

PRION-LIKE PROTEINS, PHASE SEPARATION AND NEURODEGENERATIVE DISORDERS

UAB

Universitat Autònoma de Barcelona

Marcos Gil García

Tutored by Prof. Salvador Ventura Zamora

Biochemistry Degree, Autonomous University of Barcelona, 2016

INTRODUCTION

Proteins are molecules implied in a myriad of tasks in the cell. To perform biological functions, proteins need to fold into defined three-dimensional structures. Mistakes in this complex process produce unfolded states, ultimately leading to the formation of stable **aggregates**.

The role played by intrinsically disordered proteins in cell processes is receiving increasing attention. These proteins promote phase separation states in cytoplasm and punctual mutations in their sequences are behind **neurodegenerative disorders**.

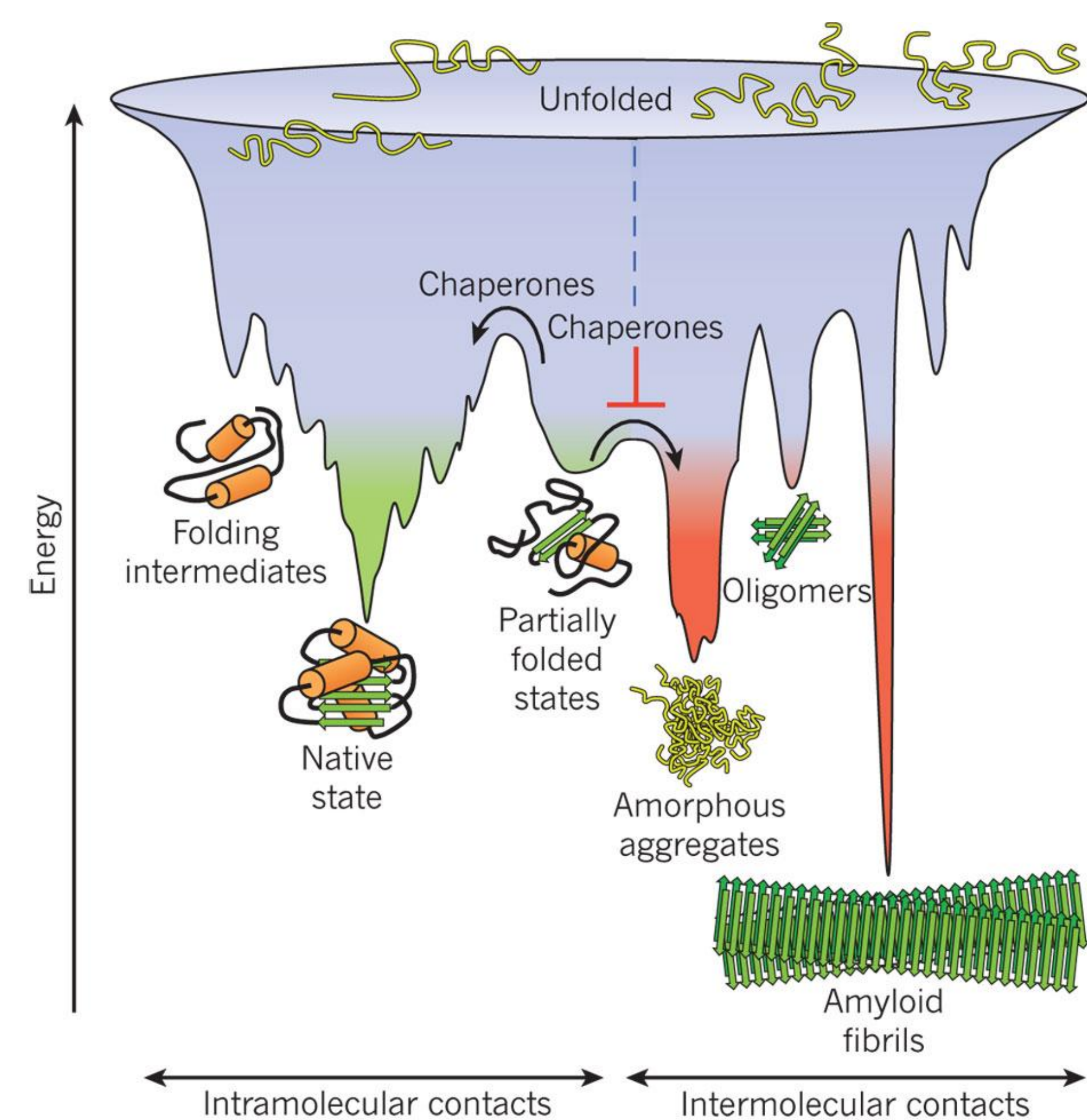


Figure 1: Folding funnel hypothesis [1].

