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# **Hylight**

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## ABOUT

Welcome to Hylight's documentation!

Hylight is a post-processing tool written in Python to simulate the luminescence of solids based on the results of *ab initio* computations.

You may be interested in reading our first paper using it on the luminescence of BaZrO<sub>3</sub>:Ti [[BaZrO3:Ti](#)].

To learn how to use Hylight, you can read the [tutorial](#). Finally there is a complete [reference](#) that goes over all individual functions implemented in the package.



## ATTRIBUTION

### 2.1 Licenses

Hylight is written and maintained by the PyDEF team. The source code for Hylight is distributed under the [GPLv3](#) license. This documentation is distributed under [CC-BY](#) license.

### 2.2 Citing Hylight

If you ever use Hylight for a published scientific work, we would ask you to cite the related paper [[Hylight](#)].





## CONTENT

### 3.1 BaZrO3:Ti study

Here are some utility functions and import that will be used throughout the study.

```
%matplotlib inline
```

```
import numpy as np
import matplotlib.pyplot as plt
from hylight.constants import *
```

```
eV1_in_nm = h_c / eV_in_J * 1e9
```

```
def best_max(raw_x, raw_y, f=0.95):
    "Fit a 2nd order polynom to get a position of the maximum without the noise."
    guess = np.max(raw_y)

    y = raw_y[raw_y > f * guess]
    x = raw_x[raw_y > f * guess]

    x1, x2, x3, x4 = np.mean(x**1), np.mean(x**2), np.mean(x**3), np.mean(x**4)

    x2y = np.mean(x**2 * y)
    xy = np.mean(x * y)
    my = np.mean(y)

    A = np.array([[x4, x3, x2],
                  [x3, x2, x1],
                  [x2, x1, 1.]])

    B = np.array([[x2y, xy, my]]).transpose()

    alpha, beta, gamma = np.linalg.solve(A, B)[:, 0]

    return -0.5 * beta / alpha, gamma - 0.25 * beta**2 / alpha

def load_exp(path, sep=";", skip=0):
    exp = np.loadtxt(path,
                     delimiter=sep,
                     skiprows=skip)
```

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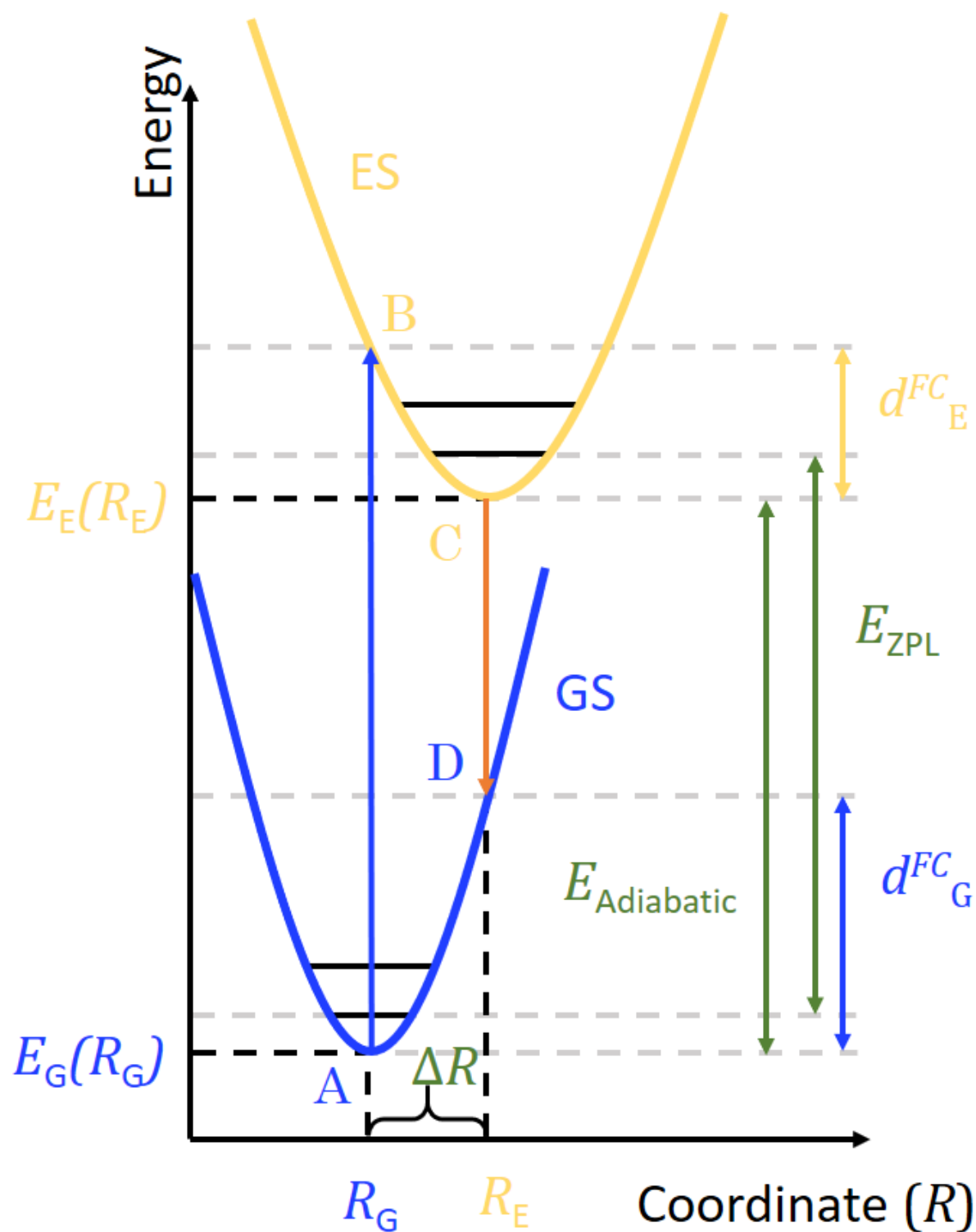
```
x, y = eVl_in_nm / exp[:, 0], exp[:, 1]
xmax, ymax = best_max(x, y)
y /= ymax
return x, y, xmax

def select_interval(x, y, emin, emax, norm=True, npoints=None):
    slice_ = (x > emin) * (x < emax)
    xs, ys = x[slice_], y[slice_] / (np.max(y[slice_]) if norm else 1.)

    if npoints is not None:
        emin = max(np.min(xs), emin)
        emax = min(np.max(xs), emax)
        xint = np.linspace(emin, emax, npoints)
        return xint, interp1d(xs, ys)(xint)

    return xs, ys
```

To perform the simulation of the spectrum of BaZrO<sub>3</sub>:Ti we need a few pieces of information. The following diagram shows the four points A, B, C and D that need to be computed in DFT. We used constrained DFT to compute B and C with an explicit hole in the VBM and an electron in the CBM. The cDFT and the regular DFT computations provides us with total energies and the positions  $R_G$  and  $R_E$  that we will use as inputs in the next section.



