

# Jean-Baptiste Pettit

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## + EDUCATION

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|-------------|--|
| 2010 – 2014 | <b>University of Cambridge</b> - PhD in statistics/bioinformatics                          |
| 2009 – 2010 | <b>INP-ENSEEIH</b> (French Grande Ecole) - Specialized master's degree in computer science |
| 2007 – 2010 | <b>INP-ENSAT</b> (French Grande Ecole) - Master's degree in life sciences engineering      |
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## + PROFESSIONAL EXPERIENCE

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|----------------------|---|
| 2015-Present         | <b>Insight Data Science, Data Engineering Fellow</b> – Palo Alto, CA <ul style="list-style-type: none"><li>- Designed and developed a data pipeline, API and website for news comments aggregation from multiple sources including RSS and Twitter Streaming feeds.</li><li>- Implemented “Big Data” pipeline to crawled news articles comments and Tweets using Spark for batch processing, Storm for Streaming, Hbase as a data store, Kafka for managing queues deployed on an AWS infrastructure.</li></ul>   |
| 2014-Present         | <b>ITK - Statistical modeling for agronomy, Developer</b> – Montpellier (France) <ul style="list-style-type: none"><li>- Participated and delivered the first release of a Decision Support Tool for agriculture to one of the world's top crop science companies.</li></ul>  |
| 2010-2014<br>4 years | <b>University of Cambridge, PhD student</b> – Cambridge (UK) <ul style="list-style-type: none"><li>- Developed a Hidden Markov Random Field clustering methodology for Genomics data (10 GB of data), published in PloS Computational Biology. Open sourced on Github</li><li>- Created of a statistical spatial localisation methodology for single cell transcriptomics assays (100 GB of data). Accepted in Nature Biotechnologies (in press). Open sourced on Github</li><li>- Released a WebGL/Javascript web application for scientific 3D data visualization (published in BMC Bioinformatics). Open sourced on Github</li></ul> |
| 2010<br>6 months     | <b>European Bioinformatics Institute, Intern</b> – Cambridge (UK) <ul style="list-style-type: none"><li>- Contributed in a JavaEE application for Systems Biology Format Conversion. Deployment of a JSP based web application for SBFC. Open sourced on SourceForge</li></ul>  |
| 2009<br>3 months     | <b>CNRS, Intern</b> – Sisyphe Laboratory – Paris (France) <ul style="list-style-type: none"><li>- Modeled (ODE based) of N<sub>2</sub>O emissions by agricultural soils. Computed GIS indexes in ArcGIS</li></ul>   |

## + SKILLS

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| Statistics  | Bayesian hierarchical models, Markov chains/random fields models, generalized linear mixed models, ODE models, Mixture models, Multivariate analysis. |
| Programming | Functional (Scala), Object Oriented (JavaEE, Python), Imperative (C), Scientific (R, Matlab)  |
| Tools       | Hadoop Map Reduce, Spark, Spark Streaming, Pig, Hive, Kafka, AWS, Hbase, Cassandra, MySQL   |
| languages   | French (native), English (fluent), German (proficient), Spanish (basic), Arabic (basic)   |

## + SIDE PROJECTS

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| Blogging  | <ul style="list-style-type: none"><li>- Paid blogger for 1 year for the leading French Tech blog Presse-citron.net</li><li>- Blogging targeted at the Hacker News community (jboggp)</li></ul>                              |
| Twitter   | <ul style="list-style-type: none"><li>- Created a series of Twitter robots (@WhoTrendedit) analysing trending topics on Twitter and linked to a website (<a href="http://whotrended.it">http://whotrended.it</a>)</li></ul> |
| Freelance | <ul style="list-style-type: none"><li>- Created the back-office employee/database/Ebay interaction system for a leading library in Paris</li></ul>  |