WHAT IS A PRION-LIKE PROTEIN?

Prion-Like proteins are a group of proteins sharing long disordered domains rich in Gly, Tyr and polar amino acids, known as **Low Complexity Domains (LC domain)**. Most of these polypeptides are **DNA or RNA binding proteins** and they are important components of **non-membrane functional structures**.

Fused in Sarcoma (FUS) protein as an example of Prion-Like protein

- ❖ Predominantly nuclear protein
- ❖ Involved in RNA homeostasis
- ❖ Related with Amyotrophic Lateral Sclerosis (ALS)

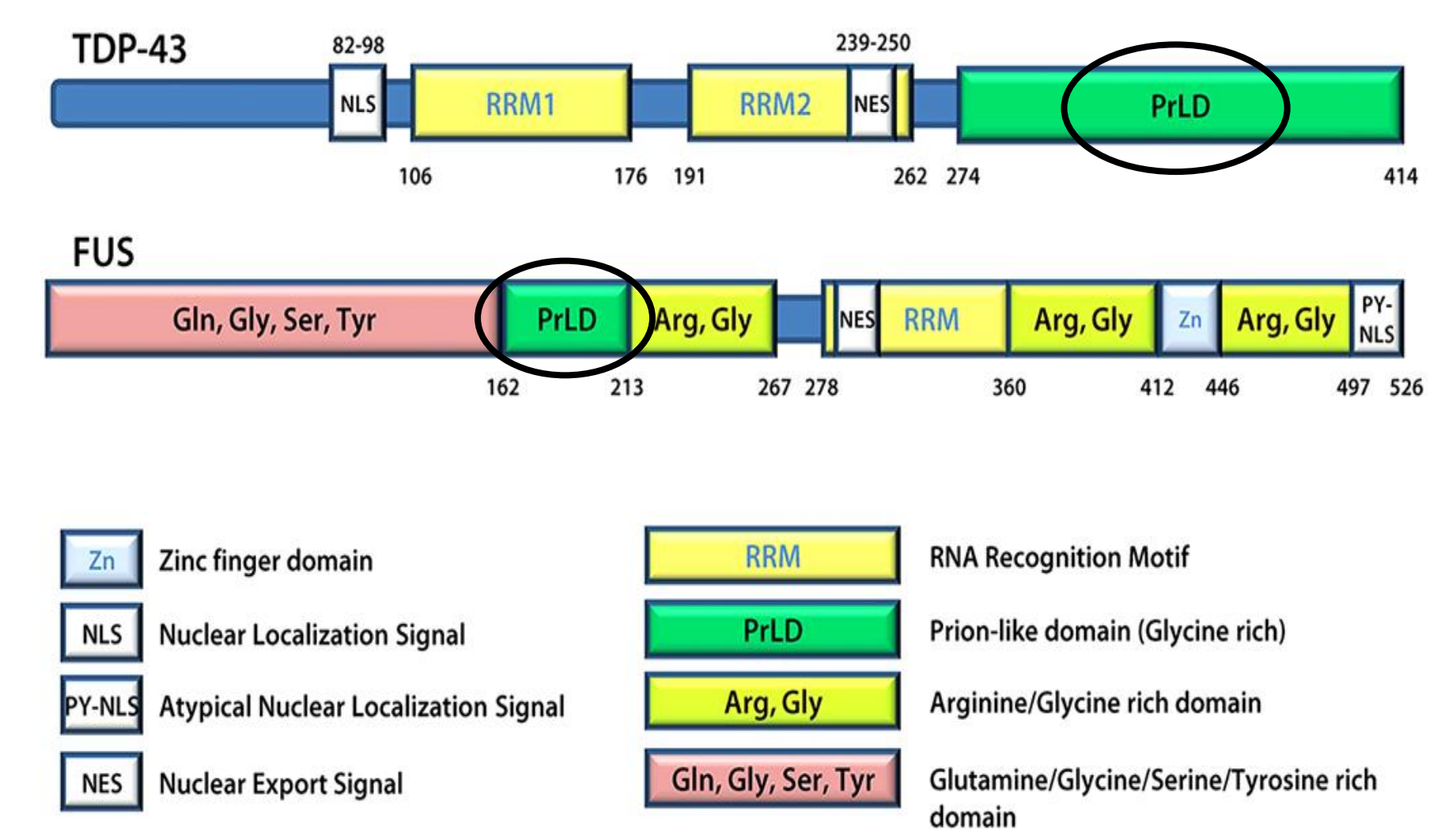


Figure 2: TDP-43 and FUS sequences representation [2].

