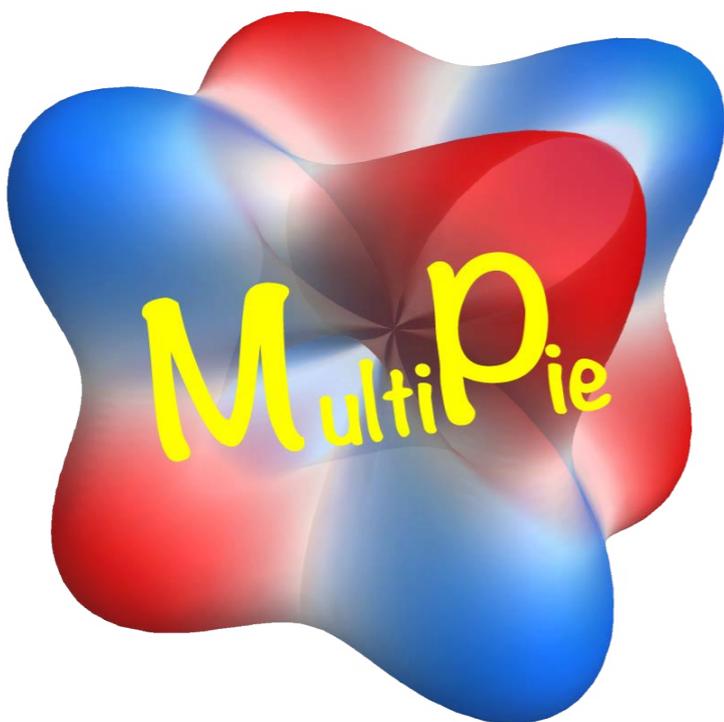


**Install Guide (in Japanese)** [https://cmt-mu.github.io/QtDraw/install\\_guide.pdf](https://cmt-mu.github.io/QtDraw/install_guide.pdf)

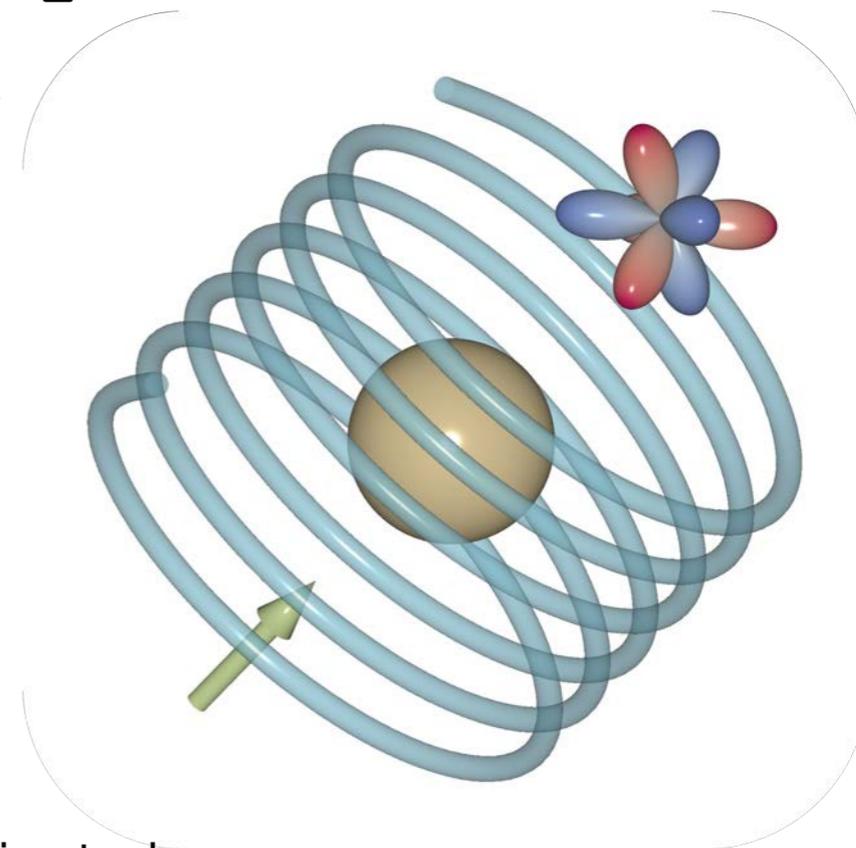
## MultiPie



- Symmetry operations for point/space groups
- Symmetry-adapted basis construction  
(*"MultiPy"* was already registered in PyPI, umm...)

<https://cmt-mu.github.io/MultiPie/>

## QtDraw



- 3D drawing tool  
(including orbital, vector stream spline curve...)
- With MultiPie, draw symmetry-related objects, projection to each Irreps. (Symmetry-Adapted)

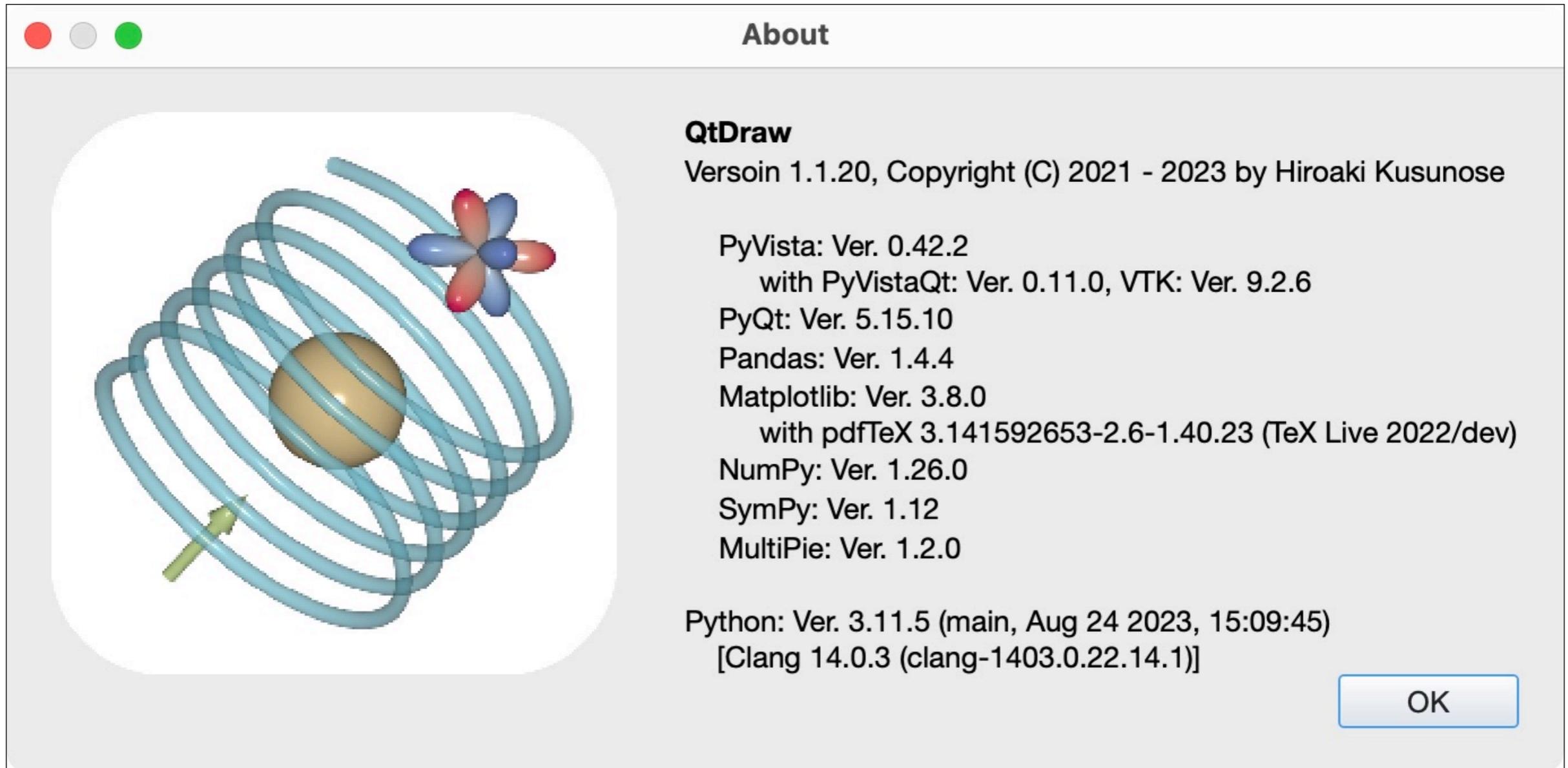
<https://cmt-mu.github.io/QtDraw/>

If you are using MultiPie and/or QtDraw in your scientific research, please help our scientific visibility by citing our work:

*Hiroaki Kusunose, Rikuto Oiwa, and Satoru Hayami, Symmetry-adapted modeling for molecules and crystals, Phys. Rev. B **107**, 195118 (2023).*

DOI: <https://doi.org/10.1103/PhysRevB.107.195118>

# Checked Library Version for QtDraw



## MacBook Air

15インチ、M2、2023

チップ Apple M2  
メモリ 8 GB  
シリアル番号   
macOS Ventura 13.5

QtDraw can read

- `.qtdw` : QtDraw file
- `.cif` : CIF file
- `.vesta` : VESTA file

# Concept of Symmetry-Adapted Basis

HK, R. Oiwa, and S. Hayami, PRB **107**, 195118 (2023)

## Neumann Principle

Every physical phenomenon manifested by a crystal must possess an equivalent or higher symmetry as the crystal itself.



