

Additional File 6: Search for the expression profile of the genes identified in the RGP and Met libraries in the proteomics and SAGE analyses performed by de Souza et al. [93]. The table lists the genes represented in the RGP and Met libraries that were detected as differentially expressed between non-tumorigenic and tumorigenic murine melanocytic cell lines in the above cited study.

Table S7: Genes from the SSH libraries also detected as differentially expressed between non-tumorigenic and tumorigenic murine melanocytic cell lines in the proteomics and SAGE analyses performed by de Souza et al. [93]*

		RGP library genes	Met Library genes
SAGE	up-regulated in the nontumoral melanocytic cell	ATF4	CALU EIF1AY TYRP1
	up-regulated in the melanoma cell line TM1	RPS27A LSM5 ACTG1	HSPA5 NPM1 DDX5
Proteomics	up-regulated in the melanoma cell lines TM1 and TM5	ACTB	NPM1

* The reference number corresponds to the number of the reference list of the main text.