

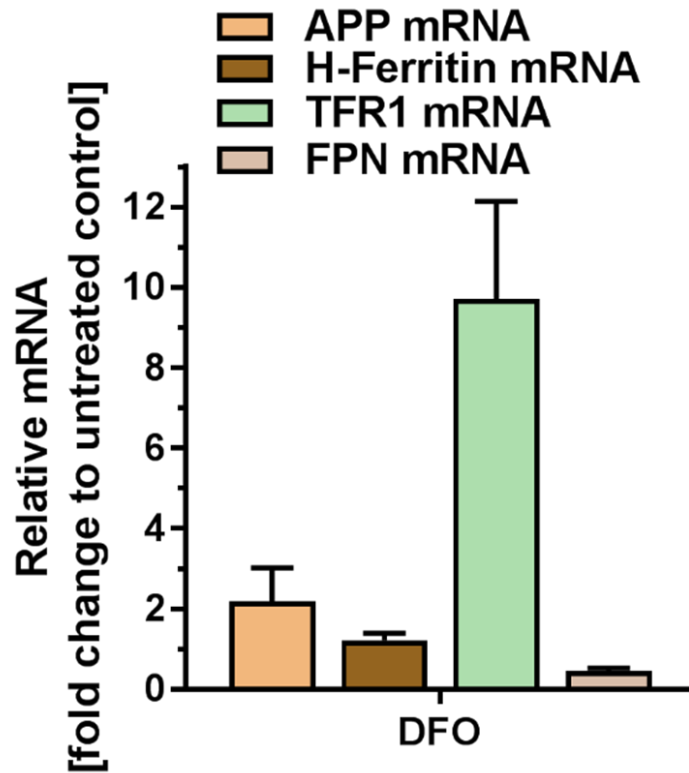
Supplemental information: Manganese causes neurotoxic iron accumulation via translational repression of Amyloid Precursor Protein (APP) and H-Ferritin

