

Lecture plan

1. Motor system

- Overview

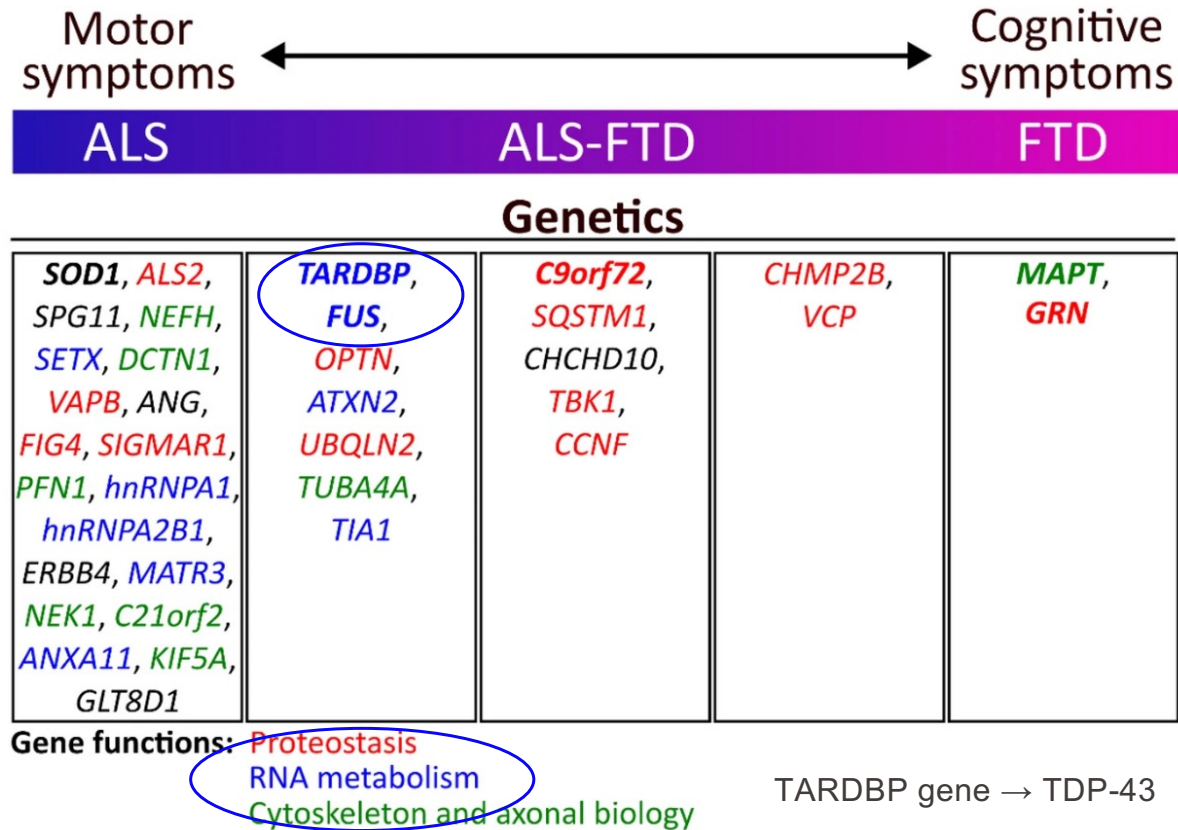
2. Motor Neuron Diseases

- Clinical presentation
- Molecular pathology

3. Amyotrophic Lateral Sclerosis

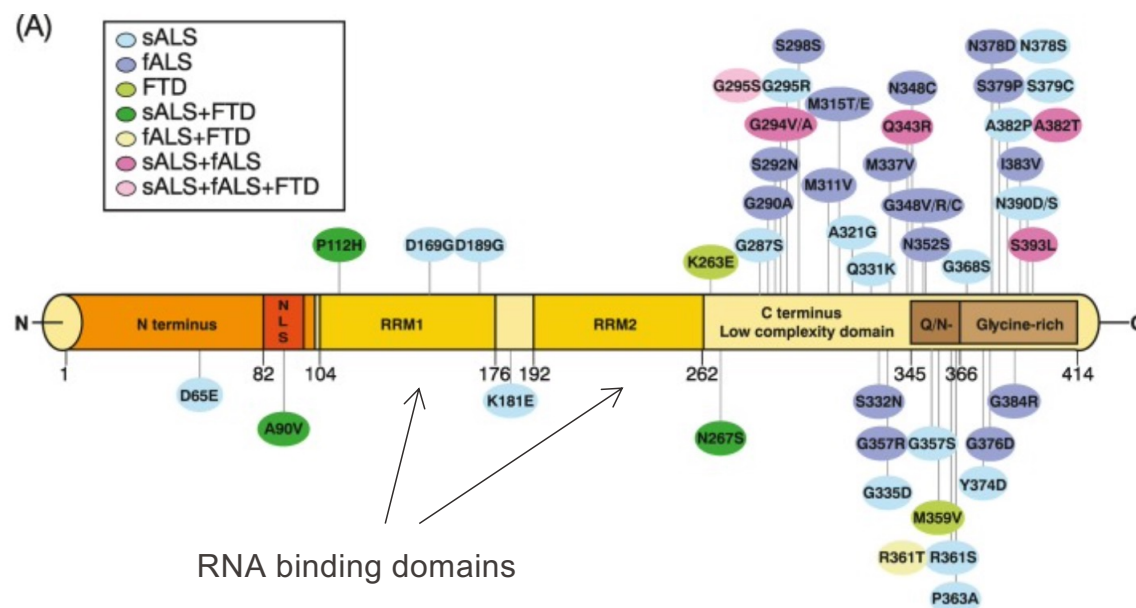
- Clinical presentation, epidemiology, etiology
- Non-cell autonomous mechanisms
- Molecular pathology: RNA metabolism

RNA metabolism is a prevalent mechanistic pathway in genes linked to ALS



TDP-43 (43 kDa trans-activating response DNA-binding protein)

- ≥ 38 mutations, <2% of ALS cases
- All but one mutation (Y374X) lead to autosomal dominant inheritance
- Classical ALS phenotype, some mutations associated with frontotemporal lobar degeneration (FTLD)
- Originally identified as a transcriptional repressor
- Two RNA recognition motifs (RRM)
- Normal nuclear distribution → **cytosolic inclusions**



