Correlation between functional annotation and topology of protein interaction network

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Abstract

The functional annotation of proteins was believed to be related to the topology of

protein-protein interaction network. People utilized the protein-protein interaction network to infer the protein function by various methods. Here we select the protein interaction data of Saccharomyces cerevisia and

calculated the correlation between functional annotation of proteins and topology of protein-protein interaction network. The result shows that the functional correlation decays exponentially with the distance between two proteins,

and beyond the characteristic distance it has no correlation.

Keyword: protein-protein interaction; network; correlation.