

Outils et bases de données

Affichage des résultats 1 à 5 sur 106 au total

STATISTICS

PELICAN (<https://gitlab.in2p3.fr/phoogel/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