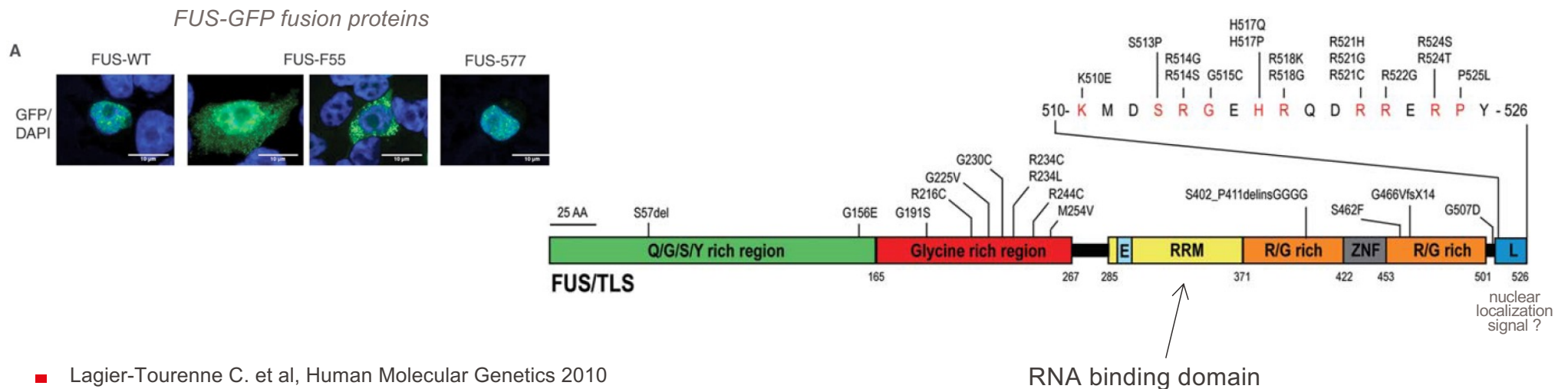
Bodanski A. et al, *Amyotrophic Lateral Sclerosis* 2010

Lagier-Tourenne C. et al, *Human Molecular Genetics* 2010

<https://doi.org/10.1016/j.tins.2021.02.008>

FUS/TLS (fused in sarcoma/translocated in liposarcoma)

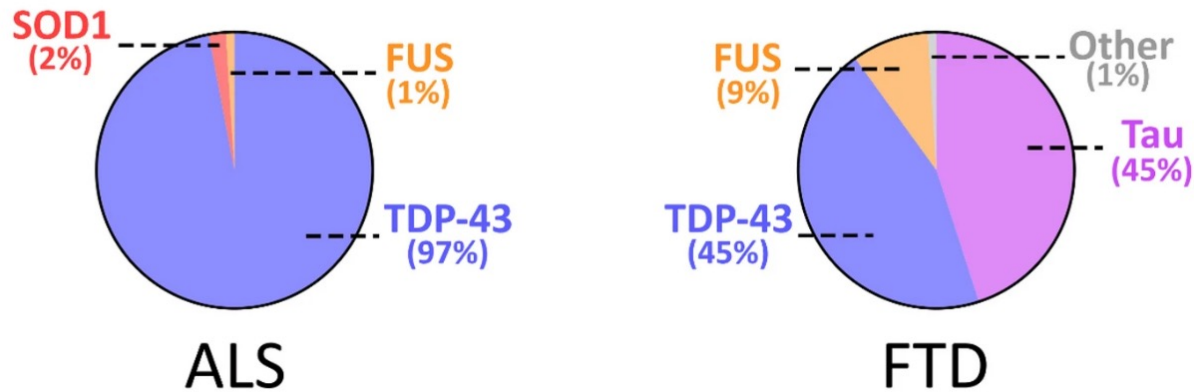
- ≥ 30 mutations, <2% of ALS cases
- Dominant inheritance except one recessive mutation (H517Q)
- Familial ALS, sometimes with early onset (<30 yrs) and rapid progression
- Some rare patients with FTLD
- One RNA recognition motif (RRM)
- Mainly nuclear protein → **cytosolic in stress/pathologic conditions**



- Lagier-Tourenne C. et al, Human Molecular Genetics 2010
- Kwiatkowski TJ Jr, et al Science 323(5918):1205-8, 2009

Pathology: TDP-43 and FUS are mislocalized and form aggregates in ALS

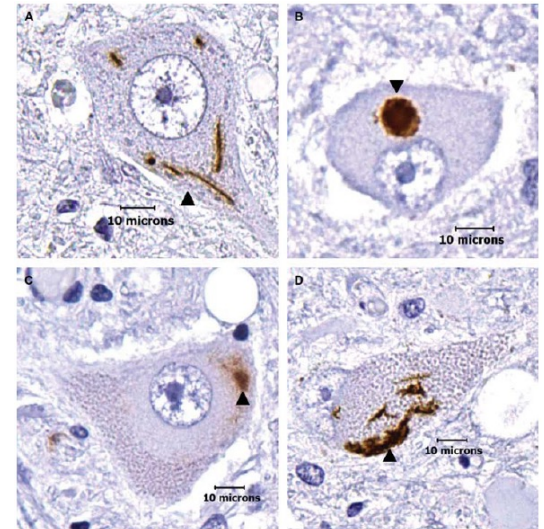
Pathological protein inclusions



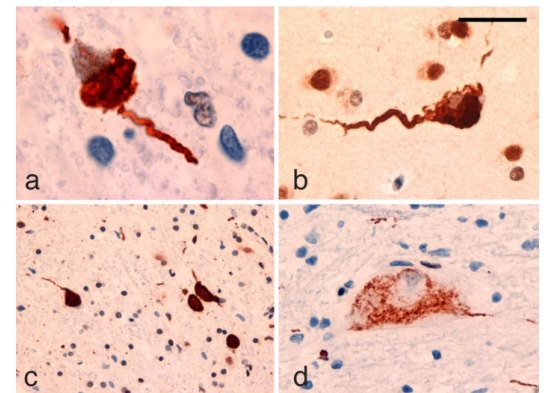
⇒ TDP-43 pathology is considered a common feature in ALS
⇒ Central mechanism?

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TDP-43 inclusions (in 97% of ALS cases)



FUS inclusions



EPFL TDP-43 pathology: mechanisms

TDP-43: loss of function or gain of a toxic activity ?

Gain of a toxic function

- Presence of TDP-43 aggregates
- Accumulation of the protein in the cytosol
- Autosomal dominant inheritance
- Experimental evidence for toxic aggregation activities

