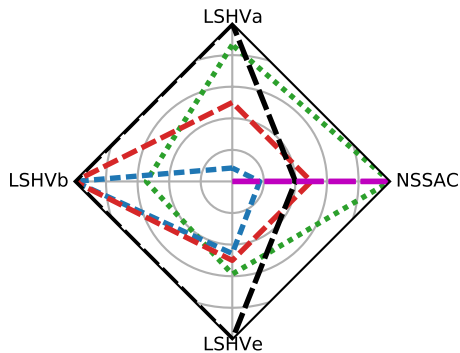
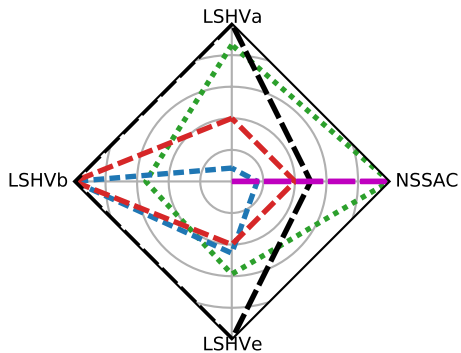


--- Unweighted UniFrac error
 --- L1 norm error
 --- Completeness
 --- Purity
 --- False positives

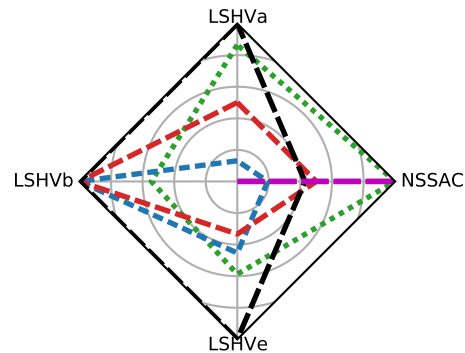
phylum



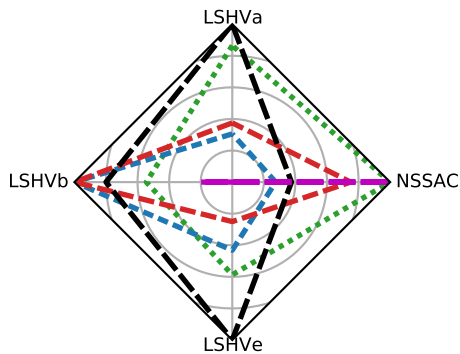
class



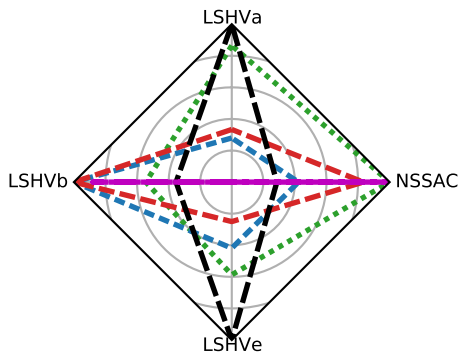
order



family



genus



species

