

Frederic COMMO, PhD

Data Scientist - Oncology & Bioinformatics

Project Leader, Precision Cancer Medicine & Molecular Screening Programs











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Frederic Commo

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Work experience

2014 - Present Data Scientist at Gustave Roussy, Villejuif, France

Project Leader, Bioinformatics for Precision Medicine & Molecular Screening Programs

• Identifying molecular alterations matching with targeted therapies. Deployment of data transfer/storage and analysis solutions for molecular screening programs.

• Developing sharable visualization tools to assist therapeutic decision-making. Interactive aCGH Viewer an end-user server solution for visualizing and interacting with genomic profiles. curve fitter, online application for curve fitting and drug response analysis (an extension of the nplr R package).

• Identifying new biomarkers and exceptional responders to treatments (ongoing research)

• Publications: Ann Oncol 2015 (1st author), Bioinformatics 2015 (1st author)

• R package as author: rCGH on Bioconductor (genomic profiling and visualization)

2012 - 2014 Data Scientist - Visiting Scientist at Sage Bionetworks, Seattle, WA

• Steering Committee & Tumour Board Committee, Precision Medicine Programs.

• Computational methods for Precision Cancer Medicine (microarrays & NGS).

• Publications: J Clin Oncol. 2013, Clin Cancer Res. 2013, Mol Cancer Ther. 2013

• R package as author: nplr on CRAN (Logistic regressions for curve fitting and drug response analysis)

2007 - 2012 Project leader at Gustave Roussy, Villejuif, France

Department of Translational Research & Inserm U981 (Pr Fabrice André)

Project management & data analysis, Precision Medicine Program SAFIR01.

Microarray Gene Expression analyses & meta-analyses.

2004-2007 **APHP non-active status** to join Compiègne University of Technology.

1993-2004 **Research Assistant**. Dpt of Pathology, TENON Hospital, Paris.

1986-93 Laboratory technician. Dpt of Pathology, Jean VERDIER Hospital, Bondy, France.

Education

2015 PhD in Oncology & Bioinformatics, University Paris-Saclay

Genomic analysis within precision medicine: optimisations & visualization tools. Supervisor: Pr. Fabrice André

Engineer's Diploma & M. Sci., Compiègne University of Technology (U.T.C.) 2007

Viva voce: Molecular portrait of ependymoma recurrences (Plos One 2010 as co-1st author).

1985 Higher Technician Degree: Lab Technician School, Pitié-Salpêtrière, Paris.

Continued Education

2013 Machine Learning, Caltech, through edX (9 weeks)

Data Analysis, John Hopkins Univ. through Coursera (8 weeks)

• Mathematical Biostatistics, John Hopkins Univ. through Coursera (7 weeks)

2012 • Python programming, MIT, Cambridge, USA through edX (15 weeks)

1995-98 • Ecole Pratique des Hautes Etudes, University Paris-VI, Paris.

• Molecular biology applied to diagnosis, University Paris-VI, Paris. 1991-92

Skills

Data Analysis

• Big Data, Statistical Inferences, Clustering, Data Modelling, Features Extraction.

Programming

• R, Python, SQL, html, CSS, javascript, Cloud Computing.

Languages

• French native, English.

Publications

- Collaboration to 48 *publications* in peer-reviewed international journals.
- Online end-user applications: aCGHviewer, curveFitter, CellExplorer

Teaching

- Statistical methods applied to genomics at *U.T.C. from 2008 to 2010*.
- Applied Statistics at *Ionis Engineering School, Paris, 2012*

Hobbies

Motorcycling, blues guitar, Diving (Advanced PADI certified).