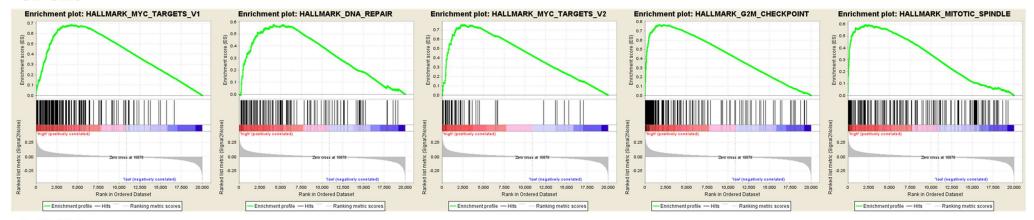
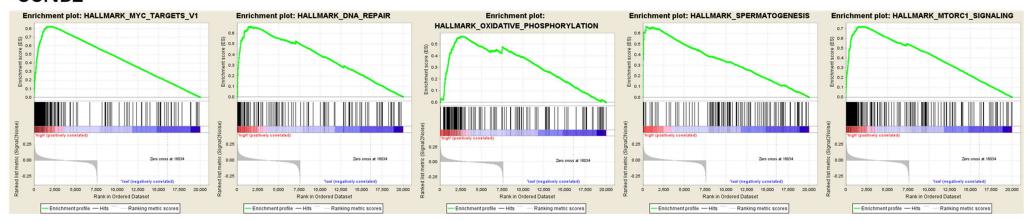


Figure S1 Qualification and statistics for the results of Western-Blot (A-B) The bar graph shows the gray value analysis of CyclinD1, and P21 for TCA8113 and CAL27 respectively. Data are mean \pm SD (error bars) from three independent experiments. (C) The bar graph shows the gray value analysis of AKT and P-AKT. Data are mean \pm SD (error bars) from three independent experiments. (D) The bar graph shows the gray value analysis of FAK and P-FAK. Data are mean \pm SD (error bars) from three independent experiments. (E) The bar graph shows the gray value analysis of P65 and P-P65. Data are mean \pm SD (error bars) from three independent experiments. (F) The bar graph shows the gray value analysis of STAT3 and P-STAT3. Data are mean \pm SD (error bars) from three independent experiments. *P<0.05, **P<0.01, ***P<0.001

