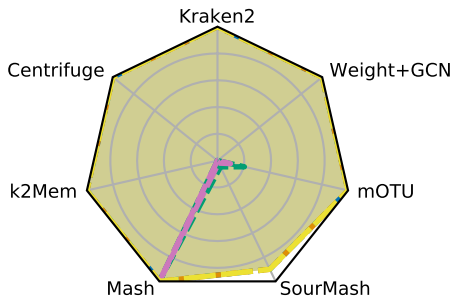
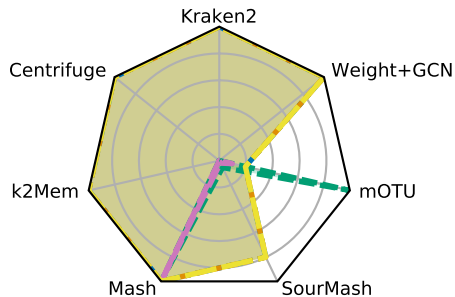


..... Completeness
 - - - - Purity
 - - - - L1 norm error
 - - - - Weighted UniFrac error
 - - - - False positives
 - - - - True positives

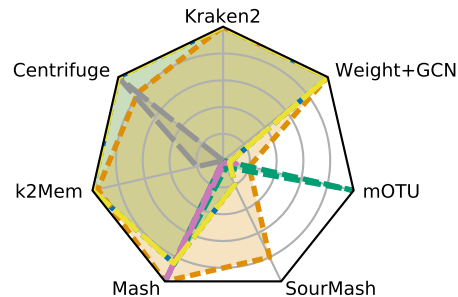
phylum



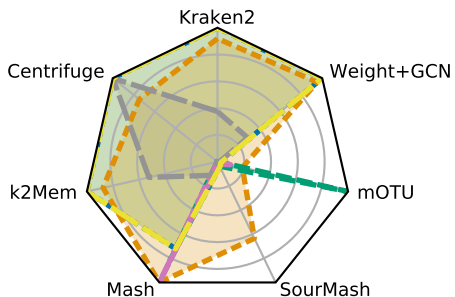
class



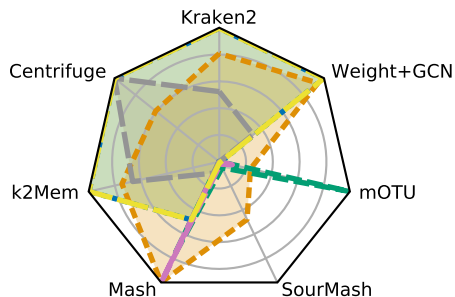
order



family



genus



species

