

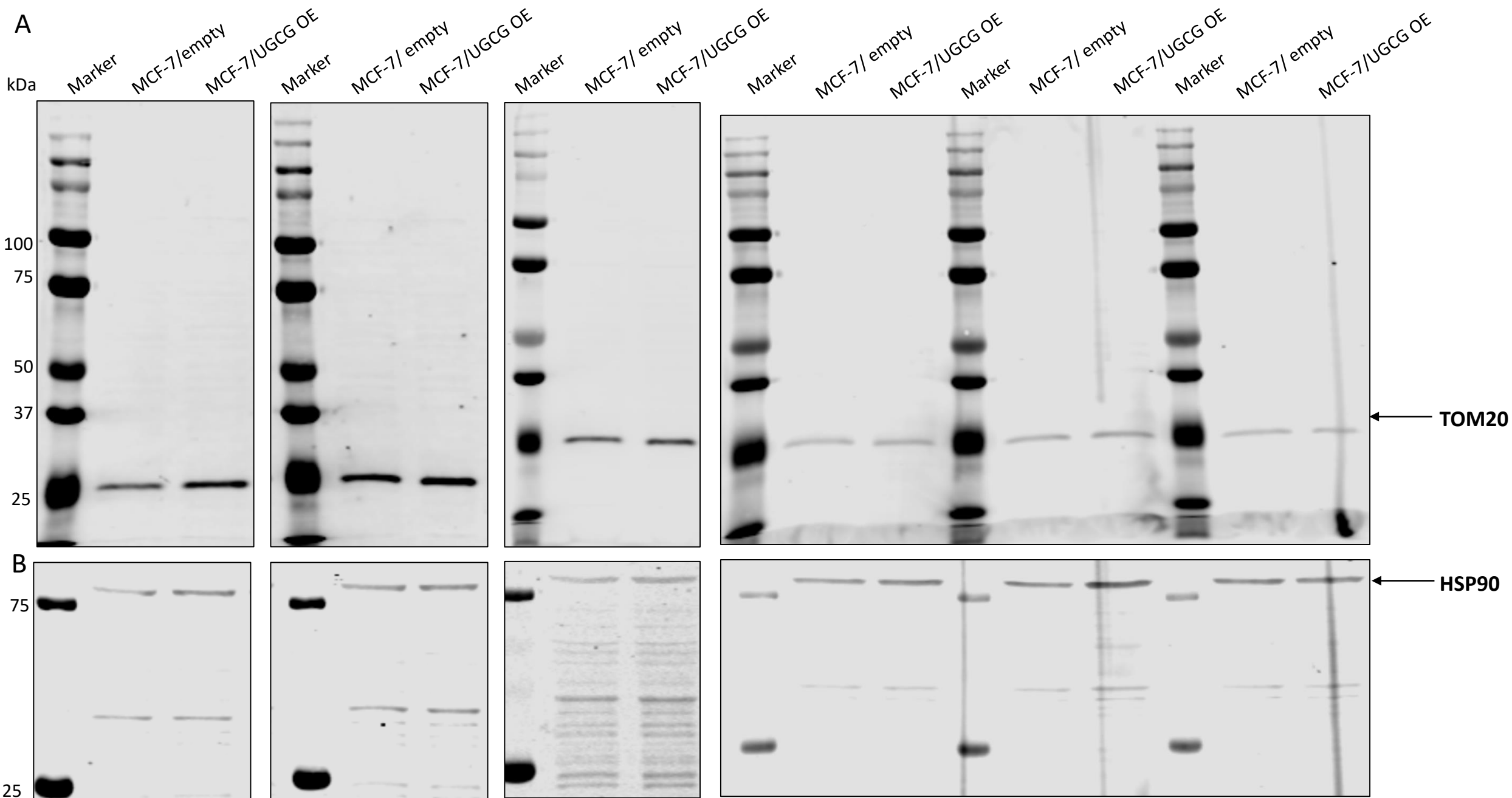
UGCG overexpression leads to increased glycolysis and increased oxidative phosphorylation of breast cancer cells

Nina Schömel¹, Lisa Gruber¹, Stephanie J. Alexopoulos², Sandra Trautmann¹, Ellen M. Olzomer², Frances L. Byrne², Kyle L. Hoehn², Robert Gurke^{1,3}, Dominique Thomas¹, Nerea Ferreirós¹, Gerd Geisslinger^{1,3}, Marthe-Susanna Wegner^{1,2}

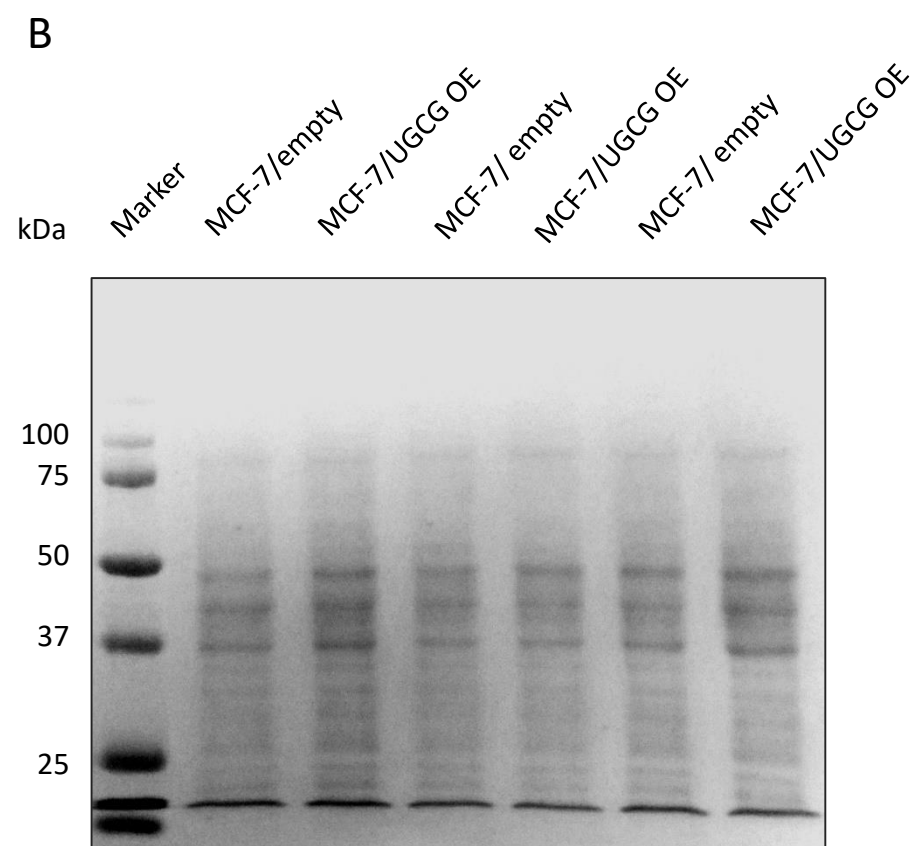
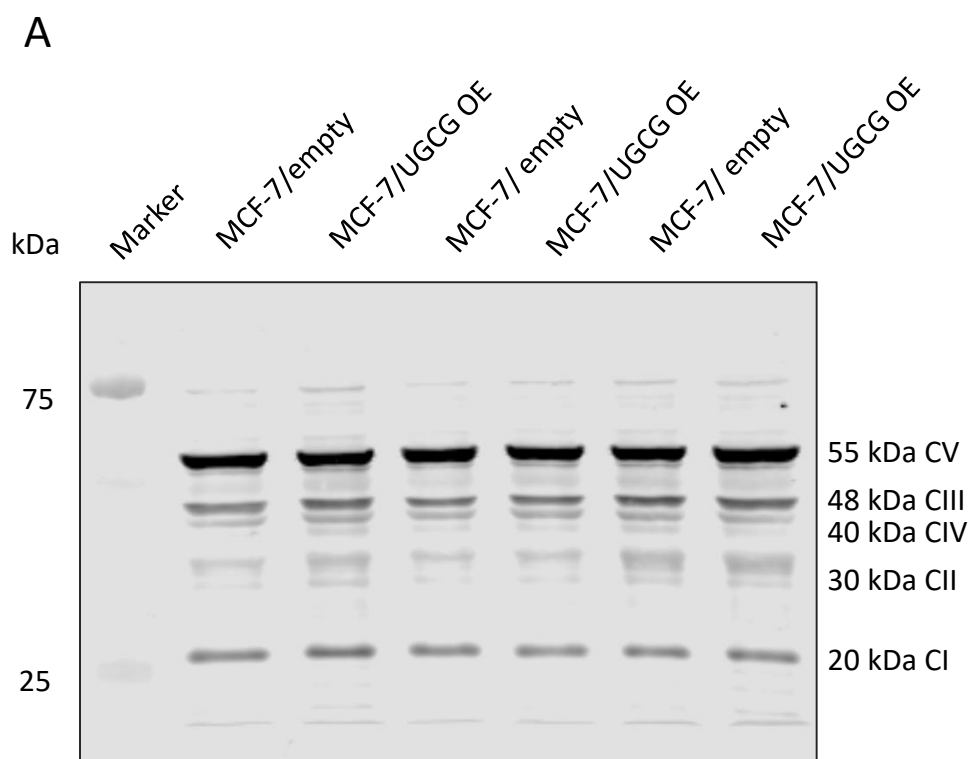
¹*pharmazentrum frankfurt/ZAFES*, Institute of Clinical Pharmacology, Johann Wolfgang Goethe University, Theodor Stern-Kai 7, 60590 Frankfurt am Main, Germany.