LIQUID-LIQUID PHASE SEPARATION

Dynamic macromolecular assemblies maintained by specific protein-protein and protein-nucleic acids **weak interactions**.

What is needed for a liquid phase separation?

- ❖ Interactions among Low Complexity Domains
- ❖ A defined protein concentration that does not compromise dynamism

Hallmarks

- Spherical shape
- Dynamic structure
- Fusion ability
- Deformable

Controversy about the role of cross- β structure

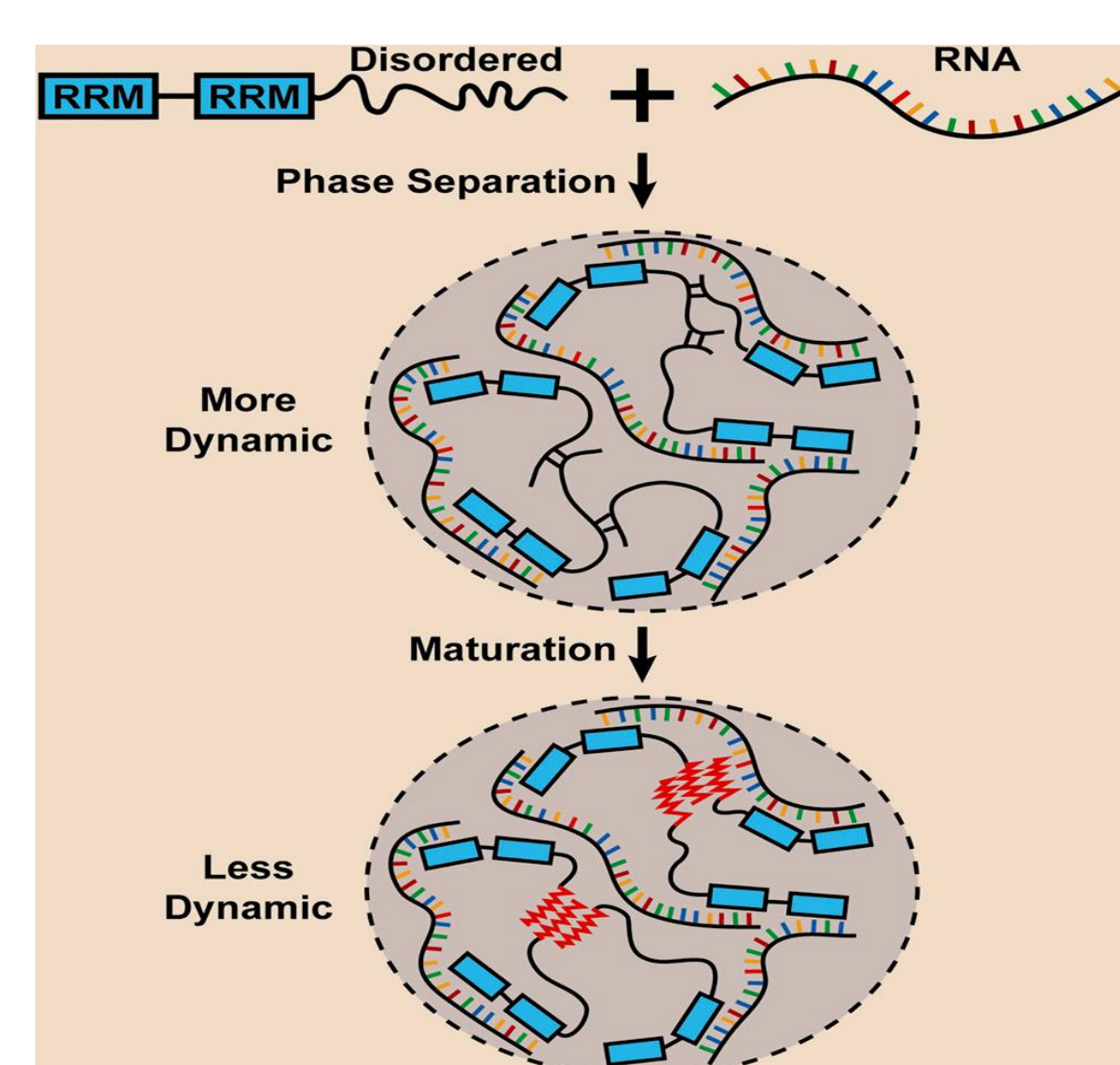
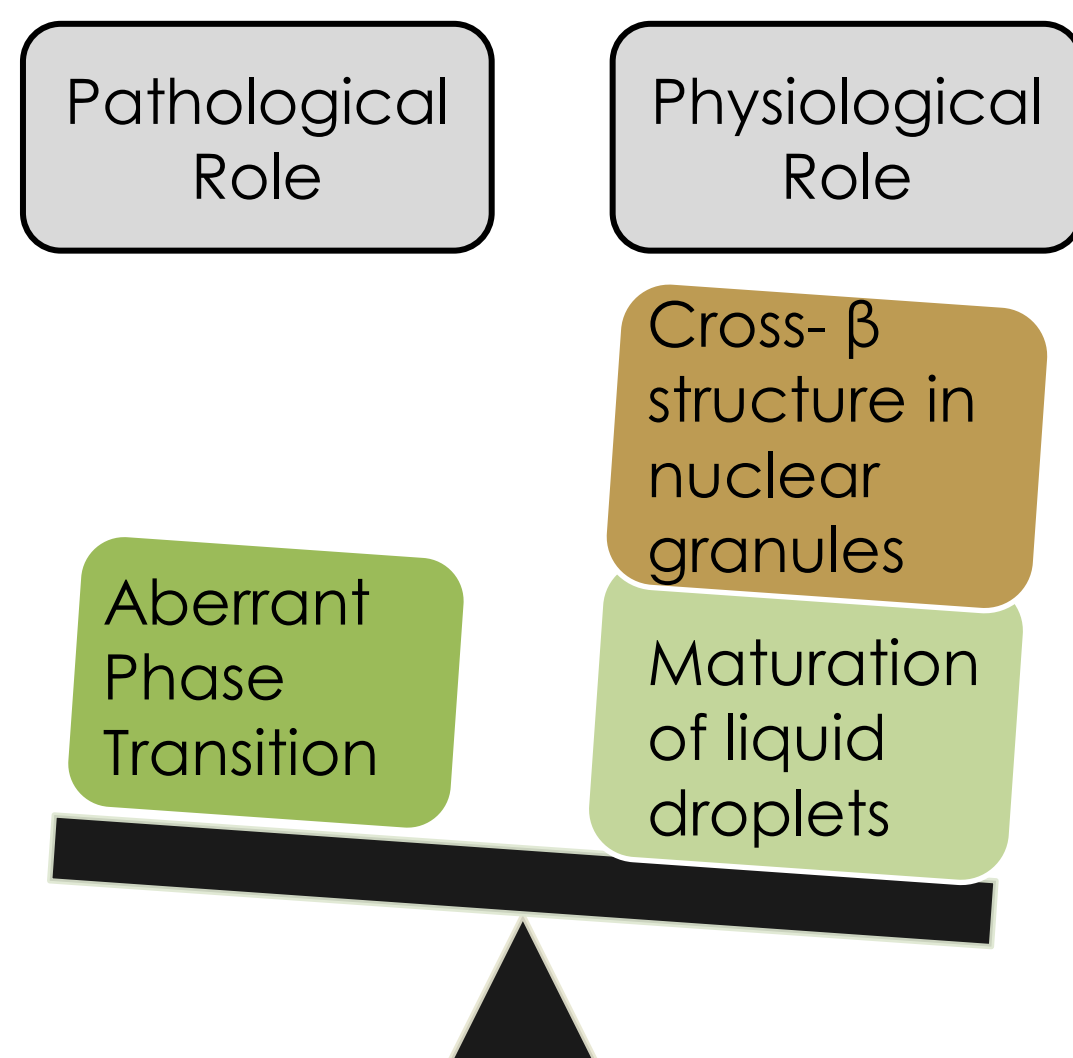


Figure 3: Liquid-liquid phase separation process [3].

ABERRANT PHASE TRANSITION

Mistakes in phase separation equilibrium produce an **aberrant transition** to a fibrous and **pathological** state.

