

Figure S1: The state of the LD of the SNPs located between the GWAS line and suggestive line (1×10^{-8} - 1×10^{-4}) on chromosome 2, based on opioid use disorder patients compared to controls from the UAE population. Block 1, 2 mapped to the *KIAA1211L* gene and is significantly associated with opioid use disorder in this cohort.

