

How to use Software:

General Instruction : This software is now immature state. I shall developed it user friendly with in very short time.

1. Install Visual Studio for c compiler
2. Download the split_trna37.pdf file and split_trna36.pdf file
3. Convert .pdf to .txt
4. Change the .txt file into .c
5. Copy split_trna37.c and split_36.c separately into different folder.
6. Create input.txt file with only the sequence of the genome.
7. Open split_trna37.c and run this program .
8. Enter the range (which is the length of the tRNA e.g 72 Enter 100)
9. Similarly for split_trna36.c
10. If any problem please contact through email urmandal@gmail.com