

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`. The central box is shaded gray, while the two target boxes are white with black borders.

src.molecule.Molecule.center  
\_of\_mass

src.molecule.get\_rotate  
\_translate