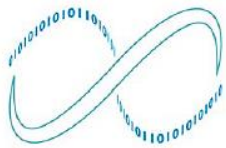




Centro di Competenza sul Calcolo Scientifico

# *Systems Biology & Systems Medicine*



*SysBioM*

*Centro Interdipartimentale  
Molecular Systems Biology*

Michele De Bortoli



*Scienze Cliniche e  
Biologiche*

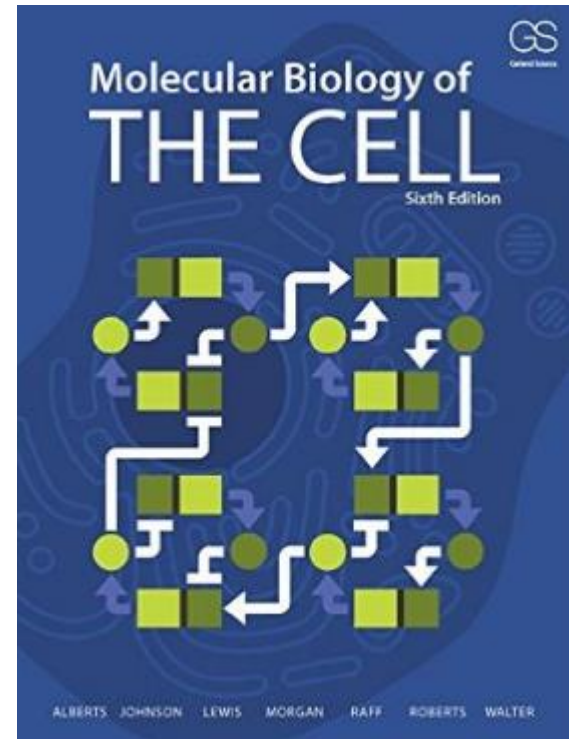
*Polo San Luigi*

*Orbassano*

*1. Sistemi complessi: programma di dottorato e Centro*

*2. Dipartimento di Scienze Cliniche e Biologiche*

Le funzioni delle cellule e degli organismi sono costruite, regolate e si perpetuano grazie all'azione di centinaia di migliaia di entità molecolari diverse (geni, RNA, proteine, metaboliti,...) che interagiscono tra loro in modo coordinato, specifico e dinamico.



Studiarne il comportamento come **sistema**, e non come singole molecole, è fondamentale.

Oggi questo è possibile, grazie all'avanzamento di:

- Tecniche analitiche parallele  
(NGS, arrays, proteomica, metabolomica,...)
- Metodi di analisi dei dati e di modellizzazione  
(sistemi complessi, neural networks, ...)
- Metodi informatici e computazionali  
(database, algoritmi di integrazione, ...)

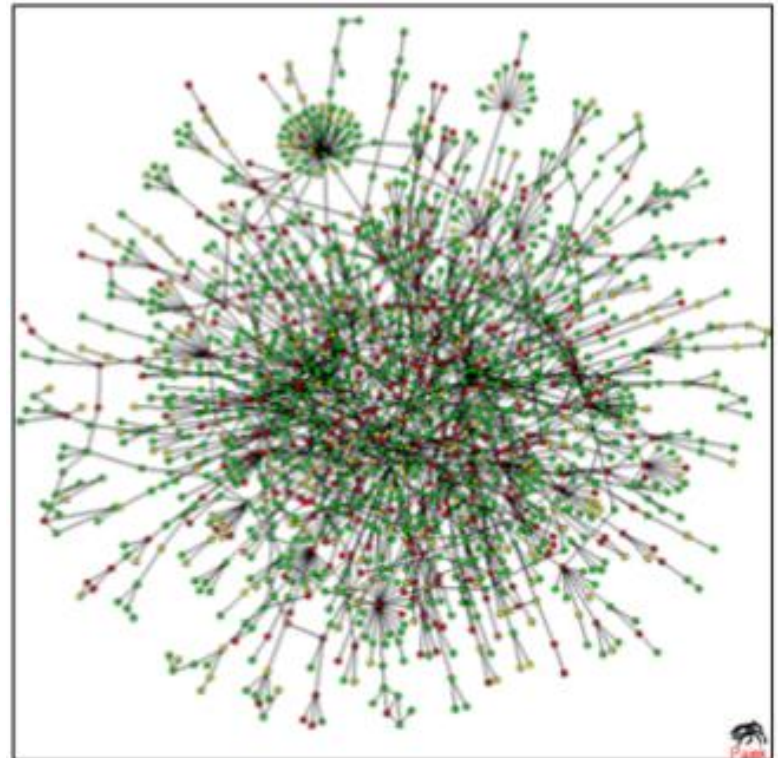
*Systems Biology* e *Systems Medicine* richiedono una profonda interazione ed integrazione tra le capacità e le conoscenze di varie discipline

Fisica, Biologia e Medicina, Matematica, Informatica, ...