(Static) macroscopic response can be classified by POINT GROUP symmetry ( $\mathbf{k} = 0 : \Gamma$  point in BZ)

## Classification for Materials

### For TP spherical systems

T : time-reversal (electric + or magnetic -)  
 P : spatial-inversion (polar or axial)  
 g : anisotropy (l, m)



### Molecule and Crystal

Subgroup of spherical systems

Linear combination of spherical base

$$Z_{l\gamma}^{(\Gamma, n)} = \sum_m c_{\gamma m}^{(\Gamma, n)} Z_{lm}$$

Type ( $Z_{lm}$ )	Symbol	T	P		Charge	g
Electric (E)	$Q_{lm}$	+	polar	$(-1)^l$	1	$Y_{lm}$
Magnetic (M)	$M_{lm}$	-	axial	$-(-1)^l$	$i(\mathbf{e}_1 \times \mathbf{e}_2) \cdot \mathbf{e}_3$	$Y_{lm}$
Magnetic-Toroidal (MT)	$T_{lm}$	-	polar	$(-1)^l$	$i$	$Y_{lm}$
Electric-Toroidal (ET)	$G_{lm}$	+	axial	$-(-1)^l$	$(\mathbf{e}_1 \times \mathbf{e}_2) \cdot \mathbf{e}_3$	$Y_{lm}$

# Construction of Basis

## Basis for Materials

Electronic degrees of freedom = (atomic d.o.f.)  $\otimes$  (site/bond d.o.f.)

$$\mathbb{Z}_\alpha = \sum_{\beta\gamma} C_\alpha^{\beta,\gamma} \mathbb{X}_\beta \otimes \mathbb{Y}_\gamma$$

"Clebsch-Gordan" coefficient

### Atomic d.o.f

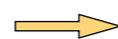
Given by quantum-mechanical expressions of complete set,  
compute matrix elements by using general formula

HK, R. Oiwa, and S. Hayami, JPSJ **89**, 104704 (2020)

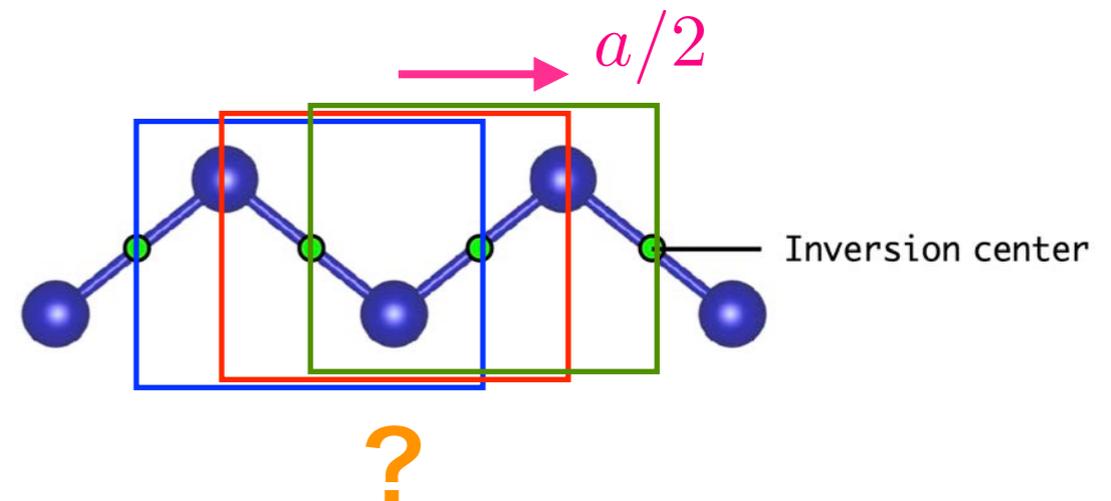
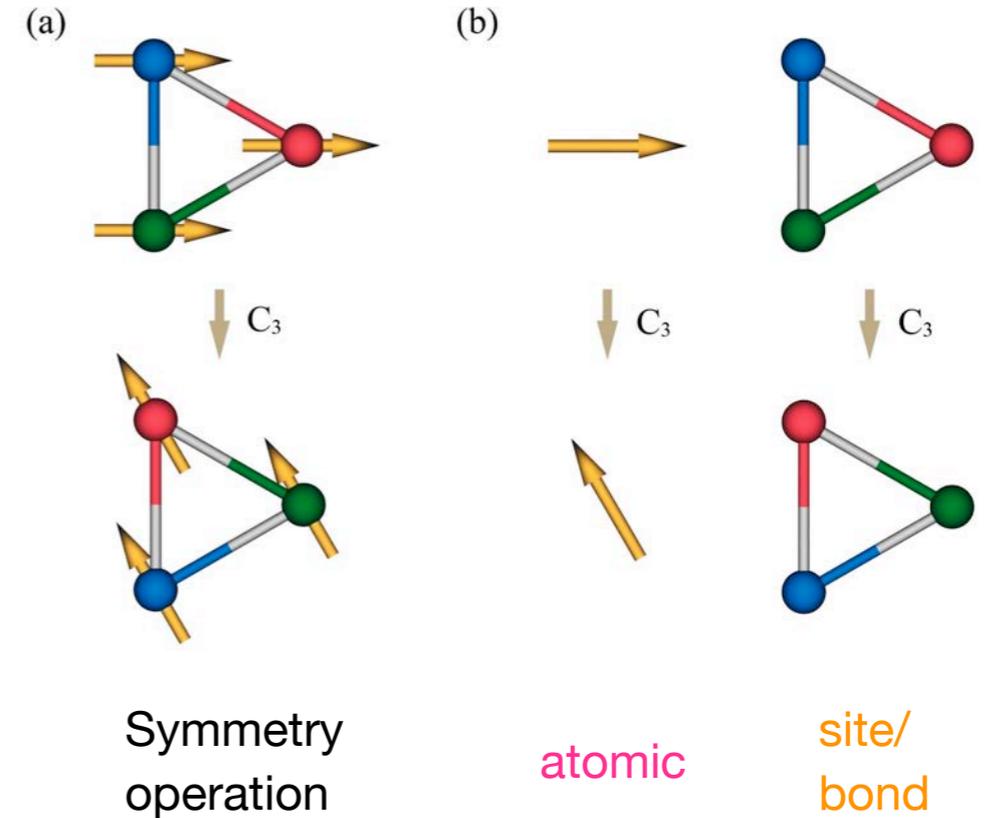
### Site/Bond d.o.f

Ambiguity

- How to choose cluster and center (translation) ?
- Non isoradius (nonsymmorphic = screw or glide) ?



Use "Virtual Cluster"



# Concept of Virtual Cluster

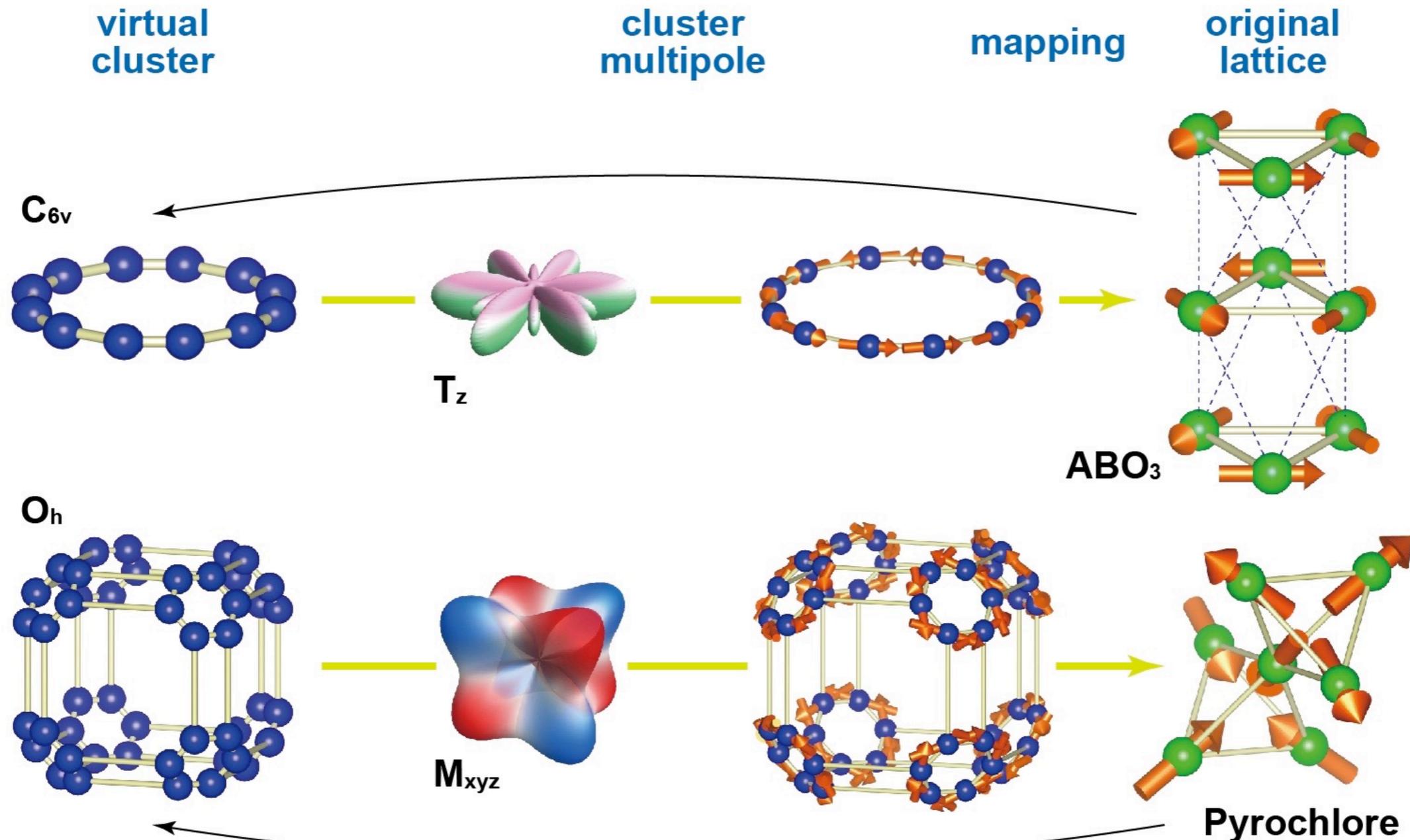
## Virtual Cluster

Site-Cluster

M.-T. Suzuki, et al., PRB **99**, 174407 (2019)

