

**Project Title: Unravelling the role of RNA-binding proteins in Polyglutamine Diseases: from pathogenesis to therapeutics**

**Supervisor: Clévio Nóbrega, PhD**  
Centre for Biomedical Research – University of Algarve  
<http://cbmr.ualg.pt/research/stemcelldevelop/the-molecular-neuroscience-and-therapy-laboratory/cdnobrega@ualg.pt> / 289800100, Ext.7591

**Luís Pereira de Almeida, PhD**  
Center for Neuroscience and Cell Biology – University of Coimbra  
[http://www.cnbc.pt/research/department\\_group\\_show.asp?iddep=1222&idgrp=1254&ldGrupo=1221&hash=33](http://www.cnbc.pt/research/department_group_show.asp?iddep=1222&idgrp=1254&ldGrupo=1221&hash=33)  
[luispa@cnc.uc.pt](mailto:luispa@cnc.uc.pt) / +239829190, Ext. 116

**Location of research lab/research center:**  
Faro, Centre for Biomedical Research – University of Algarve

**Summary: (1000 characters)**

The RNA processing events, which are highly regulated by the RNA-binding proteins (RBPs) play a major role in the regulation of brain functioning<sup>1,2</sup>. More than 50% of known RBPs are expressed in the brain and recent studies suggested that a deregulation in RBPs functions and RNA processing events might underlie the pathogenesis of several neurodegenerative diseases (NDs)<sup>3</sup>. The main goal of this project is to investigate the role of RBPs in the context of Polyglutamine (polyQ) diseases and to identify new targets for therapeutic intervention<sup>4,5</sup>. PolyQ diseases are a group of 9 incurable, inherited NDs caused by an abnormal expansion of a CAG tract, having no therapy currently available to stop or delay disease progression. The project will combine high-throughput screening (e.g. transcriptomics of polyQ patients brain samples), *in vitro* models (e.g. neurons derived from polyQ patients iPS cells) and *in vivo* models (e.g. lentiviral and transgenic polyQ mouse models). Importantly, it will be developed in collaboration with the Center for Neuroscience and Cell Biology (Portugal) and the biotech company BrainVectis (France).

**Bibliographic references:**

<sup>1</sup> Bryant CD, Yazdani N, 2016. RNA-binding proteins, neural development and the addictions. *Genes Brain Behav*, 15:169-86.

<sup>2</sup> Carmo-Silva, Nóbrega C, Pereira de Almeida L, Cavadas C, 2017, Unravelling the role of ataxin-2 in metabolism. *Trends in Endocrinology and Metabolism*, 28:309-318

<sup>3</sup> Matos CA, Pereira de Almeida L, Nóbrega C, 2017, Machado-Joseph disease/Spinocerebellar ataxia type 3: from pathogenic mechanisms to therapeutic strategies. *Journal of Neurochemistry*, *in press*.

<sup>4</sup> Nóbrega C, Carmo-Silva S, Albuquerque D, Vasconcelos-Ferreira A, Vijayakumar U-G, Mendonça L, Hirai H, Pereira de Almeida L, 2015, Reestablishing Ataxin -2 downregulates translation of mutant ataxin-3 and alleviates Machado-Joseph disease. *Brain*, 138: 3537-54.

<sup>5</sup> Nascimento-Ferreira I\*, Nóbrega C\*, Ferreira-Vasconcelos A, Onofre I, Albuquerque D, Avelaira C, Hirai H, Déglon N, Pereira de Almeida L, 2013, Beclin-1 mitigates motor and neuropathological deficits in genetic mouse models of Machado-Joseph disease. *Brain*, 136:2173-88. \*equal contribution