Loss-of-function

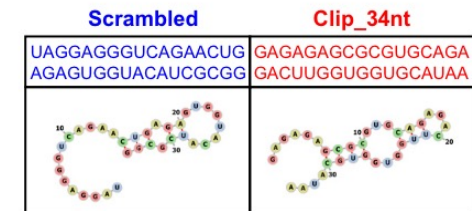
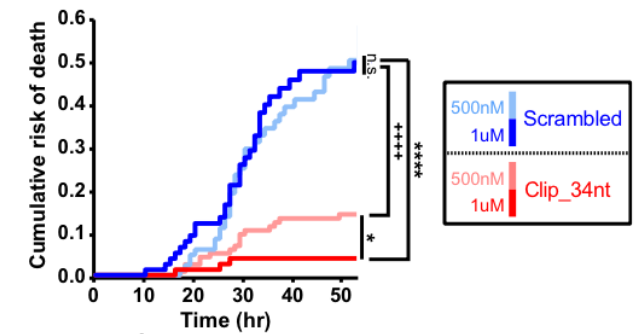
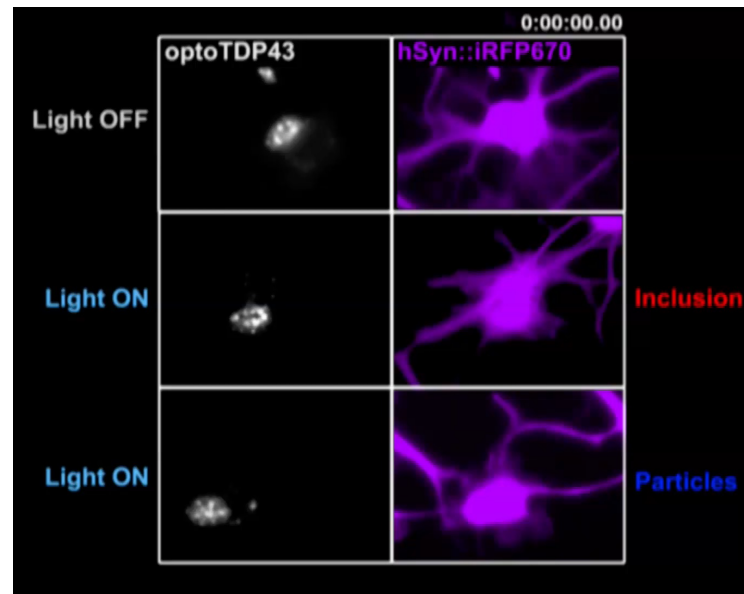
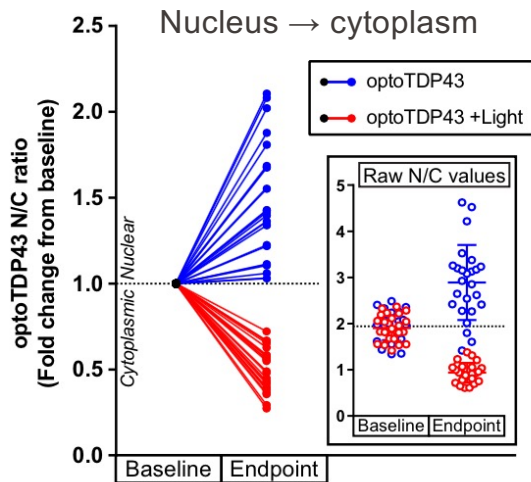
- Loss of nuclear TDP-43 (even partial loss has consequences)
- Knock-out of TDP-43 is embryonic lethal
- Very important function in RNA metabolism, mRNA transport, response to stress
- Experimental evidence for loss-of-function toxicity



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OptoTDP-43: light-inducible oligomerization

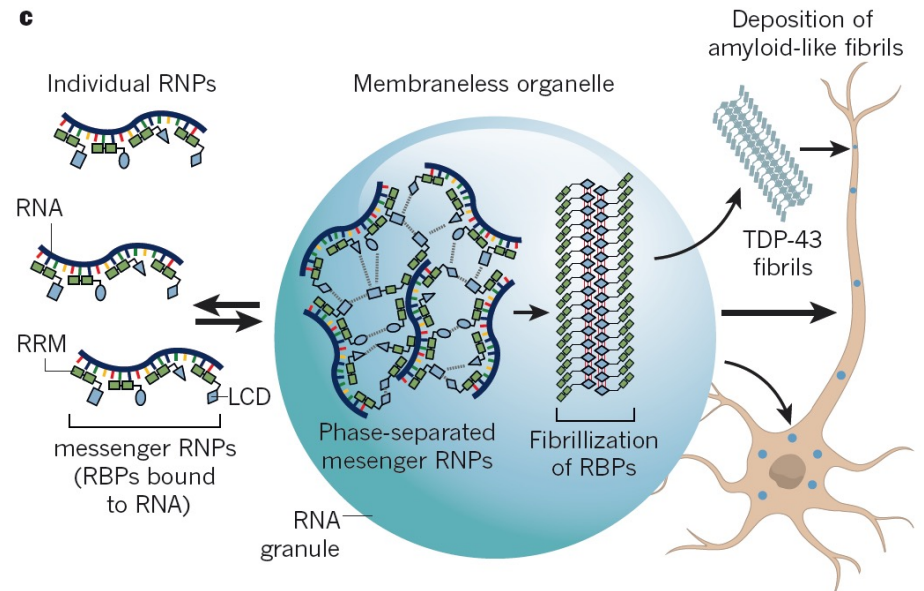
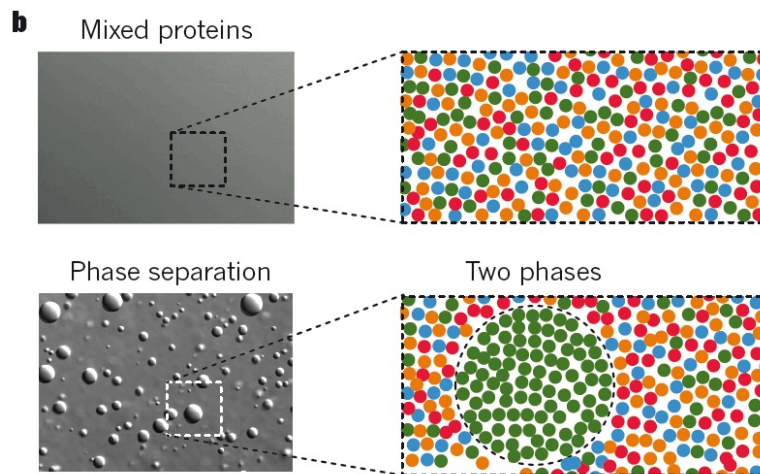
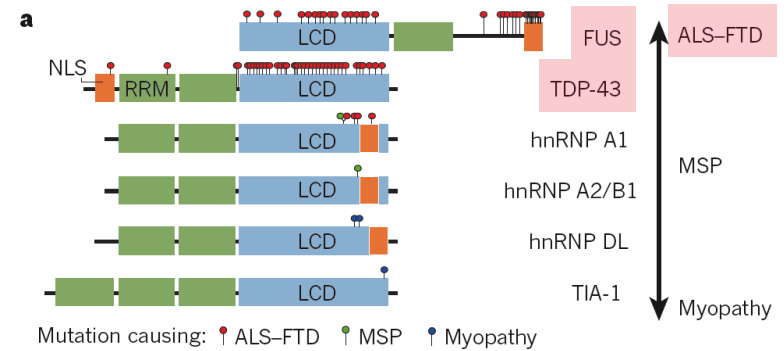
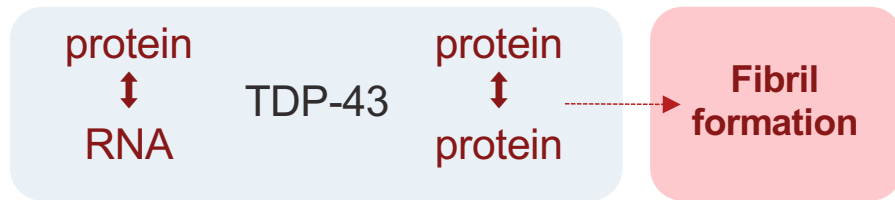
Oligomerization induces cytosolic localization of TDP-43 \Rightarrow neuronal toxicity opposed by RNA binding



