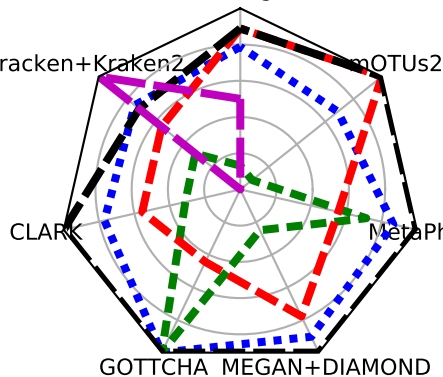


--- Weighted UniFrac error
 --- L1 norm error
 --- Completeness (recall)
 --- Purity (precision)
 --- False positives

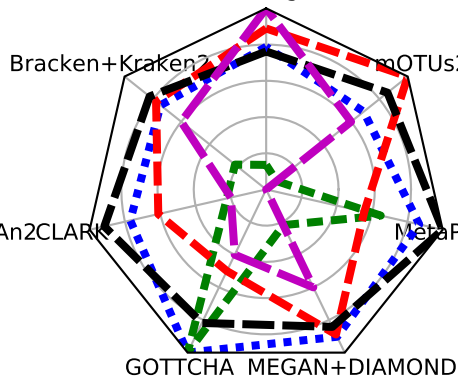
phylum

Metalign



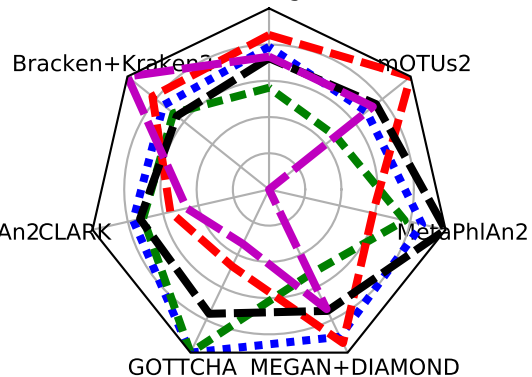
class

Metalign



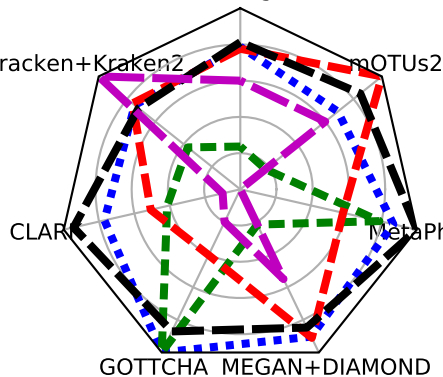
order

Metalign



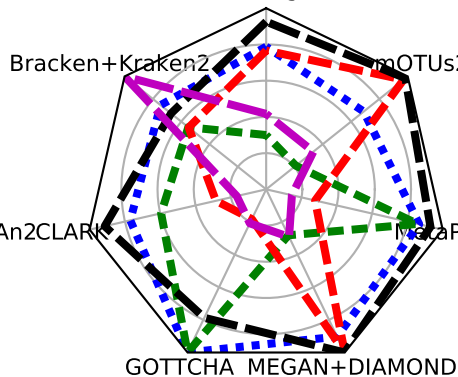
family

Metalign



genus

Metalign



species

Metalign

