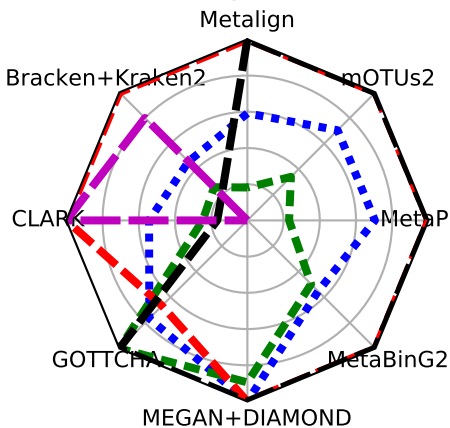
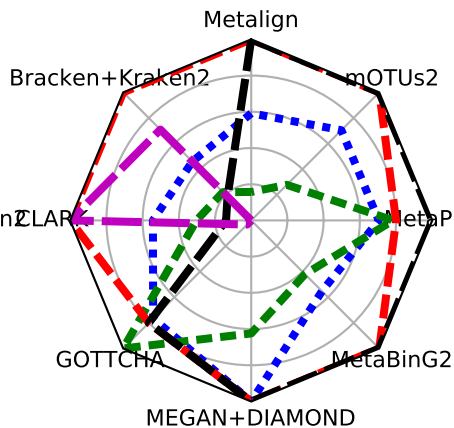


--- Weighted UniFrac error   
 --- L1 norm error   
 --- Completeness (recall)   
 --- Purity (precision)   
 --- False positives

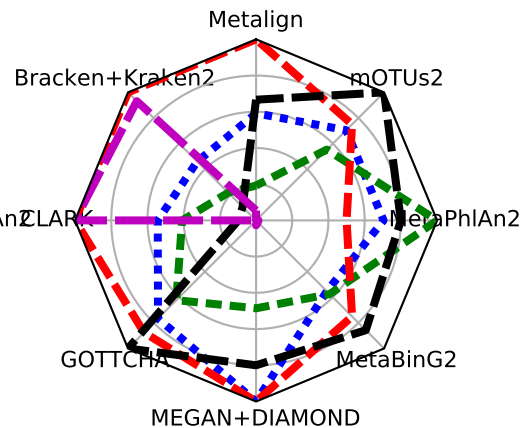
### phylum



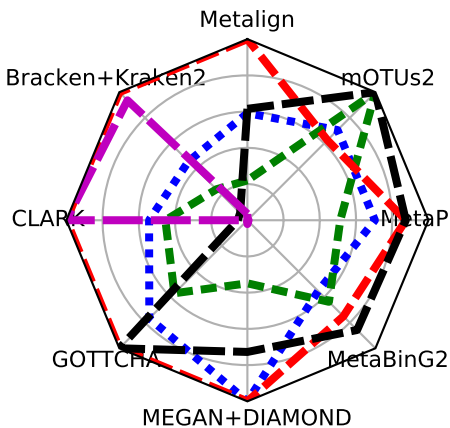
### class



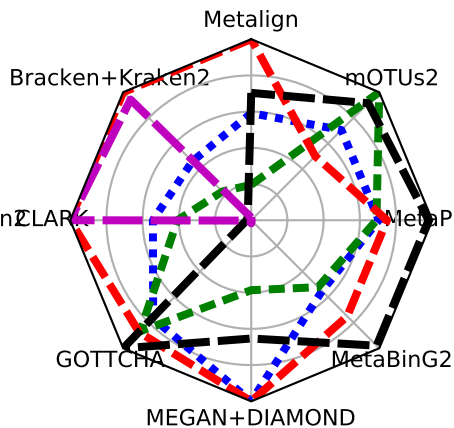
### order



### family



### genus



### species

