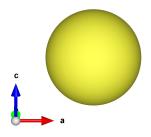
User manual for atorvi - ATomic ORbitals VIzualization package



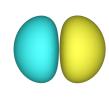






Table of contents

- Introduction
- Minimal theory: Atomic orbitals
- Installation
- Workflow
 - Step 0: Import the package
 - Step 1: Create the output file in XCrysDen format
 - Step 2: Add atoms to the file
 - * 2.1. Manual atoms addition
 - * 2.2. Reading atoms from a file
 - Step 3: Add orbitals to the file
 - * 3.1. Orbitals at arbitrary positions
 - * 3.2. Orbitals at atoms positions
 - * 3.3. Orbitals at all atoms of the same element
 - * 3.4. Mixing (or hybridization) of atomic orbitals
 - Step 4: Write the file
- Important notes
- Author

Introduction

atorvi is a Python package for creating three-dimensional visualizations of atomic orbitals in crystalline materials. These visualizations can be used in research publications, educational materials, and scientific analysis.

Common applications in condensed matter physics and solid-state chemistry include:

- 1. **Magnetic Ordering**: Visualize orbital overlap between magnetic atoms in ferromagnetic and antiferromagnetic materials to understand exchange interactions between neighboring sites.
- 2. **Chemical Bonding**: Examine hybrid orbitals (sp², sp³) in semiconductors and insulators to understand covalent bonding and electron sharing between atoms.
- 3. **Electronic Correlations**: Study electron localization in strongly correlated systems like Mott insulators, complementing computational results from DFT+U and DMFT calculations.
- 4. **Band Structure Analysis**: Map the orbital contributions to electronic bands when analyzing band structures and density of states (DOS).
- 5. **Crystal Field Theory**: Explore how crystal fields affect d-orbital splitting in transition metal compounds, helping explain their electronic, magnetic and optical behavior.

The package generates files in XCrysDen format, enabling both interactive exploration and high-resolution image export.

Minimal theory: Atomic orbitals

The atomic orbital of the hydrogen-like atom with quantum numbers n and ℓ is calculated as:

$$\psi_{n\ell}(\mathbf{r}) = R_{n\ell}(r) X_{\ell c}(\mathbf{r}),$$

where $X_{\ell c}$ are the cubic harmonics.

The radial part of the atomic orbital is calculated as:

$$R_{n\ell}(r) = \sqrt{\left(\frac{2Z}{na_0}\right)^3 \frac{(n-\ell-1)!}{2n(n+\ell)!}} e^{-Zr/na_0} \left(\frac{2Zr}{na_0}\right)^{\ell} L_{n-\ell-1}^{(2\ell+1)} \left(\frac{2Zr}{na_0}\right),$$

where: $L_{n-\ell-1}^{(2\ell+1)}$ – are the generalized Laguerre polynomials, a_0 is the Bohr radius and Z is the screened nuclear charge. We use the effective nuclear charge by Clementi $et\ al.$ to account the shielding effect of inner-shell electrons on outer-shell electrons, providing a more accurate representation of the potential energy experienced by electrons in multi-electron atoms during calculations.

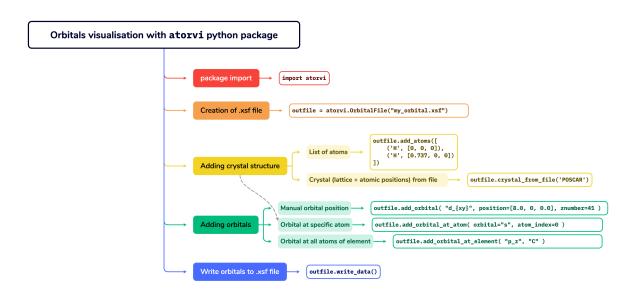
atorvi always generates orbitals of the outermost shell for a given element.

Installation

To install **atorvi**, you can use pip:

```
pip install atorvi
```

Workflow



Step 0: Import the package

In your python script or Jupyter notebook write

```
import atorvi
```

After the import, several useful objects are available for you, such as

```
atorvi.supported_orbitals,
atorvi.p_orbitals,
atorvi.d_orbitals,
atorvi.f_orbitals
```

Step 1: Create the output file in XCrysDen format

Create an instance of the OrbitalFile class, which will be used to write the .xsf file.

```
outfile = atorvi.OrbitalFile("my_orbital.xsf")
```

Step 2: Add atoms to the file

This step is optional. You can generate orbitals without adding atoms to the file.

You can add atoms to the file manually creating a non-periodic molecule, or you can read crystal structure from a (XSF/POSCAR/CIF) file.

2.1. Manual atoms addition .add_atoms() method accepts a list of tuples, where each tuple contains the atomic symbol and the coordinates of the atom. You can add atoms one-by-one or in a batch. Atomic coordinates are given in angstroms.

```
outfile.add_atoms([
    ('H', [0, 0, 0]),
    ('H', [0.737, 0, 0])
])
```

2.2. Reading atoms from a file .crystal_from_file() method accepts a path to the file with crystal structure. We use pymatgen to read the file, so this package should be installed in your environment with pip install pymatgen to use this method. Supported file formats are the same that pymatgen supports.

```
structure = outfile.crystal_from_file('./KCuF3_structure.xsf')
```

this method returns a pymatgen.core.structure.Structure object, which can be used to further manipulate the crystal structure in your script if necessary.

