Mouse Models of Mutations and Variations in Autism Spectrum Disorder-Associated Genes: Mice Expressing *Caps2/Cadps2* Copy Number and Alternative Splicing Variants

**Figure S1.** A predicted ASD-associated gene network. The network was predicted by the basic network weighting method of GeneMANIA [1] using information on co-localization, pathways, physical interactions and predicted relationships [2]. CADPS2 is highlighted by a red circle. ASD-associated genes are indicated by large black circles. Genes indicated by small gray circles are genes relevant to or functionally interacting with ASD-associated genes.
Reference


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