²School of Biotechnology and Biomolecular Sciences, University of New South Wales, Sydney, New South Wales 2052, Australia.

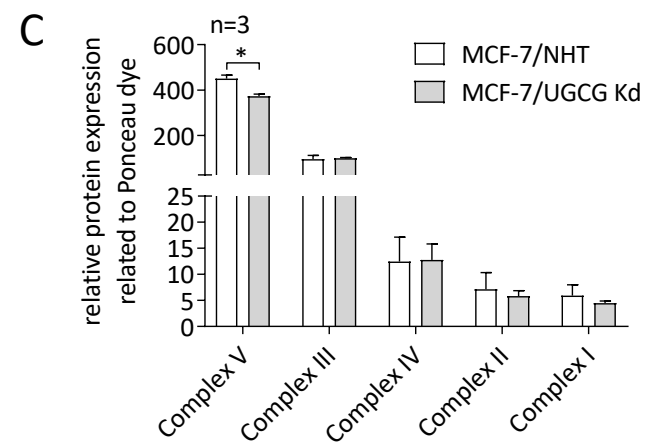
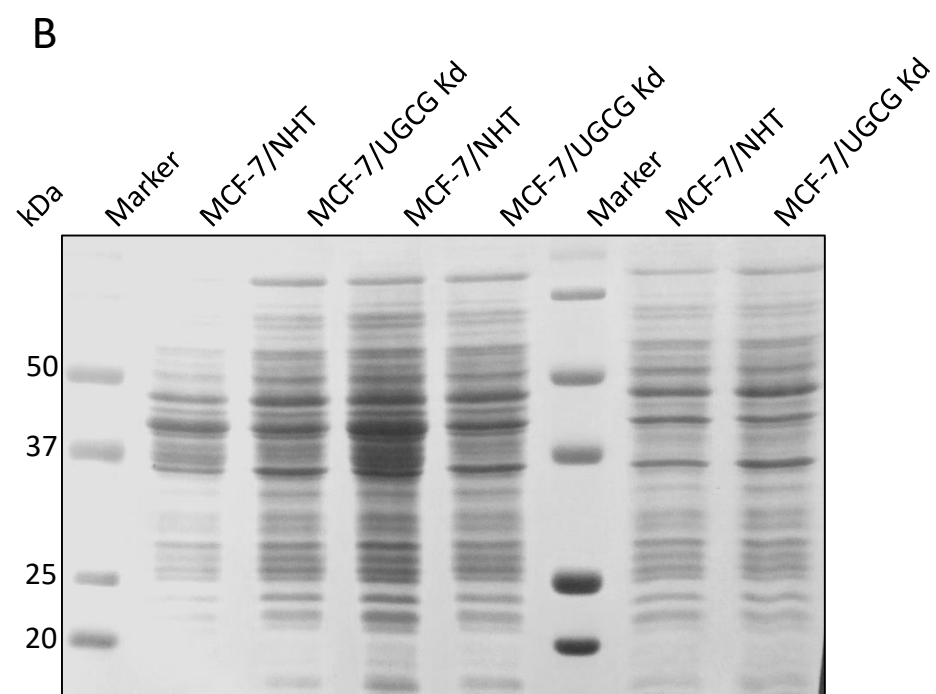
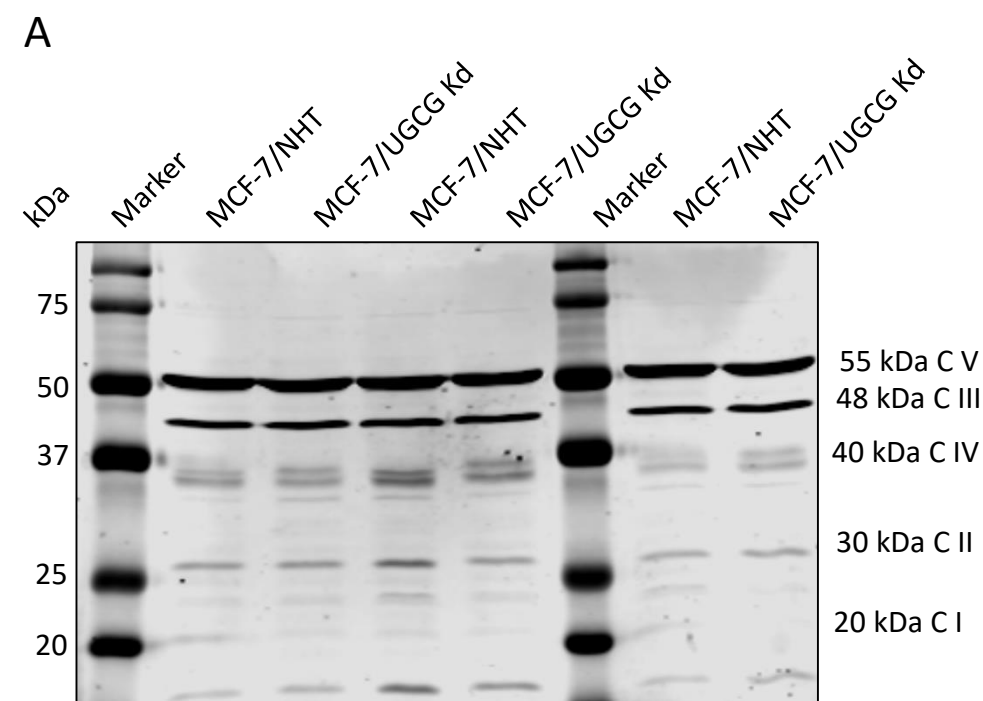
³Fraunhofer Institute for Molecular Biology and Applied Ecology IME, Project Group Translational Medicine and Pharmacology (TMP), Theodor Stern-Kai 7, 60590 Frankfurt am Main, Germany.



