



# Live imaging of RNA dynamics for genetic forms of ALS in zebrafish

Raphael Munoz-Ruiz, Doris Lou Demy, Hortense de Calbiac, Sorana Ciura, Etor Kabashi



**Raphaël Munoz-Ruiz**

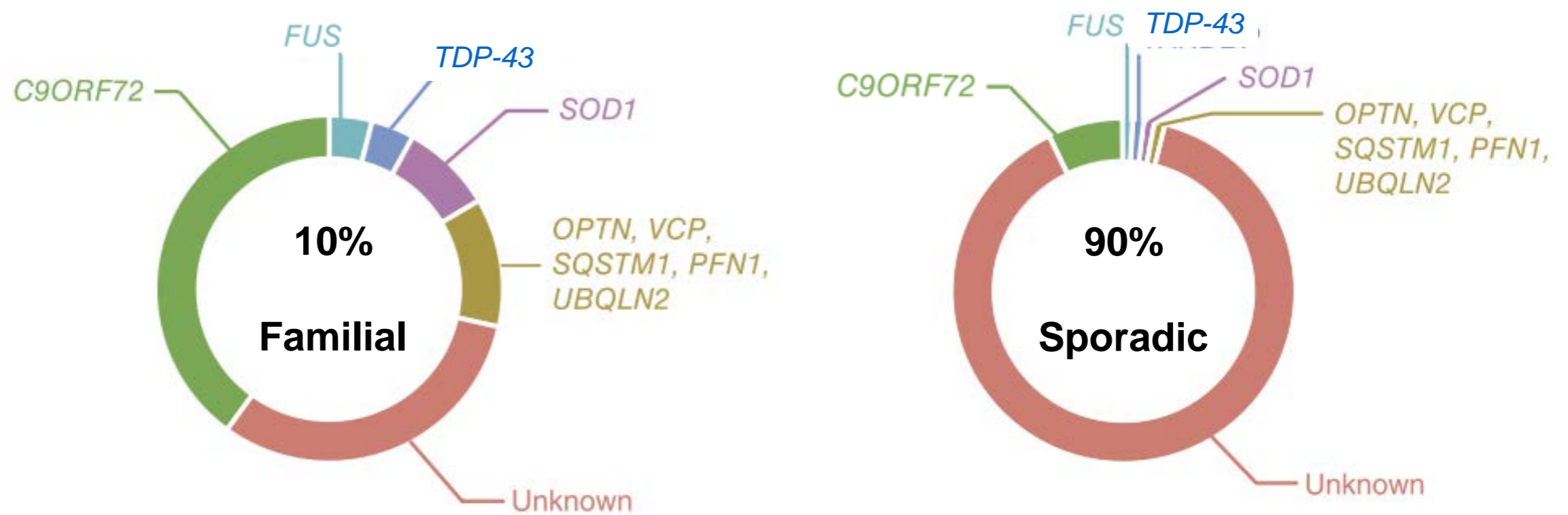
PhD student

Etor Kabashi's lab

« Treatments of ALS: from genetics to zebrafish »  
Institut du Cerveau et de la Moelle Epinière - Paris

