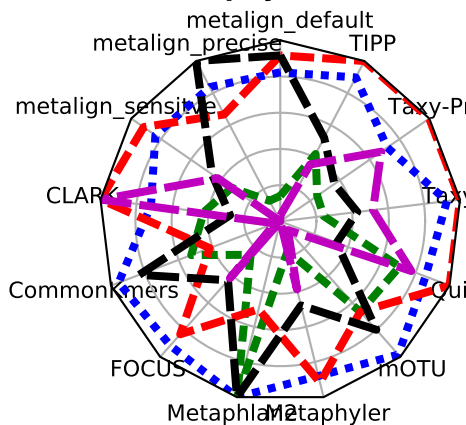
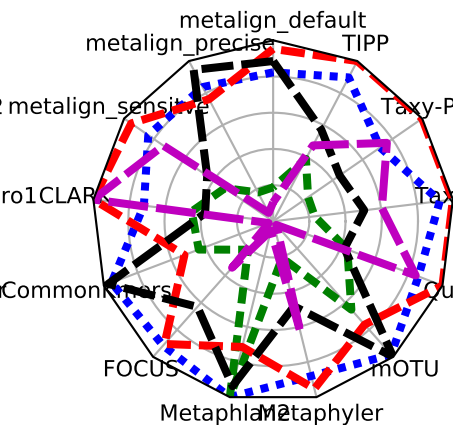


--- Weighted UniFrac error    
 --- L1 norm error    
 --- Completeness (recall)    
 --- Purity (precision)    
 --- False positives

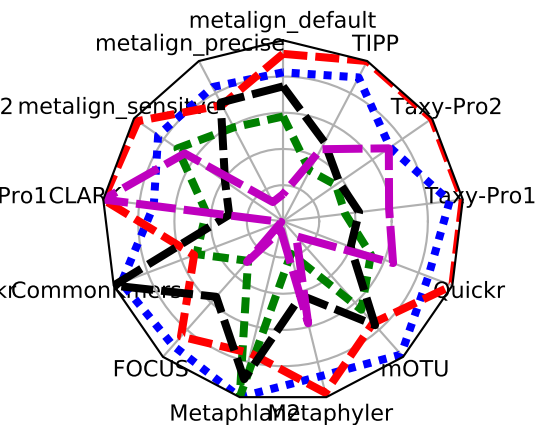
### phylum



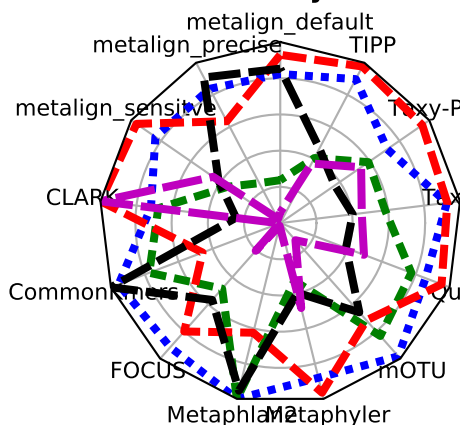
### class



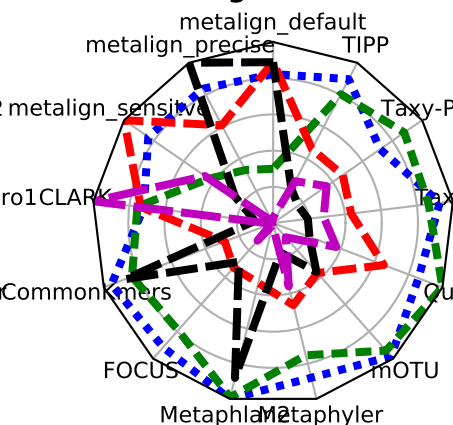
### order



### family



### genus



### species

