

# Jean-Baptiste Lugagne

Post-doctoral researcher, Biomedical Engineering, Boston University  
44 Cummington Mall  
Boston, MA 02215, United States

+1 617-642-6504  
✉ [jlugagne@bu.edu](mailto:jlugagne@bu.edu)  
🌐 [jblugagne.gitlab.io](http://jblugagne.gitlab.io)

## Education

- 2016 **PhD, Bioengineering**, U. Sorbonne Paris-Cité, France  
*Real-time control of a genetic toggle switch*  
Co-advised by Prof. Grégory Batt, INRIA & Prof. Pascal Hersen, CNRS.
- 2012 **M.S., Signal Processing Engineering**, Grenoble Institute of Technology, France  
*Stochastic simulation of enzymatic reactions under transcriptional feedback regulation*  
Co-advised by Prof. Guy-Bart V. Stan & Dr. Diego A. Oyarzún, Imperial College London.
- 2010 **B.S., Electrical Engineering**, Grenoble Institute of Technology, France

## Research experience

- 2018-  
Present **Post-doctoral researcher**, Boston University, United States  
Advisor: Prof. Mary J. Dunlop  
Control theory • Deep learning • Synthetic biology • Antibiotic resistance • Metabolic engineering  
• Optogenetics • Microfabrication & microfluidics • Image processing
- Developed a novel approach for real-time feedback control of gene expression, including for antibiotic resistance genes, in thousands of single *E. coli* cells based on deep model predictive control and optogenetics.
  - Created DeLTA, a deep learning-based software for single cell segmentation and tracking in time-lapse microscopy movies.
  - Established collaboration with Prof. Ji-Xin Cheng's group for label-free stimulated Raman spectro-microscopy imaging of bioproduction in *E. coli* cells.
- 2012-  
2017 **Graduate research assistant & Post-doctoral researcher**, CNRS & INRIA, Paris, France  
Advisors: Prof. Pascal Hersen & Prof. Grégory Batt  
Control theory • Synthetic biology • Microfabrication & microfluidics • Automated microscopy • Machine learning • 3D printing & electronic prototyping
- Controlled gene expression in a genetic toggle switch to stabilize it around its unstable equilibrium point in single cells, with real-time feedback or dynamic forcing.
  - Developed microscopy image analysis algorithm based on classification of pixel's intensity signatures in a Z-stack with support vector machines or random forests.
- 2012 **Graduate research assistant**, Imperial College, London, UK  
Advisors: Prof. Guy-Bart V. Stan & Dr. Diego A. Oyarzún  
Systems biology • Stochastic modelling of gene expression and metabolism • Control theory
- Investigated propagation of genetic and metabolic noise in a metabolic pathway under transcriptional feedback.
- 2011 **Graduate research assistant**, INRIA, Grenoble, France  
Advisor: Prof. Hidde de Jong  
Systems & synthetic biology • Deterministic modelling of gene expression • Signal processing
- Modeled and fit parameters to experimental data of spatio-temporal dynamics in a colony of synthetic *E. coli* cells coupling quorum sensing and toggle switch circuits for biosensing.

## Publications

†: co-corresponding author – \*: co-first author

### IN REVISION

**Lugagne, J.-B.**<sup>†</sup>, Blassick, C. M., Dunlop, M. J.<sup>†</sup> (2022). Deep model predictive control of gene expression in thousands of single cells. *bioRxiv*. [10.1101/2022.10.28.514305](https://doi.org/10.1101/2022.10.28.514305)

### PUBLISHED ARTICLES

Klumpe, H. E.\*, **Lugagne, J.-B.**<sup>\*†</sup>, Khalil, A. S., Dunlop, M. J.<sup>†</sup> (2023). Deep neural networks for predicting single cell responses and probability landscapes. *ACS Synthetic Biology*. [10.1021/acssynbio.3c00203](https://doi.org/10.1021/acssynbio.3c00203)

Tague, N., Lin, H., **Lugagne, J.-B.**, O'Connor, O. M., Burman, D., Wong, W. W., Cheng, J.-X., Dunlop, M. J. (2023). Longitudinal single-cell imaging of engineered strains with stimulated Raman scattering to characterize heterogeneity in fatty acid production. *Advanced Science*. [10.1002/advs.202206519](https://doi.org/10.1002/advs.202206519)

**Lugagne, J.-B.**, & Dunlop, M. J. (2022). Anticipating antibiotic resistance. *Science*. [Perspective] [10.1126/science.abn9969](https://doi.org/10.1126/science.abn9969)

Sampaio, N. M. V, Blassick, C. M., Andreani, V., **Lugagne, J.-B.**, & Dunlop, M. J. (2022). Dynamic gene expression and growth underlie cell-to-cell heterogeneity in Escherichia coli stress response. *PNAS*. [10.1073/pnas.2115032119](https://doi.org/10.1073/pnas.2115032119)