- RNA binding antagonizes TDP-43 toxicity

EPFL TDP-43: evidence for the gain of toxic activity

A class of proteins with low-complexity domains (LCD) that form membrane-less organelles



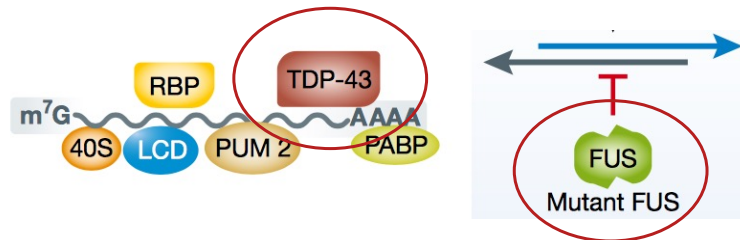
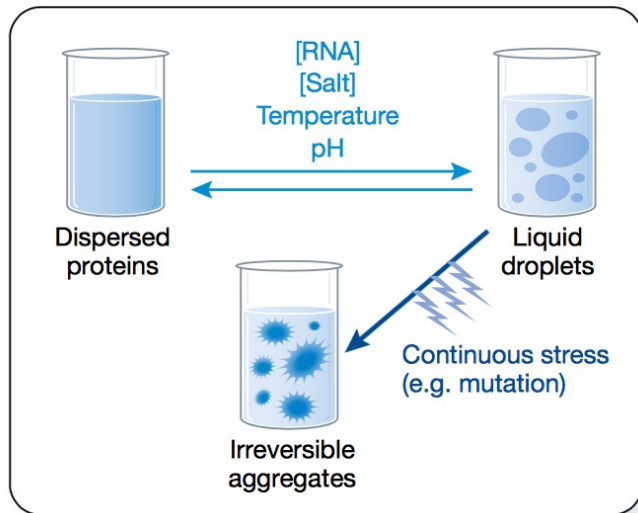
■ Taylor JP et al, Nature 2016

EPFL TDP-43: evidence for the gain of toxic activity

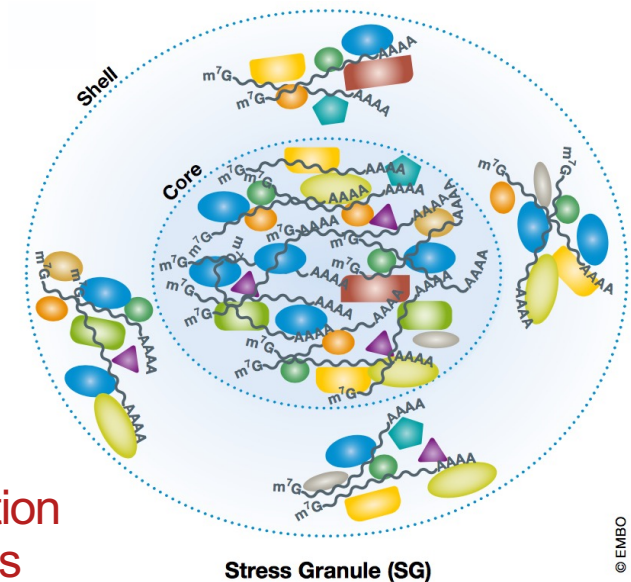
Stress granules: cytoplasmic hubs for mRNA triage and cellular stress adaptation

- RNP granules concentrating specific cellular components is a **conserved strategy** across multiple organisms and in different cellular compartments. They are formed in response to **stress** or drugs.
- Stress granules represent **assemblies of mRNPs stalled in translation initiation**.
- Stress granules are **dynamic structures**, controlled by protein chaperones, RNA helicases, and post-translational modifications.
- Mutations that alter stress granule formation contribute to some **neurodegenerative diseases** and cancers.