Which is the main source promoting an aberrant phase transition?

- ❖ Mutations in Prion-like proteins
- ❖ High concentration of Prion-like proteins inside phase separation

Low Complexity Domains of Prion-Like proteins are essential to promote phase transition

- ❖ Low Complexity Domain of TIA1 mediates aggregation reactions in the presence of b-isoX compound

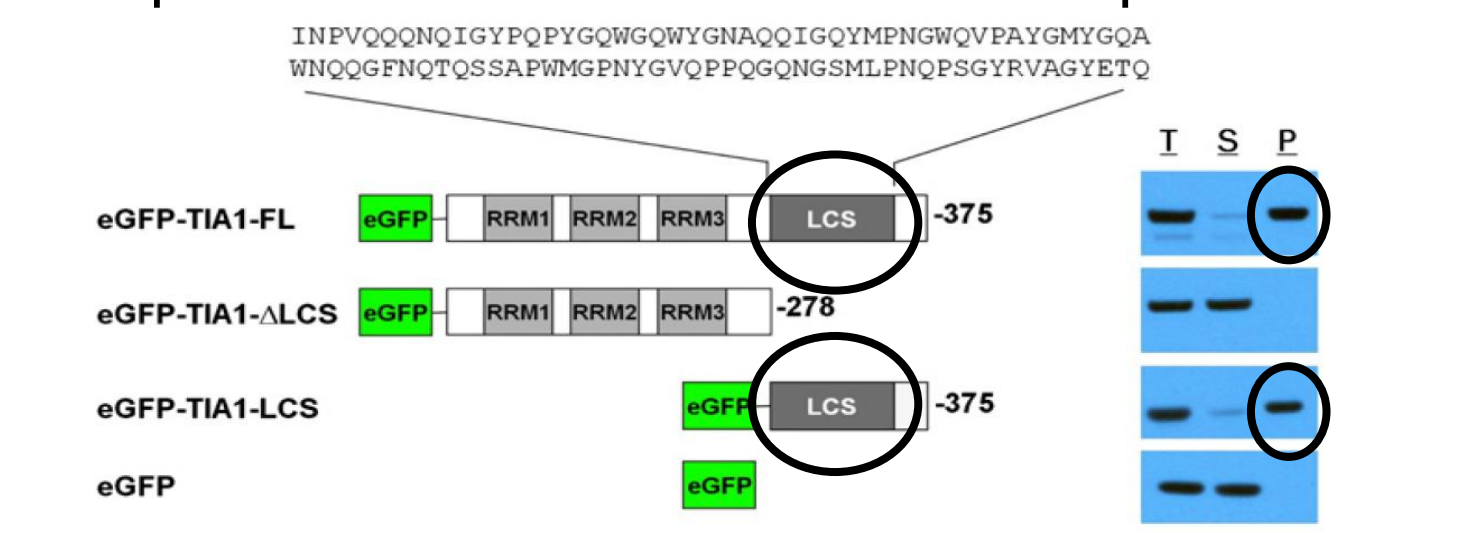


Figure 4: Different recombinant variants of TIA1-GFP localize in different fractions depending on the presence of b-isoX [4].

Punctual mutations in Low Complexity Domains promote an aberrant phase transition