O'Connor, O. M., Alnahhas, R. N., **Lugagne, J.-B.**<sup>†</sup>, & Dunlop, M. J.<sup>†</sup> (2022). DeLTA 2.0: A deep learning pipeline for quantifying single-cell spatial and temporal dynamics. *PLOS Computational Biology*. [10.1371/journal.pcbi.1009797](https://doi.org/10.1371/journal.pcbi.1009797)

Lin, H., Lee, H. J., Tague, N., **Lugagne, J.-B.**, Zong, C., Deng, F., Shin, J., Tian, L., Wong, W., Dunlop, M. J., & Cheng, J.-X. (2021). Microsecond fingerprint stimulated Raman spectroscopic imaging by ultrafast tuning and spatial-spectral learning. *Nature Communications*. [10.1038/s41467-021-23202-z](https://doi.org/10.1038/s41467-021-23202-z)

**Lugagne, J.-B.**, Lin, H., & Dunlop, M. J. (2020). DeLTA: Automated cell segmentation, tracking, and lineage reconstruction using deep learning. *PLOS Computational Biology*. [10.1371/journal.pcbi.1007673](https://doi.org/10.1371/journal.pcbi.1007673)

**Lugagne, J.-B.**, & Dunlop, M. J. (2019). Cell-machine interfaces for characterizing gene regulatory network dynamics. *Current Opinion in Systems Biology*. [10.1016/j.coisb.2019.01.001](https://doi.org/10.1016/j.coisb.2019.01.001)

**Lugagne, J.-B.**, Jain, S., Ivanovitch, P., Ben Meriem, Z., Vulin, C., Fracassi, C., Batt, G., & Hersen, P. (2018). Identification of individual cells from z-stacks of bright-field microscopy images. *Scientific Reports*. [10.1038/s41598-018-29647-5](https://doi.org/10.1038/s41598-018-29647-5)

**Lugagne, J.-B.**, Kirch, M., Köhler, A., Batt, G., & Hersen, P. (2017). Balancing a genetic toggle switch by real-time feedback control and periodic forcing. *Nature Communications*. [10.1038/s41467-017-01498-0](https://doi.org/10.1038/s41467-017-01498-0)

**Lugagne, J.-B.**, Brackx, G., Seyrek, E., Berret, J.-F., Hersen, P., & Charron, G. (2017). Assembly and characterizations of bifunctional fluorescent and magnetic microneedles displaying length tunability over one decade. *Advanced Functional Materials*. [10.1002/adfm.201700362](https://doi.org/10.1002/adfm.201700362)

Piffoux, M., Silva, A. K. A., **Lugagne, J.-B.**, Hersen, P., Wilhelm, C., & Gazeau, F. (2017). Extracellular vesicle production loaded with nanoparticles and drugs in a trade-off between loading, yield and purity: Towards a personalized drug delivery system. *Advanced Biosystems*. [10.1002/adbi.201700044](https://doi.org/10.1002/adbi.201700044)

Oyarzún, D. A., **Lugagne, J.-B.**, & Stan, G. B. V. (2014). Noise propagation in synthetic gene circuits for metabolic control. *ACS Synthetic Biology*. [10.1021/sb400126a](https://doi.org/10.1021/sb400126a)

**Lugagne, J.-B.**, Oyarzún, D. A., & Stan, G. B. V. (2013). Stochastic simulation of enzymatic reactions under transcriptional feedback regulation. In *IEEE European Control Conference 2013 (ECC13)*. [10.23919/ECC.2013.6669756](https://doi.org/10.23919/ECC.2013.6669756)

## Research presentations

### TALKS

**Lugagne, J.-B.**, Klumpe, H. E., Blassick, C. M., & Dunlop, M. J. (2023). High-throughput single-cell control using real-time feedback. *SwissUK Synbio*, Lausanne, Switzerland.

**Lugagne, J.-B.**, & Dunlop, M. J. (2022). High-throughput single-cell control using real-time feedback. *Biocontrol seminars*, Online.

**Lugagne, J.-B.**, & Dunlop, M. J. (2022). High-throughput single-cell control using real-time feedback. *Winter Q-Bio 2022*, Ko Olina, Hawaii, USA.

**Lugagne, J.-B.**, & Dunlop, M. J. (2021). Deep model predictive control of gene expression in single cells. *BDC Spring Symposium*, Boston University, USA.

**Lugagne, J.-B.**, & Dunlop, M. J. (2019). Single-cell real-time feedback control of gene expression. *BDC Symposium*, Boston University, USA.

**Lugagne, J.-B.**(2017). Balancing a genetic toggle switch by real-time feedback control and periodic forcing. ETH Zürich, Switzerland.

**Lugagne, J.-B.**(2017). Balancing a genetic toggle switch by real-time feedback control and periodic forcing. University of Edinburgh, United-Kingdom.

**Lugagne, J.-B.**(2017). Balancing a genetic toggle switch by real-time feedback control and periodic forcing. University of Cambridge, United-Kingdom.

**Lugagne, J.-B.**, Kirch, M., Köhler, A., Batt, G., & Hersen, P. (2015). Real-time control of a genetic toggle switch. *Design, Optimization and Control in Systems and Synthetic Biology*, Paris, France.

