

# Daniele Castrovilli

BioInformatic



## About me

Graduate in bioinformatics (Bachelor and master degrees) at the University of Verona acquiring skills in the field of data analysis , and their manipulation . I implemented databases for genomes and data retrieving from them with PHP, and i learned programming languages such as Python , C , PHP , SQL , Matlab . My final project was the analysis , by means of proteomic data , the murine cells infected with acute myeloid leukoencephalopathy treated with pantethine IN vivo and EX vivo. After i worked as a bioinformatics at the hospital in Verona ( AOUI ) for the analysis of data from patients with cerebrovascular insufficiency . At last i've analyzed data obtained from devices created within the group at OpenBioMedical initiative as a volunteer. As master degree thesis i worked on developing a data warehouse supporting genomic analyses, grating the storage, manipulation and data interpretation of allelic variant files. Actually i'm a business intelligence analyst at Trueblue S.R.L. working on data warehouse for pharmaceutical industries interpreting and analyzing business core data.

## Contact

Via Bolzano, 5  
Verona 37135  
Italy

+39 342-0343653

E-mail  
Web Site   
GitHub: xcloudx8 

## Education

2004–2010	<b>High school</b> degree in Scientific studies	Liceo scientifico A. Vallone, Galatina(LE)
10/2010 – 03/2013	<b>Bachelor</b> degree in Bioinformatic	Universita' degli studi di Verona, Verona
10/2014 – 03/2017	<b>Master</b> degree in Bioinformatic and medical biotechnology, Bioinformatic curriculum	Universita' degli studi di Verona, Verona

## **Skills:**

### **Languages**

- Italian mother tongue
- English fluency
- French fluency

### **Programming skills**

- ❤️ Python
- SQL
- PHP
- JAVA
- C
- Matlab
- CSS3 & HTML5

### **Software skills**

- ❤️ Atom Editor
- Github
- Matlab
- Microsoft Excel
- Pentaho
- MySQL
- PostgreSQL
- Microsoft SQL Integration Service
- Microsoft SQL Reporting Service
- Microsoft SQL Server Management

# Experience

## Working experiences

Nov 2016 - Attuale

### **Business Intelligence analyst**

Trueblue S.R.L.

Working at Trueblue S.r.l. as Business Intelligence Analyst.

With the aid of big data we give useful business information to pharmaceutical business.

Detailed achievements:

- Data manipulation
- Data interpretation
- Pharmaceutical business core management
- MS SQL Server Integration Service, MS SQL Server Reporting Service, MS SQL Server Management

## Universita' degli studi di Verona

Sept 2012-Feb 2013	<b>Probabilistic technique for studying multiple sclerosis with proteomic data</b> <small>Universita' degli studi di Verona, Bachelor degree final project</small> <i>Bioinformatic</i> We have studied the effects of the vitamin B5 on murine cell growth both IN vivo and EX vivo and studied the effects with and without the vitamin Detailed achievements: <ul style="list-style-type: none"><li>• Learned how to manage huge file in excel</li><li>• Learned how to use MATLAB for data manipulation</li></ul>
Dec 2015-Mar 2015	<b>Analysis on bacteriophages and analysis on similarities between viruses attacking same bacteria.</b> <small>Universita' degli studi di Verona</small> <i>Bioinformatic</i> In this work we have analyzed the similarities between the genomes of the viruses and the specific host bacteria , and highlight any similarities between different virus genomes but attacking the same species of bacterium. the genomes of 300 bacteria and bacteriophages in 1000 were analyzed , with automation through bash shell scripts , python and analysis through IGTtools . Detailed achievements: <ul style="list-style-type: none"><li>• Learned how to manipulate data with bash shell scripting</li><li>• Learned how to use SED, GREP, AWK for data analysis</li><li>• Learned how to read and interpret huge amount of data</li></ul>
Sept 2015-Mar 2016	<b>Report generator</b> <small>Universita' degli studi di Verona</small> <i>Bioinformatic</i> Software for creating a simple pdf file from genomic analysis with NGS for a patient report, it is written in Python combining the typographic power of Latex. Detailed achievements: <ul style="list-style-type: none"><li>• Learned how to manage an entire project in Python</li><li>• Learned how to use GITHUB</li><li>• Learned teamwork</li><li>• Learned use of web parsers</li></ul>
Sept 2015-Mar 2016	<b>Variant database</b> <small>Universita' degli studi di Verona</small> <i>Bioinformatic</i> Creation of a simple database written in MySQL and PHP Detailed achievements: <ul style="list-style-type: none"><li>• Learned better SQL language</li><li>• Learned how to use MYSQL, POSTGRESQL</li><li>• Learned PHP</li><li>• Learned HTML and CSS3 usage of data retrieval</li></ul>
Sept 2016-Mar 2017	<b>Modeling and implementing a data warehouse to support genomic analyses</b> <small>Universita' degli studi di Verona</small> <i>Bioinformatico</i> Developing of a data warehouse to help genomic analyses and to create a big data support to genomic laboratory of University of Verona. Detailed achievements: <ul style="list-style-type: none"><li>• Better skills in SQL language</li><li>• Learned use of PENTAHO SOFTWARE SUITE, POSTGRESQL</li><li>• How to read genomic data files</li><li>• Understandings of statistical data</li></ul>

# AOUI, Universitary hospital of Verona

Jul 2016-Oct 2016	<b>Bioinformatic support for data mining</b> <i>Bioinformatic</i> Manipulation, extraction and data visualization about a wide range of data, from metabolic syndrom to child obesity Detailed achievements: <ul style="list-style-type: none"><li>• Learned how to extract useful data and make it understandable with visualization</li></ul>	AOUI hospital, Verona
Apr 2013-Jul 2013	<b>Genome-wide association for metabolic syndrome</b> <i>Bioinformatic</i> Analysis with bioinformatic tools on patients affected by metabolic syndrome Detailed achievements: <ul style="list-style-type: none"><li>• Learned how to read genetic population data</li><li>• Learned basic usege of PLINK</li></ul>	AOUI hospital, Verona
May 2013-Jul 2013	<b>GWAAS Database Help</b> <i>Bioinformatic</i> Analysis and creation of genetic databases with Hapmap for GWAAS analysis Detailed achievements: <ul style="list-style-type: none"><li>• Learned how to use HapMap</li><li>• Learned better BASH SHELL SCRIPTING</li></ul>	AOUI hospital, Verona

## Volunteering

Valunteer	<b>OpenBioMedical Initiative</b> <i>Bioinformatic</i> Support on developing source code and being a Bioinformatic consultant Detailed achievements: <ul style="list-style-type: none"><li>• Learned better team management and teamworking</li></ul>	Nowhere
Volunteer	<b>Ubuntu translator / Bug Tracker</b> <i>Volunteer</i> Support on translating strings in Ubuntu OS and bug discovery. Detailed achievements: <ul style="list-style-type: none"><li>• Learned how to translate softwares</li><li>• Learned better HOW A COMMUNITY WORKS</li></ul>	Nowhere
Volunteer	<b>AISM</b> <i>Volunteer</i> Volunteer on the AISM association as care giver	Nowhere

*Disclaimers:*

*This CV has been made entirelly with LaTeX*