

Figure S1. Expression levels of predicted downstream genes of miR-613. (A) GJA1, (B) HACD3, (C) PAX7, (D) PHAX, (E) BDNF, (F) G6PD, (G) CPED1, (H) TNKS2 and (I) SLC44A1 in normal and tumor tissues. Data are presented as the mean \pm SEM. * $P < 0.05$. NS, not significant; GJA1, gap junction α -1 protein.

