



SUPPLEMENTARY FIG. S6. RQ-PCR analysis of the (A) *Wnt*, *Igf2*, and *Fgf16* (B) genes in the indicated bladder tissues. The expression level in the indicated bladder tissues is represented as % *Gapdh* (as determined using five or more independent experiments) and is shown as the mean \pm SEM ($P < 0.05$, $P < 0.01$ in comparison with IC group bladders, one-way ANOVA with Bonferroni post-test).