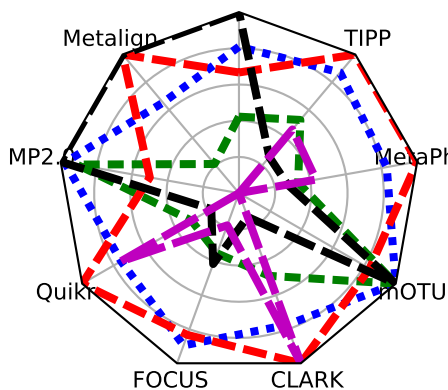


--- Weighted UniFrac error   
 --- L1 norm error   
 --- Completeness (recall)   
 --- Purity (precision)   
 --- False positives

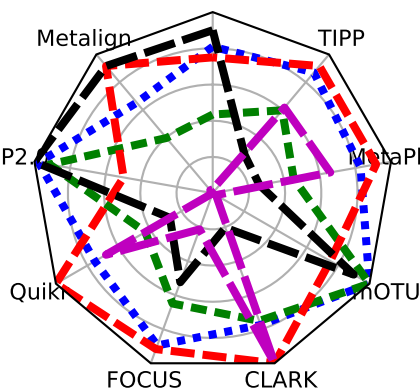
### phylum

sourmash



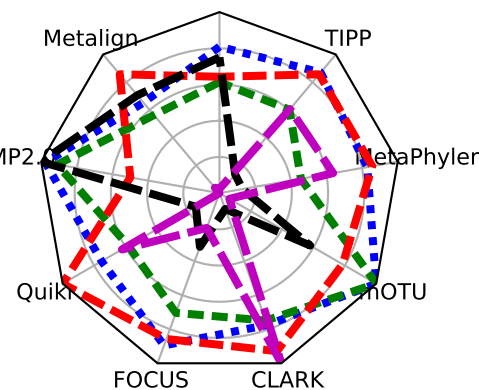
### class

sourmash



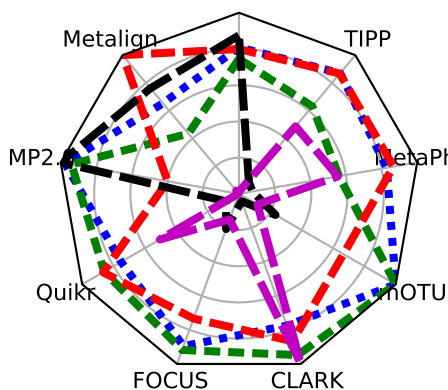
### order

sourmash



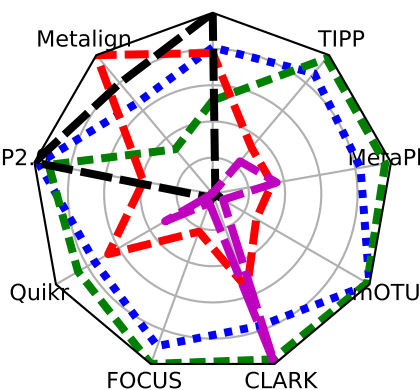
### family

sourmash



### genus

sourmash



### species

sourmash

