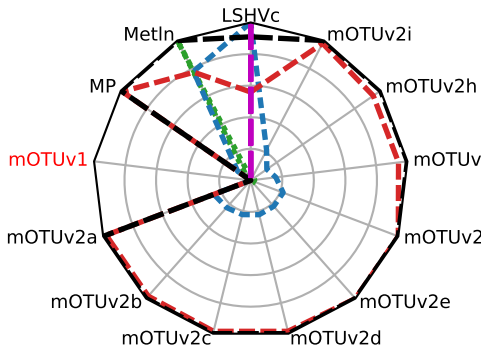
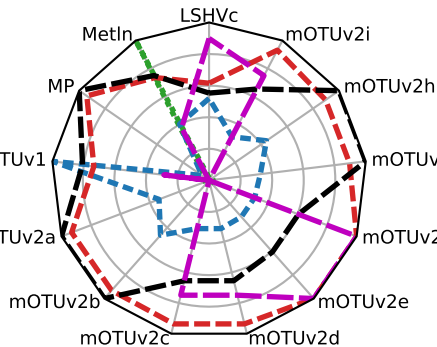


..... Unweighted UniFrac error   
 - - - L1 norm error   
 - - - Completeness   
 - - - Purity   
 - - - False positives

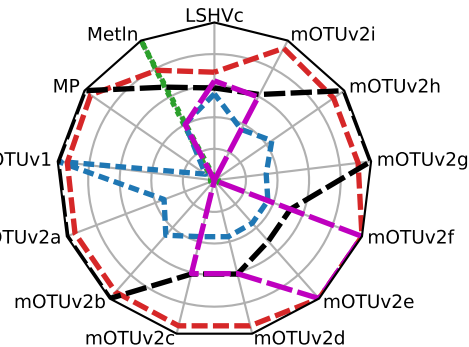
### phylum



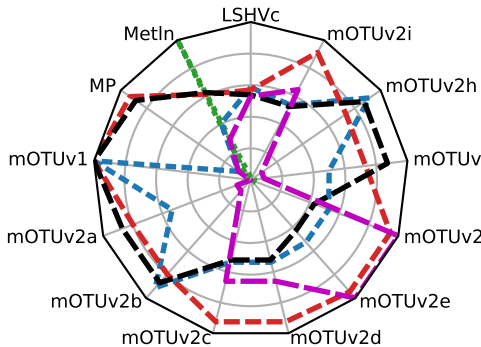
### class



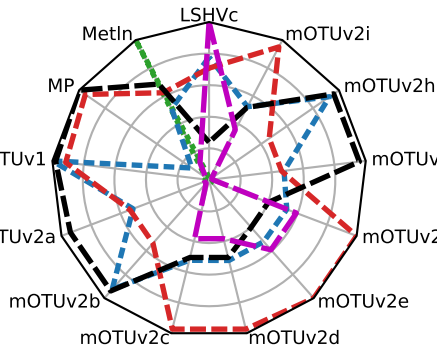
### order



### family



### genus



### species

