

Supplementary information

Radiation alters the cargo of exosomes released from squamous head and neck cancer cells to promote migration of recipient cells

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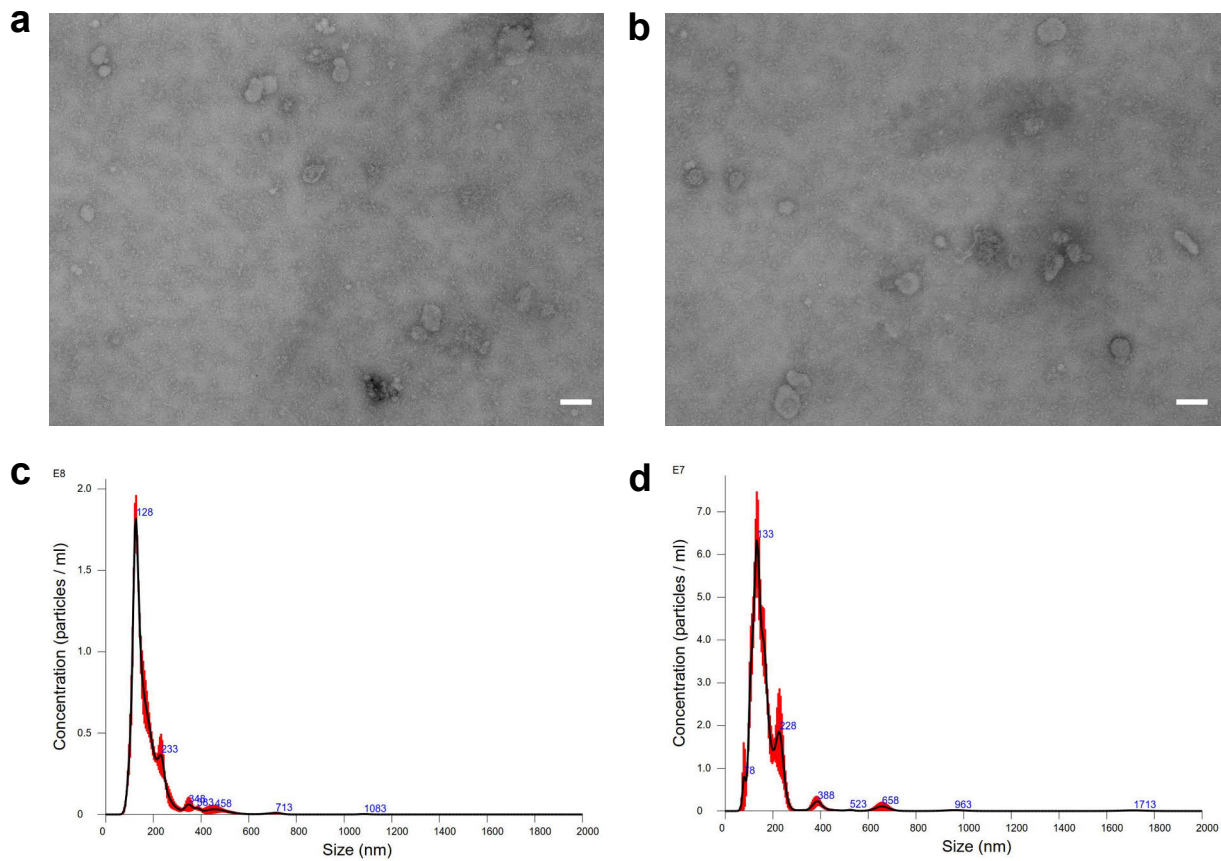
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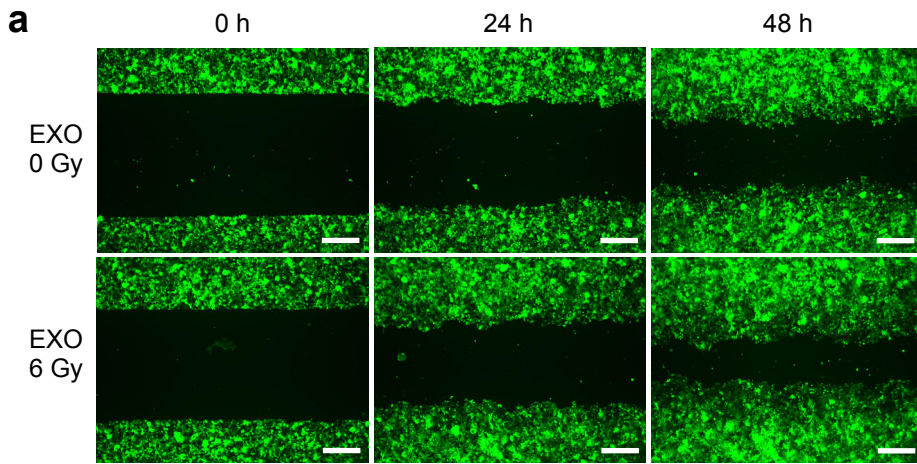
