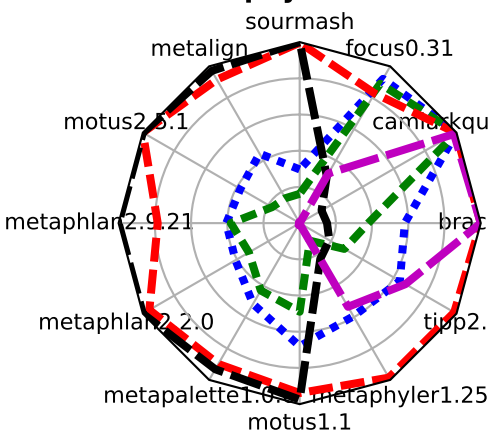
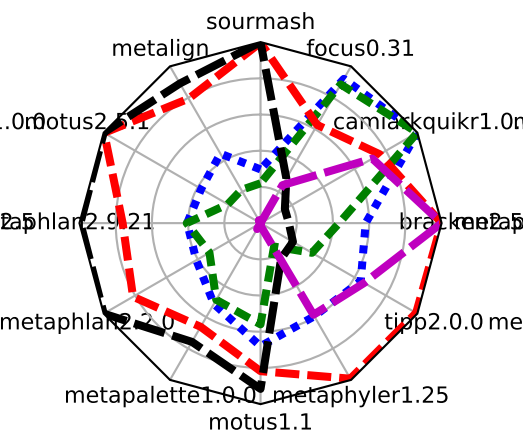


■ ■ Weighted UniFrac error    
 ■ ■ L1 norm error    
 ■ ■ Completeness (recall)    
 ■ ■ Purity (precision)    
 ■ ■ False positives

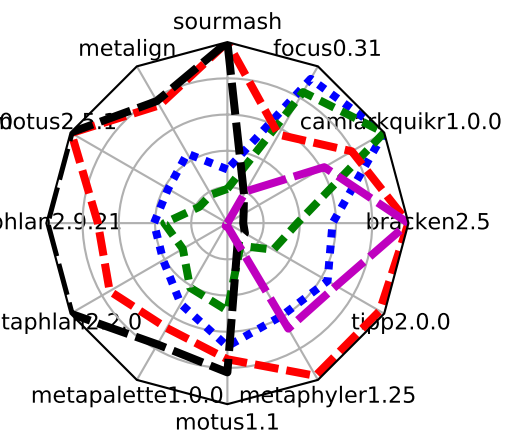
### phylum



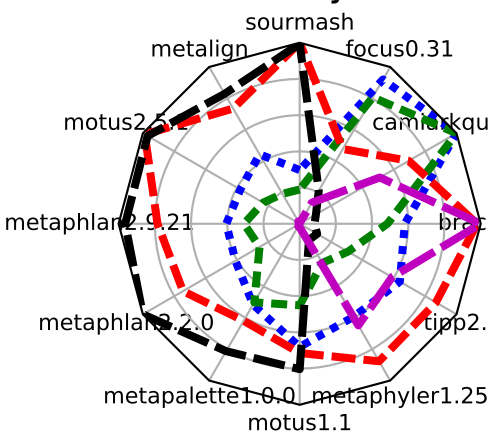
### class



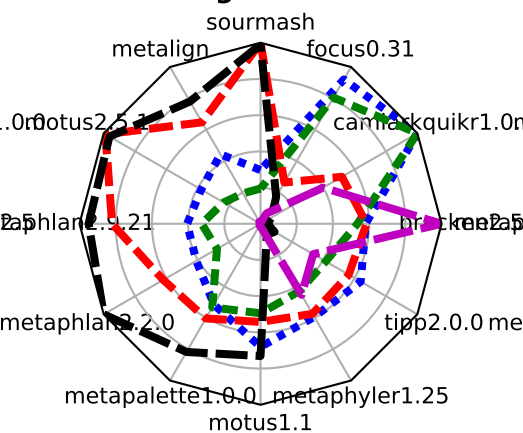
### order



### family



### genus



### species

