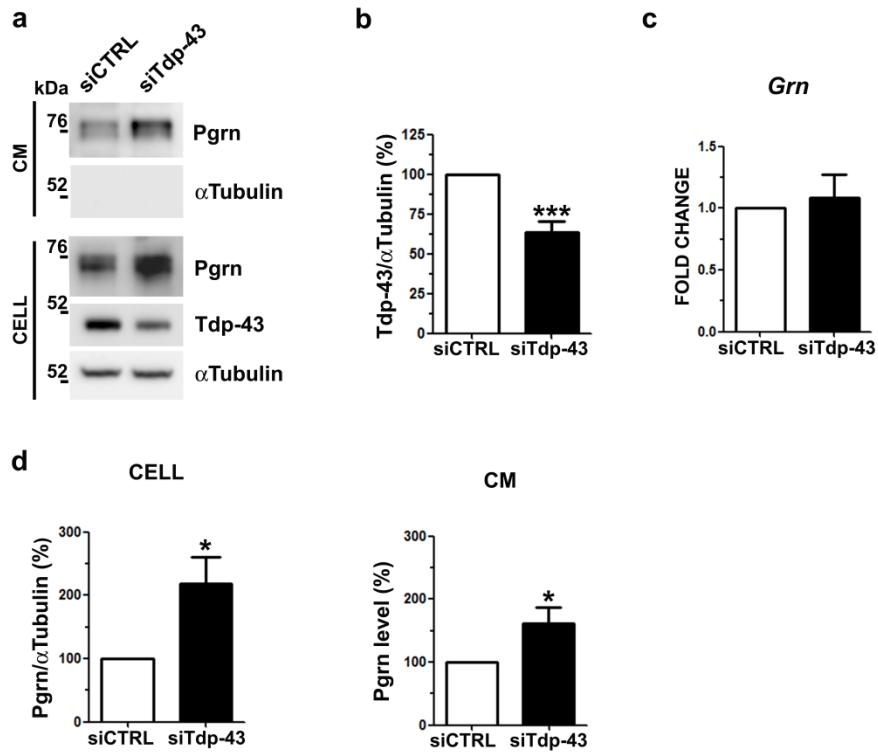


SUPPLEMENTARY MATERIALS

Supplementary Figure 1



Analysis of intracellular and secreted Pgrn levels upon Tdp-43 LOF in murine neuroblastoma N2a cells.

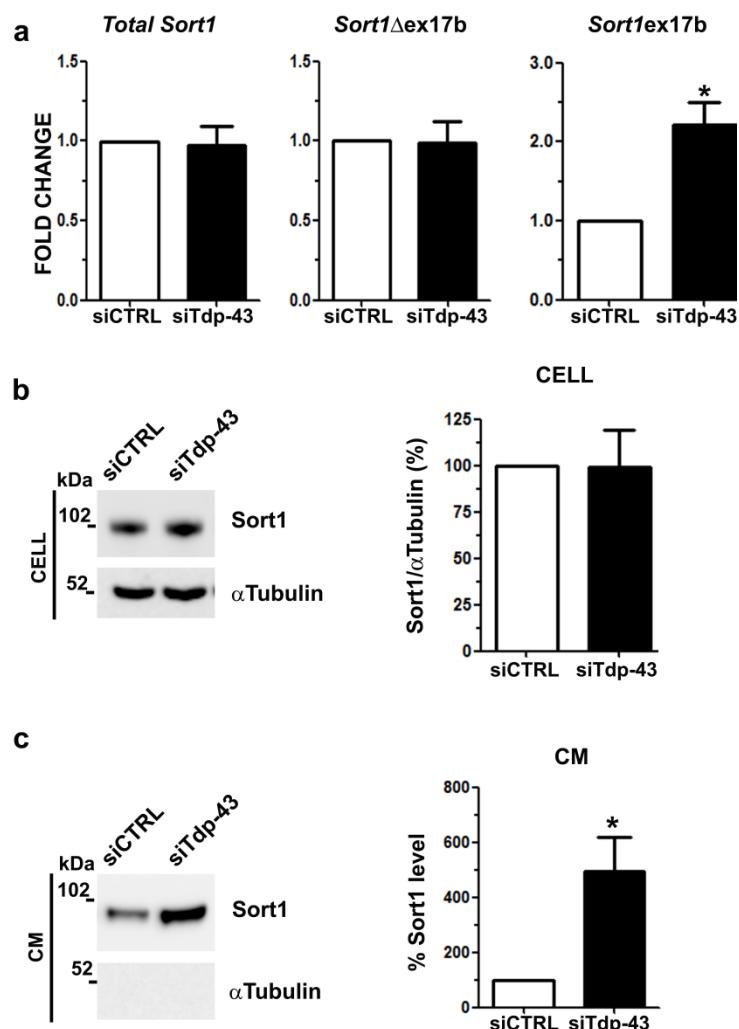
- a) Representative WB images of Pgrn levels in cell lysates (CELL) and in conditioned media (CM) upon Tdp-43 depletion in murine N2a cells. Tdp-43 immunoblot was reported to show gene silencing efficiency, α-Tubulin was used for data normalization and as negative control in conditioned media.
- b) Densitometric and statistical analyses of Tdp-43 protein levels shown in (a).
- c) Real time PCR of *Grn* gene expression upon Tdp-43 depletion.
- d) Densitometric and statistical analyses of intracellular (CELL) and secreted (CM) Pgrn levels in Tdp-43 depleted N2a cells shown in (a) (mean±s.e.m.; n=4 independent experiments; Two-tailed Unpaired *t* test; * p< 0.05, *** p<0.001.)