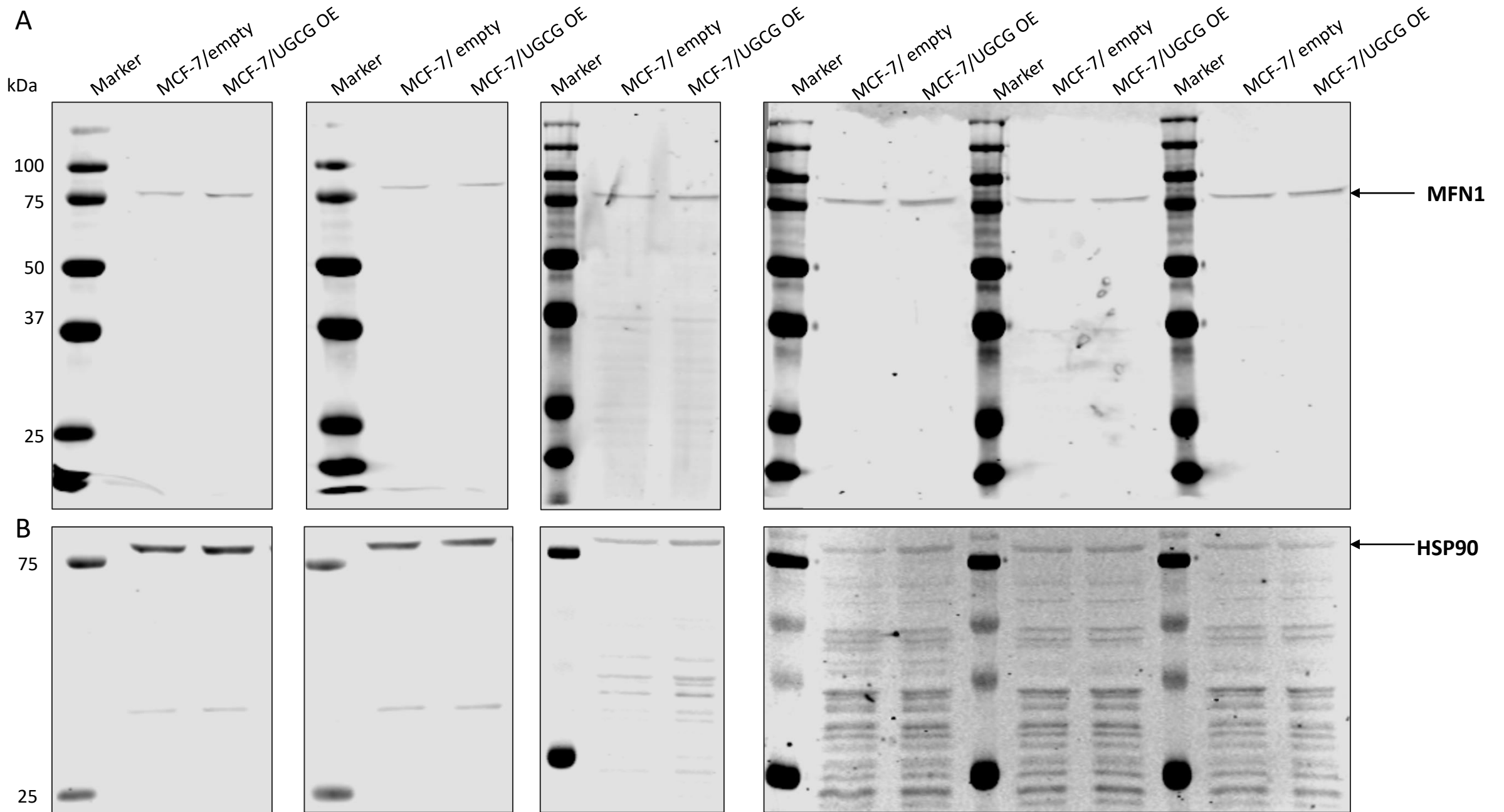
Analysis mitochondrial dynamics by Western Blot. (A) *Translocase of the outer mitochondrial membrane 20* (TOM20) protein Western Blots of all experiments. (B) *Heat shock protein 90* (HSP90) protein Western Blots of all experiments.



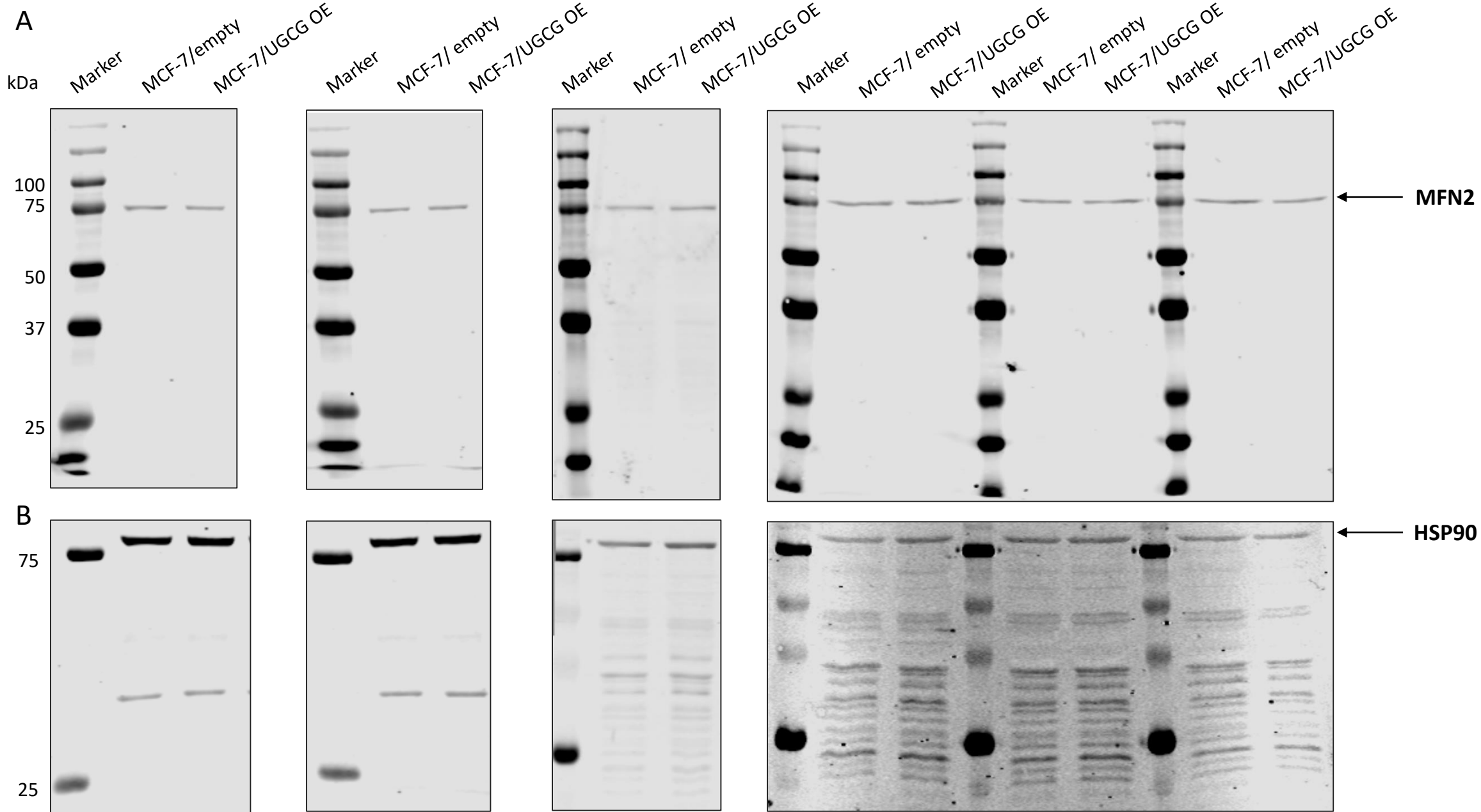
Analysis of OXPPOS protein concentrations by Western Blot. (A) OXPPOS protein Western Blots of all experiments. (B) Ponceau dye of all experiments. C = complex.