Liquid-Liquid Phase Separation

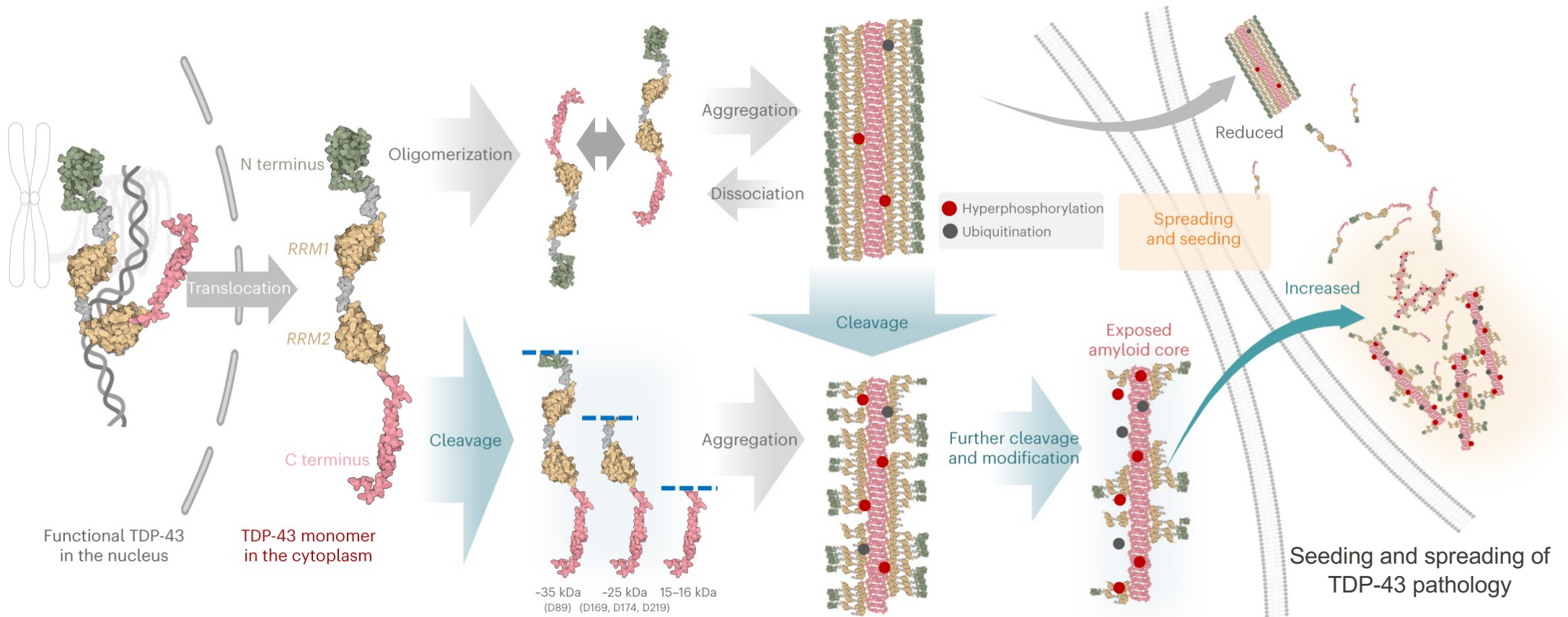


FUS and TDP43 regulate the formation and disassembly of stress granules



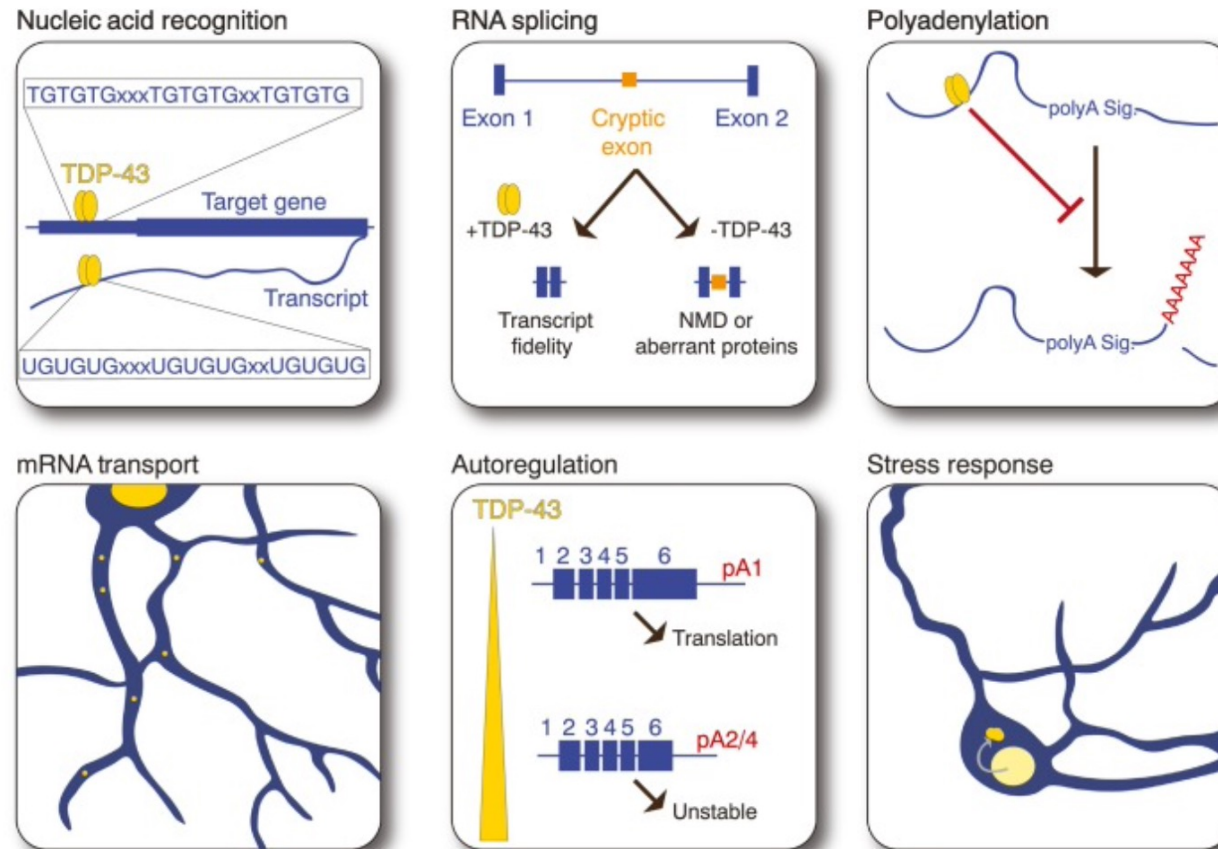
- Gao FB et al, EMBO J 2017
- Protter DSW et al, Trends in Cell Biol 2016

The path towards aggregation: mechanisms of TDP-43 filament formation



■ *Nature Neuroscience* volume 26, pages 983–996 (2023)

EPFL TDP-43: evidence for the loss of function toxicity

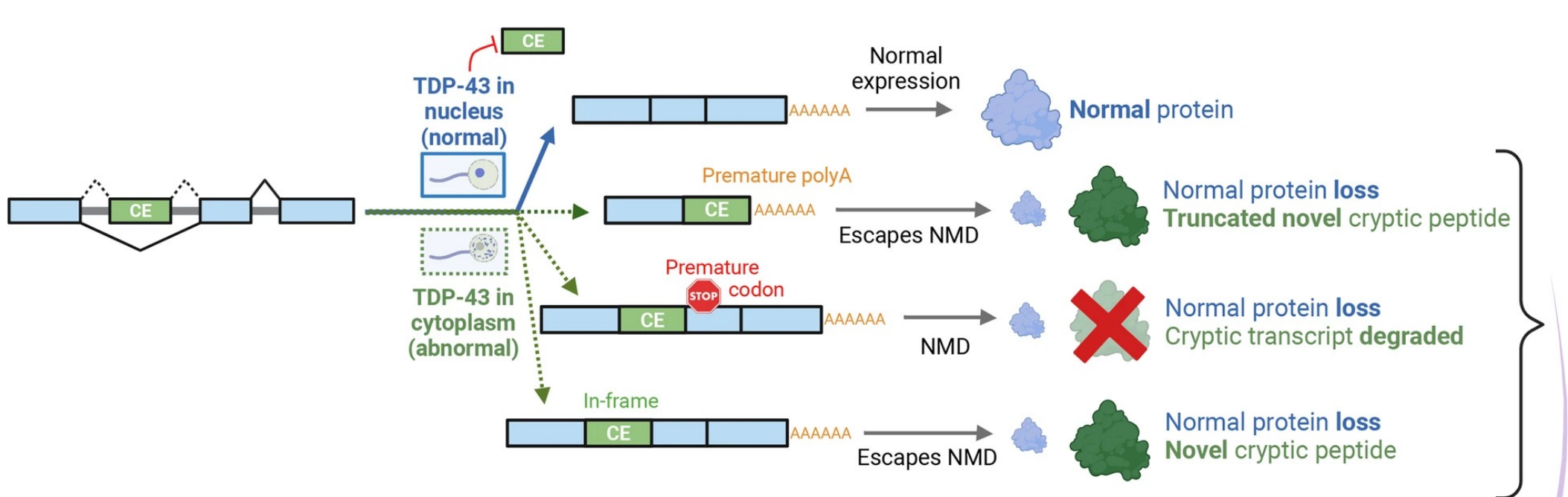


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■ <https://doi.org/10.1016/j.tins.2021.02.008>