Given the geometry  $R_G$  we also perform a DFPT computation to get the eigen values and eigenvectors of the dynamical matrix, that is we get the  $\Gamma$  vibrational modes of the crystal.

The modes are extracted from the VASP output file and stored in the OUTCAR.npz with the command line tool `hilight-modes`:

```
$ hilight-modes vasp OUTCAR OUTCAR.npz
Loaded 120 modes from OUTCAR.
Wrote OUTCAR.npz.
```

### 3.1.1 Vibronic luminescence spectrum

#### Electronic and vibrational parameters

Here we declare the inputs that will be used later.

```
fc_shift_es = 0.1091 # B-C
fc_shift_gs = 0.2871 # D-A
delta_e_em = 2.6905 # C-D

e_adia = delta_e_em + fc_shift_gs # C-A

outcar = "OUTCAR.npz" # vibrations
poscar_gs = "POSCAR" # R_G
poscar_es = "POSCAR_ES" # R_E
T = 300 # measure temperature (K)

print(f"Adiabatic energy difference: {e_adia:.3} eV")
```

```
Adiabatic energy difference: 2.98 eV
```

#### Simulating the spectrum

Here we load the experimental data that will be used for comparison with the simulation.

```
exp_e, exp_i, max_emission = load_exp("emission_300K.csv", skip=1)

print(f"Measured maximum of emission: {max_emission:.3} eV")
```

```
Measured maximum of emission: 2.73 eV
```

We now plot the spectral function to identify important modes.

```
from hilight.multi_phonons import plot_spectral_function

fig, (ax_fc, ax_s) = plot_spectral_function(outcar, poscar_gs, poscar_es, use_cm1=True,
↳ disp=5e-1, mpl_params={
    "S_stack": {"color": "orange"},
    "FC_stack": {"color": "orange"},
    "S_peaks": {"color": "blue", "lw": 1.5},
    "FC_peaks": {"color": "blue", "lw": 1.5},
})

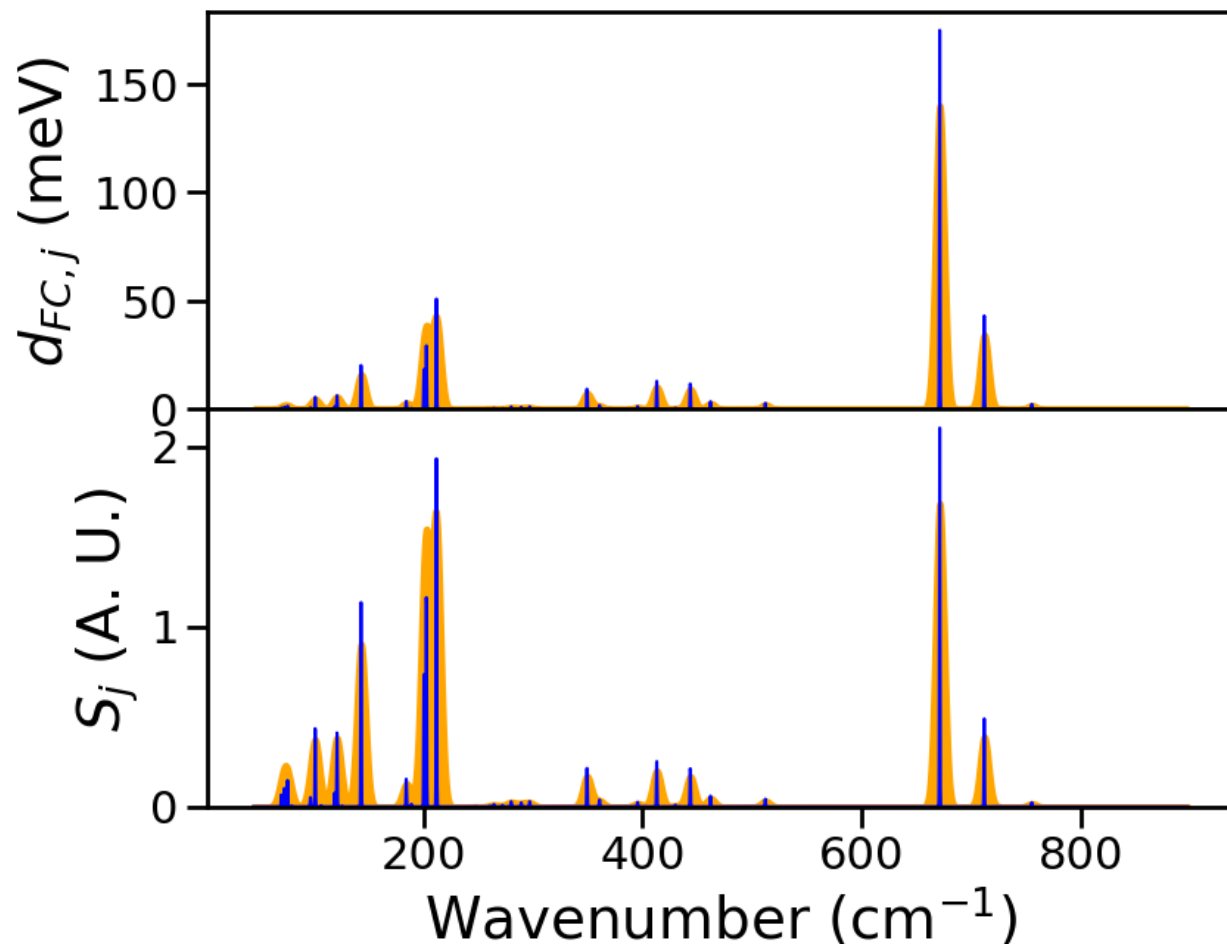
ax_fc.set_ylabel("$d_{FC,j}$ (meV)")
```

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```
ax_s.set_ylabel("$S_j$ (A. U.)")
```

```
fig.set_size_inches((9, 7))
```



We simulate the spectrum using the Mode Independent scaling approximation for the ES PES.

