



Supplementary Figure 3. GO enrichment analysis for the metatranscriptome profiles of the oral microbiome associated with cancer status. Over-represented Molecular Functions. Enriched terms obtained using GOseq were summarized and visualized as a network using REVIGO. Summarized GO terms in healthy control matching tumor sites vs. OSCC tumor sites. Bubble color indicates the user-provided p-value (darker red is closer to 0). Highly similar GO terms are linked by edges in the graph.