Supplementary Figure 6. MiR-1 targets HGF and VEGFA and these genes are up-regulated in metastases from SBNETs. Using the miRanda-mirSVR and TargetScan prediction algorithms, we confirmed our Gene Ontology (GO) analyses that miR-1 regulates (A) HGF and (B) VEGFA. We also found both these genes significantly up-regulated in metastases compared to primary SBNETs in previous gene profiling (GSE27162). P values were calculated using one-way analysis of variance (ANOVA) to compare gene levels between groups followed by Tukey’s multiple comparison tests. Scatterplots are shown for each group and the horizontal lines represent the mean gene expression level and SD (**P<0.01, ***P<0.001).