```
from hylight.multi_phonons import spectrum, ExpES, OmegaEff
import logging

logging.getLogger().setLevel(logging.INFO) # setting the logging level to INFO

e, sp = spectrum( # simulate the spectrum
    outcar,
    poscar_gs,
    poscar_es,
    e_adia,
    T,
    fc_shift_gs,
    fc_shift_es,
    ex_pes=ExpES.ISO_SCALE,
    omega_eff_type=OmegaEff.FC_MEAN,
```

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```
)

sp /= np.max(sp)  # normalize the spectrum

_, max_th = max(zip(sp, e))  # extract the maximum of emission energy
print(f"Computed maximum of emission: {max_th:.3} eV")
```

```
d_fc^e,v = 0.15494048765687785
omega_gs = 59.3441936353856 meV 478.64320119276204 cm-1
omega_es = 36.58255313775182 meV 295.0581896729496 cm-1
S_em = 10.055832452822724
d_fc^g,v = 0.407730650836752
FWHM 723.172263202136 meV
FWHM 0K 564.4130957462941 meV
```

Using a Gaussian line shape.  
Computed maximum of emission: 2.71 eV

Finally we setup a plot to show the result of the simulation against the measurement.

```
plt.figure(figsize=(9, 7))

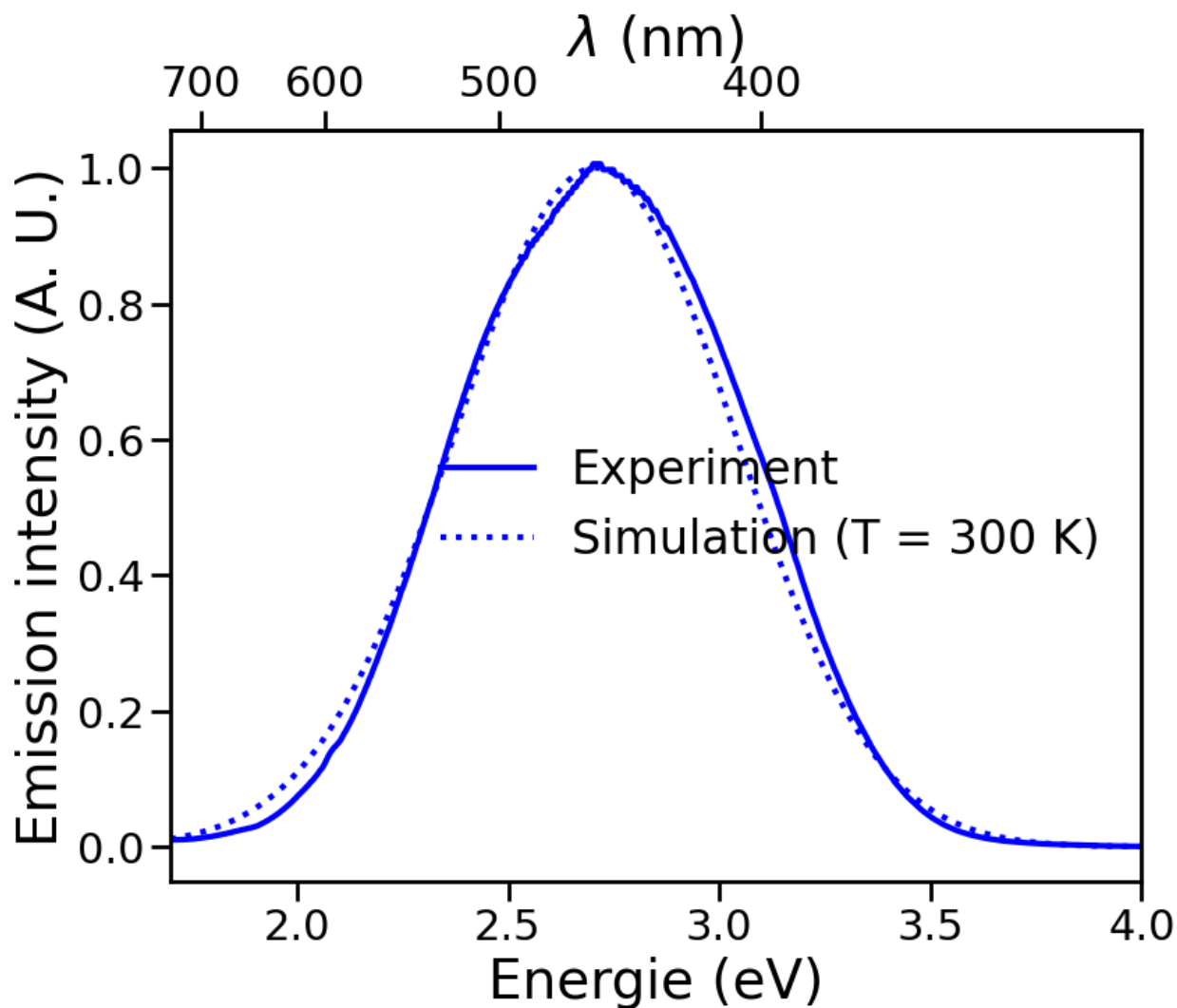
e, sp = e[1:], sp[1:]
exp_e, exp_i = exp_e[1:], exp_i[1:]
plt.plot(exp_e, exp_i, "b", lw=3, label="Experiment")
plt.plot(e, sp, ":", color="b", lw=3, label=f"Simulation (T = {T} K)")

plt.xlim(1.7, 4)
plt.xlabel("Energie (eV)")
plt.ylabel("Emission intensity (A. U.)")

ax = plt.gca()
np.seterr(divide='ignore')  # there is a division by zero occuring in the next line but
↪ it is irrelevant for our xlim
secax = ax.secondary_xaxis('top', functions=(lambda x: eV1_in_nm / x, lambda x: eV1_in_
↪ nm / x))
secax.set_xlabel("$\lambda$ (nm)")

plt.legend(prop={"size": 24})
```

```
<matplotlib.legend.Legend at 0x7f726b36eec0>
```



## 3.2 hylight package

### 3.2.1 Subpackages

`hylight.crystal` package

Submodules

`hylight.crystal.loader` module

Read vibrational modes from CRYSTAL log.

`hylight.crystal.loader.load_phonons(path)`

Load phonons from a CRYSTAL17 logfile.

