



Individual Polymorphisms Mutations by Position Subfolders



User Manual

Section II.1.A: Polymorphisms - Individual (p44, p71 - p75)

See the full manual [here](#).



Example File

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



Parameter Selection

- **Output Format:**
 - *Mutations by Position*
- **Mutations:**
 - 43,272
- **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 this, there are three sections: 'Input', 'Screening', and 'Reference'. The 'Input' section has a dropdown menu for 'Output Format' set to 'Mutations by Position'. The 'Screening' section has a 'Select ...' dropdown and three checkboxes: 'All Positions' (unchecked), 'Mutated Positions' (unchecked), and 'Subfolders' (checked). The 'Reference' section is empty. Below these sections, there is an 'Output' section with a 'Minimum Value' input field set to '43,272' and a 'Mutations' input field. At the bottom right, there are 'Calculate' and 'Cancel' buttons. Information and help icons are located at the bottom left of the main content area.