Site/Bond-Cluster

HK, R. Oiwa, and S. Hayami, PRB **107**, 195118 (2023)



# MultiPie - Example

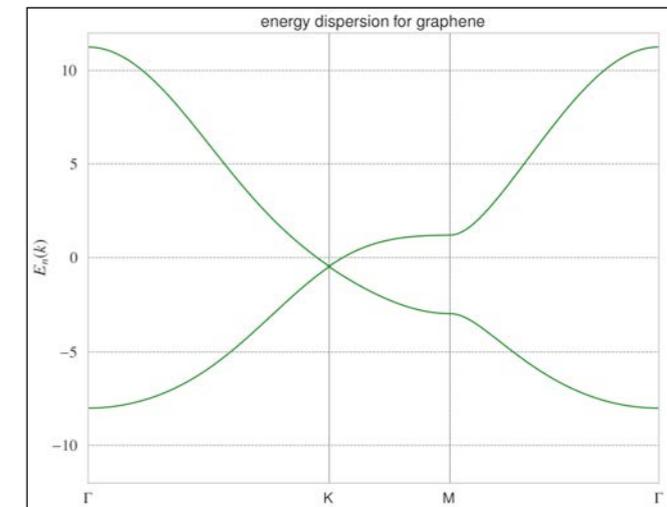
## Example for graphene

Download

<https://github.com/CMT-MU/MultiPie/tree/main/docs/example>  
and try "python create\_plot.py"

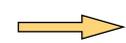
**Input file** (graphene.py : Python dict format)

```
graphene = {  
  "model": "graphene", # name of model.  
  "group": 191, # No. of space group.  
  "cell": { "c": 4 }, # set large enough interlayer distance.  
  #  
  "site": { "C": ( "[1/3,2/3,0]", "pz" ) }, # positions of C site and its orbital.  
  "bond": [ ( "C", "C", [1, 2, 3, 4, 5, 6] ) ], # C-C bonds up to 6th neighbors.  
  #  
  "spinful": False, # spinless.  
  #  
  "k_point": { "Γ": "[0, 0, 0]", "M": "[1/2, 0, 0]", "K": "[1/3, 1/3, 0]" }, # def. of k points.  
  "k_path": "Γ-K-M-Γ", # high-symmetry line.  
}
```



## Create Symmetry-Adapted Multipole Basis (SAMB)

```
create_samb graphene
```



In "graphene" folder

See, document for detailed dict structure

- |                      |                    |                      |                         |
|----------------------|--------------------|----------------------|-------------------------|
| • graphene_model.py  | (Model info.)      | • graphene_samb.tex  |                         |
| • graphene_matrix.py | (Full matrix form) | • graphene_samb.pdf  | (SAMB info. for human)  |
| • graphene_samb.py   | (SAMB info.)       | • graphene_view.qtdw | (structure QtDraw file) |

# MultiPie - Output of Full Matrix

Full matrix form file  $\longrightarrow H = \sum_j z_j \mathbb{Z}_j$

```
graphene = {
  "model": "graphene",
  "molecule": False,
  "group": ("D6h^1", "space group No. 191 : D6h^1 / P6/mmm : PG D6h"),
  "dimension": 2, # matrix size.
  "ket": ["pz@C_1", "pz@C_2"], # Hilbert space : orbital @ site.
  "cell_site": { # site : position, symmetry operations.
    "C_1": ("[1/3, 2/3, 0]", "[1,6,7,8,9,10,14,15,16,17,23,24]"),
    "C_2": ("[2/3, 1/3, 0]", "[2,3,4,5,11,12,13,18,19,20,21,22]"),
  },
  "version": "1.1.15",
  "k_point": {"Gamma": "[0, 0, 0]", "M": "[1/2, 0, 0]", "K": "[1/3, 1/3, 0]"},
  "k_path": "Gamma-K-M-Gamma",
  "A": "[[1.0, -0.5, 0.0], [0.0, 0.86602540378444, 0.0], [0.0, 0.0, 4.0]]", # unit vectors : [a1,a2,a3].
  "matrix": { # SAMB.
    # z_# : { (n1,n2,n3,a,b), matrix element }, where lattice vector, R = [n1,n2,n3], a, b : row and col. in full matrix.
    "z_001": {(0, 0, 0, 0, 0): "sqrt(2)/2", (0, 0, 0, 1, 1): "sqrt(2)/2"},
    ...
    "z_007": {(2, 2, 0, 0, 0): "sqrt(3)/6", (-2, -2, 0, 0, 0): "sqrt(3)/6",
              (2, 2, 0, 1, 1): "sqrt(3)/6", (-2, -2, 0, 1, 1): "sqrt(3)/6",
              (0, 2, 0, 1, 1): "sqrt(3)/6", (0, -2, 0, 1, 1): "sqrt(3)/6",
              (2, 0, 0, 1, 1): "sqrt(3)/6", (-2, 0, 0, 1, 1): "sqrt(3)/6",
              (0, 2, 0, 0, 0): "sqrt(3)/6", (0, -2, 0, 0, 0): "sqrt(3)/6",
              (2, 0, 0, 0, 0): "sqrt(3)/6", (-2, 0, 0, 0, 0): "sqrt(3)/6"},
  }
}
```

[graphene\\_matrix.py](#)

## Fourier transform of full matrix

$$[\mathbb{Z}_j(\mathbf{k})]_{a,b} = \sum_{n_1, n_2, n_3} z_j[(n_1, n_2, n_3, a, b)] e^{2\pi i \mathbf{k} \cdot (\mathbf{R} + \mathbf{r}_a - \mathbf{r}_b)}$$

e.g. :  $r_a = \text{cell\_site}[\text{ket}[a].\text{split}("@")[1]] [0]$   
 $r_b = \text{cell\_site}[\text{ket}[b].\text{split}("@")[1]] [0]$

# MultiPie - Output of Non-Identity Irrep.

## Input file

```
graphene = {
  ....
  "generate": {
    "time_reversal_type": "both",
    "irrep": ["A1g", "B1u", "A2g", "A2u"],
  },
  "spinful": True,
}
```

Consider "symmetry-breaking" terms

### 1. Mass term (spinless)

$$Q_{3, B_{1u}} \quad \text{"C" site-cluster}$$

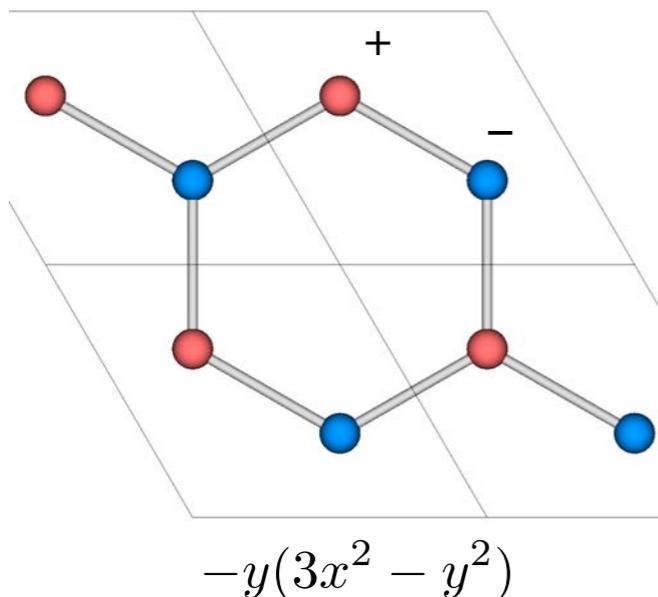
### 2. Haldane's magnetic flux term from kinetic SOC (spinless)

$$M_{1, A_{2g}} \quad \text{"C-C" 2nd neighbor bond-cluster} \quad M_z \in A_{2g}$$

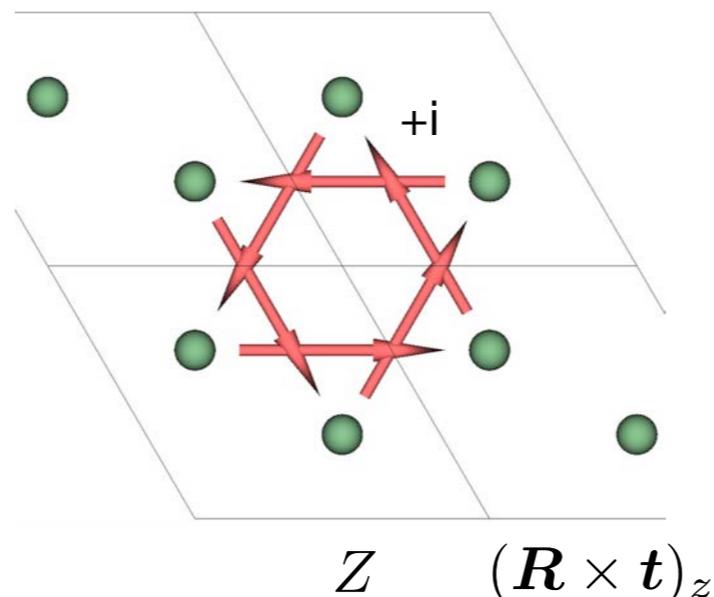
### 3. Surface Rashba term from z-polar field (spinful) $E_z \in A_{2u}$

