

SARA FONSECA COSTA

Postdoctoral Researcher

I am a data scientist/bioinformatician at the University of Lausanne.

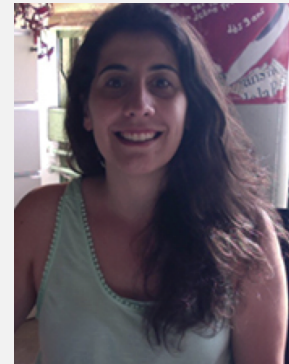
My research interests include aging, circadian clocks, genomics and transcriptomics. I'm working at the moment with single-cell transcriptomics data, in order to integrate this data type in the Bgee database. I use mainly R for data analysis and visualization.

EDUCATION

- 2013
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2017
- **University of Fribourg**
PhD Biochemistry 📍 Fribourg, Switzerland
- Thesis: Model based inference of photoperiod and age related changes in circadian oscillators.
- Advisor & Co-advisors: Prof. Dr. Urs Albrecht, Dr. Jürgen Ripperger and Prof. Dr. Daniel Wegmann
- Jan 2012
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Out 2012
- **University of Geneva**
Master thesis 📍 Geneva, Switzerland
- Thesis: Study of an automatization model to analyse genetic diversity in human populations.
- Advisors: Dr. José Nunes (UNIGE) and Prof. Dr. Miguel Rocha (UM)
- 2010
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2012
- **University of Minho**
M.S. in Bioinformatics 📍 Braga, Portugal
- 2006
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2010
- **Polytechnic Institute of Bragança**
B.S. in Biotechnology Engineering 📍 Bragança, Portugal

RESEARCH EXPERIENCE

- Jul 2018
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current
- **Postdoctoral Researcher**
University of Lausanne 📍 Lausanne, Switzerland
- Data analysis and integration of single cell RNA-Seq data in Bgee database
 - Maintenance and update of the Bgee pipeline (together with other Bgee members).
- Sep 2017
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Dec 2017
- **Postdoctoral Researcher**
Walter and Elisa Hall Institute / SIB 📍 Lausanne, Switzerland
- Advisor & Co-advisor: Prof. Dr. Terry Speed (WEHI) and Prof. Dr. Mauro Delorenzi (SIB)
- Improvement of data analysis of circadian and aging data using new statistical approaches.



CONTACT INFO

✉ sara.fonsecacosta@unil.ch

🔗 github.com/SFonsecaCosta

ORCID - <https://orcid.org/0000-0001-7794-7997>

☎ +4121 692 4221

For more information, please contact me via email.

SKILLS

Experienced in molecular biology, biochemistry, genetics and functional genomics.

Experienced in statistical analysis.

Full experience with next generation sequencing data analysis.

Highly skilled in R, Bash, LaTeX

Knowledge of: Python, Perl, C++

*This resume was made with the R package **pagedown**.*

Last updated on 2020-10-21.

PROFESSIONAL EXPERIENCE

2009
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2011

- **Student researcher**
University of Porto - Department of Molecular Pathology and Immunology, Hospital Santo Antonio
Molecular Biology and Bioinformatics (HLA OSAS)
Advisor: Prof. Dr. Berta Martins
Porto, Portugal

TEACHING EXPERIENCE

2019
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2020

2018
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2020

2018
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2019

2013
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2017

2014
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2015

2013
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2015

- **Molecular evolution (Bachelor)**
Teaching assistant
Lausanne, Switzerland
- **Molecular genetics methods (Master)**
Teaching assistant
Lausanne, Switzerland
- **Data analysis (Bachelor)**
Teaching assistant
Lausanne, Switzerland
- **TP's Biochemistry**
Teaching assistant
Fribourg, Switzerland
- **Bioinformatics practical Classes**
Teaching assistant
Fribourg, Switzerland
- **Genotyping by Southern Blot Hybridization**
Teaching assistant
Fribourg, Switzerland

ORGANISER

2020

2012

- **Gene expression made useful easily: tools and database of Bgee**
Organizing team
UNIL, Lausanne, Switzerland
- **Bioinformatics Open Days 2012**
Organizing Committee
UM, Braga, Portugal

