

Amelia Haigh, Nagore Elu, Susana R Louros

Centre for Discovery Brain Sciences, Simons Initiative for the Developing Brain | The University of Edinburgh, Scotland, UK

1. Introduction

- Fragile X Syndrome is caused by mutations in Fragile X Messenger Ribonucleoprotein 1 (*Fmr1* gene), and is the leading monogenic cause of autism spectrum disorder (Stone et al., 2023)
- The ubiquitin proteasome system (UPS) is involved in the hyperexcitability of *Fmr1*^{-/-} mice after acoustic stimulation (used to model audiogenic seizures); a phenotype that is corrected by administering the proteasome inhibitor Bortezomib (BTZ) (Louros et al., 2023)
- My project assessed how UPS dysfunction contributes to the hyperexcitability in FXS, particularly in the inferior colliculus (IC). *Fmr1*^{-/-} mice were crossed with lines expressing Ub^{G76V}-Green Fluorescent Protein (GFP), a reporter first described by Lindsten et al (2003)
- We aimed to identify the neurons where BTZ administration inhibited the activity of the proteasome, leading to the accumulation of Ub^{G76V}-GFP

2. Methodology

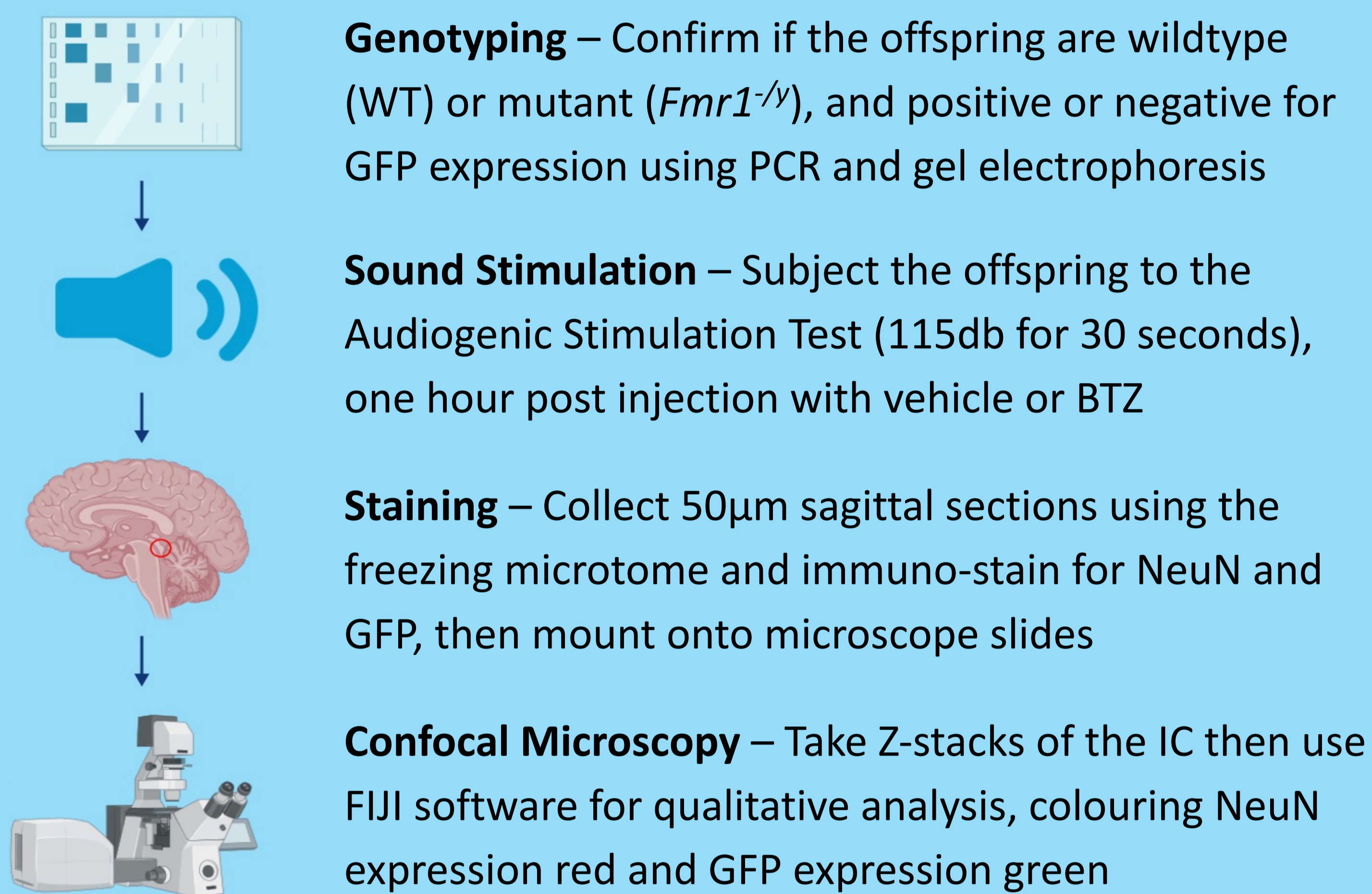


Figure 1. Schematic of methodology used to assess UPS dysfunction. Created in BioRender [10/08/24]