$$Q_{1, A_{2u}} \quad \text{"C-C" 1st neighbor bond-cluster}$$

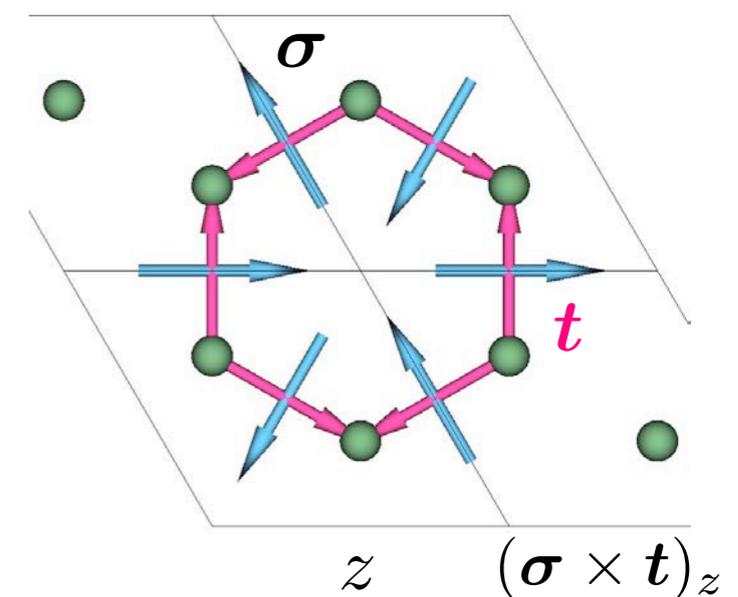
### 1. Mass term



### 2. Haldane's M-flux



### 3. Rashba



# QtDraw - Features

based on Qt Python and PyVista

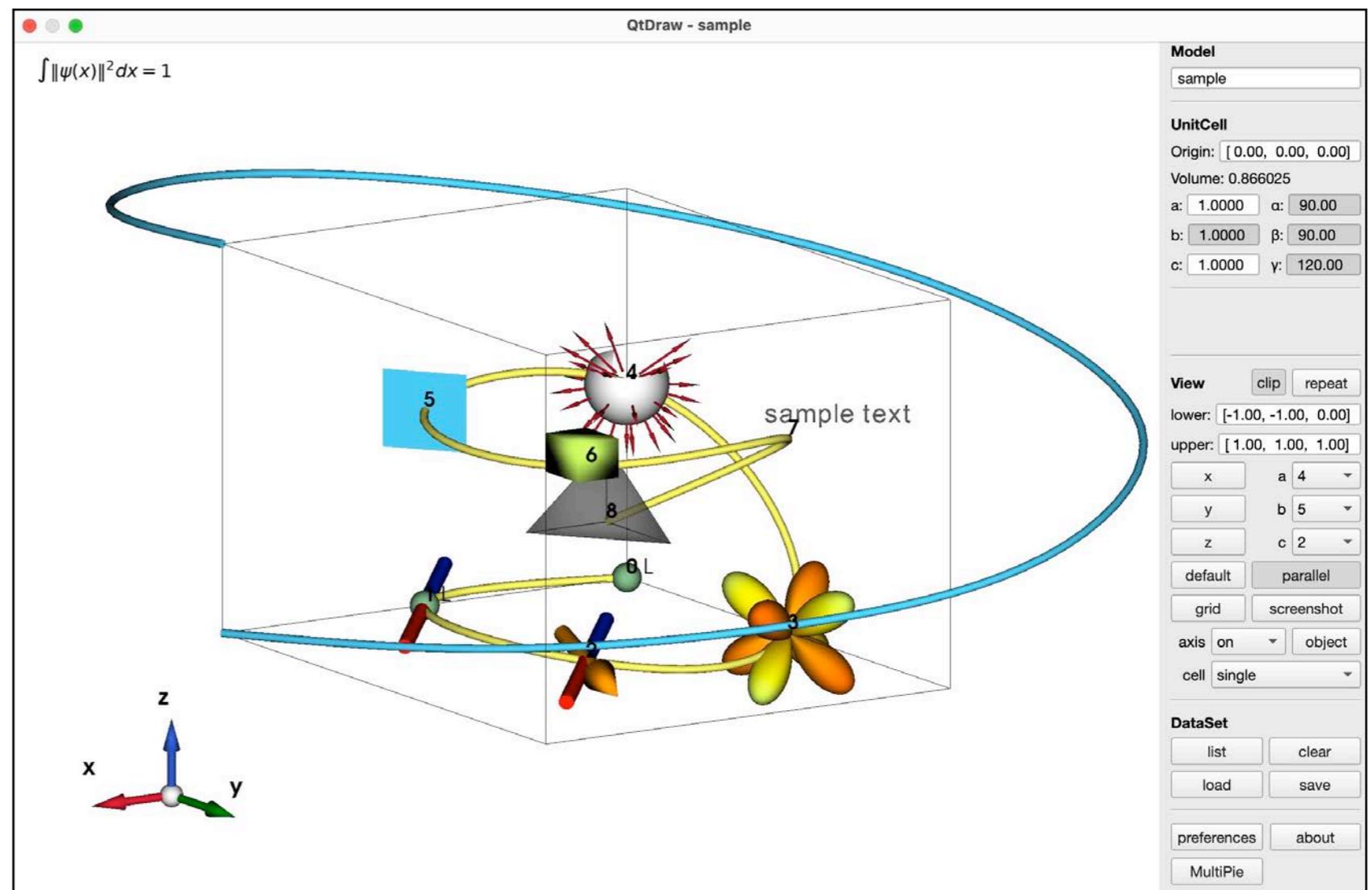
<https://www.qt.io/qt-for-python>

<https://docs.pyvista.org/version/stable/>

tested mainly on Mac (managed to work out on Windows, no check on Linux)

## Drawable objects

- Sphere (site)
- Bond (monotone/two-tone color)
- Vector
- Stream vector
- Plane
- Box
- Polygon
- Text (3d, 2d)
- Spline curve (data or function)
- Caption



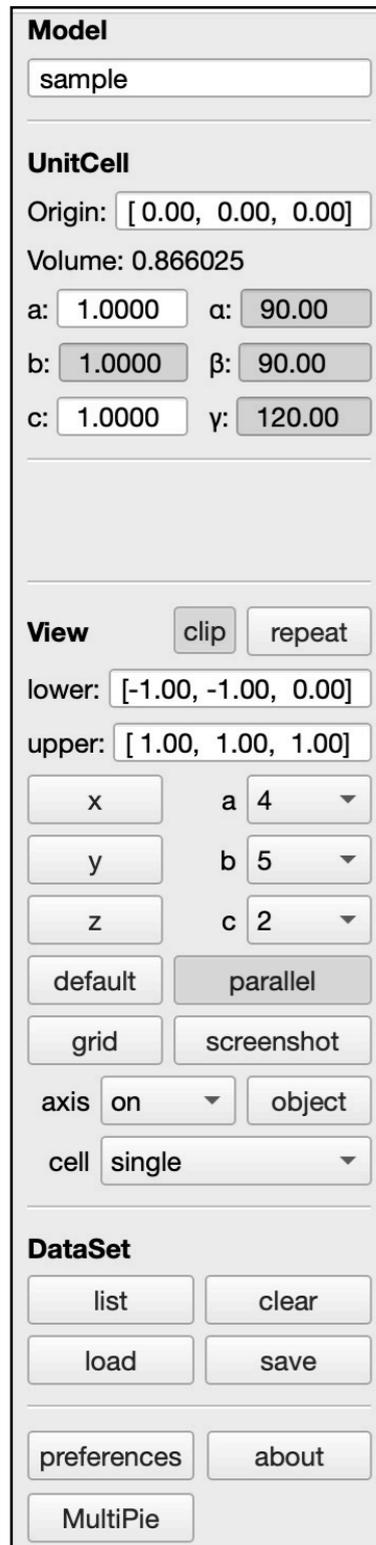
All objects can be drawn by calling **Python API** : easy to extend functionality of drawing

e.g. spin/orbital modulation patterns

# QtDraw - Main Menu

\*\*\* NO UNDO functionality at this version \*\*\*

## Menu



model name

origin (reduced)

lattice const. & angle

clip/repeat on/off

clip/repeat range

viewpoint x/y/z/default  
index (a,b,c)

parallel or perspective view

grid on/off, screenshot : [png/bmp/tif/tiff/svg/eps/ps/pdf]

axis, cell on/off

list/clear data

load/save data (.qtdw file)

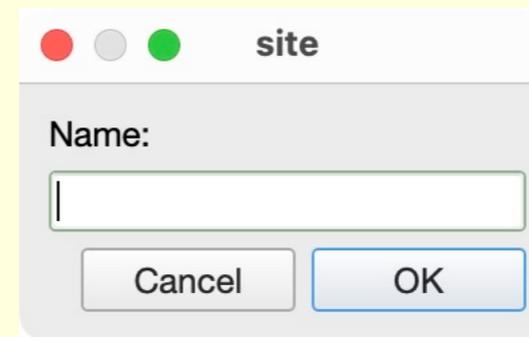
setting options

use MultiPie

• Create / add / remove objects

• Edit properties of objects

Add : create object in new group



specify  
group name

Edit properties of objects in group or individual object

Insert : insert object in selected group

Remove : remove selected object

Coordinate: reduced one with unit vectors

[x, y, z] in Cell + Cell position

Draw objects in a Cell, and then repeat



# QtDraw - Object Property

## Site

size	color
0.50	 darkseagreen

## Bond

vector	width	color	color2
[-0.5000, -0.5000, 0.0000]	1.00	 red	 blue

## Vector

vector	length	width	offset	color
[0.0000, 0.5000, 0.0000]	0.40	1.00	-0.43	 orange

cartesian

## Orbital

shape	surface	size	scale	theta0	theta1	phi0	phi1	color
<i>xyz</i>	<i>xyz</i>	0.20	<input checked="" type="checkbox"/>	0	180	0	360	 Wistia

