

# Libra/CP2K Interface

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## CP2K

- CP2K is a quantum chemistry and solid state physics software package that can perform atomistic simulations of solid state, liquid, molecular, periodic, material, crystal, and biological systems.
- CP2K provides a general framework for different modeling methods such as DFT using the mixed Gaussian and plane waves approaches GPW and GAPW.
- Supported theory levels include DFTB, LDA, GGA, MP2, RPA, semi-empirical methods (AM1, PM3, PM6, RM1, MNDO, ...), and classical force fields (AMBER, CHARMM, ...).
- CP2K can do simulations of molecular dynamics, metadynamics, Monte Carlo, Ehrenfest dynamics, vibrational analysis, core level spectroscopy, energy minimization, and transition state optimization using NEB or dimer method.

# Theory

- A molecular orbital (MO) is defined as a linear combination of atomic orbitals:

$$|\psi_n\rangle = \sum_i c_i |\phi_i\rangle$$

- The MO overlap:

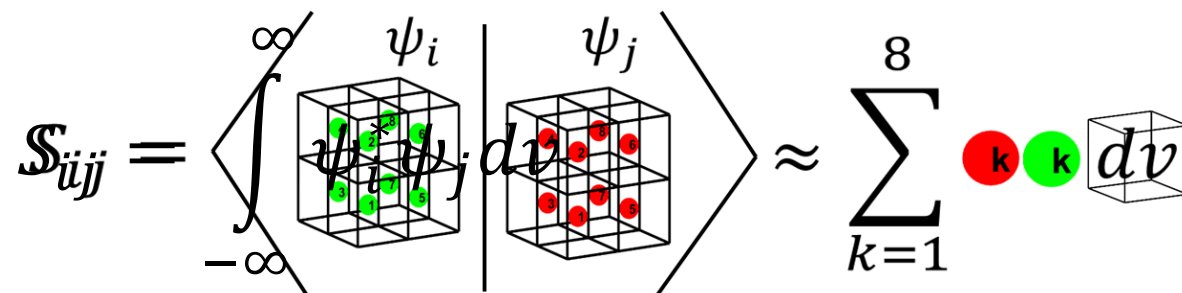
$$S_{nm} = \langle \psi_n | \psi_m \rangle = \sum_{i,j} c_i^* c_j \langle \phi_i | \phi_j \rangle$$

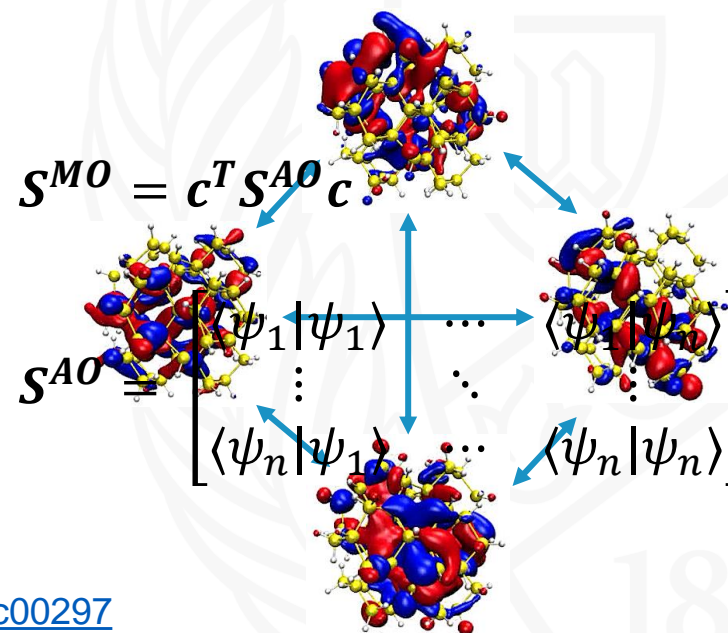
the  $\langle \phi_i | \phi_j \rangle$  is the atomic orbital overlap. So the MO overlap matrix ( $\mathbf{S}^{MO}$ ) can be written as follows where  $\mathbf{c}$  is the matrix of molecular orbital coefficients and  $\mathbf{S}^{AO}$  is the atomic orbital overlap matrix:

$$\mathbf{S}^{MO} = \mathbf{c}^T \mathbf{S}^{AO} \mathbf{c}$$

# Molecular orbital integrals

- Grid-based approach using *.cube* files
  - Easy to implement
  - Most codes can output these file
  - Not suitable for large structures with large number of states
- Analytical approach
  - Suitable for large systems and large number of states
  - One interface cannot directly be used for many codes

$$S_{ij} = \int_{-\infty}^{\infty} \psi_i \psi_j d\tau \approx \sum_{k=1}^8 \psi_i(\mathbf{k}) \psi_j(\mathbf{k}) d\tau$$




# Gaussian type orbitals (GTO)

- Atom centered basis sets

$$\varphi(r - R; n, \xi) = N(x - R_x)^{n_x}(y - R_y)^{n_y}(z - R_z)^{n_z} \times \exp(-\xi(r - R)^2)$$

$$N = \left(\frac{2\xi}{\pi}\right)^{\frac{3}{4}} (4\xi)^{(n_x+n_y+n_z)/2} \times \left((2n_x - 1)!! (2n_y - 1)!! (2n_z - 1)!!\right)^{-\frac{1}{2}}$$

