

### Section III.1: Similarity (p110 - p114)

See the full manual [here](#).

---

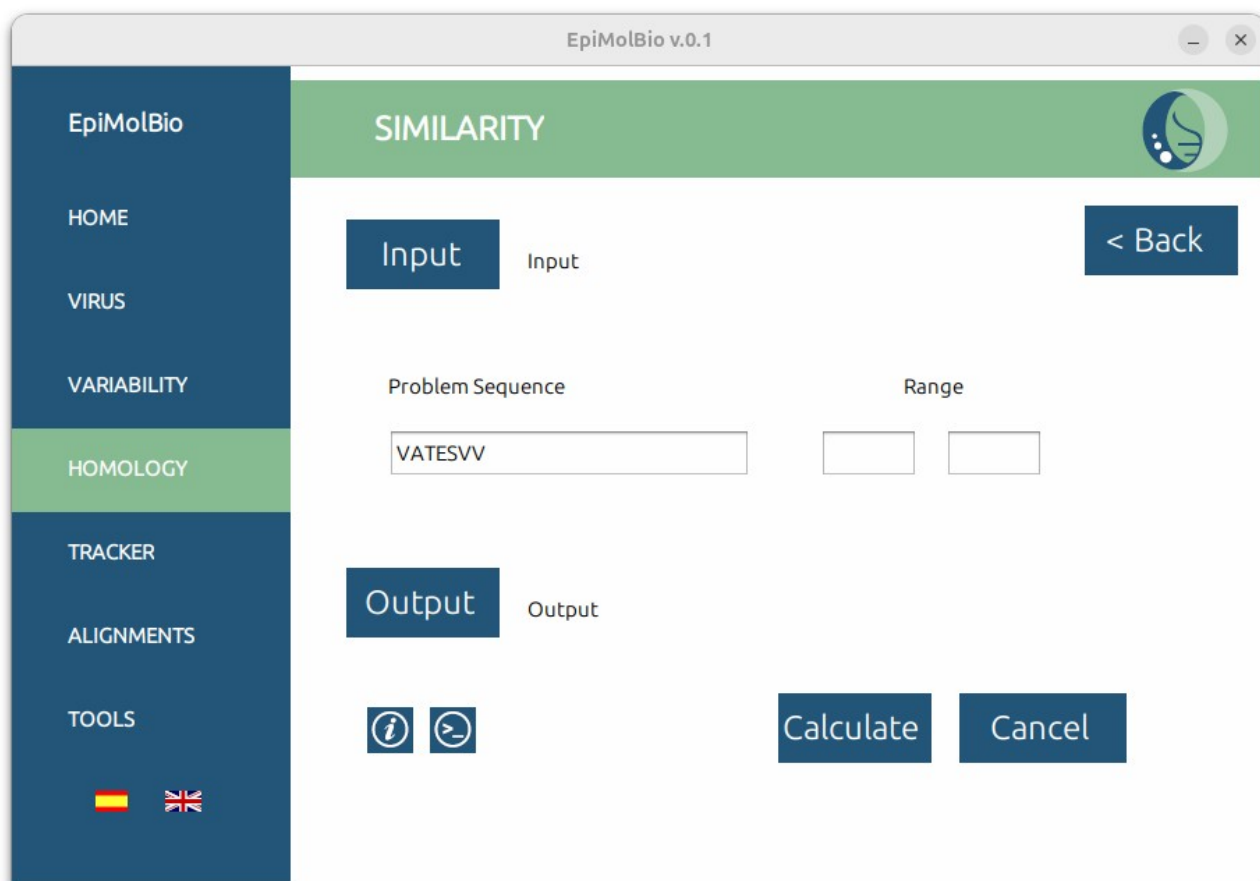
### Example File

File Name: SIMILARITY RT HIV-1 PEPTIDE VATESVV.html

---

### Parameter Selection

- **Problem Sequence:**
  - VATESVV
- **Range:**
  - Empty



The screenshot shows the EpiMolBio v.0.1 web application interface. The title bar indicates the version. The left sidebar contains navigation links: EpiMolBio, HOME, VIRUS, VARIABILITY, HOMOLOGY (highlighted in green), TRACKER, ALIGNMENTS, and TOOLS. At the bottom of the sidebar are flags for Spanish and English. The main content area is titled 'SIMILARITY' and features a green header bar with a DNA helix icon. Below the header, there are two main sections: 'Input' and 'Output'. The 'Input' section includes a 'Problem Sequence' field containing 'VATESVV' and a 'Range' section with two empty input boxes. A '< Back' button is located in the top right of the input section. The 'Output' section includes an 'Output' label, an information icon (i), a warning icon (⚠), and two buttons: 'Calculate' and 'Cancel'.