

Figure S8. Expression levels of Opto-mFGFR1 and FGFR1 in M38K^{Opto-mFGFR1}, SPC212^{Opto-mFGFR1} and hBE^{Opto-mFGFR1} cells. Total RNA from these cells and control cells was extracted, reverse transcribed and subjected to qPCR. Bars represent fold mRNA levels of Opto-mFGFR1 compared to endogenous FGFR1. Mean values ± SEM for three independent experiments each performed in duplicates are shown.