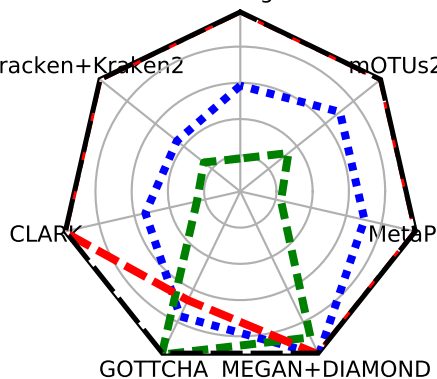


■ ■ ■ Weighted UniFrac error
 ■ ■ ■ L1 norm error
 ■ ■ ■ Completeness (recall)
 ■ Purity (precision)
 ■ ■ ■ False positives

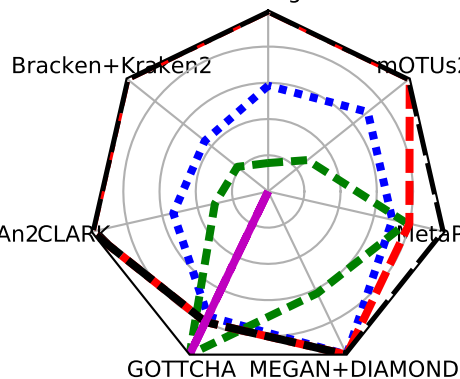
phylum

Metalign



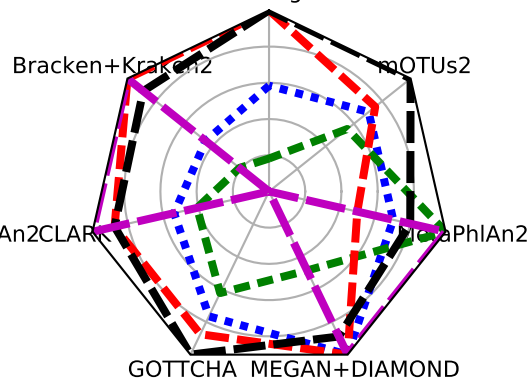
class

Metalign



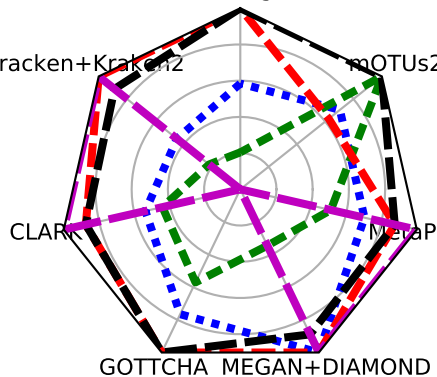
order

Metalign



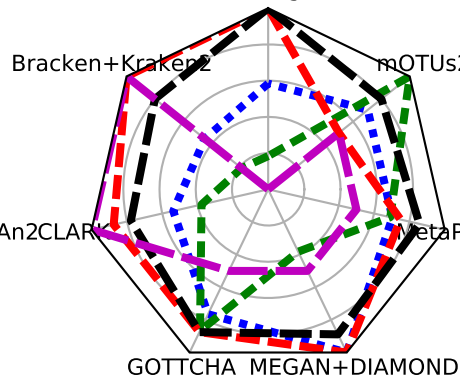
family

Metalign



genus

Metalign



species

Metalign