See example of reading structure from file in examples/structure_from_file/

Step 3: Add orbitals to the file

You can add to the file the following orbitals:

```
print(atorvi.supported_orbitals)

['s',
    'p_z', 'p_x', 'p_y',
    'd_{3z^2-r^2}', 'd_{xz}', 'd_{yz}', 'd_{xy}', 'd_{x^2-y^2}',
    'f_{z^3}', 'f_{xz^2}', 'f_{yz^2}', 'f_{xyz}', 'f_{z(x^2-y^2)}',
    'f_{x(x^2-3y^2)}', 'f_{y(3x^2-y^2)}']
```

3.1. Orbitals at arbitrary positions You can add an orbital at the arbitrary position using .add_orbital(orbital, position, znumber, coeff) method. This method accepts the following arguments:

```
orbital: str
The type of orbital (e.g., "s", "p_x", "d_{xy}", etc.).

position: list, optional
The position of the orbital in angstroms (default is [0.0, 0.0, 0]).

znumber: int, optional
The atomic number of the element (default is 8 i.e. Oxygen).

coeff: float, optional
The coefficient of the orbital (default is 1.0).

"""
```

Example:

```
outfile.add_orbital("d_{3z^2-r^2}", position=[0, 0, 0], znumber=41)
```

See also examples/d_orbitals/ and examples/f_orbitals/ for more examples.

3.2. Orbitals at atoms positions If you have created some atoms at the Step 2, you can add an orbital at the position of the *i*-th atom using .add_orbital_at_atom(orbital, atom_index, coeff) method. This method accepts the following arguments:

```
orbital : str
   The type of orbital (e.g., "s", "p_x", "d_{xy}", etc.).
atom_index : int
   The index of the atom in the system.
coeff : float, optional
   The coefficient of the orbital (default is 1.0).
"""
```

Example:

```
atoms = [
    ('Cu', [0.0,0.0,0.0]),
    ('Cu', [2.0,0.0,0.0]),
    ]

outfile.add_atoms(atoms)

outfile.add_orbital_at_atom('d_{xz}', 0)
outfile.add_orbital_at_atom('d_{yz}', 1)
```

See also examples/orbital ordering for more examples.

3.3. Orbitals at all atoms of the same element You can add an orbital at all atoms of the same element using .add_orbital_at_element(orbital, element, coeff) method. This method accepts the following arguments:

```
"""
orbital : str
    The type of orbital (e.g., "s", "p_x", "d_{xy}", etc.).
element : str
    The symbol of the element.
coeff : float, optional
    The coefficient of the orbital (default is 1.0).
"""
```

Example:

```
outfile.add_orbital_at_element("p_z", "C")
```

See also examples/orbital_ordering/ for more examples.

3.4. Mixing (or hybridization) of atomic orbitals You can get hybridized (sp² or sp³ or *etc*) orbitals combining the atomic orbitals with specific coefficients using the coeff parameter of the .add_orbital* methods. For example, the sp³ hybridized orbital is: $\frac{1}{2}s + \frac{1}{2}p_x + \frac{1}{2}p_y + \frac{1}{2}p_z$.

One can get this with:

```
outfile.add_atoms([("C", [0, 0, 0])])

outfile.add_orbital_at_atom("s", 0, coeff=0.5)
outfile.add_orbital_at_atom("p_x", 0, coeff=0.5)
outfile.add_orbital_at_atom("p_y", 0, coeff=0.5)
outfile.add_orbital_at_atom("p_z", 0, coeff=0.5)
```

See examples/sp3_hybridization/ for more examples.

Using the same coeff parameter, you can also get molecular bonding and antibonding orbitals. For example, the antibonding molecular orbital of hydrogen molecule is: $\varphi_{ABO} = \frac{1}{\sqrt{2}} s_A - \frac{1}{\sqrt{2}} s_B$:

```
outfile.add_atoms([
    ('H', [0, 0, 0]),
    ('H', [0.737, 0, 0])
])
outfile.add_orbital_at_atom('s', 0, coeff = 0.707)
outfile.add_orbital_at_atom('s', 1, coeff = -0.707)
```

See also examples/H2_molecule/.

Step 4: Write the file

Just call .write_data() method to write the .xsf file with the structure and the orbitals. There is optional parameter squared=False for this function. If squared=True, the value of the orbital is squared. This is useful for visualizing the density of the orbital.

Example:

```
outfile.write_data()
```

Important notes

- These are atomic orbitals of a hydrogen-like atom. Their radial part is more or less close to the radial part of multi-electron atom, thanks to the screened nuclear charge we use. But in molecules or solid state the radial part could differ a lot. Therefore, use obtained pictures with caution only for qualitative description and illustrations.
- It is a good idea to vary the isosurface value in your plotting software to get the best picture of orbitals.
- The 3D data generated by atorvi uses the regular cubic mesh with the 0.05 Å grid spacing by default. You can tune this parameter at the very first step with grid_step=0.05 parameter of the OrbitalFile class:

```
outfile = atorvi.OrbitalFile("my_orbital.xsf", grid_step=0.1)
```

A smaller grid_step will give you a more smooth picture, but will also increase the size of the output file. And vice versa.

Author

atorvi is developed and maintained by Dmitry Korotin. Contributions, suggestions, and feedback are welcome to help improve the project.