Supplementary Figure 2

Human and murine GRN 3'UTR sequence alignment and the putative TDP-43 binding region.

The human (NM_002087.3) and murine (NM_008175.5) *GRN* 3'UTR sequence alignment performed by the pairwise sequence alignment EMBOSS Needle software. Red boxes were used to highlight the putative binding site for TDP-43, containing the (TG)₆ motif (dark red box), in the murine sequence. Blue boxes indicate the TDP-43 binding region in the human *GRN* 3'UTR sequence predicted by RBPmap software setting a stringent Z-score ≥ 5 . The (TG)₅ motif is highlighted (dark blue box). The primer sequences used for cloning of the murine and human 3'UTR sequences are indicated.

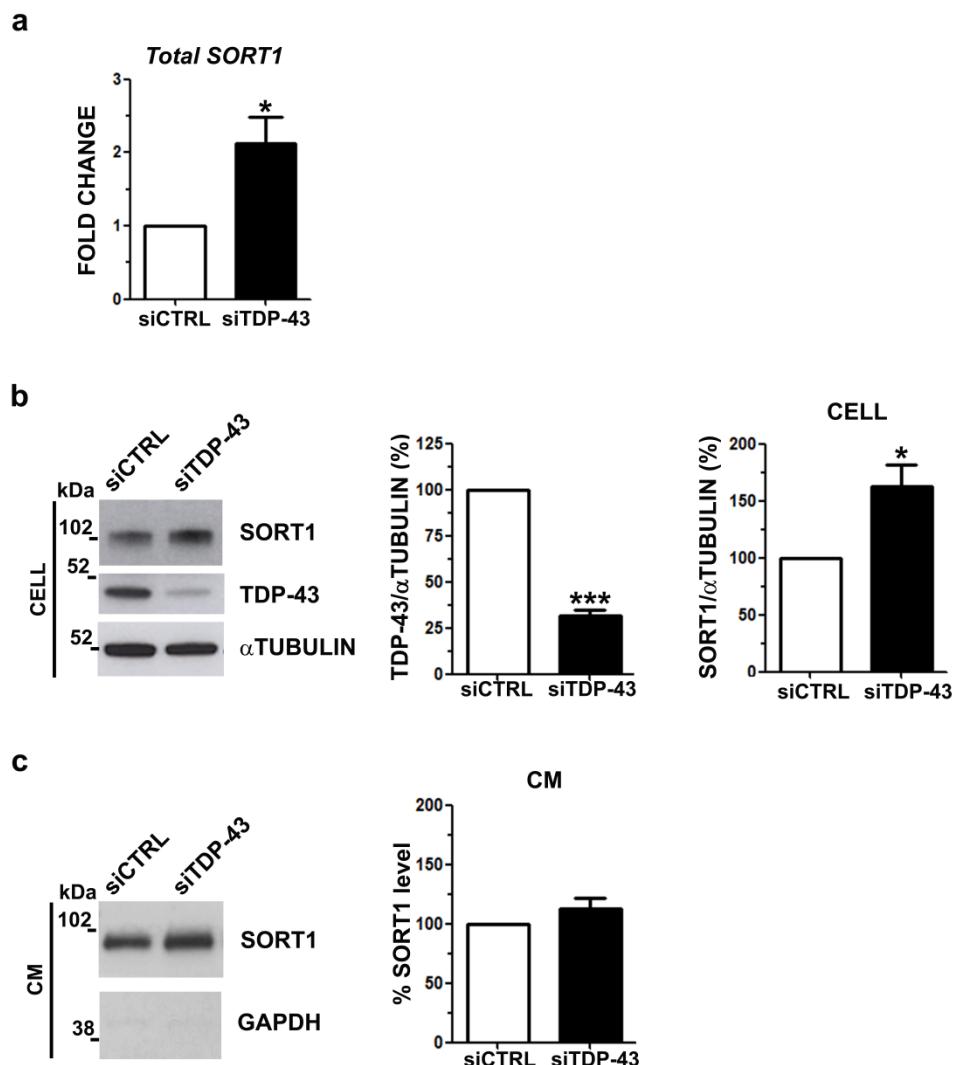
Supplementary Figure 3



Sortilin1 splicing and protein levels upon Tdp-43 LOF in murine neuroblastoma N2a cells.

