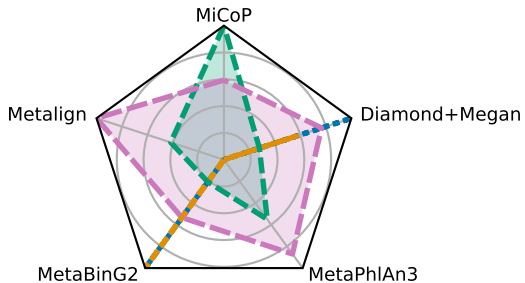
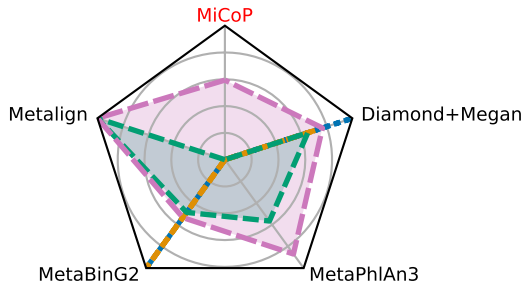


..... Completeness - - - Purity - - - L1 norm error - - - Weighted UniFrac error

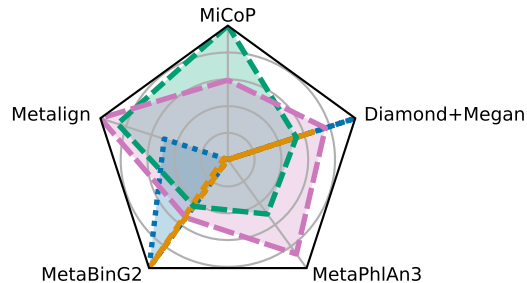
phylum



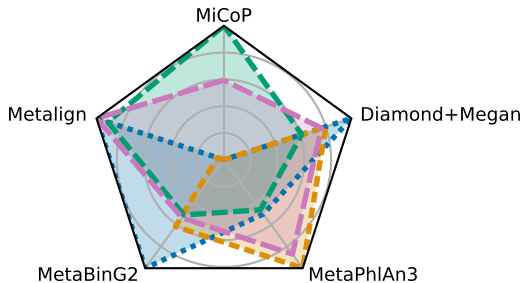
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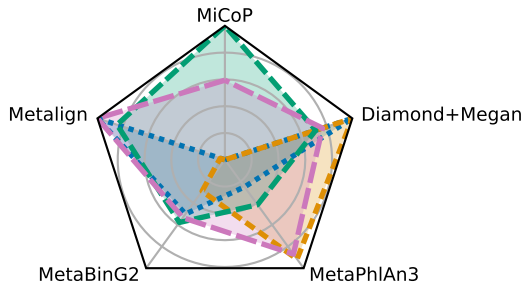
order



family



genus



species

