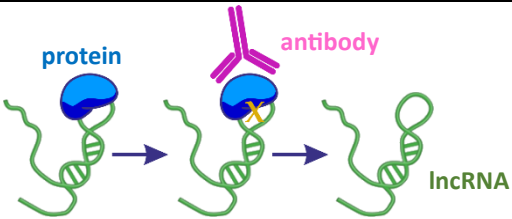


## 2027 Internship Offer

**Master 1:** YES – Duration: 4 months

**Master 2:** YES – Duration: 5 months

Team, Contact	Robert FEIL, Institute of Molecular Genetics of Montpellier (IGMM), <b>Isabel CHILLON</b> ( <a href="mailto:isabel.chillon@igmm.cnrs.fr">isabel.chillon@igmm.cnrs.fr</a> )
Title	Regulatory roles of long non-coding RNAs (lncRNAs) during cellular stress
Research Themes and questions	<p>This project aims to investigate how specific long non-coding RNAs (lncRNAs) regulate gene expression under conditions of cellular stress, thereby restoring homeostasis.</p> <p>In particular, the student will contribute to characterizing the direct protein partners of the lncRNA Meg3 that are engaged during the DNA damage response in neurons, which the lab has recently identified.</p> <p>The results of this internship will help us understand the role of lncRNAs in dysregulated pathways involved in neurodegenerative diseases.</p>
Methods and experimental approaches	<p><b>Model system:</b> primary neuronal cells differentiated from mouse embryonic stem cells.</p> <p><b>Experimental approaches.</b> The student will apply biochemical and functional assays to validate lncRNA-protein interactions and assess their functional significance:</p> <ul style="list-style-type: none"> <li>- eCLIP-RT-qPCR: to validate lncRNA-protein interactions identified by a screening assay.</li> <li>- Detection of the marker <math>\gamma</math>H2AX by immunostaining and flow cytometry: to identify interactions involved in DNA damage pathways.</li> <li>- Apoptosis and cell cycle analysis measured by flow cytometry: to interrogate interactions involved in proliferation and programmed cell death.</li> </ul>
Illustration	<p>eCLIP-RT-qPCR</p> 

2-3 Publications	<ol style="list-style-type: none"><li data-bbox="502 280 1390 539">1. López-Perrote A, Martín-Cuevas EM, Mérida-Cerro JA, Aicart-Ramos C, González-Corpas A, González J, Le Coq J, Boskovic J, <b>Chillón I</b>#, Llorca O#, Moreno-Herrero# F and Huarte M# (2026) CONCR lncRNA organizes a 3'-end structural domain that engages DDX11 for DNA replication and sister chromatid cohesion. <i>Mol Cell</i>. Accepted. #Co-corresponding authors.</li><li data-bbox="502 589 1390 772">2. Assenza S, Blanco C, Favard C, Carnesecchi J, Marcia M and <b>Chillón I</b> (2025) Conserved architecture of a functional lncRNA-protein interaction in the DNA damage response pathway. <i>bioRxiv</i> 2025.2007.2024.666508. <a href="https://www.biorxiv.org/content/10.1101/2025.07.24.666508v1">https://www.biorxiv.org/content/10.1101/2025.07.24.666508v1</a></li></ol>
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