**Returns**

(phonons, pops, masses) phonons: list of `hylight.mode.Mode` instances pops: population for each

atom species masses: list of SI masses

`hilight.crystal.loader.normalize(name)`

## Module contents

### hilight.findiff package

#### Submodules

#### hilight.findiff.collect module

Grab forces from a collection of single point computations and compute hessian.

`hilight.findiff.collect.dropwhile_err(pred, it, else_err)`

itertools.dropwhile wrapper that raise `else_err` if it reach the end of the file.

`hilight.findiff.collect.get_forces(n, path)`

Extract `n` forces from an OUTCAR.

`hilight.findiff.collect.get_forces_and_pos(n, path)`

Extract `n` forces and atomic positions from an OUTCAR.

`hilight.findiff.collect.get_ref_info(path)`

Load system infos from a OUTCAR.

This is an ad hoc parser, so it may fail if the OUTCAR changes a lot.

#### Returns

(atoms, ref, pops, masses) atoms: list of species names pos: positions of atoms pops: population  
for each atom species masses: list of SI masses

`hilight.findiff.collect.process_phonons(outputs, ref_output, basis_source=None, amplitude=0.01,  
nproc=1, symm=True)`

Process a set of OUTCAR files to compute some phonons using Force based finite differences.

#### Parameters

- **outputs** – list of OUTCAR paths corresponding to the finite displacements.
- **ref\_output** – path to non displaced OUTCAR.
- **basis\_source** – read a displacement basis from a path. The file is a npy file from numpy's save. If `None`, the basis is built from the displacements. If not `None`, the order of outputs *must* match the order of the displacements in the array.
- **amplitude** – amplitude of the displacement, only used if `basis_source` is *not* `None`.
- **nproc** – number of parallel processes used to load the files.
- **symm** – If `True`, use symmetric differences. OUTCARs *must* be ordered as `[+delta_1, -delta_1, +delta_2, -delta_2, ...]`.

#### Returns

the same tuple as the `load_phonons` functions.

Remark: When using non canonical basis (displacements are not along a single degree of freedom of a single atom at a time) it may be important to provide the basis explicitly because it will avoid important rounding errors found in the OUTCAR.



**hilight.findiff.gen module**

Generate a collection of positions for finite differences computations.

`hilight.findiff.gen.gen_disp(ref, basis, amplitude=0.01, symm=False)`

Iterate over displaced Poscar instances from a given set of directions.

Takes a reference position and displaces it into directions from basis to produce a set of positions for finite differences computations.

**Parameters**

- **ref** – reference Poscar instance
- **basis** – numpy array where each row is a direction.
- **amplitude** – (optional, 0.01) amplitude of the displacements in Angstrom.
- **symm** – (optional, False) when True, generate both ref+delta and ref-delta POSCARs for each direction.

`hilight.findiff.gen.random_basis(n, seed=0)`

Generate a random basis matrix or rank n.

**Parameters**

- **n** – Rank of the basis
- **seed** – (optional, 0) randomness seed.

**Returns**

a (n, n) orthonormal numpy array.

`hilight.findiff.gen.save_disp(ref, basis, disp_dest='.', amplitude=0.01, symm=False)`

Produces displaced POSCARs from a given set of directions.

Takes a reference position and displaces it into directions from basis to produce a set of positions for finite differences computations.

**Parameters**

- **ref** – reference Poscar instance
- **basis** – numpy array where each row is a direction.
- **disp\_dest** – (optional, ".") directory name or callable that determine the destination of the POSCARs.
- **amplitude** – (optional, 0.01) amplitude of the displacements in Angstrom.
- **symm** – (optional, False) when True, generate both ref+delta and ref-delta POSCARs for each direction.

`hilight.findiff.gen.save_disp_from_basis(source, basis_source, disp_dest='.', amplitude=0.01, symm=False)`

Produces displaced POSCARs from a given set of directions.

Takes a reference position and displaces it into directions from basis\_source to produce a set of positions for finite differences computations.

**Parameters**

- **source** – reference POSCAR

- **basis\_source** – Name of the basis set file. It is a npy file made with numpy's save function. Each row is a direction.
- **disp\_dest** – (optional, ".") directory name or callable that determine the destination of the POSCARs.
- **amplitude** – (optional, 0.01) amplitude of the displacements in Angstrom.
- **symm** – (optional, False) when True, generate both ref+delta and ref-delta POSCARs for each direction.

```
hilight.findiff.gen.save_random_disp(source, disp_dest='.', basis_dest='basis.npy', amplitude=0.01,  
                                     seed=0, symm=False)
```

Produces displaced POSCARs in random directions.

Takes a reference position and displaces it into random directions to produce a set of positions for finite differences computations. It also save the set of displacements in basis\_dest.

#### Parameters

- **source** – reference POSCAR
- **disp\_dest** – (optional, ".") directory name or callable that determine the destination of the POSCARs.
- **basis\_dest** – (optional, "basis.npy") Name of the basis set file. It is a npy file made with numpy's save function.
- **amplitude** – (optional, 0.01) amplitude of the displacements in Angstrom.
- **seed** – (optional, 0) random seed to use to generate the displacements.
- **symm** – (optional, False) when True, generate both ref+delta and ref-delta POSCARs for each direction.

## Module contents

A module for finite differences computations.

## hilight.jmol package

### Module contents

Write Jmol files.

```
hilight.jmol.export(dest, mode, compression=8, **opts)
```

Export a mode to JMol zip format.

#### Parameters

- **dest** – path to the JMol zip file.
- **mode** – the mode to export.
- **compression** – (optional) zipfile compression algorithm.
- **unitcell** – lattice vectors as a 3x3 matrix where vectors are in rows.
- **bonds** – a list of (*sp\_a*, *sp\_b*, *min\_dist*, *max\_dist*) where species names are strings of names of species and \*\_dist are interatomic distances in Angstrom.

- **atom\_colors** – a list of (sp, color) where sp is the name of a species and color is the name of a color or an HTML hex code (example “#FF0000” for pure red).

`highlight.jmol.export_disp(dest, struct, disp, compression=8, **opts)`

Export a difference between two positions to JMol zip format.

