

A Computational Approach inspired by Simulated Annealing to study the Stability of Protein Interaction Networks in Cancer and Neurological Disorders

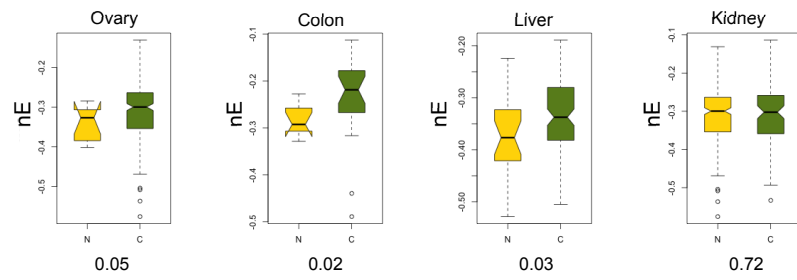
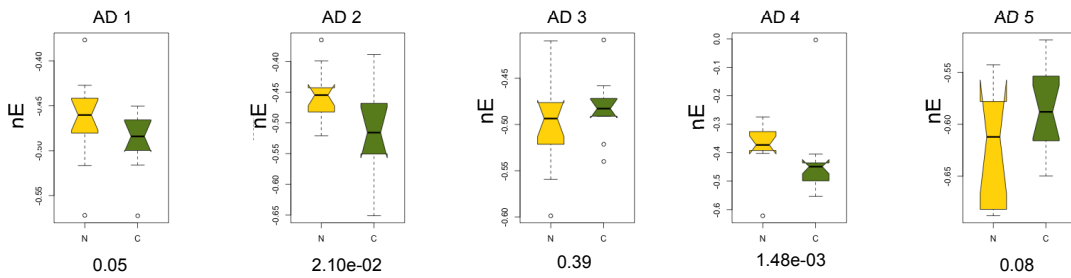
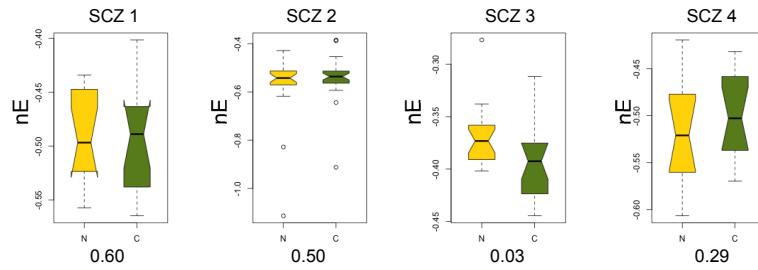
Data Mining and Knowledge Discovery

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Online Resource 4 nE distribution mapping all the genes in **(a)** normal (N) and cancer (C) states (Ovarian, Colon, Liver and Kidney). **(b)** Normal (N) and AD (C). **(c)** Normal (N) and SCZ disease (C) state in the HPRD network. The Wilcoxon-rank p-value is presented below the x-axis