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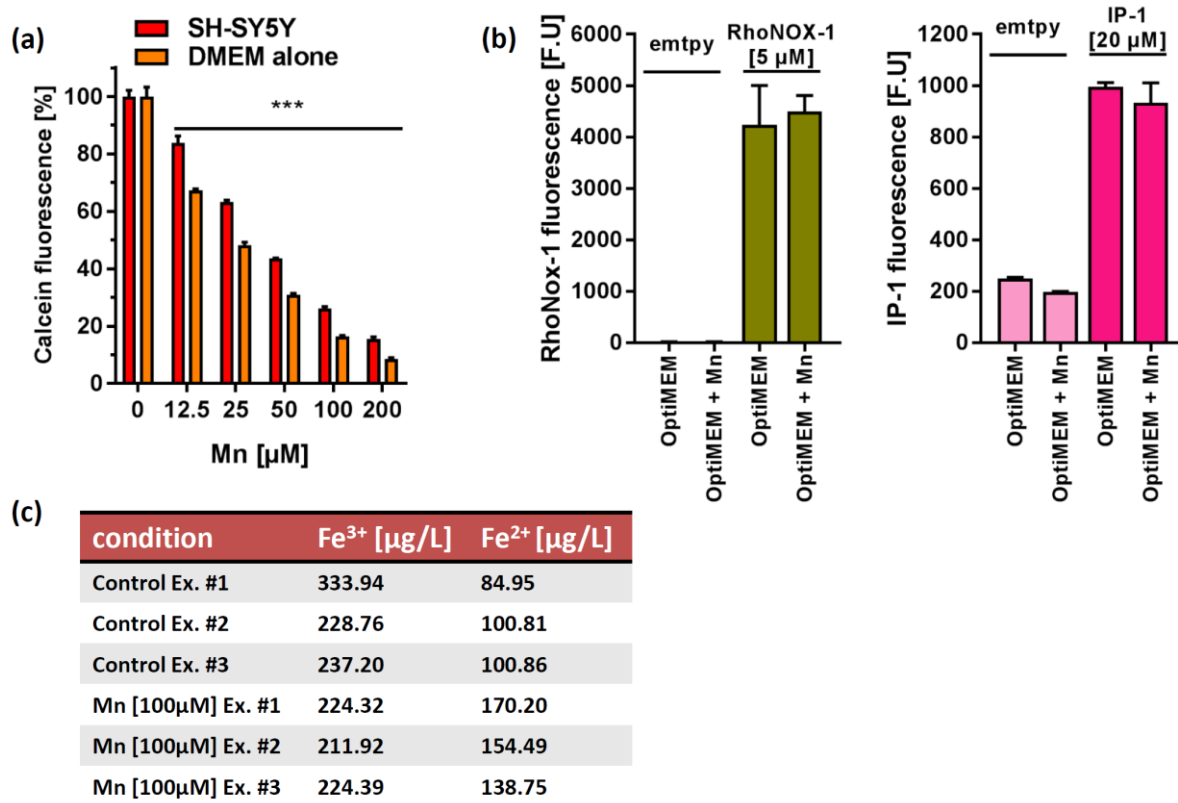
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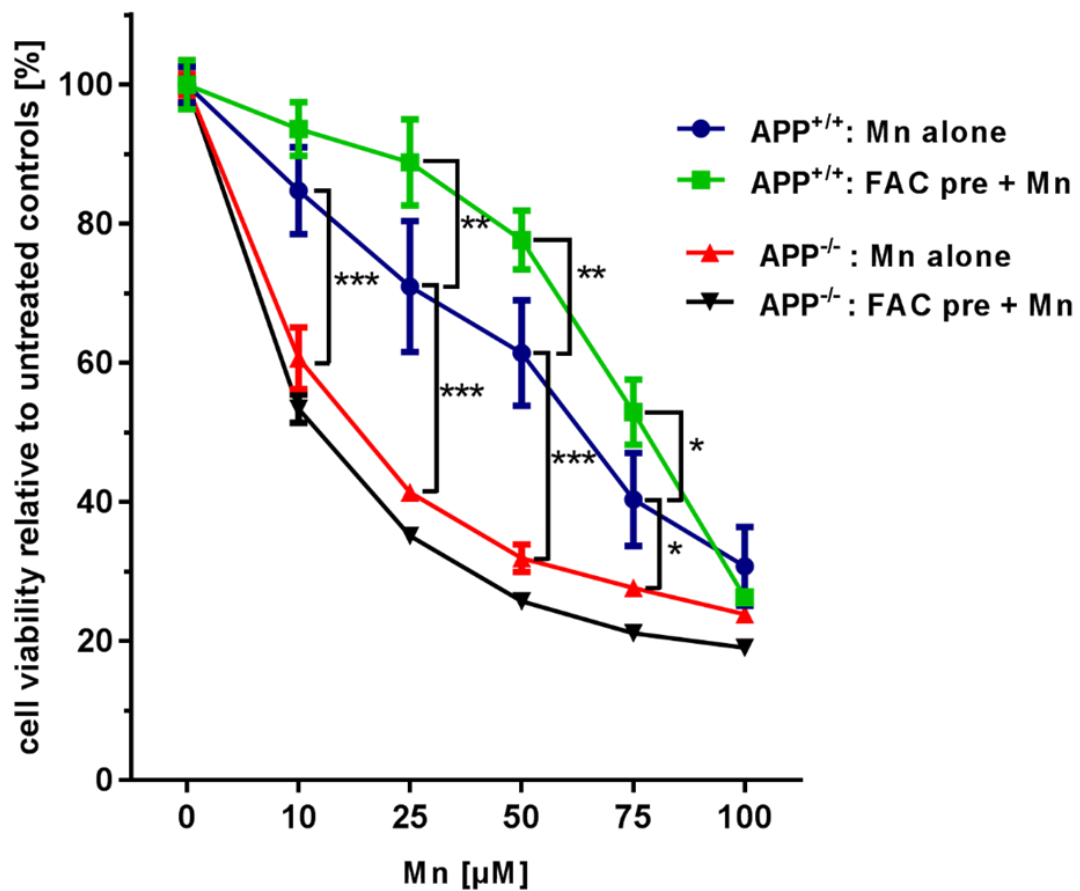
Supplemental Fig. 1 (corresponding to Fig. 2): RT-PCR analysis of indicated iron responsive target genes (APP, H-Ferritin, TFR1 and FPN) from SH-SY5Y cells incubated for 24 h with 100 μ M DFO. All data are normalized to housekeeping gene β -actin and presented as relative fold-induction compared to untreated controls (n=4). N number indicates number of replicates. Data shown represent mean \pm SD.

