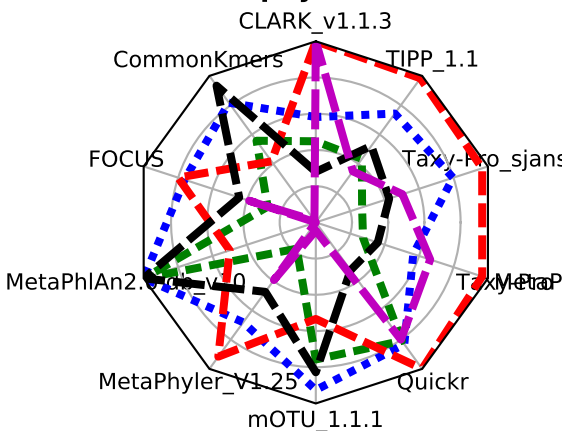
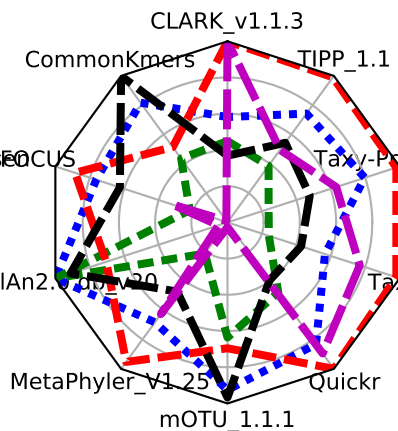


■ Weighted UniFrac error
 ■ L1 norm error
 ■ Completeness (recall)
 ■ Purity (precision)
 ■ False positives

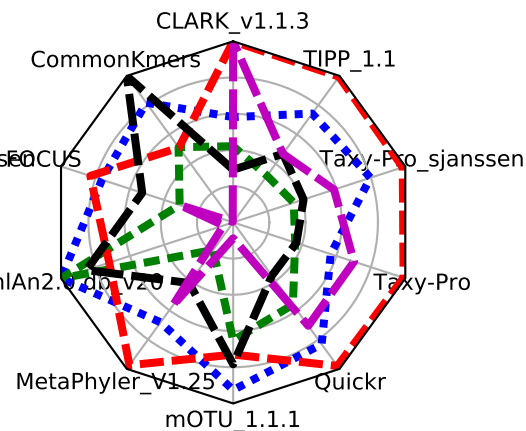
phylum



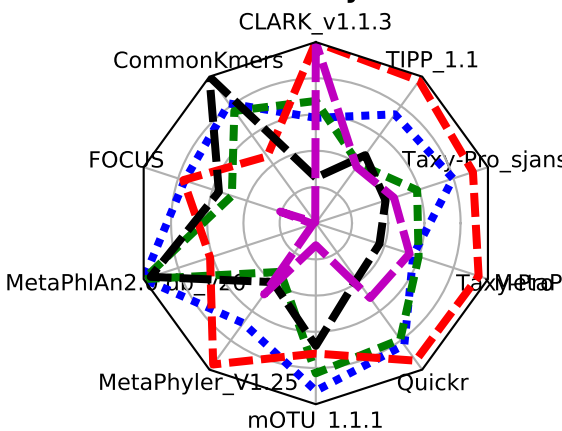
class



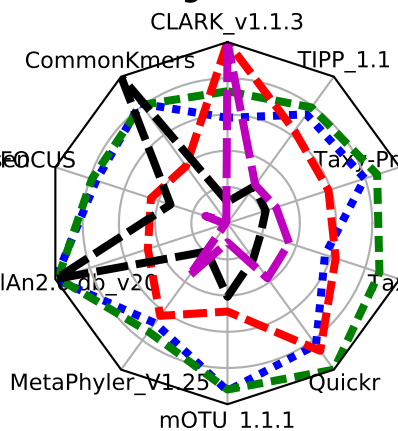
order



family



genus



species

