EMview: A Multiple Sequence Alignment View Based on Excel

Inputs:

Sequence database search: BLAST, FASTA suites.

Multiple sequence alignment: CLUSTAL, HSSP, MSF, FASTA, PIR, MAF.

If provided VCF or other SNP file, it will be import to alignment

Outputs:

Emview, HTML, FASTA, CLUSTAL, MSF, PIR, RDB (tab-separated).

The program will run step by step

From ab1 file to Emview to other format