L'elaborazione si basa su:

- Teoria delle reti
- Modellizzazione quantitativa
- Ontologie

*Centro Interdipartimentale  
Molecular Systems Biology*



# Dottorato in Sistemi complessi nelle Scienze della Vita

## PhD Programme: Complex Systems in Life Sciences

input

### LM

Biologia  
Biotecnologia  
Medicina  
Fisica  
Matematica  
Informatica  
Ingegneria

Laureati stranieri

### Collegio:

Medicina, Biologia,  
Fisica, Informatica,  
Matematica,  
Ingegneria, esterni

Didattica formale e  
seminariale, con  
programmi di  
azzerramento

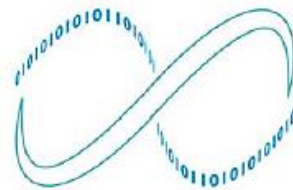
### Thesis:

- Regulatory Genomics
- Cellular regulatory pathways
- Bioinformatics
- Systems Biology
- Molecular networks
- Systems Medicine
- Modelling of biological phenomena
- Epidemiology
- Vaccinology

output

La disponibilità di una risorsa di super-calcolo scientifico in Open-Lab è fondamentale per il successo delle ricerche che impegnano i nostri studenti

... e non solo!



*SysBioM*

*Centro Interdipartimentale  
Molecular Systems Biology*

Polo Universitario del San Luigi di Orbassano

Ospedale – Università - NICO



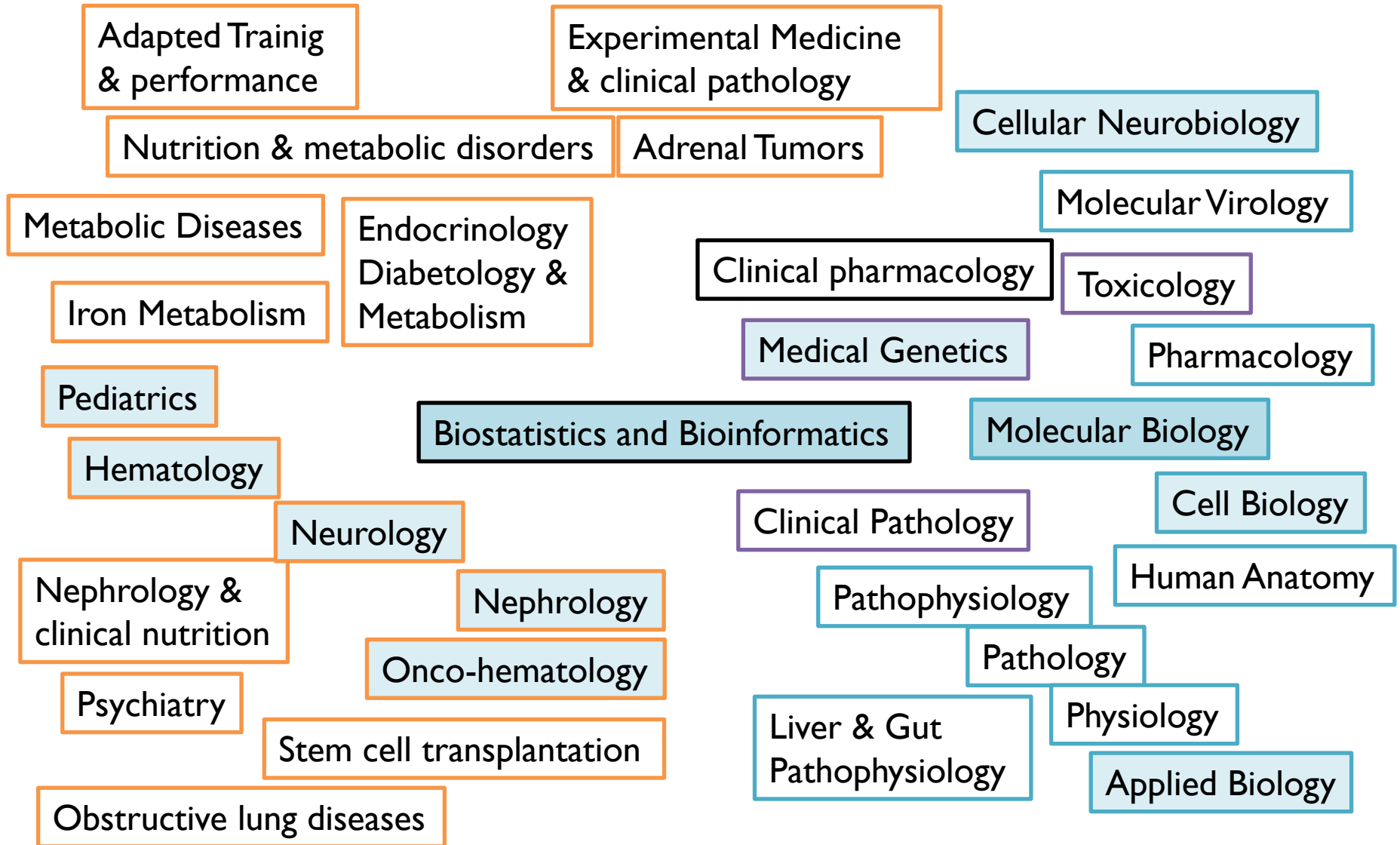
Dipartimento di Oncologia  
Dipartimento di Neuroscienze