3. Ub^{G76V}-GFP reporter is functional

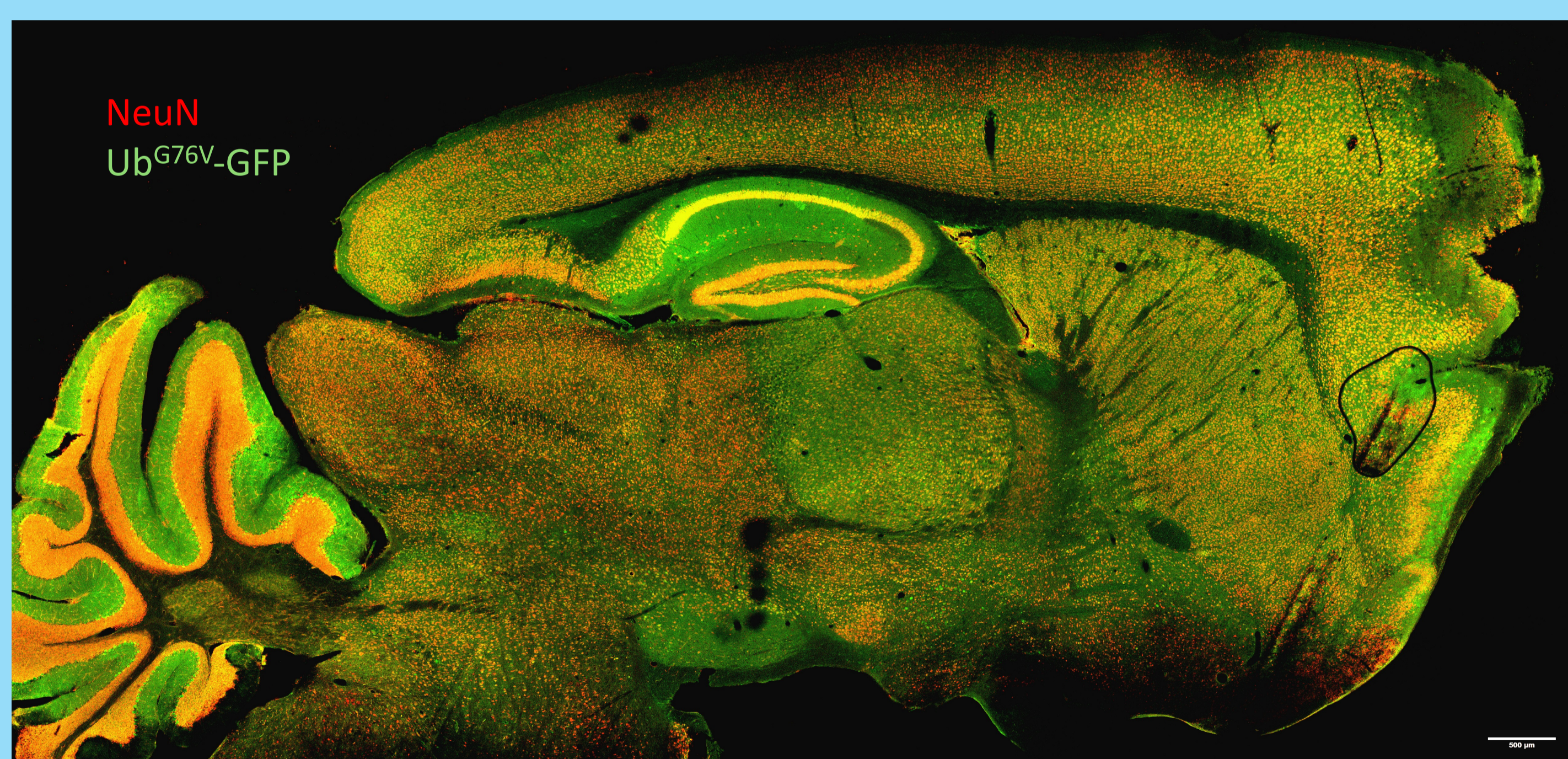


Figure 2. Neuronal expression of the proteasome reporter in the new FX-Ub^{G76V}-GFP mouse line

