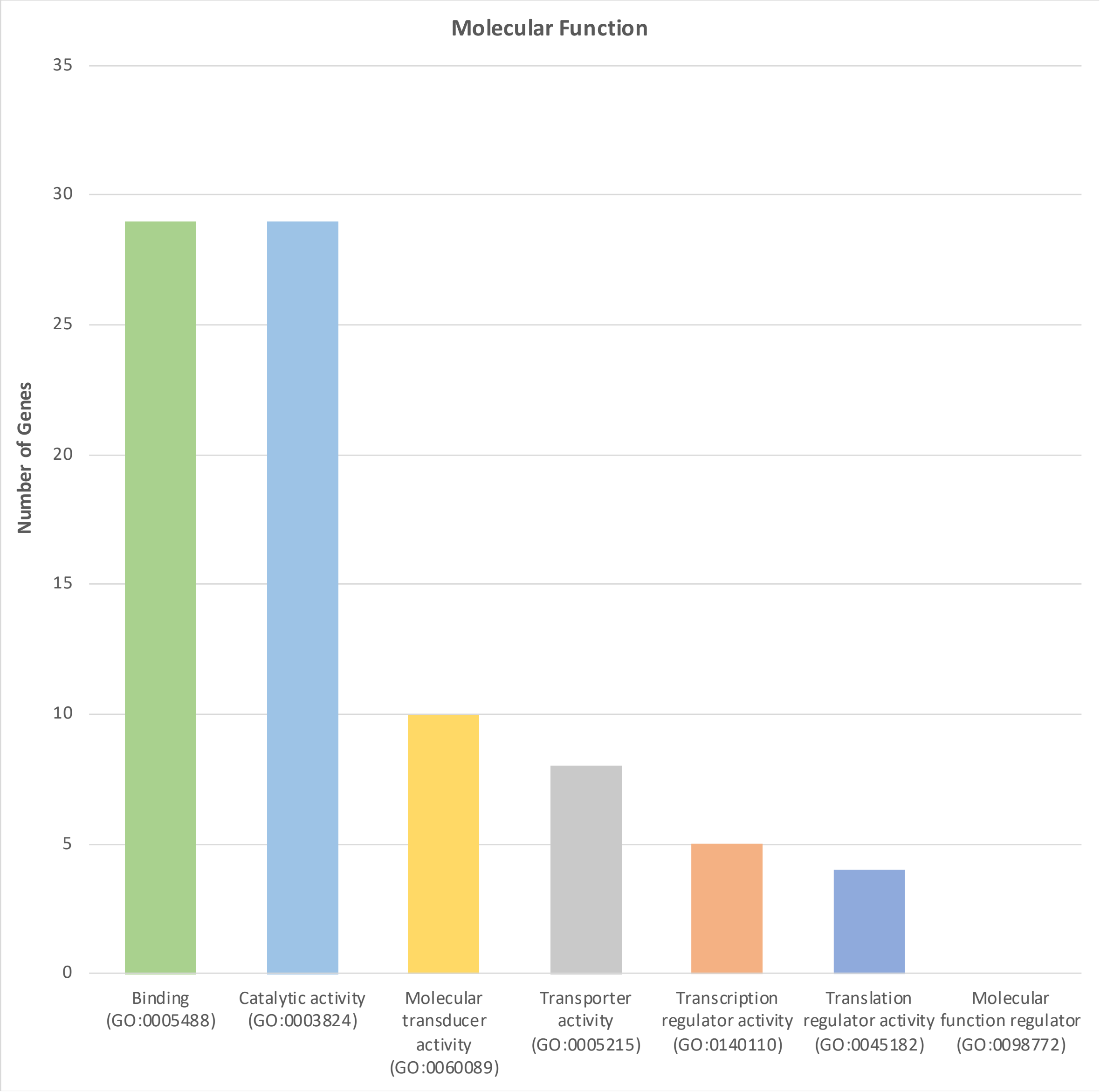

Figure S1. "Molecular Function" GO enrichment of the 147 miRNA-targeted genes identified through miRCancerdb.

