INPHINIT DOCTORAL CONTRACT



CALL FOR CANDIDATES INCOMING

Bioinformatics for accurate genetic diagnosis and personalized medicine

CENTRE: IIS ARAGÓN
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POSITION DESCRIPTION

Personalized Medicine is an urgent need for people as each of us carries a slightly different genetic information and accumulates fortuitous mutations.

Our research group is expert in proteins. We study proteins in the laboratory and in the computer. Our goals are:

- 1-To understand the phenotypic effect of mutations at the protein level in order to develop accurate bioinformatics tools to assist medical doctors in delivering accurate genetic diagnostic.
- 2-To develop new drugs to provide cure for people in need of personalized treatments (against resistant bacteria, rare inherited metabolic diseases or a variety of amyloidosis).

In this project, we will focus in developing accurate bioinformatics tools for genetic diagnosis. A combined project including computational drug discovery work on selected diseases can be offered for students so inclined.

The successful candidate will get involved in:

- 1-developing & applying Bioinformatics tools (e.g. MD simulations) for the accurate interpretation of human genetic variations.
- 2-evaluating novel SNVs found in neonatal screening programs run in Spain and France.
- 3-interpreting the deleterious "variome" of a few well-selected genetic disorders.
- 4-(personal interest existing) improving new drugs discovered and developed in our group.

Full time contract: 3 years

FELLOWSHIP PROGRAM Doctorate INPhINIT Incoming

Pre-doctoral contract for earlystage researchers of any nationality to pursue PhD studies on life sciences in Health Research Institute

IIS Aragón (Zaragoza, Spain) www.inphinitlacaixa.org

6 February 2019

Deadline for submitting applications.

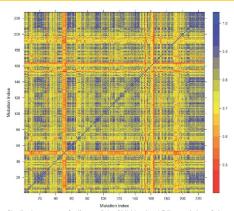
20 February 2019 Deadline for submitting language certificate.

23 April 2019 Notification of the shortlist results.

28 and 29 May 2019 Face-to-face interviews in Barcelona.

4 June 2019 Publication of final list of selected candidates.

From 4 June 2019 to 5 July 2019 Matching research centre – fellow.



Similarity map of all possible SNV in the LR5 module of the LDL receptor obtained from Molecular Dynamics simulations, from Human Molecular Genetics, 2016, 25: 1233-1246