

## Research Project

MHV - Investigating the function of GABAb receptor-interacting proteins by FRET assays

## Third-party funded project

**Project title** MHV - Investigating the function of GABAb receptor-interacting proteins by FRET assays **Principal Investigator(s)** Jacquier, Valérie; Bettler, Bernhard;

Organisation / Research unit

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GABA $_B$  receptors are the principal inhibitory G-protein coupled receptors in the brain and hold considerable promise as therapeutic targets for the treatment of neurological and psychiatric disorders, including anxiety, depression, addiction, epilepsy, and chronic pain. They are composed of two different subunits, GABA $_{B1}$  and GABA $_{B2}$ .

For a long time, it was believed that the different responses observed in the brain in response to  $\mathsf{GABA}_B$  receptor activation reflected the presence of different  $\mathsf{GABA}_B$  receptor subtypes. However, we now know that the only molecular diversity in the  $\mathsf{GABA}_B$  system arises from the two isoforms of the  $\mathsf{GABA}_{B1}$  subunit:  $\mathsf{GABA}_{B1a}$  and  $\mathsf{GABA}_{B1b}$ . Surprisingly, these two isoforms have very similar pharmacological and biophysical properties *in vitro*, which makes it difficult to explain the pharmacological and kinetic differences of native  $\mathsf{GABA}_B$  responses.

A family of four sequence-related proteins were recently found to interact with the receptor and to modify the  $GABA_B$ -mediated response when coexpressed with the receptor *in vitro*. The present project aims at elucidating the molecular events underlying the kinetics changes that these auxiliary proteins impose on the  $GABA_B$  receptor response. Assays based on Fluorescence Resonance Energy Transfer (FRET) will be used to investigate whether these proteins interact with different components of the  $GABA_B$ -mediated signalling cascade, and how these interactions are modulated upon receptor activation.

These experiments will allow us to provide a mechanistic explanation of how these proteins modify kinetic and pharmacological properties of  $\mathsf{GABA}_B$  responses and contribute to the functional heterogeneity observed with native  $\mathsf{GABA}_B$  responses. Identification of the function of these proteins will spark drug discovery efforts, as this provides an opportunity for a more selective interference with the  $\mathsf{GABA}_B$  receptor system.

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