Supplementary Figure S6. (A, B) Correlations between the indicated mRNA as determined using the co-expression tool in cBioPortal in the Skin Cutaneous Melanoma dataset (TCGA, PanCancer Atlas, 448 patients) or the Metastatic Melanoma dataset (DFCI, Science 2015, 110 patients). Data are represented by scatter plots showing the Spearman's correlation.