

Unweighted UniFrac error

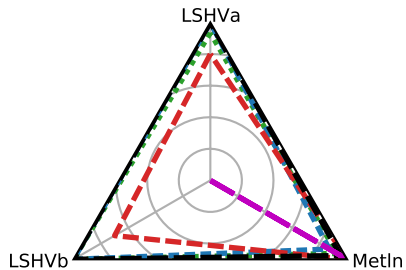
L1 norm error

Completeness

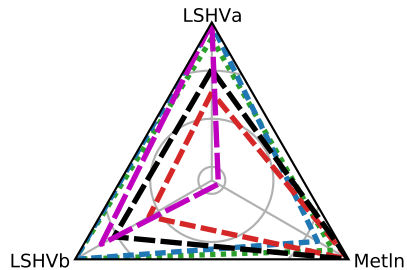
Purity

False positives

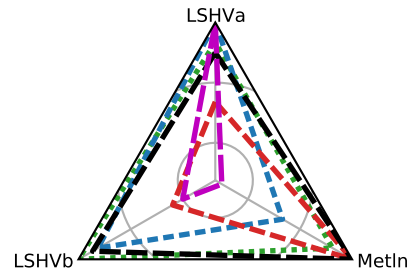
### phylum



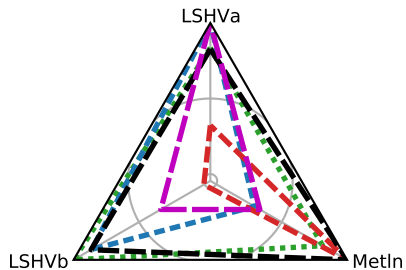
### class



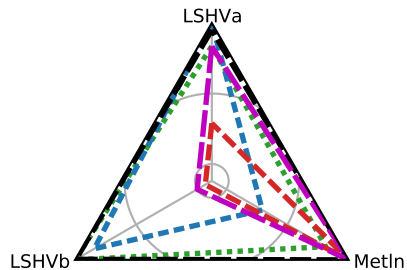
### order



### family



### genus



### species