$\theta$  range

$\phi$  range

## Stream

shape	vector	size	v_size	width	scale	theta	phi	theta0	theta1	phi0	phi1	color	component
1	$\begin{pmatrix} x \\ y \\ z \end{pmatrix}$	0.10	0.15	1.00	<input type="checkbox"/>	4	8	0	120	0	270	 coolwarm	abs

stream  
on shape

vector at  $r$

# arrow  
in  $(\theta, \phi)$

$\theta$  range

$\phi$  range

color  
based on

# QtDraw - Object Property

---

## Plane

normal	x	y	color
[1.00, 1.00, 0.00]	0.20	0.20	 sky

reduced      reduced

## Box

a1	a2	a3	edge	wireframe	width	color
[0.1000, 0.0000, 0.0000]	[0.0000, 0.1000, 0.0000]	[0.0000, 0.0000, 0.1000]	<input type="checkbox"/>	<input type="checkbox"/>	1.00	 honeydew

reduced      reduced      reduced

## Polygon

point	connection	edge	wireframe	width	color
[0.0000, 0.0000, 0.0000]	[0, 1, 2]				
[0.2000, 0.0000, 0.0000]	[0, 1, 3]	<input checked="" type="checkbox"/>	<input type="checkbox"/>	2.00	 aluminum
[0.0000, 0.2000, 0.0000]	[1, 2, 3]				
[0.0000, 0.0000, 0.2000]	[2, 0, 3]				

reduced      connecting point #s

## Text3d

text	size	depth	normal	offset	color
sample text	1.00	3.00	[4.00, 5.00, 2.00]	[0.1000, 0.1000, 0.1000]	 iron

reduced      reduced

# QtDraw - Object Property

## Spline

point	width	n_interp	closed	natural	color
[0.0000, 0.0000, 0.0000]					
[0.5000, 0.0000, 0.0000]					
[0.5000, 0.5000, 0.0000]					
[0.0000, 0.5000, 0.0000]					
[0.0000, 0.0000, 0.5000]	1.00	100	<input type="checkbox"/>	<input checked="" type="checkbox"/>	banana
[0.5000, 0.0000, 0.5000]					
[0.5000, 0.5000, 0.5000]					
[0.0000, 0.5000, 0.5000]					
[0.2500, 0.2500, 0.2500]					

natural spline

# interpolation

reduced

## Spline\_t

expression	t_range	width	n_interp	closed	natural	color
$\begin{pmatrix} \cos(2\pi t) \\ \sin(2\pi t) \\ t \end{pmatrix}$	[0.0000, 1.1000, 0.1000]	1.00	100	<input type="checkbox"/>	<input checked="" type="checkbox"/>	sky

natural spline

# interpolation

reduced

## Caption

caption	space	size	bold	color
0				
1	caption			
2	text			
3				
4	0	18	<input checked="" type="checkbox"/>	licorice
5				
6	Text (2d)			
7				
8				

position	relative	caption	size	color	font
[0.02, 0.95]	<input checked="" type="checkbox"/>	$\int \psi(x)^2 dx = 1$	8	licorice	arial

position 2d (origin at left-top)

text (with simple LaTeX)

# QtDraw with MultiPie Enhanced (QtDraw<sup>+</sup>)

## Additional Panel

by pushing "MultiPie" in Menu

Group operations become available !

space/point group

product of irrep.

draw harmonics

active tensor

matrix element

draw VC at Wyckoff

space group  191. D6h<sup>1</sup> (P6/mmm)

symmetry operation  Wyckoff position

symmetric    anti-sym.  -

harmonics  rank  irrep. decomp.

response tensor  polar

atomic multipole  bra-ket

virtual cluster

decompose harmonics to PG

SITE: [x,y,z], BOND: [tail];[head] / [vector]@[center] / [start]:[vector]

sympy-style expression can be used

object drawing

SITE: draw equivalent sites.

1. input representative SITE, + ENTER.

draw sites by symmetry operation (SO)

BOND: draw equivalent bonds.

1. input representative BOND, + ENTER.

draw bonds by SO

VECTOR: draw vectors at equivalent sites or bonds.

1. choose type, 2. input vector [x,y,z] # representative SITE/BOND, + ENTER.

draw same vectors at positions by SO

type of MP

ORBITAL: draw orbitals at equivalent sites or bonds.

1. choose type, 2. input orbital (xyz polynomial) # representative SITE/BOND, + ENTER.

draw same orbitals at positions by SO

type of MP

POINT-GROUP HARMONICS: draw point-group harmonics at equivalent sites or bonds.

1. choose (type,rank,irrep.), 2. input representative SITE/BOND, + ENTER.

draw PG harmonics

⇒ used expression is shown (in LaTeX form).

expression

LaTeX

WYCKOFF: find wyckoff position and local symmetry. 1. input representative SITE/BOND, + ENTER.

⇒ wyckoff position and its local symmetry are shown.

find Wyckoff and local symmetry

⇒ Wyckoff position

local symmetry

type, rank, PG-MP

definition of PG-MP

# QtDraw+ - Drawing with Symmetry Operation

## Bond expression

Tail-Head :  $[1/2, 1/2, 0] ; [0, 0, 0]$   
 Vector-Center :  $[1/2, 1/2, 0] @ [0, 0, 0]$   $\longrightarrow$  object is drawn at bond center  
 Start-Vector :  $[1/2, 1/2, 0] : [0, 0, 0]$

## Basis drawing

object drawing    basis drawing

SITE: draw site-cluster basis.  
 1. input representative SITE, + ENTER,  
 ⇒ 2. choose basis, 3. push `draw`.

draw site-cluster basis

[ 1/2, 1/2, 0 ]

⇒  draw

BOND: draw bond-cluster basis.  
 1. input representative BOND, + ENTER,  
 ⇒ 2. choose basis, 3. push `draw`.

draw bond-cluster basis

[ 1/2, 1/2, 0 ] @ [ 1/4, 1/4, 0 ]

⇒  draw

VECTOR: draw symmetry-adapted vector.  
 1. choose type, 2. input representative SITE/BOND, + ENTER,  
 ⇒ 3. choose (type,basis), 4. push `draw` or 3. input linear combination (LC), + ENTER or 3. push `modulation`.

draw symmetry-adapted vector

Q  [ 1/2, 1/2, 0 ]

⇒ Q  draw

LC (Q01+Q02)/sqrt(2) draw linear combination of symmetry-adapted vectors

modulation Q,G  draw modulation of symmetry-adapted vectors

ORBITAL draw symmetry-adapted orbital.  
 1. choose (type,rank), 2. input representative SITE/BOND, + ENTER,  
 ⇒ 3. choose (type,basis), 4. push `draw` or 3. input linear combination (LC), + ENTER or 3. push `modulation`.

draw symmetry-adapted orbital

Q 1  [ 0, 0, 0 ]; [ 1/2, 1/2, 0 ]

⇒ Q  draw

LC (Q01+Q02)/sqrt(2) draw linear combination of symmetry-adapted orbitals

modulation Q,G  draw modulation of symmetry-adapted orbitals

HOPPING: draw hopping direction.  
 1. input representative BOND, + ENTER.

draw hopping direction

[ 0, 0, 0 ]; [ 1/2, 1/2, 0 ]

# QtDraw+ - Crystal Structure Drawing

## Example Draw "graphene"

1. Choose space group

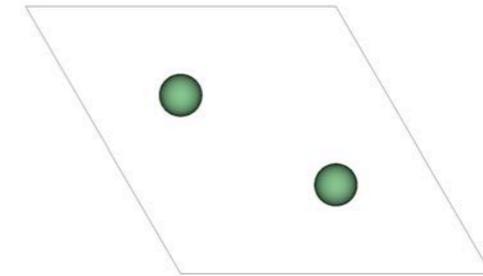
space group    hexagonal    191. D6h<sup>1</sup> (P6/mmm)

2. Input site

object drawing    basis drawing

SITE: draw equivalent sites.  
1. input representative SITE, 2. ENTER.

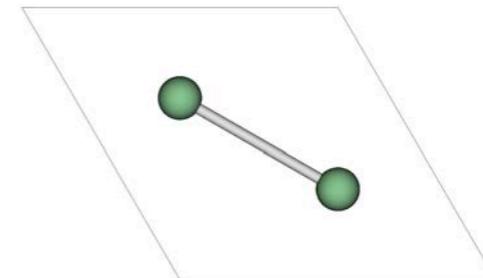
[ 1/3, 2/3, 0 ]



3. Input bond

BOND: draw equivalent bonds.  
1. input representative BOND, 2. ENTER.

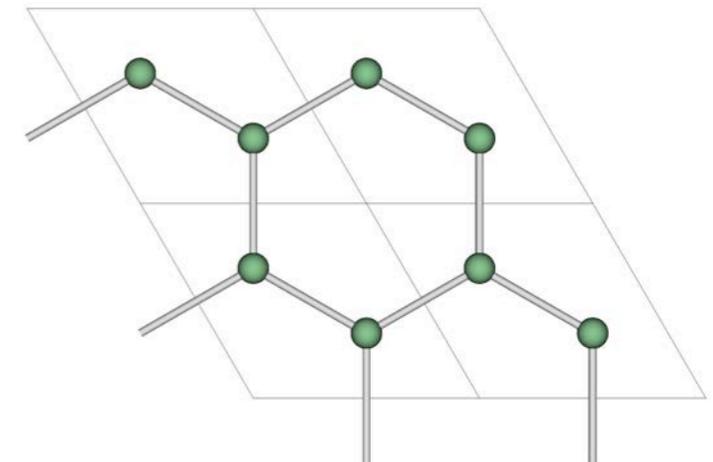
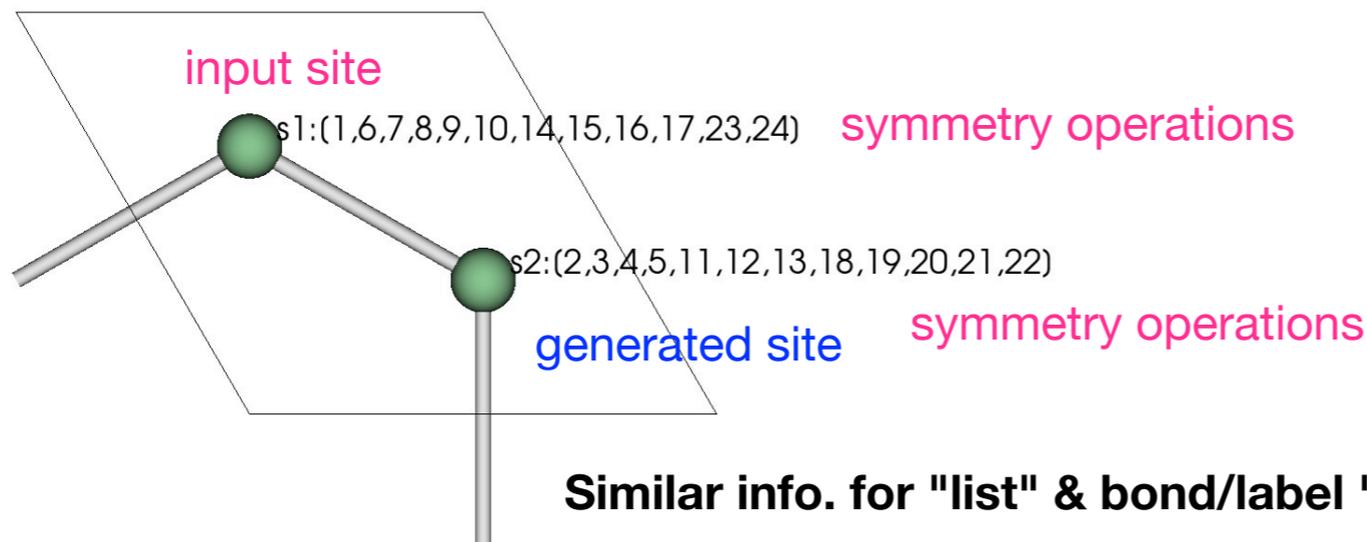
[ 2/3, 1/3, 0 ]; [ 1/3, 2/3, 0 ]