		50	60	70	80	90			
Human APP mRNA (46)	GGGGCCCC	GGAGACGGCGGC	GTGGCGG	CGCGGG	CAGAGCAAGGACGGCGCGGATC		(-43)	AUG	
Mouse APP mRNA (33)	GGGGCCACC	GGAGACGGCGGC	CCGGAC	GC	CACAGCCAGGCGCGCGGATC		(-45)	AUG	
Rat APP mRNA (33)	ACCGGAGAC	GGCGCGCGCGCGG	CGACGAC	CGCGGAC	CACAGTCAGGGTGGCGCGGATC		(-43)	AUG	
Human FPN (99)	GCTTCCATA	AAGGCTTTGCCTT	TCCA	ACTTCAGCT	CAGTGT	TAGCTAAGTTGGAAA	(-94)	AUG	
Human L-Fer (3)	GTTTCGGCGGT	CCCGCGGTCTG	TCTCTTG	CTTCAA	CAGTGT	TTGGACGGAA	CAGATC	(-139)	AUG
Human H-Fer (5)	TCTTCGCCGA	GAGTCGTCGGG	TTTCTG	CTTCAA	CAGTGT	TTGGACGGAA	CCGGC	(-149)	AUG
Consensus			G-----G-----	C-----	CA--GN--G---	G---C---C			
					*	**	**	*	
Conserved C	+76			82-87	Loop (wrt human APP mRNA)				
	+33			39-44	Loop (wrt human L-Ferritin mRNA)				

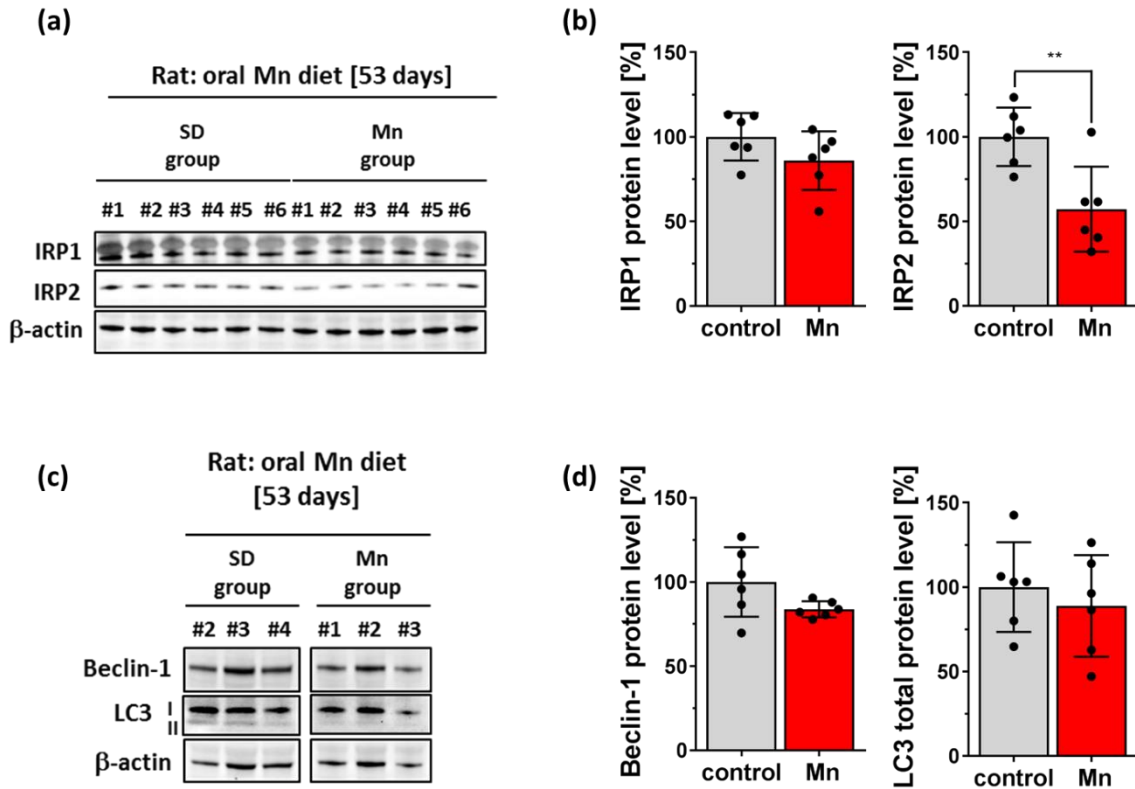
Supplemental Fig. 2 (corresponding to Fig. 3): Bioinformatics alignment of the 5'untranslated regions of APP mRNAs when aligned to the ferritin canonical IREs AGU and AGA tri-loops as a potential IRP 1 binding site (bold lettering).



