



epiPATH

Molecular Epidemiology Database for Infectious Pathogens

Please, login to access this application

Username

Password

Log In

Note: This server uses Session Cookies. Please, turn on "Accept Cookies" at the preferences menu of your browser. (example @ mozilla: tools -> cookie manager -> allow session cookies from this site)

About epiPATH

epiPATH is a bioinformatics platform to store and analyze evolutionary, population and epidemiological data from infectious pathogens and the sequences obtained in our laboratory. Now, we have stored about 11950 sequences of HCV but in a nearly future we will introduce information about other pathogens.

The web interface gives access to data and its management. Actually, the tools implemented are those to insert, update, search and delete information in the database server. Moreover, we are going to add analysis tools to study our data stored. To access all this tools and data you must have authorization with an username and password. If you want to have access, please contact with the system administrator.

epiPATH distribution

epiPATH database schema, toolscripts and web interface are free software. [Learn more]

Institut Cochin de Biologie Evolutive

User identification



epiPATH

Molecular Epidemiology Database for Infectious Pathogens

Welcome, alicia

Logout

This bioinformatics platform provides a web interface to insert, update, delete and search data in the epiPATH database. Download the epiPATH Manual.

Data

[Insert Data](#)

[Update Data](#)

[Search Data](#)

[Delete Data](#)