

**Flavia Villani**  
*Curriculum Vitae*

---

<b>Personal Data</b>	Place and Date of Birth:	Salerno   14 November 1995	
	phone:	+39 3288064640	
	email:	flavia.villani14@gmail.com	
	page GitHub:	<a href="https://github.com/Flavia95">https://github.com/Flavia95</a>	
<b>Education</b>	Oct 2018-July 2020	<b>Master's degree in Medical Biotechnology (LM-9)</b> Federico II (NA) "Population genomics analyses on pangenome graphs"	University of Naples
	Oct 2014-Feb 2018	<b>Bachelor in Biological Sciences (L-13)</b> "L'enzima glucocinasi e la cura del diabete di tipo II"	University of Salerno (SA)
<b>Training activity</b>	Apri 2020	<b>COVID-19 Biohackathon / Pangenome Assembly</b> Virtual-USA, Japan, Germany	Course
	Nov 2019	<b>Analysis of scRNA-seq using R</b> NETTABB-University of Salerno, Italy	Course
	July 2019	<b>Linux Shell, Git and Python</b> Software Carpentry Workshop and Elixir-ITA / National Research Council, Napoli, Italy	Course
	Dec 2017	<b>EndNote Web, citation and bibliography program</b> University of Salerno, Italy	Course
<b>Research activity</b>	2019-2020	<b>Master's degree</b> in 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"><li>• Study and application of programming languages (python, bash, awk, R).</li><li>• Use of high-performance machines with associated queuing systems (condor).</li><li>• Elaboration of a library in Python for the analysis of population genetics on pangenomic data (<a href="https://github.com/Flavia95/VGpop">https://github.com/Flavia95/VGpop</a>).</li></ul>	
	2017	<b>Training for Bachelor in Biological Science</b> at Clinical analysis laboratory Biolabor-SRL (SA)	
		<ul style="list-style-type: none"><li>• Knowledge of laboratory equipment and analytical methods. Use of clinical medicine machinery based on the Lambert-Beer law to determine clinical parameters.</li><li>• 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.</li></ul>	

<b>Congress Seminar</b>	Apr 2020	<b>Graphical pangenomics in the time of COVID19</b> Virtual, Helsinki	Seminar
	Feb 2020	<b>Modeling neurogenesis with human-derived cerebral organoids</b> Seminar National Research Council, Naples, Italy	
	Oct 2019	<b>Ginecologia e Donna Un viaggio di tutta una vita</b> SIGO-Naples, Italy	Congress
	Nov 2019	<b>NETTAB/BBCC2019</b> University of Salerno, Italy	Congress
	Apr 2016	<b>OGM: fatti, misfatti e prospettive</b> University of Salerno, Italy	Seminar
<b>Publications</b>	Jordan M Eizenga, Adam M Novak, Emily Kobayashi, <i>Flavia Villani</i> , Cecilia Cisar, Simon Heumos, Glenn Hickey, Vincenza Colonna, Benedict Paten, Erik Garrison (16 July 2020). Efficient dynamic variation graphs. Bioinformatics.		
<b>Certificates</b>	21 Apr 2018	<b>ESB Level 1 Certificate in ESOL International All Modes (B2 CEFR)</b> English language certification Candidate Number: IT-0042278	
	28 Mar 2015	<b>Primo soccorso B.L.S.D (Basic Life Support with defibrillation) e ostruzione delle vie aeree</b> Salerno, Italy	Certificate BLSD
<b>Languages</b>	English - (B2)		
<b>Computer Skills</b>	<i>Programming:</i> Python, R, Bash, LaTeX. <i>Computer Software:</i> Linux, Windows.		
<b>Other information</b>	<ul style="list-style-type: none"> <li>Quote in the newspaper "Le Monde": (<a href="https://www.lemonde.fr/blog/binaire/2020/05/06/sars-cov-2-et-covid-19-on-va-jouer-sur-les-mots/">https://www.lemonde.fr/blog/binaire/2020/05/06/sars-cov-2-et-covid-19-on-va-jouer-sur-les-mots/</a>)</li> </ul>		