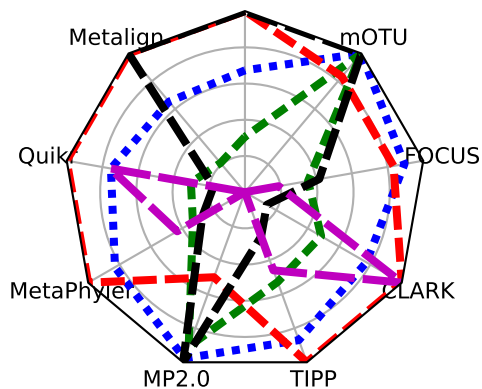


--- Weighted UniFrac error
 --- L1 norm error
 --- Completeness (recall)
 --- Purity (precision)
 --- False positives

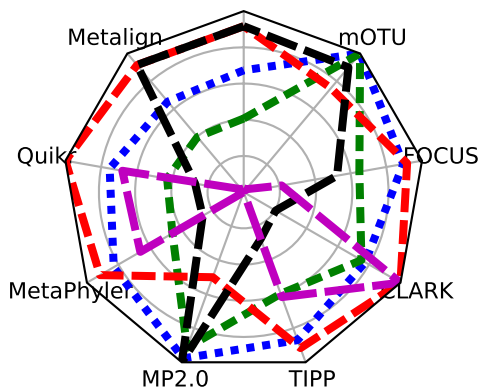
phylum

sourmash



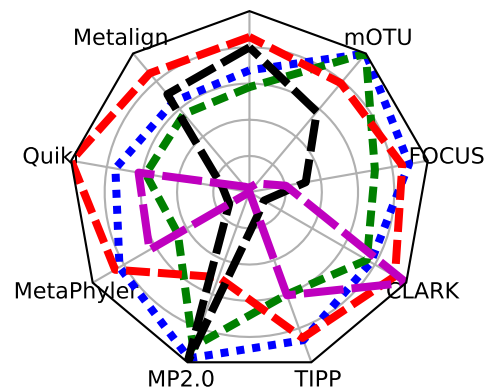
class

sourmash



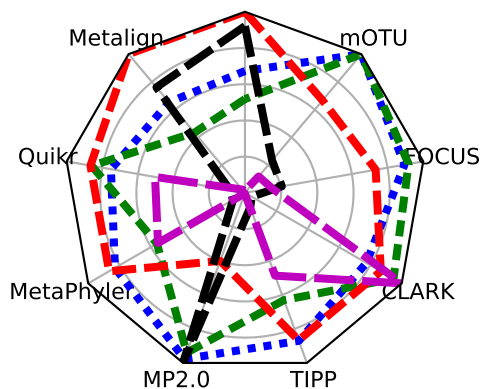
order

sourmash



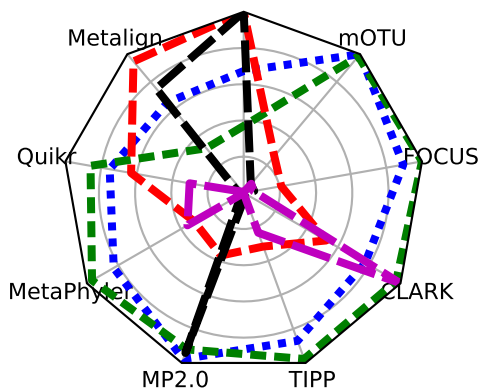
family

sourmash



genus

sourmash



species

sourmash