EPFL TDP-43: evidence for the loss of function toxicity

TDP-43-mediated incorporation of cryptic exons: implications for ALS-FTD



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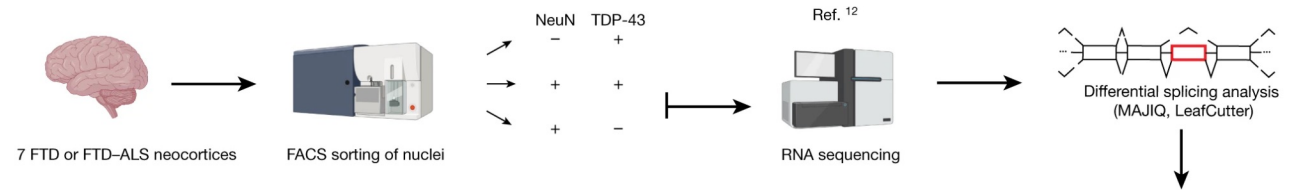
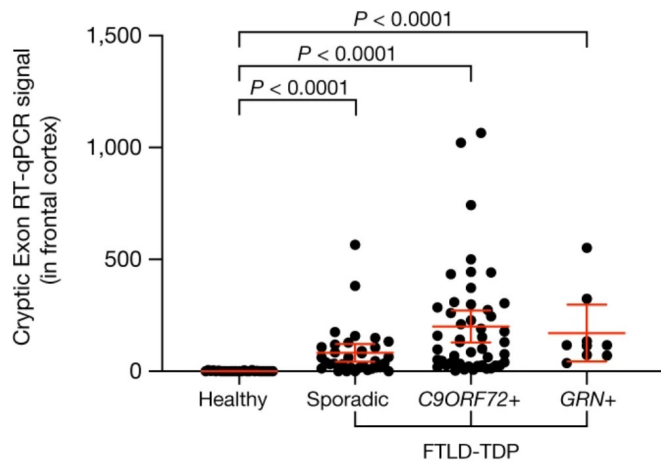
■ Mol Neurodegeneration 18, 16 (2023). <https://doi.org/10.1186/s13024-023-00608-5>

Role in disease?

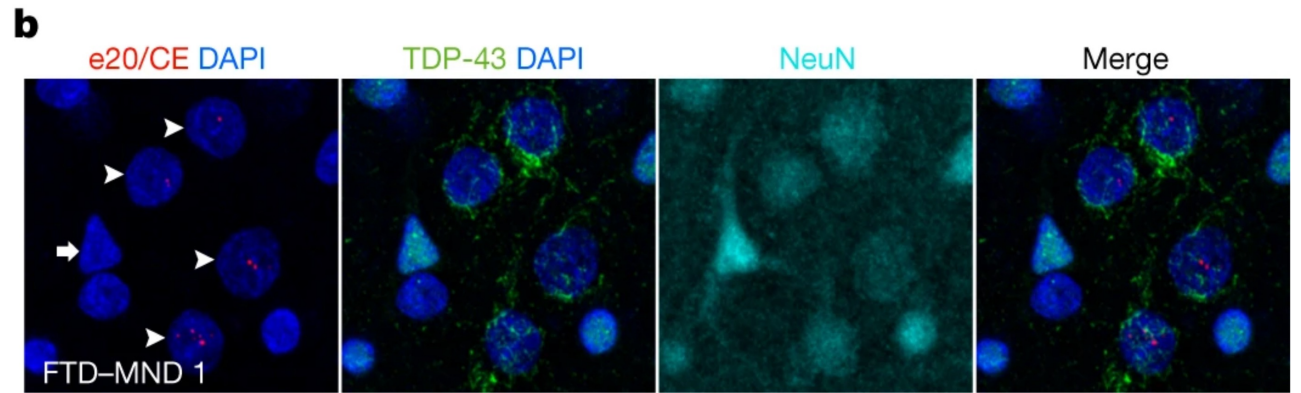
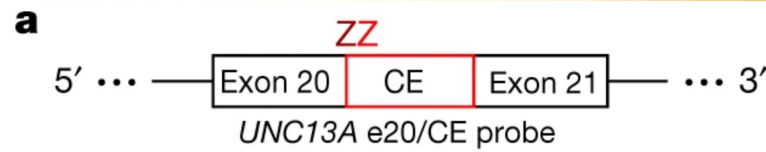
EPFL TDP-43: evidence for the loss of function toxicity

Loss of nuclear TDP-43 is associated with cryptic exon splicing in patients with FTD and MND

Cryptic exon in UNC13A mRNA



AKT3	C2CD5	CYFIP2	FAM193A	KALRN	MARK3	PLEKHA1	PTPRT	SEPT7P2	STXB5L	UNC79
ANK3	CACNA1E	CYP2C8	FKBP15	KCNQ2	MNAT1	PLEKHA5	RAP1GAP	SETD5	SYNE1	UQCRC2
ANKRD20A5P	CADPS	DLGAP4	HP1BP3	KIF3A	NCAM1	POLDIP3	RAPGEF4	SH3KBP1	SYT7	VWA8
APLP2	CAMK2B	EIF4G3	ICA1	KMT2C	NFIX	PPP6R3	RAPGEF6	SHISA9	TRAPPC12	WHSC1L1
ARHGEF11	CEP290	EPB41L1	IMMT	LOC100630923	PACRGL	PRUNE2	RNASET2	STMN2	TRRAP	ZNF138
ATP8A2	CNOT6	ERC2	IQCB1	MADD	PDE2A	PTPRD	SCN1A	STXB1	UNC13A	ZNF292



