

Expanded View Figures

Figure EV2. Validation of O-GlcNAcylated modification on TDP-43.

A Mass spectra data show the original peptide sequences containing O-GlcNAcylation sites of TDP-43. Top: T199 site; bottom: S212, T233 sites.

B Summary of three identified O-GlcNAcylated peptides of TDP-43 with significant ion scores by mass spectra analysis. "A score" represents the fidelity of MS.

Source data are available online for this figure.

Α T199 Intensity (%) 100**1** У5 644.39 3541.62 2808.43 У13 1520.79 Y27 3312.54 Intensity Y26 У14 1683.85 ¥10 1249,68 5 У7 904.54 Y25 3065.47 У6 757.47 У17 2047.03 Va Y11 1348.74 Y20 Y22 2479.22 2721.36 Y23 h1 Y12 b11 1898.94 У16 m/z _____ m/z 1000 1500 2500 3000 3500 500 FSqYGDVMDVFIPKPFR R + LE + D + ET + M + D + bMax yMax D I + F + v +7+D-IG + F-- सम्ब Et Intensity (%) m/z 1000 2000 3000 S212, T233 y¹⁸ b., Inter 100 Y10 Y12 1147.60 1346.70 947.49 Intensity у7 787.46 Y11 _{¥16} 1817.89 ¥9 ^y17 1888.93 У<u>1</u>3 1459.78 95 601.39 147.11 ^y15 1702.87 m/z -----bMax -----yMax 30YGDVmDVFIPKPFRAFAFV tFADDQIAQSLcGEDLIIK--Intensity (%) m/z 1000 2000 В Peptide sequence A score O-GIcNAc site R.EFFS(GIcNAc)QYGDVMDVFIPKPFRAFAFVT 55.4 S212 (GICNAC)FADDQIAQSLCGEDLIIK.G T233 95.3 C.T(GIcNAc)EDMTEDELREFFSQYGDVMDVFIP

T199

67.7

Figure EV2.

KPFR.A





Cells expressing the indicated forms of GFP-TDP-43 were subjected to immunoprecipitation, and the O-GlcNAcylated levels of TDP-43 were examined. Source data are available online for this figure.

Figure EV4. O-GlcNAcylation deficient mutations do not affect the RNAbinding ability of TDP-43.

RNA-EMSA was performed using total cell lysate from the indicated GFP-TDP-43 transfected HEK 293T cells and biotin-labeled IL-6 RNA probe (top panel). The indicated GFP-TDP-43 proteins were examined by immunoblotting (bottom panel).

Source data are available online for this figure.





Figure EV5. TDP-43 affects OGT mRNA splicing.

A CLIP-Seq data show the RNA-binding sites on OGT of TDP-43. The numbers shown in y-axis represent the intensity of signals, and the numbers shown in x-axis represent exons of OGT.

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B RNA splicing status of OGT at exon 7 was examined by PCR analysis using specific primers flanking exon 7. The number represented the ratios of exon 7 exclusion (-) vs. inclusion (+) (top panel). TDP-43, OGT, and OGA protein levels were examined by immunoblotting (bottom panel).

Source data are available online for this figure.