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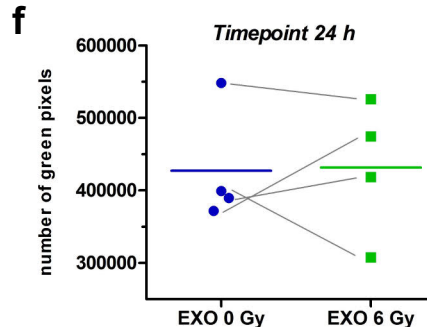
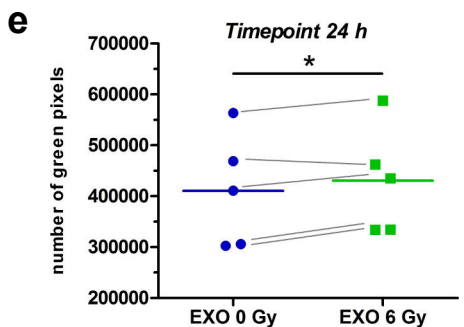
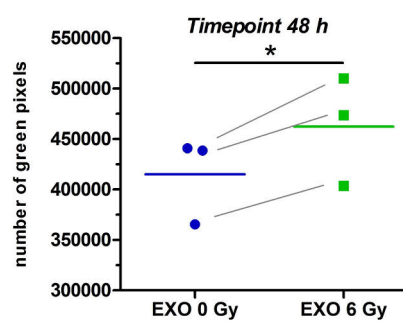
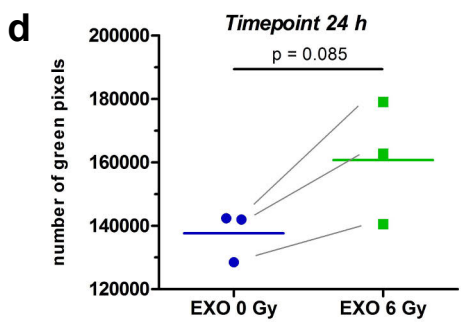
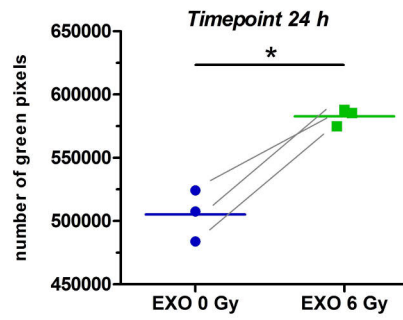
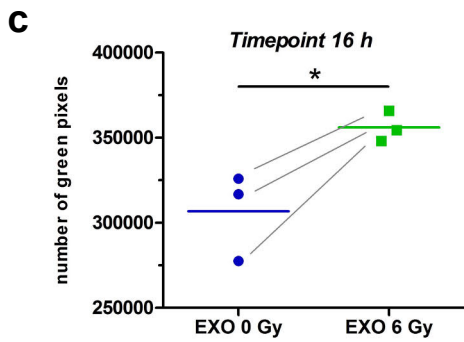
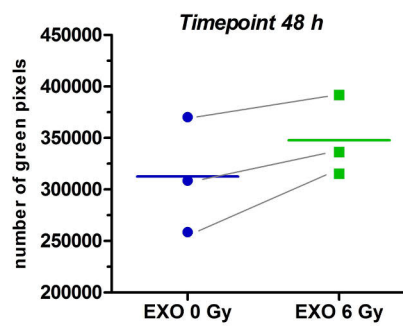
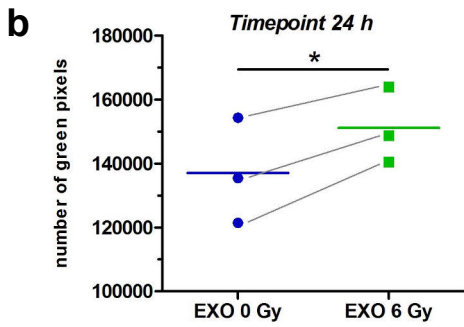