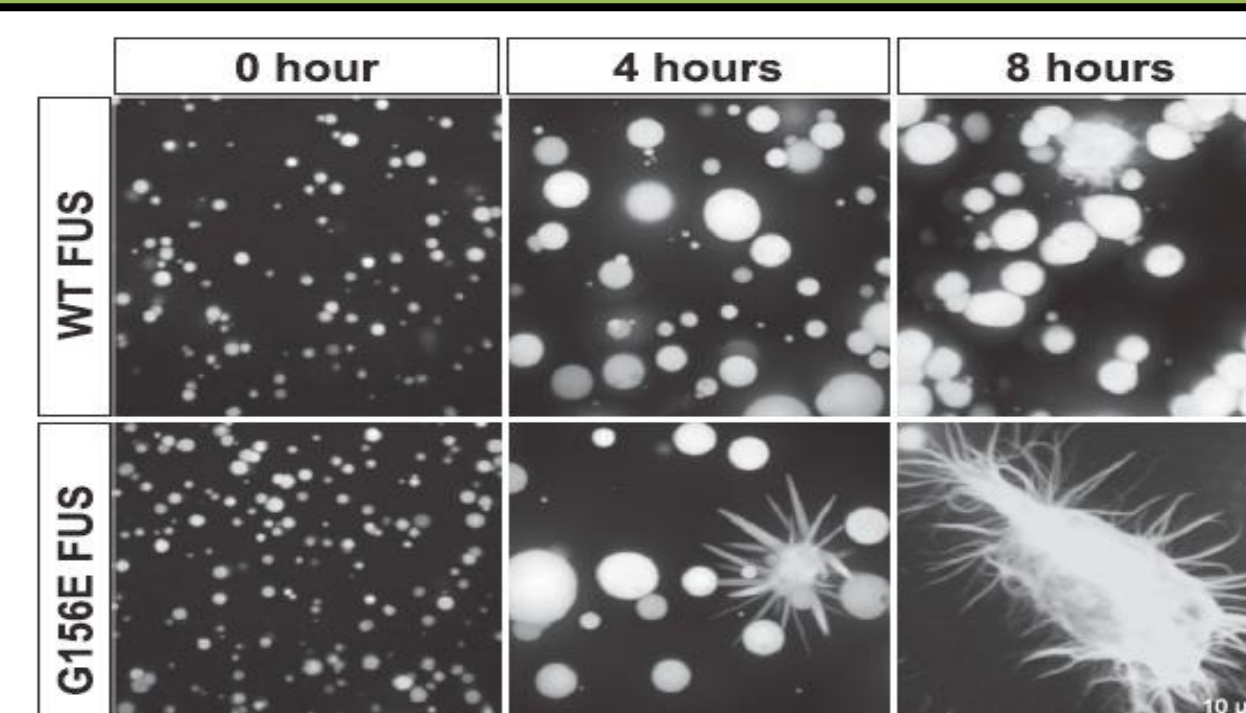


Figure 5: The aberrant phase transition is clear in G156E FUS sample at 8h [5].

