

## **Supplementary Materials**

### **DCTN1 binds to TDP-43 and regulates TDP-43 aggregation**

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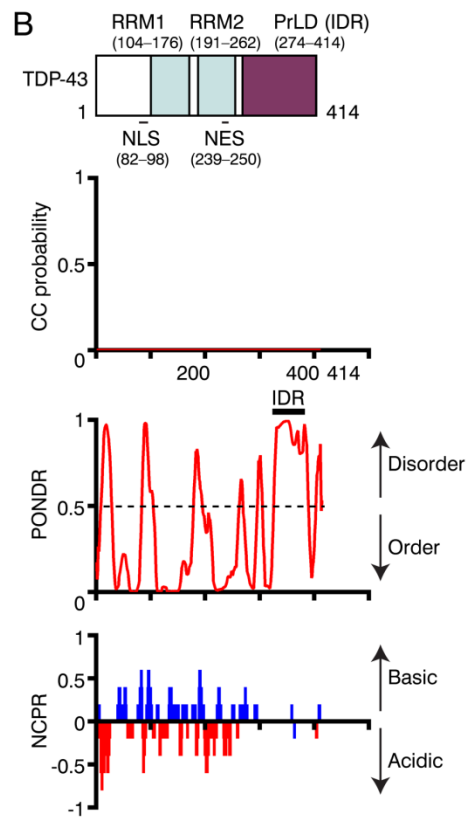
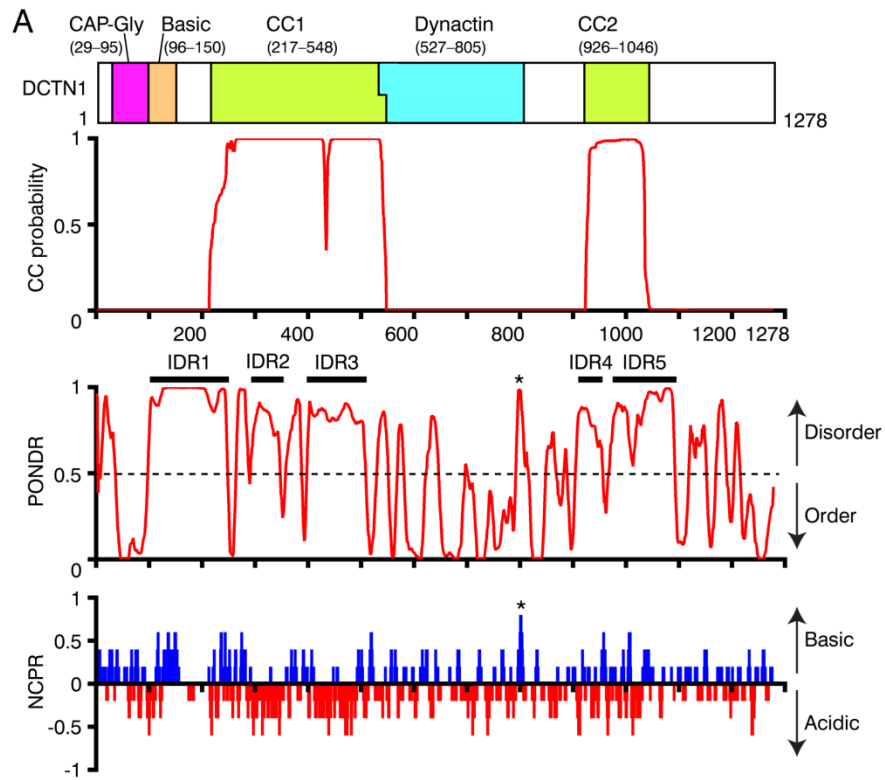
Movie S3

