Monomerization of TDP-43 in ALS

Literature Seminar #1 B4 Youhi Hwang 2024/12/5(Thu)

Introduction

- TDP-43 as a Leading Cause of ALS
- TDP-43 Dimerizes or Multimerizes in physiological conditions

Main

TDP-43 monomerization/multimerization status in CNS tissues of patients with ALS
TDP-43 monomerization induces cytoplasmic mislocalization
TDP-43 monomer aggregate inhibits normal TDP-43 function
Spliceosomal integrity is essential for TDP-43 dimerization/multimerization

TDP-43 monomerization precedes its pathological changes

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Amyotrophic Lateral Sclerosis (ALS)



From a website of Active Home Health & Hospice



Taylor, J., Brown, R. & Cleveland, D. *Nature* **539**, 197–206 (2016).



TDP-43 :ubiquitously expressed and predominantly nuclear RNA binding protein shuttling between the nucleus and the cytoplasm, which is encoded by the *TARDBP* gene

Pathological inclusions in ALS



TDP-43 redistribution to cytoplasmic inclusions (arrows) in spinal motor neurons in ALS



Taylor, J., Brown, R. & Cleveland, D Nature **539**, 197–206 (2016).

Ling, S. C. , et al. Neuron , 2013, 79, 417

TDP-43

Molecular mechanisms underlying TDP-43 pathogenesis



Tamaki, Y.; Urushitani, M. Int. J. Mol. Sci. 2022, 23, 12508.

•TDP-43 plays a critical role in multiple aspects of RNA metabolism, such as mRNA splicing.

 Pathological hallmarks in patients with ALS :cytoplasmic mislocalization, aggregation, hyperphosphorylation of TDP-43

 The exact molecular mechanism of TDP-43–mediated neurodegeneration remains unclear.



 Cleaved C-terminal fragments of TDP-43 accumulate in the brain and the spinal cord of patients with ALS.

- •Almost all ALS-linked mutations are located in the C-terminal domain(CTD).
- \rightarrow The CTD is a critical component of TDP-43 pathology in ALS.

• The functions of the N-terminus of TDP-43 remain largely unknown, but recent studies have highlighted the functional significance of **TDP-43 NTD**.

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Many proteins form dimeric/multimeric complexes



 Proteins in biological systems often bind other biomolecules.

Dimerization/multimerization
 can confer several different
 structural and functional
 advantages to proteins.

Kosuke Hashimoto et al 2011 Phys. Biol. 8 035007

Distribution of homooligomeric states in a non-redundant set of Protein Data Bank (PDB) structures.

The dimerization of TDP-43

The constitutive expression of an 86-kDa TDP-43 in human cell lines in culture



(exposed for 24 h to the vehicle (DMSO) (*lane 1*) or 1-μM MG-132 (*lane 2*)) The effects of transient expression of TDP-43 siRNA



The lanes (1−3) represent non-transfected cells, th∉2010). cells transfected with the vector of a scrambled sequence, and the cells transfected with the vector of TDP-43 siRNA, respectively.

•A discernible amount of the 86-kDa TDP-43-immunoreactive band is expressed in all the cells examined.

 Transient expression of the siRNA vector targeted specifically to TDP-43 substantially reduced the expression of both 43- and 86-kDa TDP-43.

 \rightarrow suggesting that the 86-kDa protein is composed of TDP-43

The dimerization of TDP-43

Coimmunoprecipitation with Flag-tagged proteins Fig. 4



Flag-tagged FL TDP-43 (*lanes 1* and 3) or GFP (*lanes 2* and 4) The lanes (1–4) represent (1 and 2) the immunoprecipitates and (3 and 4) the corresponding input controls.

The proteins binding to tagged TDP-43 always included the endogenous FL TDP-43 protein.

→suggesting that TDP-43 intrinsically forms the dimer.

Shiina, Y., *et al. Cell Mol. Neurobiol.* **30**, 641–652 (2010).

The dimerization of TDP-43

The domains involved in the intermolecular interaction of TDP-43 proteins Fig. 1





Western blot with anti-TDP-43 antibody (*lanes 1, 3, 5, 7, 9, 11, 13,* and *15*) or anti-Flag M2 antibody (*lanes 2, 4, 6, 8, 10, 12, 14,* and *16*). Extracted from HEK293 cells.

