



Figure S2. Likelihood ratios of IS, CN, co-evolution, and sequence conservation scores. Likelihood ratio statistically evaluate how well different evolutionary approaches (IS, CN, co-evolution, and sequence conservation score) discriminate central cavity residues from non-cavity residues for each of the following percentile groups. Red, green, blue, and yellow indicate the average likelihood ratio of IS, CN, co-evolution, and sequence conservation in the given percentile cutoff, respectively.