ADRIÀ MITJAVILA VENTURA

Bioinformatician at Istituto Europeo di Oncologia

Biotechnologist living around computers, interested in data analysis to support experimental research as well as in pure (bio)computational experiments and in software/pipeline development.

EXPERIENCE Bioinformatician 07/2019 Istituto Europeo di Oncologia Present • Analysis and integration of RNA-seq, ChIP-seq and ATAC-seq data. • Development of an ATAC-seq pipeline for Snakemake. Master's thesis student & research assistant 09/2017 Instituto de Biomedicina de Sevilla 10/2018 • Analysis of MNase-seq data. • Performance and analysis of MNase-qPCR experiments. Internship & Bachelor's thesis student 09/2016 Microbial. Sistemes i aplicacions analítiques, S.L. 06/2017

EDUCATION

2018

Present

2017

2018

2012

2017

MSc in Bioinformatics and Biostatistics

MSc in Molecular Genetics and Biotechnology

Universitat Oberta de Catalunya

• qPCR of intestinal biopsies to detect/quantify bacterial genres.

Universidad de Sevilla

Seville, Spain

• Master's thesis: Influence of Xrn1 in nucleosome positioning across Saccharomyces <u>cerevisiae</u> genome and its effects in transcription.

• Statistical analysis of high-throughput sequencing data to quantify bacteria species.

BSc in Biotechnology

Girona, Spain Universitat de Girona

• Bachelor's thesis: Butyrate-induced changes in the diversity of intestinal mucosaassociated microbiota in colorectal cancer patients submitted to a lateral ileostomy.



CONTACT

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SKILLS

Omics data analysis:

Microarray RNA-seq ChIP-seq ATAC-seq MNase-seq

Programming/scripting:

R/Bioconductor Python Bash

Other:

Snakemake Conda Docker Git

Languages:

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

References

Available upon request

