



**Vito Flavio Licciulli** obtained a degree in Computer Science at University of Bari in 1992 (110/110). Since 1994 he is involved in Bioinformatics with the Bari Bioinformatics Group and he is currently a researcher at the C.N.R.- Istitute for Biomedical Technologies (ITB), Bari (Italy).

His main research activities are:

- design, implementation and maintenance of specialized biological databases developed in collaboration with internal or other research groups;
- management and integration of large amounts of biological data (BigData) through the development and implementation of data-warehouses and other data integration tools;
- development of bioinformatics pipelines for the analysis of Next Generation Sequencing (NGS) data, in particularly for trascryptomics studies, both coding and non-coding;
- development of WEB platforms and interfaces for the pipelines and software developed in the institute.

He is author/coauthor of more than 30 papers in international journals in the field of Bioinformatics, he has been involved in more 20 national and international project both as participant and as a Principal Investigator, and he has played in several lectures in Bioinformatics courses/masters.

He is also responsible of database and computer server management at ITB-CNR-Bari institute.

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