

## **Ana Rita Matos**

Project title: The primary genetic cause and beyond: the life-time risk and phenotype modifiers in HDGC

| Duration               | 4 weeks   |
|------------------------|---|
| Short Bio              | I am a PhD student at the Institute for Research and Innovation in Health (i3S) in Porto, Portugal. My current research focusses on Hereditary Diffuse Gastric Cancer (HDGC), a deadly cancer syndrome associated with E-cadherin (CDH1) germline alterations. The research aims to understand why some CDH1 mutation carriers develop cancer early while others remain disease-free.   |
| Home<br>Institution    | 13S – Institute for Research and Innovation in Health, Porto,<br>Portugal   |
| Host institution       | Centre Léon Bérard, Lyon, France  |
| Project<br>description | This project focuses on HDGC syndrome and aims to understand the molecular mechanisms behind the varying risks within families carrying CDH1 mutations. The incomplete penetrance in HDGC families necessitates mutation-based risk assessments. The research is dedicated to predicting cancer development in CDH1 mutation carriers, enhancing decision-making for at-risk HDGC families. We seek to generate lifetime-risk estimations based on intra-familial or mutations-specific data. The project benefits from a 120-individual HDGC family's long-term study, with half carrying a CDH1-causing founder mutation. During the fellowship, I aimed to utilize a software package for cancer risk estimation developed by Youenn Drouet (Postdoc at Host Institute). |
| Personal<br>statement  | Throughout the course of this fellowship, I discovered a newfound interest in bioinformatics and statistical analysis. Although I initially had a basic understanding of the tools and knowledge required to achieve our proposed tasks, the robust training I received equipped me with the necessary skills to  |
|                        | delve into these complex fields.  |

In collaboration with:







This experience not only broadened my expertise but also sparked a passion for the intersection of genetics, bioinformatics, and statistical analysis, laying the foundation for future pursuits in this dynamic and interdisciplinary field.

I thoroughly enjoyed my experience during this fellowship. From day one, I felt consistently welcomed and embraced by Dr. Youenn Droet and his team. I was fortunate to have a host supervisor who was not only highly knowledgeable but also exceptionally supportive. Their availability and willingness to accommodate all my questions created a conducive environment for learning and collaboration, making my time in the fellowship immensely enjoyable and enriching.





