

forcebalance.molecule.is  
\_charmm\_coord

```
graph LR; A[forcebalance.molecule.is_charmm_coord] --> B[forcebalance.molecule.isint]; A --> C[forcebalance.molecule.isfloat];
```

A diagram illustrating a branching structure. On the left, a gray rectangular box contains the text "forcebalance.molecule.is\_charmm\_coord". Two blue arrows originate from the right side of this box. The top arrow points to a white rectangular box containing the text "forcebalance.molecule.isint". The bottom arrow points to another white rectangular box containing the text "forcebalance.molecule.isfloat".

forcebalance.molecule.isint

forcebalance.molecule.isfloat