$$\Phi_i(\vec{r}) = \varphi(r; n_i, \xi) \cdot Y_{n_i, m_i}(\theta, \phi)$$

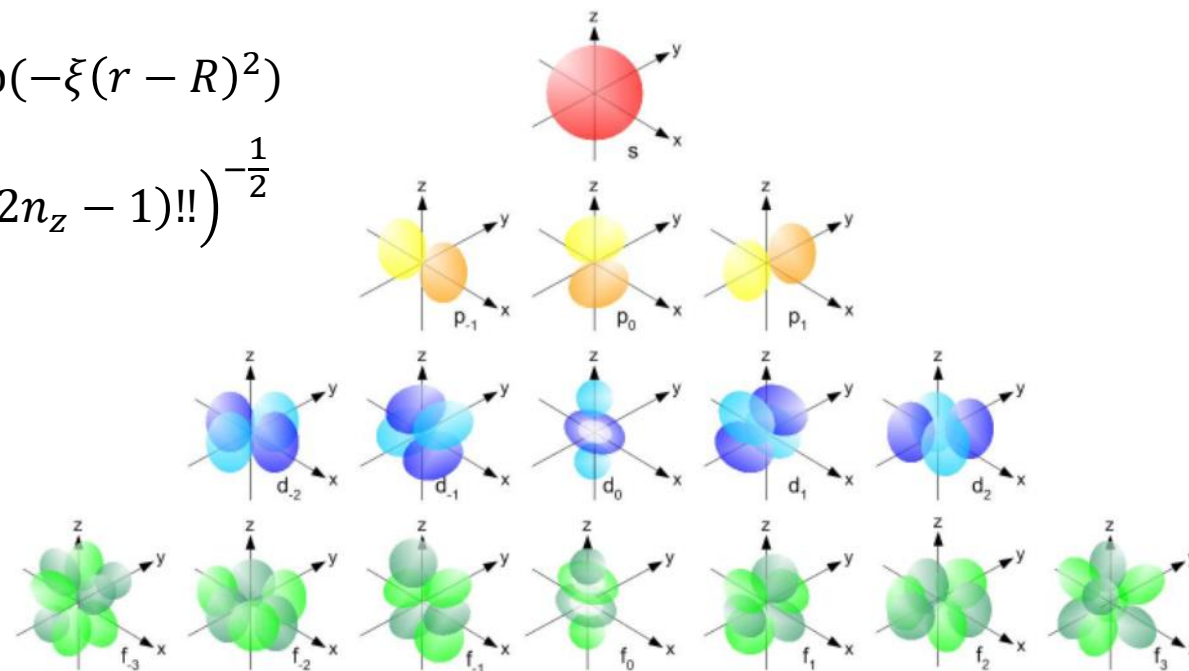
$n$  shows the angular momentum value:

s-orbital  $n = 0$

p-orbital  $n = 1$

d-orbital  $n = 2$

f-orbital  $n = 3$



# Recurrence relations between GTOs

$$\langle s|s \rangle = \left(\frac{\pi}{\xi}\right)^{\frac{3}{2}} \exp(-\xi(A-B)^2)$$

$$\langle p_i|s \rangle = (P_i - A_i)\langle s|s \rangle$$

$$\langle p_i|p_j \rangle = (P_j - B_j)\langle p_i|s \rangle + \frac{\delta_{ij}}{2\xi}\langle s|s \rangle$$

$$\langle d_{ij}|s \rangle = (P_j - A_j)\langle p_i|s \rangle + \frac{\delta_{ij}}{2\xi}\langle s|s \rangle$$

$$\langle d_{ij}|p_k \rangle = (P_k - B_k)\langle d_{ij}|s \rangle + \frac{\delta_{ik}}{2\xi}\langle p_j|s \rangle + \frac{\delta_{jk}}{2\xi}\langle p_i|s \rangle$$

$$\langle d_{ij}|d_{kl} \rangle = (P_l - B_l)\langle d_{ij}|p_k \rangle + \frac{\delta_{il}}{2\xi}\langle p_j|p_k \rangle + \frac{\delta_{jl}}{2\xi}\langle p_i|p_k \rangle + \frac{\delta_{kl}}{2\xi}\langle d_{ij}|s \rangle$$

$i, j, k, l = x, y, z$

Obara, and Saika, J. Chem. Phys. 1986, 84, 3963-3974.

Obara, and Saika, J. Chem. Phys. 1988, 89, 1540-1559.

Libint, Version 2.6.0 Edward F. Valeev, <http://libint.valeev.net>.



Open-source and OpenMP  
parallelized code for computing  
integrals between GTOs

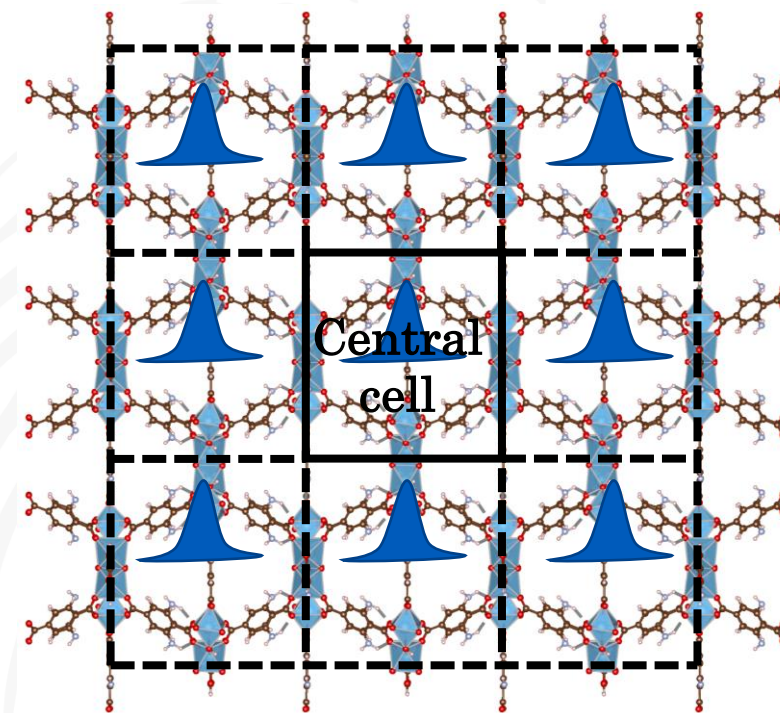
# Periodic basis functions

- The Bloch function for K-point in a periodic structure is defined as:

$$\beta_a^k(r) = \frac{1}{\sqrt{N}} \sum_R \varphi_a(r - R) e^{ikR}$$

- Overlaps between Bloch functions of two different K-points:

$$\begin{aligned}
 S_{a,b}^k &= \langle \beta_a^k | \beta_b^{k'} \rangle = \frac{1}{N} \int dr \sum_{R,R'} e^{-ikR} \varphi_a^*(r - R) e^{ik'R'} \varphi_b(r - R') \\
 &= \frac{1}{N} \int dr \sum_{R,R'} e^{i(kR - k'R')} \varphi_a^*(r - R) \varphi_b(r - R')
 \end{aligned}$$



