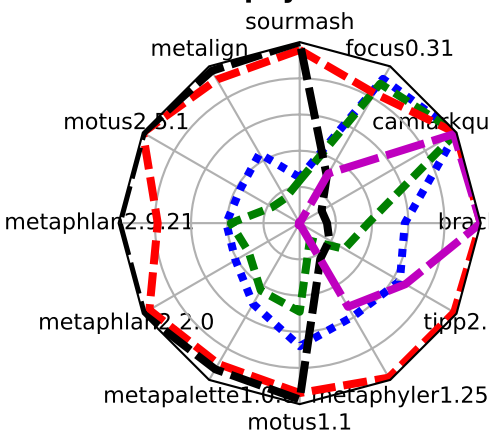
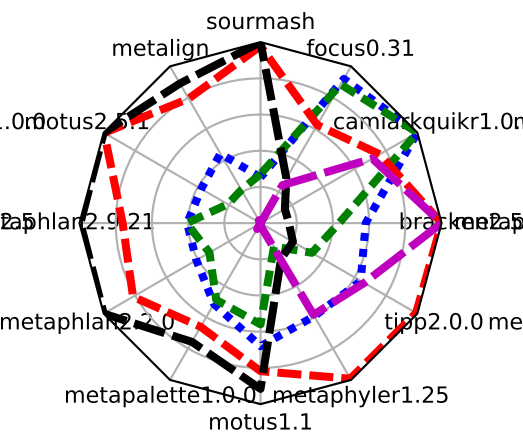


■ ■ Weighted UniFrac error
 ■ ■ L1 norm error
 ■ ■ Completeness (recall)
 ■ ■ Purity (precision)
 ■ ■ False positives

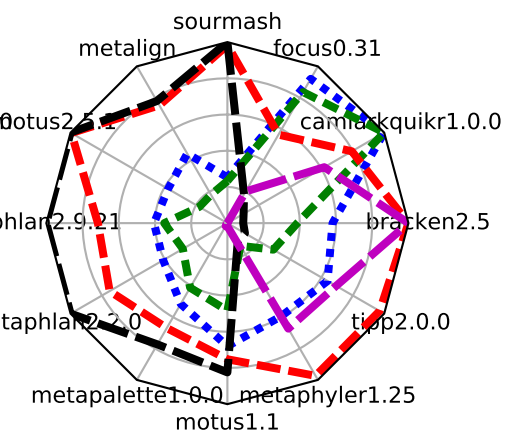
phylum



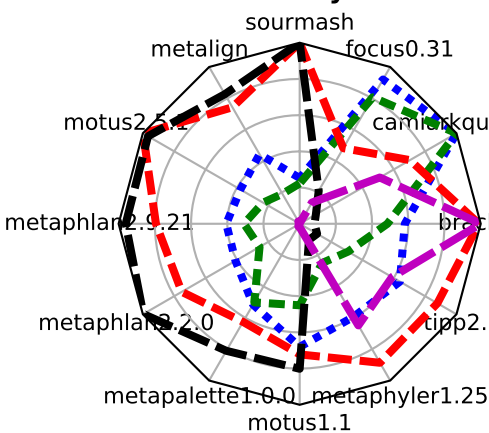
class



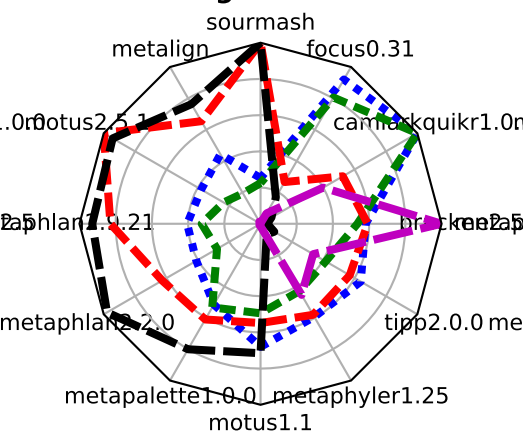
order



family



genus



species