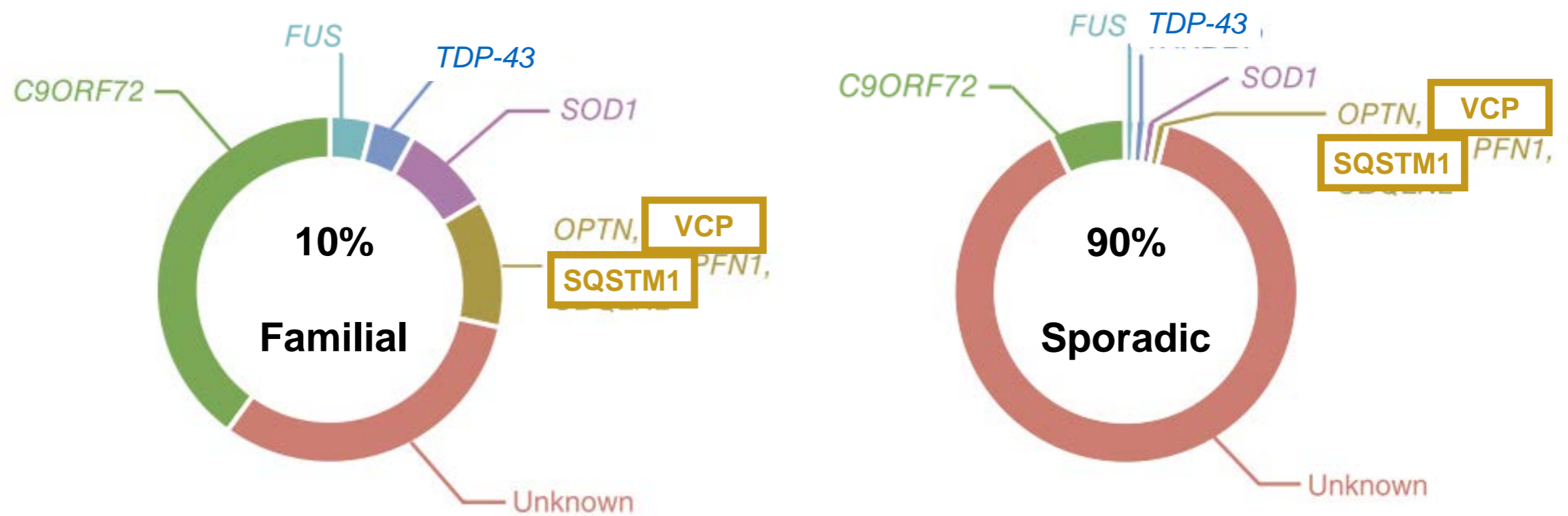


# The Genetics of ALS



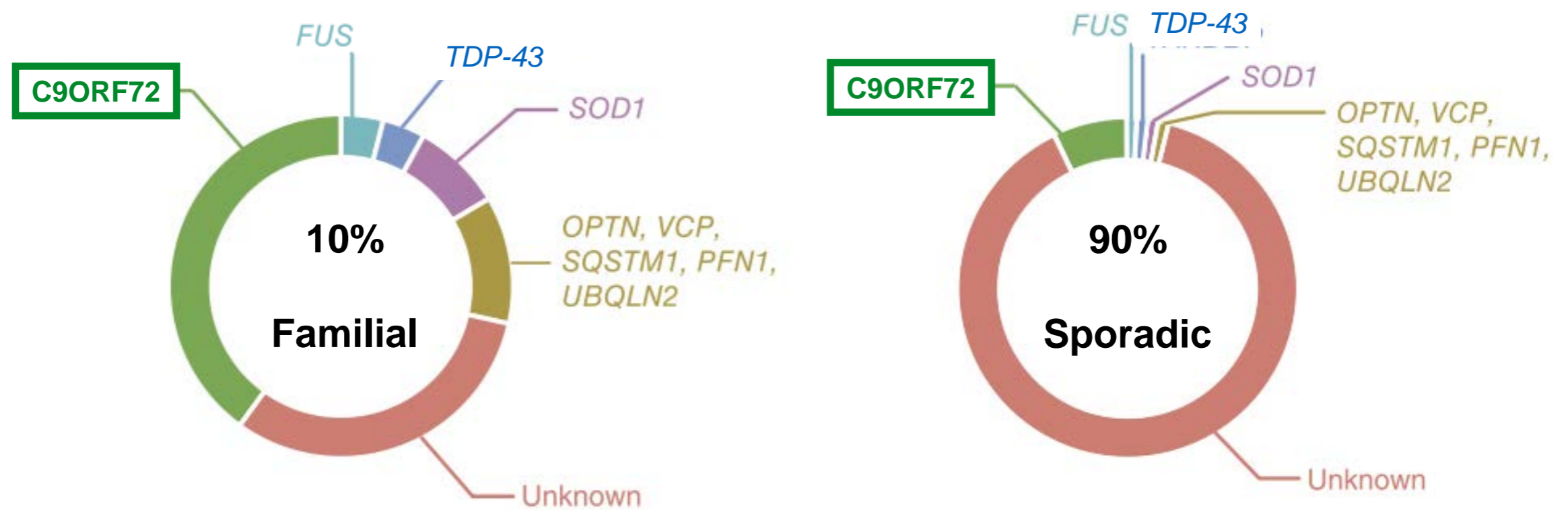
Modified from **Renton** et al. 2014

# The Genetics of ALS



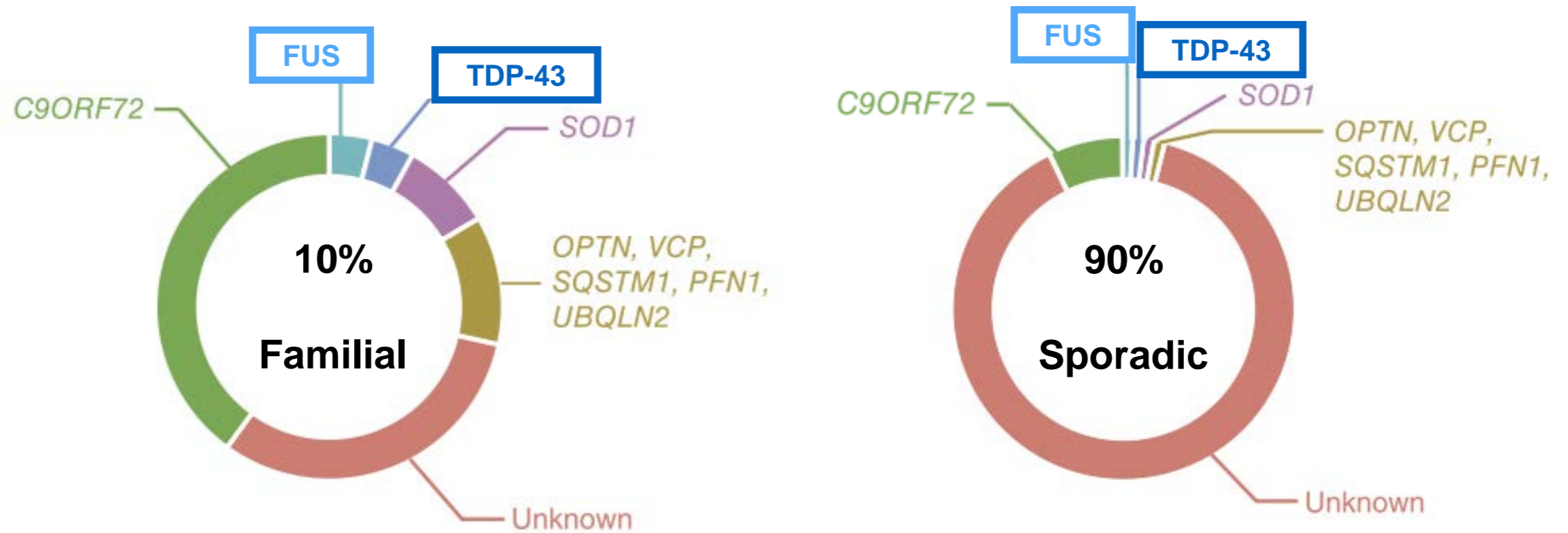
Modified from **Renton** et al. 2014

# The Genetics of ALS



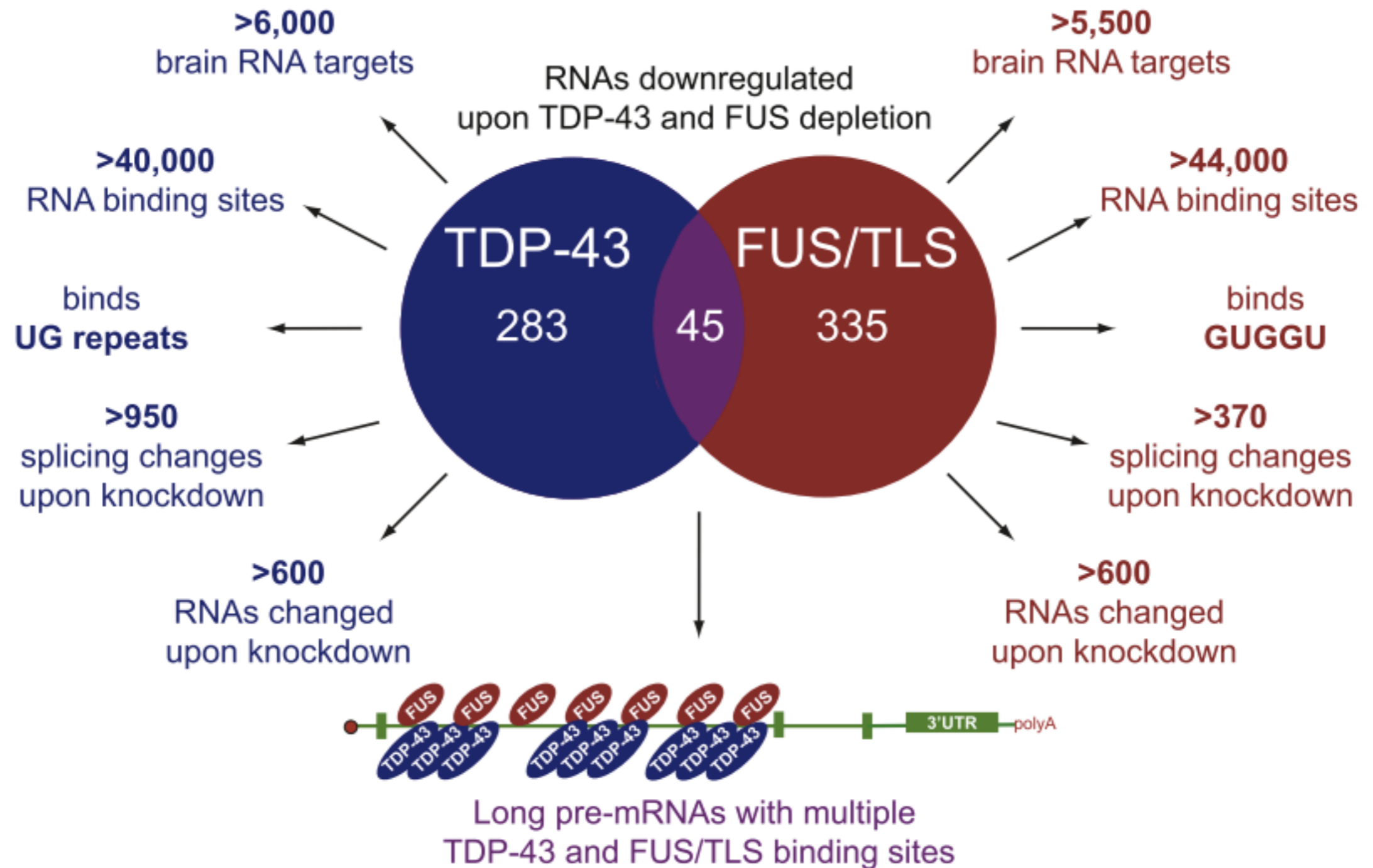
Modified from **Renton** et al. 2014

