Supplementary information

Parkin is transcriptionally regulated by ATF4: evidence for an interconnection between mitochondrial stress and ER stress

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Supplemental Figure 1:

Mitochondrial stress induced by CCCP activates ATF4-dependent transcription.

HEK293T cells were transfected with the ATF4RE-luc reporter contruct containing a confirmed ATF4-binding site (A), or with park-luc (B), containing the putative ATF4-binding site of the parkin promoter. One day after transfection the cells were exposed to CCCP (10 μ M for 24 h). Shown is the fold induction of luciferase activity in CCCP-treated cells in comparison to non-treated control cells.

Supplemental Figure 2:

c-Jun knockdown efficiency corresponding to the experiment shown in Fig. 5 D. mRNA levels were determined by quantitative RT-PCR using c-Jun-specific primers.

Supplemental Figure 3:

Parkin-deficiency decreases cell viability in response to ER stress. HEK293T cells (A) or SH-SY5Y cells (B) were transfected with parkin-specific or control siRNA duplexes. Three days later the cells were stressed with TG (10 μ M) or TM (2 μ g/ml) for 16 h. Cellular viability was determined by the MTT assay. Shown is the relative viability of cells transfected with parkin siRNA in comparison to control siRNA-transfected cells treated with ER stressors based on triplicates of at least 3 independent experiments. The parkin knockdown efficiency was verified by Western blotting using the anti-parkin mAb PRK8 (A, lower panel) or by quantitative RT-PCR using parkin-specific primers (B, lower panel).

Supplemental Figure 4:

CHOP, phospho-c-Jun and phospho-JNK are sensitive to the severity of ER stress and are increased in parkin-deficient cells. A, B. Levels of CHOP, phospho-c-Jun and phospho-JNK increase with the severity of ER stress in SH-SY5Y cells. SH-SY5Y cells were treated

with thapsigargin (TG) at the indicated concentrations for 16 h to analyze CHOP levels (A) or 1 h to determine the levels of phospho-c-Jun and phospho-JNK (B) by Western blotting. Loading was controlled by re-probing the blots for β -actin. **C**, **D**. CHOP, phospho-c-Jun and phospho-JNK levels are increased in parkin-deficient patient fibroblasts. Skin fibroblasts from a patient with pathogenic mutations in the parkin gene leading to a loss of parkin expression, were incubated with TG at the indicated concentrations for 16 h to analyze CHOP levels (C) or 1 h to determine the levels of phospho-c-Jun and phospho-JNK (D) by Western blotting. As a control, age- and gender-matched skin fibroblasts were analyzed in parallel. Loading was controlled by re-probing the blots for β -actin. The two JNK-immunoreactive bands represent phosphorylated JNK1 (46 kDa) and JNK2/3 (54 kDa), respectively.

Supplemental Figure 5:

ER stress luciferase reporter constructs cloned to quantify the ER stress response.

Binding of ER stress-specific transcription factors to specific sites and UPR transducers involved are schematically shown. ERSE: ER stress response element, UPRE: unfolded protein response element, ERSE II: ER stress response element II, ATF4RE: ATF4 response element.

Supplemental Figure 6:

CHOP suppresses the activation of the park-luc reporter construct in response to ER stress. HEK293T cells were co-transfected with the park-luc reporter plasmid and ATF4, CHOP (1), or GFP (as a control). 8 h after transfection cells were incubated with 1 μM TG for 14 h. Shown is the fold induction of luciferase activity in comparison to GFP-expressing control cells based on triplicates of at least 3 independent experiments (left panel). Expression levels of ATF4 and CHOP were analyzed by immunoblotting using the anti-ATF4 pAb C-20 or the anti-CHOP sc-7351 mAb (Santa Cruz Biotechnology, Santa Cruz, CA, USA) (right panel). Loading was controlled by re-probing the blots for β -actin. *** p < 0.001.

Reference:

1. Al Sarraj J, Vinson C, Thiel G. Regulation of asparagine synthetase gene transcription by the basic region leucine zipper transcription factors ATF5 and CHOP. Biol Chem. 2005 Sep;386(9):873-879.







0,4

0,2

0

control parkin siRNA siRNA

Supplemental Figure 4, Bouman et al.