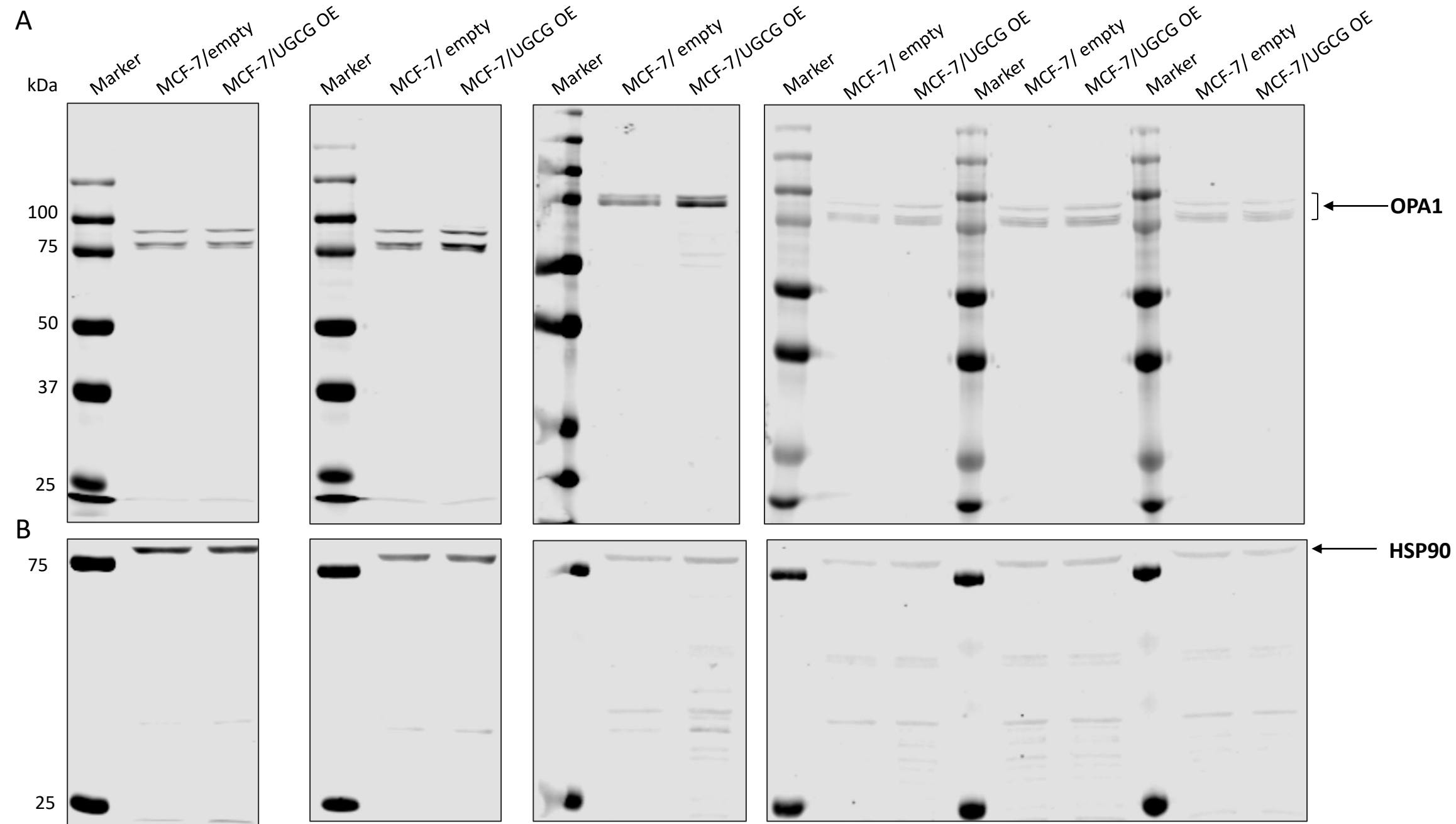
Determination of OXPHOS protein concentrations by Western Blot analysis. MCF-7/UGCG knockdown (Kd) cells were generated as described previously [6]. NHT = non-human target control cells. C = complex. (A) OXPHOS protein Western Blots of all experiments. (B) Ponceau dye of all experiments. (C) Densitometrical analysis of OXPHOS complexes I-V protein concentrations by Western blot analysis. Protein expression is related to Ponceau dye. Data are presented as a mean of $n = 3 \pm \text{SEM}$. , *** $p \leq 0.001$..



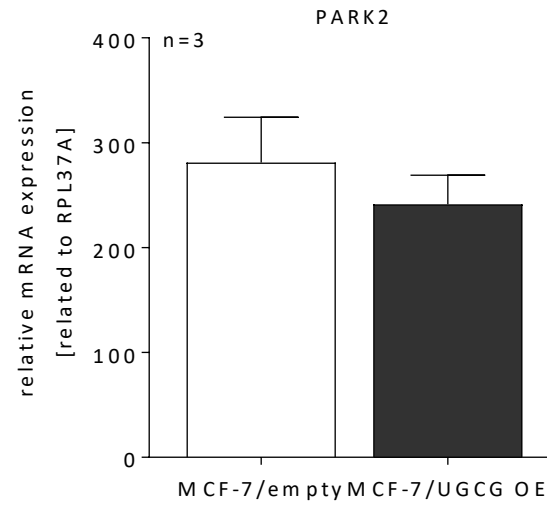
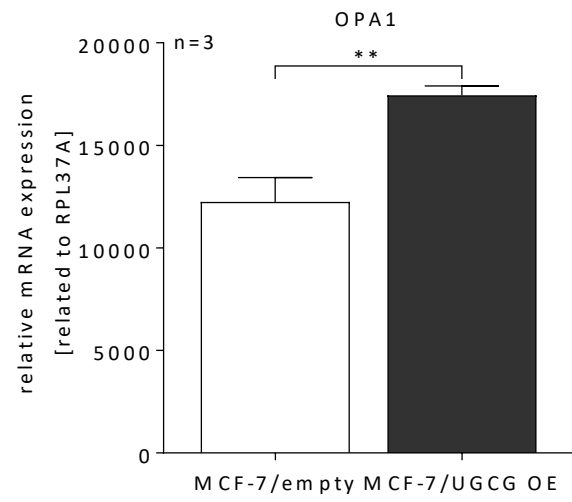
Analysis mitochondrial dynamics by Western Blot. (A) *Mitofusin 1* (MFN1) protein Western Blots of all experiments. (B) *Heat shock protein 90* (HSP90) protein Western Blots of all experiments.



