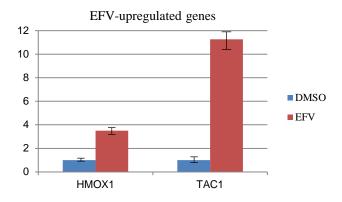
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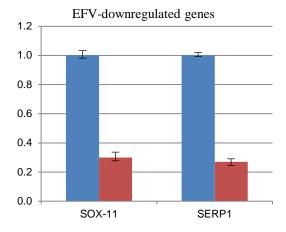


Figure S1: Real Time PCR analysis of gene expression in EFV-treated A-375 cells. The histograms represent the fold-increase or decrease in mRNA levels for the indicated genes, selected from the list in Additional file 1, in EFV-treated cells relative to levels measured in the DMSO samples (set as reference). Bars represent the standard error of the mean expression level and delimit the calculated maximum and minimum level of expression. Triplicates for each sample were analyzed in two independent experiments.

