Clémence Réda | Marie Skłodowska-Curie Postdoctoral Fellow @ Universität Rostock

(updated on 22/01/2025)

Systembiologie und Bioinformatik Rostock, Ulmenstraße 69, 18059 Rostock, Germany

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Professional Experience

Research Positions

Tenured research fellow | Chargé de recherche de classe normale — **France**
Centre national de la recherche scientifique (CNRS), *UMR 8197* — *From 02/2025*

Marie Skłodowska-Curie Postdoctoral Fellow | **Pr. Olaf Wolkenhauer** — **Rostock, Germany**
Universität Rostock *(SBI Rostock)* — *05/2023–02/2025*
Development of the RECeSS project, focusing on the development of new, improved techniques for drug development based on collaborative filtering approaches.
Skills Collaborative Filtering · Python (Programming Language) · Applied Machine Learning

MSC PF (research visit) | **Pr. Hisashi Kashima & Dr. Koh Takeuchi** — **Kyoto, Japan**
Kyoto University *(Machine Learning and Data Mining Research Laboratory)* — *12/2024–01/2025*
Joint project on diversity for recommendations in collaborative filtering (part of the RECeSS project).
Skills Collaborative Filtering · Python (Programming Language) · Applied Machine Learning

MSC PF (secondment) | **Dr. Jill-Jênn Vie** — **Saclay, France**
Inria Saclay *(SODA team)* — *07/2023–10/2023*
Design of the JELI algorithm, a collaborative filtering approach integrating graph priors to enable explicit interpretability, and application to drug repurposing (part of the RECeSS project).
Skills Factorization Machines · Knowledge Graph · Interpretability

Postdoctoral position | **Pr. Andrée Delahaye-Duriez** — **Paris, France**
Neurodiderot *(UMR 1141)* — *09/2022–03/2023*
Development and implementation of the NORDic pipeline for Boolean networks, Prefiguration of the multiomics workflow for the RHU FAME project led by Pr. Élie Azoulay.
Skills Systems Biology · Programming · Interdisciplinary Research · Bioinformatics

PhD position | **Pr. Andrée Delahaye-Duriez & Dr. Émilie Kaufmann** — **Paris, France**
Neurodiderot *(UMR 1141)* & ***SCOOL*** *(UMR 9189)* — *09/2019–09/2022 (36 months)*
Combination of gene regulatory networks and sequential machine learning for drug repurposing.
Skills Systems Biology · Multi-Armed Bandits · Interdisciplinary Research · Bioinformatics

Master internship | **Pr. Andrée Delahaye-Duriez & Dr. Émilie Kaufmann** — **Paris, France**
Neurodiderot *(UMR 1141)* & ***SCOOL*** *(UMR 9189)* — *03/2019–08/2019 (4 months)*
Design of a drug repurposing method through a bandit algorithm combined with the prediction of transcriptomic states by a gene regulatory network. Application to the prediction of new anti-epileptics.
Skills Interdisciplinary Collaboration · Interdisciplinary Research · Statistical Learning · Project Design · Bioinformatics

Predoctoral internship | **Dr. Bartek Wilczyński** — **Warsaw, Poland**
Regulomics team *(MIM UW)* — *10/2017–07/2018 (10 months)*
Proof-of-concept on the explicit inclusion of biological interactions in gene regulatory networks and its impact on inference and simulation of transcriptomic regulation. Led to a publication in Journal of Theoretical Biology (DOI : 10.1016/j.jtbi.2019.110091).
Skills Network Analysis · Epigenetics · Python (Programming Language) · Systems Biology · Scientific Presentation

Master internship | **Dr. Nicholas Luscombe & Dr. Garth Ilsley** — **Onna-son, Japan**
Genomics and Regulatory Systems Unit *(OIST)* — *02/2017–07/2017 (5 months)*
Design and implementation of a single-cell RNA sequencing clustering method taking into account intergene expression dependencies using a probabilistic model ; implementation in R Shiny of a web application for the visualisation and preliminary analysis of single-cell RNA sequencing data. Application to transcriptomic data analysis in ciona (*Ciona intestinalis*).
Skills Benchmarking · R Shiny · Unsupervised Learning · Data Visualization · Python (Programming Language)

