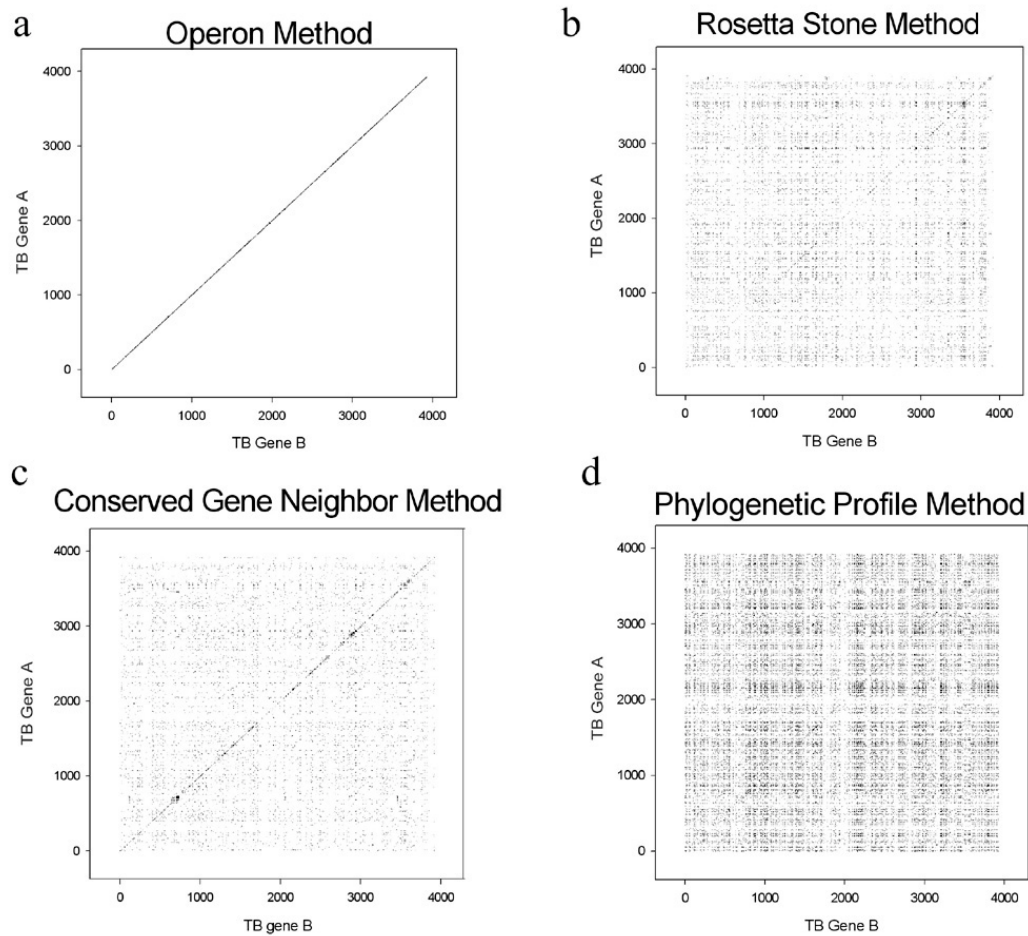


Supplementary Material



Supplementary Material Figure 1. Genome-wide functional linkage maps representing functional linkages inferred by each of the individual computational methods. The genome-wide functional linkage maps depicted here represent functional linkages inferred by each of the individual computational methods. **(a)** Operon method (100bp threshold). **(b)** Rosetta Stone method. **(c)** Conserved Gene Neighbor method. **(d)** Phylogenetic Profile method