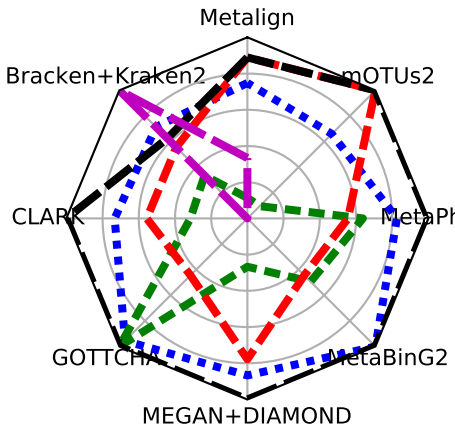
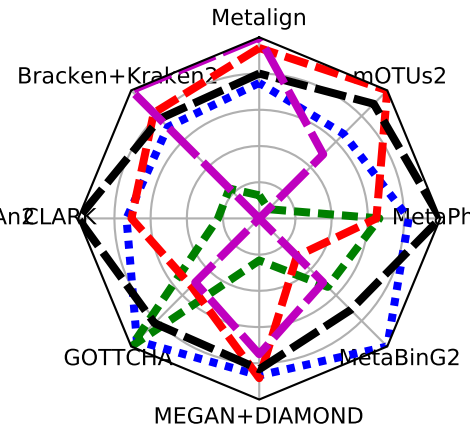


--●-- Weighted UniFrac error
 --■-- L1 norm error
 --■-- Completeness (recall)
 --■-- Purity (precision)
 --■-- False positives

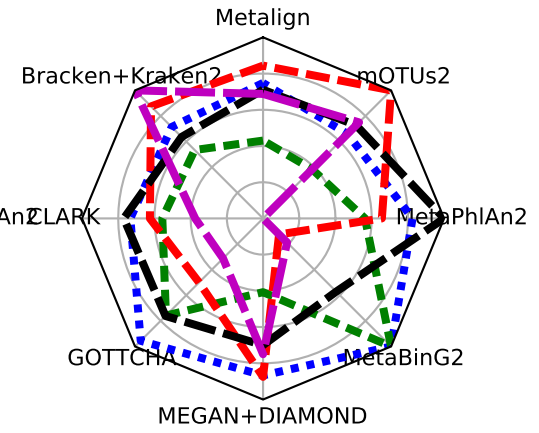
phylum



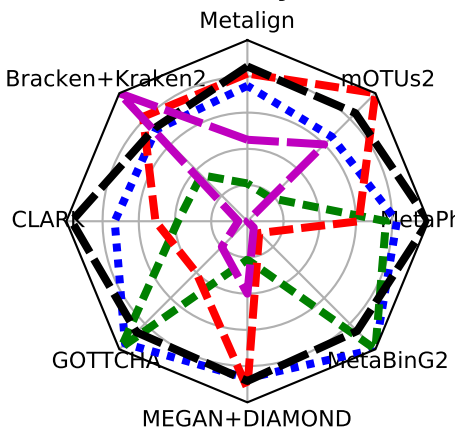
class



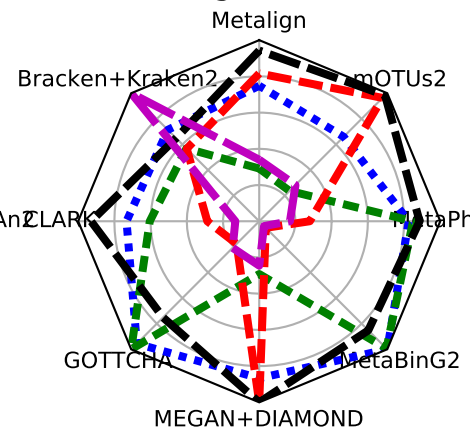
order



family



genus



species

