

## Section VI.2.C: Partial Sequence Filtering (p181 - p186)

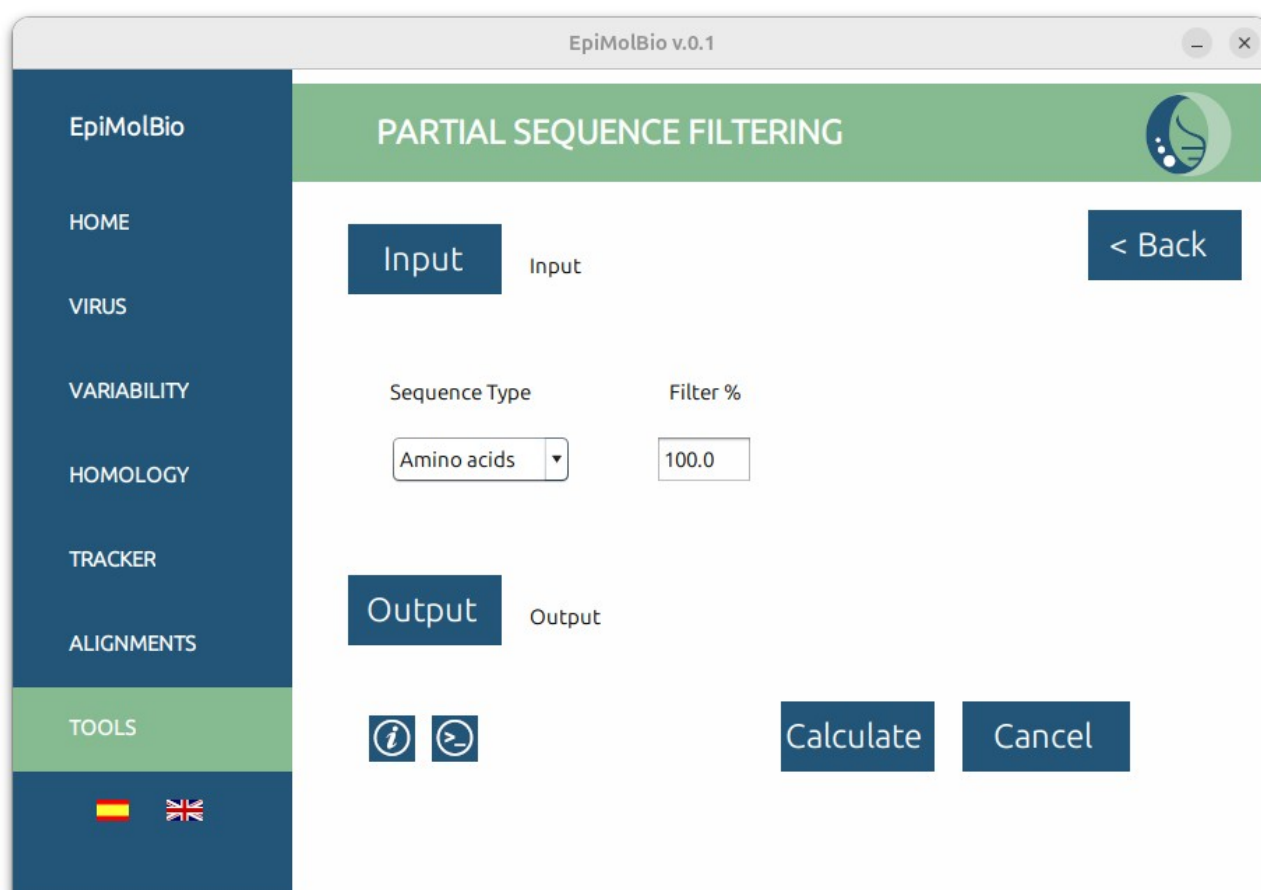
See the full manual [here](#).

### Example File

**File Name:** PARTIAL FILTER RT HIV-1 AA\_A.fasta, PARTIAL FILTER RT HIV-1 AA\_B.fasta, PARTIAL FILTER RT HIV-1 AA\_C.fasta, PARTIAL FILTER RT HIV-1 AA\_D.fasta, LOST SEQUENCES.html

### Parameter Selection

- **Sequence Type:**
  - *Amino acids*
- **Filter %:**
  - 100.0



The screenshot shows the EpiMolBio v.0.1 web application interface. The main title is "PARTIAL SEQUENCE FILTERING". The interface is divided into a left sidebar and a main content area. The sidebar contains navigation links: EpiMolBio, HOME, VIRUS, VARIABILITY, HOMOLOGUE, TRACKER, ALIGNMENTS, TOOLS, and language flags (Spanish and English). The main content area has a green header bar with the title and a logo. Below the header, there are two tabs: "Input" and "Output". The "Input" tab is active, showing a "Sequence Type" dropdown menu set to "Amino acids" and a "Filter %" input field set to "100.0". There is a "< Back" button in the top right corner. Below the input fields, there are two buttons: "Calculate" and "Cancel". At the bottom left of the main content area, there are two icons: an information icon (i) and a help icon (?).