



Supplemental Figure 4. Colocalization Diagnostic Plots. (A, B) Genome wide association study (GWAS) of schizophrenia (3) sites were plotted against transcript quantitative trait loci (tQTL; PsychEncode; 73) sites for two *MAD1L1* transcripts (yellow) and (C) one *MRM2* transcript (blue). (D) Z-Score imputation for *MRM2* transcript.