#### Parameters

- **dest** – path to the JMol zip file.
- **struct** – the reference position (a `highlight.struct.Struct` instance).
- **disp** – an array of displacements.
- **compression** – (optional) zipfile compression algorithm.
- **unitcell** – lattice vectors as a 3x3 matrix where vectors are in rows.
- **bonds** – a list of (*sp\_a*, *sp\_b*, *min\_dist*, *max\_dist*) where species names are strings of names of species and \*\_dist are interatomic distances in Angstrom.
- **atom\_colors** – a list of (sp, color) where sp is the name of a species and color is the name of a color or an HTML hex code (example “#FF0000” for pure red).

`highlight.jmol.format_v(v)`

`highlight.jmol.write_jmol_options(f, opts)`

Write options in the form of a JMol script.

#### Parameters

- **f** – a file like object.
- **opts** – a set of options (see export).

`highlight.jmol.write_xyz(f, atoms, ref, delta)`

Write the coordinates and displacements of in the JMol xyz format.

#### Parameters

- **f** – a file-like object to write to.
- **atoms** – the list of atom names
- **ref** – an array of atomic positions
- **delta** – an array of displacements

## highlight.phonopy package

### Submodules

#### highlight.phonopy.loader module

Module to load phonons frequencies and eigenvectors from phonopy output files.

It always uses PyYAML, but it can also need h5py to read hdf5 files.

**class** `highlight.phonopy.loader.Struct`(*pops*: 'list[int]', *lattice*: 'NDArray[float]', *masses*: 'list[float]', *atoms*: 'list[str]', *ref*: 'NDArray[float]')

Bases: object

```
atoms: list[str]

classmethod from_yaml_cell(cell: dict) → Struct

lattice: ndarray[Any, dtype[float]]

masses: list[float]

pops: list[int]

ref: ndarray[Any, dtype[float]]

highlight.phonopy.loader.get_struct(phyaml: str) → Struct

highlight.phonopy.loader.load_phonons(dir_: str) → tuple[list[highlight.mode.Mode], list[int], list[float]]

highlight.phonopy.loader.load_phonons_bandsh5(bandh5: str, phyaml: str, op=<built-in function open>) →
    tuple[list[highlight.mode.Mode], list[int], list[float]]

highlight.phonopy.loader.load_phonons_bandyaml(bandy: str) → tuple[list[highlight.mode.Mode], list[int],
    list[float]]

highlight.phonopy.loader.load_phonons_qpointsh5(qph5: str, phyaml: str, op=<built-in function open>) →
    tuple[list[highlight.mode.Mode], list[int], list[float]]

highlight.phonopy.loader.load_phonons_qpointsyaml(qpyaml: str, phyaml: str) →
    tuple[list[highlight.mode.Mode], list[int], list[float]]
```

## Module contents

### highlight.vasp package

#### Submodules

#### highlight.vasp.common module

Common utilities to read and write VASP files.

```
class highlight.vasp.common.Poscar(cell_parameters, species, species_names=None)
```

Bases: [Struct](#)

```
classmethod from_file(filename)
```

Load a POSCAR file

#### Parameters

**filename** – path to the file to read

#### Returns

a Poscar object.

```
to_file(path='POSCAR', cartesian=True)
```

Write a POSCAR file

The property system\_name may be set to change the comment at the top of the file.

#### Parameters

- **path** – path to the file to write

- **cartesian** – if True, write the file in cartesian representation, if False, write in fractional representation

### hilight.vasp.loader module

Read vibrational modes from VASP files.

`hilight.vasp.loader.load_phonons(path)`

Load phonons from a OUTCAR.

Remark: This function is a bit heavy because of text parsing. You may want to use `hyligh-modes` to parse one for all the file and later load that prepared file using `npz.load_phonons` instead.

#### Returns

(phonons, pops, masses) phonons: list of `hilight.mode.Mode` instances pops: population for each atom species masses: list of SI masses

`hilight.vasp.loader.load_poscar(path)`

Read the positions from a POSCAR.

#### Returns

a `np.ndarray((natoms, 3), dtype=float)`

`hilight.vasp.loader.load_poscar_latt(path)`

Read the positions and the lattice parameters from a POSCAR.

#### Returns

a (`np.ndarray((natoms, 3), dtype=float)`, `nd.array((3, 3), dtype=float)`) first element is the set of positions second element is the lattice parameters

### hilight.vasp.utils module

Pervasive utilities for `hilight.vasp` submodule.

`hilight.vasp.utils.make_finite_diff_poscar(outcar, poscar_gs, poscar_es, A=0.01, load_phonons=<function load_phonons>, bias=0, mask=None)`

Compute positions for evaluation of the curvature of the ES PES.

#### Parameters

- **outcar** – the name of the file where the phonons will be read.
- **poscar\_gs** – the path to the ground state POSCAR
- **poscar\_es** – the path to the excited state POSCAR, it will be used as a base for the generated Poscars
- **A** – (optional, 0.01) the amplitude of the displacement in Å.
- **load\_phonons** – (optional, `vasp.loader.load_phonons`) the procedure use to read outcar

#### Returns

(mu, pes\_left, pes\_right) mu: the effective mass in kg pes\_left: a Poscar instance representing the left displacement pes\_right: a Poscar instance representing the right displacement

## Module contents

### 3.2.2 Submodules

#### 3.2.3 `hylight.constants` module

A collection of useful physics constants.

#### 3.2.4 `hylight.loader` module

Wrapper for the npz loader.

#### 3.2.5 `hylight.mode` module

Vibrational mode and related utilities.

**class** `hylight.mode.Mode`(*atoms, n, real, energy, ref, eigenvector, masses*)

Bases: `object`

The representation of a vibrational mode.

It stores the eigenvector and eigendisplacement. It can be used to project other displacement on the eigenvector.

**huang\_rhys**(*delta\_R*)

Compute the Huang-Rhys factor

$$S_i = 1/2 \frac{\omega}{\hbar} [(M^{1/2T} \Delta R) \cdot \gamma_i]^2 = 1/2 \frac{\omega}{\hbar} \sum_i m_i^{1/2} \gamma_i \Delta R_i^2$$

##### Parameters

**delta\_R** – displacement in SI

**project**(*delta\_Q*)

Project delta\_Q onto the eigenvector

**project\_coef2**(*delta\_Q*)

Square lenght of the projection of delta\_Q onto the eigenvector.

**project\_coef2\_R**(*delta\_R*)

Square lenght of the projection of delta\_R onto the eigenvector.

**to\_jmol**(*\*\*opts*)

Write a mode into a Jmol file.

See `hylight.jmol.export()` for the parameters.

