
APPLICATIONS

With this research and its conclusions, scientists can:

- Better understand the origin of HIV-1 and HIV-2
- Better understand spread of the virus
- Create medicine and treatments aimed at certain unchanging or weak areas of HIV

On a pure side, this research has helped in:

- Creating a strong hypothesis (the tree) for the evolutionary history of the taxa
 - How they are related
 - How they evolved
- Potentially guiding our search for new subtypes of HIV