4. BTZ administration increases GFP expression in *Fmr1*^{-/-} IC

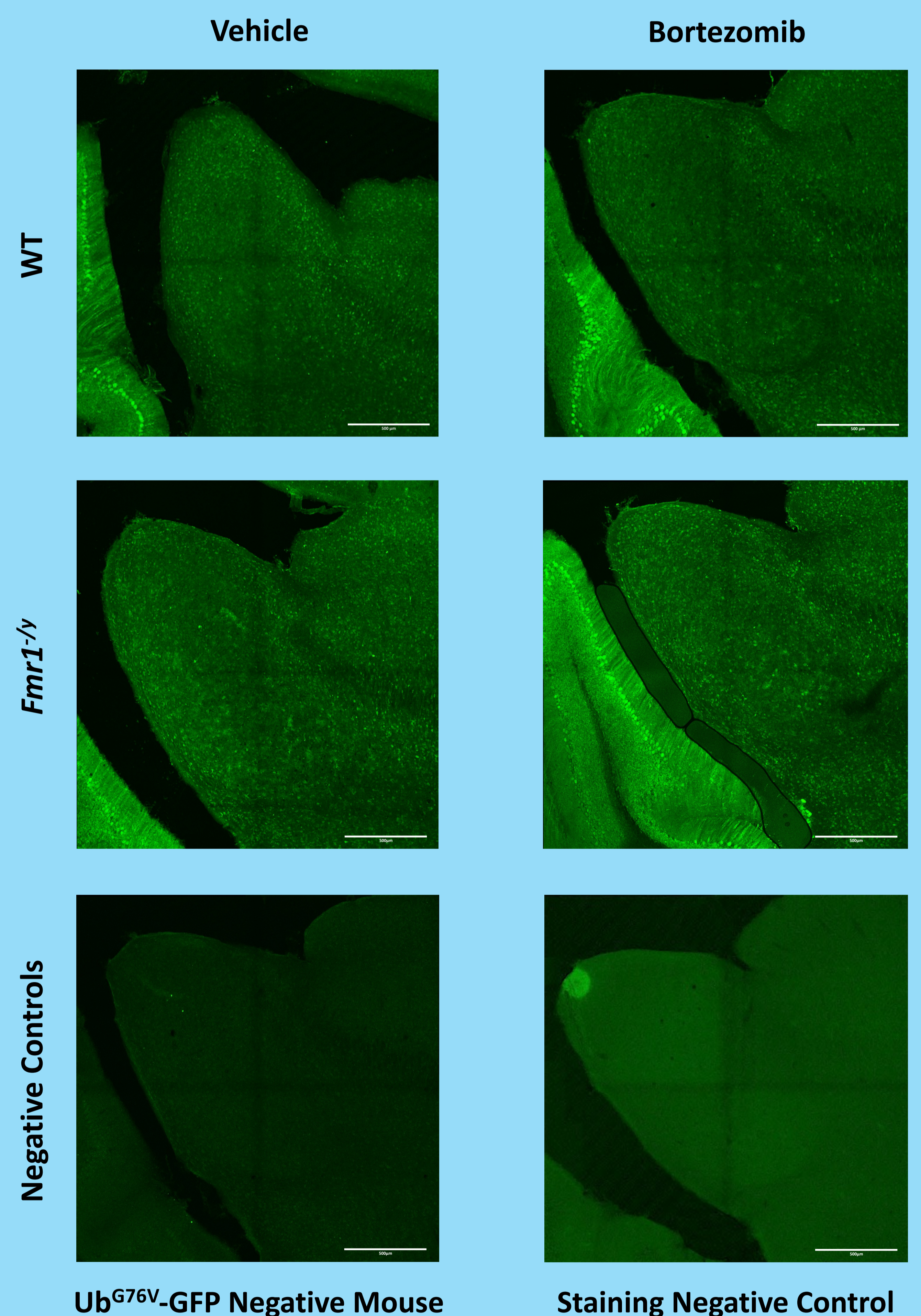


Figure 3. Ub^{G76V}-GFP expression in the inferior colliculus (IC) of wild-type (WT) and *Fmr1*^{-/-} mice, following injection with vehicle or Bortezomib. Proteasome inhibition causes a visible increase in Ub^{G76V}-GFP expression in the *Fmr1*^{-/-} mouse

5. Future Directions

- *Fmr1*^{-/-} mice exhibit a higher expression of GFP, indicating proteasomal inhibition causes impaired GFP degradation
- Future experiments should use Imaris software to quantitatively analyse GFP expression, and determine whether the differences between WT and *Fmr1*^{-/-} colonies are significant
- Tissue sections should be triple immunostained for NeuN, GFP, and cFos (neurone activation marker). Observing co-localisation between cFos and GFP expression would indicate that proteasomal activity is more inhibited in hyperactivated neurones, generating a novel target for the treatment of Fragile X Syndrome

6. References

- Lindsten, K., Menéndez-Benito, V., Masucci, M. G., & Dantuma, N. P. (2003). A transgenic mouse model of the ubiquitin/proteasome system. *Nature biotechnology*, 21(8), 897–902, <https://doi.org/10.1038/nbt851>
- Louros, S. R., Seo, S. S., Maio, B., Martinez-Gonzalez, C., Gonzalez-Lozano, M. A., Muscas, M., Verity, N. C., Wills, J. C., Li, K. W., Nolan, M. F., & Osterweil, E. K. (2023). Excessive proteostasis contributes to pathology in Fragile X syndrome. *Neuron*, 111(4), 508–525.e7, <https://doi.org/10.1016/j.neuron.2022.11.012>
- Stone, W. L., Basit, H., Shah, M., & Los, E. (2023). Fragile X Syndrome. In *StatPearls*. StatPearls Publishing.