Bachelor internship | **Dr. Macha Nikolski & Dr. Mathieu Raffinot** **Bordeaux, France**
Centre de Bioinformatique de Bordeaux *(Université de Bordeaux)* *05/2016–07/2016 (2 months)*
Design and implementation of compared analyses of taxonomic trees built from metagenomic data. Application to the analysis of data from intestinal guts of children afflicted with cystic fibrosis at Hôpital Pellegrin in Bordeaux.
Skills Metagenomics · Phylogenetics · Supervised Learning · Unsupervised Learning · Python (Programming Language)

Teaching & Mentoring Experiences

Biostatistics, programming and bioinformatics **Université Paris Cité**
Doctorant Contractuel avec Mission d'Enseignement (DCME) (Teaching Assistant) *09/2020–09/2021 (64 hours)*
References: Dr. Anne Badel & Pr. Olivier Taboureau

Co-supervision of a PhD **SBI Rostock**
Joint supervision of Orell Trautmann with Pr. Olaf Wolkenhauer (50%) *08/2024–present*
Design of a knowledge graph-based algorithm predicting drug combinations.

Co-supervision of a Masters-equivalent internship and a PhD **SBI Rostock**
Joint supervision of Rahul Bordoloi with Pr. Olaf Wolkenhauer (33%) *09/2023–present*
Development of a linear discriminant algorithm on multivariate temporal data: paper.

Co-supervision of a Master's degree internship **Inserm Neurodiderot**
Joint supervision of Adrien Dufour with Pr. Andrée Delahaye-Duriez (25%) *02/2020–07/2020 (6 months)*
Identification of functional families of migroglia cells from targeted single-cell RNA sequencing data of inflammatory microglia at a developmental stage: paper.

Co-supervision of a Masters's degree project **ENS Paris-Saclay**
Joint supervision of Ariane Alix with Dr. Émilie Kaufmann (50%) *11/2019–01/2020 (2 months)*
Proposal of a project on the adaptation of a published drug-target prediction method to drug repurposing using collaborative filtering in the course *Graphs in Machine Learning* taught by Dr. Michał Valko in Master Vision Apprentissage (**MVA** 2020).

Education

Université Paris Cité, Inserm UMR 1141 & CNRS UMR 9189
PhD in Genetics *09/2019 – 09/2022*
Doctorate Degree in Science. Title: **Combination of gene regulatory networks and sequential machine learning for drug repurposing**, supervised by Pr. Andrée Delahaye-Duriez (Inserm UMR 1141) & Dr. Émilie Kaufmann (CNRS UMR 9189).
***Viva*: 09/09/2022**.

École Normale Supérieure[†] **(ENS) Paris-Saclay** **(ex-École Normale Supérieure de Cachan)**
M2 Master Vision, Apprentissage (MVA) *09/2018 – 09/2019*
Master's degree in Machine Learning. (*summa cum laude*, Grade: 16.17/20, no ranking)

ENS Paris-Saclay
M1 Master Parisien en Recherche en Informatique (MPRI) *09/2016 – 09/2017*
Master's degree in Computer Sciences. (*summa cum laude*, Grade: 16.72/20, rank: 3/25)

École Normale Supérieure de Cachan
L3 Licence informatique fondamentale ENS Cachan *09/2015 – 09/2016*
Bachelor's degree in Computer Sciences. (*cum laude*, Grade: 14.64/20, rank: 10/26)

[†] *École Normale Supérieures are selective French schools for research and teaching.*

Funding and Awards as Principal Recipient

Accessit from the Societe Savante Francophone d'Apprentissage Machine **SSFAM**
PhD award (award list) *2024*

Marie Skłodowska-Curie Postdoctoral Fellowship 2022 **Horizon 2020**
Postdoctoral grant *2023–2025 (2 years)*
RECeSS project, Project ID: 101102016.