4. Push "clip" and "repeat"



5. Push "repeat" and "list" & site/label "on"



# QtDraw+ - SAMB Drawing

## Draw SAMB for "graphene"

### 1. Draw "mass term"

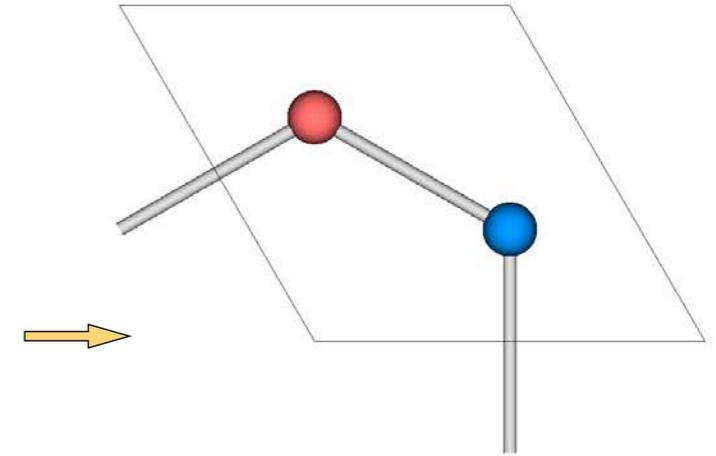
object drawing   basis drawing   **draw weight of sites**

SITE: draw site-cluster basis.  
 1. input representative SITE, + ENTER,  
 ⇒ 2. choose basis, 3. push `draw`.

[ 1/3, 2/3, 0 ]

⇒

Z MP basis   atomic x site-cluster



### 2. Draw "Rashba SOC"

HOPPING: draw hopping direction.  
 1. input representative BOND, + ENTER.   **draw hopping direction**

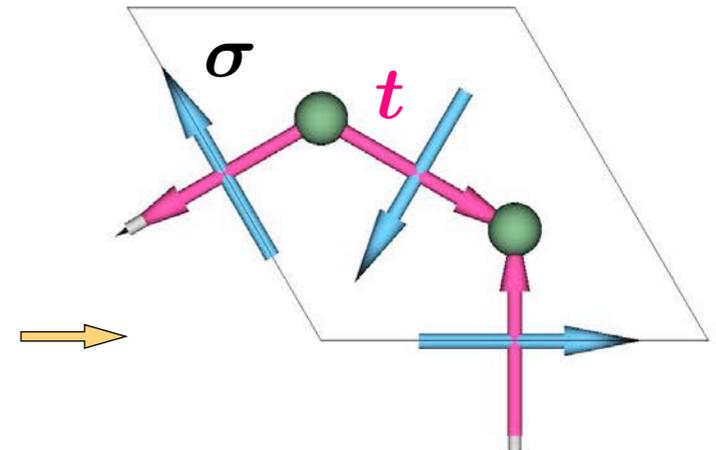
[ 1/3, 2/3, 0 ] ; [ 2/3, 1/3, 0 ]

VECTOR: draw symmetry-adapted vector.  
 1. choose type, 2. input representative SITE/BOND, + ENTER,   **draw spin direction**  
 ⇒ 3. choose (type,basis), 4. push `draw` or 3. input linear combination (LC), + ENTER or 3. push `modulation`.

M   [ 1/3, 2/3, 0 ] ; [ 2/3, 1/3, 0 ]

⇒

Z MP basis   atomic x bond-cluster



### 3. Draw "cluster quadrupole order"

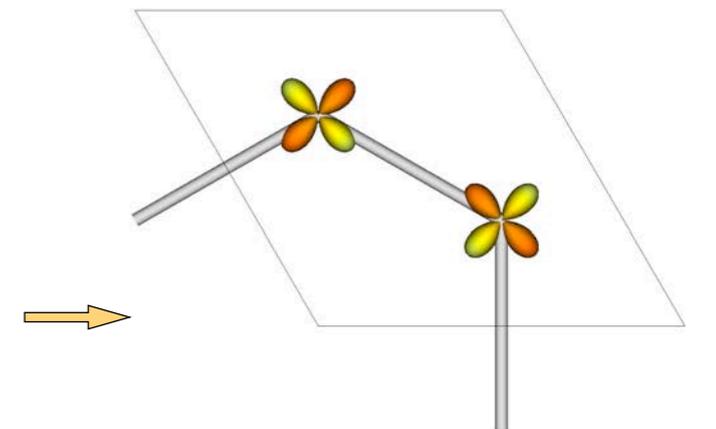
ORBITAL draw symmetry-adapted orbital.   **draw orbitals**

1. choose (type,rank), 2. input representative SITE/BOND, + ENTER,  
 ⇒ 3. choose (type,basis), 4. push `draw` or 3. input linear combination (LC), + ENTER or 3. push `modulation`.

Q   2   [ 1/3, 2/3, 0 ]

⇒

Electric  $E_{1u}$  order



# QtDraw+ - Modulation Drawing

## SAMB modulation

$$\{ \mathbf{X}_\eta(\mathbf{p}_s; \mathbf{R}) \} = \sum_i c_i \mathbb{X}_i \cos[\mathbf{k}_i \cdot (\mathbf{R} + \mathbf{p}_s) - \frac{\pi}{2} n_i]$$

coeff
k
cell plus set
phase

n = 0 : cos

n = 1 : sin

for given [[X, c, k, n]]

## Draw SAMB modulation for "graphene"

### 1. Simple AFM (single-k)

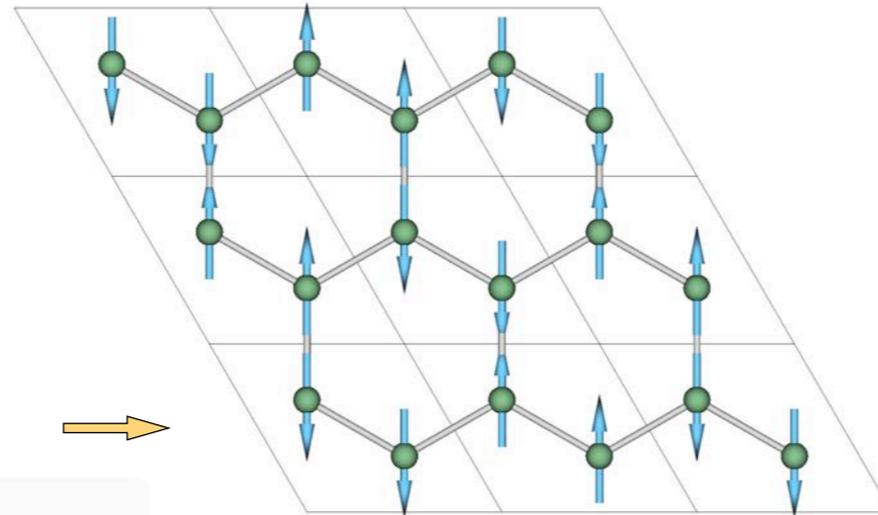
### 2. Vortex-like AFM (triple-k)

(a) First, draw sites & bonds, and repeat in range [0,0,0]-[3,3,1]

(b) Construct vector SAMB

VECTOR: draw symmetry-adapted vector.  
 1. choose type, 2. input representative SIT  
 ⇒ 3. choose (type,basis), 4. push 'draw' o

M [ 1/3, 2/3, 0 ]



(c) Open modulation panel, and edit

modulation T,M

Modulation - vector

basis	coeff	k	phase
M02	7/10	(1/2, 1/2, 0)	cos

add remove

lower [0,0,0] repeat [3,3,1]

Apply Cancel OK

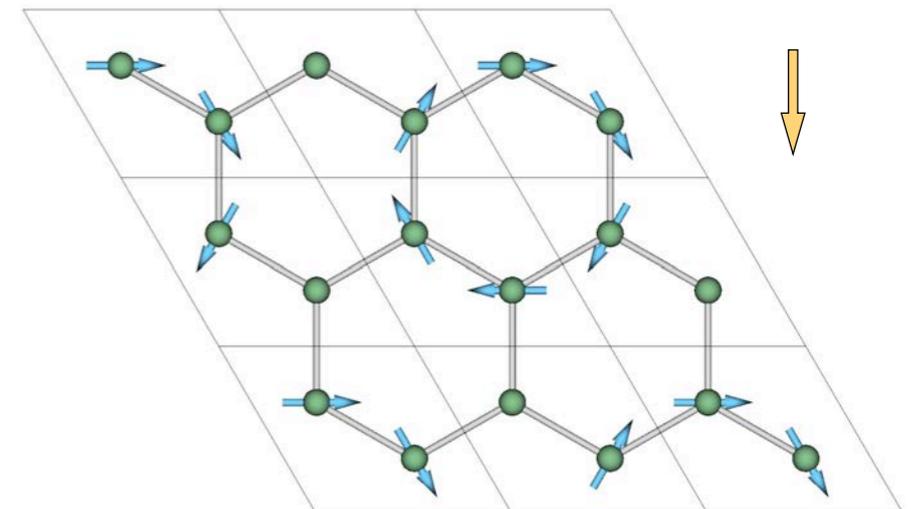
Modulation - vector

basis	coeff	k	phase
M03	√3/6	(0, 1/2, 0)	cos
M02	1/4	(1/2, 0, 0)	cos
M03	√3/12	(1/2, 0, 0)	cos
M04	√3/12	(1/2, 1/2, 0)	cos
M05	1/4	(1/2, 1/2, 0)	cos

add remove

lower [0, 0, 0] repeat [3, 3, 1]

Apply Cancel OK



# QtDraw+ - SAMB Name Ambiguity

## Caveat

Sometime, MP base belonging to the same irreps. but with different rank cannot be distinguished !  
Then, the displayed rank and type of MP differs for essentially the same basis.