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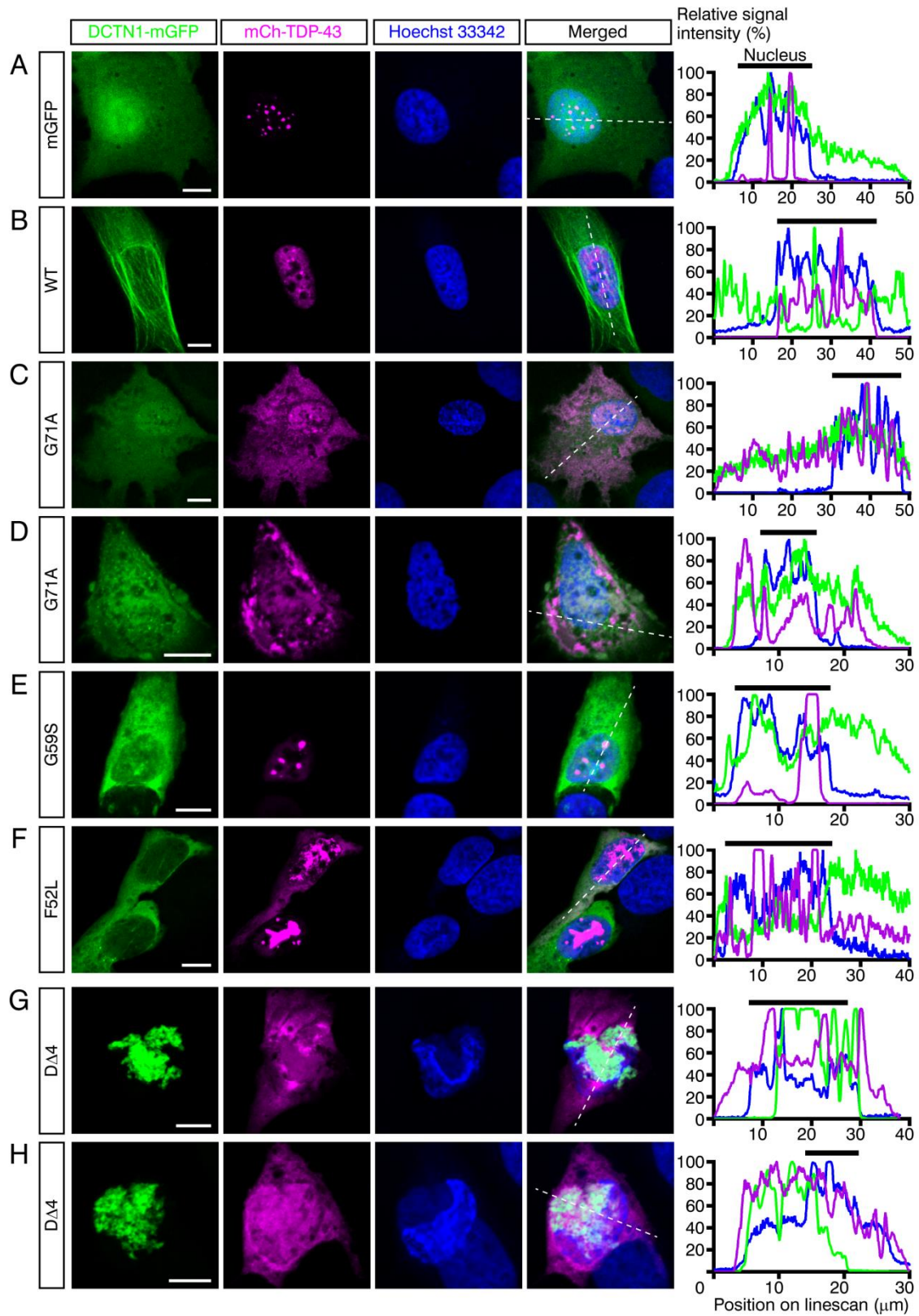
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