# The Genetics of ALS



Modified from **Renton** et al. 2014

# TDP-43 and FUS: RNA-binding proteins

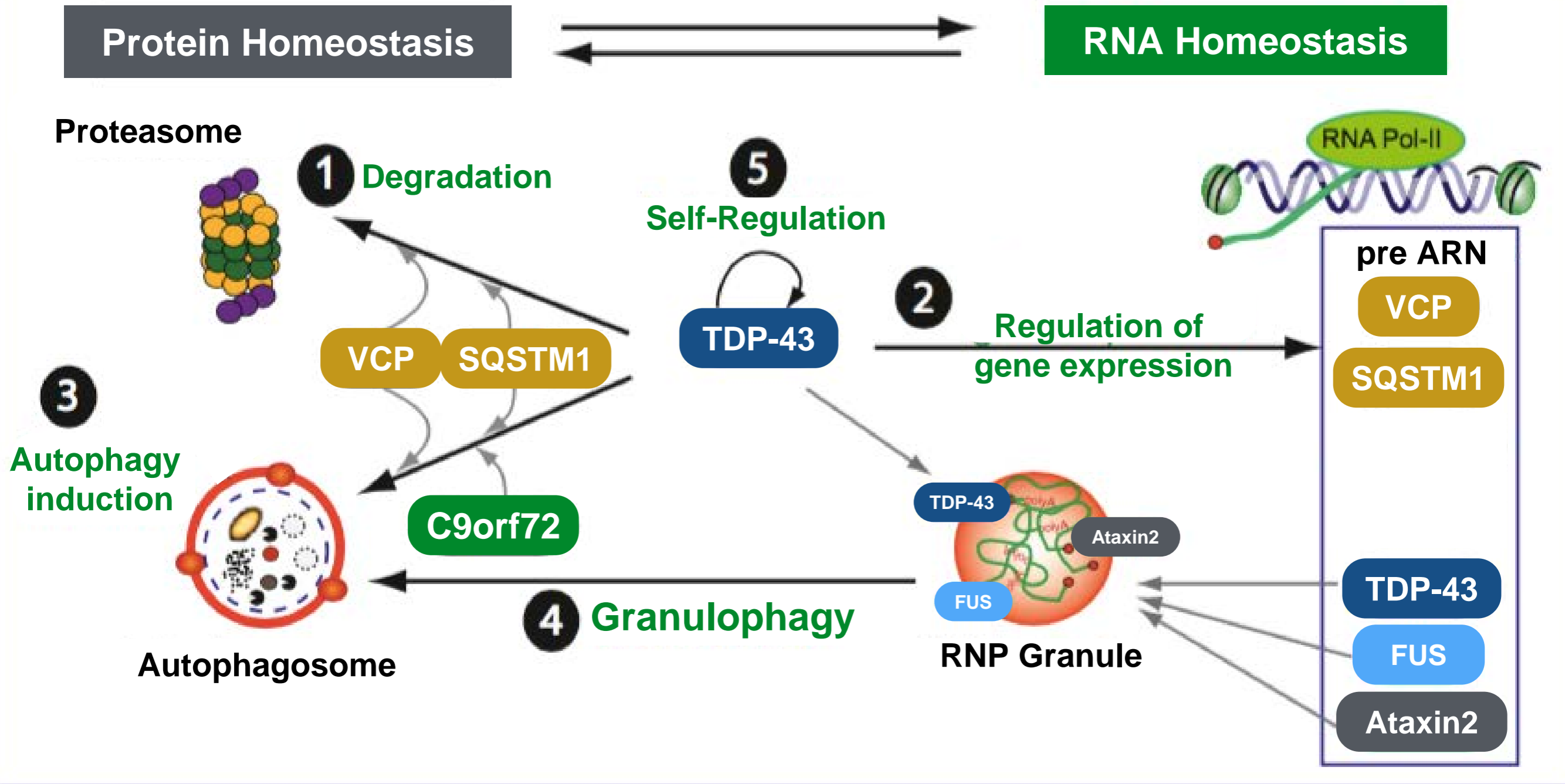


(from **Ling** et al. 2013)



# Protein and RNA homeostasis

## TDP-43 self-regulatory mechanism

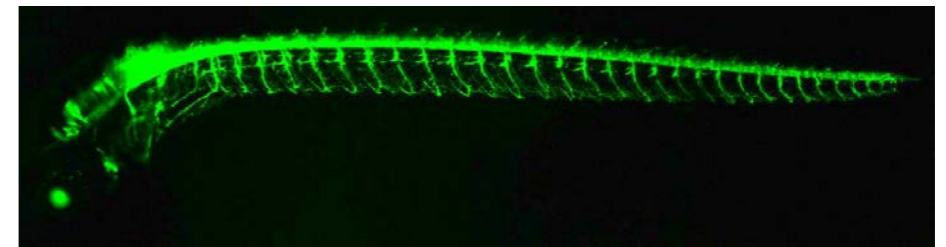
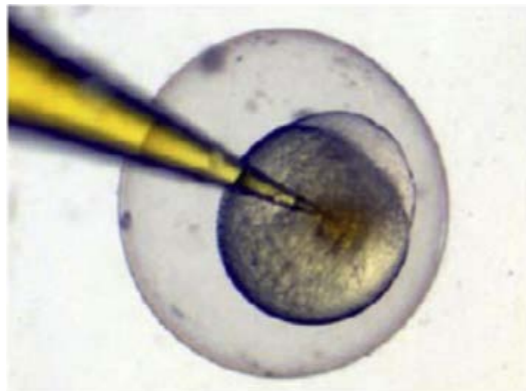