**Dipartimento di Scienze Cliniche e Biologiche**

*Corso di Laurea in Medicina e Chirurgia  
Corsi di laurea delle professioni sanitarie  
Master in Cellular & Molecular Biology*








Orange: clinical . Blue:basic. Blue shadow grades indicate involvement in computing.



## Key Research Areas

### Rare diseases

The last day of February every year is **Rare Disease Day**. In January 2016, 10 new projects kicked off focusing on the development of new treatments for rare diseases. The projects are funded under Horizon 2020, the EU's funding programme for research and innovation (2014 – 2020).

Read more about these projects  321 KB

Many rare diseases cause chronic health problems or are even life-threatening. Genetic factors play a role in a majority of these diseases. The impact on the quality of life of affected patients, of whom many are children, is significant.

In the European Union, a disease is considered rare when it affects not more than 1 person in 2.000. This low prevalence is the common feature shared by all rare diseases, which altogether affect all biological systems. This nevertheless means that between 6 000 and 8 000 different rare diseases affect or will affect an estimated 30 million people in the European Union.

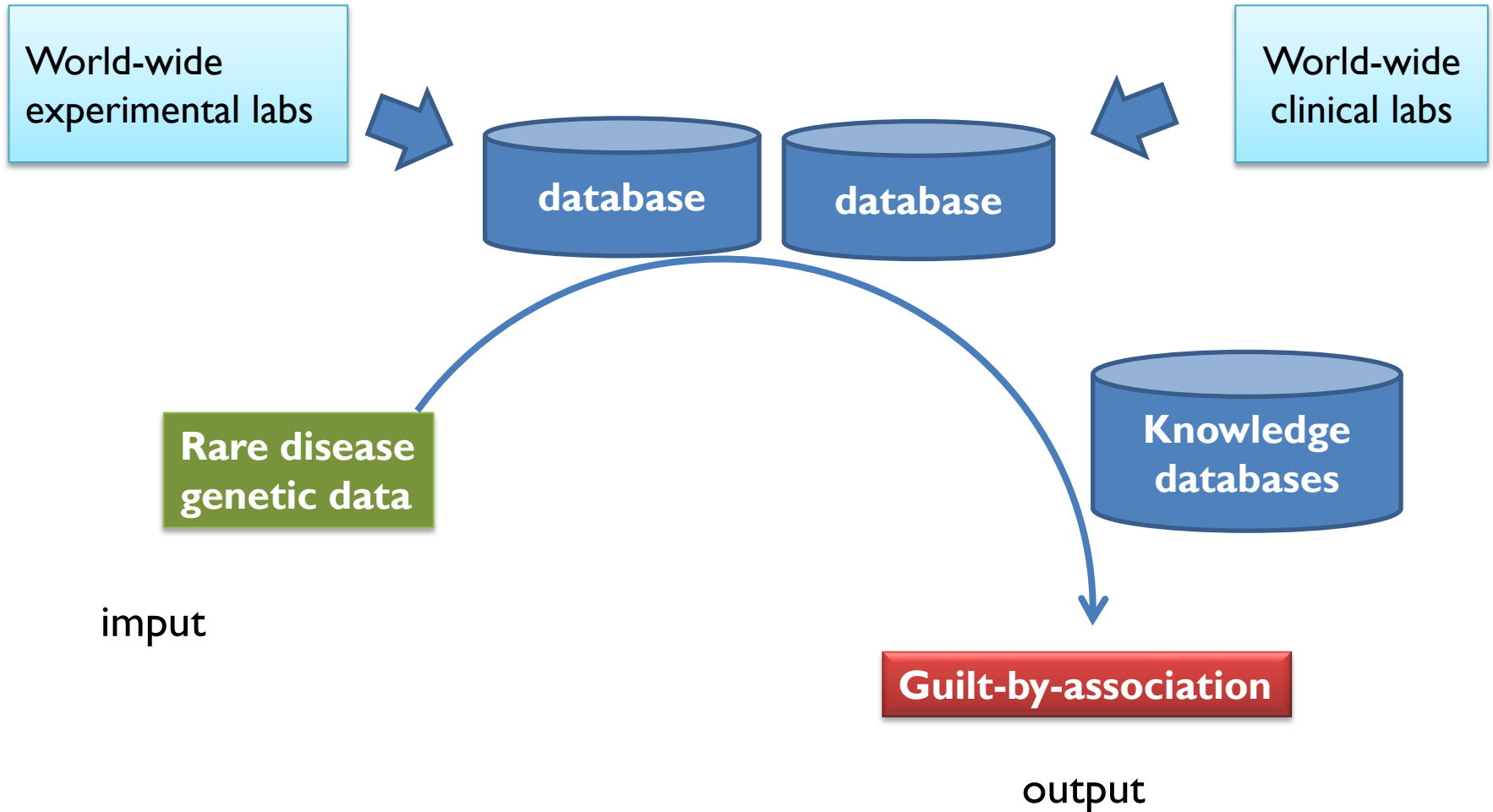
### Links

- [Horizon 2020](#)
- [International Rare Diseases Research Consortium \(IRDiRC\)](#)
- [E-RARE ERANET Co-fund](#)
