

src.molecule.Molecule.align

```
graph LR; A[src.molecule.Molecule.align] --> B[src.molecule.Molecule.center_of_mass]; A --> C[src.molecule.get_rotate_translate];
```

The diagram illustrates the dependencies of the `src.molecule.Molecule.align` method. A central box on the left, labeled `src.molecule.Molecule.align`, has two arrows pointing to the right. The top arrow points to a box labeled `src.molecule.Molecule.center_of_mass`, and the bottom arrow points to a box labeled `src.molecule.get_rotate_translate`. This indicates that the `align` method relies on both the `center_of_mass` property and the `get_rotate_translate` method.

src.molecule.Molecule.center_of_mass

src.molecule.get_rotate_translate