# *In vivo* study of RNA dynamics: the zebrafish model

- Large progeny generating model
- Vertebrate
- Transparency at embryonic stages
- Great range of genetic tools

=> Generate pathological models

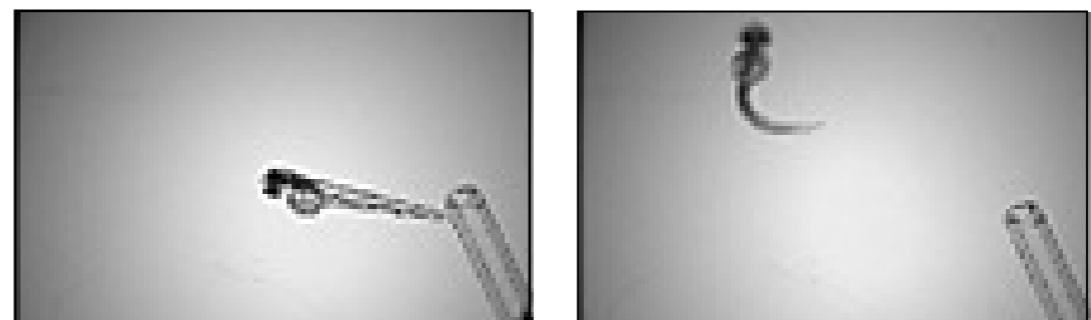
**Injections**



Hb9-GFP transgenic line

=> Unveil genetic interactions

**Touch evoked escape response (TEER)**

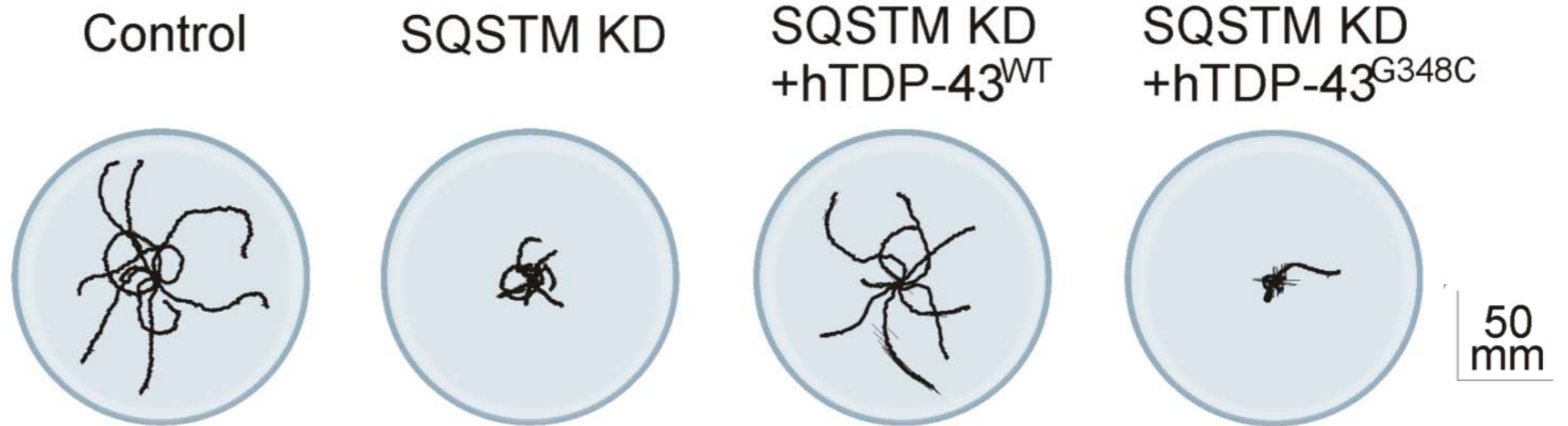


