



## Individual Polymorphisms Multiple Mutations Subfolders



### User Manual

#### Section II.1.A: Polymorphisms - Individual (p44, p66 - p70)

See the full manual [here](#).



### Example File

File Name: SUBFOLDERS OUTPUT MULTIPLE MUTATIONS RT HIV-1.csv



### Parameter Selection

- **Output Format:**
  - *Multiple Mutations*
- **Mutations:**
  - *K43E,P272A*
- **All/Mutated/Subfolders:**
  - *Subfolders*

The screenshot shows the EpiMolBio v.0.1 web application interface. The left sidebar contains navigation links: EpiMolBio, HOME, VIRUS, VARIABILITY (highlighted), HOMOLOGUE, 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 the title, there are three input fields: 'Input', 'Output Format' (set to 'Multiple Mutations'), and 'Screening' (set to 'Select ...'). A 'Reference' field is also present. Below these, there are checkboxes for 'All Positions', 'Mutated Positions', and 'Subfolders' (checked). A 'Minimum Value' field is set to 'Mutations' with a value of 'K43E,P272A'. At the bottom, there are 'Output' and 'Input' buttons, an 'i' icon, a '>-' icon, and 'Calculate' and 'Cancel' buttons.