Supplementary Figure S1. Characterization of isolated HNSCC exosomes. (a) Electron microscopy of BHY (EXO 0 Gy) exosomes (scale bar: 100 nm). **(b)** Electron microscopy of FaDu (EXO 0 Gy) exosomes (scale bar: 100 nm). **(c)** Size distribution of BHY (EXO 0 Gy) exosomes measured with NanoSight technology. **(d)** Size distribution of FaDu (EXO 0 Gy) exosomes measured with NanoSight technology.

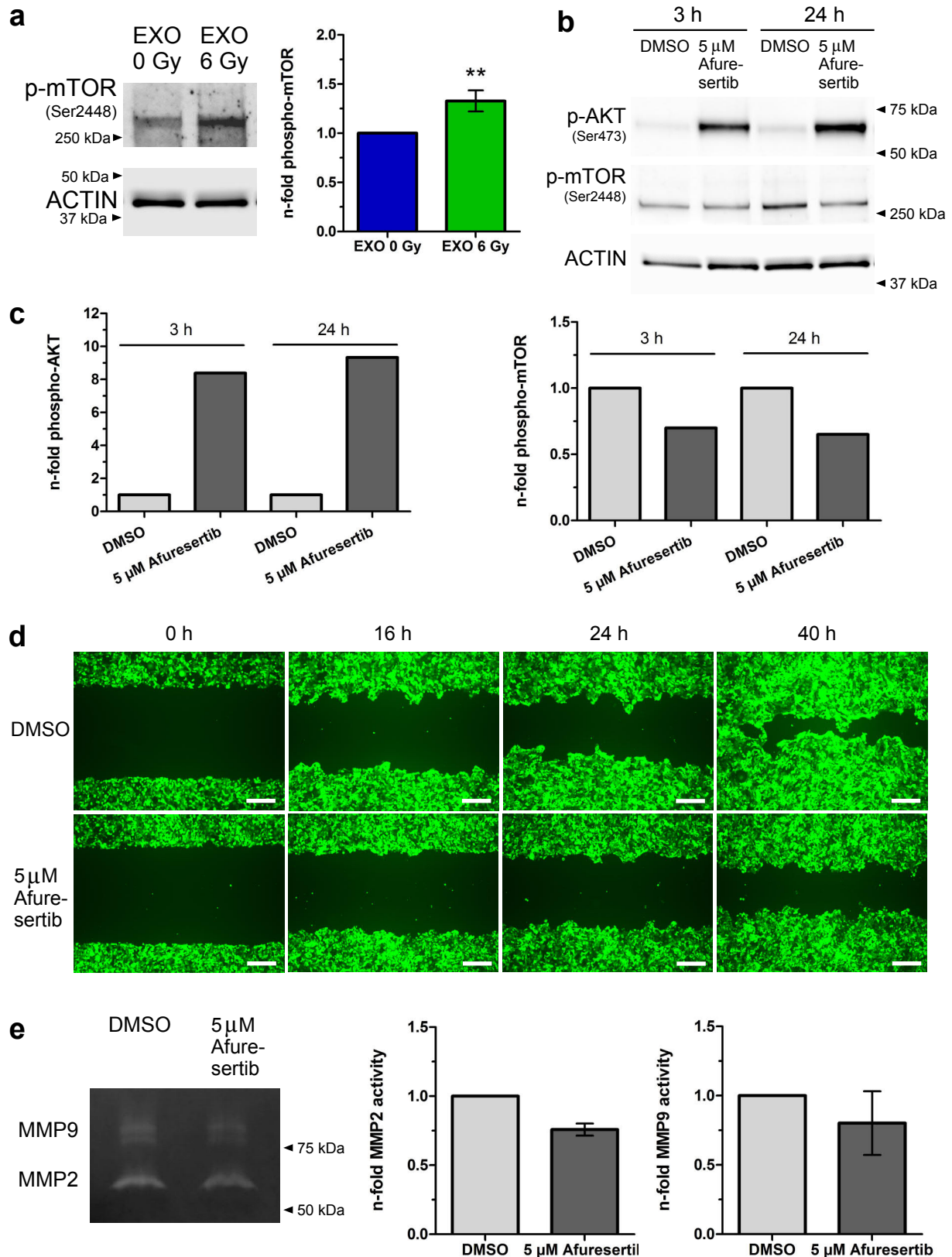


Supplementary Figure S2. Effect of exosomes isolated after irradiation of tumour and non-tumour cell lines on HNSCC migration. (a, b)

Exemplary wound healing of FaDu-GFP cells after 24 and 48 hours (scale bar: 500 μ m). Cells were either preincubated with exosomes from non-irradiated (EXO 0 Gy; upper row) or from 6 Gy irradiated FaDu cells (EXO 6 Gy; lower row).

