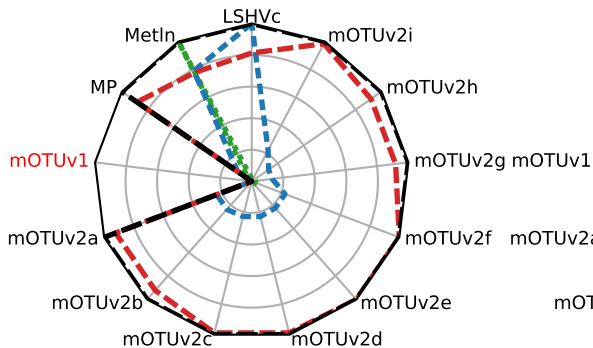
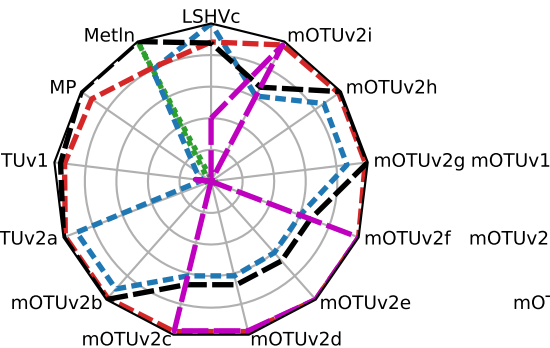


..... Unweighted UniFrac error
 - - - L1 norm error
 - - - Completeness
 - - - Purity
 - - - False positives

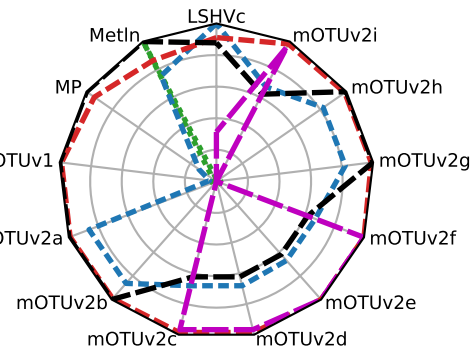
phylum



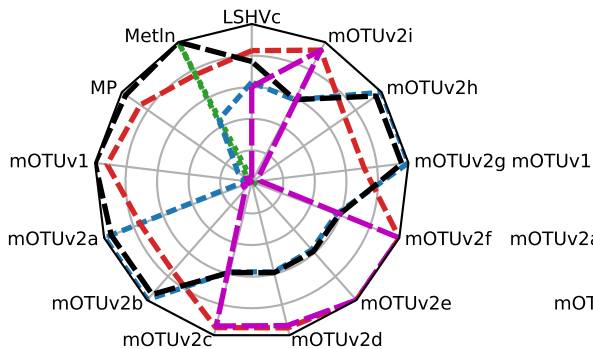
class



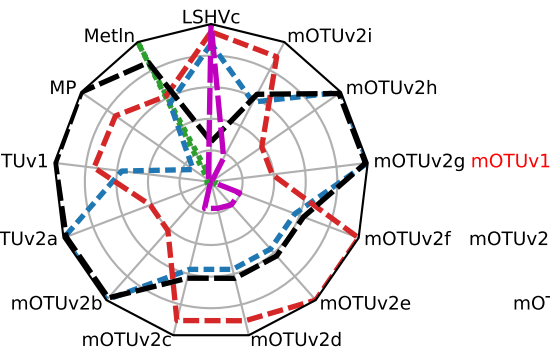
order



family



genus



species