MIL-125-NH<sub>2</sub>

## How Libra computes the MO overlap?

- Libra uses ‘*molden*’ file formats to read the molecular orbital coefficients, energies, occupation, spin, atomic coordinates, and basis set although it can use other file formats as well including *MOLog* printed out by CP2K.
- Here, we’ll show an example of the inner functions that compute the MO overlap matrix for one geometry. Note that the workflow is not like this and you only need to specify a couple of variables. The following shows how one can work with the Libra functions, for example, if one intends to write an on-the-fly computation of the NACs.



```

import os
import numpy as np
import matplotlib.pyplot as plt
from liblibra_core import *
from libra_py import CP2K_methods, moldern_methods, data_conv, units
    
```

```

# Molden file name
moldern_file_name = 'test.moldern'
# Number of processors
nprocs = 16
# Spherical or Cartesian coordiantes? Spherical!
is_spherical = True
    
```

```

# The integration shells and angular momentum values
shell_1, l_vals =
moldern_methods.moldern_file_to_libint_shell(moldern_file_name,
is_spherical)
    
```

```

# All of the eigenvectors and energies of the system
eig_vect_1, energies_1 =
moldern_methods.eigenvectors_moldern(moldern_file_name, nbasis(shell_1),
l_vals)
    
```

```

# Resorting the moldern indices
new_indices = CP2K_methods.resort_molog_eigenvectors(l_vals)
eigenvectors_1 = []
for j in range(len(eig_vect_1)):
    # the new and sorted eigenvector
    eigenvector_1 = eig_vect_1[j]
    eigenvector_1 = eigenvector_1[new_indices]
    # append it to the eigenvectors list
    eigenvectors_1.append(eigenvector_1)
eigenvectors_1 = np.array(eigenvectors_1)
    
```

```

# Alpha and Beta spin eigenvectors
# alpha -> even indices
alpha_eig_vects = eigenvectors_1[0::2]
alpha_eig_vals = energies_1[0::2]
# beta -> odd indices
beta_eig_vects = eigenvectors_1[1::2]
beta_eig_vals = energies_1[1::2]
# Compute AO overlap matrix
AO_S = compute_overlaps(shell_1, shell_1, nprocs)
# Converting to numpy array
AO_S = data_conv.MATRIX2npararray(AO_S)
    
```

```

print('The shape of the AO matrix...\n', AO_S.shape)
S_alpha = np.linalg.multi_dot([alpha_eig_vects, AO_S,
alpha_eig_vects.T])
S_beta = np.linalg.multi_dot([beta_eig_vects, AO_S,
beta_eig_vects.T])
    
```

# Periodic calculations

```
AO_S = compute_overlaps(shell_1,shell_1,nprocs)
if is_periodic:
    cell = []
    cell.append(params['A_cell_vector'])
    cell.append(params['B_cell_vector'])
    cell.append(params['C_cell_vector'])
    cell = np.array(cell)*units.Angst
    # Generating translational vectors
    translational_vectors = params['translational_vectors']
    for i1 in range(len(translational_vectors)):
        translational_vector = np.array(translational_vectors[i1])
        print(F'Computing the AO overlaps between R({translational_vector[0]},{translational_vector[1]},
            {translational_vector[2]}) and R(0,0,0)')
        shell_1p, l_vals =
        molder_methods.molder_file_to_libint_shell(molder_filename,is_spherical,is_periodic,cell,
            translational_vector)
        AO_S += compute_overlaps(shell_1,shell_1p, nprocs)
```

# What do we need to specify in the input?

```
params['nprocs']
params['mpi_executable']
params['istep']
params['fstep']
params['lowest_orbital']
params['highest_orbital']
params['isxTB']
params['isUKS']
params['is_periodic']
if params['is_periodic']:
    params['A_cell_vector']
    params['B_cell_vector']
    params['C_cell_vector']
    params['periodicity_type'] # example: 'XYZ'
    origin = [0,0,0]
    params['translational_vectors'] =
    CP2K_methods.generate_translational_vectors(
    origin, [2,2,2], params['periodicity_type'])
params['is_spherical']
params['remove_molden']
params['res_dir']
params['all_pdosfiles']
params['all_logfiles']
```

```
params['cp2k_exe']
params['cp2k_ot_input_template'] # just for xTB
params['cp2k_diag_input_template']
params['trajectory_xyz_filename']
```

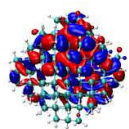
```
# For cube visualization
params['cube_visualization']
params['vmd_input_template']
params['states_to_plot']
params['plot_phase_corrected']
params['vmd_exe']
params['tachyon_exe']
params['x_pixels']
params['y_pixels']
params['image_format']
params['remove_cube']
params['all_images']
```

```
step2.run_cp2k_libint_step2(params)
```