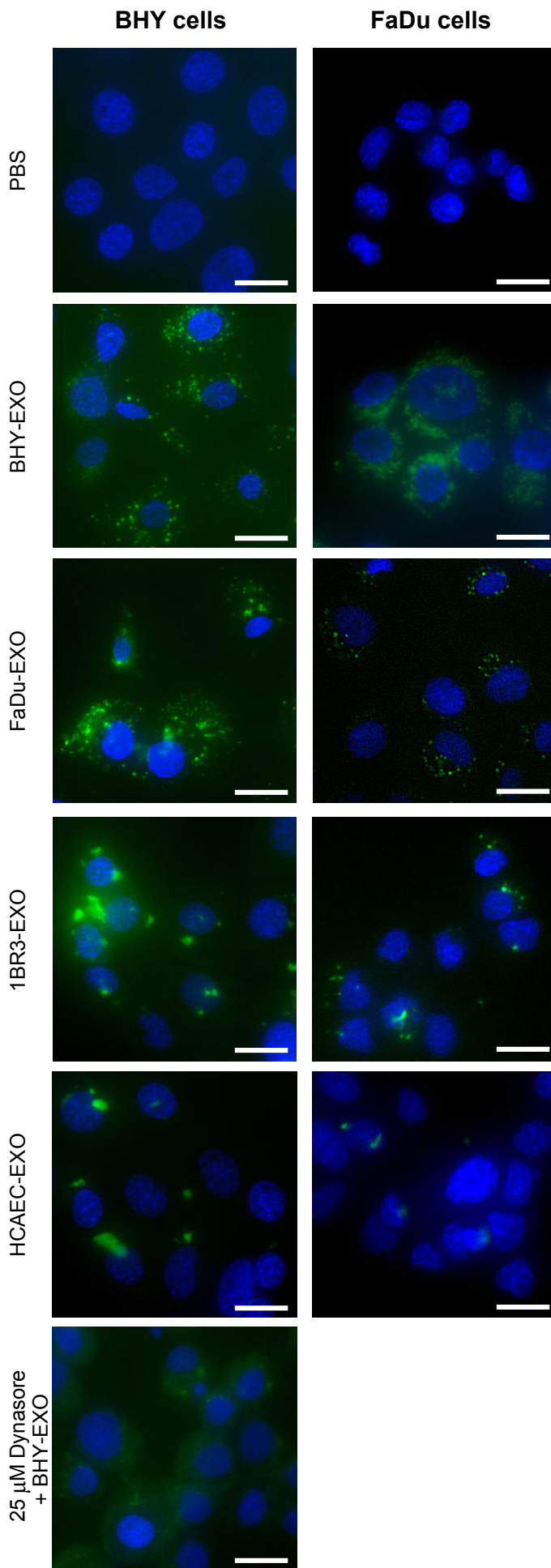


(c) BHY-GFP cells were either preincubated with exosomes from non-irradiated (EXO 0 Gy) or 6 Gy (EXO 6 Gy) irradiated FaDu cells. (d) FaDu-GFP cells were either preincubated with exosomes from non-irradiated (EXO 0 Gy) or 6 Gy (EXO 6 Gy) irradiated BHY cells. (e) BHY-GFP cells were either preincubated with exosomes from non-irradiated (EXO 0 Gy) or 6 Gy (EXO 6 Gy) irradiated IBR3 cells. (f) BHY-GFP cells were either preincubated with exosomes from non-irradiated (EXO 0 Gy) or 6 Gy (EXO 6 Gy) irradiated HCAEC cells. Quantification of the wound healing capacity with the Image Colour Analyser after 16, 24 and 48 hours [n \geq 3; two-sided, paired t-test; p-value < 0.05].



Supplementary Figure S3. Effect of the AKT-inhibitor Afuresertib on molecular targets

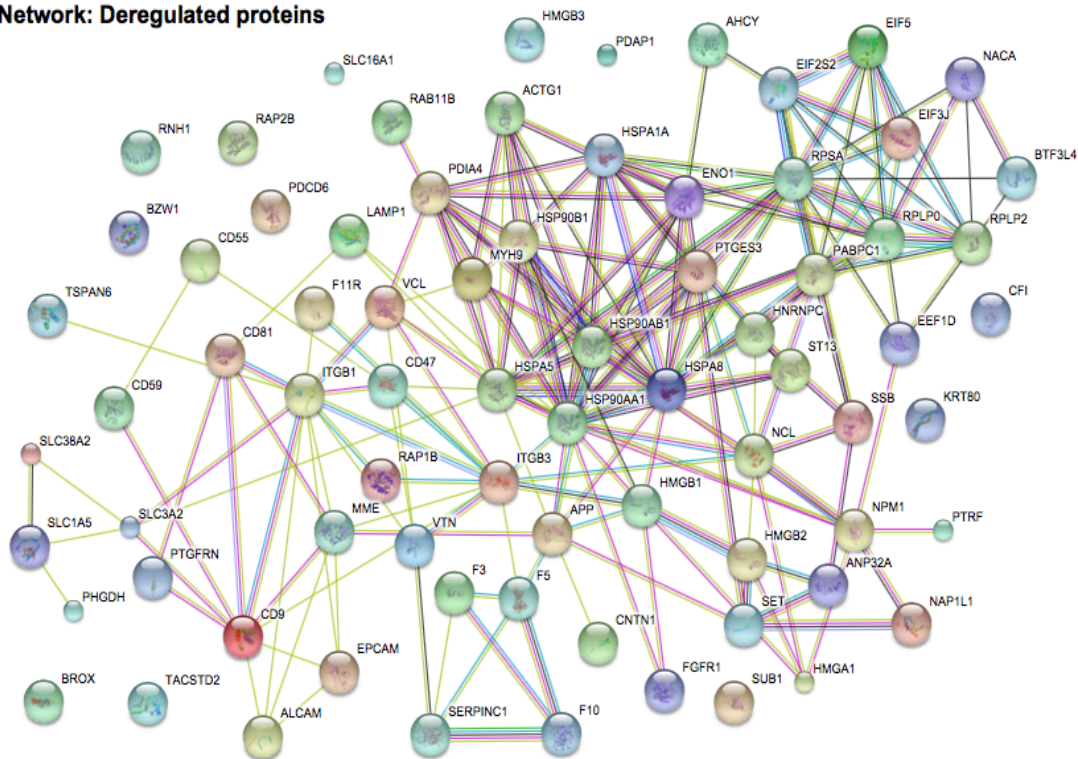
and migration. (a) Western blot of phospho-mTOR (Ser2448) of cells which were incubated for 3 hours with exosomes isolated either from irradiated cells (EXO 6 Gy) or from non-irradiated cells (EXO 0 Gy). Normalization was performed to ACTIN and to cells treated with exosomes from non-irradiated cells (EXO 0 Gy). Cropped blots are displayed [n = 4; \pm SD; two-sided, one-sample t-test; p-value < 0.01]. **(b, c)** Western blot of phospho-AKT (Ser473) and phospho-mTOR (Ser2448) of cells which were treated with 5 μ M Afuresertib or DMSO for 3 or 24 hours. Normalization was performed to ACTIN and to control cells treated with DMSO. Cropped blots are displayed [n = 1]. **(d)** Exemplary wound healing of BHY-GFP cells after treatment with 5 μ M of the AKT-inhibitor Afuresertib, or with the equal volume of DMSO. The pictures were taken 16, 24 and 40 hours after migration start (scale bar: 500 μ m). **(e)** MMP2 and MMP9 activity in the supernatants 24 hours after treatment of BHY cells with 5 μ M Afuresertib or DMSO. Normalization was performed to control cells treated with DMSO. Cropped gels are displayed [n = 2; \pm SD].



