Davide Baldazzi PERSONAL INFORMATION Via Fratelli Pinardi 8, 40128 Bologna (Italy) (+39) 3932929283 🔀 davide.baldazzi8@unibo.it Skype davide.baldazzi Sex Male | Date of birth 22/03/1993 | Nationality Italian Presently, I am a student of a four year PhD course in Data Science and Computation PERSONAL STATEMENT at the University of Bologna. My scholarship is due to a collaboration between a cancer research center (IRCCS-CRO; http://www.cro.sanita.fvg.it) located in Aviano (IT) and the University of Bologna. I'm currently supervised by Prof. Rita Casadio, group leader of the Biocomputing Group in Bologna, and by Dr. Roberta Maestro, leader of the Oncogenetic and Oncogenomic department at IRCCS-CRO. I'm involved in different projects including: Analysis of NGS data from cancer patients and specifically Whole Exome Sequencing Data (WES) and RNA Sequencing Data (RNA-Seq) Identification of novel human genomic structural variation at the transcriptomic level Structural Bioinformatics Database generation and curation Functional annotation of enzymes My goal is to become an expert in order to perform high-quality research in the field of Bioinformatics and develop methods useful to the whole community by focusing on health-related problems and stratified medicine. EDUCATION AND TRAINING PhD in Data Science and Computation 01/11/2017-Present EQF level 8 University of Bologna, Bologna (Italy) Scholarship: provided by IRCCS-CRO located in Aviano (IT). Tutor: Prof. Rita Casadio, leader of the Biocomputing Group (University of Bologna) Workplace: Biocomputing Group; FABIT department (Via San Giacomo 9/2 - 40126 Bologna, BO, Italy) Project: Development of a model for data standardisation and integration from different resources. A federated database about sarcomas (Research topics: Personalized Medicine)

01/10/2015–26/07/2017 Master Degree in Bioinformatics

University of Bologna, Bologna (Italy)

Thesis title: The relation between protein function and structure: functional annotation based on structural domains

Degree Mark: 110/110 cum Laude

Main Subjects:

- Programming
- Structural Bioinformatics

EQF level 7

- Computational Biology
- Proteomics
- Genomics
- Phylogenetics

01/10/2012–22/07/2015 Bachelor Degree in Biotechnology

University of Bologna, Bologna (Italy)

Thesis title: "Studio della relazione tra funzione, struttura e sequenza in alcuni tipi di deidrogenasi"

Degree mark: 104/110

Main subjects:

- Biochemistry
- Biology (molecular biology, microbiology, structural biology)
- Genetics (classical genetics, molecular genetics)
- Biodiversity and Evolution
- Biotechnologies (industrial, genetics, molecular, medical)

INTERNATIONAL RESEARCH __ EXPERIENCES

13/06/2018-11/09/2018 Scholarship

"Study and development of computational methodology for the interpretation of cancer genome"

La Jolla, San Diego, CA 92093, US

Position: Recipient of a fellowship as Visiting Graduate.

The project has been developed under the lead of Dr. Hannah Carter at UCSD (University of California, San Diego).

Italian referent: Dr. Emidio Capriotti

PERSONAL SKILLS

Mother tongue(s) Italian

Other language(s)

guage(s)	UNDERSTANDING		SPEAKING		WRITING	
	Listening	Reading	Spoken interaction	Spoken production		
English	B2	B2	B2	B2	B2	
	assessed by CLA (Centro Linguistico di Ateneo), University of Bologna in 2015					

Levels: A1 and A2: Basic user - B1 and B2: Independent user - C1 and C2: Proficient user Common European Framework of Reference for Languages

Job-related skills

Application of Statistical Methods

DB has got a vast knowledge of statistical methods. He is proficient with R programming language for statistical computation with a focus on genomic related analysis.

Data analysis and visualisation

EQF level 6

DB is experienced with data integration and annotation from different resources/databases with manual curation through literature mining. He is also capable of handling structural alignment and sequence alignment methods, performing building by homology and developing HMM from alignment profiles.

DB is also familiar with the analysis of Big Data being able to deal with them taking advantage of complex informatic architectures.

- Machine Learning
 - DB is familiar with the development of the following machine learning approaches
 - Support Vector Machines
 - Artificial Neural Networks
 - Hidden Markov Models
- Systems Biology

DB is expert on the computational and mathematical analysis for modelling complex biological networks

Database generation and curation

DB is experienced in the development of databases and their integration in a web service with and without the usage of a pre-built framework.

Digital skills

SELF-ASSESSMENT						
Information processing	Communicatio n	Content creation	Safety	Problem solving		
Proficient user	Proficient user	Proficient user	Independent user	Proficient user		

Digital skills - Self-assessment grid

DB is capable of working with a variety of technology and software solutions. He has got a solid background and advanced knowledge in development programming languages such as:

- Python
- Javascript
- Shell Scripting

DB is also familiar with markup languages like:

- HTML
- Markdown
- LaTeX

As already stated, DB is also expert on database creation, curation and management taking advantages of dedicated database management tools:

- PostgreSQL
- MySQL

ORGANIZING ACTIVITIES 21st Bologna Winter School – What can we learn from protein structure? 17/02/2020-21/02/2020 http://www.biocomp.unibo.it/~school2020/ Local Committee 20th Bologna Winter School – Data Science for Bioinformatics 18/02/2019-22/02/2019 http://www.biocomp.unibo.it/~school2019/ Local Committee 19th Bologna Winter School – Big Data and Bioinformatics 12/02/2018-16/02/2018 http://www.biocomp.unibo.it/~school2018/ Local Committee FEBS 2018 - The molecular basis of diseases: Can we infer phenotypes 23/05/2018-25/05/2018 from protein variant analysis? Bologna, 40124, BO, Italy Local Committee **TEACHING ACTIVITIES** A.Y. 2020-2021 Academic Tutor (60 hours) Bologna, 40125, BO, Italy "Laboratory of Bioinformatics 1"; Master Degree in Bioinformatics; University of Bologna DB supervised the technical part of the course. He helped students with the installation of the bioinformatics tools required for the final project and taught how to use them. DB collected and provided a first evaluation of the final projects for the course. A.Y. 2019-2020 Academic Tutor (60 hours) Bologna, 40125, BO, Italy "Laboratory of Bioinformatics 1"; Master Degree in Bioinformatics; University of Bologna DB supervised the technical part of the course. He helped students with the installation of the bioinformatics tools required for the final project and taught how to use them. DB collected and provided a first evaluation of the final projects for the course. An Introduction to R – How to perform Differential Expression Analysis September 2019 (15 hours) CRO, Aviano, 33081, PN, Italy DB held a practical course about R, a programming language and software environment for statistical computing. He taught the basic syntax of R and how to perform differential gene expression analysis using dedicated additional packages. A.Y. 2018-2019 Academic Tutor (60 hours) Bologna, 40125, BO, Italy "Laboratory of Bioinformatics 2"; Master Degree in Bioinformatics; University of Bologna DB supervised the technical part of the course. He helped students with the installation of the bioinformatics tools required for the final project and taught how to use them. DB collected and provided a first evaluation of the final projects for the course.

PUBLICATIONS

ORCID ID 0000-0001-5731-3663

05/2021 BENZ WS: the Bologna ENZyme Web Server for four-level EC number First Author Baldazzi, D; Castrense, S.; Martelli, P.L.; Casadio, R.;*Nucleic Acids Research*; 2021 DOI: 10.1093/nar/gkab328

Journal Impact Factor: 11.501 (2021)

10/2020 Following Author Following Author Following Author Tumor genotype, location and malignant potential shape the immunogenicity of primary, untreated Gastrointestinal Stromal Tumors Gasparotto, D.; Sbaraglia, M.; Rossi, S.; Baldazzi, D.; Brenca, M.; Mondello, A.; Nardi, F.; Racanelli, D.; Cacciatore, M.; Dei Tos, A.; Maestro, R.; *JCI Insight* 2020. DOI: 10.1172/jci.insight.142560 Journal Impact Factor: 6.205 (2020)

07/2020Highlighting Human Enzymes Active in Different Metabolic Pathways and
Diseases: The Case Study of EC 1.2.3.1 and EC 2.3.1.9
Babbi, G.; Baldazzi, D.; Savojardo, C.; Pier Luigi, M.; Casadio, R.; Biomedicines 2020, 8, 250.
DOI: 10.3390/biomedicines8080250
Journal Impact Factor: 3.600 (2020)

04/2020 Next-Generation Sequencing Approaches for the Identification of Co-First Author Pathognomonic Fusion Transcripts in Sarcomas: The Experience of the Italian ACC Sarcoma Working Group. Racanelli D, Brenca M, Baldazzi D, et al.; *Front Oncol.* 2020;10:489. DOI: 10.3389/fonc.2020.00489 Journal Impact Factor: 4.250 (2020)

ABSTRACTS

05/2020 Following Author	Epitheliod sarcoma: Molecular insights into proximal versus classic variant. Frezza A.M., Sigalotti L., Del Savio E., Baldazzi D., Sbaraglia M., Righi A., Gambarotti M., Barisella M., Brich S., Dei Tos A.P., Stacchiotti S., and Maestro R.; <i>Journal of Clinical</i> <i>Oncology</i> 2020, 38, no. 15_suppl
	DOI: 10.1200/jco.2020.38.15_suppl.e23552
	Journal Impact Factor: 18.230 (2020)
10/2019 Following Author	Myoepithelial tumours of soft tissues and extraskeletal myxoid chondrosarcomas feature a distinct transcriptional pattern Racanelli D., Stacchiotti S., Brenca M., Sbaraglia M., Fassetta K., Baldazzi D., Piccinin S., Brich S., Casali P.G., Collini P., Dagrada G.P., Fiore M., Gronchi A., Astolfi A., Pantaleo M.A., Righi A., Pilotti S., Dei Tos A.P., Maestro R.; <i>Annals of Oncology</i> 2019, Vol 30, Suppl. 5, V704

DOI: 10.1093/annonc/mdz283.054

Journal Impact Factor 10.570 (2020)

OTHERS	
PRIN; 2017 Researcher	Cutting Edge Analytical Chemistry Methodologies and Bio-Tools to Boost Precision Medicine in Hormone-Related DIseases Scientific Coordinator: Mondello, L. Scientific Director: Guardigli, M. PRIN (Progetto di Ricerca di Interesse Nazionale); 36 months.
ADDITIONAL WORK EXPERIENCE	
2010-2015	Lifeguard Summer Job
2011-2013	Professional Athlete Sport Club: Romagna Rugby Football Club