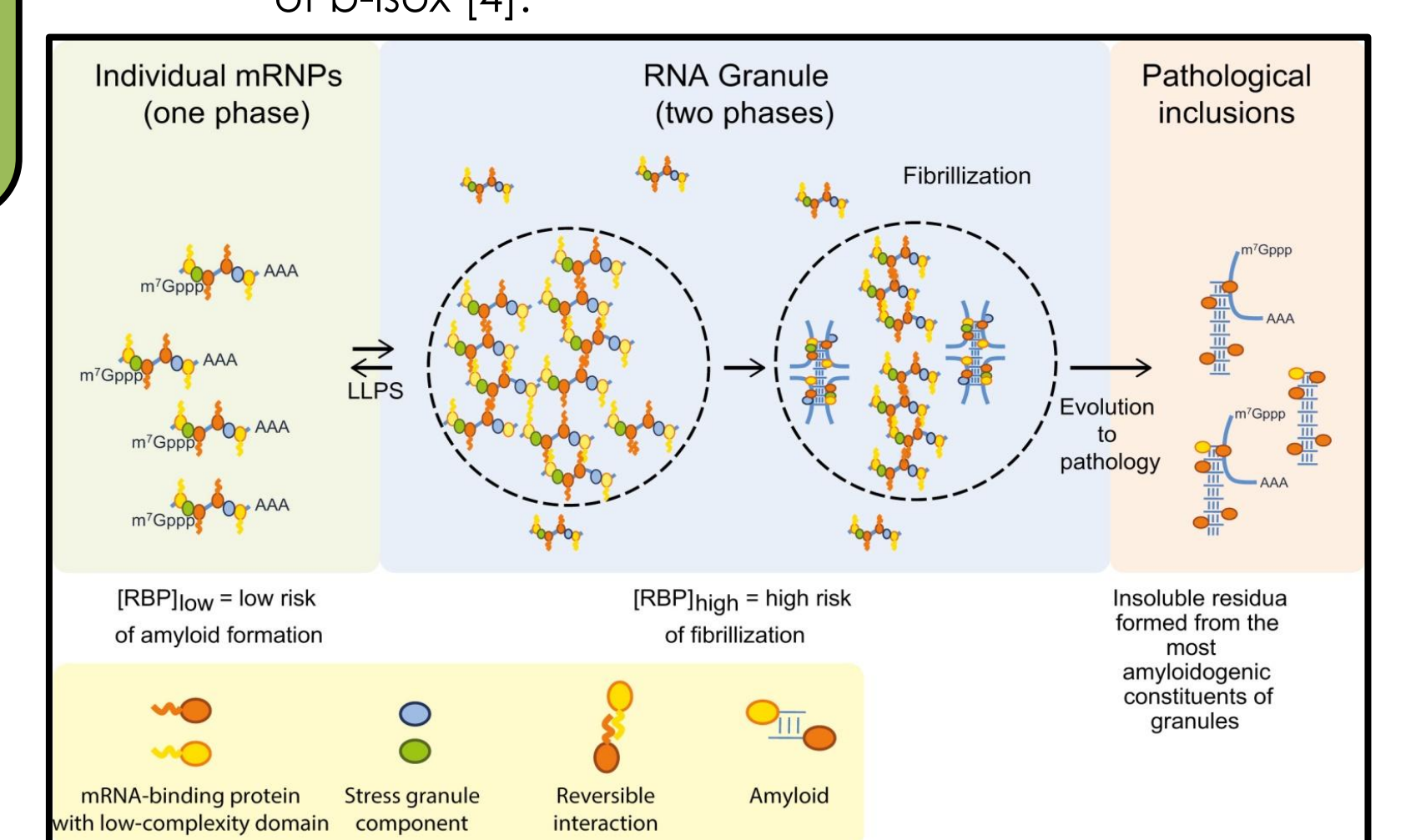


Figure 6: The alteration of liquid-like state equilibrium leads to pathological conformations [6].

NEUROPATHOLOGY OF PRION-LIKE PROTEINS

Mutations in Prion-like proteins are involved in the onset of **neurodegenerative disorders**.

- ❖ Punctual mutations in FUS protein are related with the onset of ALS

Failure in FUS-dependent DNA repair causes neurodegeneration

FUS-R521C mutant decreases dendrite growth

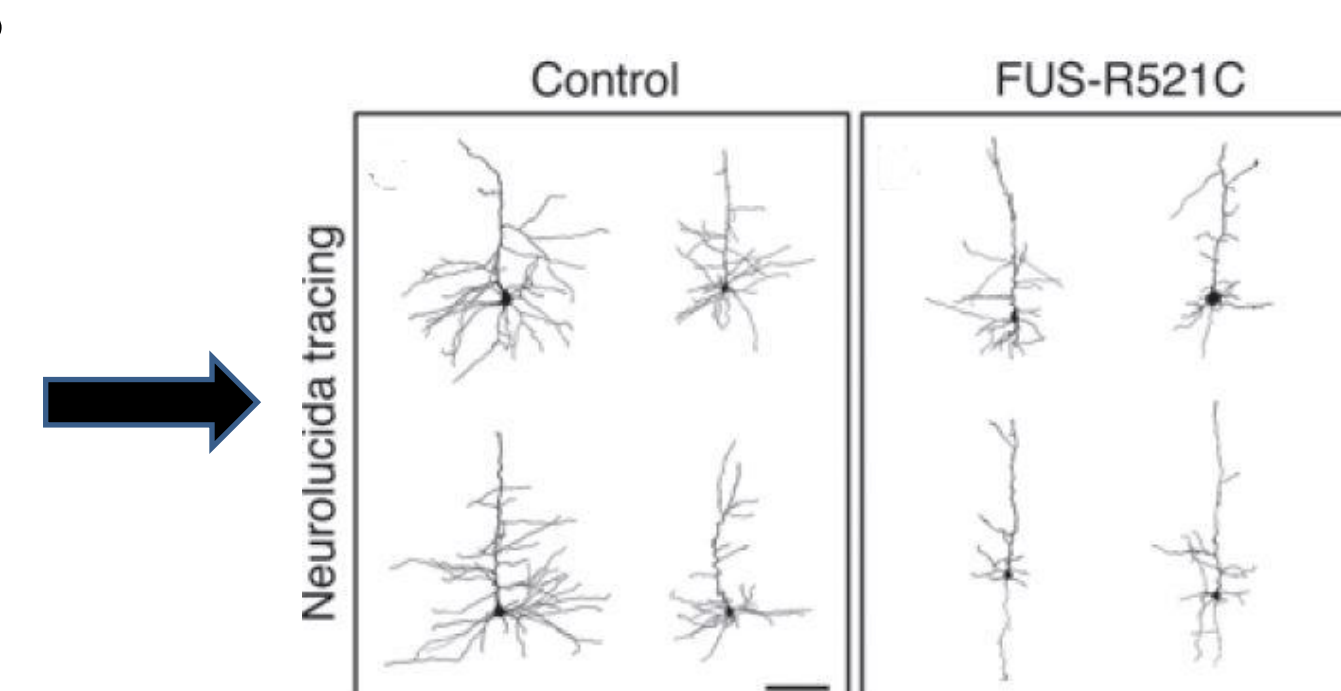


Figure 7: Effects of FUS-R521C in dendritic arborization are clear in cortical neurons [7].

