

Courtesy translation of D.R. n. 046/2023

For more details on the selection process, please refer to the Italian version of D.R. n. 046/20232 available at http://www.hunimed.eu/it/lavora-con-noi/

SELECTION PROCEDURE FOR RESEARCH FELLOWSHIP

Research Program Title	Single-cell dissection of the cellualr ecosystem in hepatocellular carcinoma
Tutor	Prof. Luigi Maria TERRACCIANO
Scientific Area	06 – Medical Sciences
Gross amount of the fellowship	30.000 Euro
Duration of the fellowship	12 months
Objectives of the research	We hypothesize that by combining the resolution of single cell RNA sequencing (scRNA-seq)in situ visualization and the power afforded by large cohorts of bulk RNA- sequencing of HCCs we will be able to better characterize HCC from both tumor-centric and tisseu microenvironment (TME)-centric perspectives, and to identify somatic genetic alterations that may modulate the TME in HCC. The aims of this project are 1) to dissect the cellular composition of HCC, 2) to define transcriptional cell states of the tumor cells that are associated with distinct TME patterns in HCC and 3) to identify genetic features in HCC that may modulate TME heterogeneity and composition. For aim 1 we will use scRNA-seq and in situ visualization to determine the identity, abundance, cell type-specific gene signatures and spatial distribution of the various major and rare populations in HCC. To extend the findings by scRNA-seq to large HCC cohorts in aims 2 and 3, we will optimize and apply in silico virtual microdissection to define the molecular and cellular subclasses of HCC from both tumor-centric and TME-centric perspectives and to identify the genetic features that may modulate the TME.



Activities to be carried out	 Mutational analysis and sequencing of DNA and RNA in fresh and paraffin samples; primers design and Sanger direct sequencing; estabilishment of primary cell coltures; preparation of HCC organoids from liver biopsy.
Work place	PIEVE EMANUELE - Milan
Mandatory requirements	 PhD in Genetics, Biostatistical sciences, Biomedical Engineering, Biological Sciences, Bioinformatics sciences or related discipline Professional CV suitable to the carrying out of the research activities outlined above
Selection process	Application for admissions must be submitted at the following link: https://pica.cineca.it/humanitas No hard copy of the application must be sent by post. At first access, applicants need to register by clicking on "Register" and completing the requested data. If applicants already have LOGINMIUR credentials, they do not need to register again. They must access with their LOGINMIUR username and password in the relevant field LOGINMIUR. Applicants must enter all data necessary to produce the application and attach the required documents in PDF format.
Selection criteria	Selection criteria are predetermined by the Selection Committee. As part of the selection process, the Committee will evaluate the curriculum, titles and publications presented by the candidate and will consider, in particular: - Minimal/theoretical experience in NGS sequencing of DNA and RNA; - Experience in molecular biology (i.e. cell colture); - At least elementary competence in bioinformatics methods; - Experience in biology of hepatobiliary cancer;



- Ability to work in group and follow young PhD
students;
- Advanced knowledge of written and spoken
English.

FURTHER INFORMATION:

In the event of any conflict between Job Opening text and Italian D.R. text, the Italian version will prevail.

For more details on the selection process please refer to the **D.R. n. 046/2023** (http://www.hunimed.eu/it/lavora-con-noi/) or send an inquiry to ufficiodocenti@hunimed.eu or telephone +39 02.8224.5642/5421.