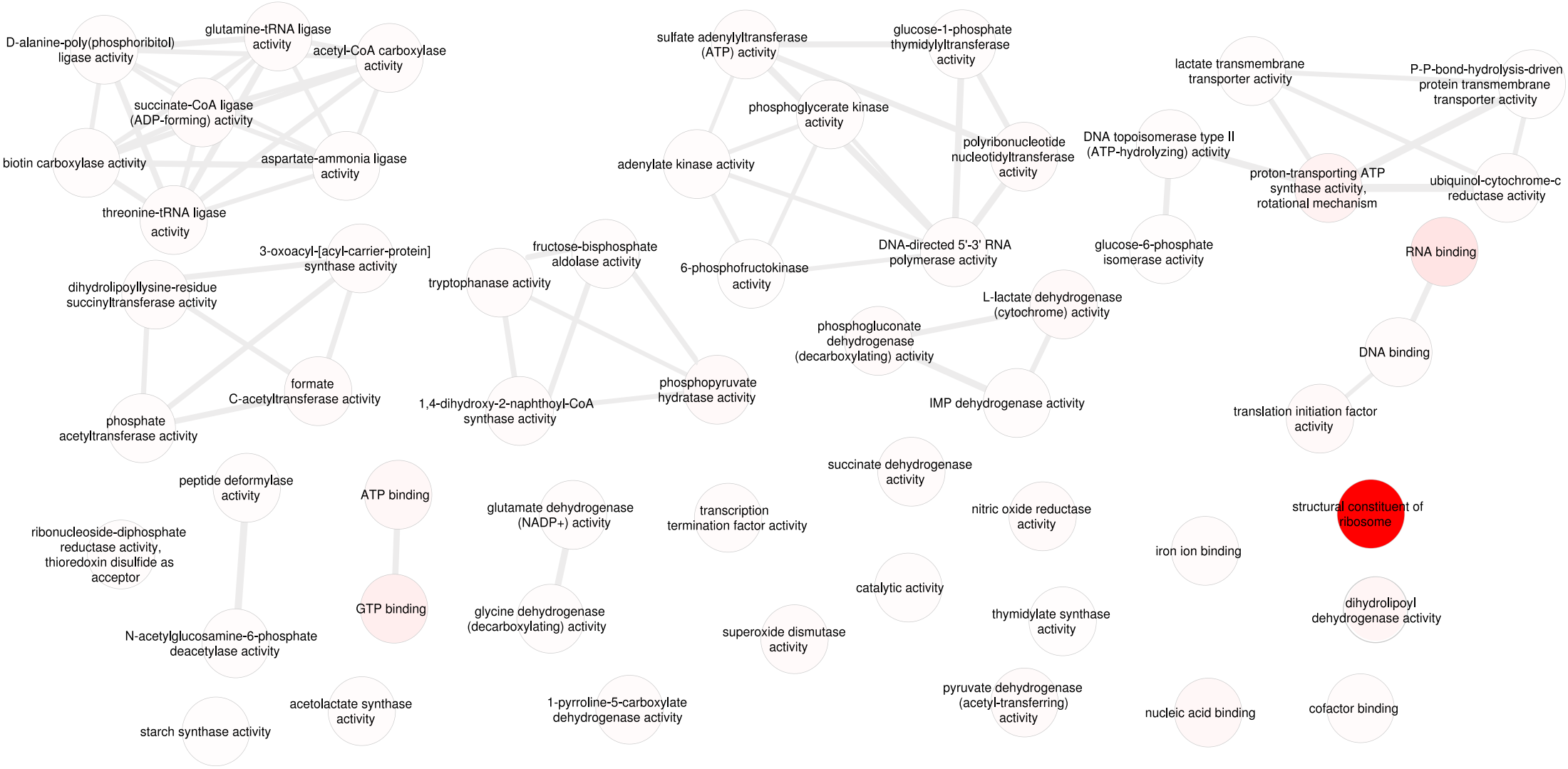

Supplementary Figure 5. GO enrichment analysis for the metatranscriptome profiles of the oral microbiome associated with cancer status.
Over-represented Molecular Functions. Enriched terms obtained using GOseq were summarized and visualized as a network using REVIGO. Summarized GO terms in OSCC tumor-adjacent vs. buccal sites from healthy control patients.
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.