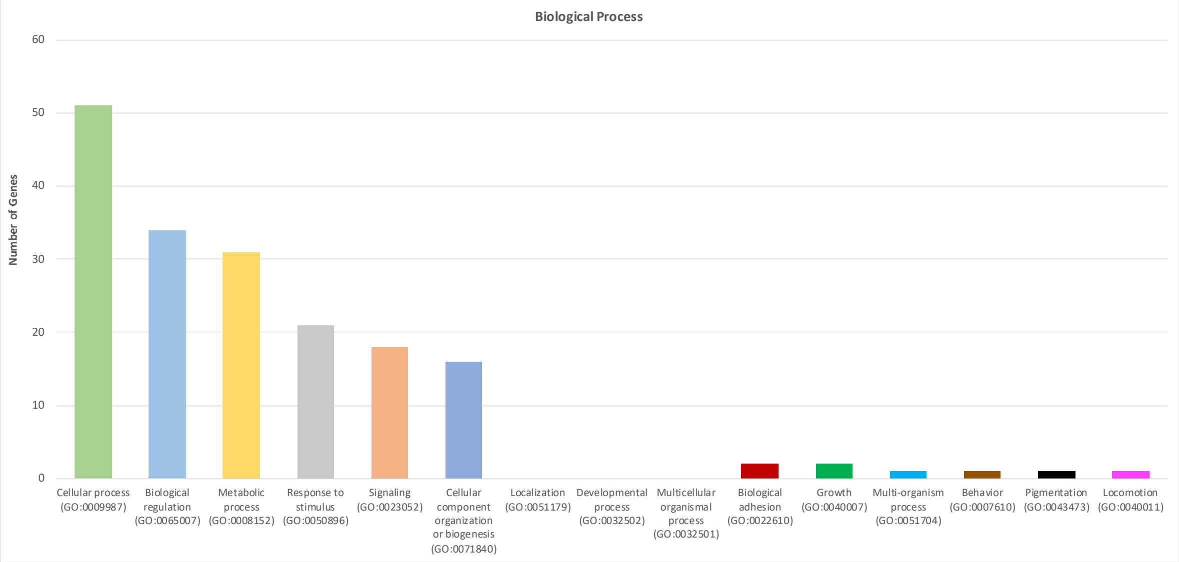

Figure S2. "Biological Process" GO enrichment of the 147 miRNA-targeted genes identified through miRCancerdb.

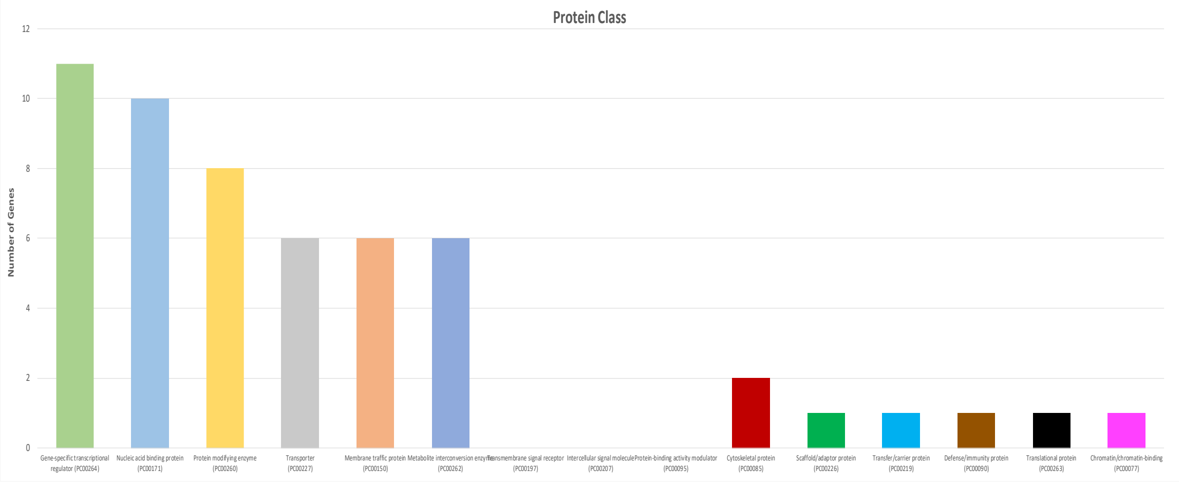

Figure S3. "Protein Class" GO enrichment of the 147 miRNA-targeted genes identified through miRCancerdb.