2-3 dpf

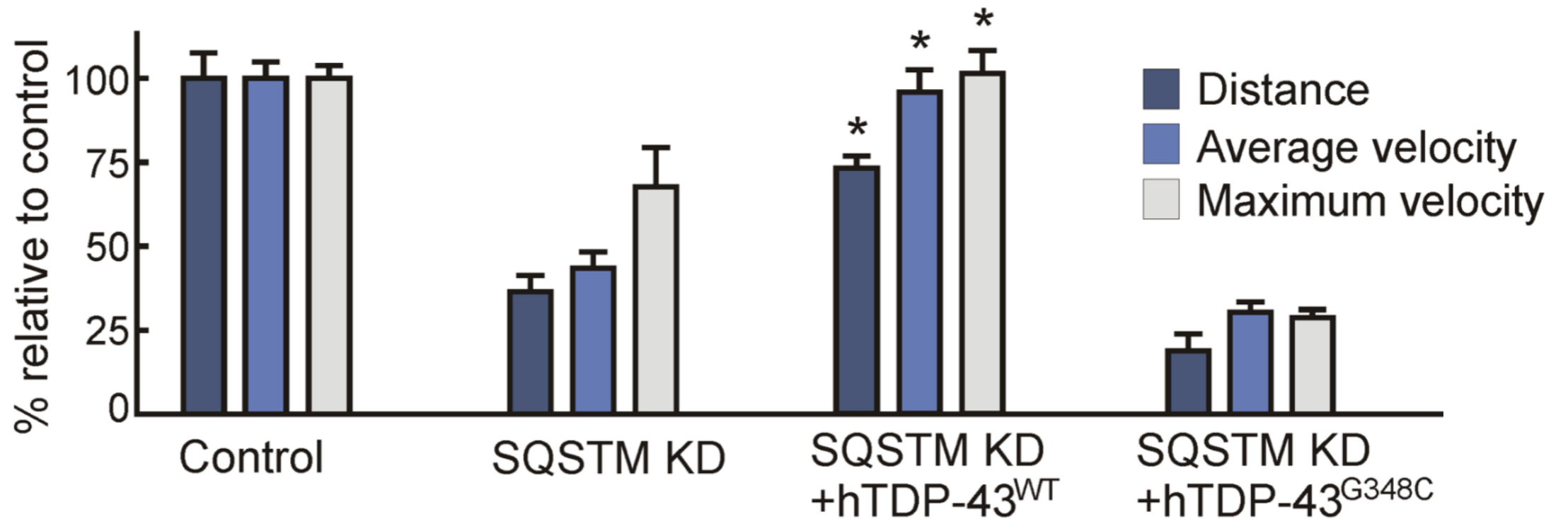


# Genetic interaction between TDP-43 and SQSTM1

Escape response



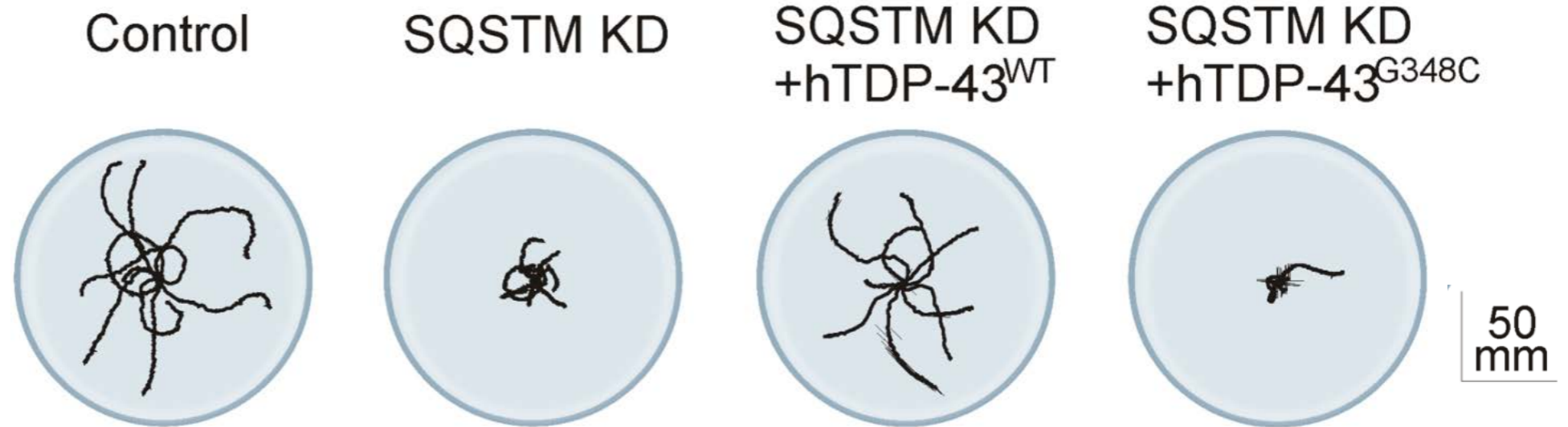
Motor phenotype



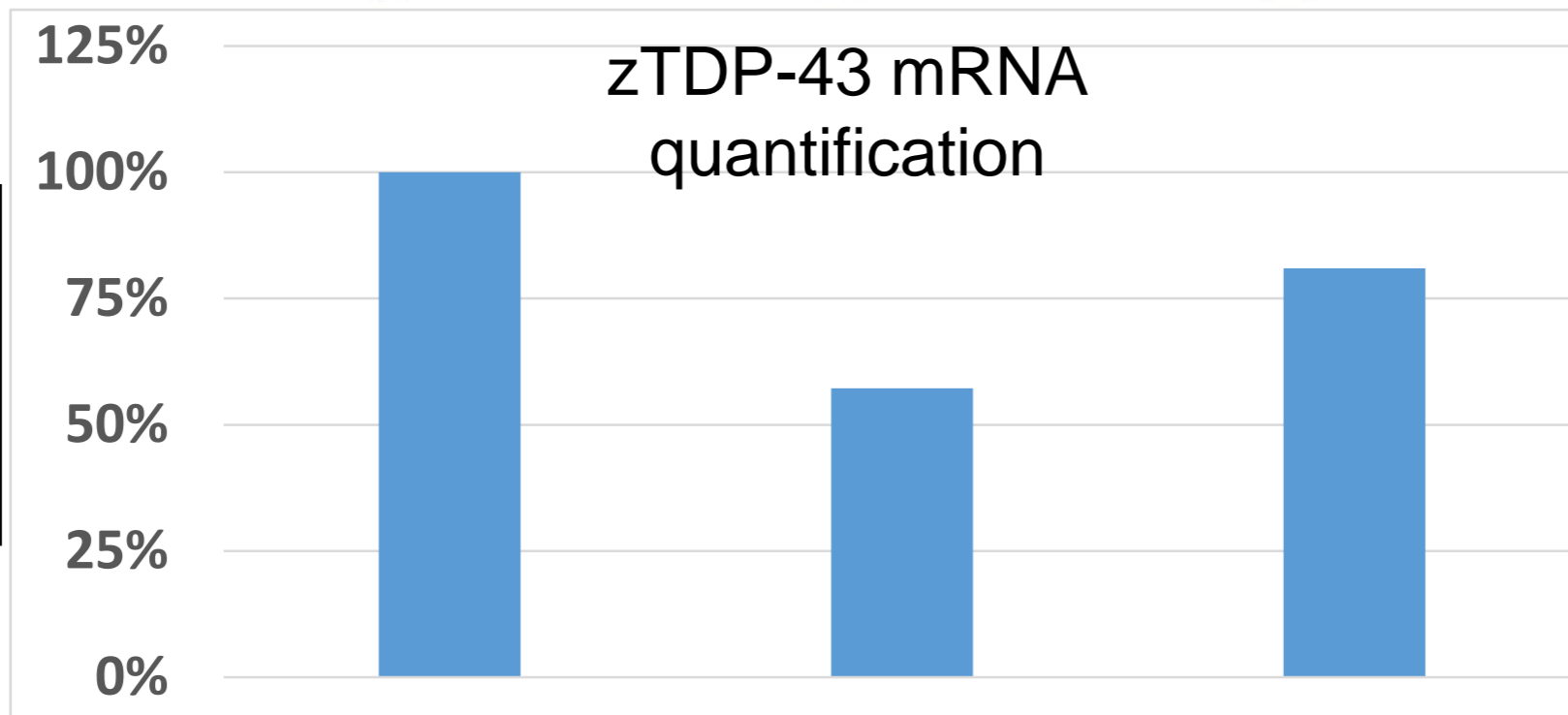
**KD of SQSTM1 leads to reduced motricity**  
**=> significantly ameliorated by human WT TDP-43 RNA**