## Example

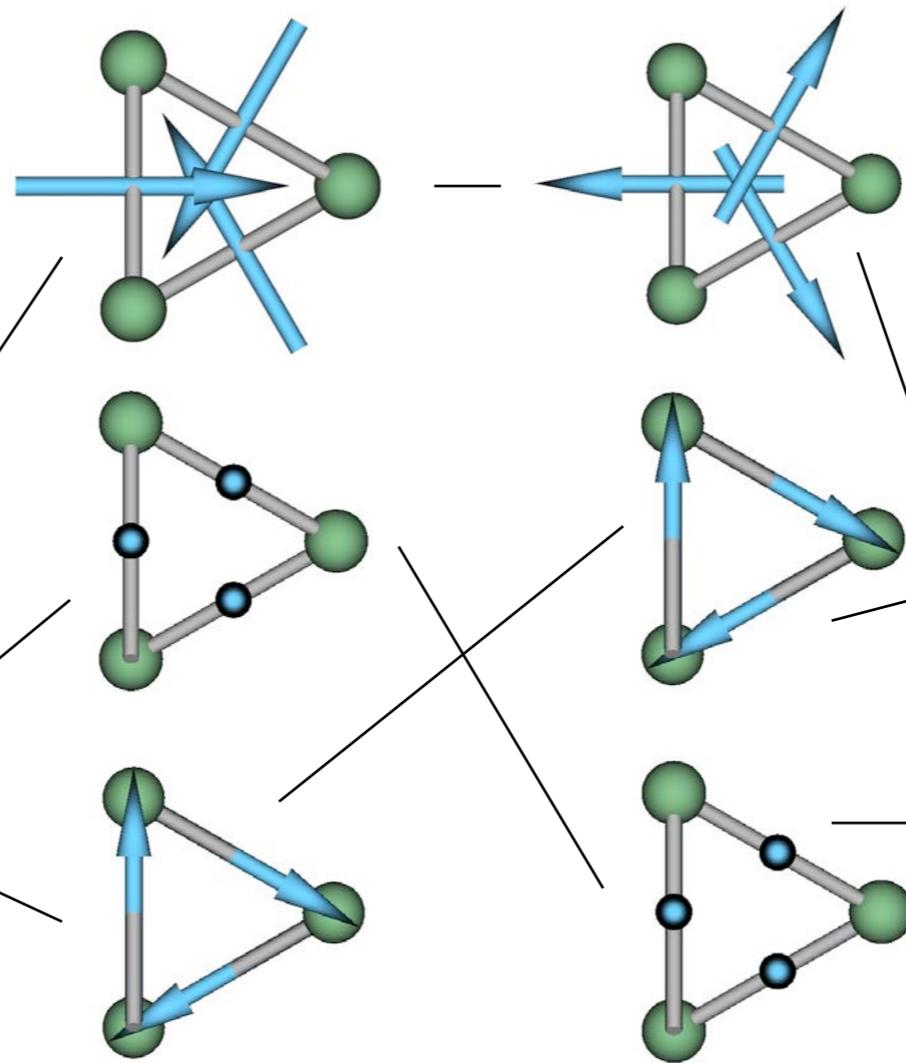
space group  152. D3<sup>4</sup>

M-vector on [0.3, 0, 1/3]; [0, 0.3, 2/3]

or

[-0.3, -0.3, 0]; [0.3, 0, 1/3]

Q(1,A2,,) = Ma(1,E,) x Tb(1,E,)
Q(1,E,,0) = Ma(1,A2,) x Tb(1,E,)
Q(1,E,,1) = Ma(1,A2,) x Tb(1,E,)
G(0,A1,,) = Ma(1,A2,) x Tb(1,A2,)
G(2,A1,,) = Ma(1,A2,) x Tb(1,A2,)
G(2,E,1,0) = Ma(1,A2,) x Tb(1,E,)
G(2,E,1,1) = Ma(1,A2,) x Tb(1,E,)
G(2,E,2,0) = Ma(1,E,) x Tb(1,E,)
G(2,E,2,1) = Ma(1,E,) x Tb(1,E,)
T(2,A1,,) = Ma(1,E,) x Qb(2,E,1)
T(2,E,2,0) = Ma(1,E,) x Qb(2,E,1)
T(2,E,2,1) = Ma(1,E,) x Qb(2,E,1)
M(1,A2,,) = Ma(1,A2,) x Qb(0,A1,)
M(1,A2,,) = Ma(1,E,) x Qb(2,E,1)
M(1,E,,0) = Ma(1,E,) x Qb(0,A1,)
M(1,E,,0) = Ma(1,A2,) x Qb(2,E,1)
M(1,E,,1) = Ma(1,E,) x Qb(0,A1,)
M(1,E,,1) = Ma(1,A2,) x Qb(2,E,1)



Q(1,A2,,) = Ma(1,E,) x Tb(1,E,)
Q(1,E,,0) = Ma(1,A2,) x Tb(1,E,)
Q(1,E,,1) = Ma(1,A2,) x Tb(1,E,)
G(0,A1,,) = Ma(1,A2,) x Tb(1,A2,)
G(2,A1,,) = Ma(1,A2,) x Tb(1,A2,)
G(2,E,1,0) = Ma(1,A2,) x Tb(1,E,)
G(2,E,1,1) = Ma(1,A2,) x Tb(1,E,)
G(2,E,2,0) = Ma(1,E,) x Tb(1,E,)
G(2,E,2,1) = Ma(1,E,) x Tb(1,E,)
T(1,A2,,) = Ma(1,E,) x Qb(1,E,)
T(1,E,,0) = Ma(1,A2,) x Qb(1,E,)
T(1,E,,1) = Ma(1,A2,) x Qb(1,E,)
M(0,A1,,) = Ma(1,E,) x Qb(1,E,)
M(1,A2,,) = Ma(1,A2,) x Qb(0,A1,)
M(1,E,,0) = Ma(1,E,) x Qb(0,A1,)
M(1,E,,1) = Ma(1,E,) x Qb(0,A1,)
M(2,E,2,0) = Ma(1,E,) x Qb(1,E,)
M(2,E,2,1) = Ma(1,E,) x Qb(1,E,)

# Application - SAMB Decomposition

## Local Susceptibility

space group  152. D3<sup>4</sup> cf. Te

$$\chi_i^{\alpha\beta} = a_0 Q_0 + \sum_{\alpha} b_{\alpha} Q_{2,\alpha}$$

electric symmetric tensor = site-cluster mono/quadrupole

View    
 lower:   
 upper:

Site : [0.3, 0, 1/3]

Bond : [0.3, 0, 1/3]; [0, 0.3, 2/3]

ORBITAL draw symmetry-adapted orbital.

1. choose (type,rank), 2. input representative SITE/BOND, + ENTER,

⇒ 3. choose (type,basis), 4. push `draw` or 3. input linear combination (LC), + ENTER or 3. push `modulation`.

Q  [0.3, 0, 1/3]

Q(1,A2,,) = Qa(2,E,1) x Qs(1,E,)

Q(1,E,,0) = Qa(2,A1,) x Qs(1,E,)

Q(1,E,,1) = Qa(2,A1,) x Qs(1,E,)

**1** Q(2,A1,,) = Qa(2,A1,) x Qs(0,A1,)

Q(2,E,1,0) = Qa(2,E,1) x Qs(0,A1,)

Q(2,E,1,1) = Qa(2,E,1) x Qs(0,A1,)

Q(2,E,2,0) = Qa(2,E,2) x Qs(0,A1,)

Q(2,E,2,1) = Qa(2,E,2) x Qs(0,A1,)

**2** Q(3,A1,,) = Qa(2,E,2) x Qs(1,E,)

Q(3,A2,2,) = Qa(2,E,2) x Qs(1,E,)

