



Supplemental Figure 2A. *FERMT2* LOAD GWAS locus. Genome browser view of SSV deletion variant (rs1277882435) disruption of the *TAL1* transcription factor. Tracks include (upper to lower): gene structure (GENCODE V41); candidate cCCREs (ENCODE) TFs from the JASPAR core collection; and SSVs from short genetic variants (dbSNP153). The deletion disrupts the *TAL1* TF. Displayed genomic region includes the GWAS SNP (rs17125924) and the indel variant (rs1277882435) located in the enhancer element (E1714539).

