

Unweighted UniFrac error

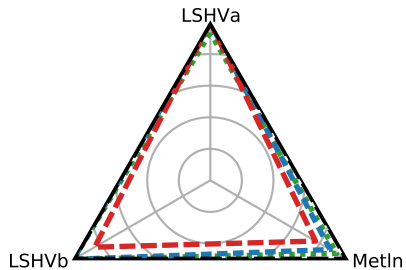
L1 norm error

Completeness

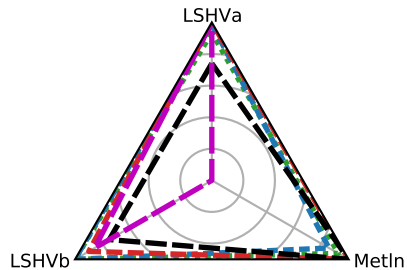
Purity

False positives

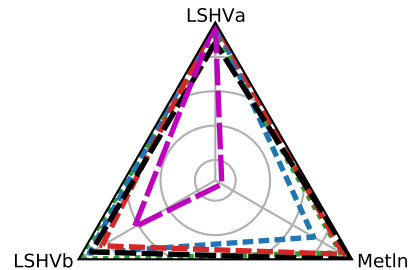
phylum



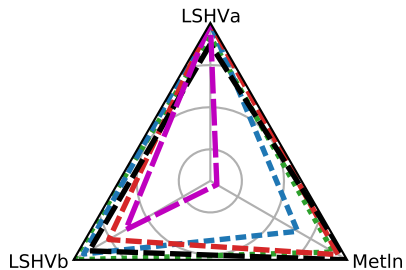
class



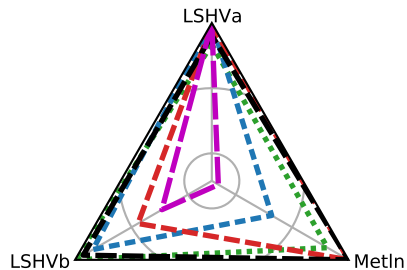
order



family



genus



species

