Supplementary materials: Targeted Therapy-Resistant Melanoma Cells Acquire Transcriptomic Similarities with Human Melanoblasts

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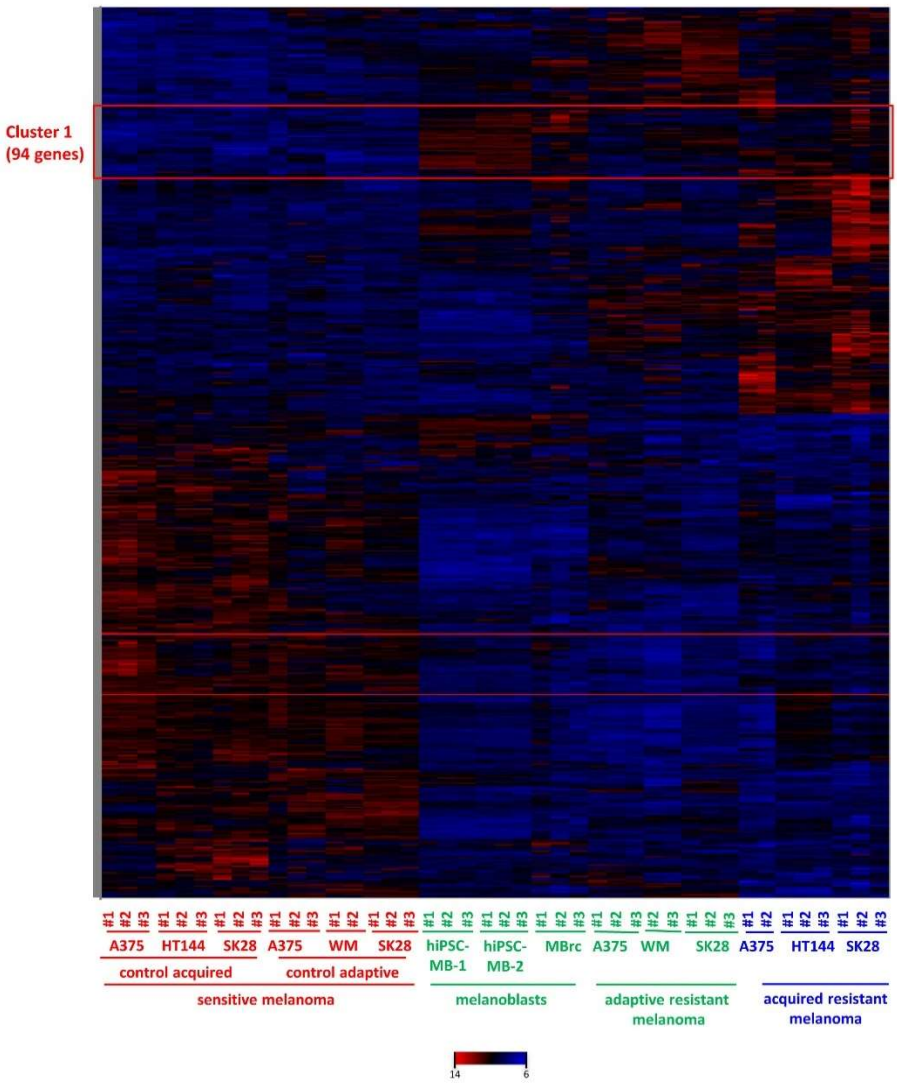

Figure S1. Gene clustering from melanoblast samples, sensitive, adaptive and acquired resistant melanoma samples. Cluster 1 represents upregulated gene in all melanoblast samples and all resistant melanoma samples.

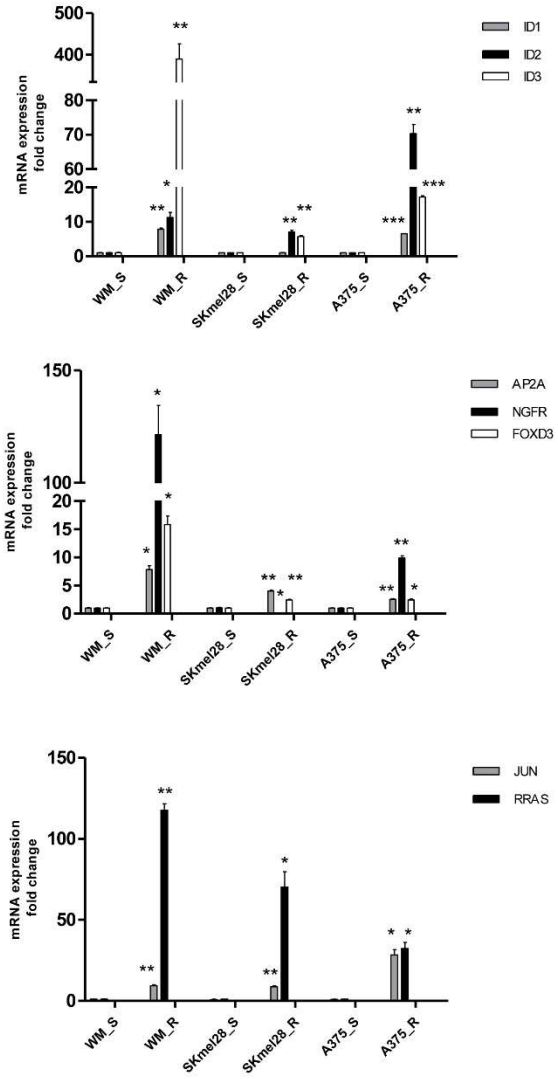

Figure S2: Real-time qPCR analysis of candidate genes' expression: *ID1, ID2, ID3, FOXD3, NGFR, JUN,* and *RRAS*, in sensitive and resistant melanoma cell lines A375, SKmel28, and WM266-4 Data represent a mean of three independent experiments ± SEM.