PEER REVIEW PUBLICATIONS

2020

- **The Bgee suite: integrated curated expression atlas and comparative transcriptomics in animals**
Nucleic Acids Research, <https://doi.org/10.1093/nar/gkaa793>
Frederic B Bastian, Julien Roux, Anne Niknejad, Aurélie Comte, Sara S Fonseca Costa, Tarcisio Mendes de Farias, Sébastien Moretti, Gilles Parmentier, Valentine Rech de Laval, Marta Rosikiewicz, Julien Wollbrett, Amina Echchiki, Angélique Escoriza, Walid H Gharib, Mar Gonzales-Porta, Yohan Jarosz, Balazs Laurency, Philippe Moret, Emilie Person, Patrick Roelli, Komal Sanjeev, Mathieu Seppey, Marc Robinson-Rechavi

Practical available:
<https://github.com/SFonsecaCosta/Bgee>

Info available:
<http://hdl.handle.net/1822/33574>

- 2020 ● **Single-cell transcriptomics allows novel insights into aging and circadian processes**
Briefings in Functional Genomics,
<https://doi.org/10.1093/bfpg/elaa014>
Sara S Fonseca Costa, Marc Robinson-Rechavi and Jürgen A Ripperger
- 2017 ● **Normalisation against Circadian and Age-Related Disturbances Enables Robust Detection of Gene Expression Changes in Liver of Aged Mice**
PloS one 12 (1), <https://doi.org/10.1371/journal.pone.0169615>
Sara S Fonseca Costa, Daniel Wegmann, Jürgen A Ripperger
- 2017 ● **Internal normalization of nascent RNA sequencing experiments to identify age-related changes of the mouse liver transcriptome**
Genomics and Computational Biology 3 (2),
<https://doi.org/10.18547/gcb.2017.vol3.iss2.e34>
Sara S Fonseca Costa and Jürgen A. Ripperger
- 2016 ● **REV-ERB α influences the stability and nuclear localization of the glucocorticoid receptor**
Journal of cell science 129 (21), <https://doi.org/10.1242/jcs.190959>
Takashi Okabe, Rohit Chavan, Sara S Fonseca Costa, Andrea Brenna, Jürgen A Ripperger, Urs Albrecht
- 2016 ● **Liver-derived ketone bodies are necessary for food anticipation**
Nature communications 7 (1), 1-10,
<https://doi.org/10.1038/ncomms10580>
Rohit Chavan, Céline Feillet, Sara S Fonseca Costa, James E Delorme, Takashi Okabe, Jürgen A Ripperger, Urs Albrecht
- 2015 ● **Impact of the circadian clock on the aging process**
Frontiers in neurology 6, 43,
<https://doi.org/10.3389/fneur.2015.00043>
Sara S Fonseca Costa and Jürgen A. Ripperger



BOOK CHAPTERS

- 2020 ● **Chromatin Immunoprecipitation (ChIP) from Mouse Liver Nuclei**
Springer US - [DOI:10.1007/978-1-0716-0381-9](https://doi.org/10.1007/978-1-0716-0381-9)
Sara S. Fonseca Costa and Jürgen A. Ripperger



OPEN SOURCE CONTRIBUTIONS

- 2019 ● **BgeeCall R Package**
Contribution
Julien Wollbrett[aut], Julien Roux[aut], Sara Fonseca Costa[ctb], Marc Robinson Rechavi[ctb], Frederic Bastian[aut]

<https://www.bioconductor.org/packages/release/workflows/html/Bgee-Call.html>

CONFERENCE PRESENTATIONS

- 2019 ● **Expression calls from RNA-seq using BgeeCall R package**
Talk + poster presentation, Bioconductor conference - eurobioc2019
📍 UCLouvain, Brussels, Belgium
Sara Fonseca Costa
- 2016 ● **Internal normalization of nascent RNA sequencing experiments to identify age-related changes of the mouse liver transcriptome**
Talk presentation, 2nd Challenges in Computational Biology: Gene Expression Data Analysis
📍 Mainz, Germany
Sara S Fonseca Costa

LANGUAGE

- **Portuguese**
Native
- **English**
Fluente
- **French**
Basic
- **Spanish**
Basic