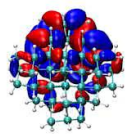
# Molecular orbitals visualization

Highest Occupied Molecular Orbital

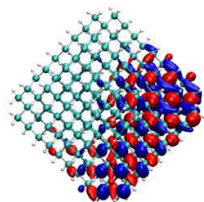
Lowest Unoccupied Molecular Orbital



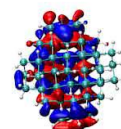
Si<sub>59</sub>H<sub>60</sub>



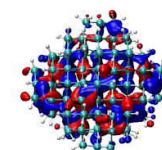
Si<sub>123</sub>H<sub>100</sub>



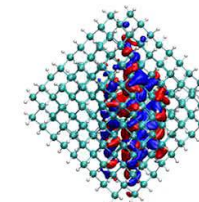
Si<sub>265</sub>H<sub>140</sub>



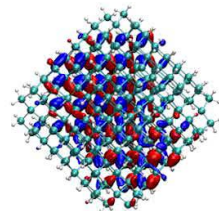
Si<sub>59</sub>H<sub>60</sub>



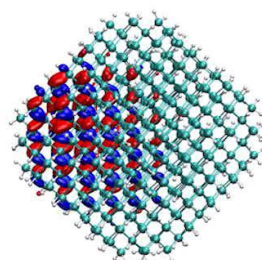
Si<sub>123</sub>H<sub>100</sub>



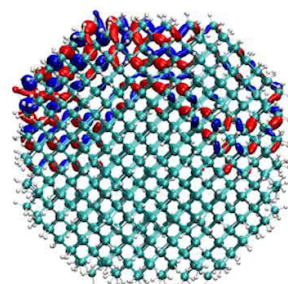
Si<sub>265</sub>H<sub>140</sub>



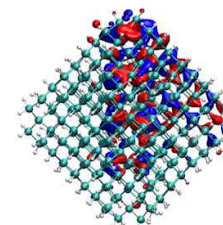
Si<sub>329</sub>H<sub>172</sub>



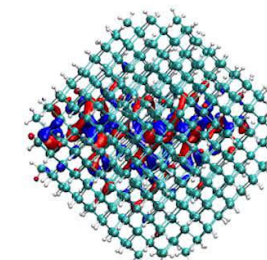
Si<sub>501</sub>H<sub>228</sub>



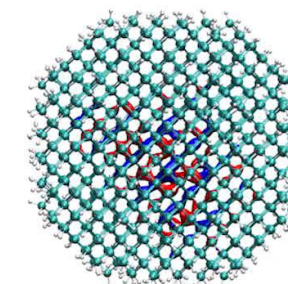
Si<sub>1009</sub>H<sub>412</sub>