Research

Preprints

An Anytime Algorithm for Good Arm Identification
M. Jourdan & C. Réda
, *Under review, DOI: 10.48550/arXiv.2310.10359*

Peer-Reviewed Scientific Journals

2025

Multivariate Functional Linear Discriminant Analysis for the Classification of Short Time Series with Missing Data
R. Bordoloi, C. Réda, O. Trautmann, S. Bej, O. Wolkenhauer
, *In press, Machine Learning, DOI: 10.48550/arXiv.2402.13103*

Comprehensive evaluation of pure and hybrid collaborative filtering in drug repurposing
C. Réda, J.-J. Vie, O. Wolkenhauer
, *Scientific Reports, 15, 2711, DOI: 10.1038/s41598-025-85927-x*

Joint Embedding-Classifier Learning for Interpretable Collaborative Filtering
C. Réda, J.-J. Vie, O. Wolkenhauer
, *BMC Bioinformatics, 26, 26, DOI: 10.1186/s12859-024-06026-8*

Neonatal inflammation impairs developmentally-associated microglia and promotes a highly reactive microglial subset
A. Dufour*, A. Heydari-Olya*, S. Foulon*, C. Réda, et al.
, *Brain, Behavior, and Immunity, DOI: 10.1016/j.bbi.2024.09.019*

2024

stanscofi and benchscofi: a new standard for drug repurposing by collaborative filtering
C. Réda, J.-J. Vie, O. Wolkenhauer
, *Journal of Open Source Software, 9(93):5973, DOI: 10.21105/joss.05973*

2023

NORDic: a Network-Oriented package for the Repurposing of Drugs
C. Réda & A. Delahaye-Duriez
, *Journal of Open Source Software, 8(90):5532, DOI: 10.21105/joss.05532*

2021

Machine learning applications in drug development
C. Réda, É. Kaufmann & A. Delahaye-Duriez
, *Computational and Structural Biotechnology Journal, 18:241-252, DOI: 10.1016/j.csbj.2019.12.006*

2020

Automated inference of gene regulatory networks using explicit regulatory modules
C. Réda & B. Wilczyński
, *Journal of Theoretical Biology, 486:110091, DOI: 10.1016/j.jtbi.2019.110091*

2019

Identification de cibles thérapeutiques et repositionnement de médicaments par analyses de réseaux géniques
A. Delahaye-Duriez, C. Réda & P. Gressens
, *Médecine/Sciences, 35:515-518, DOI: 10.1051/medsci/2019108*

Peer-Reviewed Conference Proceedings

2022

Near-optimal Collaborative Learning in Bandits
C. Réda, S. Vakili, É. Kaufmann
, *Proceedings of the 36^{th} Conference on Advances in Neural Information Processing Systems (**NeurIPS** 2022)*
HAL: 03825099 [*Selected as Oral*]

Prioritization of Candidate Genes Through Boolean Networks
C. Réda, A. Delahaye-Duriez
, *Proceedings of the 20^{th} International Conference on Computational Methods in Systems Biology (**CMSB** 2022)*
Springer:89-121 [*Best Student Paper Award*]

2021

Dealing With Misspecification In Fixed-Confidence Linear Top-m Identification
C. Réda, A. Tirinzoni & R. Degenne
, *Proceedings of the 35^{th} Conference on Neural Information Processing Systems (**NeurIPS** 2021), 34, HAL: 03409205*

Top-m identification for linear bandits
C. Réda, É. Kaufmann & A. Delahaye-Duriez
*, Proceedings of the 24^{th} International Conference on Artificial Intelligence and Statistics (**AISTATS** 2021), 130*
HAL: 03172145

Oral Communications at International Conferences

C. Réda. Benchmarking collaborative filtering approaches to drug repurposing
e:Med Meeting 2023 on Systems Medicine (Berlin, Germany) *10/10/2023*

C. Réda. Near-optimal Collaborative Learning in Bandits
35^{th} *International Conference on Advances in Neural Information Processing Systems (New Orleans, USA)* *07/12/2022*

C. Réda. Prioritization of Candidate Genes Through Boolean Networks
20^{th} *International Conference on Computational Methods in Systems Biology (Bucharest, Romania)* *14/09/2022*

C. Réda. Gene network oriented drug discovery: automated inference of Boolean networks (...)
13^{th} *Conference on Dynamical Systems Applied to Biology and Natural Sciences (held virtually)* *10/02/2022*

C. Réda. Dealing With Misspecification In Fixed-Confidence Linear Top-m Identification
NeurIPS@Paris 2021 *(Paris, France)* *08/12/2021*