**Lugagne, J.-B.**, Oyarzún, D. A., & Stan, G. B. V. (2013). Stochastic simulation of enzymatic reactions under transcriptional feedback regulation. *IEEE European Control Conference 2013 (ECC13)*, ETH Zürich, Switzerland.

### CONFERENCE POSTERS

**Lugagne, J.-B.**, Tague, N., & Dunlop, M. J. (2019) Single-cell, dynamic interrogation of antibiotic resistance acquisition. *Q-Bio 2019*, San Francisco, California, USA.

**Lugagne, J.-B.**, Tague, N., & Dunlop, M. J. (2019) Dynamic control of antibiotic resistance related genes with optogenetics. *EBRC Retreat*, Boston, Massachusetts, USA.

**Lugagne, J.-B.**, Kirch, M., Köhler, A., Batt, G., & Hersen, P. (2017). Balancing a genetic toggle switch by real-time feedback control and periodic forcing. *International Workshop on Control Engineering and Synthetic Biology*, London, United-Kingdom.

**Lugagne, J.-B.**, Kirch, M., Köhler, A., Batt, G., & Hersen, P. (2015). Real-time control of a genetic toggle switch. *Who Am I Colloquium*, Deauville, France.

## Grant writing

National Science Foundation grant [2032357](https://www.nsf.gov/awardsearch/showAward.do?awardNumber=2032357)

“Single-cell feedback, optogenetics, and deep learning to control gene expression in bacteria”

- Came up with complete concept for grant.
- Wrote entire draft of proposal.

- Worked with Dr. Dunlop on grantsmanship and refining project objectives.
- Responded to reviewer feedback.
- Grant was funded.

Department of Energy grant [DE-SC0019387](#)

“High-throughput chemical imaging for optimizing biofuel synthesis using synthetic biology”

- Wrote all of one objective (out of three).
- Generated preliminary data demonstrating feasibility of project goals.
- Worked with collaborators and project PIs to develop proposal ideas.
- Grant was funded.

National Science Foundation grant [2143289](#)

“Transitions: Deep Learning Models for Microbial Image Analysis and Time-Series Predictions”

✂

National Institutes of Health grant [R01AI102922](#)

“Cell-to-cell heterogeneity and the emergence of antibiotic resistance”

- Provided feedback and input on early drafts.
- Both grants were funded.

## Teaching ✂ mentoring experience

- Guest lecturer in BE404 Control Theory, Boston University, Under-graduate and graduate level.
- Guest lecturer in BE403 Signal Processing, Boston University, Under-graduate and graduate level.
- Supervision of 2 Master’s students theses on mathematical modelling of real-time control of a genetic toggle switch.
- Supervision of 2 Master’s students projects on machine learning based image analysis of microscopy images.
- Co-supervision of a Master’s student thesis on biological laboratory design and social interactions within a research group.
- Supervision of 5 PhD rotation projects, on a range of topics including CRISPR-based genomic integration, deep learning-based image processing, or synthetic circuit design.
- Supervision of 2 Senior design project teams, for the development of automated sample preparation devices, and for a cloud-computing based web interface for microscopy image analysis.
- Supervision of International Genetically Engineered Machine competition (iGEM) Grenoble team 2012, especially for mathematical modelling of gene expression.

## Summary of qualifications ✂ proficiencies

### TECHNICAL SKILLS

Deep model-predictive control • Reinforcement learning

Microbiology • Molecular biology • Optogenetics

Deterministic ✂ stochastic modelling of genetic and metabolic networks

Computer vision • Deep learning

Microfabrication ✂ microfluidics • Microscopy • Automation • Instrumentation

3D printing • Rapid prototyping • Electronics  
Python • Matlab • C/C++ • Tensorflow & Pytorch • Git • Unix/Linux

#### LEADERSHIP & COMMUNICATION

10+ years of scientific project management, as part of a team, a larger collaboration, or as main researcher, from conception to publication and follow-up studies.

10+ years of scientific communication, by presenting at conferences and writing journal articles.

Collaboration on open-source and open data research projects and active support for the community, for example on a software for microscopy time-lapse analysis ([DeLTA](#)).

Co-organizer of the [Biocontrol seminars](#), a series of monthly online seminars featuring international speakers.

## References

Prof. Mary J. Dunlop – Boston University – Post-doctoral advisor – [mjdunlop@bu.edu](mailto:mjdunlop@bu.edu)

Prof. Pascal Hersen – CNRS & Curie Institute – PhD advisor – [pascal.hersen@curie.fr](mailto:pascal.hersen@curie.fr)

Prof. Grégory Batt – INRIA & Pasteur Institute – PhD advisor – [gregory.batt@inria.fr](mailto:gregory.batt@inria.fr)

Prof. Ji-Xin Cheng – Boston University – Post-doctoral collaborator – [jxcheng@bu.edu](mailto:jxcheng@bu.edu)