# Genetic interaction between TDP-43 and SQSTM1

Escape response



qPCR targeting zTDP-43 RNA

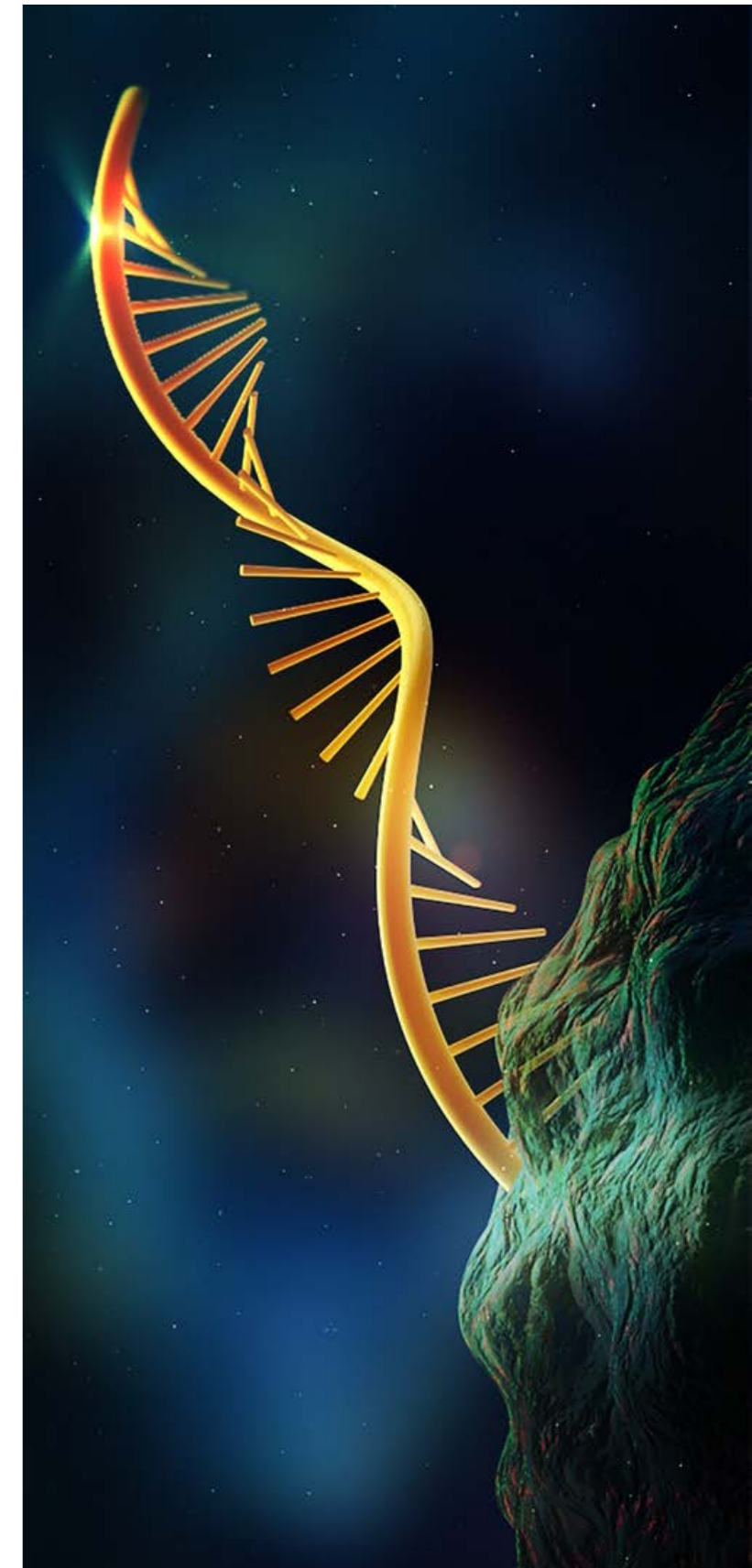


**KD of SQSTM1 leads to reduced zTDP-43 mRNA levels => significantly ameliorated by human WT TDP-43 RNA**

# Studying RNA metabolism in ALS

Understand RNA-binding proteins central role and their interactions with key ALS transcripts (SQSTM1, C9orf72, VCP) in motor neurons

- Precise **localization** and colocalization
- **Interaction partners**
- **Quantification**
- **Dynamics** through live-imaging time-lapse
- **Axonal transport**
- **Healthy VS Pathological condition**

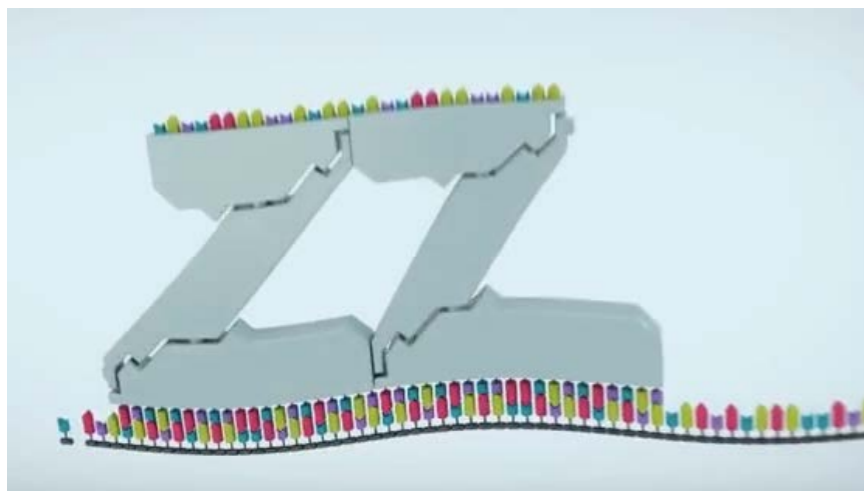


# Investigating RNA metabolism: Precise *in situ* hybridization

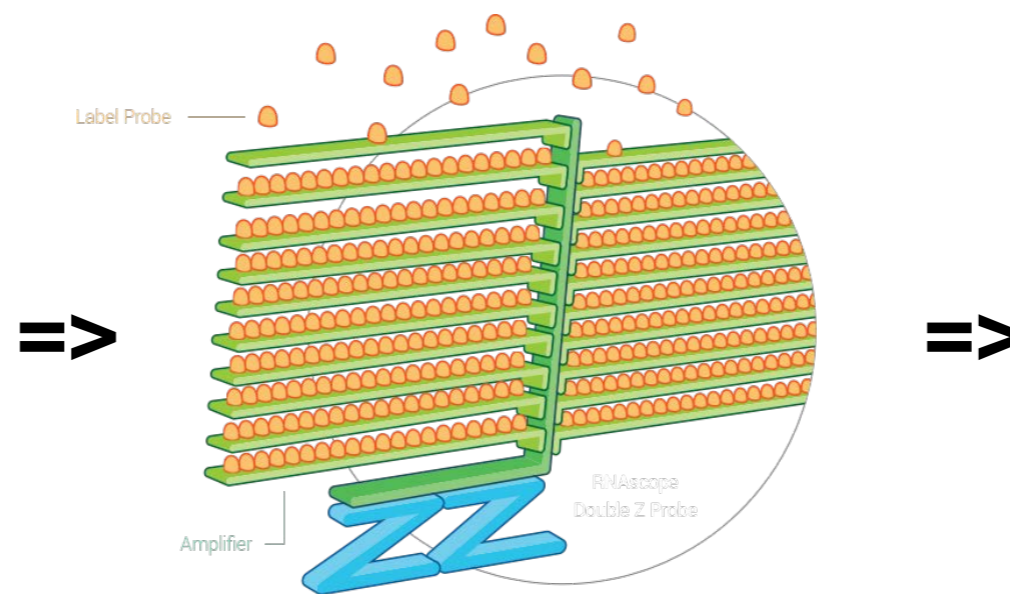
Control FISH with ACD's RNAscope:

