

Archivo de Ejemplo

Nombre del archivo: CONTAR SECUENCIAS MUTADAS RT VIH-1 AA.csv

Selección de Parámetros

- **Formato:**
 - *Secuencias Mutadas*
- **Referencia:**
 - *PISPIETVPVKLPGMDGPKVKQWPLTEEKIKALVEICTEMEKEGKISKIGPENPYNTPVFAIKKKD
STKWRKLVDFRELNKRTQDFWEVQLGIPHPAGLKKKKSXTVLDVGDAYFSVPLDEDFRKYTAFTI
PSINNETPGIRYQYNVLPQGWWKGSPAIFQSSMTKILEPFRKQNPDIVIYQYMDDLTVGSDLEIGQ
HRTKIEELRQHLLRWGLTTPDKKHQKEPPFLWMGYELHPDKWTVQPIVLPEKDSWTVNDIQKL
VGKLNWASQIYPGIKVRQLCKLLRGTKALTEVIPLTEEALELAENREILKEPVHGVYDPSKDLIA
EIQKQGQGWQTYQIYQEPFKNLKTGKYARMRGAHTNDVKQLTEAVQKITTESIVIWGKTPKFKL
PIQKETWETWWTEYWQATWIPEWEFVNTPLVKLWYQLEKEPIVGAETF*
- **AA/NT:**
 - AA



The screenshot shows the EpiMolBio v.0.1 web application interface. The main header is green with the title 'CONTAR SECUENCIAS' and a logo. A dark blue sidebar on the left contains navigation links: EpiMolBio, INICIO, VIRUS, VARIABILIDAD, HOMOLOGÍA, RASTREADOR, ALINEAMIENTOS, HERRAMIENTAS (highlighted), and flags for Spanish and English. The main content area has a white background. At the top, there are buttons for 'Entrada' and '< Atrás'. Below, there are two sections: 'Formato' with a dropdown menu set to 'Secuencias Mutadas', and 'Referencia' with a text input field containing a long protein sequence. To the right of the 'Referencia' field are two checkboxes: 'AA' (checked) and 'NT' (unchecked). Below these, there are buttons for 'Salida' and 'Salida'. At the bottom, there are two buttons: 'Calcular' and 'Cancelar'. There are also two small circular icons (one with an 'i' and one with a magnifying glass) and a 'Calcular' button.