**3** G(2,A1,,) = Qa(2,E,1) x Qs(1,E,)

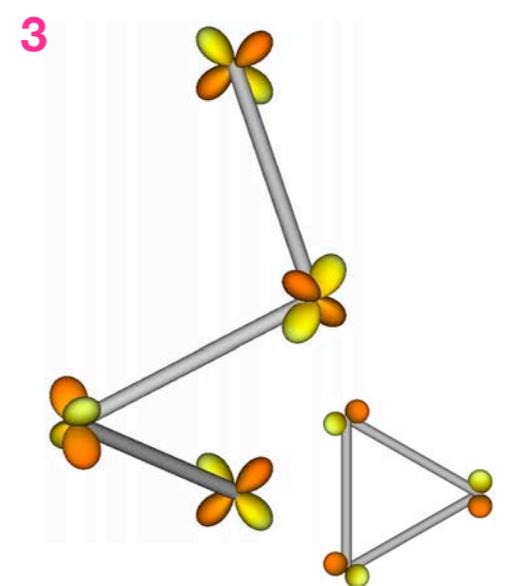
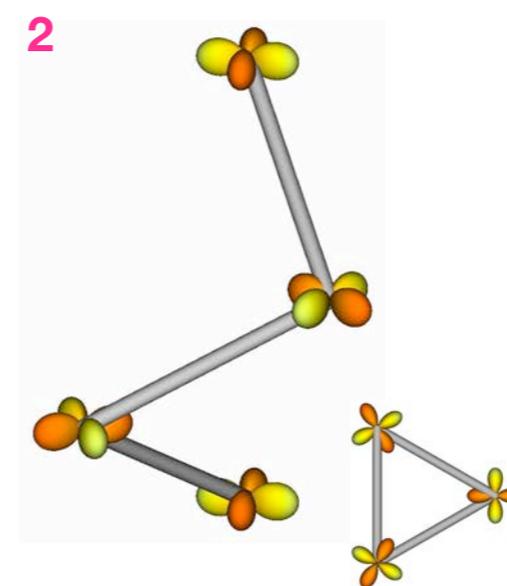
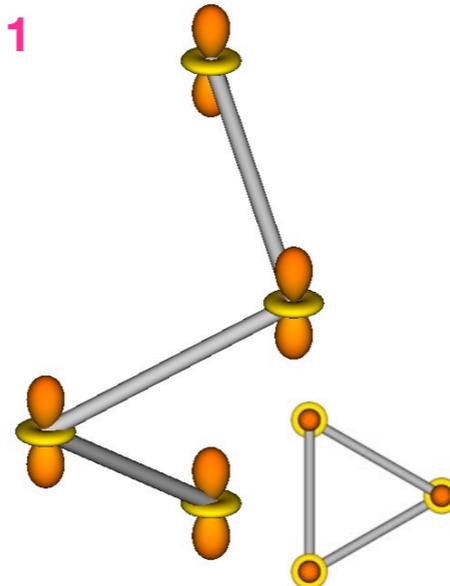
G(2,E,1,0) = Qa(2,A1,) x Qs(1,E,)

G(2,E,1,1) = Qa(2,A1,) x Qs(1,E,)

G(2,E,2,0) = Qa(2,E,1) x Qs(1,E,)

G(2,E,2,1) = Qa(2,E,1) x Qs(1,E,)

1 (monopole) + 3 identity irrep. (A<sub>1</sub>)



# Application - SAMB Decomposition

## Stiffness of Phonon Te

$$V = \frac{1}{2} g_{ij}^{\alpha\beta} (x_i^\alpha - x_{0i}^\alpha)(x_j^\beta - x_{0j}^\beta)$$

$$g_{ij}^{\alpha\beta} = a_0 Q_0 + \sum_{\alpha} b_{\alpha} Q_{2,\alpha}$$

electric symmetric tensor = bond-cluster mono/quadrupole

ORBITAL draw symmetry-adapted orbital.

1. choose (type,rank), 2. input representative SITE/BOND, + ENTER,

⇒ 3. choose (type,basis), 4. push `draw` or 3. input linear combination (LC), + ENTER or 3. push `modulation`.

Q 2 [0.3, 0, 1/3]; [0, 0.3, 2/3]

**1**  $Q(0,A1,,) = Qa(2,E,1) \times Qb(2,E,1)$

**2**  $Q(2,A1,,) = Qa(2,A1,) \times Qb(0,A1,)$

$Q(2,E,1,0) = Qa(2,E,1) \times Qb(0,A1,)$

$Q(2,E,1,0) = Qa(2,A1,) \times Qb(2,E,1)$

$Q(2,E,1,1) = Qa(2,E,1) \times Qb(0,A1,)$

$Q(2,E,1,1) = Qa(2,A1,) \times Qb(2,E,1)$

$Q(2,E,2,0) = Qa(2,E,2) \times Qb(0,A1,)$

$Q(2,E,2,0) = Qa(2,E,1) \times Qb(2,E,1)$

$Q(2,E,2,1) = Qa(2,E,2) \times Qb(0,A1,)$

$Q(2,E,2,1) = Qa(2,E,1) \times Qb(2,E,1)$

$G(1,A2,,) = Qa(2,E,1) \times Qb(2,E,1)$

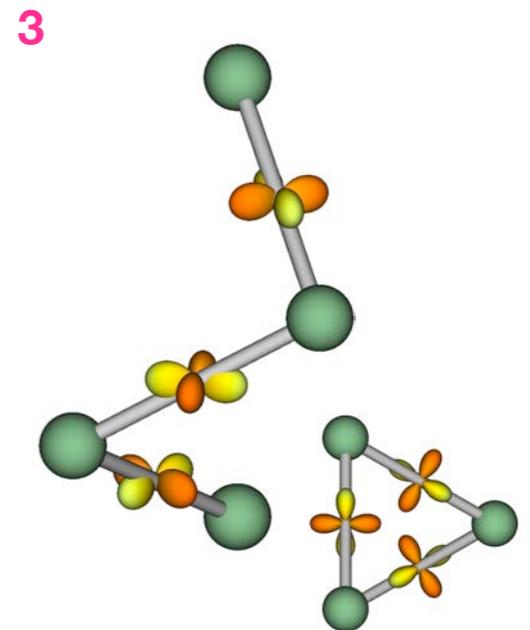
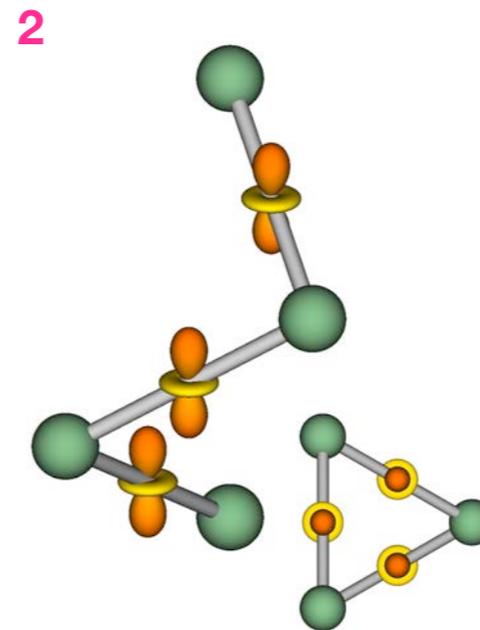
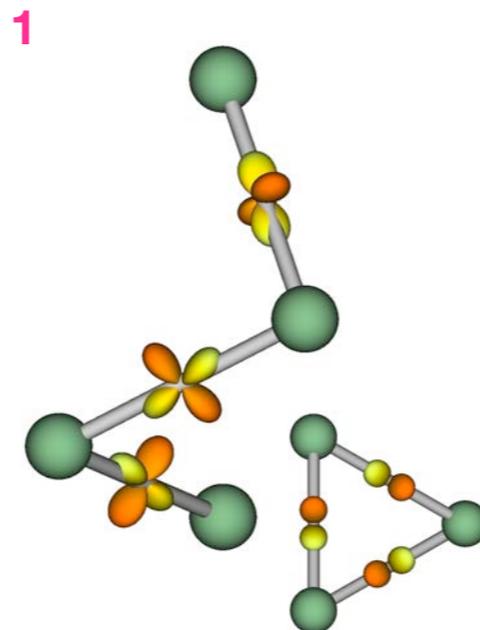
$G(1,E,,0) = Qa(2,A1,) \times Qb(2,E,1)$

$G(1,E,,1) = Qa(2,A1,) \times Qb(2,E,1)$

**3**  $G(3,A1,,) = Qa(2,E,2) \times Qb(2,E,1)$

$G(3,A2,2,) = Qa(2,E,2) \times Qb(2,E,1)$

1 (monopole) + 3 identity irrep. (A<sub>1</sub>)



# Application - SAMB Decomposition

## Other Physical Quantities

Type	Expression	Correspondence	SAMB
Electric potential	$\phi q$	$q \rightarrow Q_{0,0}^{(a)}$	(E) Atomic (s) & Site-cluster
Crystal field	$\phi_{lm} Q_{lm}$	$Q_{lm} \rightarrow Q_{lm}^{(a)}$	(E) Atomic & Site-cluster
Zeeman term	$-h^a m^a$	$m^a \rightarrow M_{1m}^{(a)}$	(M) Atomic (spinful) & Site-cluster
Spin-orbit int.	$\zeta l^a \sigma^a$	$l^a, \sigma^a \rightarrow M_{1m}^{(a)}$	(E) Atomic (p-spinful) & Site-cluster
Density-density int.	$V_{ij} n_i n_j$	$n_i n_j \rightarrow Q_{0,0}^{(a)}$	(E) Atomic (s) & Bond-cluster
Elastic energy	$\epsilon_{ij}^{ab} u_i^a u_j^b$	$u_i^a u_j^b \rightarrow Q_{0,0}^{(a)}, Q_{2m}^{(a)}$	(E) Atomic (p) & Bond-cluster
Exchange int.	$J_{ij}^{ab} S_i^a S_j^b$	$S_i^a S_j^b \rightarrow Q_{0,0}^{(a)}, Q_{2m}^{(a)}$	(E) Atomic (p) & Bond-cluster
DM int.	$D_{ij}^c \epsilon_{abc} S_i^a S_j^b$	$\epsilon_{abc} S_i^a S_j^b \rightarrow G_{lm}^{(a)}$	(ET) Atomic (p) & Bond-cluster
Real hopping	$t_{ij} c_i^\dagger c_j + \text{H.c.}$	$c_i^\dagger c_j + \text{H.c.} \rightarrow Q_{lm}^{(b)}$	(all) Atomic & Bond-cluster
Imaginary hopping	$i t_{ij} c_i^\dagger c_j + \text{H.c.}$	$i c_i^\dagger c_j + \text{H.c.} \rightarrow T_{lm}^{(b)}$	(all) Atomic & Bond-cluster

Various quantities can be expressed by using SAMB