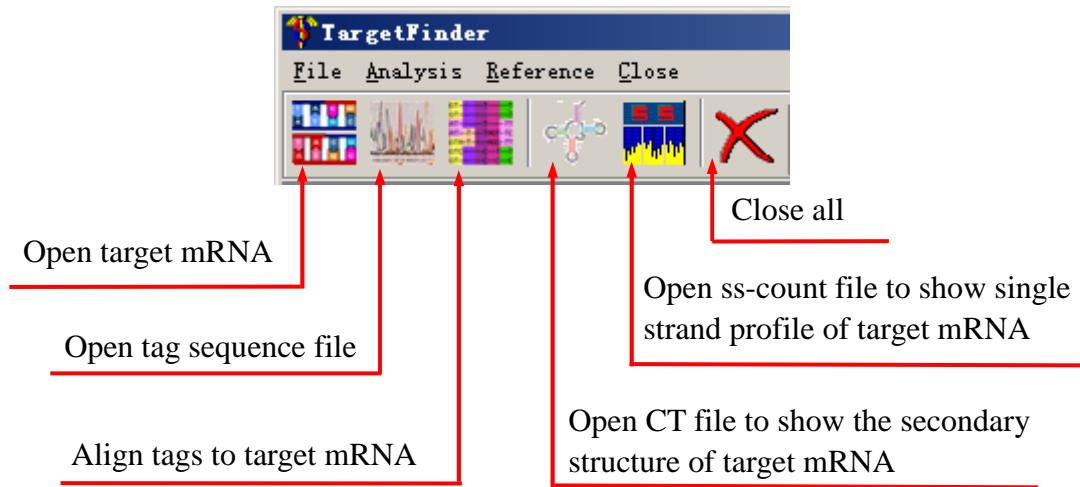


TargetFinder 1.0

1. Speed buttons



2. Sample data file


TarGene.fa — sample target mRNA sequence file in FASTA format


Tags.txt — sample tags sequence file


SS_count.txt — sample ss-count file


Structure.ct — sample secondary structure file in CT format


3. Quick tour

Step 1. Click “” button to open the sample target mRNA file “*TarGene.fa*”

Step 2. Click “” button to open the sample tags file “*Tags.txt*”

Step 3. Click “” button to align the tags to target mRNA

Step 4. Click “” button to open the sample structure file “*Structure.ct*” of target mRNA

Step 4. Click “” button to open the sample ss-count file “*SS_count.txt*” of target mRNA