



Istituto di Informatica  
e Telematica



Consiglio Nazionale  
delle Ricerche

# Bioinformatic Research at IIT: the Highlights

Marco Pellegrini

Istituto di Informatica e Telematica  
CNR

# People

Filippo Geraci

Mauro Leoncini

Manuela Montangero

**Marco Pellegrini**

Maria Elena Renda

Alessio Vecchio

Giovanni Manzini

Romina D'Aurizio

Davide Verzotto

Francesco Russo



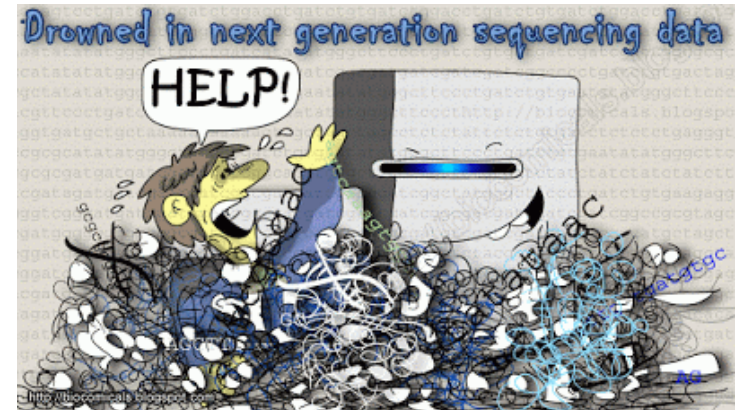
# Bioinformatics at IIT-CNR

- **Biological sequences analysis/classification**
  - Gene and microRNA expression data analysis
  - Motif and Tandem Repeat identification/extraction
  - Next generation sequencing: CNV detection
- **Diseases classification and gene expression profiling**
  - Biological networks (PPI)
  - Prediction of Genes/microRNA markers in Tumors
- **Web tools development**
  - for the analysis and visualization of biological sequences and Bio-networks (AMIC@, Core&Peel, Excavator2, Dot2Dot,..)



***In collaboration with:***

*IFC - CNR, Pisa  
Dept. of Computer Engineering,  
Univ. of Modena & Reggio Emilia*



<http://bioalgo.iit.cnr.it>



# Bioinformatics: focused projects



## Recent Projects

- **Analisi del ruolo di alcuni complessi miRNA/mRNA nel tumore della prostata .**  
Funded by *ITT (Istituto Toscano Tumori)*  
Partners: IIT e IFC
- **Eligibility of miRNAs modified by docetaxel in prostate cancer cells to plasma biomarkers in patients responsive and no more responsive to docetaxel.**  
Funded by *ITT (Istituto Toscano Tumori)*  
Partners: IIT e IFC
- **REPEATALS: Analisi di sequenze tandem repeats polimorfici nella Sclerosi Laterale Amiotrofica (SLA) .**  
Funded by *Arisla (Associazione italiana per la Ricerca sulla Sclerosi Laterale Amiotrofica)*  
Partners: IIT, Univ. Novara, ITB (Istituto di Tecnologie Biomediche, MI)
- **PRIN2015: "The role of tandem repeats in neurodegenerative diseases: a genomic and proteomic approach"**  
Funded by *MIUR* . Partners: IIT, Univ. Novara, U. Insubria.



# Bioinformatics



## Strategic Factors:

- Strong links with Biologists (IFC CNR , U. of Eastern Piedmont, IEO Istituto Oncologico Europeo)
- Tuscany Bioinformatics Days – Bioinformatiha
- Editorial Activity: **Special Issues** of Frontiers in Bioengineering and Biotechnology (Nature Group) .
- M.P. Associate Editor of: *Frontiers in Genetics* and *BMC Cancer*.



# Thesis Topics

- Algorithms for Long-reads Sequencing: Using Single-Cell Sequencing to Improve Structural Variants Detection in Clinical Practice
- Precision Oncology:
  - Multi-Omics data Analysis for Biomarkers Discovery in Cancer Research
  - Integrative Analysis of multi-omics data to study Tumor Heterogeneity
- Computational Genomics: Study of the impact of Genomic Structural Variations on dysregulation of coding and non-coding genes.
- Genomics and Informatics: Development of tools for Copy Number Variants annotation

Romina D'Aurizio, PhD  
romina.daurizio@iit.cnr.it



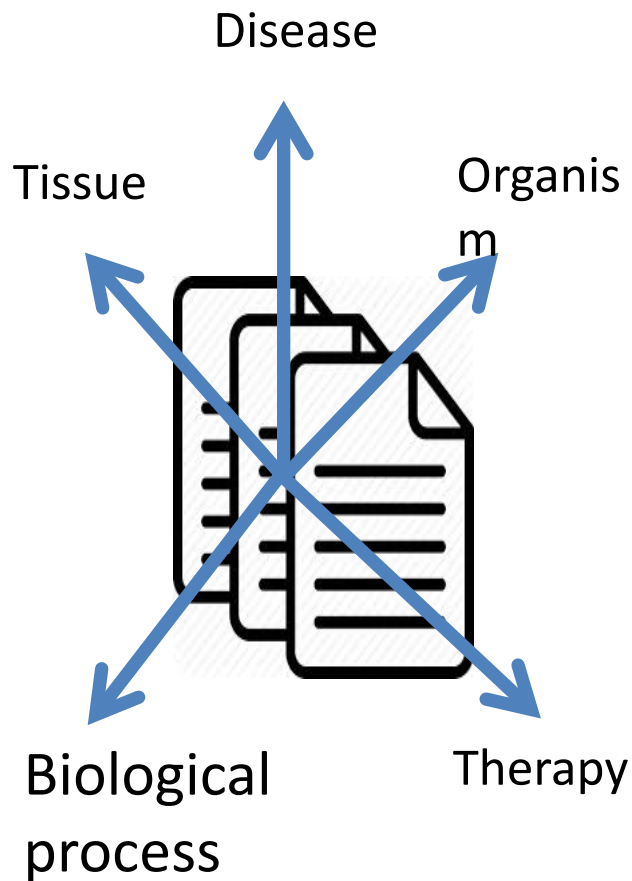
# Thesis Topics

- - Tandem repeat and SV analysis in human neurological disorders
- Integrated data mining methods for biological sequence analysis
- Metagenomic sequence classification

Contact : [davide.verzotto@iit.cnr.it](mailto:davide.verzotto@iit.cnr.it)



# Information retrieval on biomedical documents



- Goals:
  - Use machine learning and AI to extract categorical information from biomedical texts
  - Use categorical information to build a social network of concepts
  - Extract new knowledge from the network
- Contact:
  - [filippo.geraci@iit.cnr.it](mailto:filippo.geraci@iit.cnr.it)





# Thesis Topics

- Network based drug repositioning.
- Detection of borderline CNV (small copy number variations) in cancer genomes.

Contact :  
[marco.pellegrini@iit.cnr.it](mailto:marco.pellegrini@iit.cnr.it)



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