Supplementary Figure S4. Exosomes from tumour and non-tumour cells transfer proteins to recipient BHY and FaDu cells. Exosomal proteins of BHY (EXO-BHY), FaDu (EXO-FaDu), 1BR3 (EXO-1BR3), HCAEC (EXO-HCAEC) cells and PBS as negative control were stained with CFSE and subsequently transferred onto recipient BHY and FaDu cells. In addition BHY cells were pretreated with 25 μ M Dynasore for 1 hour with subsequent transfer of BHY exosomes (25 μ M Dynasore + BHY-EXO). The protein uptake was monitored after 24 hours of exosome exposure (scale bar: 25 μ m).

Process analysis with String

Network: Deregulated proteins



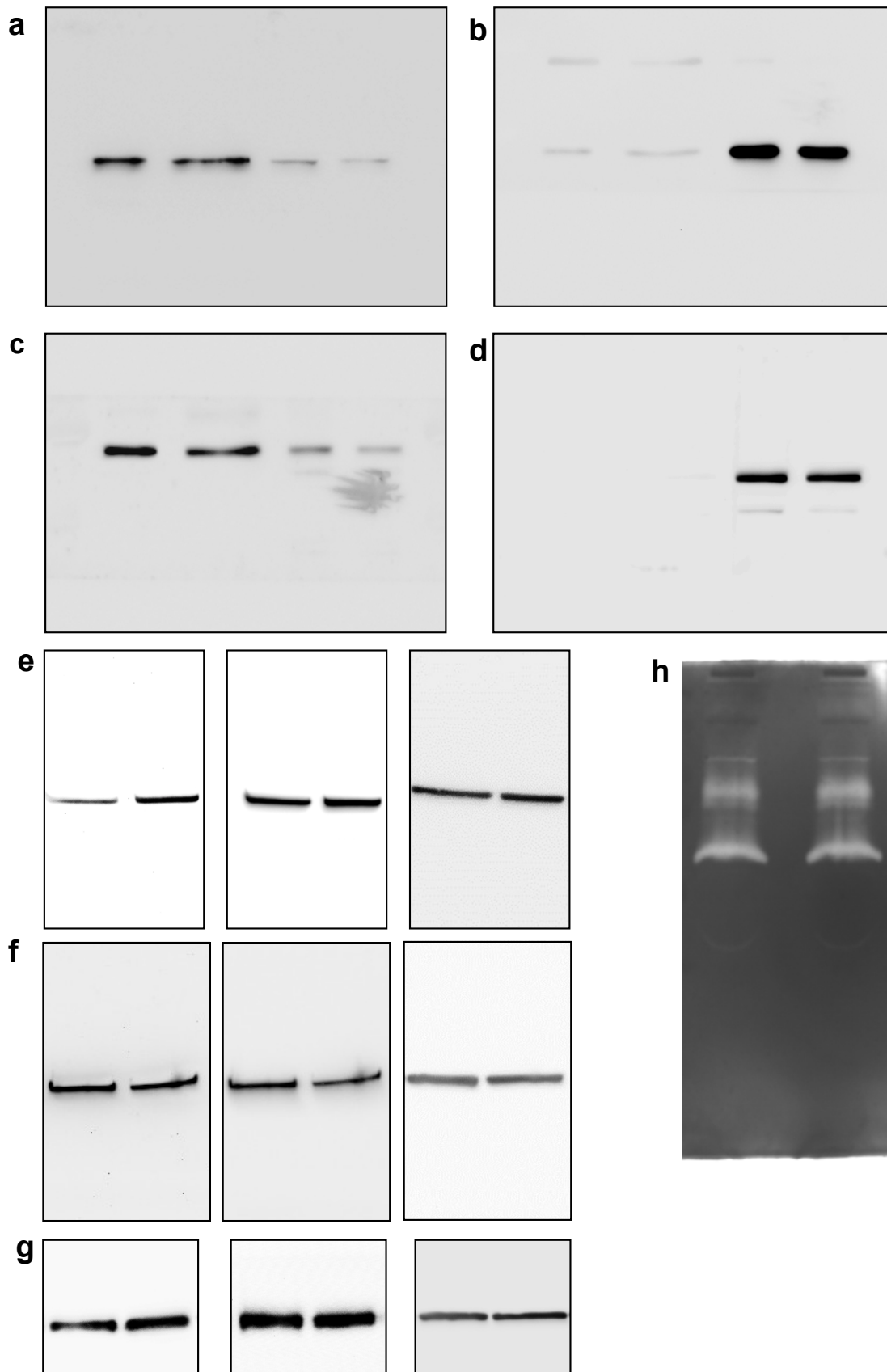
Edge legend

<ul style="list-style-type: none"> ■ from curated databases ■ experimentally determined ■ gene neighborhood ■ gene co-occurrence ■ textmining ■ co-expression ■ protein homology 	<ul style="list-style-type: none"> } Known interactions } Predicted interactions
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Network Stats

number of nodes	75
number of edges	196
average node degree	5.23
avg. local clustering coefficient	0.474
expected number of edges	77
PPI enrichment p-value	0

Supplementary Figure S5. Interaction network of radiation-regulated exosomal proteins. *In silico* STRING analysis of deregulated exosomal proteins to determine protein interaction. The minimum required interaction score was set to medium confidence (0.004) and the analysis revealed a high degree of interaction indicated by the PPI enrichment p-value $< 1 \times 10^{-15}$.



Supplementary Figure S6. Full-length western blots and gels. (a) Western blot Alix (Fig 1 b) (b) Western blot GAPDH (Fig 1 b) (c) Western blot TSG101 (Fig 1 b) (d) Western blot Calnexin (Fig 1 b) (e) Western blot phospho-mTOR (Ser2448), mTOR and Actin (Fig 4 a) (f) Western blot phospho-mTOR (Ser2448), mTOR and Actin (Fig 4 b) (g) Western blot phospho-rpS6 (Ser240/244), rpS6 and Actin (Fig 4 c) (h) Zymography (Fig 4 d).