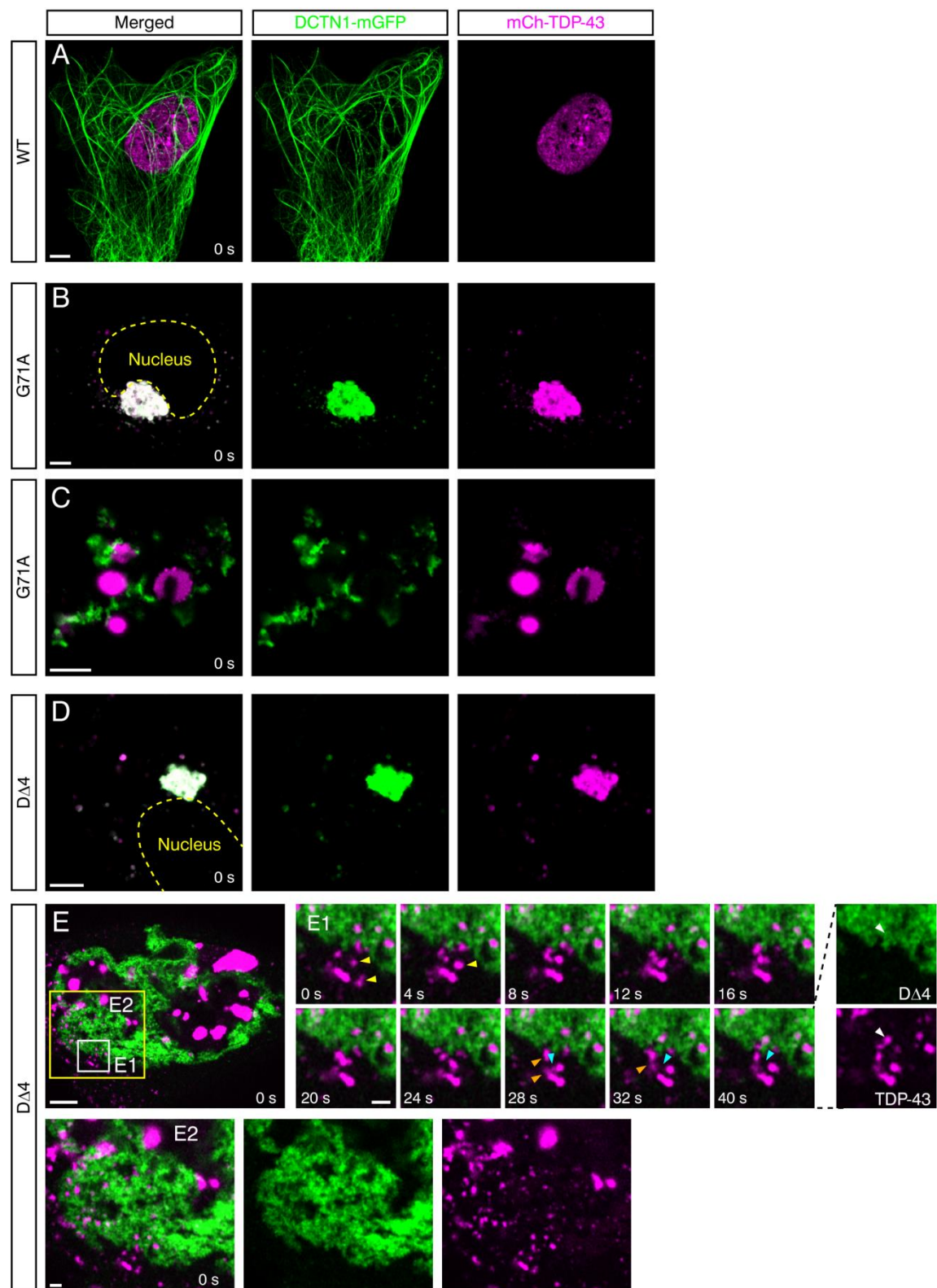
Original blot and gel images (Supporting Information)



