

# Curriculum Vitae

## Behnam Yousefi

Birth date: 15 Jan 1991

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### Education

#### PhD Student in Computational Biology

Computational Biology Department, Institut Pasteur, Paris, France;

École Doctorale Complexité du Vivant, Sorbonne Université, Paris, France

Feb. 2020-2023.

Thesis: Integrative modeling of cancer drug response and auto-immune disease base on network-based machine learning techniques (under the supervision of Dr. Benno Schwikowski).

Secondments:

- BIO3 – Laboratory for Systems Medicine, KU Leuven, Leuven, Belgium (4 month)  
Subject: Medical applications of individual specific networks (under the supervision of Prof. Kristel van Steen)
- InSilicoTrials company, Trieste, Italy (3 months)  
Subject: Market analysis of compounds for neurodegenerative diseases (under the supervision of Dr. Roberta Bursi and Dr. Fienne Sips)

#### Master in Biomedical Engineering, Bioelectric

Iran University of Science and Technology (IUST), Tehran, Iran, Sept. 2015- Feb. 2019.

**GPA: 18.18/20** (*Top Student*, the average GPA in Biomedical Engineering Department is 16.10/20)

Thesis: Diagnosis and treatment of obstructive sleep apnea based on machine learning techniques (in Iran Neural Technology Research Center under the supervision of Prof. Abbas Erfanian).

#### Bachelor in Electrical Engineering, Electronics

Azad University (Abhar branch), Abhar, Zanjan, Iran, 2009-2014.

**GPA: 19.14/20** (*Top Student*)

Thesis: Design and implementation of a hardware based on AVR microcontroller for data transfer in control systems.

#### High School Diploma in Physics and Mathematics

Exemplary School of Imam Sadegh, Abhar, Zanjan, Iran. 2005-2008.

**GPA: 17.91/20**

### Honors and Awards

Awarded the **Marie Skłodowska-Curie Actions fellowship** for PhD studies, 2020.

**Top Student** Award of biomedical engineering department, Iran University of Science and Technology, 2016.

**Ranked in the top 1%** of the National University Entrance Examination, 2015.

**Top Student** Award of electronics engineering department, Azad University, 2011.

**Top Student** Award of electronics engineering department, Azad University, 2010.

Accepted in the first stage of National Astronomy Olympiad, 2007.

### Experimental and Professional Experience

**Machine learning:** supervised and unsupervised machine learning, deep learning, graph neural networks, recommender systems, consensus clustering.

**Network analysis:** graph representation learning, multilayer and heterogeneous graph analysis, graph-based clustering.

**Data analysis:** data preprocessing and normalization, statistical tests, low dimensional representation.

**Systems and signals modelling:** noise reduction, adaptive filters, spectral analysis, wavelet analysis.

**Chaos and nonlinear dynamics:** phase space reconstruction, Poincare section, complexity and entropy measures.

**Bioinformatics:** network biology, gene expression data (RNAseq and microarray) analysis, gene module detection, gene enrichment analysis.

**Neural data analysis:** spike sorting, spike train and LFP signal analysis, neural decoding,

**Visualization:** efficient visualization of computational pipeline and results.

### Other Skills

**Scientific software and languages:**

R, Python, MATLAB, LabVIEW, C/C++, HTML, CSS, Bash, SLURM, Reproducibility techniques (git, Docker, Conda)

**Languages:**

English (full professional), Persian (native), French (beginner).

### Publication

Farzaneh Firoozbakht, **Behnam Yousefi**, Benno Schwikowski, “An Overview of Machine Learning Methods for Monotherapy Drug Response Prediction”, briefings in bioinformatics, 2021 (DOI: [10.1093/bib/bbab408](https://doi.org/10.1093/bib/bbab408))

**Behnam Yousefi**, Benno Schwikowski, Kristel Van Steen, “Individual-specific network inference to capture dynamic microbiome interactions” (submitted)

**Behnam Yousefi**, Farzaneh Firoozbakht, Benno Schwikowski, “Robust Stratification Through Consensus Clustering in Bioinformatics” (under preparations)

### Posters and Abstracts

1- ASHG (American Society of Human Genetics) conference-2021: “Individual Specific Networks Hot Zone Detection”

2- IGES (International Genetic Epidemiology Society) conference-2021: “Novel analysis pipeline for microbiome data via individual-specific networks”

3- IGES (International Genetic Epidemiology Society) conference-2022: “Individual-specific Networks and Representation Learning to Capture Dynamics of Microbiome interactions”