C. Réda. Automated inference of gene regulatory networks using explicit regulatory modules
Journées Ouvertes de Biologie, Informatique et Mathématique (JOBIM) 2020 (held virtually) *02/07/2020*

Poster Presentations at International Conferences

C. Réda. JELI: an interpretable embedding-learning recommender system for drug repurposing
ECCB 2024 (Turku, Finland) *09/2024*

C. Réda. JELI: an interpretable embedding-learning recommender system for drug repurposing
JOBIM 2024 (Toulouse, France) *06/2024*

C. Réda. Towards a large-scale benchmark of collaborative filtering in drug repurposing
SMPGD 2024 (Paris, France) *02/2024*

C. Réda. Drug repurposing in breast cancer by combining bandit algorithms and Boolean networks (...)
ISMB/ECCB 2023 *(Lyon France)* *07/2023*

C. Réda. Prioritization of Candidate Genes Through Influence Maximization
Journées Ouvertes de Biologie, Informatique et Mathématique (JOBIM 2022, Rennes, France) *07/2022*

C. Réda. Dealing With Misspecification In Fixed-Confidence Linear Top-m Identification
35^{th} *International Conference on Advances in Neural Information Processing Systems (NeurIPS 2022, held virtually)* *12/2021*

C. Réda. Top-m identification for linear bandits
24^{th} *International Conference on Artificial Intelligence and Statistics (AISTATS 2021, held virtually)* *04/2021*

Open-Source Softwares & Datasets

Softwares

2024

Joint Embedding-classifier Learning for improved Interpretability (JELI)
C. Réda
, Zenodo, DOI: 10.5281/zenodo.12193722, GitHub: `recess-eu-project/JELI`
Python package implementing an explicitly interpretable collaborative filtering

2023

BENCHmark for drug Screening with COllaborative FIltering (benchscofi)
C. Réda
, Zenodo, DOI: 10.5281/zenodo.8241505, GitHub: `recess-eu-project/benchscofi`
Python package implementing algorithms and methods from the state-of-the-art in drug repurposing with collaborative filtering

STANdard for drug Screening by COllaborative FIltering (stanscofi)
C. Réda
, Zenodo, DOI: 10.5281/zenodo.8038847, GitHub: `recess-eu-project/stanscofi`
Python package for the automation of the training and validation of drug repurposing with machine learning

Network Oriented Repurposing of Drugs (NORDic)
C. Réda
, Zenodo, DOI: 10.5281/zenodo.7239047, GitHub: `clreda/NORDic`
Python package for the inference, analysis of Boolean networks & application to drug repurposing

Datasets

2023

PREDICT
C. Réda
, *Zenodo, DOI: 10.5281/zenodo.7982964*
Large drug repurposing dataset with open-source generation

TRANSCRIPT
C. Réda
, *Zenodo, DOI: 10.5281/zenodo.7982969*
Drug repurposing dataset on transcriptomic data with open-source generation

Commitment to Popularization of Sciences and Law Making

Popularization of Sciences

11/21/2024: Popularization paper (in French) on drug repurposing aimed at medical practitioners: C. Réda, B. Villoutreix and A. Delahaye-Duriez. **Repositionnement de médicaments** *In* La Revue du Praticien, 21 novembre 2024, 74(9);942-6 (link)

05/2023–hiatus: **Created and published on** RECeSS project blog: progress reports on the RECeSS project and introductory blog posts on drug repurposing and collaborative filtering.

12/2016–09/2018: **Published on** Tryalgo [in French] : series of blog posts on known algorithms with concrete applications, aimed at high school and college students (approx. $2,400$ unique monthly users ; two of these posts constitute the Top-2 most visited pages.

10/2016: **Published on** Binaire (blog on Computer Science affiliated with French newspaper *Le Monde*) **and** The Conversation [in French] : "A.P.B. : La vie après le bac" (conjointy written with Serge Abiteboul). Explanation of the algorithm of Gale-Shapley which has been in use in a previous version of the French national web application for high school students' applications to college

Popularization of Law-Making

12/2016–09/2018: **Published on** Réfléchir.fr [in French] : series of blog posts on laws passed since 2017 in France: explanation of their content and their consequences (534 followers on February, 24 2021).