## IGF-1 contributes to the expansion of melanoma-initiating cells through an epithelial-mesenchymal transition process

**Supplementary Materials** 



**Supplementary Figure S1: IGF-1 signaling pathways are attenuated in the C10 clone.** (A) Levels of phosphorylated AKT and ERK were analyzed in the C10 clone, B16-F10<sup>CT</sup> and B16-F10<sup>WT</sup> cells, by immunoblotting. The phosphorylation signals were quantified as pAKT and pERK versus total  $\beta$ -actin, and ERK. Data are expressed as the mean  $\pm$  SEM of 4 independent experiments. C10 cells had lower levels of phosphorylated forms of AKT (0.48  $\pm$  0.05 versus 0.96  $\pm$  0.11 and 1  $\pm$  0, p < 0.05, n = 4) and ERK-1/2 MAPK (0.45  $\pm$  0.07 versus 1.34  $\pm$  0.39 and 1  $\pm$  0, p < 0.05, n = 4) than B16-F10CT and B16-F10WT cells. (B) IGF-1R expression was analyzed in the total lysates of B16-F10WT and B16-F10CT cells and of the four IGF-1dull clones (A6, C10, E11, F9), by western blotting.



Supplementary Figure S2: IGF-1R neutralization decreases the expression of EMT and stemness markers in B16-F10<sup>CT</sup> and B16-F10<sup>WT</sup> cells. Flow cytometry analysis of the CD44 and CD29 mesenchymal markers and of the Oct-3/4 and SOX2 transcription factors in the C10 clone, and in B16-F10<sup>CT</sup> and B16-F10<sup>WT</sup> cells with and without neutralizing IGF-1R antibody (0.2  $\mu$ g/mL) treatment for 48 h.



Supplementary Figure S3: IGF-1R neutralization decreases the drug efflux pump activity of B16-F10<sup>CT</sup> and B16-F10<sup>WT</sup> cells. (A) Side population (SP) analysis. C10, B16-F10<sup>CT</sup> and B16-F10<sup>WT</sup> cells with and without neutralizing IGF-1R antibody (0.2  $\mu$ g/mL) treatment for 48 h were stained with Hoechst 33342 dye in the presence or absence of Ko143 (1  $\mu$ M) and the SP population was analyzed by flow cytometry. (B) Analysis of the intracellular accumulation of MIT in B16-F10<sup>WT</sup>, B16-F10<sup>CT</sup> and C10 cells with and without neutralizing IGF-1R antibody treatment for 48 h. Mitoxantrone efflux and ABCG2 activity were determined by flow cytometry, by measuring mitoxantrone accumulation in the presence of the ABCG2 inhibitor Ko143 (1  $\mu$ M). Percentages of cells displaying MIT efflux are shown at the top left of each histogram.



Supplementary Figure S4: ABCG2 protects B16-F10<sup>WT</sup> and B16-F10<sup>CT</sup> from MIT-induced cell death. Cells were treated with 2  $\mu$ M mitoxantrone in the presence or absence of the ABCG2 inhibitor Ko143 (1  $\mu$ M) and cell viability was determined 48 h later in the FDA assay. Data are expressed as the mean ± SEM. The addition of Ko143 to the culture medium greatly increased cell death in B16-F10CT (48.5 ± 2.5% to 65.0 ± 2.0%, *p* < 0.05, *n* = 3) and B16-F10WT (36.5 ± 7.5% to 72.0 ± 6.0%, *p* < 0.05, *n* = 3) cells treated with 2  $\mu$ M MIT, whereas it had no significant impact on the death of C10 cells.