Shiina, Y., et al. Cell Mol. Neurobiol. 30, 641–652 (2010).

•N-terminal half domain is sufficient for the intermolecular interaction of TDP-43, possibly by constructing a conformationally ordered interacting domain.

• TDP-43 seems to natively dimerize or exist in a monomer-dimer equilibrium.

The multimerization of TDP-43





Fig.1



Immunoblots of human fibroblast fractions *C : cytoplasm N : nuclear

Afroz, et al. Nat. Commun. 8, 45 (2017).

- TDP-43 forms physiological multimers in human tissues.
- Multimeric TDP-43 species are detected predominantly in the nucleus.

TDP-43 multimerization impedes pathologic aggregation

Fig. 6



Crystals of TDP-43 N-terminal domain(NTD) showing helical filaments with single molecules arranged in headto-tail fashion

Tripartite GFP complementation assay(in mouse C17.2 cells) Afroz, et al. Nat. Commun. 8, 45 (2017).

•NTD-mediated TDP-43 multimerization impedes inter-molecular LCD interactions by distantly locating them from each other and antagonizes pathologic protein aggregation.

TDP-43 multimerization is crucial for RNA splicing regulation



• Downregulation of TDP-43 resulted in missplicing of specific RNA targets, which was rescued by co-transfection with wild type TDP-43.

• The multimerization-deficient mutants failed to rescue RNA missplicing.

 \rightarrow NTD-driven TDP-43 multimerization is essential for its function in RNA splicing.

Role of TDP-43 multimerization in physiological conditions & in disease



• Within the nucleus, TDP-43 multimerizes via the NTD.

• TDP-43 multimerization prevents irreversible aggregation by spatially separating the LCDs.

• Disturbance in the equilibrium between the oligomeric and monomeric TDP-43 in the cytoplasm may result in proteolytic cleavage of exposed monomeric TDP-43 and generation of C-terminal fragments lacking the NTD, which may initiate formation of irreversible pathologic aggregates via the LCD.

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TDP-43 dimerization/multimerization in patients with ALS



Representative immunoblots (IB) of DSG–cross-linked postmortem brain samples & Quantification of the dimer/monomer ratio of TDP-43



Representative immunoblots (IB) of sarkosyl-insoluble samples from postmortem spinal cord tissues & correlation dot plots of the dimer/monomer ratio (relative to the mean level of the control samples) of TDP-43 and relative levels of sarkosyl-insoluble TDP-43



Representative immunoblots (IB) of DSG–cross-linked samples from postmortem spinal cord tissues & Quantification of the dimer/monomer ratio of TDP-43

• The dimer/monomer ratio of TDP-43 was reduced in the affected brain & spinal cord tissues of ALS cases.

• The TDP-43 dimer/monomer ratio was negatively correlated with the insoluble TDP-43 levels in the spinal cord sample

 \rightarrow TDP-43 dimerization/multimerization is impaired in the CNS tissue of patients with ALS

 \rightarrow TDP-43 dimerization/multimerization is more impaired in lesions with more severe TDP-43 pathology.

Kotaro Oiwa *et al. , Sci.Adv*.**9**,eadf6895(2023).

TDP-43 monomer constitutes pathological inclusions in ALS motor neurons



(A) Three-dimensional models of the monomeric and dimeric NTD of TDP-43 as predicted by AlphaFold2. The critical amino acid of the anti–TDP-43 monoclonal antibody (E2G6G) epitope, Leu⁴¹, is indicated.



E2G6G strongly stained the pathological aggregates of TDP-43 in the motor neurons from ALS cases.
 →TDP-43 monomer constitutes the pathological inclusions in ALS motor neurons.

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Monomerization of TDP-43 induces its cytoplasmic mislocalization



(E) Experimental protocol
(F) representative images of human iPSC– derived motor neurons infected with lentivirus expressing HB9^{e438}::TDP-43^{WT}- or HB9^{e438}::TDP-43^{6M}-mCherry-IRES-Venus.
(G) Box and whisker plots of the cytoplasmic/nuclear ratios of TDP-43-mCherry fluorescence quantified from (F)

> Kotaro Oiwa *et al.*, *Sci.Adv.***9**,eadf6895(2023).