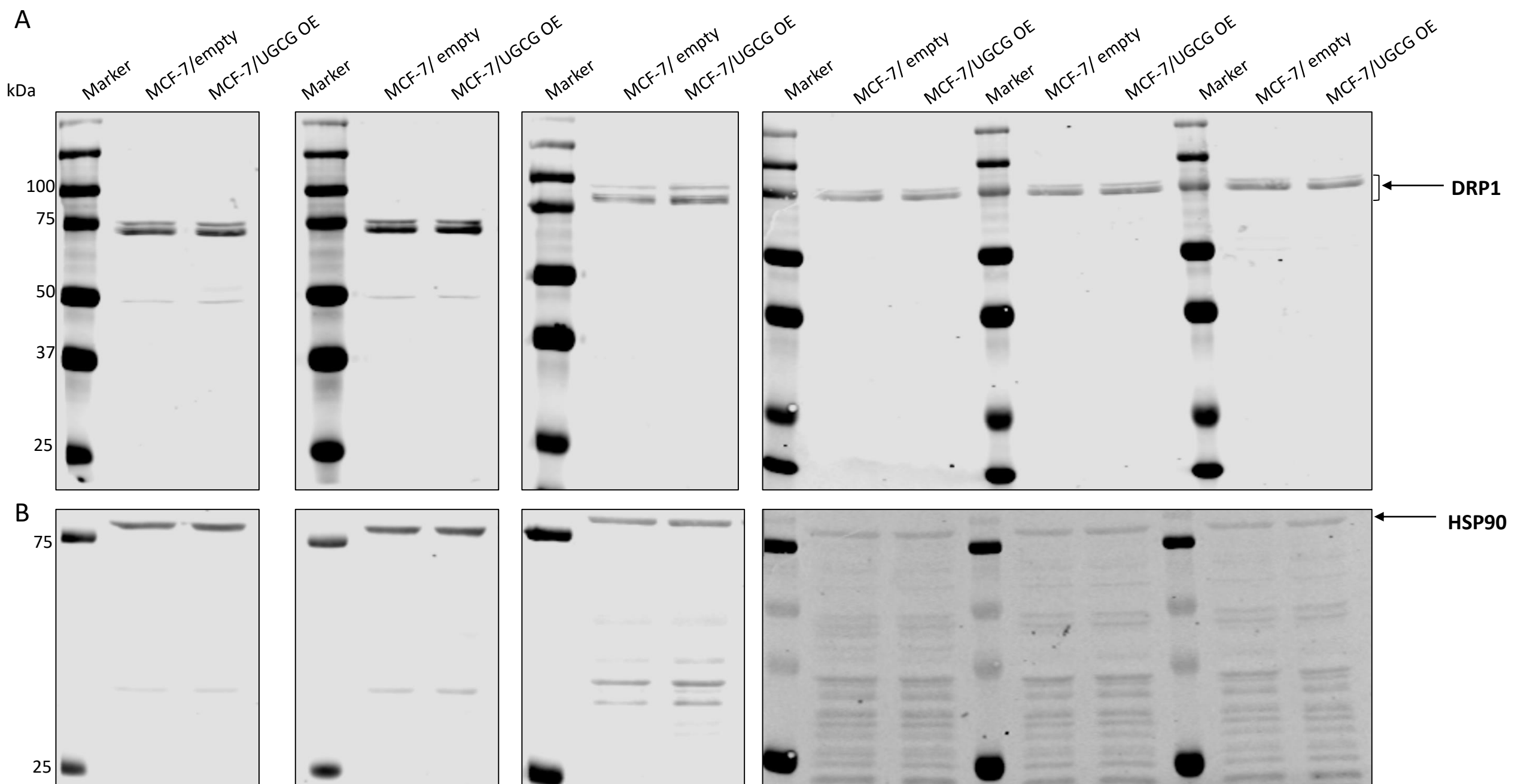
Analysis mitochondrial dynamics by Western Blot. (A) *Mitofusin 2* (MFN2) protein Western Blots of all experiments. (B) *Heat shock protein 90* (HSP90) protein Western Blots of all experiments.



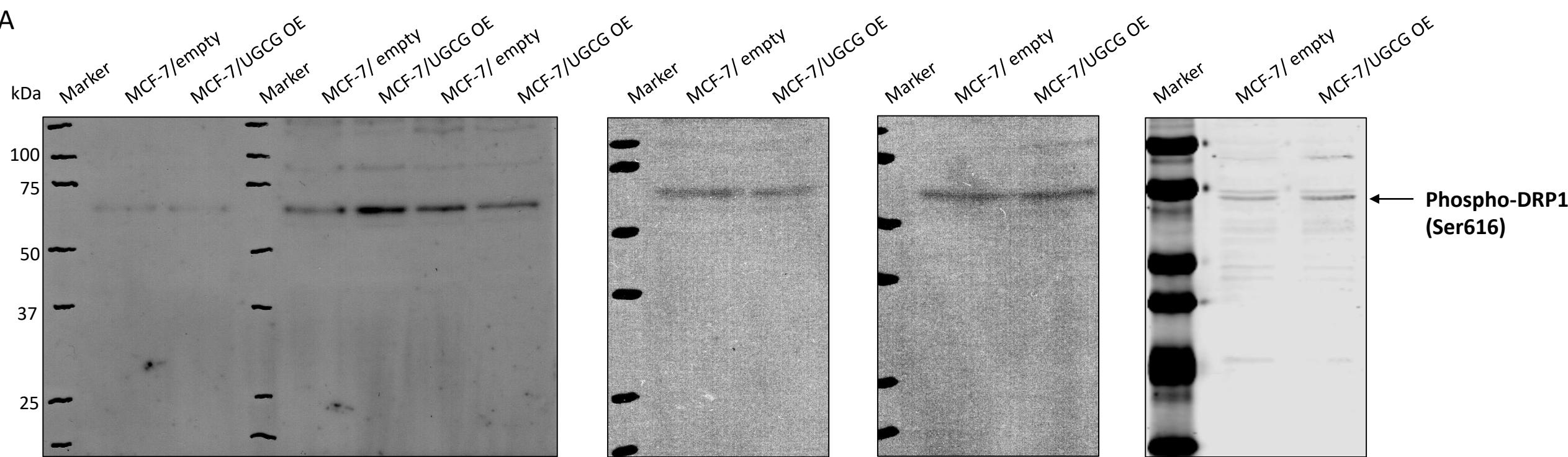
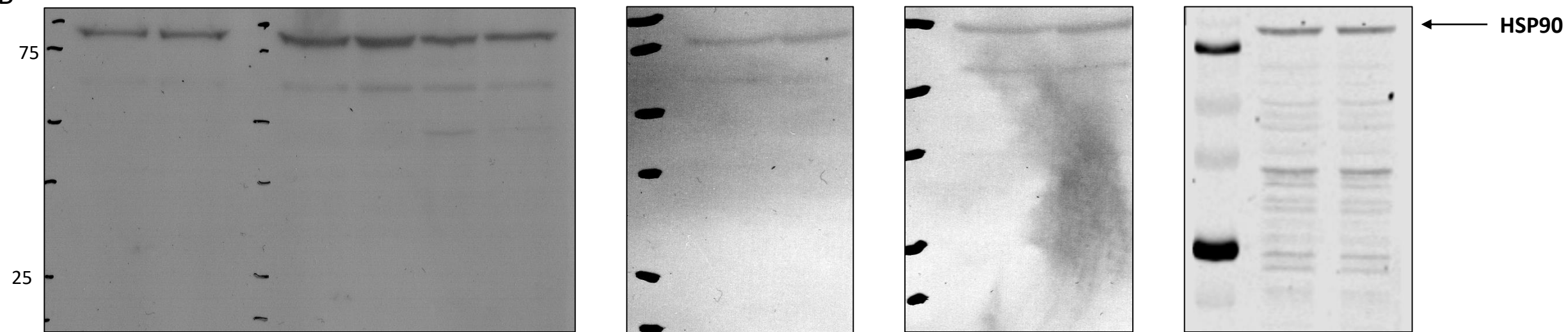
Analysis mitochondrial dynamics by Western Blot. (A) *Optic atrophy type 1* (OPA1) protein Western Blots of all experiments. (B) *Heat shock protein 90* (HSP90) protein Western Blots of all experiments.



