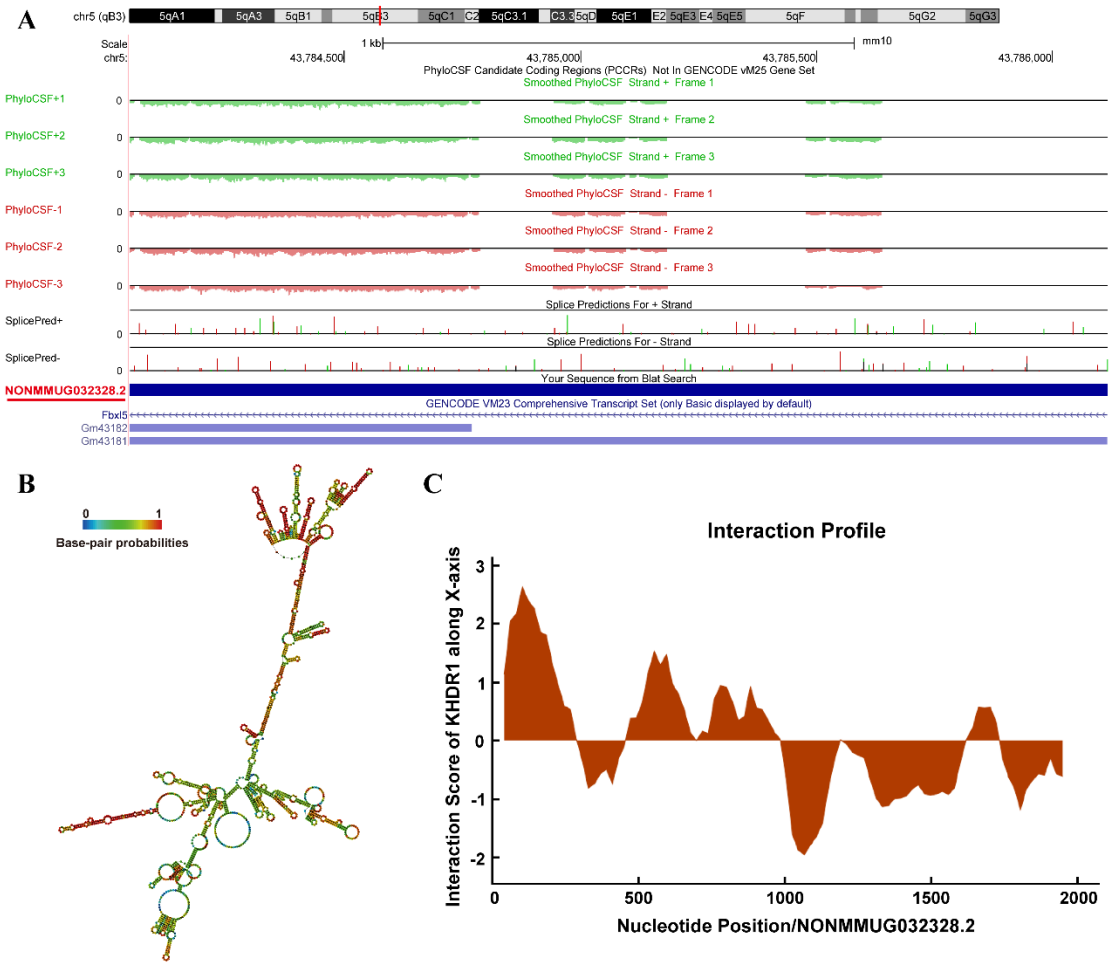



**Figure S4.** Bioinformatics analysis of NONMMUG032328.2. (A) The genomic location and coding potential of NONMMUG032328.2 in the mouse genome were characterized. (B) Secondary structure prediction of NONMMUG032328.2. Shown is an optimal minimum free energy structure (MFE= -611.70 kcal/mol) (C) KHDR1 showed strong binding to NONMMUG032328.2.