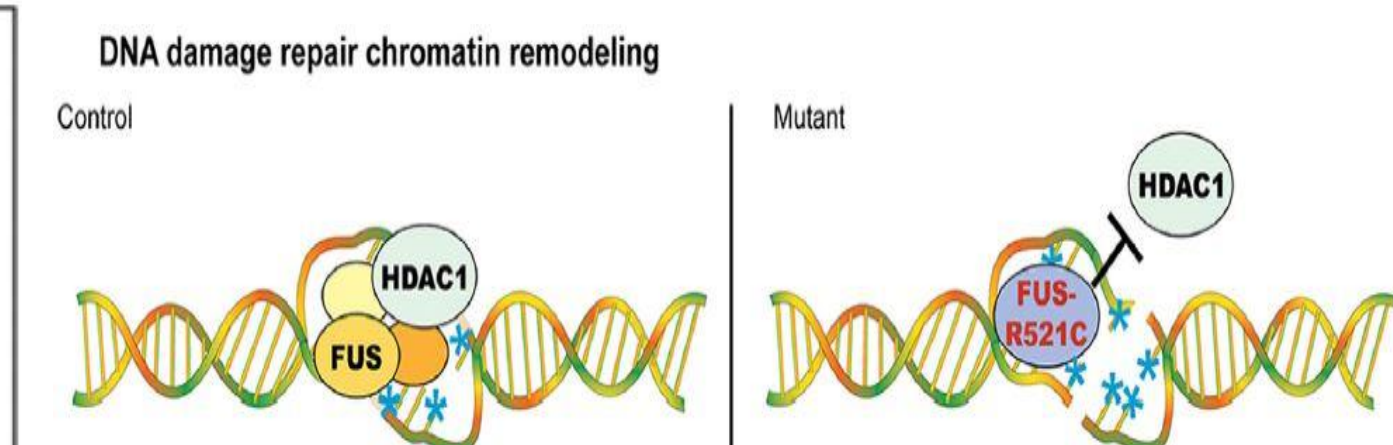


Figure 8: Mechanisms of FUS and HDAC1 in DNA repair [7].

CONCLUSION

In contrast to what is usually assumed, the neurotoxic mechanism behind the aberrant phase transition of prion-like proteins is **not caused by the intrinsic toxicity of the amyloid-like aggregates**, but rather results from the role of RNA-binding proteins in transcriptional and DNA repair processes. **Amyloid-like deposits inhibit the correct function of RNA-binding proteins**, decreasing cell fitness and viability.

This novel mechanism **will change the way we look at neurodegenerative disorders**, likely opening **novel therapeutic opportunities**.

REFERENCES

- Harit, F.U. et al., 2011. Molecular chaperones in protein folding and proteostasis. *Nature*, 475(7356), pp.324-332.
- Aulas, A. et al., 2015. Alterations in stress granule dynamics driven by TDP-43 and FUS: a link to pathological inclusions in ALS? *Frontiers in Cellular Neuroscience*, 9(October), pp.1-13.
- Lin, Y. et al., 2015. Formation and Maturation of Phase-Separated Liquid Droplets by RNA-Binding Proteins. *Molecular Cell*, 60(2), pp.208-219.
- Kato, M. et al., 2012. Cell-free formation of RNA granules: Low complexity sequence domains form dynamic fibers within hydrogels. *Cell*, 149(4), pp.753-767.
- Patel, A. et al., 2015. A Liquid-to-Solid Phase Transition of the ALS Protein FUS Accelerated by Disease Mutation. *Cell*, 162(5), pp.1066-1077.
- Mallix, A. et al., 2015. Phase Separation by Low Complexity Domains Promotes Stress Granule Assembly and Drives Pathological Fibrillization. *Cell*, 163(1), pp.123-133.
- Qiu, H. et al., 2014. ALS-associated mutation FUS-R521C causes DNA damage and RNA splicing defects. *The Journal of Clinical Investigation*, 124(3), pp.981-999.