Si<sub>329</sub>H<sub>172</sub>



Si<sub>501</sub>H<sub>228</sub>



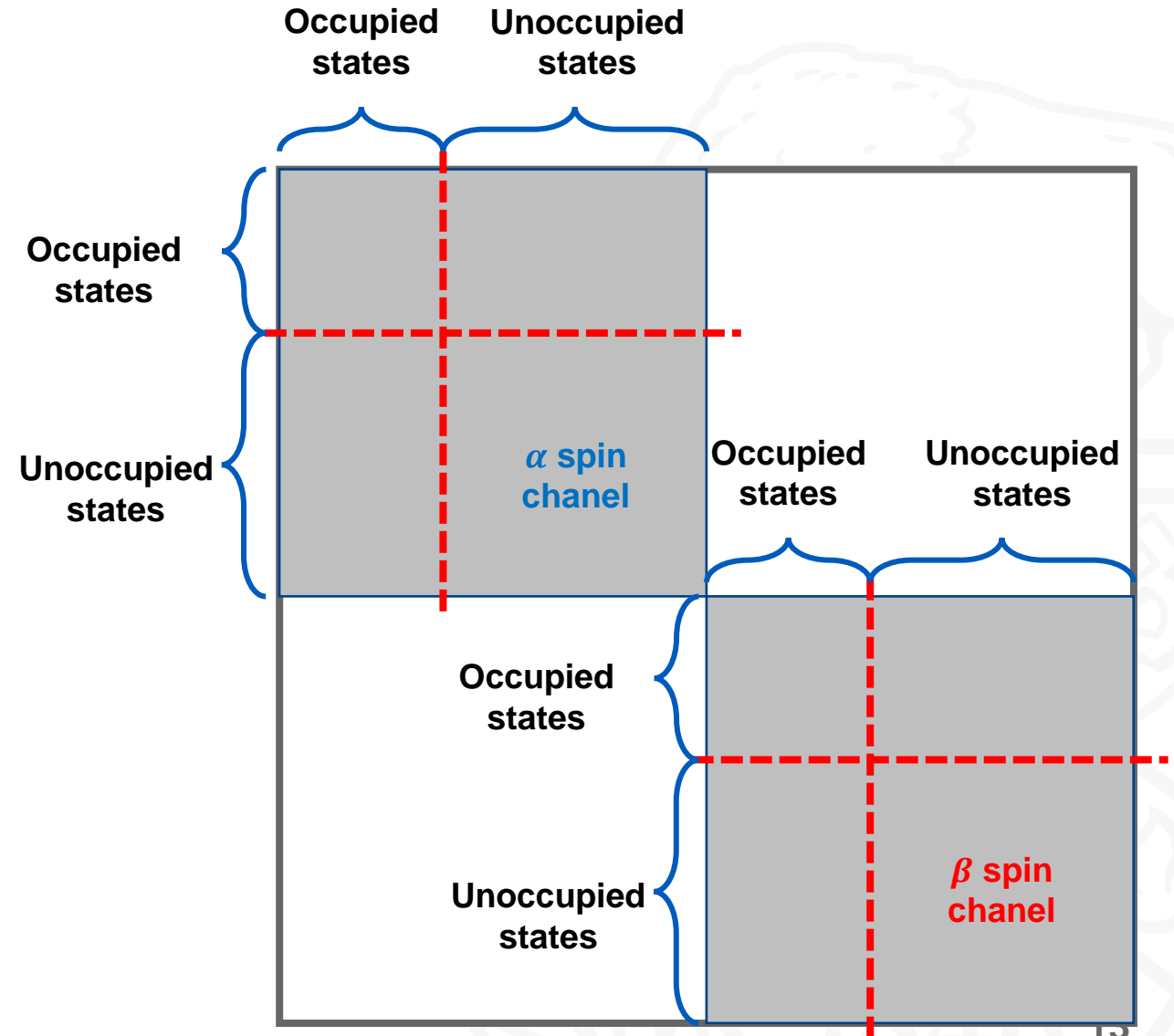
Si<sub>1009</sub>H<sub>412</sub>

0

0

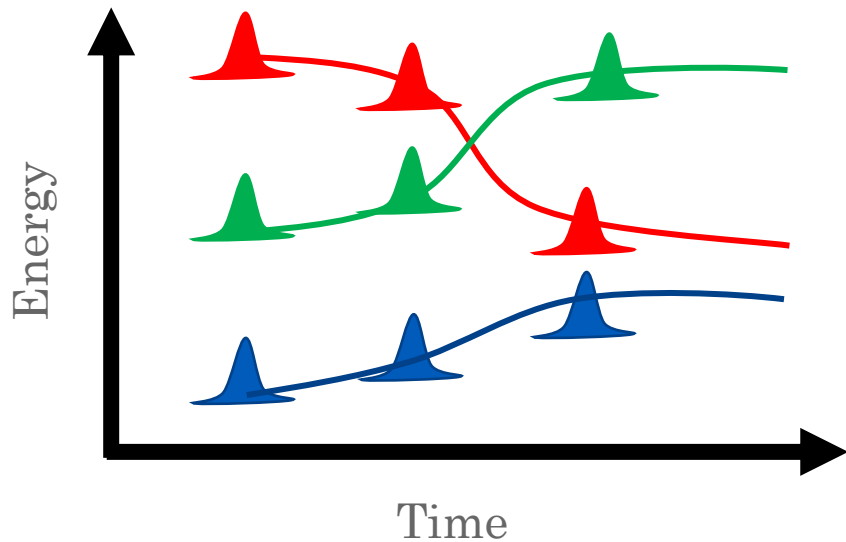
# Storing data

- Libra saves MO overlaps in 2-spinor format
- With no spin-orbit coupling, the second and third block of the matrix is zero.
- To efficiently storing the data, we use `scipy.sparse` library of Python



# Corrections to time-overlap matrix

- State tracking
  - Min-cost algorithm
  - Stochastic state-tracking

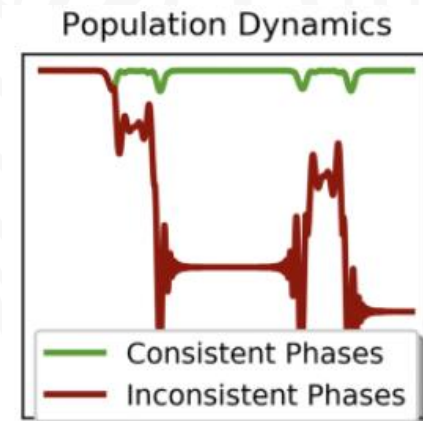


- Phase correction

$$F_i(t + \Delta t) = \frac{\langle \Psi_i(t) | \Psi_i(t + \Delta t) \rangle}{\|\Psi_i(t)\| \|\Psi_i(t + \Delta t)\|}$$

The phase-corrected wave function at time  $t + \Delta t$  is:

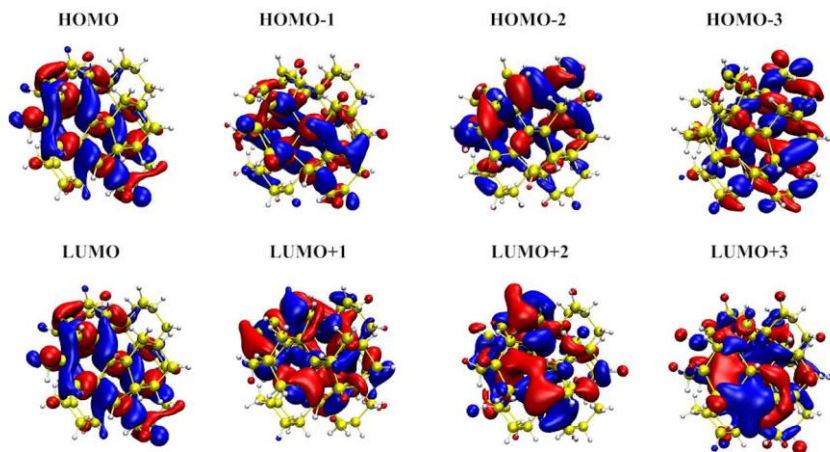
$$\bar{\Psi}_i(t + \Delta t) = F_i^*(t + \Delta t) \Psi_i(t + \Delta t)$$



