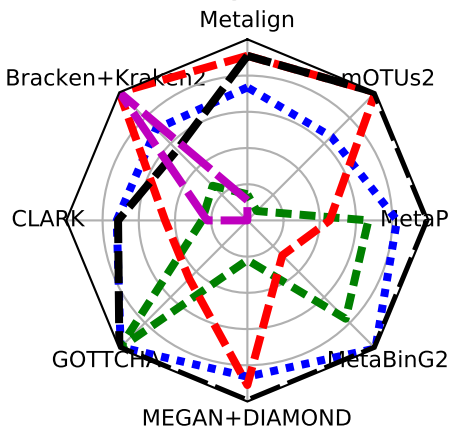
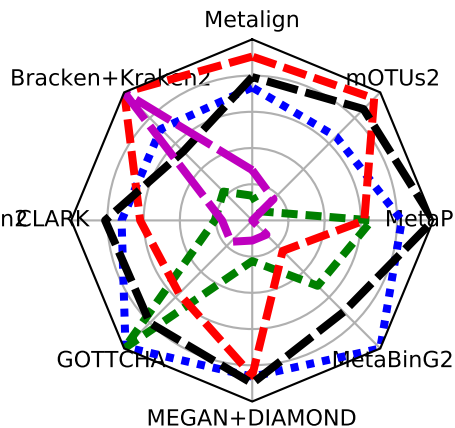


--- Weighted UniFrac error
 --- L1 norm error
 --- Completeness (recall)
 --- Purity (precision)
 --- False positives

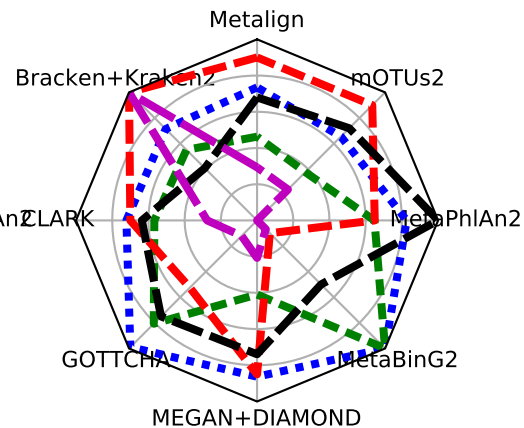
phylum



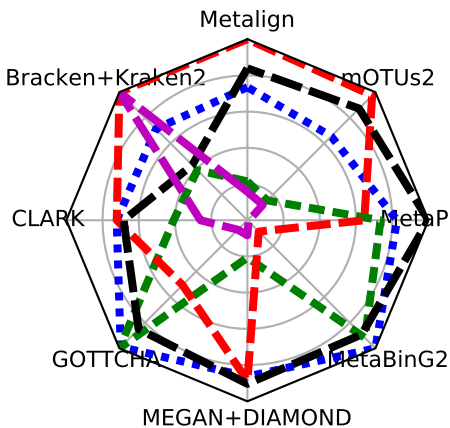
class



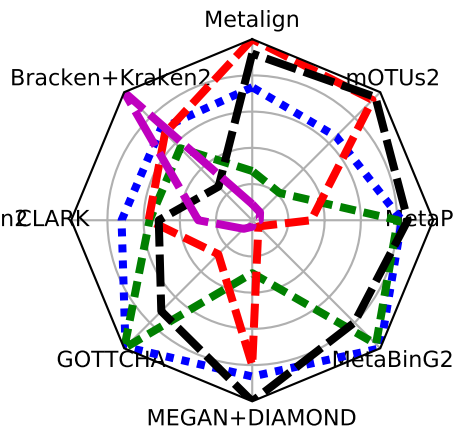
order



family



genus



species

