

Individual Polymorphisms Mutations Table Subfolders

User Manual

Section II.1.A: Polymorphisms - Individual (p44, p52 - p59)

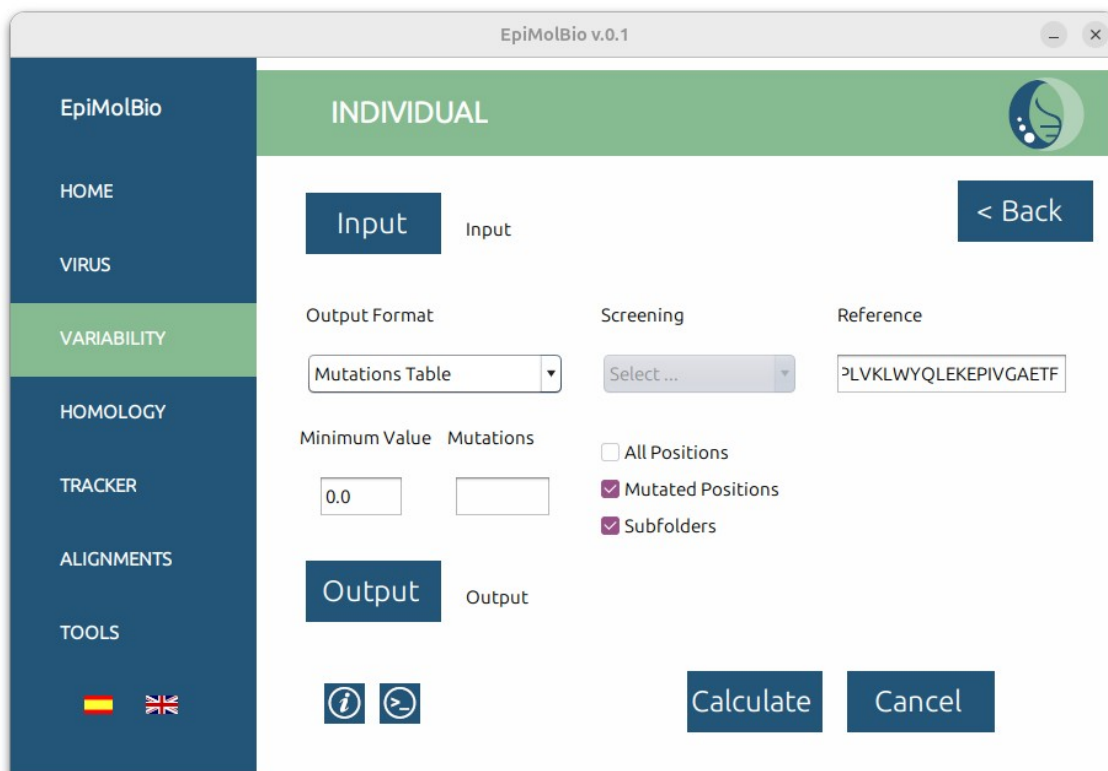
See the full manual [here](#).

Example File

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

Parameter Selection

- **Output Format:**
 - *Mutations Table*
- **Reference:**
 - *PISPIETVPVKLPGMDGPKVKQWPLTEEKIKALVEICTEMEKEGKISKIGPENPYNTPVFAIKKK
DSTKWRKLVDFRELNKRTQDFWEVQLGIPHPAGLKKKSVTVLDVGDAYFSVPLDEDFRKYTAF
TIPSINNETPGIRYQYNVLPQGWKGSPAIFQSSMTKILEPFRKQNPDIVIYQYMDDLYVGSdleIG
QHRTKIEELRQHLLRWGLTTPDKKHQKEPPFLWMGYELHPDKWTVQPIVLPEKDSWTVNDIQ
KLVGKLNWASQIYPGIKVRQLCKLLRGTKALTEVIPLTEEALELAENREILKEPVHGVYYDPSKDLI
AEIQKQGQGQWTYQIYQEPFKNLKTGKYARMGAHTNDVKQLTEAVQKITTESIVIWGKTPKF
KLPIQKETWETWWTEYWQATWIPEWEFVNTPLVKLWYQLEKEPIVGAETF*
- **Minimun Value:**
 - 0.0
- **All/Mutated/Subfolders:**
 - *Mutated Positions*
 - *Subfolders*



The screenshot shows the EpiMolBio v.0.1 web application interface. The left sidebar contains navigation links: HOME, VIRUS, VARIABILITY (highlighted), HOMOLOG, TRACKER, ALIGNMENTS, and TOOLS. At the bottom of the sidebar are flags for Spain and the UK. 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 Table'. The 'Screening' section has a dropdown menu for 'Select ...' and checkboxes for 'All Positions' (unchecked), 'Mutated Positions' (checked), and 'Subfolders' (checked). The 'Reference' section has a text input field containing 'PLVKLWYQLEKEPIVGAETF'. At the bottom, there are 'Calculate' and 'Cancel' buttons, along with information and help icons.