

Alma Mater Studiorum Università di Bologna Archivio istituzionale della ricerca

Hyperactivation of the Hippo signalling in the Gaucher disease

Published Version:

This is the submitted version (pre peer-review, preprint) of the following publication:

Silvia Strocchi, D.M. (2018). Hyperactivation of the Hippo signalling in the Gaucher disease.

Availability: This version is available at: https://hdl.handle.net/11585/649689 since: 2018-11-14
Published:
DOI: http://doi.org/
Terms of use:
Some rights reserved. The terms and conditions for the reuse of this version of the manuscript are

This item was downloaded from IRIS Università di Bologna (https://cris.unibo.it/). When citing, please refer to the published version.

Hyperactivation of the Hippo Signalling in the Gaucher Disease

Silvia Strocchi¹, Daria Messelodi², Andrea Pession^{2,3}, Daniela Grifoni¹

- 1. Cance REvolution Lab, Department of Pharmacy and Biotechnology, University of Bologna, Bologna, Italy
- 2. Medical and Surgical Department, University of Bologna, Bologna, Italy
- 3. Inter-Departmental Centre for Cancer Research "Giorgio Prodi", University of Bologna, Bologna, Italy

In the last 10 years, hundreds of studies have focused on the so-called "Hypo-Hippo" condition, associated with cancer development, where the downstream effector of the Hippo signalling pathway YAP (Yki in flies) is activated and upregulates its target genes, leading to unrestrained proliferation. More recently, a "Hyper-Hippo" condition has been associated with neurodegenerative phenotypes. In the few studies so far available, hyper-activation of upstream components of the pathway has been found to cause YAP/Yki inactivation and to block growth signals. This results in neuroinflammation and neuronal cell death both in mammals and in *Drosophila*. The Gaucher Disease (GD), described as the most common lysosomal disorder, arises from mutations in the GBA1b gene, which encodes the βglucocerebrosidase acid enzyme. In more than 90 percent of patients, these mutations are responsible for a systemic disorder that can be treated with a substitutive enzymatic therapy. Few patients, often affected by the same mutation as the systemic Gaucher patients, develop a neuronopathic form, with neuroinflammation and neuronal cell death, and lack effective therapy. This observation opens to the possible involvement of other genes and pathways in this form of GD. Taking advantage of a *Drosophila* Gaucher-like model based on dGBA1b knock-out $(GBA1b^{KO})$, we analyzed the Hippo pathway activity in the mutant context. We found deregulation of some Yki target molecules, such as CycE, dIAP and MYC, which were all severely reduced both in terms of transcript and protein in the GBA1b^{KO} context, suggesting a cell growth impairment. We also analyzed upstream components and found that Fat, an atypical cadherin upstream of the kinase complex, is up-regulated. In the light of this, we decided to examine some soluble factors known to be negatively regulated by Fat and involved in glial and synapse development, the *Drosophila* glypicans Dally and Dally-like. As expected, we found them downregulated in the GBA1b^{KO} background, and we do believe this may be correlated with the neurological damage characteristic of the neuronopathic Gaucher. Preliminary functional data will be presented which support our current hypothesis.