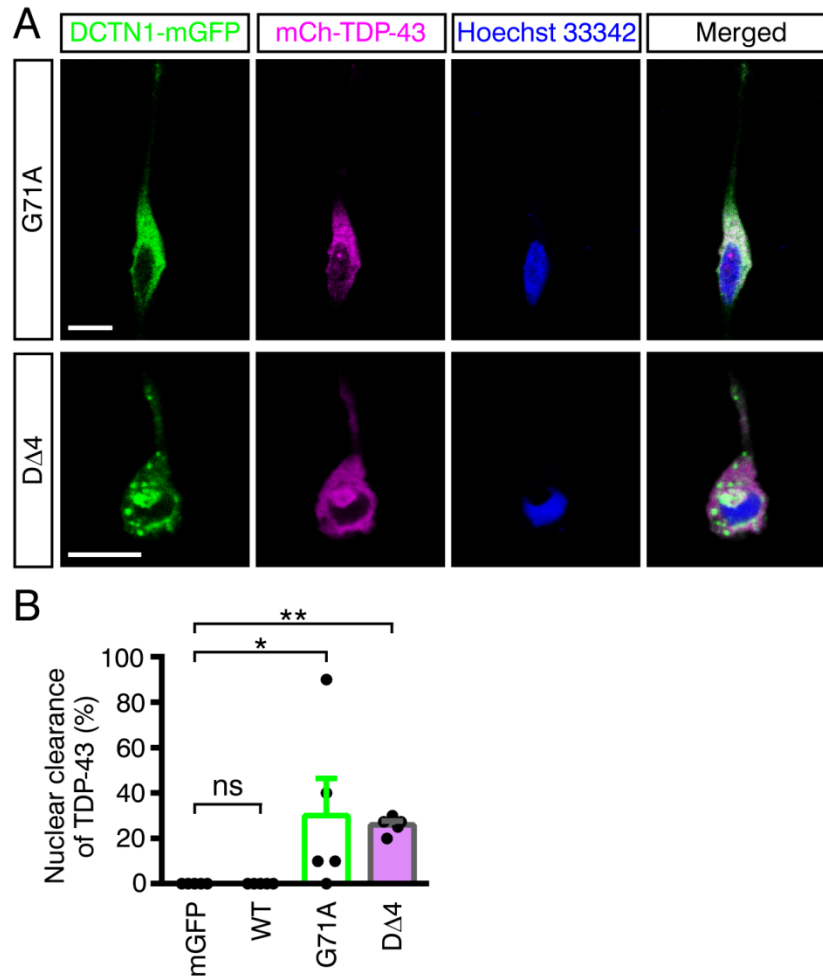
**Figure S1.** Bioinformatics analyses on DCTN1 and TDP43. **(A)** Analyses on DCTN1. Results of coiled-coil (CC) prediction by Multicoil2, disordered domain prediction by PONDR (predictor of natural disordered regions), and NCPR (net charge per residue; 5-amino acid window) analyses are shown. Black bars in the PONDR graph show predicted intrinsically disordered regions (IDRs). In NCPR, charged residues are shown in blue (basic: Arg and Lys) and red (acidic: Asp and Glu). **(B)** Analyses on TDP-43. Results of CC prediction, PONDR, and NCPR are presented.



