

MIX

```
python mix.py --hg_fa_fp ../../HOMD/14.5/homd.fa --hg_tax_fp  
../../HOMD/14.5/homd.tax --mock_fa_fp  
../../BEI MOCK/HM-783D/mock.fa --mock_tax_fp  
../../BEI MOCK/HM-783D/mock.tax
```

1. Match taxonomy by fuzzy seach.
 - <https://github.com/seatgeek/fuzzywuzzy>
2. Color print
 - <https://pypi.python.org/pypi/termcolor>
3. Since we only have taxonomy label at genus and species level for mock database, we will convert HOMD/GG taxonomy to match mock taxonomy if matched or selected to be matched.

Written with [StackEdit](#).