

## Section IV.2: Tracker Flanking (p135 - p140)

See the full manual [here](#).

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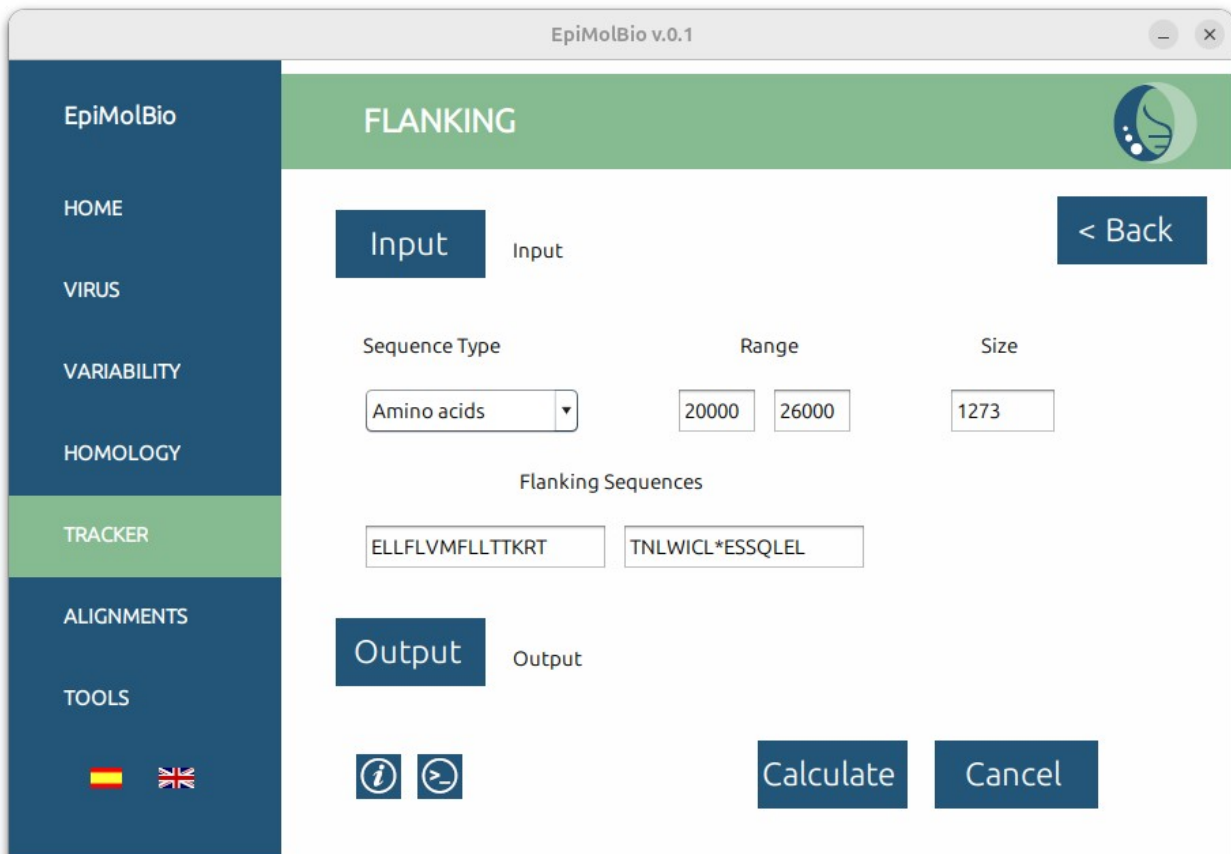
### Example File

File Name: TRACKED FLANKING SARS-COV-2 (SPIKE).fasta

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### Parameter Selection

- **Sequence Type:**
  - *Amino acids*
- **Range:**
  - 20000 - 26000
- **Size:**
  - 1273
- **Flanking Sequences:**
  - *ELLFLVMFLLTTKRT - TNLWICL\*ESSQLEL*



The screenshot shows the EpiMolBio v.0.1 FLANKING interface. The window has a title bar with the text "EpiMolBio v.0.1" and standard window controls. On the left is a dark blue sidebar with navigation links: EpiMolBio, HOME, VIRUS, VARIABILITY, HOMOLOGUE, TRACKER (highlighted in green), ALIGNMENTS, and TOOLS. At the bottom of the sidebar are flags for Spain and the United Kingdom. The main content area has a green header with the word "FLANKING" and a circular logo. Below the header, there are input fields for "Sequence Type" (set to "Amino acids"), "Range" (20000 to 26000), and "Size" (1273). There are also input fields for "Flanking Sequences" containing "ELLFLVMFLLTTKRT" and "TNLWICL\*ESSQLEL". At the bottom, there are buttons for "Calculate" and "Cancel", along with information and help icons. A "< Back" button is located in the top right corner of the main area.

EpiMolBio v.0.1

FLANKING

Input Input < Back

Sequence Type Range Size

Amino acids 20000 26000 1273

Flanking Sequences

ELLFLVMFLLTTKRT TNLWICL\*ESSQLEL

Output Output

Calculate Cancel