**to\_traj**(*duration, amplitude, framerate=25*)

Produce a ase trajectory for animation purpose.

##### Parameters

- **duration** – duration of the animation in seconds
- **amplitude** – amplitude applied to the mode in Å (the modes are normalized)
- **framerate** – number of frame per second of animation

`hilight.mode.dynamical_matrix(phonons)`

Retrieve the dynamical matrix from a set of modes.

Note that if some modes are missing the computation will fail.

**Parameters**

**phonons** – list of modes

`hilight.mode.get_HR_factors(phonons, delta_R_tot, mask=None)`

Compute the Huang-Rhys factors for all the real modes with energy above bias.

**Parameters**

- **phonons** – list of modes
- **delta\_R\_tot** – displacement in SI
- **mask** – a mask to filter modes based on their energies.

`hilight.mode.get_energies(phonons, mask=None)`

Return an array of mode energies in SI

`hilight.mode.rot_c_to_v(phonons)`

Rotation matrix from Cartesian basis to Vibrational basis (right side).

### 3.2.6 hilight.mono\_phonon module

Simulation of spectra in 1D model.

`hilight.mono_phonon.compute_spectrum(e, e_zpl, S, sig, e_phonon_g)`

Compute a spectrum from 1D model with experimental like inputs

**Parameters**

- **e** – a numpy array of energies to compute the spectrum at
- **e\_zpl** – energy of the zero phonon line, in eV
- **S** – the emission Huang-Rhys factor
- **sig** – the lineshape standard deviation

**E\_phonon\_g**

the ground state phonon energy

`hilight.mono_phonon.huang_rhys(stokes_shift, e_phonon)`

Huang-Rhys factor from the Stokes shift and the phonon energy.

`hilight.mono_phonon.sigma(T, S_em, e_phonon_g, e_phonon_e)`

Temperature dependant standard deviation of the lineshape.

**Parameters**

- **T** – temperature in K
- **S\_em** – emission Huang-Rhys factor
- **e\_phonon\_g** – energy of the GS PES vibration (eV)
- **e\_phonon\_e** – energy of the ES PES vibration (eV)

```
hilight.mono_phonon.spectrum(e_zpl, T, fc_shift_g, fc_shift_e, e_phonon_g, hard_osc=False, n_points=5000,  
                             e_min=0, e_max=5)
```

Compute a spectrum from a single vibrational mode and some energetic terms.

#### Parameters

- **e\_zpl** – energy of the zero phonon line, in eV
- **T** – temperature in K
- **fc\_shift\_g** – Franck-Condon shift of emission in eV
- **fc\_shift\_e** – Franck-Condon shift of absorption in eV
- **e\_phonon\_g** – Mode energy in ground state in eV
- **hard\_osc** – boolean, use the hard oscillator mode: vibration mode has the same energy in GD and ES
- **n\_points** – number of points in the spectrum
- **e\_min** – energy lower bound for the spectrum, in eV
- **e\_max** – energy higher bound for the spectrum, in eV

### 3.2.7 hilight.multi\_phonons module

Simulation of spectra in nD model.

```
class hilight.multi_phonons.ExPES(wm)
```

Bases: object

Mode of approximation of the ES PES curvature.

*ISO\_SCALE*: We suppose eigenvectors are the same, but the frequencies are scaled by a constant factor.

*SINGLE\_ES\_FREQ*: We suppose all modes have the same frequency (that should be provided).

*FULL\_ND*: (Not implemented) We know the frequency of each ES mode.

**FULL\_ND**: *ExPES* = <hilight.multi\_phonons.ExPES object>

**ISO\_SCALE**: *ExPES* = <hilight.multi\_phonons.ExPES object>

**SINGLE\_ES\_FREQ**: *ExPES* = <hilight.multi\_phonons.ExPES object>

```
class hilight.multi_phonons.LineShape(value)
```

Bases: Enum

Line shape type.

**GAUSSIAN** = 0

**LORENTZIAN** = 1

**NONE** = 2

```
class hilight.multi_phonons.Mask(intervals)
```

Bases: object

An energy based mask for the set of modes.



**accept**(*value*)

Return True if value is not under the mask.

**add\_interval**(*interval*)

Add a new interval to the mask.

**as\_bool**(*ener*)

Convert to a boolean *np.ndarray* based on *ener*.

**classmethod from\_bias**(*bias*)

Create a mask that reject modes of energy between 0 and *bias*.

**Parameters**

**bias** – minimum of accepted energy (eV)

**Returns**

a fresh instance of *Mask*.

**plot**(*ax, unit*)

Add a graphical representation of the mask to a plot.

**Parameters**

- **ax** – a matplotlib *Axes* object.
- **unit** – the unit of energy to use (ex: `hylight.constant.eV_in_J` if the plot uses eV)

**Returns**

a function that must be called without arguments after resizing the plot.

**reject**(*value*)

Return True if *value* is under the mask.

**class** `hylight.multi_phonons.OmegaEff`(*value*)

Bases: Enum

Mode of computation of a effective frequency.

*FC\_MEAN*:

$$\Omega = \frac{\sum_j \omega_j d_j^{\text{FC}}}{\sum_j d_j^{\text{FC}}}$$

Should be used with *ExpES.ISO\_SCALE* because it is associated with the idea that all the directions are softened equally in the excited state.

*HR\_MEAN*:

$$\Omega = \frac{d^{\text{FC}}}{S_{\text{tot}}} = \frac{\sum_j \omega_j S_j}{\sum_j S_j}$$

*HR\_RMS*:

$$\Omega = \sqrt{\frac{\sum_j \omega_j^2 S_j}{\sum_j S_j}}$$

*FC\_RMS*:

$$\Omega = \sqrt{\frac{\sum_j \omega_j^2 d_j^{\text{FC}}}{\sum_j d_j^{\text{FC}}}}$$

Should be used with `EXPES.SINGLE_ES_FREQ` because it makes sense when we get only one  $\Omega_{\text{eff}}$  for the excited state (this single effective frequency should be computed beforehand)

`ONED_FREQ`:

$$\Omega = \frac{(\Delta Q^T D \Delta Q)}{\Delta Q^2}$$

Correspond to the actual curvature in the direction of the displacement.

`FC_MEAN = 0`

`FC_RMS = 3`

`HR_MEAN = 1`

`HR_RMS = 2`

`ONED_FREQ = 4`

`hilight.multi_phonons.compute_delta_R(poscar_gs, poscar_es)`

Return  $\Delta R$  in Å.

