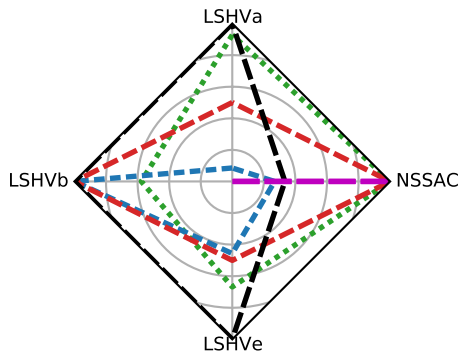
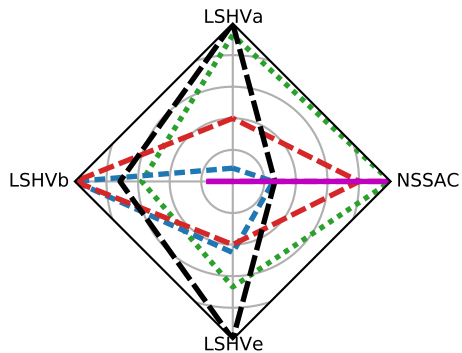


Unweighted UniFrac error L1 norm error Completeness Purity False positives

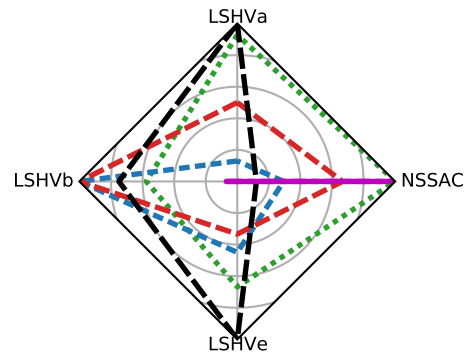
phylum



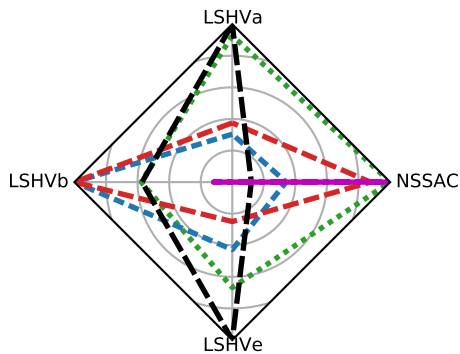
class



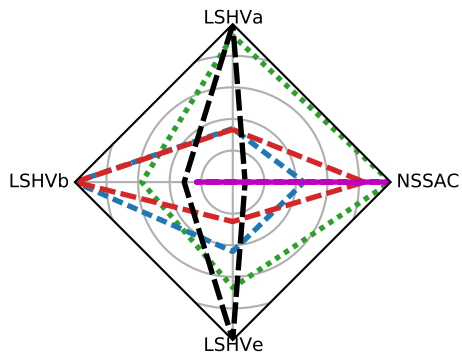
order



family



genus



species

