Figure S6  Human *DPYSL2* 3'-flanking cNCR (3P-cNCR) construct containing a SZ-associated SNP rs73229635 (p-value<10^{-5} in large AJ collection) showed enhancer function and difference in driving luciferase expression between constructs containing Wt allele and Risk allele: A) A construct (blue bar) containing SNP rs73229635 in 3'-flanking (red thin line) shown on the UCSC genome browser; B) 3P-cNCR luciferase assay results in cells