

Flavia Villani
Curriculum Vitae

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| Personal Data | Place and Date of Birth: Salerno 14 November 1995 phone: +39 3288064640 email: flavia.villani14@gmail.com page GitHub: https://github.com/Flavia95 |
| Education | <div>Oct 2018-July 2020 Master's degree in Medical Biotechnology (LM-9) University of Naples Federico II (NA) "Population genomics analyses on pangenome graphs"</div> <div>Oct 2014-Feb 2018 Bachelor's degree in Biological Sciences (L-13) University of Salerno (SA) "L'enzima glucocinasi e la cura del diabete di tipo II"</div> |
| Training activity | <div>Apri 2020 COVID-19 Biohackathon/Pangenome https://github.com/virtual-biohackathons/covid-19-bh20/wiki/Pangenome Course Virtual</div> <div>Nov 2019 Analysis of scRNA-seq using R Course NETTABB-University of Salerno, Italy</div> <div>July 2019 Linux Shell, Git and Python Course Software Carpentry Workshop and Elixir-ITA / National Research Council, Napoli, Italy</div> <div>Dec 2017 EndNote Web, citation and bibliography program Course University of Salerno, Italy</div> |
| Research activity | <div>2019-2020 Master's degree in a laboratory of computational biology at National Research Council Institute of Genetics and Biophysics, IGB-CNR (NA). Thesis title "Population genomics analyses on pangenome graphs"<ul style="list-style-type: none">• Study and application of programming languages (python, bash, awk, R).• Use of high-performance machines with associated queuing systems (condor).• Elaboration of a library in Python for the analysis of population genetics on pangenomic data (https://github.com/Flavia95/VGpop).</div> <div>2017 Training for Bachelor in Biological Science at Clinical analysis laboratory Biolabor-SRL (SA)<ul style="list-style-type: none">• Knowledge of laboratory equipment and analytical methods. Use of clinical medicine machinery based on the Lambert-Beer law to determine clinical parameters.• Determination of blood count, chemical-physical and parasitological examinations of faeces, glycated hemoglobin, ESR, antibiogram, urin analysis, microscopic observation of urinary sediments, tests for the determination of blood group, HIV tests, HPV, hormonal tests and observation of serum protein.</div> |

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| Congress Seminar | Sep 2020 | T2T-HPRC Scalable variant detection in pangenome models, Virtual | Poster |
| | Sep 2020 | T2T-HPRC Easy and Efficient Variation Graphs, Virtual | Poster |
| | Apr 2020 | Graphical pangenomics in the time of COVID19 Virtual, Helsinki | Seminar |
| | Feb 2020 | Modeling neurogenesis with human-derived cerebral organoids Seminar National Research Council, Naples, Italy | |
| | Oct 2019 | Ginecologia e Donna Un viaggio di tutta una vita SIGO-Naples, Italy | Congress |
| | Nov 2019 | NETTAB/BBCC2019 University of Salerno, Italy | Congress |
| | Apr 2016 | OGM: fatti, misfatti e prospettive University of Salerno, Italy | Seminar |

Publications Jordan M Eizenga, Adam M Novak, Emily Kobayashi, *Flavia Villani*, Cecilia Cisar, Simon Heumos, Glenn Hickey, Vincenza Colonna, Benedict Paten, Erik Garrison (16 July 2020). Efficient dynamic variation graphs. Bioinformatics.
<https://doi.org/10.1093/bioinformatics/btaa640>

Certificates 21 Apr 2018 **ESB Level 1 Certificate in ESOL International All Modes (B2 CEFR)**
English language certification
Candidate Number: IT-0042278
28 Mar 2015 **Primo soccorso B.L.S.D (Basic Life Support with defibrillation) e ostruzione delle vie aeree**
Salerno, Italy Certificate BLSD

Languages English - (B2)

Computer Skills *Programming:* Python, R, Bash, LaTeX.
Computer Software: Linux, Windows.

Other information

September-October 2020 Helper in "Corso di Genomica e Proteomica computazionale per biologi sperimentali"

June-August 2020 Helper in GSOC - Parallel Graph Traversal for Variation Graphs project: <https://gsocgraph.blogspot.com/2020/08/final-week-recap-of-my-gsoc-experience.html>

April 2020 Quote in the newspaper "Le Monde": (<https://www.lemonde.fr/blog/binaire/2020/05/06/sars-cov-2-et-covid-19-on-va-jouer-sur-les-mots/>)