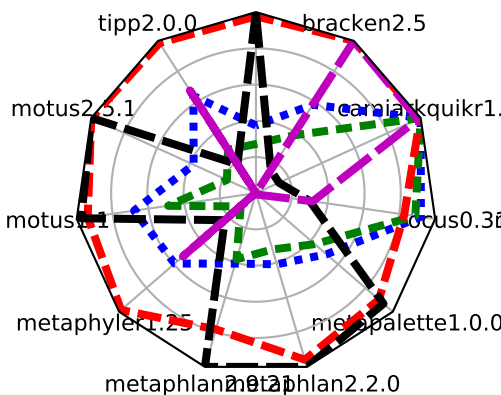


■ ■ Weighted UniFrac error
 ■ ■ L1 norm error
 ■ ■ Completeness (recall)
 ■ ■ Purity (precision)
 ■ ■ False positives

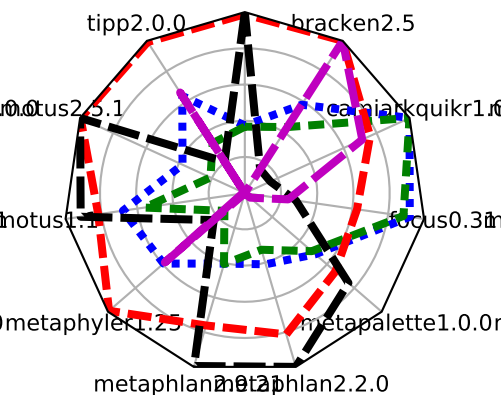
phylum

sourmash



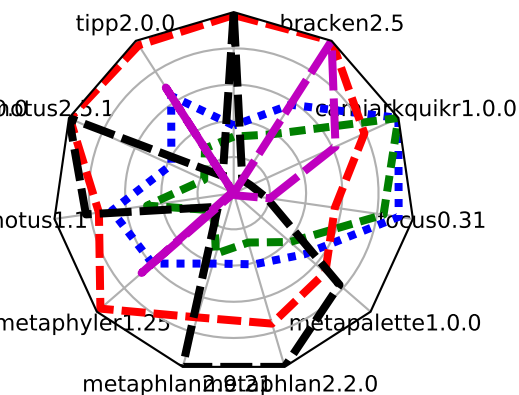
class

sourmash



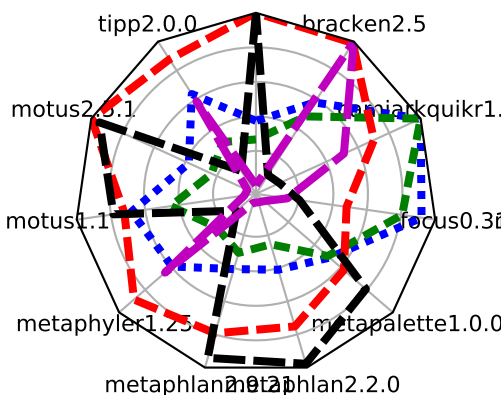
order

sourmash



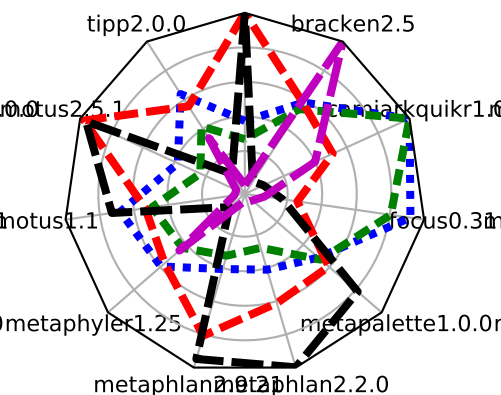
family

sourmash



genus

sourmash



species

sourmash

