T-REKS

Unix commands :

```
#Get help
java -jar T-Reks.jar -h

#Launch Graphical user interface (GUI)
java -jar T-Reks.jar

#Launch T-REKS with default parameter
java -jar T-Reks.jar -infile=fastaFile.fasta
```

For reliable large-scale analysis, we suggest to use T-REKS with the following multiple sequence alignment tools :

- [MUSCLE](#) (v3.6)
- [CLUSTAL W](#) (v1.83)

These programs are included in T-REKS.zip

```
##############################################################################
              Launch T-REKS with MUSCLE and CLUSTAL W algorithms.
    → Absolute path is required to launch T-REKS with MUSCLE and CLUSTAL W.
##############################################################################

Example :
java -jar T-Reks.jar -infile=file.fasta -msaMode=x -muscle=/home/user/muscle
-clustal=/home/user/clustalw
```