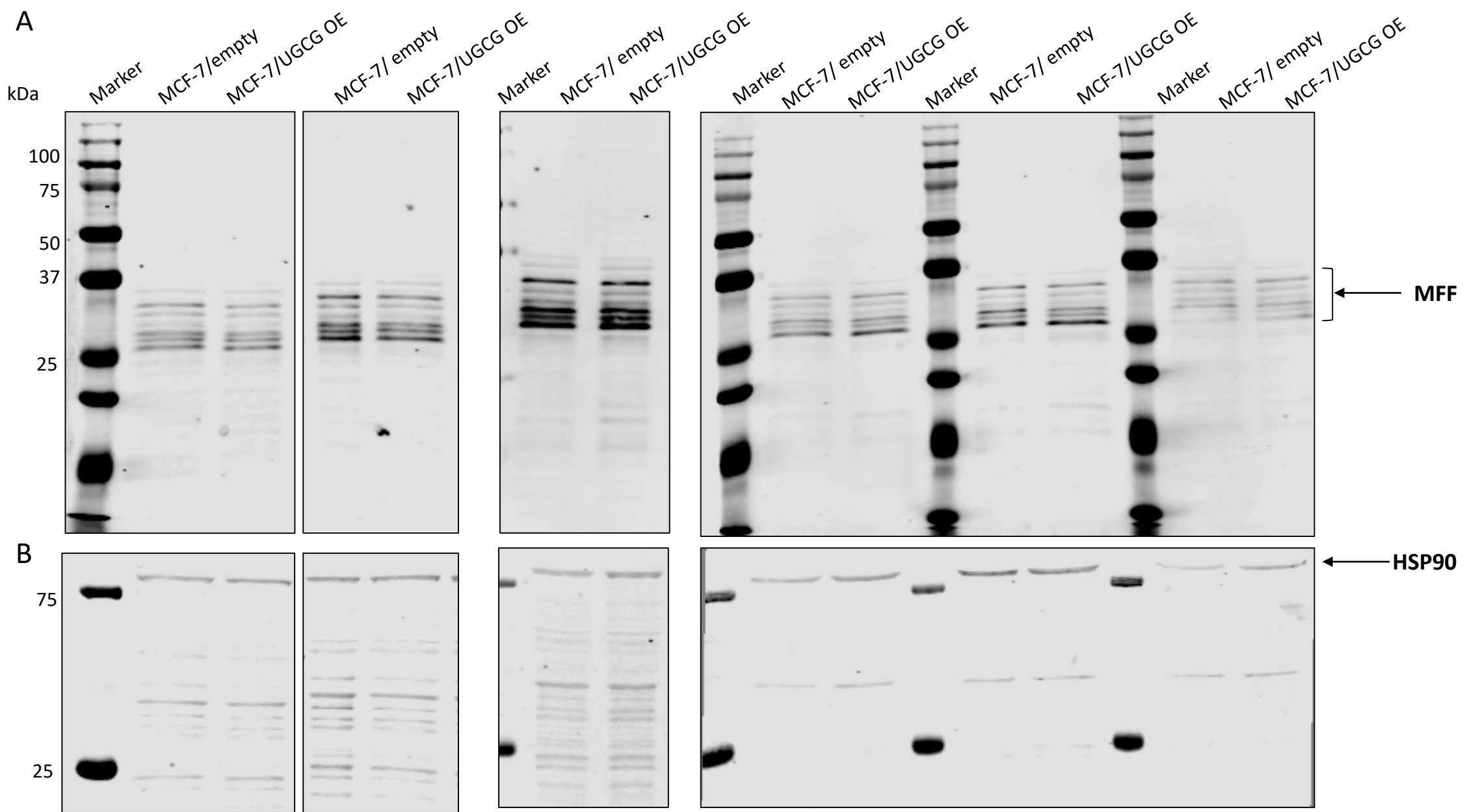
Analysis of *optic atrophy type 1* (OPA1) and *Parkin 2* (PARK2) mRNA expression by qRT-PCR. mRNA expression is related to the housekeeper RPL37A. Data are presented as a mean of $n = 3 \pm \text{SEM}$. ** $p \leq 0.01$.



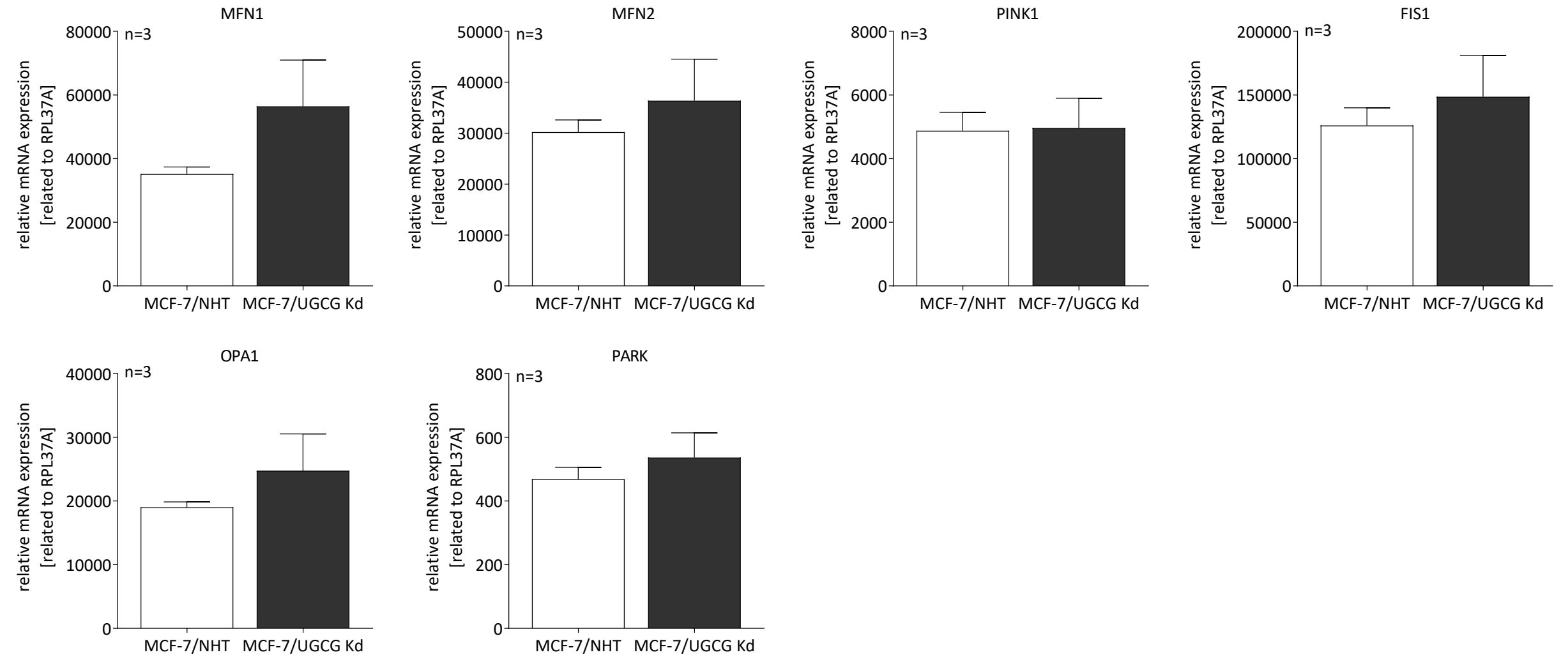
Analysis mitochondrial dynamics by Western Blot. (A) *Dynamin related protein1* (DRP1) protein Western Blots of all experiments. (B) *Heat shock protein 90* (HSP90) protein Western Blots of all experiments.

