

ADRIÀ MITJAVILA VENTURA

MSc | Bioinformatician at *Institut Josep Carreras*

Biotechnologist with training in Molecular Genetics and Bioinformatics. Interested in omics data analysis to study the mechanisms of gene regulation in mammals. See more about me on [my website](#).



EXPERIENCE

08/2021
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Present

Bioinformatician

Institut Josep Carreras

📍 Badalona, Spain

Study of PIWI-interacting RNAs (piRNAs), their biogenesis and variation across species. | Development of a pipeline for processing small RNA-seq data focused to analyze piRNAs.

07/2019
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07/2021

Bioinformatician

Istituto Europeo di Oncologia

📍 Milan, Italy

Study of Wnt/ β catenin signaling pathway -and others- in the context of colorectal cancer and stem cell development. | Analysis and integration of RNA-seq, ChIP-seq and ATAC-seq data.

09/2017
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10/2018

Master's thesis student & Research assistant

Instituto de Biomedicina de Sevilla

📍 Seville, Spain

Study of Xrn1 and Ccr4 roles in yeast nucleosome architecture and gene transcription. | Analysis of MNase-seq data. | Performance and analysis of MNase-qPCR experiments.

09/2016
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06/2017

Internship & Bachelor's thesis student

Microbial. Sistemas i aplicacions analítiques, S.L.

📍 Girona, Spain

Microbiological and molecular analyses for pathogen detection and food forensics. | Study of butyrate effects on the ileal microbiota of colectomized colorectal cancer patients.



EDUCATION

2018
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2022

MSc in Bioinformatics and Biostatistics

Universitat Oberta de Catalunya (UOC)

📍 Online

Overall grade: 8.94 | Master's thesis: *Analysis of the variation in piRNA expression across three Mus species* (thesis grade: 9.78/10, with honours).

2017
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2018

MSc in Molecular Genetics and Biotechnology

Universidad de Sevilla (US)

📍 Seville, Spain

Overall grade: 9.33/10 | Master's thesis: *Influence of Xrn1 in nucleosome positioning across Saccharomyces cerevisiae genome and its effects in transcription* (thesis grade: 9.2/10).

2012
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2017

BSc in Biotechnology

Universitat de Girona (UdG)

📍 Girona, Spain

Overall grade: 8.12/10 | Bachelor's thesis: *Butyrate-induced changes in the diversity of intestinal mucosa-associated microbiota in colorectal cancer patients submitted to a lateral ileostomy* (thesis grade: 9.4/10)



PUBLICATIONS

Journal
article

Begley *et al.* Xrn1 influence on gene transcription results from the combination of general effects on elongating RNA pol II and gene-specific chromatin configuration. *RNA Biology* 2020. doi: [10.1080/15476286.2020.1845504](https://doi.org/10.1080/15476286.2020.1845504)

CONTACT

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📞 0000-0002-0337-1929

SKILLS

Bioinformatics:

Microarray, (small) RNA- / ChIP- / ATAC- / MNase-seq

Informatics:

R/Bioconductor, R Markdown, Shiny, Bash, Snakemake, Python, Git, HPC, Conda, Docker, Singularity

Experimental:

(RT)-(q) PCR, primer design, electrophoresis, Western Blot, MNase-digestion, bacterial culture

Languages:

Catalan (Native), Spanish (Native), English (C1 - IELTS 7.5), Italian (Basic)

CODE

[ggmitji](#)
[plotmics](#)
[chromHMMviewR](#)
More on [my GitHub profile](#)

AWARDS

[INPhINIT Incoming PhD Fellowship](#) (LaCaixa, 2022-2025)

[Master's thesis, with honours](#) (UOC, 2022)

[Eliteathlete](#) (BOE, 2015-2021)

[World Rowing U23 Champion](#) (FISA, 2016)

References available upon request.
Build with [pagedown](#), code [here](#).
Last updated on 2022-06-09.