- a)** Real time PCR of total and splicing isoforms of *Sort1* in Tdp-43 knocked-down murine neuroblastoma N2a cells shown in Supplementary Figure 1. **b)** Representative WB images and densitometric analysis of total intracellular (CELL) Sort1 protein levels upon Tdp-43 knock-down in N2a cells. α -Tubulin was used for data normalization. **c)** Representative WB and densitometry data of Sort1 levels in the conditioned media (CM) (mean \pm s.e.m.; n=4 independent experiments; Two-tailed Unpaired *t* test; * p< 0.05).

Supplementary Figure 4



Sortilin1 splicing and protein levels upon TDP-43 *LOF* in human neuroblastoma SK-N-BE cells.

- a)** Real time PCR of total *SORT1* in condition of TDP-43 gene silencing in human neuroblastoma SK-N-BE cells (mean \pm s.e.m.; n=4 independent experiments; Two-tailed Unaired *t* test; * p< 0.05).
- b)** Representative WB images and densitometric analysis of total *SORT1* protein levels upon TDP-43 knock-down in SK-N-BE cells. TDP-43 immunoblot was performed to assess gene silencing efficiency and α -Tubulin was used for data normalization (mean \pm s.e.m.; n=4 independent experiments; Two-tailed Unpaired *t* test; * p< 0.05; *** p< 0.001).
- c)** Representative WB and densitometry data of *SORT1* levels in the conditioned media (CM). GAPDH immunoblot was performed as negative control (mean \pm s.e.m.; n=3 independent experiments; Two-tailed Unpaired *t* test).

Supplementary Table S1. List of the primary antibodies used for immunofluorescence (IF) and Western blot (WB) assays.

Antibody	Source	Assay
GFP (1:2000)	Roche_11814460001	IF/WB
PGRN (mouse) (1:1000)	R&D Systems_AF2557	WB
PGRN (human) (1:300)	Invitrogen_40-3400	WB
TDP-43 (1:1000)	Protein Tech_10782-2-AP	WB
SORT1 (1:300)	R&D Systems_MAB3154	WB
α Tubulin (1:500)	Sigma-Aldrich_T6199	WB
POLDIP3 (1:500)	Cell Signaling_5439	WB
FLAG (1:1000)	F3165 (Sigma-Aldrich)	IP
goat anti-mouse IgG	sc-2025 (Santa Cruz)	IP

Supplementary Table S2. List of primers sequences used for real time PCR.

Gene	Foward primer	Reverse primer	Ref
<i>murine GRN</i>	CCAACTACAGCTGCTGTAAC	CTCGTTATTCTAGGCCATGTG	[7]
<i>murine SORT1 (Total)</i>	CGTGTTCCCTGGAGGACTTCCT	TTCAGGCTGCTCCACGCCT	[21]
<i>murine SORT1ex17b</i>	AAATCCCAGGAGACAAATGC	GAGCTGGATTCTGGGACAAG	[21]
<i>murine SORT1Δex17b</i>	CCCCACAAAGCAGAATTCCAAGTC	TGACAAGCATCAGTCCCACGAT	[21]
<i>murine RPL10a</i>	GAAGAAGGTGCTGTCTGGC	TCGGTCATCTCACGTGGC	[7]
<i>human GRN</i>	CAGCTACAGCTGCTGCCGTC	CTCAGTGGTGGGCCATTG	-
<i>human SORT1 (Total)</i>	GGCCTGTGGGTGTCCAAGAA	GCAGGAGCCATTGCATAGGTT	[21]
<i>human SORT1ex17b</i>	AATCCAGCTCTGCCTCCTCT	TCCCACGATGGCCAGGATAA	[21]
<i>human SORT1Δex17b</i>	TGGGGTAAATCCAGTCGAG	GACTTGGATTCTGTTTCCGGAC	[21]
<i>human RPL10a</i>	GAAGAAGGTGTTATGTCTGG	TCTGTCATCTCACGTGAC	[20]