A**B**

Analysis mitochondrial dynamics by Western Blot. (A) *Phospho-dynamin related protein1* (Phospho-DRP1) (Ser616) protein Western Blots of all experiments. (B) *Heat shock protein 90* (HSP90) protein Western Blots of all experiments.

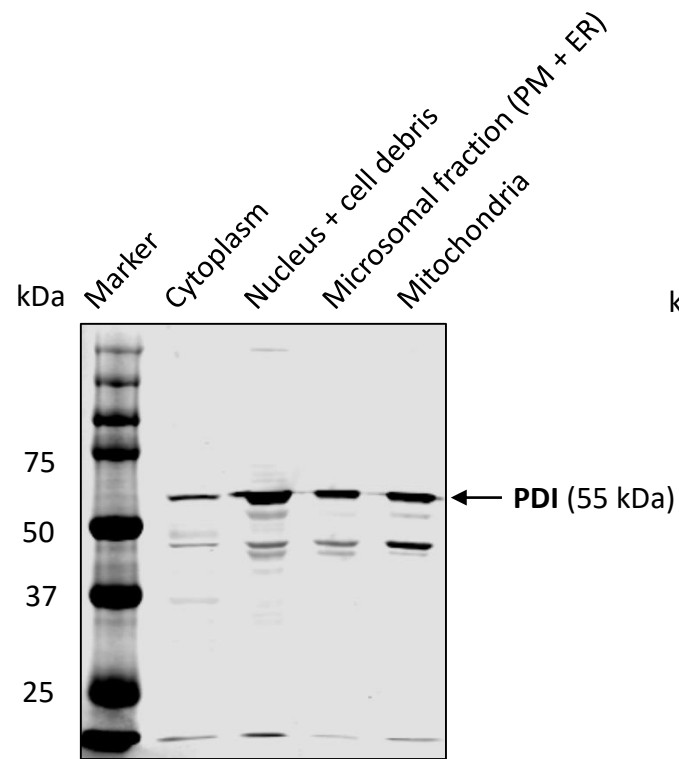


Analysis mitochondrial dynamics by Western Blot. (A) *Mitochondrial fission factor* (MFF) protein Western Blots of all experiments. (B) *Heat shock protein 90* (HSP90) protein Western Blots of all experiments.

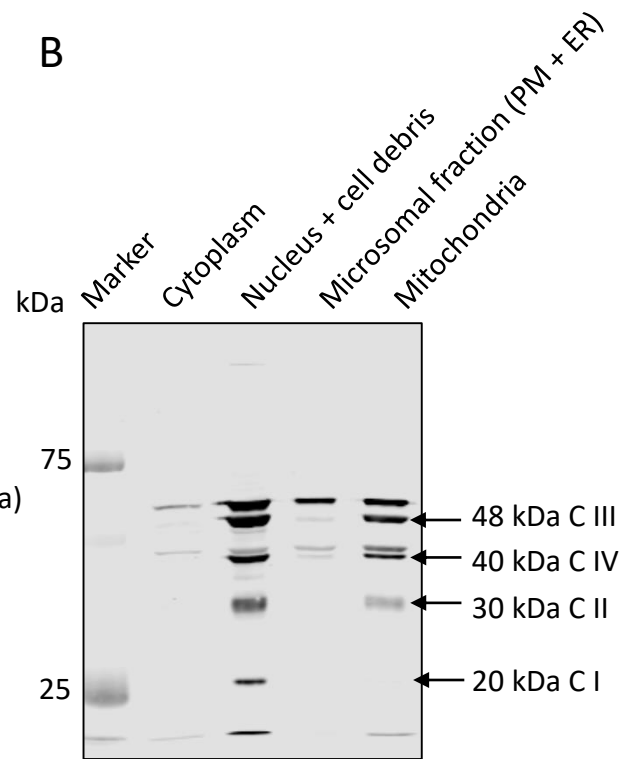


Analysis of mRNA expression of fusion and fission proteins in MCF-7/UGCG Kd cells by qRT-PCR. NHT = non-human target control, MFN1 = mitofusin 1, MFN2 = mitofusin 2, PINK1 = PTEN-induced kinase 1, FIS1 = mitochondrial fission protein 1, OPA1 = optic atrophy 1, PARK = parkin 2. mRNA expression is related to the housekeeper RPL37A. Data are presented as a mean of $n = 3 \pm$ SEM.

