

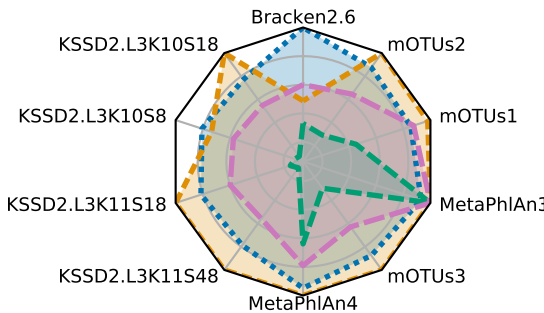
..... Completeness

----- Purity

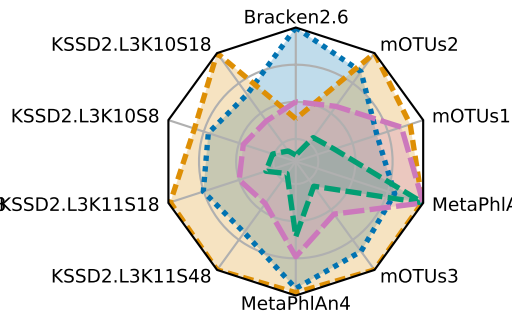
----- L1 norm error

----- Weighted UniFrac error

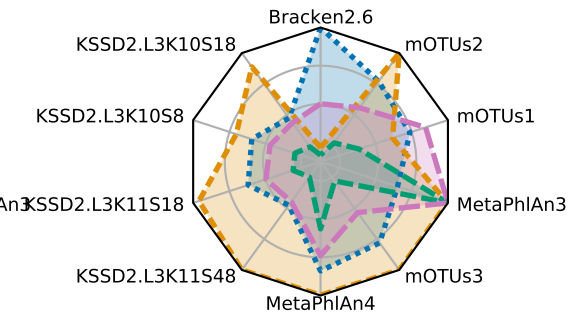
phylum



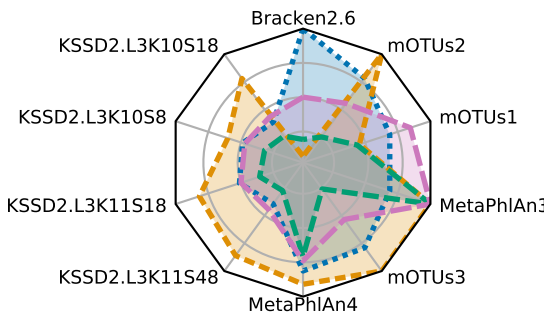
class



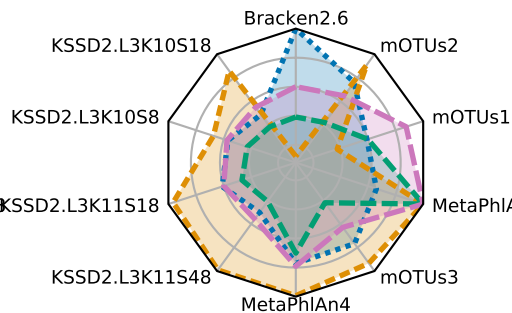
order



family



genus



species