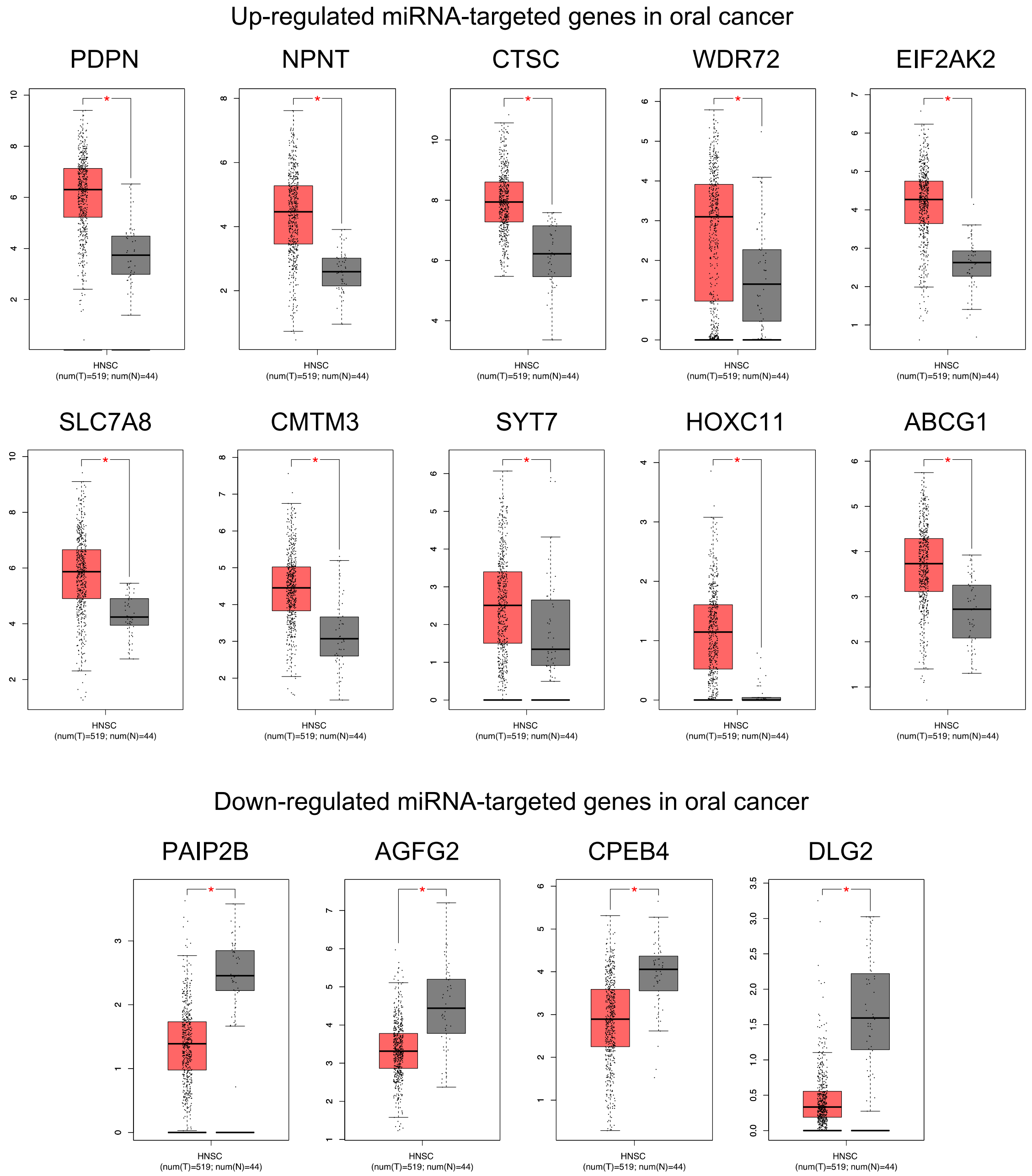

Figure S4. GEPIA analysis of the miRNA-targeted genes differentially expressed between oral cancer and healthy donors.

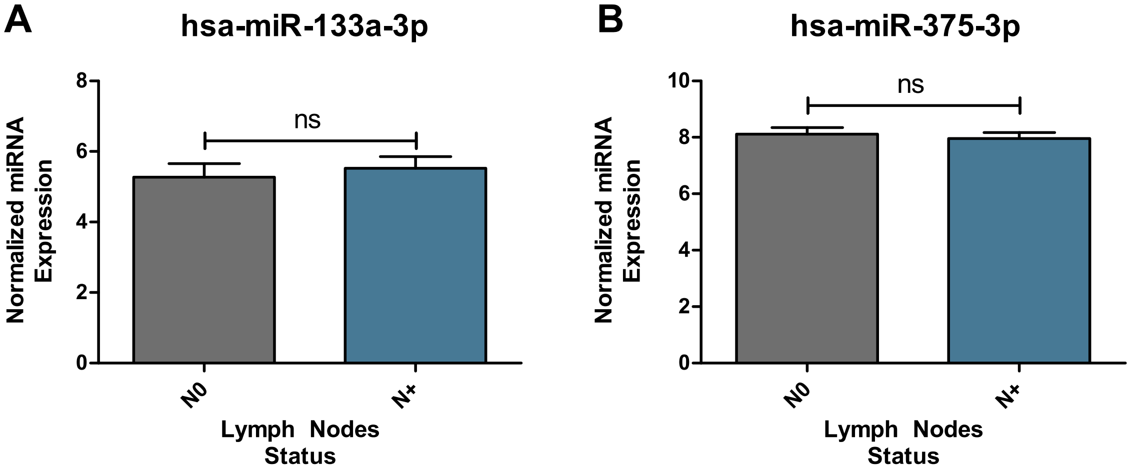

Figure S5. Panel A) No statistical difference observed for hsa-miR-133a-3p expression between oral cancer patients with positive lymph nodes and oral cancer patients without lymph nodes involvement according to the TCGA HNSC data; Panel B) No statistical difference observed for hsa-miR-375-3p expression between oral cancer patients with positive lymph nodes and oral cancer patients without lymph nodes involvement according to the TCGA HNSC data.