Davide **BRESSAN** PhD Student | Bioinformatician/Computational biologist

in linkedin.com/in/davide-bressan **O** github.com/DavideBrex

🛿 +39 3450928188 🛛 @ davide.bressan-1@unitn.it

v Via Montecorno 6A, Sona - VR, Italy

☑ davidebrex.github.io/



I am a PhD Student in the Chiacchiera Lab at the Department of Cellular, Computational and Integrative Biology, University of Trento, Italy. Trained as computer scientist and bioinformatician, I am now working as computational biologist in the field of cancer research. My PhD focuses on the epigenetics of liver regeneration and tumorigeneisis.

WORK EXPERIENCE

Current November 2020	 PhD Student - Doctoral Program in Biomolecular Sciences, UNIVERSITY OF TRENTO, Italy > PhD student in the Fulvio Chiacchiera group (Laboratory of Stem Cells and Cancer Genomics). > Transdisciplinary program in Computational Biology RNA-Seq ChIP-Seq Genetics Cancer R
Current November 2022	 Teaching assistant, course of Scientific Programming for Data Science, UNIVERSITY OF TRENTO, Italy Preparation and delivery of computer laboratory lectures for the Master in Data Science. Exam preparation Python Data structures Algorithms Student supervision
July 2020 March 2020	 Intern Bioinformatician, WINCLOVE PROBIOTICS B.V, Amsterdam, Netherlands > Trainee in the company's research and innovation team. > Analysis of metagenomic data from patients with inflammatory bowel disease using machine learning algorithms Metagenomics Machine learning Python
July 2020	Teaching assistant, course of Fundamentals of Bioinformatics, VRIJE UNIVERSITEIT, Amsterdam,
_	Netherlands
March 2020	 Guidance and evaluation of students in the first year of the Master in Bioinformatics and System Biology Python Student supervision
February 2020 August 2019	 Intern Bioinformatician, NETHERLANDS CANCER INSTITUTE (NKI), Amsterdam, Netherlands Internship as a bioinformatician in the group of Prof. Daniel Peeper (Department of Molecular Oncology and Immunology) Development of an R program for the analysis of splice variants. Thesis tityle: "The role of transcript variants in immunotherapy resistant melanoma". RNA-Seq R Melanoma

📂 Formation

Master of Science in Bioinformatics and System Biology - Universiteit van Amsterdam - Vrije Universiteit 2018-2020 Amsterdam - Netherlands

Bachelor's Degree in Bioinformatics – University of Verona - Italy 2015-2018

S LANGUAGE

Italian	$\bullet \bullet \bullet \bullet \bullet$
English	$\bullet \bullet \bullet \bullet \bigcirc$
Dutch	$\bullet \circ \circ \circ \circ$



- > Adaptability
- > Presentation and reporting
- > Autonomy

PUBLICATIONS AND POSTERS

INCREASED GENOMIC INSTABILITY AND RESHAPING OF TISSUE MICROENVIRONMENT UNDERLIE ONCOGENIC PROPERTIES OF ARID1A MUTATIONS

Science Advances - DOI: 10.1126/sciadv.adh4435 Co-first author

Arid1A Liver Cancer Mutagenesis Bioinformatics

${\rm Rar}\gamma{\rm -}{\rm Foxa1}$ signaling promotes luminal identity in prostate progenitors and is disrupted in prostate cancer

Preprint on biorxiv - DOI: https://doi.org/10.1101/2024.03.06.583256 Co-author

Ar Foxal Prostate Cancer Bioinformatics

Increased genomic instability and reshaping of tissue microenvironment drive ARID1A-dependent cancer formation

Oral Talk at SIBBM 2023 - Frontiers in Molecular Biology June 2023, Bari, Italy mSWI/SNF Liver Cancer Arid1A Bioinformatics

The role of mSWI/SNF complex in preserving liver tissue homeostasis and preventing hepatocellular carcinoma formation

Poster presented at the 33° PEZCOLLER SYMPOSIUM - Presenting author June 2022, Trento, Italy mSWI/SNF [Liver Cancer]

${\sf Role of the } {\sf mSWI/SNF} \ {\sf chromatin remodeling \ complex \ in \ preserving \ liver \ homeostasis}$

Poster presented at the EACR conference - Presenting author June 2022, Seville, Spain mSWI/SNF Liver Cancer

S References

Fulvio Chiacchiera

Assistant Professor, UNIVERSITY OF TRENTO

- @ fulvio.chiacchiera@unitn.it
- **L** +39 0461 283934

Oscar krijgsman

Principle Bioinformatician, NEOGENE THERAPEUTICS

@ o.krijgsman@nki.nl