**Table S1. Basic data of sequenced patients**

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| patients | Gender | Age | Tumor site | Differentiation | Lymphatic metastatic |
| 1 | Male | 46 | Right tougue | High | Yes |
| 2 | Female | 52 | Left tougue | Moderate | Yes |
| 3 | Male | 56 | Right tougue | Moderate | No |
| 4 | Female | 43 | Right tougue | High | No |

# Table S2. High throughput sequencing results – Differential gene analysis

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| Gene name | Log2(Metastatic Yes vs No) | P value | Log2 (CAL27 ER vs wt) | P value | Log2 (HN-6ER vs wt) | P value |
| LCN2 | 1.81 | <0.001 | 1.57 | <0.001 | 1.40 | <0.001 |
| PRAL | 1.28 | <0.001 | 4.09 | <0.001 | 1.20 | <0.05 |
| TNFSF18 | 3.23 | <0.001 | 4.01 | <0.001 | 1.28 | <0.001 |
| SHC3 | -2.40 | <0.001 | -1.05 | <0.05 | -1.51 | <0.01 |

**Table S3 Association of Lymph node metastasis in OSCC tongue xenografts**

|  |  |  |
| --- | --- | --- |
| Group | Lymph node metastasis | P Value |
| Yes | No |
|  1 |  |  |  |
| Ctrl-OV | 2 | 2 |  |
| LCN2-OV | 4 | 0 | 0.429 |
|  2 |  |  |  |
| Ctrl-sh | 2 | 2 |  |
| shLCN2 | 0 | 4 | 0.429 |
|  3 |  |  |  |
| shLCN2 | 0 | 4 |  |
| LCN2-OV | 4 | 0 | 0.029 |
|  4 |  |  |  |
| Ctrl-sh | 2 | 2 |  |
| Ctrl-OV | 2 | 2 | 1.000 |
|  |  |