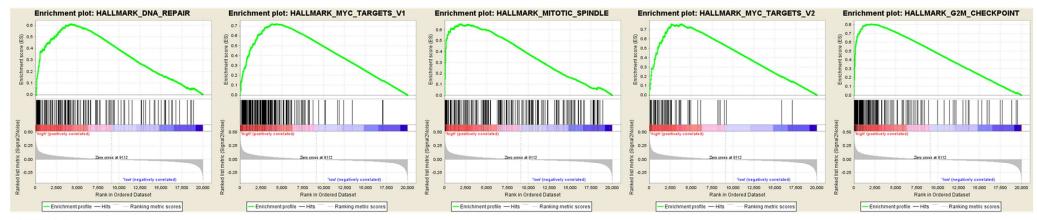
CDC20



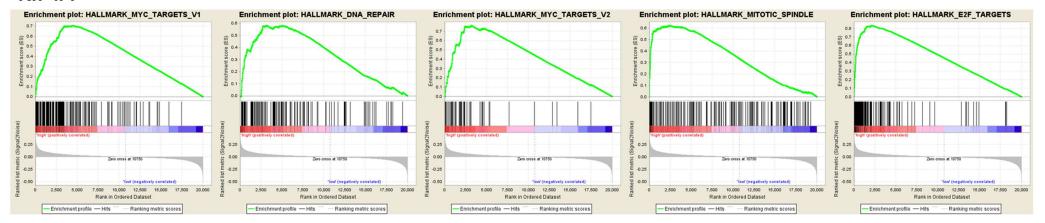
CCNB2



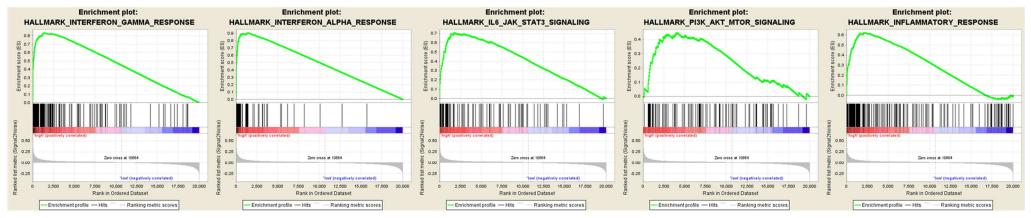
BUB1



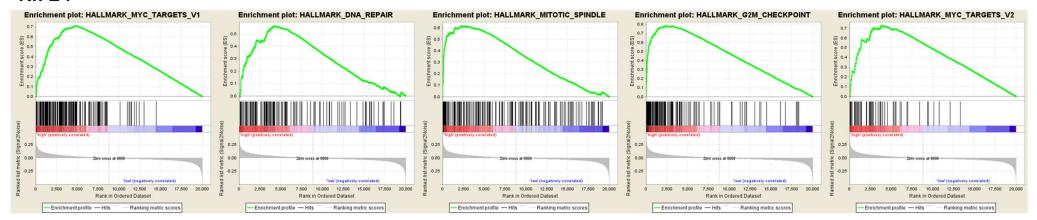
KIF4A



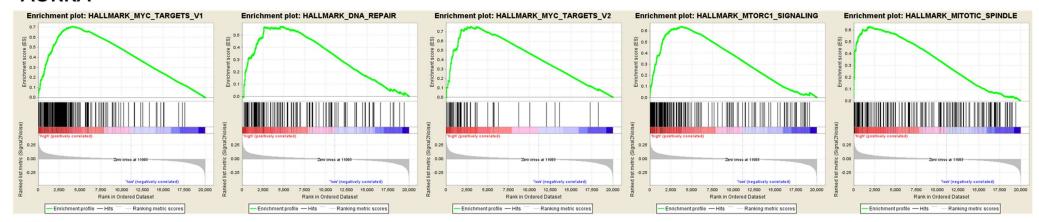
OAS1



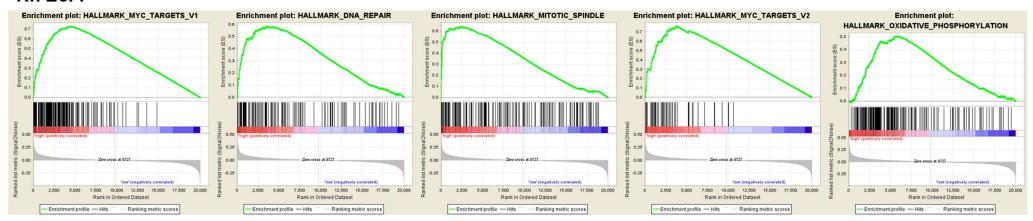
KIF14



AURKA



KIF20A



FOXM1

0.00

-0.25

0.00

-0.25

Rank in Ordered Dataset

—Enrichment profile — Hits — Ranking metric scores

7,500 10,000 12,500 15,000

Rank in Ordered Dataset

- Enrichment profile - Hits - Ranking metric scores

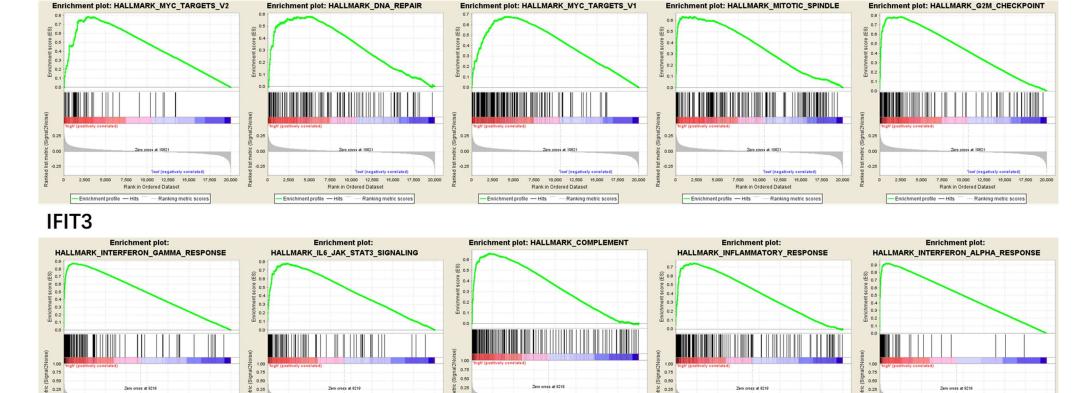


Figure S2 The GSEA analysis of the 10 hub genes to revealed their function. The dataset GSE30784 which contains 167 OSCC samples were downloaded from GEO database to perform GSEA analysis. The ten hub genes were divided into two groups according to their expression respectively (84 high expression group and 83 low expression group). The hallmark gene sets (h.all.v6.1.symbols.gmt) were used to show the function of the ten hub genes.

- Ranking metric scores

- Enrichment profile - Hits

0.00

-0.25

0.00

-0.25

7.500 10.000 12.500 15.000

Rank in Ordered Dataset

- Enrichment profile - Hits

7.500 10.000 12.500 15.000

— Ranking metric scores

- Enrichment profile - Hits

0.00

S -0.25