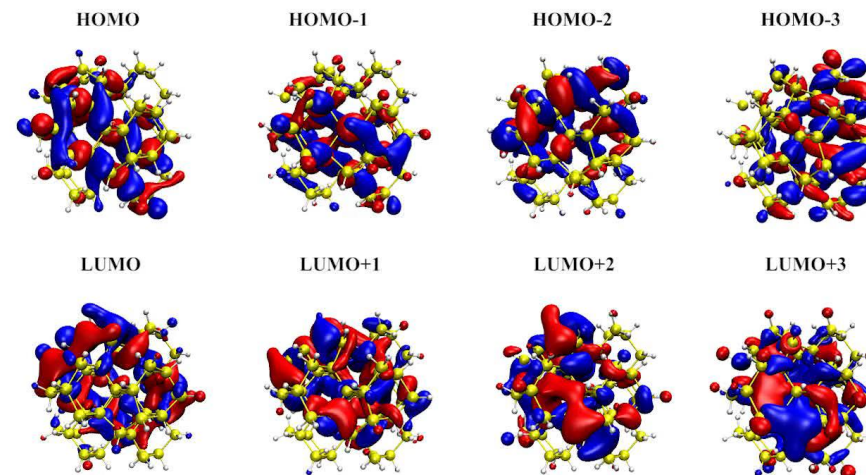
# Corrections to time-overlap matrix

- Phase correction

Phase-uncorrected orbitals



Phase-corrected orbitals

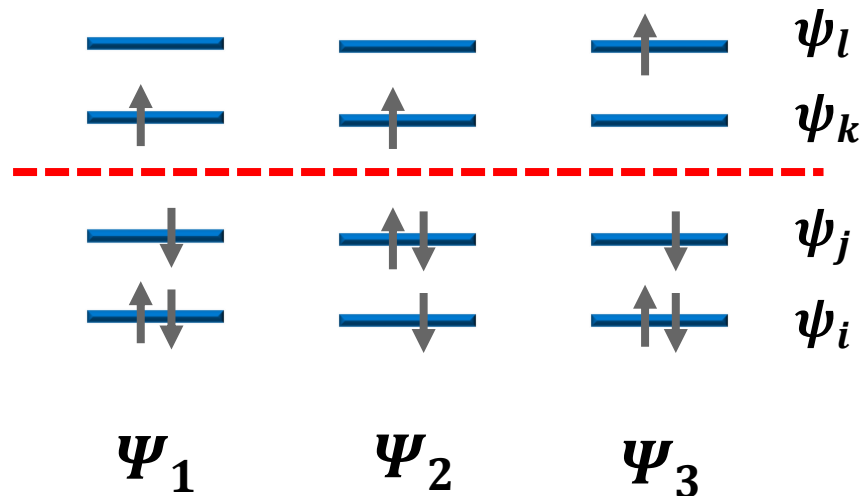


# Excited states basis

- In quantum mechanics, electronic wave function has the antisymmetric property which can be expressed in form of Slater determinant.
- For an N-electron system:

$$\Psi(x_1, \dots, x_N) = \frac{1}{\sqrt{N!}} |\psi_1(x_1)\psi_2(x_2) \dots \psi_K(x_N)\rangle = \frac{1}{\sqrt{N!}} \begin{vmatrix} \psi_1(x_1) & \dots & \psi_K(x_1) \\ \vdots & \ddots & \vdots \\ \psi_1(x_N) & \dots & \psi_K(x_N) \end{vmatrix}$$

- Single-particle excitations

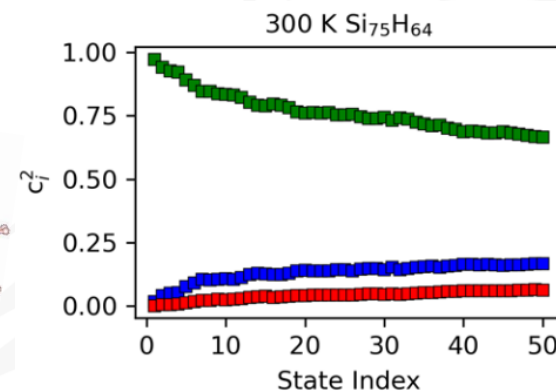
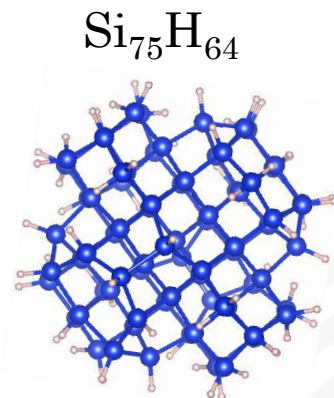
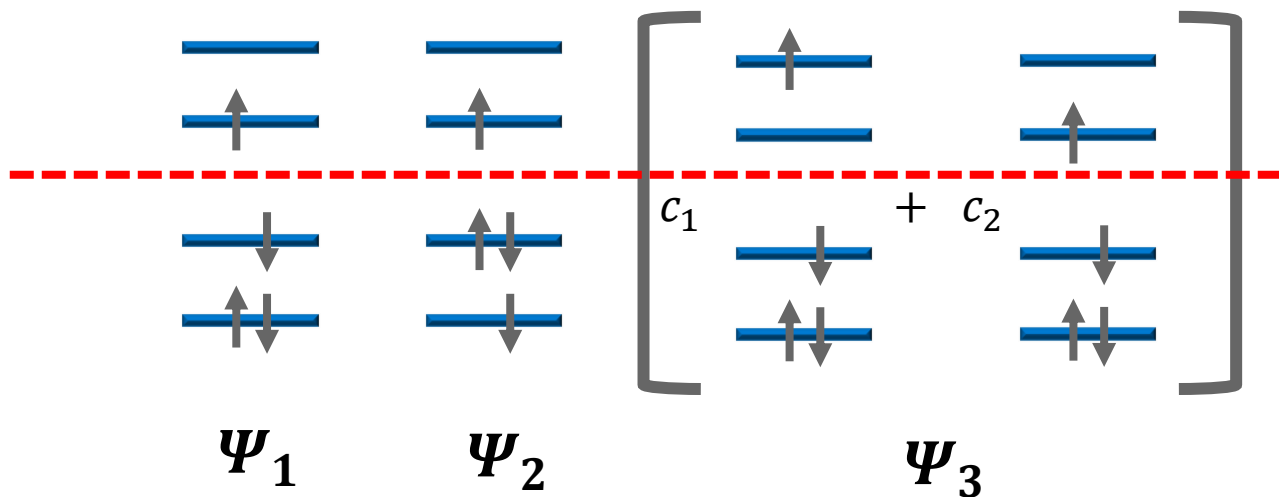


$$\langle \Psi_1 | \Psi_2 \rangle = \det \begin{vmatrix} \langle \psi_l | \psi_l \rangle & \langle \bar{\psi}_l | \psi_l \rangle & \langle \psi_k | \psi_k \rangle & \langle \bar{\psi}_j | \psi_k \rangle \\ \langle \psi_l | \bar{\psi}_l \rangle & \langle \psi_l | \psi_l \rangle & \langle \psi_k | \bar{\psi}_j \rangle & \langle \bar{\psi}_j | \bar{\psi}_l \rangle \\ \langle \psi_k | \psi_l \rangle & \langle \psi_k | \psi_i \rangle & \langle \psi_j | \psi_j \rangle & \langle \bar{\psi}_j | \psi_i \rangle \\ \langle \psi_k | \bar{\psi}_j \rangle & \langle \psi_i | \bar{\psi}_j \rangle & \langle \psi_k | \psi_j \rangle & \langle \bar{\psi}_j | \psi_j \rangle \end{vmatrix}$$