 The mislocalization of the NDD TDP-43 mutant into the cytoplasm was confirmed in human induced pluripotent stem cell (iPSC)-derived motor neurons
 →Indicating that the monomerization of TDP-43 promotes its cytoplasmic mislocalization in human motor neurons.

Nxf1 mediates cytoplasmic transport of TDP-43 monomers

Α



(A) Neuro2a cells transiently expressing TDP-43^{6M}-mCherry were treated with control siRNA or siRNAs targeting Xpo1 or Nxf1, followed by nucleocytoplasmic fractionation. GAPDH and fibrillarin were used as the cytoplasmic and nuclear markers, respectively.

(B) Relative cytoplasmic/nuclear ratios of TDP-43-mCherry in (A)

*C : cytoplasm N : nuclear

Xpo1: exportin-1 Nxf1: nuclear RNA export factor-1

- The suppression of Nxf1 substantially prevented the mislocalization of TDP-43^{6M}.
- The affinity of TDP-43^{6M} for Nxf1 was higher than that of TDP-43^{WT}.

→suggesting that Nxf1 recognizes TDP-43 monomers rather than dimers and mediates their nuclear export.



(E) TDP-43-3 × FLAG WT or 6M mutant was transiently transfected in the Neuro2a cells. An anti-FLAG antibody was used for immunoprecipitation of the lysates. Bound endogenous Nxf1 was detected by immunoblotting with an anti-Nxf1 antibody.

(F) Relative Nxf1 levels bound to TDP-43-3 \times FLAG

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TDP-43 monomerization induces insoluble aggregates



(E) Representative immunoblots of Neuro2a cells transiently expressing TDP-43-3 × FLAG WT and mutants. Proteins sequentially extracted with tris-HCl (TS), 1% Triton X-100 (TX), and 1% sarkosyl (Sar), as well as a sarkosyl-insoluble fraction (ppt), were analyzed by immunoblotting. (F) Quantification of the percentage of each fraction relative to the total amount of TDP-43-3 × FLAG in (E).



(G) Representative immunoblots of the sequentially extracted lysates from Neuro2a cells transiently expressing TDP-43^{6M}-3×FLAG full-length (FL) and deletion mutants. (H) Quantification of the percentage of each fraction to the total amount of TDP-43-3×FLAG in (G).

- All NDD-TDP-43 variants except TDP-43^{ΔNLS} were significantly enriched in the sarkosyl-insoluble fraction than TDP-43^{WT}.
- TDP-43^{6M} revealed that the CTD was the key domain for aggregation.

Ε

 \rightarrow indicating that NDD-TDP-43 forms insoluble aggregates through CTD.

TDP-43 monomer inhibits the physiological functions of endogenous TDP-43



(E) Exon skipping assay for *CFTR* exon 9 (top). Quantification of the exclusion/inclusion ratio of *CFTR* exon 9 (relative to mock-transfected cells) was performed from the band intensities (bottom). (F & G) Quantification of RNA levels of *Tardbp* intron 7 (F) & *Sort1* (G) with indicated exon inclusion. Neuro2a cells were transiently transfected with TDP-43, control siRNAs, or siRNAs targeting TDP-43. (I) Representative immunoblots showing endogenous TDP-43 and mCherry levels in cytoplasmic (Cyto), nuclear (Nuc), and 1% Triton X-100 insoluble fractions (Insol) of Neuro2a cells transiently expressing mCherry or TDP-43^{6M}-mCherry. (J) Quantification of the percentage of endogenous TDP-43 in each fraction to the total endogenous TDP-43 in (I).

- TDP-43^{6M} lost its ability to skip exon 9 of the *CFTR* gene and to regulate the splicing of *Tardbp, Sort1, Poldip3*.
- TDP-43^{6M} showed a phenotype similar to that obtained after siRNA-mediated TDP-43 knockdown.
- The overexpression of TDP-43^{6M} reduced the amount of endogenous TDP-43 in the nucleus and induced its accumulation in the insoluble fraction.

→suggesting that monomeric TDP-43 aggregates induce cytotoxicity by losing the function as splicing regulator & sequestrating the endogenous TDP-43 into the aggregates.

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Spliceosomal integrity is essential for TDP-43 dimerization/multimerization



(A) Schematic illustration of TDP-DiLuc, a NanoBiT-based complementation assay to monitor the TDP-43 dimerization/multimerization. (C) Quantification of the TDP-43 dimerization under stress conditions.



