



# Frederic COMMO, PhD

## Data Scientist – Oncology & Bioinformatics

Project Leader, Precision Cancer Medicine & Molecular Screening Programs



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## Work experience

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

Project Leader, Precision Cancer Medicine & Molecular Screening Programs

- Identifying molecular alterations matching with targeted therapies.  
Deployment of data transfer/storage and analysis solutions for molecular screening programs at Gustave Roussy.
- Developing sharable visualization tools to assist therapeutic decision-making.  
*Interactive aCGH Viewer* an end-user server solution for visualising and interacting with genomic profiles.  
*curveFitter*, 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 (1<sup>st</sup> author), *Bioinformatics* 2015 (1<sup>st</sup> 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*

### 2007 Engineering Diploma & M. Sci., Compiègne University of Technology (U.T.C.)

*Viva voce: Molecular portrait of ependymoma recurrences (Plos One 2010 as co-1<sup>st</sup> 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.

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

## Skills

- Data Analysis** • Big Data, Statistical Inferences, Clustering, Data Modelling, Features Extraction.
- Programming Languages** • R, Python, SQL, html, CSS, javascript, Cloud Computing.
- 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).