

Genomic Analyses Reveal Mutational Signatures and Frequently Altered Genes in Esophageal Squamous Cell Carcinoma

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(The American Journal of Human Genetics 96, 597–611; April 2, 2015)

In the originally published version of this article, Figure 5C mistakenly included the image of KYSE150 instead of KYSE140. Here we have included the correct image for ZNF750-si KYSE140 cells in Figure 5C. The authors regret the error.

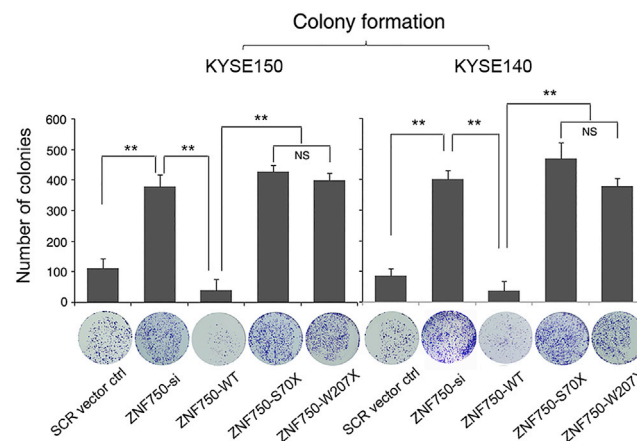


Figure 5C. ZNF750 Acts as a Tumor Suppressor in ESCC

Endogenous ZNF750 was stably knocked down in KYSE150 and KYSE140 cells and then forced to encode FLAG-tagged wild-type ZNF750 (ZNF750-WT) or the p.Ser70* (ZNF750-S70X) or p.Trp207* (ZNF750-W207X) variant. Cell proliferation was monitored by colony-formation assay. SCR indicates scramble control. The data represent the mean \pm SD; three independent experiments were performed in triplicate. Data were statistically analyzed with a two-sided t test. ** $p < 0.01$, * $p < 0.05$.

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<https://doi.org/10.1016/j.ajhg.2020.07.008>

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