

# Outils et bases de données

Affichage des résultats 1 à 5 sur 106 au total

## STATISTICS

### **PELICAN** (<https://gitlab.in2p3.fr/phoogole/pelican>)

Type : package

Statut : supported

Contact : Veber P. , Boussau B.

Search for genotype-phenotype associations at the genome scale, across species.

## PHYLOGENY

### **Phylter** (<https://damiendevienne.github.io/phylter/>)

Type : package

Statut : supported

Contact : de Vienne D.

An R package for analyzing, visualizing and filtering phylogenomic datasets. Reads a collection of...

## PHYLOGENY

### **Thirdkind** (<https://github.com/simonpenel/thirdkind/wiki>)

Type : package

Statut : supported

Contact : Penel S.

Thirdkind is a software producing svg files displaying phylogenetic reconciliations, including 3...

## OTHER

### **DDI-predictor** (<https://www.ddi-predictor.org/>)

Type : database

Statut : supported

Contact : Tod M.

Quantitative prediction of the impact on drug exposure of drug-drug interactions mediated by...

## COMPARATIVE GENOMICS

### **HTdetect** (<https://github.com/l-modolo/htdetect>)

Type : package

Statut : supported

Contact : Lerat E., Modolo L.

Horizontal Transfert detection from pairwise genome alignment