

nele Castrovilli

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Education

Universita' degli studi di Verona

Verona (VR)

MASTER DEGREE IN BIOINFORMATICS AND MEDICAL BIOTECHNOLOGIES — CURRICULUM IN BIOINFORMATICA

10/2014 - 03/2017

Universita' degli studi di Verona

Verona (VR)

BACHELOR DEGREE IN BIOINFORMATICS

10/2010 - 03/2013

Liceo scientifico A. Vallone

Galatina (LE)

SCIENTIFIC HIGH SCHOOL 2004-2010

Skills

Programming Python, Java, C, SQL, LaTeX, Matlab, Bash Shell

Web HTML5, CSS, Javascript

Operative systems Mac Os, Linux, Windows

Pentaho BI, MS SQL Server Management studio, SSRS, SSIS, Github, MySQL, PostreSQL, OracleDB, Atlassian JIRA, Atlassian Softwares

Confluence, Power BI

Languages Italian (Madrelingua), French (B1), English (B1)

Experience _____

Trueblue S.R.L. Verona, VR

Nov. 2016 - PRESENT BUSINESS INTELLIGENCE ANALYST

• Data warehouse maintainer and developer

- · Data warehouse developer for ETL process and report creation for a rich and usable data
- Using MsSql Server suite (SSRS, SSIS, MS SQL Management Studio 2008 2012

2016), Microsoft Power BI.

Standard compliant UAT document creation for projects

Melegatti S.P.A. Verona, VR

QUALITY CONTROL

Quality control over production chain and products sell outside Universitá degli studi di Verona

Verona Set. 2016 - Mar. 2017

Oct, 2013 - Dec, 2013

· Modeling and implementing a data warehouse to support genomic analyses

· Developed a data warehouse for genomes in order to create a process from uploading sequences, developing ETL stream and dashboard

creation with useful data

Verona

BIOINFORMATIC Lug. 2016 - ott. 2016

· Bioinformatic support for data mining

Developed an ETL process with python in order to extract meaningfull data from excel file format.

Universitá degli studi di Verona

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Verona

BIOINFORMATIC Set. 2015 - Mar. 2016

· Report generator

· Combining python and latex creating a useful tool capable of creating automatically a report with extracted data

Verona

BIOINFORMATIC Dic. 2015 - Mar. 2015

- · Similarity analysis between bacteriophages
- · Developed a tool capable of analyze and observe differences between genome sequences of bacteria and viruses

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Verona

BIOINFORMATIC Apr. 2013 - Luq. 2013

- · Genome wide association study for metabolic syndrome
- Using PLINK and bash shell programming we have developed a mechanism capable of automatic data analysis from data of metabolic syndrome patients

Universitá degli studi di Verona

Verona

BIOINFORMATIC Set. 2012 - Feb. 2013

- · Probabilistic techniques for studying Multiple Sclerosis with proteomic data
- · With MatLab we have developed a system capable of analyzing data and showing results with richfull graphs.

Certifications.

Python for Data Science and Machine Learning BootCamp

UDEMY Sept. 2017

Url: ude.my/UC-HI5Q6CLE Certification code: UC-HI5Q6CLE

JavaScript Essentials

UDEMY Sept. 2017

Url: ude.my/UC-3SODGSAGCertification code: UC-3SODGSAG

Advanced databases and SQL querying

Udemy Sept. 2017

Url: ude.my/UC-HOBIL3T9Certification code: UC-HOBIL3T9

Extracurricular Activity

AISM Verona (VR)

VOLONTARIATO Nov. 2012 - PRESENT

· Volunteer at AISM Verona, aiding people affected by Multiple Sclerosis. Collecting funds selling AISM's Apples and Gardenias

Open BioMedical initiative

Мемвек *Арт 2014 - Gen 2017*

· Bioinformatic support in OpenBioMedical initiative organization deveoping useful tools and supporting the creation of different DB's.

Ubuntu Bug tracker member

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Member Sep. 2010 - Actual

• In freetime i usually discover, report and resolve bugs for Ubuntu developer team

RunLover Everywhere

RUNNER Jun. 2010 - PRESENT

• I love run in free time using this practice as a tool to de-stress my body and mind.

Musician Everywhere

Drummer Forever

• Playing some drums as solo or with a band.