**Figure S2.** Effects of missense mutant DCTN1 and DA4 on TDP-43 mislocalization and aggregation. Maximum-intensity projections of deconvoluted z-stack confocal images of U2OS cells that co-expressed mGFP (**A**), wild-type (WT) (**B**) or mutant (**C-H**) DCTN1-mGFP and mCherry-tagged WT TDP-43 are shown. The transfected cells were cultured under non-stressed conditions, fixed three days after transfection, and subjected to confocal microscopy. In (F), two representative cellular phenotypes of DCTN1<sup>F52L</sup>-mGFP- and mCherry-TDP-43-coexpressing cells are shown: nuclear aggregation and ubiquitous cytoplasmic distribution of TDP-43. Scale bars, 10  $\mu$ m. The right graphs show linescans of the cells along the white broken lines.



**Figure S3.** Super-resolution live-cell imaging of dynamics of wild-type (WT) or mutant DCTN1-mGFP (p.G71A or D $\Delta$ 4) and mCherry-TDP-43 (WT) in U2OS cells. **(A)** A snapshot of U2OS cells co-expressing WT DCTN1-mGFP (shown in green) and mCherry-TDP-43 (WT; purple). **(B,C)** Snap shots of aggregates of mutant DCTN1 (p.G71A)-mGFP and mCherry-TDP-43 (WT). In (C), the nucleus is located above the top-left corner. **(D,E)** Snap shots of aggregates of mutant DCTN1 (D $\Delta$ 4)-mGFP and mCherry-TDP-43 (WT). E1: time-lapse imaging of the white-boxed area in (E) for 40 s. mCherry-TDP-43 aggregates dynamically undergo remodeling through repeated contacts among neighboring aggregates of DCTN1 (D $\Delta$ 4)-mGFP and mCherry-TDP-43, as well as fission/fusion cycles. Two fusion processes between mCherry-TDP-43 aggregates are marked with yellow and orange arrowheads, while a fission process of mCherry-TDP-43 aggregates is marked with cyan arrowheads (before and after the event). An obviously found colocalization of DCTN1 with TDP-43 is marked by white arrowheads (the right panels). E2: High-power views of the yellow-boxed area in (E). Scale bars, 5  $\mu$ m (A-E) and 1  $\mu$ m (E1 and E2). See Movie S1–S6.



**Figure S4.** Nuclear clearance of TDP-43 in iPSC-derived neurons co-expressing mutant DCTN1 and wild-type TDP-43. **(A)** Examples of nuclear clearance of mCherry-TDP-43 in mutant DCTN1-mGFP (p.G71A or DΔ4)-coexpressing neurons. Single-section confocal images of deconvoluted z-stacks are shown. Scale bars, 10  $\mu$ m. **(B)** Quantitative analysis of nuclear clearance of TDP-43.  $n = 5$  experiments (10–12 cells per experimental group). \*  $p < 0.05$ ; \*\*  $p < 0.01$  by two-tailed Mann-Whitney test. ns: not significant.



A		571	
Hs	KAHAKAIE	<b>M</b>	ELRQMEVAQANRH
M571T	KAHAKAIE	<b>T</b>	ELRQMEVAQANRH
Mm	KAHAKAIE	<b>M</b>	ELRQMEVAQANRH
Bt	KAHAKAIE	<b>M</b>	ELRQMEVAQANRH
Gg	KAHAKAIE	<b>M</b>	ELRQMEVQQANRH
Xl	KAHAKAIE	<b>M</b>	ELRKMEVTQANRH
Dr	KAYAKAIE	<b>M</b>	ELRKMEVSQANRQ
Dm	KAYTRAIDV	<b>Q</b>	LRQIELSQANEH

B		785	
Hs	LQGGQEATDIALLL	<b>R</b>	DLETSCS
R785W	LQGGQEATDIALLL	<b>W</b>	DLETSCS
Mm	LQGGQEATDIALLL	<b>R</b>	DLETSCS
Bt	LQGGQEASDIALLL	<b>R</b>	DLETSCS
Gg	LQAGQEASDLAILL	<b>K</b>	DLETSCS
Xl	LHAGQESSDFAILLK	<b>D</b>	DLETSCS
Dr	LQAGQEGAELCVLLK	<b>D</b>	LDLTSCG
Dm	LSDTAIAKVIIQEAG	<b>T</b>	SDSVL

C		997	
Hs	VQTRLEETQALLRKKEKEFEET	<b>R</b>	
R997W	VQTRLEETQALLRKKEKEFEET	<b>W</b>	
Mm	VQTRLEETQALLRKKEKDFEET	<b>R</b>	
Bt	VQTRLEETQALLRKKEKEFEET	<b>R</b>	
Gg	IQTKLDEETQTLKKKEKEFEET	<b>T</b>	
Xl	IQTKLEETQTVLKKKEKEFEET	<b>T</b>	
Dr	IQTKLDEALNTLKKKEKEFEET	<b>T</b>	
Dm	MQIRKDLAEKKLSVLQNE---	<b>R</b>	Y