Supplemental Fig. 3 (corresponding to Fig. 4): (a) SH-SY5Y cells or complete DMEM media alone were incubated with increasing Mn doses for 24 h, loaded with calcein-AM and analyzed for fluorescence. Data are presented as relative fluorescence compared to untreated fluorescence (n=6). (b) Cell-free background fluorescence of OptiMEM media ± 100 μM Mn (empty) or OptiMEM media ± 100 μM Mn loaded with either 5 μM RhoNOX-1 (left panel) or 20 μM IP-1 (right panel). Data are presented as absolute fluorescence units (F.U.) (n=3). (c) Absolute Fe²⁺ and Fe³⁺ values assayed by IC-ICP-MS and CE-ICP-MS from SH-SY5Y lysates treated with DMEM media or 100 μM Mn. Corresponding relative Fe³⁺ to total iron and Fe²⁺ to total iron ratio are shown in Fig. 4b. Data are presented as mean ± SD. N number indicates number of replicates. Differences were calculated using either two-way ANOVA followed by Bonferroni post-hoc analyses. (*, p < 0.05; **, p < 0.01; ***, p < 0.001).



Supplemental Fig. 4 (corresponding to Fig. 5): Immortalized murine embryonal fibroblasts (MEFs) were isolated from wild-type mice (APP^{+/+}) or APP-knockout (APP^{-/-}) mice. APP^{+/+} and APP^{-/-} MEFs were either pretreated with 100 μM FAC or media for 24 h and subsequently incubated with increasing doses of Mn for additional 24 h and analyzed for cell viability (n=3). Data are presented as mean ± SD. N number indicates number of replicates. Differences were calculated using two-way ANOVA followed by Bonferroni post-hoc analyses. (*, p < 0.05; **, p < 0.01; ***, p < 0.001).



Supplemental Fig. 5 (corresponding to Fig. 6): Corresponding western blots of IRP1 and IRP2 from whole brains lysates of either Sprague-Dawley rats treated with standard diet (SD group) or Mn (500 mg/kg) for 53 days. β -actin served as loading control. **(b)** Densitometric quantifications were normalized to β -actin (each group: n=6). **(c)** Representative western blots of autophagy marker Beclin-1 and LC3. β -actin served as loading control. **(d)** Densitometric quantifications were normalized to β -actin (each group: n=6). N number indicates number of individual animals. Data are presented as mean \pm SD. Differences were calculated using unpaired t-test.