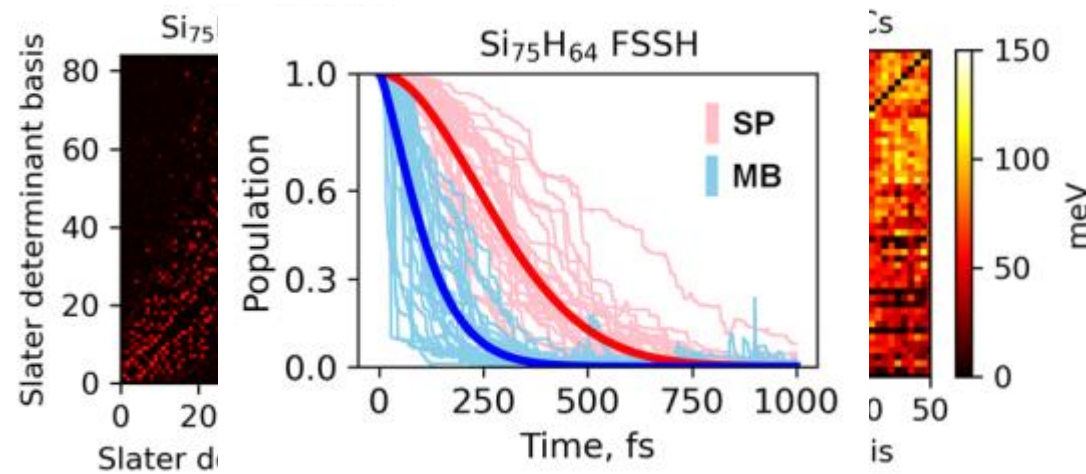


# Excited states basis

- Many-body (TD-DFT) excitation



Excitation energy

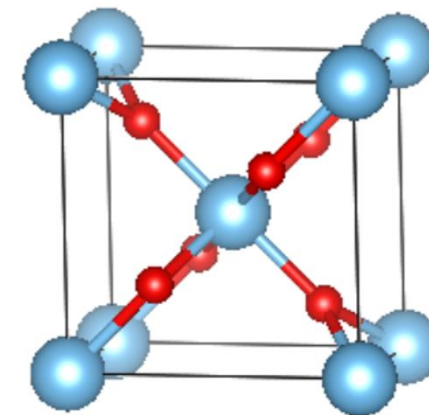
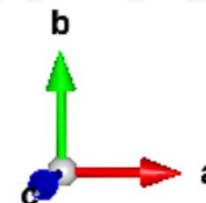


# NAC in KS basis

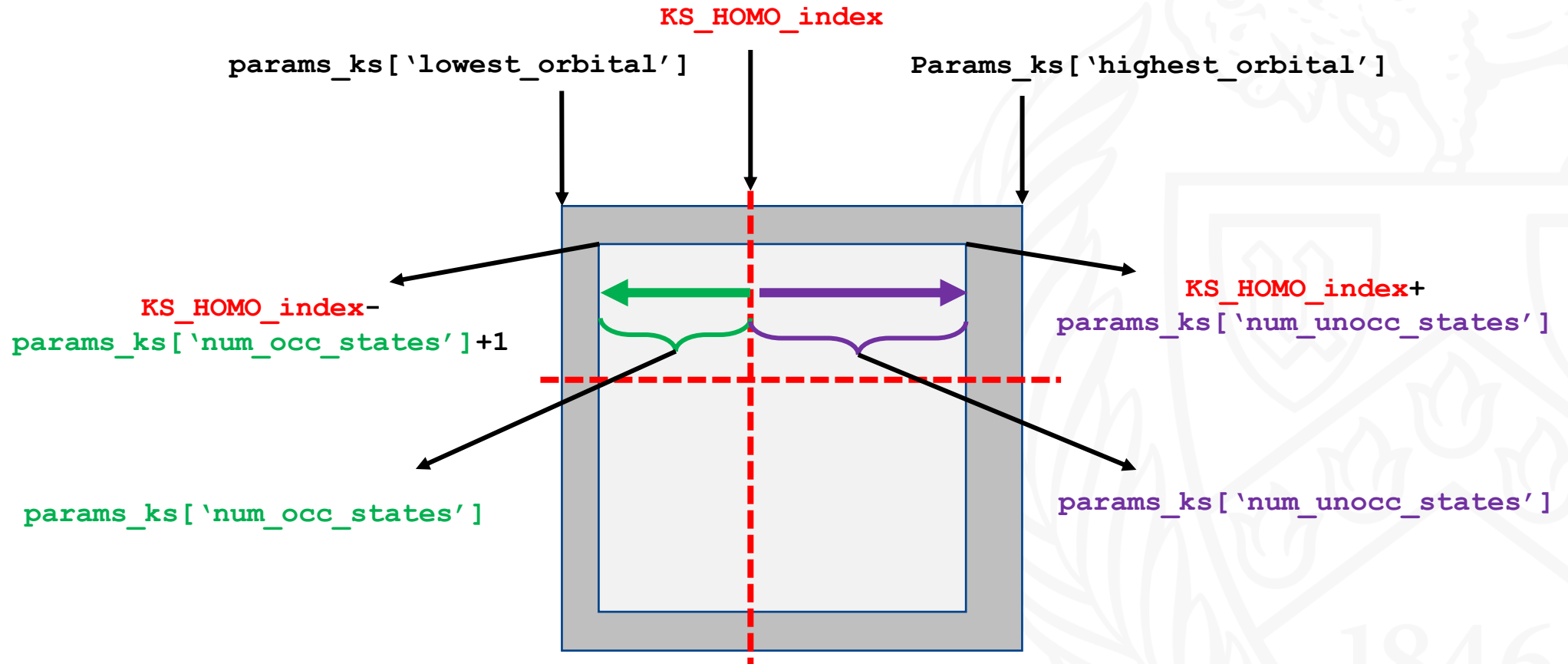
```

params_ks = { 'lowest_orbital': 24-10, 'highest_orbital': 24+11,
'num_occ_states': 10, 'num_unocc_states': 10, 'use_multiprocessing': True,
'nprocs': 8, 'time_step': 1.0, 'es_software': 'cp2k', 'path_to_npz_files':
os.getcwd()+'/res', 'logfile_directory': os.getcwd()+'/all_logfiles',
'path_to_save_ks_Hvibs': os.getcwd()+'/res-ks-DFT', 'start_time': 1200,
'finish_time': 1401, 'apply_phase_correction': True,
'apply_orthonormalization': True, 'do_state_reordering': 2,
'state_reordering_alpha':0, 'nac_algo': 0 }
# For KS states - Applying correction to KS overlaps and computing the
NACs in KS space
step3.run_step3_ks_nacs_libint(params_ks)
    
```

