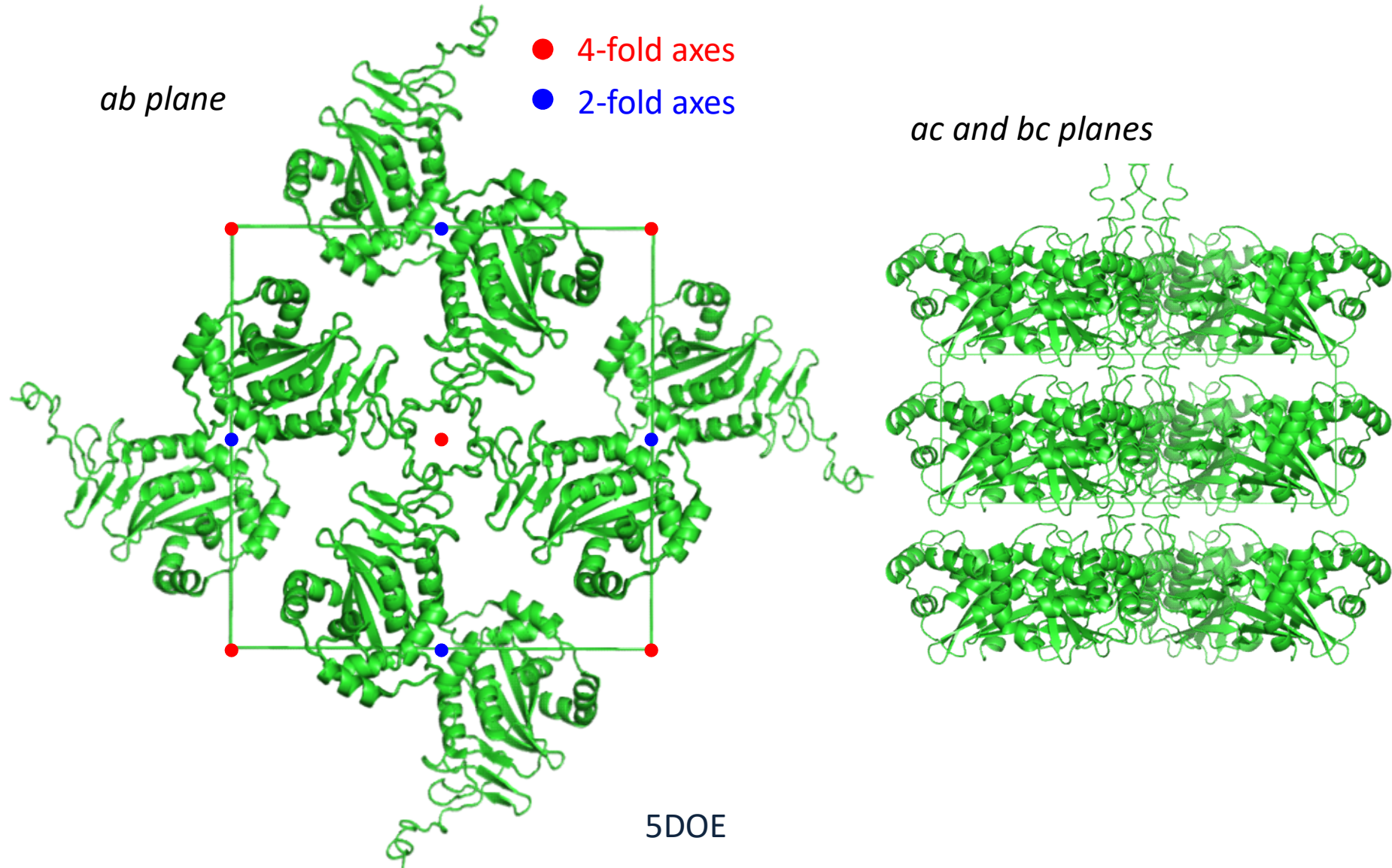




# Choice of the origin

Case in point: lattice with 4-fold axis along  $c$  (tetragonal lattice)

Tetragonal:  
 $a=b, c$   
 $\alpha=\beta=\gamma=90^\circ$



# Fractional coordinates

Each symmetry operation can be represented by:

(1) a square matrix  $\mathbf{S}$  and (2) a translation vector  $\mathbf{t}$  that transform a generic point  $\mathbf{x}$  with coordinates  $(x,y,z)$  in its symmetric  $\mathbf{x}'$  with coordinates  $(x',y',z')$ :  $\mathbf{x}' = \mathbf{S} \cdot \mathbf{x} + \mathbf{t}$

$$\begin{pmatrix} x' \\ y' \\ z' \end{pmatrix} = \begin{bmatrix} S_{11} & S_{12} & S_{13} \\ S_{21} & S_{22} & S_{23} \\ S_{31} & S_{32} & S_{33} \end{bmatrix} \cdot \begin{pmatrix} x \\ y \\ z \end{pmatrix} + \begin{pmatrix} t_x \\ t_y \\ t_z \end{pmatrix}$$

The coordinate system of this equation, however, is not the orthogonal system in which we describe the atomic structure with distances in Å (i.e. in the pdb file).

The matrices  $\mathbf{S}$  and vectors  $\mathbf{t}$  are defined in the coordinate system of each specific lattice, i.e. having the  $x,y,z$  directions along the translational vectors that define the lattice. In addition, this coordinate system has **fractional coordinates**, that are calculated as fractions of the unit cell parameters in each direction.

Example: the coordinate system of a monoclinic lattice has 2 directions ( $x$  and  $z$ ) that are not perpendicular, but form an angle equal to  $\beta$ .

Transformation: 4-fold axis along  $c$  direction of a tetragonal lattice

$$\begin{pmatrix} x' \\ y' \\ z' \end{pmatrix} = \begin{bmatrix} 0 & 1 & 0 \\ -1 & 0 & 0 \\ 0 & 0 & 1 \end{bmatrix} \cdot \begin{pmatrix} x \\ y \\ z \end{pmatrix} + \begin{pmatrix} 0 \\ 0 \\ 0 \end{pmatrix}$$

Transformation:  $2_1$  screw axis along the  $b$  direction

$$\begin{pmatrix} x' \\ y' \\ z' \end{pmatrix} = \begin{bmatrix} -1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & -1 \end{bmatrix} \cdot \begin{pmatrix} x \\ y \\ z \end{pmatrix} + \begin{pmatrix} 0 \\ 1/2 \\ 0 \end{pmatrix}$$