

Figure 5. Candidate genes, pathways and mechanisms.

(A) Top Candidate Genes and relationships to DARPP-32. All the Category I genes, all the genes with 5 out of 6 lines of evidence, and the genes with 4 out of 6 lines of evidence that have a known relationship with DARPP-32 are illustrated. Genes in plain text are increased by methamphetamine, and in genes in italics are decreased by methamphetamine.

(B) Gene Ontology (GO) analysis-derived model of biological processes and mechanisms. Numbered categories refer to GO analysis categories from Table 4.

See next page.