**Parameters**

- **poscar\_gs** – path to ground state positions file.
- **poscar\_es** – path to excited state positions file.

**Returns**

a np.array of (n, 3) shape where n is the number of atoms.

`hilight.multi_phonons.compute_spectrum(phonons, delta_R, zpl, fwhm, e_max, resolution_e, bias=0, mask=None, window_fn=<function hamming>, pre_convolve=None, shape=LineShape.GAUSSIAN)`

Compute a luminescence spectrum with the time-dependant formulation with an arbitrary linewidth.

**Parameters**

- **phonons** – list of modes
- **delta\_R** – displacement in Å
- **zpl** – zero phonon line energy in eV
- **fwhm** – zpl lineshape full width at half maximum in eV or None if fwhm is None: the raw spectrum is not convoluted with a line shape if fwhm < 0: the spectrum is convoluted with a lorentizan line shape if fwhm > 0: the spectrum is convoluted with a gaussian line shape
- **e\_max** – max energy in eV (should be at least > 2\*zpl)
- **resolution\_e** – energy resolution in eV
- **bias** – ignore low energy vibrations under bias in eV
- **window\_fn** – windowing function in the form provided by numpy (see numpy.hamming)
- **pre\_convolve** – (float, optional, None) if not None, standard deviation of the pre convolution gaussian

**Returns**

(energy\_array, intensity\_array)

```

highlight.multi_phonons.compute_spectrum_soft(phonons, delta_R, zpl, T, fc_shift_gs, fc_shift_es, e_max,
                                             resolution_e, mask=None, window_fn=<function
                                             hamming>, pre_convolve=None,
                                             shape=LineShape.GAUSSIAN,
                                             omega_eff_type=OmegaEff.FC_MEAN, result_store=None,
                                             ex_pes=<highlight.multi_phonons.ExPES object>,
                                             correct_zpe=False)

```

Compute a spectrum without free parameters.

#### Parameters

- **phonons** – list of modes
- **delta\_R** – displacement in Å
- **zpl** – zero phonon line energy in eV
- **T** – temperature in K
- **fc\_shift\_gs** – Ground state/absorption Franck-Condon shift in eV
- **fc\_shift\_es** – Excited state/emission Franck-Condon shift in eV
- **e\_max** – max energy in eV (should be at least  $> 2 \cdot \text{zpl}$ )
- **resolution\_e** – energy resolution in eV
- **bias** – ignore low energy vibrations under bias in eV
- **window\_fn** – windowing function in the form provided by numpy (see numpy.hamming)
- **pre\_convolve** – (float, optional, None) if not None, standard deviation of the pre convolution gaussian
- **shape** – ZPL line shape.
- **omega\_eff\_type** – mode of evaluation of effective frequency.
- **result\_store** – a dictionary to store some intermediate results.
- **ex\_pes** – mode of evaluation of the ES PES curvature.
- **correct\_zpe** – correct the ZPL to take the zero point energy into account.

#### Returns

(energy\_array, intensity\_array)

```

highlight.multi_phonons.compute_spectrum_width_ah(phonons_gs, phonons_es, delta_R, zpl, T, e_max,
                                                  resolution_e, bias=0, mask=None,
                                                  window_fn=<function hamming>,
                                                  pre_convolve=None, shape=LineShape.GAUSSIAN,
                                                  result_store=None, correct_zpe=False)

```

Luminescence spectrum with a line width that takes the ES PES curvature into account.

Uses the full Dushinsky matrix for the computation of the line width, but assume the Dushinsky matrix to be the identity in the computation of the FC integrals (thus ignoring mode mixing).

#### Parameters

- **phonons** – list of modes
- **delta\_R** – displacement in Å
- **zpl** – zero phonon line energy in eV
- **T** – temperature in K

- **fc\_shift\_gs** – Ground state/absorption Franck-Condon shift in eV
- **fc\_shift\_es** – Exceited state/emmission Franck-Condon shift in eV
- **e\_max** – max energy in eV (should be at least  $> 2 \times \text{zpl}$ )
- **resolution\_e** – energy resolution in eV
- **bias** – ignore low energy vibrations under bias in eV
- **window\_fn** – windowing function in the form provided by numpy (see `numpy.hamming`)
- **pre\_convolve** – (float, optional, None) if not None, standard deviation of the pre convolution gaussian
- **shape** – ZPL line shape.
- **omega\_eff\_type** – mode of evaluation of effective frequency.
- **result\_store** – a dictionary to store some intermediate results.
- **ex\_pes** – mode of evaluation of the ES PES curvature.
- **correct\_zpe** – correct the ZPL to take the zero point energy into account.

**Returns**

(energy\_array, intensity\_array)

`hilight.multi_phonons.duschinsky(phonons_a, phonons_b)`

Dushinsky matrix from b to a  $\$S_{\{a \text{ gets } b\}}$ .

`hilight.multi_phonons.dynmatshow(dynmat, blocks=None)`

Plot the dynamical matrix.

**Parameters**

- **dynmat** – numpy array representing the dynamical matrice in SI.
- **blocks** – (optional) a list of coloured blocks in the form *(label, number\_of\_atoms, color)*.

`hilight.multi_phonons.effective_phonon_energy(omega_eff_type, hrs, es, masses, delta_R=None)`

Compute an effective phonon energy in eV following the strategy of `omega_eff_type`.

**Parameters**

- **omega\_eff\_type** – The mode of evaluation of the effective phonon energy.
- **hrs** – The array of Huang-Rhyes factor for each mode.
- **es** – The array of phonon energy in eV.
- **masses** – The array of atomic masses in atomic mass unit.
- **delta\_R** – The displacement between GS and ES in Å. It is only required if `omega_eff_type` is `ONED_FREQ`.

**Returns**

The effective energy in eV.

`hilight.multi_phonons.fc_spectrum(phonons, delta_R, n_points=5000, disp=1)`

`hilight.multi_phonons.freq_from_finite_diff(left, mid, right, mu, A=0.01)`

Compute a vibration energy from three energy points.

**Parameters**

- **left** – energy (eV) of the left point

- **mid** – energy (eV) of the middle point
- **right** – energy (eV) of the right point
- **mu** – effective mass associated with the displacement from the middle point to the sides.
- **A** – amplitude (A) of the displacement between the middle point and the sides.

