

Recherche avancée

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CDYL2 Epigenetically Regulates MIR124 to Control NF-κB/STAT3-Dependent Breast Cancer Cell Plasticity (<https://hal.archives-ouvertes.fr/hal-02991271>)

Maha Siouda, Audrey Dujardin, Laetitia Barbolat-Boutrand, Marco Mendoza-Parra, Benjamin Gibert, Maria Ouzounova, Jebrane Bouaoud, Laurie Tonon, Marie Robert, Jean-Philippe Foy, Vincent Lavergne, Serge Manie, Alain Viari, Alain Puisieux, Gabriel Ichim, Hinrich Gronemeyer, Pierre Saintigny, Peter Mulligan

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MOSAIC_SSD: A new web tool for species sensitivity distribution to include censored data by maximum likelihood (<https://hal.inrae.fr/hal-02967115>)

Guillaume Kon Kam King, Philippe Veber, Sandrine Charles, Marie Laure Delignette-Muller

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Hierarchical modelling of species sensitivity distribution: development and application to the case of diatoms exposed to several herbicides (<https://hal.inrae.fr/hal-02967089>)

Guillaume Kon Kam King, Floriane Larras, Sandrine Charles, Marie Laure Delignette-Muller

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Evolution of the primate β -globin gene region (<https://hal.inrae.fr/hal-02965509>)

P. Savatier, G. Trabuchet, C. Faure, Y. Chebloune, M. Gouy, G. Verdier, V.M. Nigon

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Genome-wide studies highlight indirect links between human replication origins and gene regulation. (<https://hal.science/hal-00332341>)

Jean-Charles Cadoret, Françoise Meisch, Vahideh Hassan-Zadeh, Isabelle Luyten, Claire Guillet, L. Duret, Hadi Quesneville, Marie-Noëlle Prioleau

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Differential retention of metabolic genes following whole-genome duplication (<https://hal.science/hal-00428358v2>)

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The relationship among gene expression, the evolution of gene dosage, and the rate of protein evolution (<https://hal.science/hal-00539474v2>)

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Occurrence of a non deleterious gene conversion event in the BRCA1 gene (<https://univ-lyon1.hal.science/hal-02018994>)

C. Tessereau, M. Leone, M. Buisson, L. Duret, O. M. Sinilnikova, S. Mazoyer

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Evolution of gene neighborhoods within reconciled phylogenies (<https://hal.science/hal-00733178>)

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TPMS: a set of utilities for querying collections of gene trees. (<https://hal.science/hal-00972257>)

Thomas Bigot, Vincent Daubin, Florent Lassalle, Guy Perri  re

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Genome-scale phylogenetic analysis finds extensive gene transfer among fungi (<https://hal.science/hal-01218565>)

G.J. Szollosi, Adrián Arellano Davin, Eric Tannier, Vincent Daubin, Bastien Boussau

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- > Actualités
(357)
- > Agenda
(1228)
- > Annuaire des membres
(247)
- > Département
(5)
- > Equipe
(17)
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(2)
- > Page de base
(94)
- > (-)
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(5278)

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- > Pertinence
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