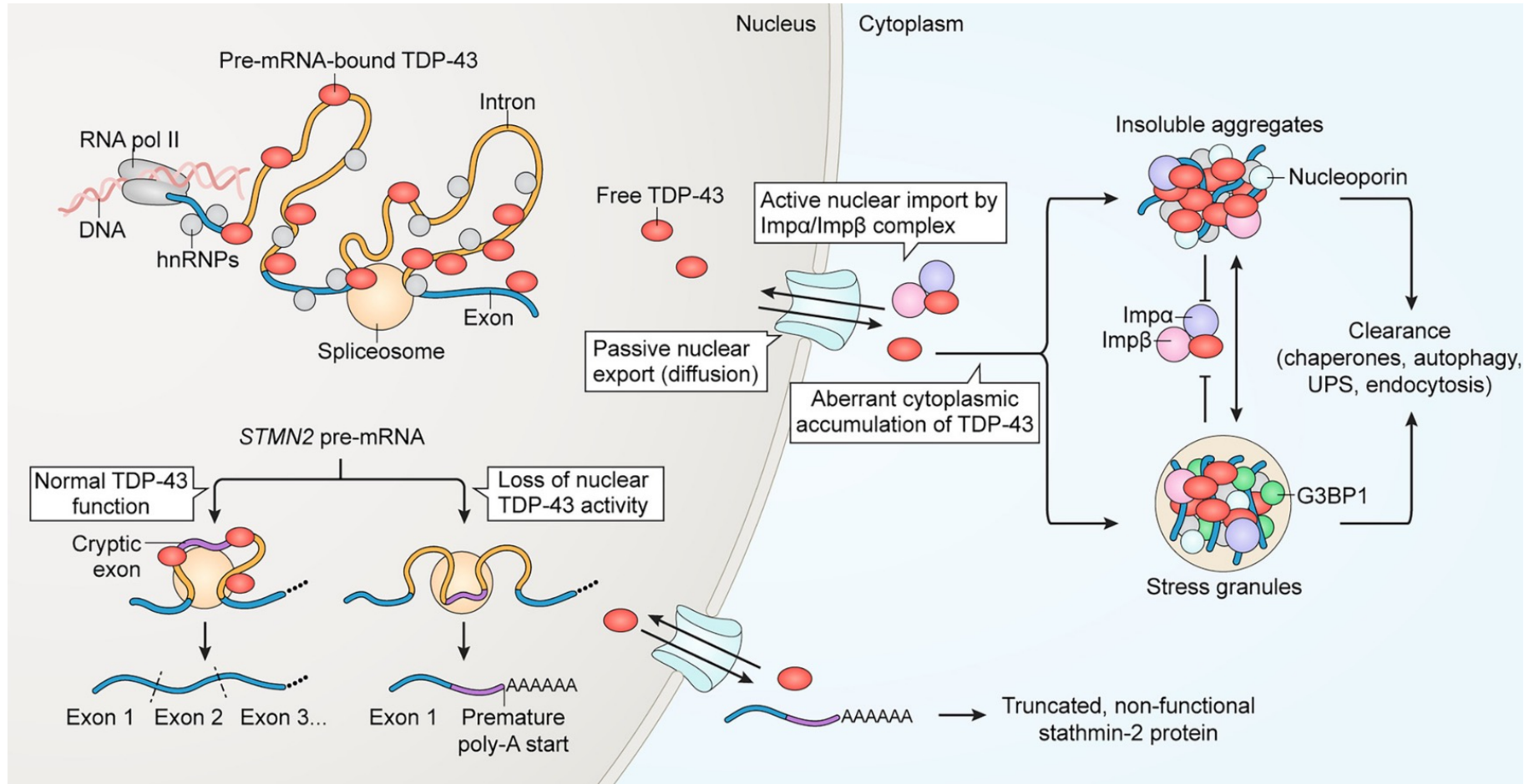
Splicing alterations in 100s → 1000s of genes related to TDP-43!

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Nature volume 603, pages 124–130 (2022)

EPFL TDP-43: evidence for the loss of function toxicity

Role of TDP-43 in RNA metabolism



Motor neuron diseases: question 10

TDP-43 biology in normal and diseased conditions shows that pathology affects the protein distribution across 4 different compartments:

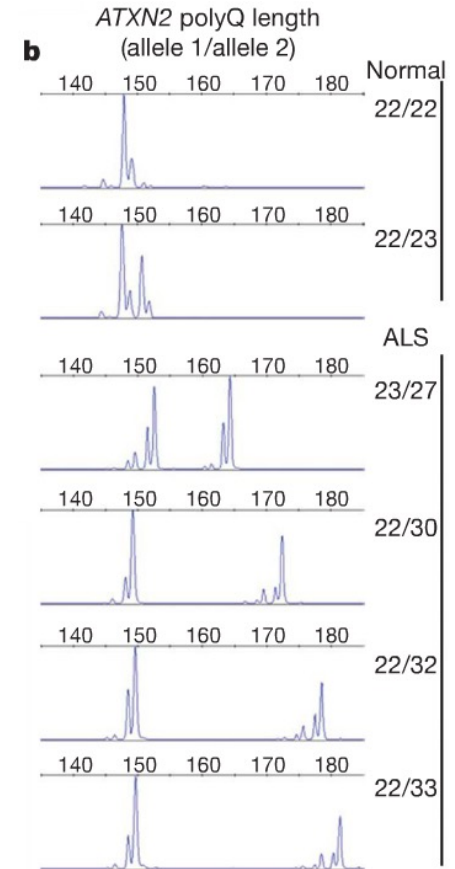
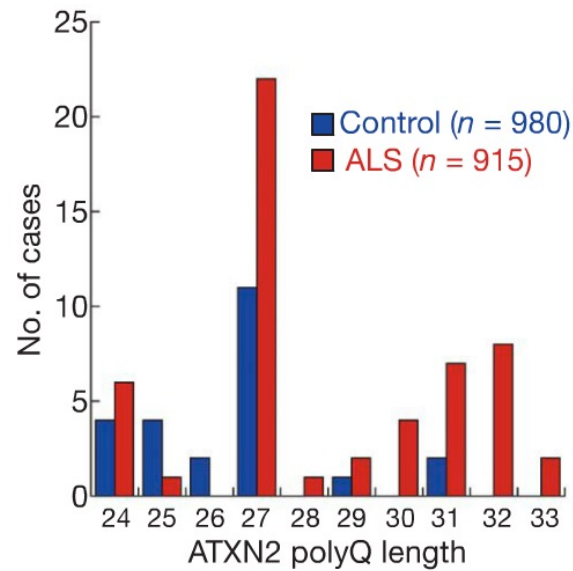


Based on these observations, would you think that neurodegeneration is caused by...

- A. the changes in the dynamic transfer across these compartments.
- B. the formation of protein aggregates, the main pathological hallmark.
- C. the loss of the effects of nuclear TDP-43 on RNA and gene expression.
- D. the combination of all of the mechanisms mentioned above.

