

Individual Polymorphisms Markers 90%

User Manual

Section II.1.A: Polymorphisms - Individual (p44, p60 - p65)

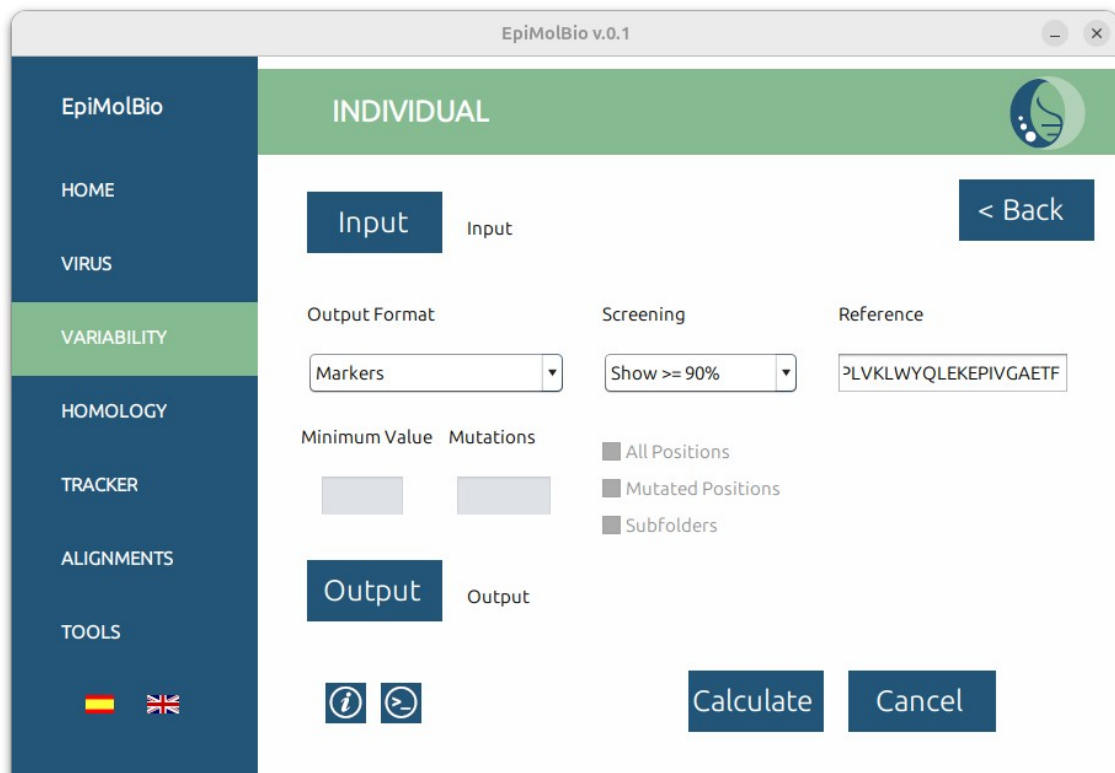
See the full manual [here](#).

Example File

File Name: MARKERS RT HIV-1 90%.html

Parameter Selection

- **Output Format:**
 - *Markers*
- **Screening:**
 - *Show >= 90%*
- **Reference:**
 - *PISPIETVPVKLPGMDGPKVKQWPLTEEKIKALVEICTEMEKEGKISKIGPENPYNTPVFAIKKKD
STKWRKLVDFRELNKRTQDFWEVQLGIPHPAGLKKKKSVTVLDVGDAYFSVPLDEDFRKYTAFTI
PSINNETPGIRYQYNVLPQGWWKGSPAIFQSSMTKILEPFRKQNPDIVIYQYMDDLTVGSDLEIGQ
HRTKIEELRQHLLRWGLTTPDKKHQKEPPFLWMGYELHPDKWTVQPIVLPEKDSWTVNDIQKL
VGKLNWASQIYPGIKVRQLCKLLRGTKALTEVIPLTEEALELAENREILKEPVHGVYDPSKDLIA
EIQKQGQGWQTYQIYQEPFKNLKTGKYARMRGAHTNDVKQLTEAVQKITTESIVIWGKTPKFKL
PIQKETWETWWTEYWQATWIPEWEFVNTPLVVKLWYQLEKEPIVGAETF*



The screenshot shows the EpiMolBio v.0.1 web application interface. The left sidebar contains navigation links: HOME, VIRUS, VARIABILITY (highlighted), HOMOLOGU, TRACKER, ALIGNMENTS, and TOOLS. At the bottom of the sidebar are flags for Spanish and English. The main content area is titled 'INDIVIDUAL' and features a 'Back' button. Below this, there are three sections: 'Input', 'Output Format', and 'Screening'. The 'Input' section has a text box for the input sequence. The 'Output Format' section has a dropdown menu set to 'Markers'. The 'Screening' section has a dropdown menu set to 'Show >= 90%'. The 'Reference' section has a text box containing the reference sequence. Below these sections are checkboxes for 'All Positions', 'Mutated Positions', and 'Subfolders'. At the bottom, there are 'Calculate' and 'Cancel' buttons, along with information and help icons.