(A) Representative confocal images of HeLa cells transiently expressing TDP-43-3 × FLAG WT or 6M stained with anti-SMN and anti-FLAG antibodies, along with line scan analysis for colocalization of TDP-43 and SMN. Scale bar, 10 μ m. (B) The percentage of the TDP-43-positive Gems per total Gems was quantified in 100 Gems from three independent experiments.

 Dimerization/multimerization of TDP-43 was impaired to different degrees by all of the examined stresses.

TDP-43^{6M} lost its localization to the Gems.

Hypothesis : Abnormalities in the Gems under stress conditions might be responsible for TDP-43 monomerization.

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Spliceosomal integrity is essential for TDP-43 dimerization/multimerization



(**D**) Representative confocal images of HeLa cells treated with the indicated reagents followed by staining with anti-SMN. Arrowheads indicate SMN-positive Gems. (**E**) Quantification of Gems per cell in (D). (**F**) Representative confocal images of SH-SY5Y cells transiently transfected with control siRNA or siRNA targeting SNRNP70. Cells treated with ActD (5 μ g/ml; 3 hours) served as the positive control. Arrowheads indicate SMN-positive Gems. (**G**) Quantification of Gems per cell in (F). (**H**) Quantification of the relative DiLuc luminescence in SH-SY5Y cells transiently transfected with TDP-DiLuc and control siRNA or siRNA targeting SNRNP70. Cells treated with ActD (5 μ g/ml; 3 hours) served as the positive control.

Kotaro Oiwa et al. , Sci.Adv.9, eadf6895(2023).

•All stresses decreased the number of Gems represented.

• Disrupting the spliceosome by the siRNA-mediated knockdown of SNRNP70 decreased the number of Gems and TDP-43 dimerization/multimerization.

→suggesting that loss of spliceosome integrity by some stress, such as transcriptional inhibition, reduces Gems, which damages TDP43 dimerization/multimerization.

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в

D

F

Cytosol/nuclear

Insoluble/soluble

012

012 4

Time (hours

Time (hours)

Time (hours)

(A) Experimental protocol of time course analysis of Neuro2a cells treated with ActD (1 μ g/ml) for the indicated time. (B) DiLuc luminescence of the Neuro2a cells stably expressing TDP-DiLuc treated with ActD $(1 \mu g/ml)$ for the indicated time (relative to mean levels at time zero). Triplicate samples were analyzed in three biologically independent experiments. (D) Time course analysis of the cytoplasmic/nuclear ratios of mCherry-TDP-43^{WT} fluorescence quantified in 50 cells from (C) from two biologically independent experiments (F) Time course analysis of the insoluble/soluble TDP-43 ratios.

Kotaro Oiwa et al., Sci.Adv.9, eadf6895(2023).

 TDP-43 monomerization preceded TDP-43 pathological changes, such as cytoplasmic mislocalization & reduced solubility

 \rightarrow suggesting that the monomerization of TDP-43 NTD precedes the loss of TDP-43 from the nucleus and the formation of aggergates

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Summary



Kotaro Oiwa et al. , Sci.Adv.9, eadf6895(2023).

OTDP-43 exists in a monomer-dimer/multimer equilibrium in cells.

OThe NTD of TDP-43 mediates the formation of physiological homodimers, which are essential for its physiological splicing function. OThe RNA export factor Nxf1 facilitates the nuclear export of TDP-43 monomers.

× Various types of stresses impair TDP-43 dimerization/multimerization and increase amounts of TDP-43 monomers, likely by inducing spliceosomal defects

Monomeric TDP-43 lost its ability to regulate mRNA splicing.

× Excess of TDP-43 monomers mislocalized in the cytoplasm aggregates with endogenous TDP-43.

Perspectives for diagnosis & treatment

• Focusing on the functional dimerization/multimerization of TDP-43 may be important to develop early diagnostic markers of TDP-43 proteinopathy including ALS.

- Keeping Spliceosomal Integrity(Further research is required)
- Decomposing TDP-43 monomers or aggregates in cytoplasm
- Reducing the ability of TDP-43 to aggregate
- Returning TDP-43 monomers from cytoplasm into nuclear
- Reducing Nxf1 activity of binding to TDP-43 monomers and exporting them from nuclear into cytoplasm