- [EU policies for rare diseases](#)

### Video

- [In search of IRDiRC](#)

# Rare disease: genetics and molecular characterization



# Single clinical case requires computational power

- girl with uncorrect 9p-syndrome diagnosis by karyotype
- Reassessment: balanced translocation
- Further analysis with NGS techniques  
complex chromosomal rearrangement (>30 breakpoints)

Breakpoints, epigenetics mapping point to pathways of neurodevelopment

- Mate-pair sequencing of the patient DNA
- WGS on trio DNA (mother-father-patient)
- RNA-sequencing on trio blood

<b>Assay</b>	<b>Sample</b>	<b>Fastq file size (GB)</b>	<b>N°Reads (Million)</b>
Mate-Pair Sequencing	Child	18	57
Whole Genome Seq	Child	315	908
Whole Genome Seq	Father	(106 the BAM file)	1369
Whole Genome Seq	Mother	272	784
RNA-Seq	Child	11	55
RNA-Seq	Father	7	39
RNA-Seq	Mother	9	46

# NGS-based rare-disease-causing-gene discovery

Whole Exome Sequencing: ~1% protein coding portion of the human genome (**the exome**) is sequenced by NGS.

## Autism sequencing consortium



Large international effort (Autism Sequencing Consortium, New York, USA) - exome sequencing of 50,000 trios (one proband and parents). Our group contributes 500 trios.

Need: analysis of WES data aligned to the reference human exome (.bam files).

## Project

“*Machine Learning Techniques in Food Risk Assessment*”  
funded by the European Food Safety Agency.

