**Description about INPUT files**

**Input file name:** miRNA\_RBP\_binding\_location\_for\_distance\_calculation.csv

Please see the description of the OUTPUT files of the Perl program which calculate the position of the miRNA/RBP binding sequence.

Input file name: m6A\_modifiaction\_location\_for\_distance\_calculation.csv

Please see the description of the OUTPUT files of the Perl program which calculate the position of the m6A modification site.

**Description about OUTPUT files**

**Name of the file:** distance\_between\_m6A\_and\_miR\_or\_RBP\_binding\_site.csv

Description of the output file’s columns are as follows:

Column 1: Name of the transcript

Column 2: The numbers of nucleotide between central adenine(A) residue of m6A site and the central nucleotide of miRNA target sites and RBP binding sites.

Column 3: Length of 3’UTR of the mRNA.