Rutile  $\text{TiO}_2$   
Single Unit Cell



# NAC in KS basis

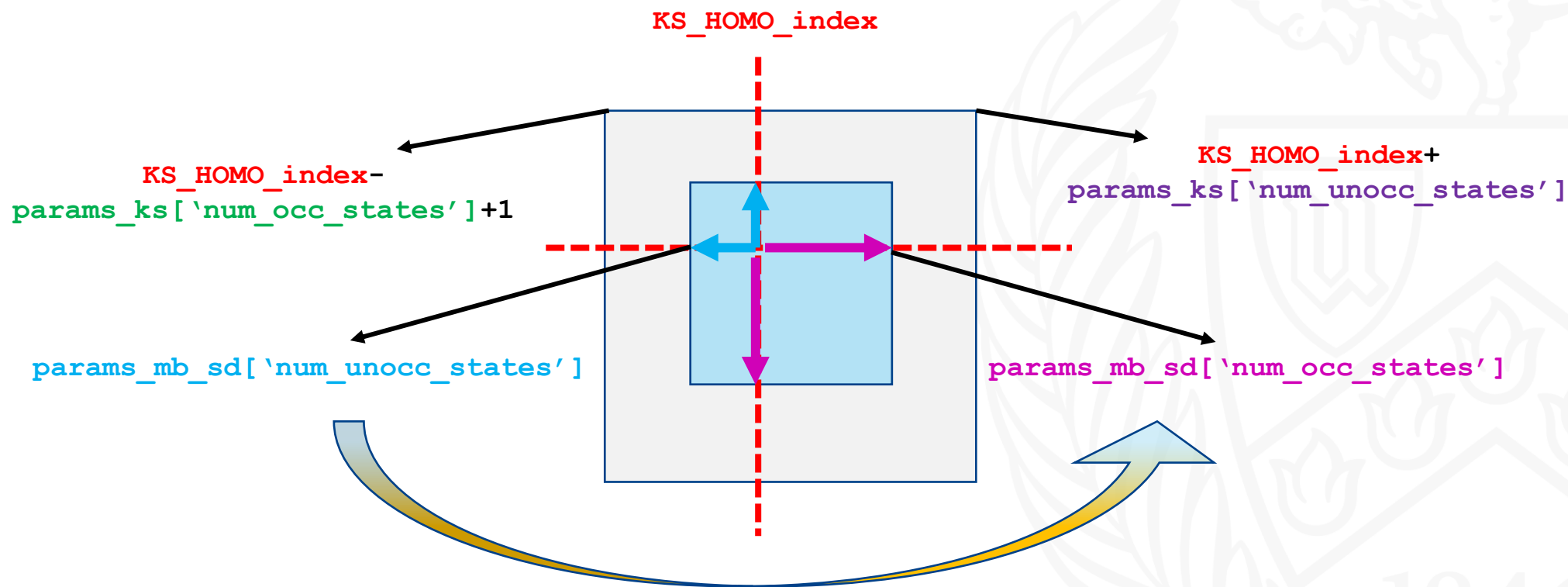


# NAC in excited states basis

```
params_mb_sd = {  
    'lowest_orbital': 24-params_ks['num_occ_states']+1, 'highest_orbital': 24+params_ks['num_unocc_states'],  
    'num_occ_states': 10, 'num_unocc_states': 10,  
    'isUKS': 0, 'number_of_states': 10, 'tolerance': 0.01, 'verbosity': 0,  
    'use_multiprocessing': True, 'nprocs': 12,  
    'is_many_body': True, 'time_step': 1.0, 'es_software': 'cp2k',  
    #'path_to_npz_files': os.getcwd()+'/../7_step2_cp2k/1_DFT/2_hpc/1_example_TiO2/res',  
    'path_to_npz_files': os.getcwd()+'/res-ks-DFT',  
    'logfile_directory': os.getcwd()+'/../7_step2_cp2k/1_DFT/2_hpc/1_example_TiO2/all_logfiles',  
    'path_to_save_sd_Hvibs': os.getcwd()+'/res-mb-sd-DFT',  
    'outdir': os.getcwd()+'/res-mb-sd-DFT',  
    'start_time': 1200, 'finish_time': 1401, 'sorting_type': 'identity',  
    'apply_phase_correction': True, 'apply_orthonormalization': True,  
    'do_state_reordering': 2, 'state_reordering_alpha': 0, 'nac_algo': 0  
}
```

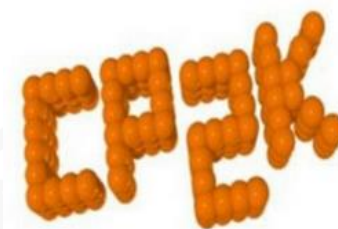
```
step3.run_step3_sd_nacs_libint(params_mb_sd)
```

# NAC in excited states basis



## Summary

- All these methodologies are implemented and available in Libra software package
  - Open-source code for quantum dynamics methodologies such as trajectory surface hopping
  - The underlying code is written in C++ for faster computation and the functions can be called from Python
  - Libint is used for computation of overlaps between GTOs
  - Sparse representation of the overlap matrices using `scipy.sparse` library in Python
  - High-throughput computation for generating the overlap matrices
  - Applicable to large systems in different electronic structure calculations frameworks such as density functional tight-binding
  - It is interfaced with many quantum chemistry codes such as CP2K, Quantum ESPRESSO, and Gaussian but the Libint interface is only available for CP2K code.



# Thank You!

Questions?