

# Daniele 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

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

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

# Experience \_\_\_\_\_

Trueblue S.R.L. Verona, VR

Business Intelligence analyst Nov. 2016 - PRESENT

- Data warehouse maintainer and developer
- Data warehouse developer for ETL process and report creation for a rich and usable data
- Using MsSql Server suite

#### Universitá degli studi di Verona

Verona

BIOINFORMATIC

Modeling and implementing a data warehouse to support genomic analyses

• Developed a data warehouse for genomes in order to create a process from upoloading sequences, developing ETL and fill dashboard with useful data

#### Universitá degli studi di Verona

Verona

BIOINFORMATIC

Lug. 2016 - ott. 2016

Set. 2016 - Mar. 2017

- Bioinformatic support for data mining
- Developed an ETL process with python in order to extract meaning ful data from excel file format.

#### Universitá degli studi di Verona

Verona

BIOINFORMATIC

BIOINFORMATIC

Set. 2015 - Mar. 2016

Dic. 2015 - Mar. 2015

Report generator

· Combining python and latex, we have created a useful tool capable of creating automatically a report with useful and meaningfull data

#### Universitá degli studi di Verona

Verona

· Similarity analysis between bacteriophages

· Developed a tool capable of analyze and observe differences between genome sequences of bacteria and viruses

JULY 29, 2017

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.

# **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**

· 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.