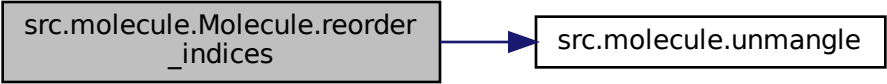


src.molecule.Molecule.reorder  
\_indices



```
graph LR; A["src.molecule.Molecule.reorder_indices"] --> B["src.molecule.unmangle"]
```

A diagram showing a call to `src.molecule.Molecule.reorder_indices` in a gray box, with a blue arrow pointing to a white box containing `src.molecule.unmangle`.

src.molecule.unmangle