Figure 3. Category I and II candidate genes probability. (A) Pyramid generated by empirical tabulation of independent
converging lines of evidence. Plain text - increased by methamphetamine. Italics – decreased by methamphetamine
(B) Comparison of different target brain regions in terms of average number of lines of evidence per candidate gene.

A

5/6: TAC1 PENK DARPP-32 *MEF2C* *TBR1 CCK* GLUL

4/6: *CDK5R1* LYPLA1 MOBP SYT1 CLDN11 *CAMKK2*

3/6: GPR88 PITPNB RAB7 ADAM10 TMEFF1 *BTBD3* GAT3 *SRRM2*

2/6: MT1 *HIP14 CLSTN1* RNF138 SSR3 UGT8 CNOT7 *NCALD NPTX1* PRMT1 ROXAN *PITX2 MEG3*

EST (AA921069) GORASP2 PSME1 *DLC2* EST (D10627) *EST (AI844183) EST(AI853550)*

B