D		1249	
Hs	MGKVTFSCAAGFGQRHRLVLTQ	<b>T</b>	
T1249I	MGKVTFSCAAGFGQRHRLVLTQ	<b>I</b>	
Mm	MGKVTFSCAAGLGQRHRLVLTQ	<b>T</b>	
Bt	MGKVTFSCAAGLGQRHRLVLTQ	<b>T</b>	
Gg	CGRWCSSSRARASPPASACSPP	<b>C</b>	
Xl	IGKVTLSCQPGQGQIHKLVLTQ	<b>T</b>	
Dr	VSKLMIPCARGKEQKHTLVLSQ	<b>M</b>	
Dm	ASDILTEYLQRKPHRATHGQFA	<b>T</b>	

**Figure S5.** Amino acid sequence conservation in the dynactin domain and C-terminal region of DCTN1 across several animal species. (A-D) Four segregated sites of missense mutations from patients with ALS are shown; (A) p.M571T, (B) p.R785W, (C) p.R997W, and (D) p.T1249I. The positions of putative ALS-linked mutations are highlighted in bold. Hs: *Homo sapiens*; Mm: *Mus musculus*; Bt: *Bos Taurus*; Gg: *Gallus gallus*; Xl: *Xenopus laevis*; Dr: *Danio rerio*; Dm: *Drosophila melanogaster*.

**Movie S1.** A representative case of super-resolution live-cell imaging of a U2OS cell co-expressing DCTN1-mGFP (wild-type [WT]; shown in green) and mCherry-TDP-43 (WT; purple). The example is from Figure S3A. Images were taken every 6 s, and are shown at 3 frames per second.

**Movie S2.** Super-resolution live-cell imaging of a U2OS cell co-expressing DCTN1-mGFP (p.G71A; shown in green) and mCherry-TDP-43 (WT; purple). The example is from Figure S3B. A large cytoplasmic aggregate, near the nucleus, in which DCTN1<sup>G71A</sup>-mGFP and mCherry-TDP-43 colocalized, was found. Images were taken every 6 s, and are shown at 3 frames per second.

**Movie S3.** Super-resolution live-cell imaging of a U2OS cell co-expressing DCTN1-mGFP (p.G71A; shown in green) and mCherry-TDP-43 (WT; purple). The example is from Figure S3C. Cytoplasmic DCTN1<sup>G71A</sup> aggregates partially colocalized with or repeatedly contacted those of TDP-43. Images were taken every 6 s, and are shown at 3 frames per second.

**Movie S4.** Super-resolution live-cell imaging of a U2OS cell co-expressing DCTN1-mGFP (DΔ4; shown in green) and mCherry-TDP-43 (WT; purple). The example is from Figure

S3D. A large cytoplasmic aggregate in which D $\Delta$ 4 and TDP-43 colocalized was observed.

Images were taken every 4 s, and are shown at 3 frames per second.

**Movie S5.** Super-resolution live-cell imaging of a U2OS cell co-expressing DCTN1-mGFP (D $\Delta$ 4; shown in green) and mCherry-TDP-43 (WT; purple). A whole-cell view; the same cells as that shown in Figure S3E was live-imaged. Aggregates of mCherry-TDP-43 interacted with massive aggregates of D $\Delta$ 4-mGFP. Images were taken every 4 s, and are shown at 3 frames per second.

**Movie S6.** Super-resolution live-cell imaging of a U2OS cell co-expressing DCTN1-mGFP (D $\Delta$ 4; shown in green) and mCherry-TDP-43 (WT; purple). A higher-power view of the cell presented in Movie S5. Microaggregates of mCherry-TDP-43 repeatedly contacted the aggregates of neighboring aggregates of mCherry-TDP-43 and those of DCTN1 (D $\Delta$ 4), and underwent cycles of fission and fusion, via a liquid-liquid phase separation (LLPS)-mediated process. mCherry-TDP-43 aggregates frequently appeared to grow in such a manner. Images were taken every 4 s, and are shown at 1.5 frames per second. See Figure S3E (E1).