



Figure S1 Power-law degree distributions of Human PPI networks reconstructed from HIPPIE with confidence score threshold of 0.68 (A) and 0.77 (B). The x-axis indicates the degrees (numbers of interactions) of genes in log10-based transformation, and the y-axis shows the percentage of genes (log10-based) with the corresponding degree. The red lines exhibit the line of the best fit for these distributions, with degree exponents of 1.49 and 1.67, respectively.