- Single-molecule precision

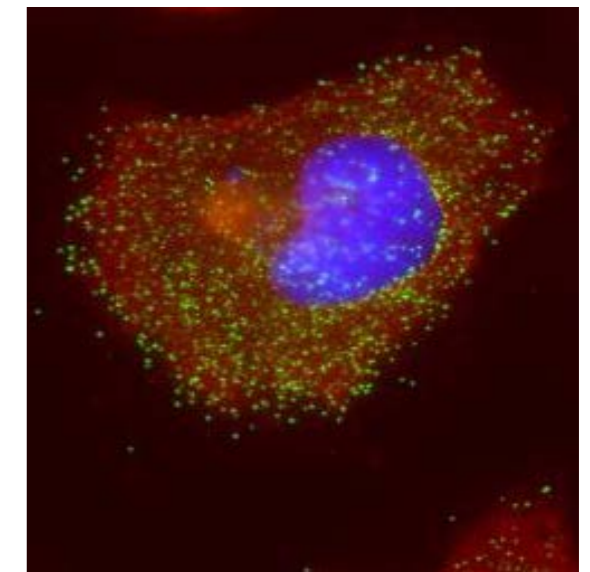
- IHC colocalization



Double Z probe fixation



Signal amplification and  
fluorescent labeling

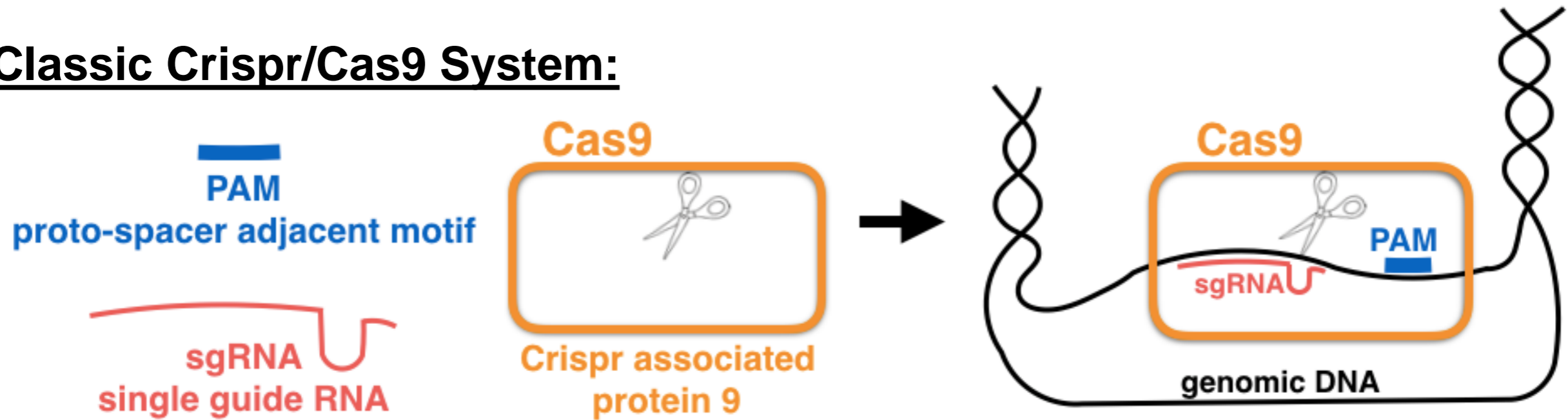


Imaging

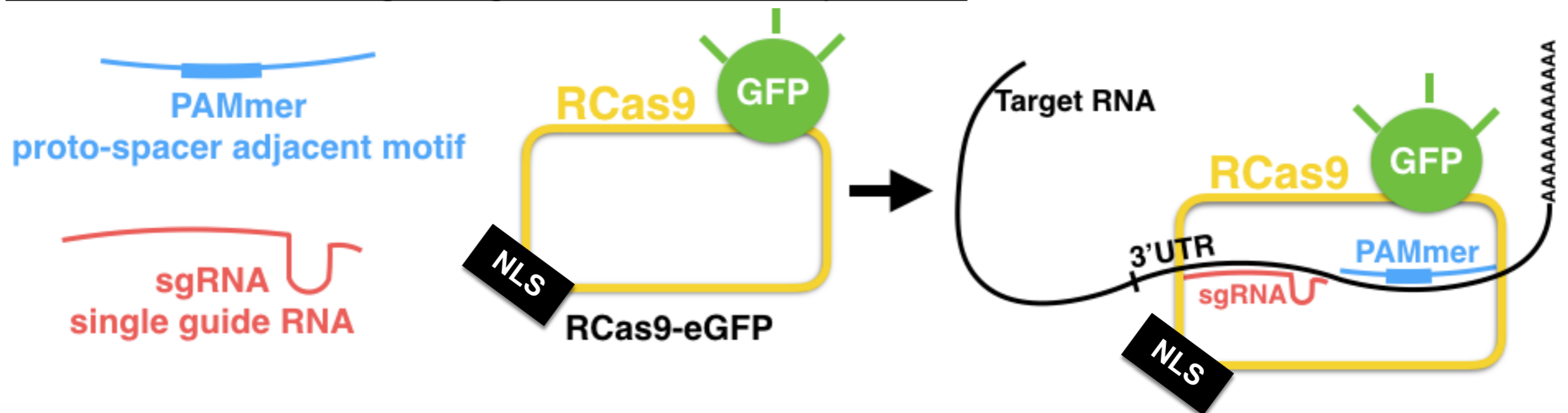


# *In vivo* study of RNA dynamics: a modified CRISPR/Cas9 system

## Classic Crispr/Cas9 System:



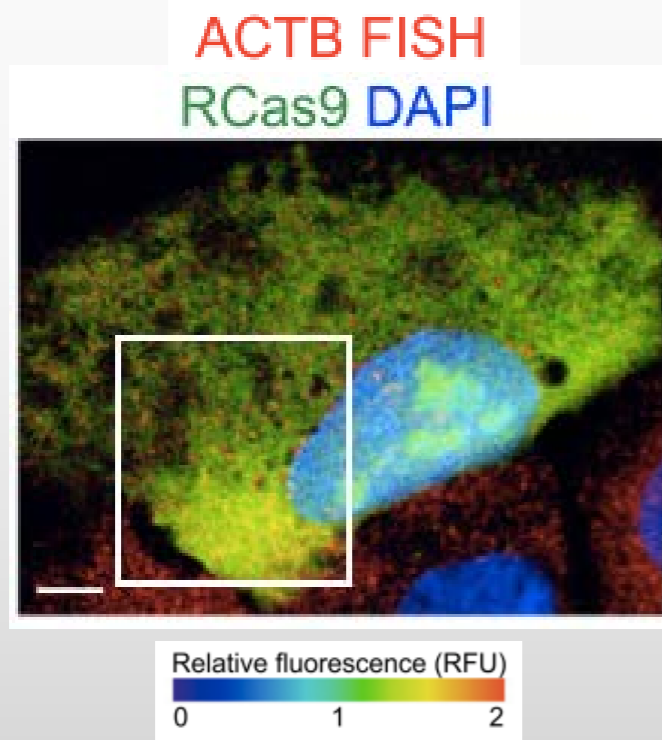
## Modified RNA targeting Crispr/Cas9 System:





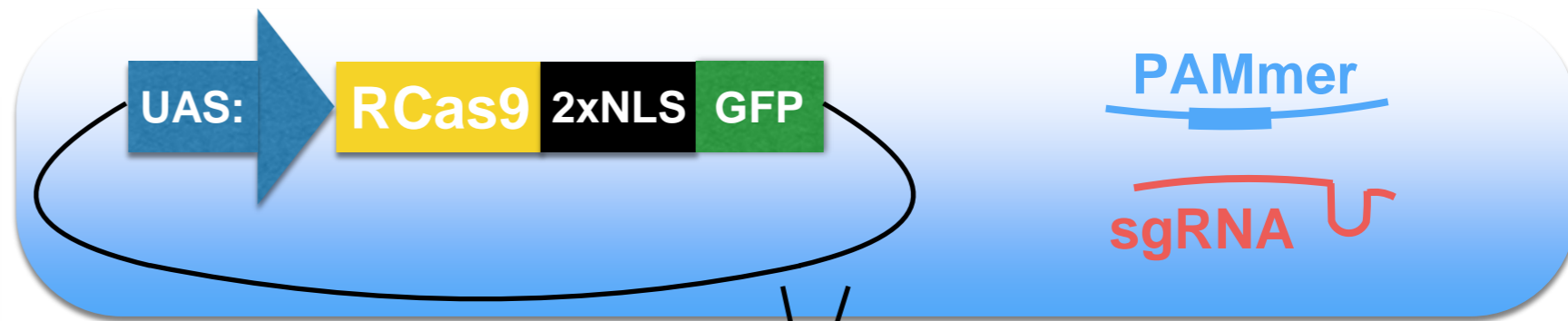
# The RNA targeting CRISPR/Cas9

## In vitro use of the technique:

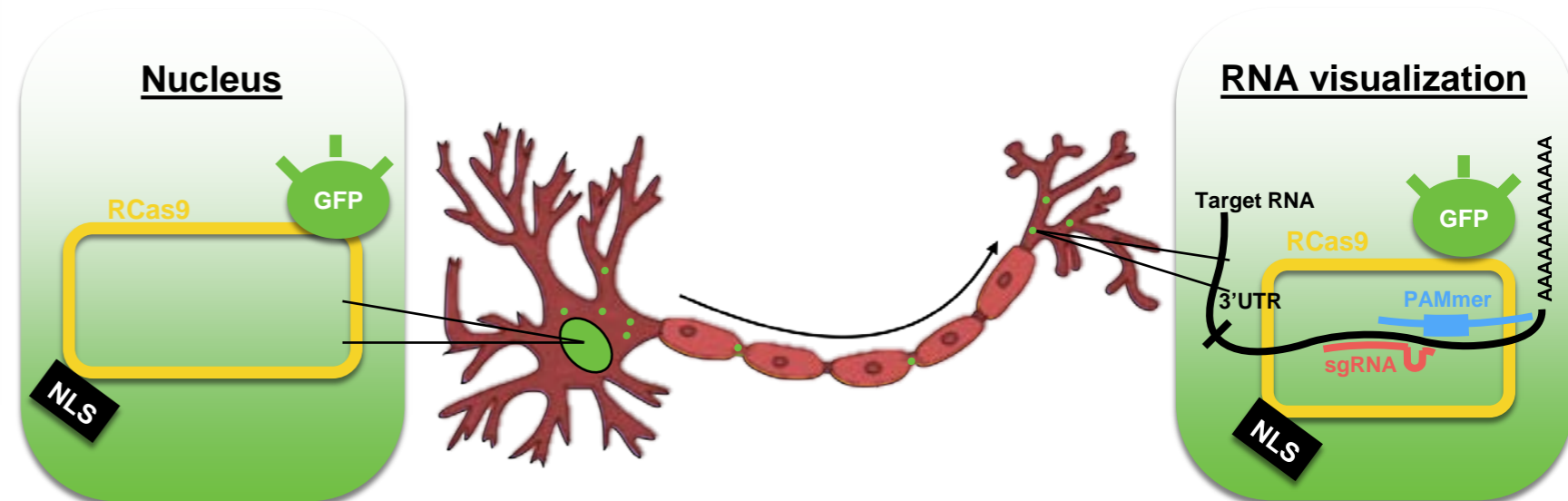
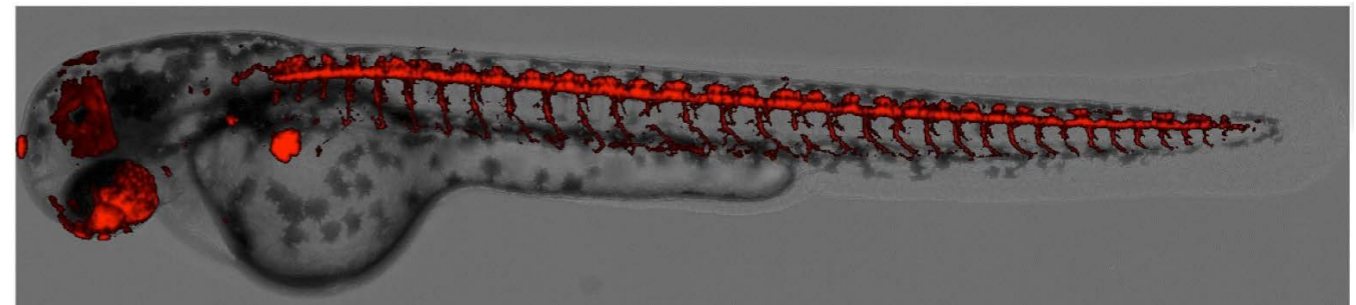


- Targets endogenous RNAs
- No secondary effects
- Localization is highly correlated with control FISH

(from Nelles et al. 2016)



Mnx1:Gal4/UAS:mRFP



**NLS** Nuclear Localisation Signal

**FISH:** Fluorescent *in situ* Hybridization

# Future directions

## **Conclusions:**

- Zebrafish models combines genetic and imaging tools
- TDP-43 and SQSTM1 functionally interact
- First *in vivo* of a modified CRISPR/Cas9 to target RNA

## **Perspectives:**

- Central role of RNA-binding proteins in ALS
- RNA homeostasis defects in pathogenicity
- Precise the level of interactions between key factors

# Acknowledgements

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Hortense de Calbiac

Doris Lou Demy

Anca Marian

Corinne Besnard-Guerin



## Edor Kabashi's Team

and thank you to:    Hervé Tostivint

François Giudicelli

... also featuring  
Maria-Letizia  
Campanari !



**UPMC**  
SORBONNE UNIVERSITÉS





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Visiting fellow  
program



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