Dealing with  
Extensive Literature  
Searches

mapping the evolution of hypothetically  
any general topic over time

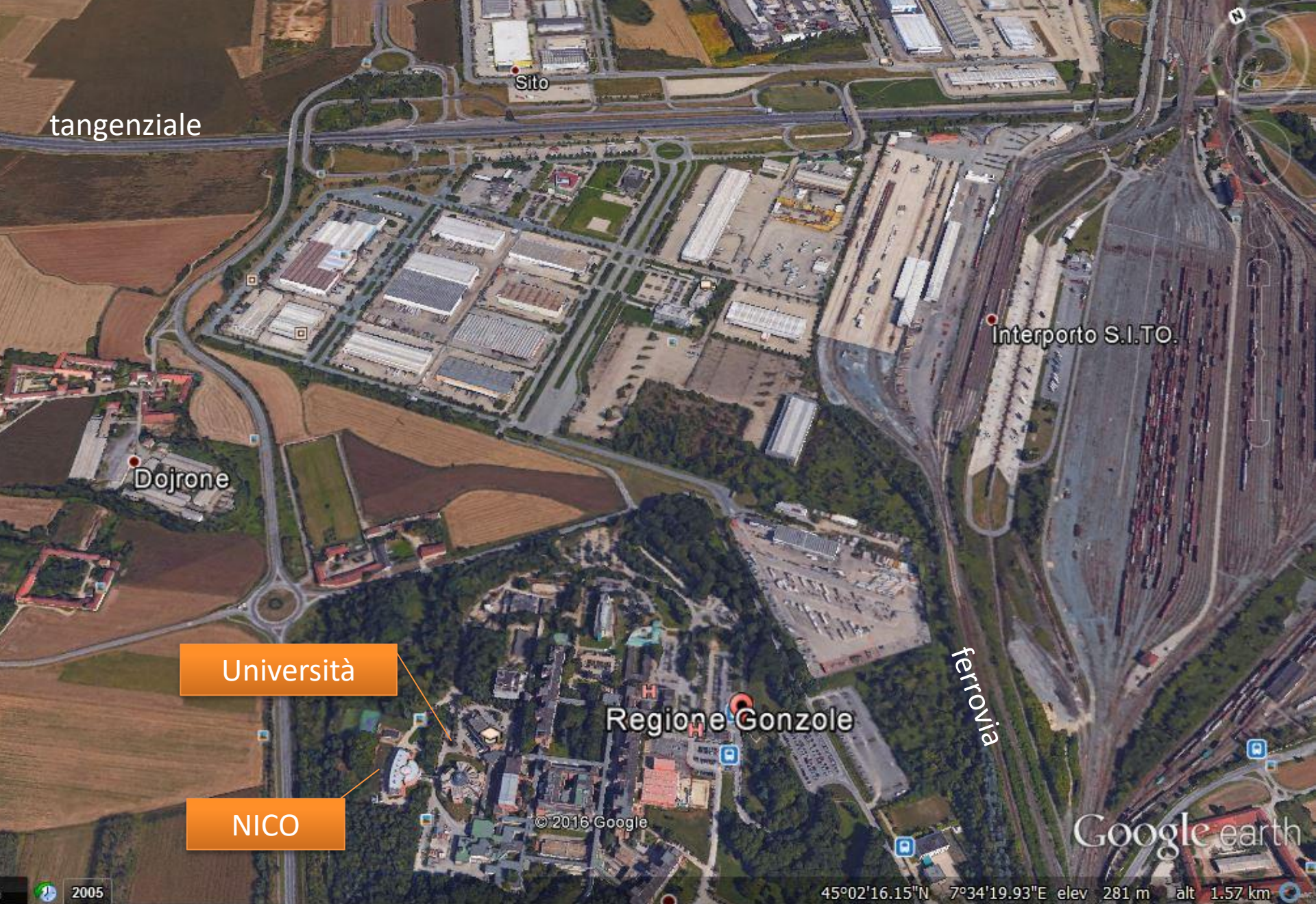
Machine Learning algorithms  
Super Learner solutions

active learning algorithms  
for unsupervised pattern

Current test: a bulk of 2,655,365  
abstracts related to Machine Learning  
Techniques retrieved from bibliographic  
databases.

Dr. Paola Berchialla - Unit of Biostatistics

Department of Clinical and Biological Sciences, University of Torino



tangenziale

Sito

Dojrone

Interporto S.I.TO.

Università

NICO

Regione Gonzole

ferrovia

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Google earth

2005

45°02'16.15"N 7°34'19.93"E elev 281 m alt 1.57 km

Grazie !