A

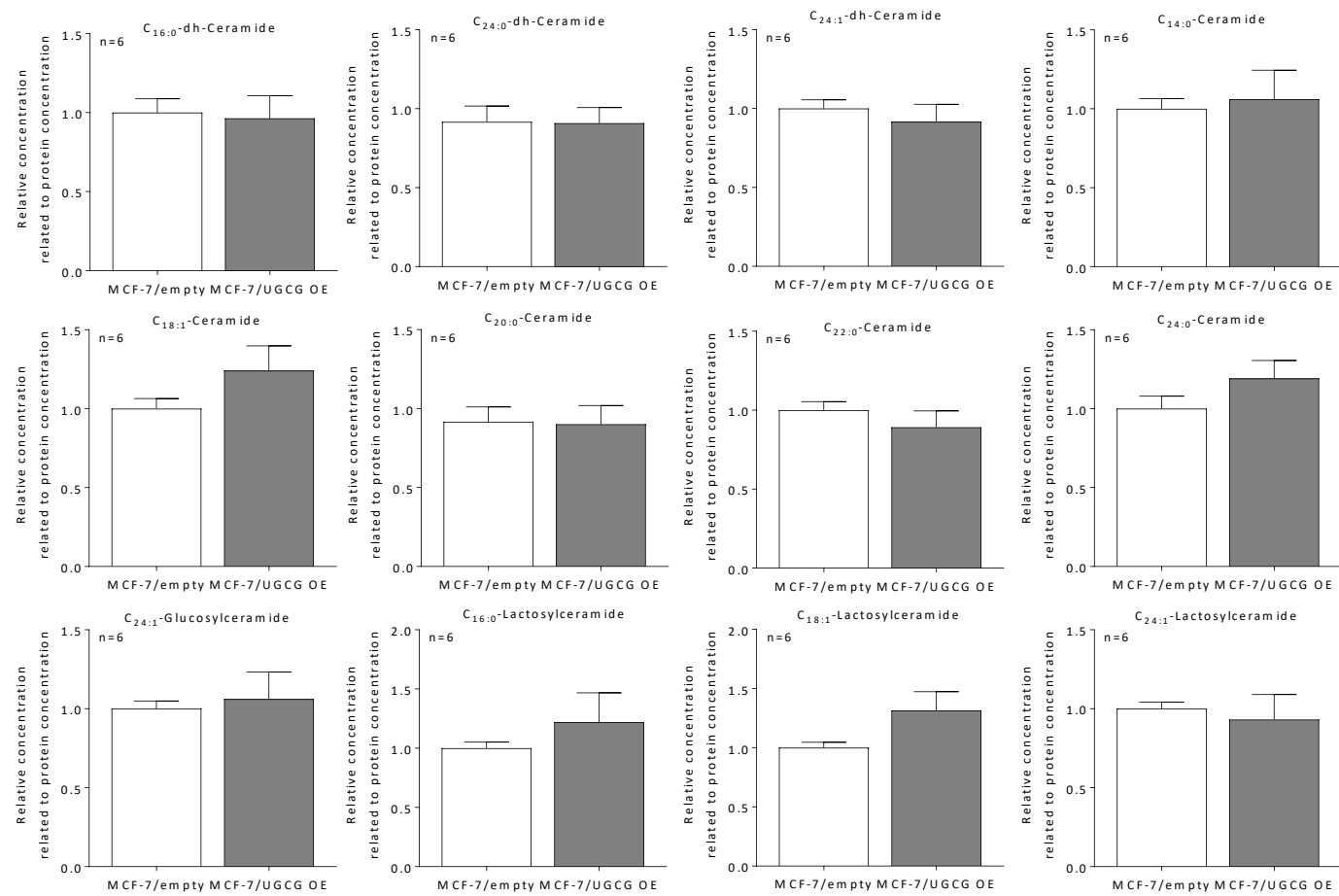


B



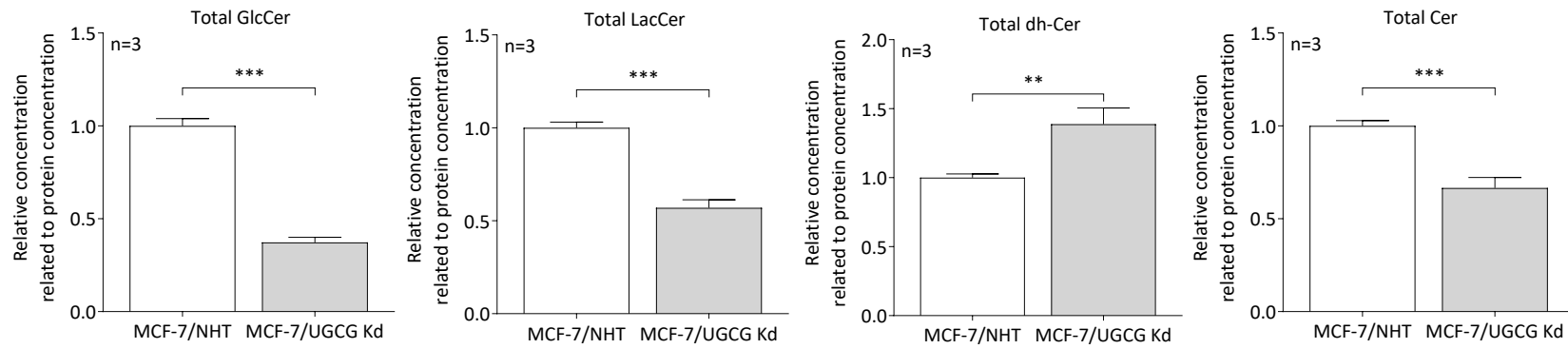
ER/mitochondria fraction verified by PDI and OXPHOS protein detection. (A) Representative image of anti-*protein disulfide isomerase* (PDI; ER marker) and (B) anti-Oxphos Rodent Western Blot Cocktail (mitochondria marker). PM = plasma membrane, ER = endoplasmic reticulum, C = Complex.

Material and Methods: Western blot analysis of purity of ER/mitochondria fractions. Isolated fractions were separated by 12 % SDS-PAGE as described in section *Protein concentration determination by Western blot analysis*. Anti-PDI antibody (ab2792, Abcam, Cambridge, UK) (1:250 in blocking solution) was applied overnight at 4 °C. The IRDye®680 conjugated secondary antibody (LI-COR Biosciences, Bad Homburg, Germany), and anti-OXPHOS rodent Western Blot Cocktail were applied as described in section *Protein concentration determination by Western blot analysis*.

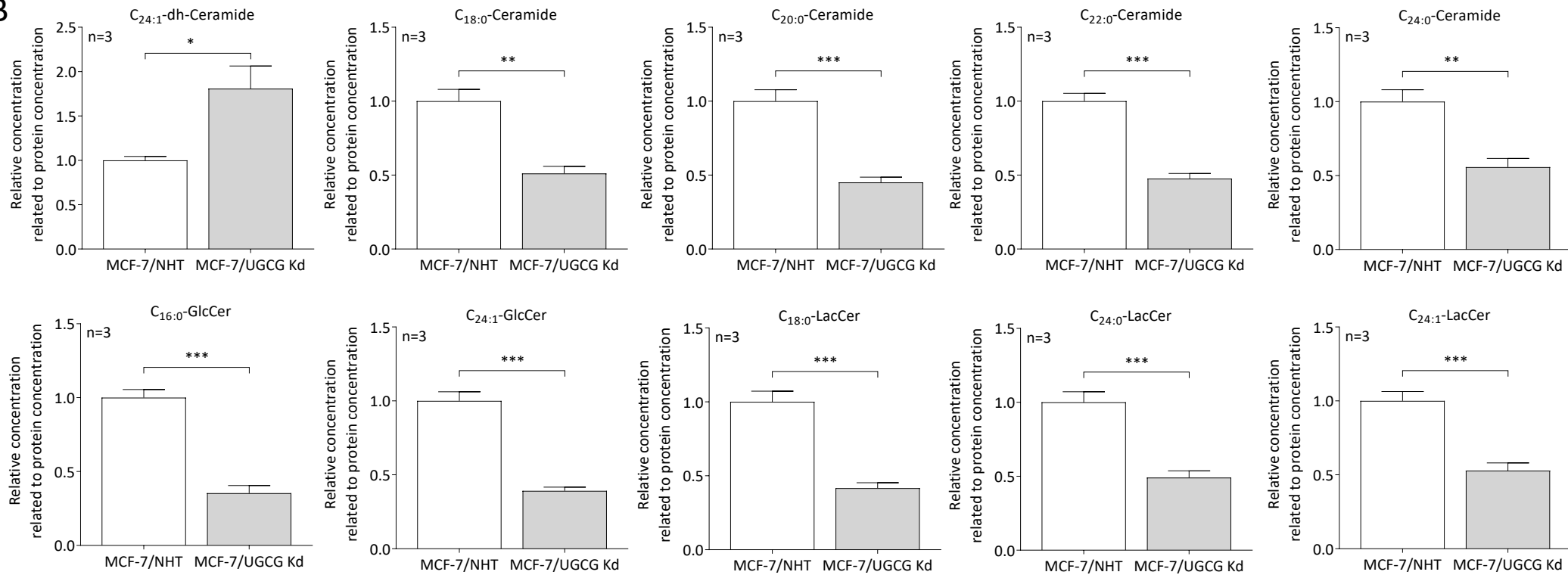


Sphingolipid levels of mitochondria/ER fractions, which were not changed following UGCG overexpression. Levels were related to MCF-7/empty. dh-Ceramide = dihydro-ceramide. Data are presented as a mean of n = 6 ± SEM.

A



B



Sphingolipid levels of mitochondria/ER fractions following UGCG Kd. NHT = non-human target control. Levels were related to MCF-7/NHT. (A) Total glucosylceramide (GlcCer), lactosylceramide (LacCer), duhydro-ceramide (dh-Cer) and ceramide levels. (B) Analytes, which were significantly altered in MCF-7/UGCG Kd cells as compared to control cells. Data are presented as a mean of $n = 3 \pm \text{SEM}$. * $p \leq 0.05$, ** $p \leq 0.01$, *** $p \leq 0.001$.