**Supplementary Table 1. GO biological process enrichment analyses of differentially expressed genes for oral tongue squamous cell carcinoma.**

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| --- | --- | --- |
| **GO Term** | **Count** | ***P* Value** |
| GO:0006614~SRP-dependent cotranslational protein targeting to membrane | 28 | 4.86×10-26 |
| GO:0019083~viral transcription | 28 | 8.62×10-24 |
| GO:0006413~translational initiation | 30 | 9.63×10-24 |
| GO:0000184~nuclear-transcribed mRNA catabolic process, nonsense-mediated decay | 28 | 4.94×10-23 |
| GO:0006364~rRNA processing | 30 | 4.01×10-18 |

Note: GO, Gene Ontology.