Ataxin 2: CAG repeats as a risk factor for sporadic ALS

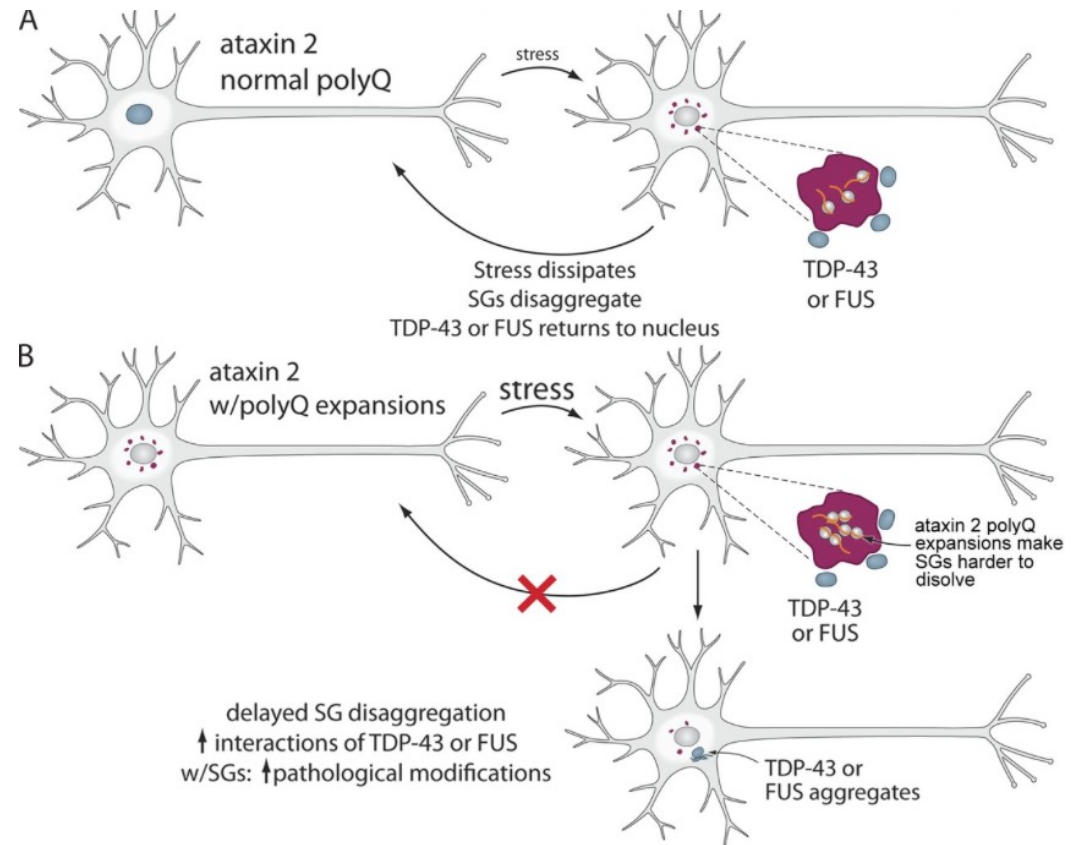
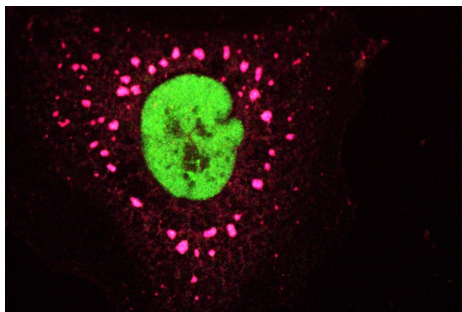
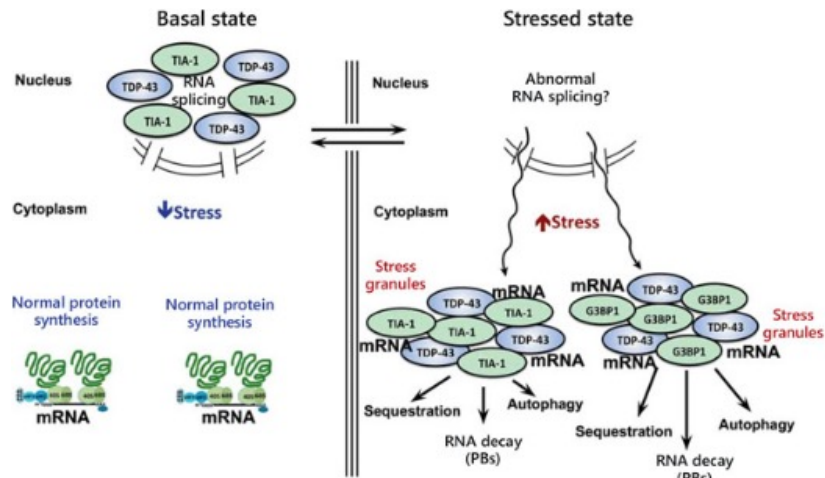
- >34 CAG repeats in ATXN2 → spinocerebellar ataxia type 2
- Ataxin 2 is part of a complex with TDP-43



PolyQ lengths ≥ 27 are significantly enriched in ALS vs. controls

▪ Elden AC, et al Nature 2010

A role for Ataxin 2 in stress granule dynamics



Motor neuron diseases: question 11

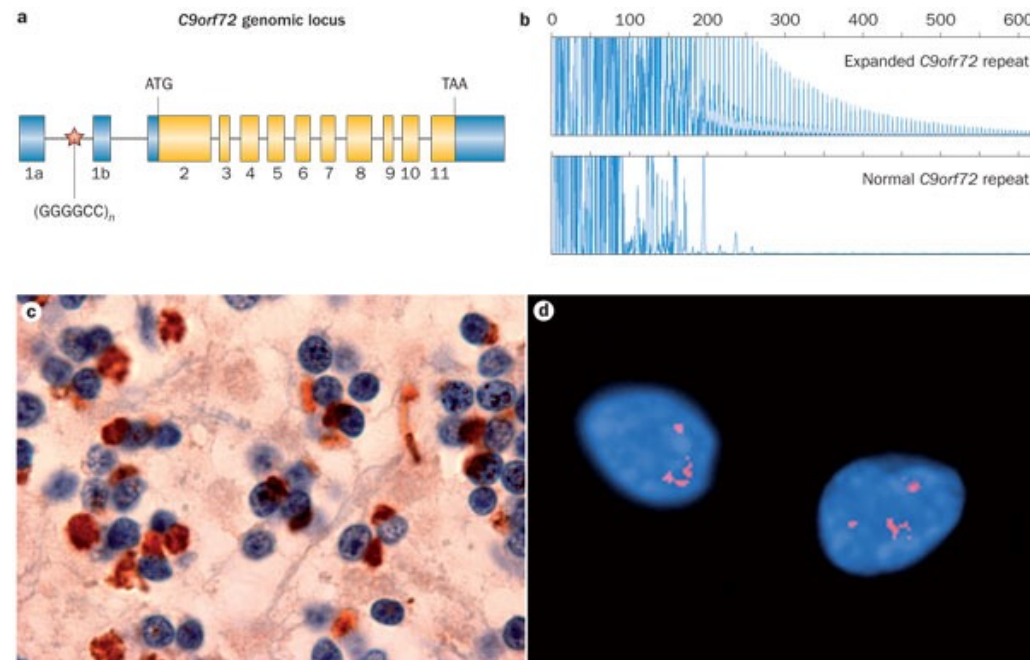
You want to start a research program to explore why motoneurons are particularly vulnerable to perturbations of RNA metabolism in ALS.

What would be your work hypothesis (one possibility)?

- A. Motoneurons have high gene expression levels in general, therefore they are sensitive to changes in RNA metabolism
- B. Splicing is important in motoneurons because they need a broader variety of proteins
- C. Expression of miRNA is more important in this cell type than in other cell types
- D. These cells are highly specialized and therefore the distribution of RNA in specific cell compartments is very important
- E. There is one specific RNA species which is perturbed and causes disease. It needs to be found.

C9orf72

- Non-coding hexanucleotide repeat expansion [GGGGCC]_n
- Normal allele: 2-25 **intronic** repeats (majority <8)
- Tens to thousands of repeats: association with **ALS/frontotemporal dementia**
- 5% of sALS, 20-40% of fALS (founder effects?)
- Possible implication in Huntington's and Alzheimer's disease



Ubiquitin-
positive
aggregates

RNA foci
(GGCCCC
probe)

- Rademakers R. et al., Nat Rev Neurol 2012

