

## Find and Replace

### User Manual

#### Section VI.1.D: Find and Replace (p167 - p171)

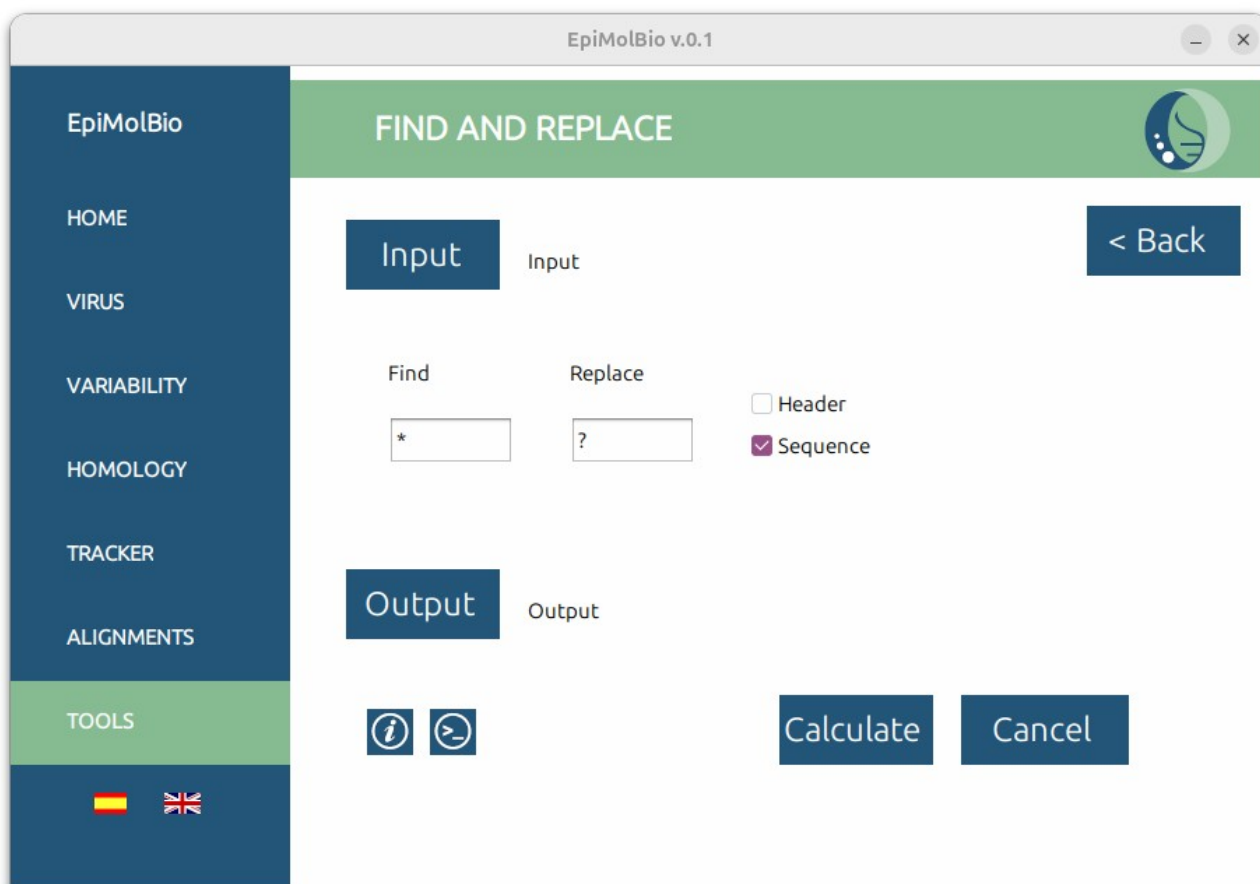
See the full manual [here](#).

### Example File

**File Name:** REPLACE RT HIV-1 AA\_A.fasta, REPLACE RT HIV-1 AA\_B.fasta, REPLACE RT HIV-1 AA\_C.fasta, REPLACE RT HIV-1 AA\_D.fasta

### Parameter Selection

- Find:
  - \*
- Replace:
  - ?
- Header/Sequence:
  - Sequence



The screenshot shows the EpiMolBio v.0.1 application window. The title bar reads "EpiMolBio v.0.1". The interface has a dark blue sidebar on the left with the following menu items: EpiMolBio, HOME, VIRUS, VARIABILITY, HOMOLOGUE, TRACKER, ALIGNMENTS, TOOLS (highlighted in green), and flags for Spain and the UK. The main content area has a green header bar that says "FIND AND REPLACE" with a circular logo on the right. Below the header, there is an "Input" section with a text box containing "\*" and a "Replace" section with a text box containing "?". To the right of these text boxes are two checkboxes: "Header" (unchecked) and "Sequence" (checked). There is a "< Back" button in the top right corner. Below the input fields is an "Output" section with a text box. At the bottom of the main area are two buttons: "Calculate" and "Cancel". There are also two small circular icons (one with an 'i' and one with a minus sign) to the left of the "Calculate" button.