High-Coverage Whole-Exome Sequencing Identifies Candidate Genes for Suicide in Victims with Major Depressive Disorder

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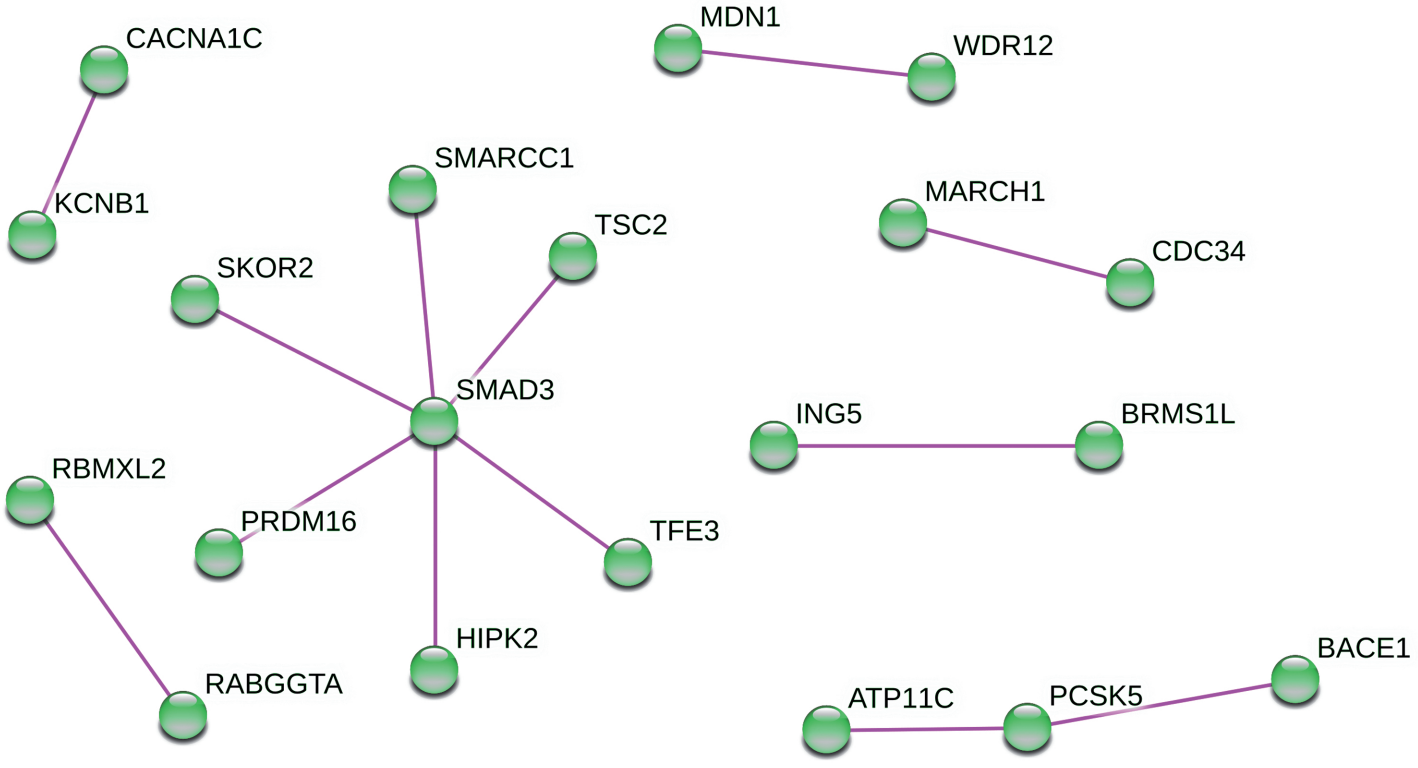
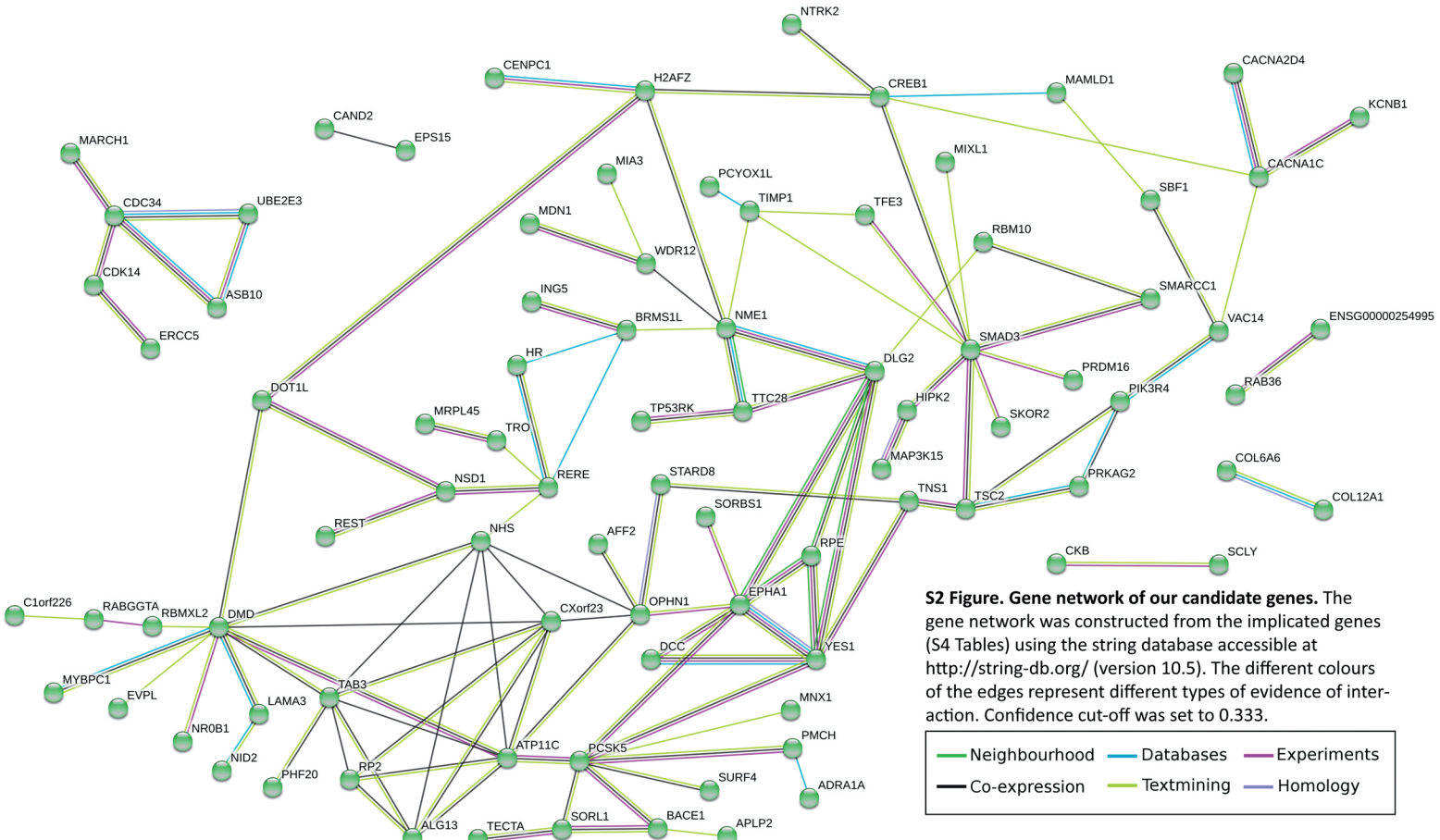


S1 Figure. Gene network of candidate genes based on experimental evidence of interaction. The gene network was constructed from all the implicated genes (S4 Table) using the string database accessible at http://string-db.org/ (version 10.5). All edges represent experimental evidence of interaction. Confidence cut-off was set to 0.333.



S2 Figure. Gene network of our candidate genes. The gene network was constructed from the implicated genes (S4 Tables) using the string database accessible at http://string-db.org/ (version 10.5). The different colours of the edges represent different types of evidence of interaction. Confidence cut-off was set to 0.333.

Neighbourhood · Databases · Experiments · Co-expression · Textmining · Homology