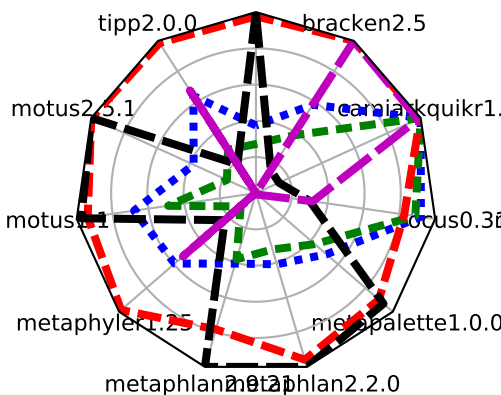


■ ■ ■ Weighted UniFrac error    
 ■ ■ ■ L1 norm error    
 ■ ■ ■ Completeness (recall)    
 ■ Purity (precision)    
 ■ ■ ■ False positives

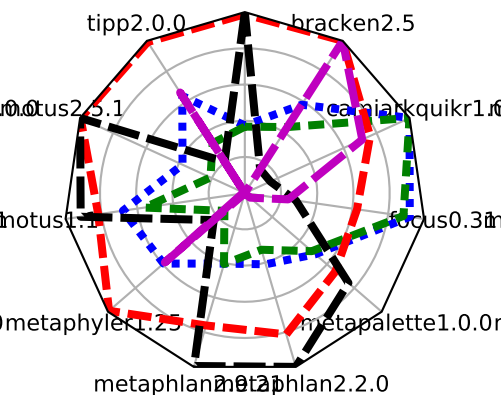
### phylum

sourmash



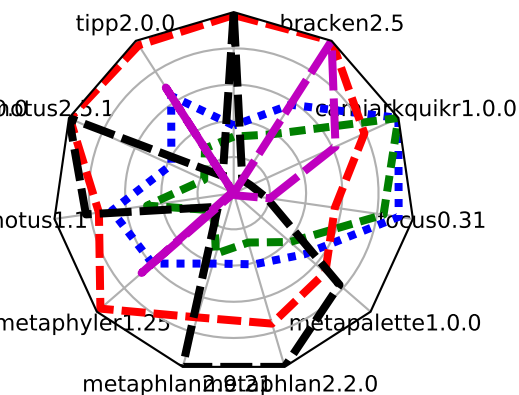
### class

sourmash



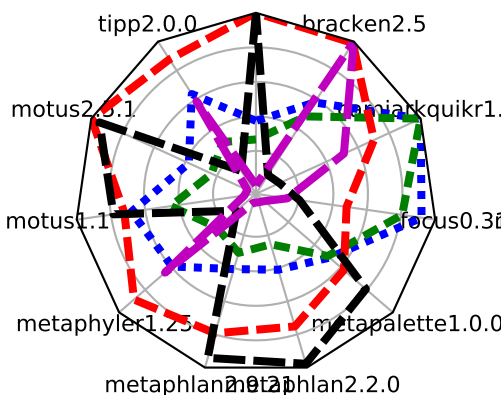
### order

sourmash



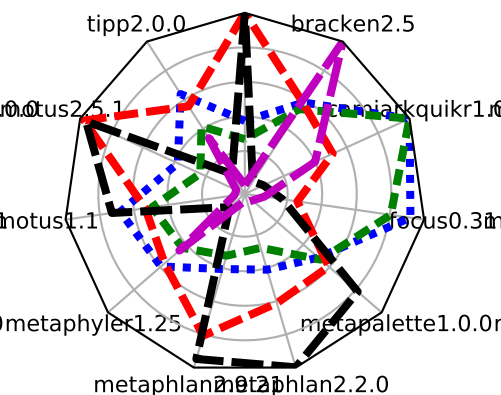
### family

sourmash



### genus

sourmash



### species

sourmash