`hilight.multi_phonons.hr_spectrum(phonons, delta_R, n_points=5000, disp=1)`

`hilight.multi_phonons.plot_spectral_function(mode_source, poscar_gs, poscar_es,  
load_phonons=<function load_phonons>,  
use_cm1=False, disp=1, mpl_params=None,  
mask=None)`

Plot a two panel representation of the spectral function of the distortion.

#### Parameters

- **mode\_source** – path to the mode file (by default a pickle file)
- **poscar\_gs** – path to the file containing the ground state atomic positions.
- **poscar\_es** – path to the file containing the excited state atomic positions.
- **load\_phonons** – a function to read mode\_source.
- **use\_cm1** – use cm1 as the unit for frequency instead of meV.
- **disp** – standard deviation of the gaussians in background in meV.
- **mpl\_params** – dictionary of kw parameters for pyplot.plot.
- **mask** – a mask used in other computations to show on the plot.

#### Returns

(figure, (ax\_FC, ax\_S))

`hilight.multi_phonons.rect(n)`

A dummy windowing function that works like numpy.hamming, but as no effect on data.

`hilight.multi_phonons.sigma_full_nd(T, delta_R, modes_gs, modes_es, mask=None)`

Compute the width of the ZPL for the ExPES.FULL\_ND mode.

#### Parameters

- **T** – temperature in K.
- **delta\_R** – distortion in Å.
- **modes\_gs** – list of Modes of the ground state.
- **modes\_es** – list of Modes of the excited state.

#### Returns

np.array with only the width for the modes that are real in ground state.

`hilight.multi_phonons.sigma_hybrid(T, S, e_phonon, e_phonon_e)`

Compute the width of the ZPL for the ExPES.SINGLE\_ES\_FREQ mode.

`hilight.multi_phonons.spectrum(mode_source, poscar_gs, poscar_es, zpl, T, fc_shift_gs=None,  
fc_shift_es=None, e_max=None, resolution_e=0.0001, bias=0,  
mask=None, load_phonons=<function load_phonons>,  
pre_convolve=None, shape=LineShape.GAUSSIAN,  
omega_eff_type=OmegaEff.FC_MEAN, result_store=None,  
ex_pes=<hilight.multi_phonons.ExPES object>, correct_zpe=False)`

Compute a complete spectrum without free parameters.

**Parameters**

- **mode\_source** – path to the vibration computation output file (by default a pickled file)
- **path\_struct\_gs** – path to the ground state relaxed structure file (by default a POSCAR)
- **path\_struct\_es** – path to the excited state relaxed structure file (by default a POSCAR)
- **zpl** – zero phonon line energy in eV
- **T** – temperature in K
- **fc\_shift\_gs** – Ground state/absorption Franck-Condon shift in eV
- **fc\_shift\_es** – Exited state/emmission Franck-Condon shift in eV
- **e\_max** – max energy in eV (should be at least  $> 2 \cdot \text{zpl}$ )
- **resolution\_e** – energy resolution in eV
- **bias** – ignore low energy vibrations under bias in eV
- **pre\_convolve** – (float, optional, None) if not None, standard deviation of the pre convolution gaussian
- **load\_phonons** – a function that takes mode\_source and produce a list of phonons. By default expect an mode\_source.

**Returns**

(energy\_array, intensity\_array)

### 3.2.8 hylight.npz module

Serialization of modes to numpy zipped file.

`hylight.npz.archive_modes(modes, dest, compress=False)`

Store modes in dest using numpy's npz format.

**Parameters**

- **modes** – a list of Mode objects.
- **dest** – the path to write the modes to.

**Returns**

the data returned by load\_phonons.

`hylight.npz.load_phonons(source)`

Load modes from a Hylight archive.

`hylight.npz.load_phonons_npz(source)`

`hylight.npz.load_phonons_pickle(source, gz=False)`

Load modes from a pickled file.

`hylight.npz.pops_and_masses(modes)`

### 3.2.9 hylight.struct module

A generic representation of a crystal cell.

**class** `hylight.struct.Struct(cell_parameters, species, species_names=None)`

Bases: `object`

A general description of a periodic crystal cell.

Store all the required infos to describe a given set of atomic positions.

**property** `atoms`

List the species names in an order matching *self.raw*.

**copy()**

**property** `raw`

Return an array of atomic positions.

This can be modified overwritten, but not modified in place.

**property** `system_name`

The name of the system, eventually generated from formula.

Can be overwritten.

### 3.2.10 hylight.utils module

Pervasive utilities.

**exception** `hylight.utils.InputError`

Bases: `ValueError`

An exception raised when the input files are not as expected.

`hylight.utils.gaussian(e, sigma, standard=True)`

A Gaussian function.

**Parameters**

- **e** – mean
- **sigma** – standard deviation
- **standard** – if True the curve is normalized to have an area of 1 if False the curve is normalized to have a maximum of 1

`hylight.utils.gen_translat(lattice: ndarray)`

Generate all translations to adjacent cells

**Parameters**

**lattice** – `np.ndarray([a, b, c])` first lattice parameter

`hylight.utils.make_cell(val)`

An helper function to create a mutable cell/box.

**Parameters**

**val** – initial value

**Returns**

a function `c`

**Example:**

```
>>> c = make_cell(42)
>>> c()
42
>>> c(23)
>>> c()
23
```

`hilight.utils.measure_fwhm(x, y)`

Measure the full width at half maximum of a given spectrum.

Warning: It may fail if there are more than one band that reach half maximum in the array. In this case you may want to use `select_interval` to make a window around a single band.

**Parameters**

- **x** – the energy array
- **y** – the intensity array

**Returns**

FWHM in the same unit as x.

`hilight.utils.periodic_diff(lattice, ref, disp)`

Compute the displacement between `ref` and `disp`, accounting for periodic conditions.

`hilight.utils.select_interval(x, y, emin, emax, normalize=False, npoints=None)`

Extract an interval of a spectrum and return the windows `x` and `y` arrays.

**Parameters**

- **x** – x array
- **y** – y array
- **emin** – lower bound for the window
- **emax** – higher bound for the window
- **normalize** – if true, the result y array is normalized
- **npoints** – if an integer, the result arrays will be interpolated to contains exactly `npoints` linearly distributed between `emin` and `emax`.

**Returns**

(`windowed_x`, `windowed_y`)

### 3.2.11 Module contents

The Hylight package.

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`hylight.setup_logging()`



## INDICES AND TABLES

- `genindex`
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## BIBLIOGRAPHY

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