

Sequence Search FASTA

User Manual

Section VI.1.C: Sequence Search (p162 - p166)

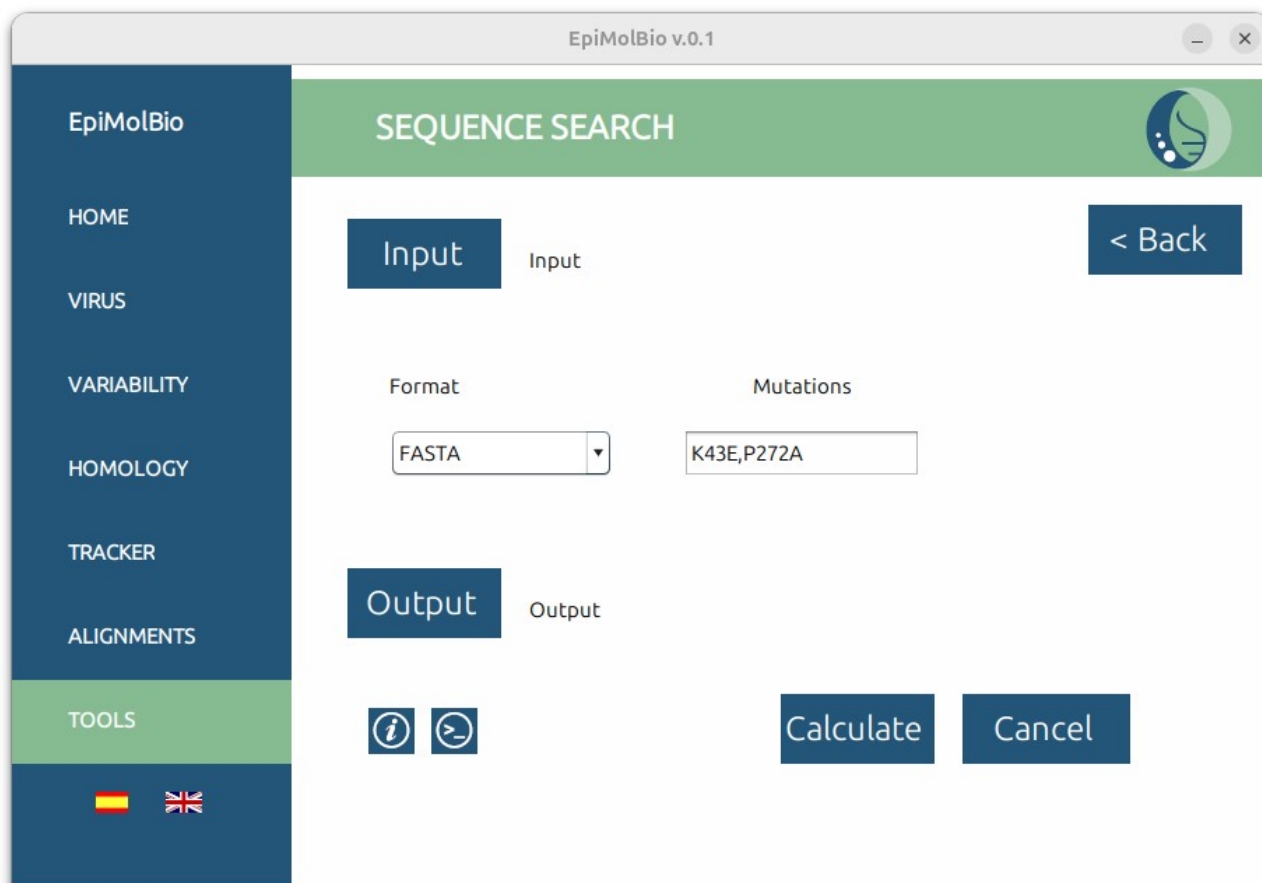
See the full manual [here](#).

Example File

File Name: SEQUENCE SEARCH FASTA RT HIV-1 AA_B.fasta, SEQUENCE SEARCH FASTA RT HIV-1 AA_C.fasta

Parameter Selection

- **Format:**
 - FASTA
- **Mutations:**
 - K43E,P272A



The screenshot shows the EpiMolBio v.0.1 web application interface. The left sidebar contains navigation links: EpiMolBio, HOME, VIRUS, VARIABILITY, HOMOLOGUE, TRACKER, ALIGNMENTS, TOOLS, and language flags (Spanish and English). The main content area is titled "SEQUENCE SEARCH" and features a "Back" button. Below the title, there are two input fields: "Format" (set to FASTA) and "Mutations" (set to K43E,P272A). At the bottom, there are "Calculate" and "Cancel" buttons, along with information and help icons.

EpiMolBio v.0.1

SEQUENCE SEARCH

Input Input

Format Mutations

FASTA K43E,P272A

Output Output

Calculate Cancel