Integration of TGF-ß and Jagged1/Notch Signalling in Epithelial-to-Mesenchymal Transition

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Supplementary Figure S1

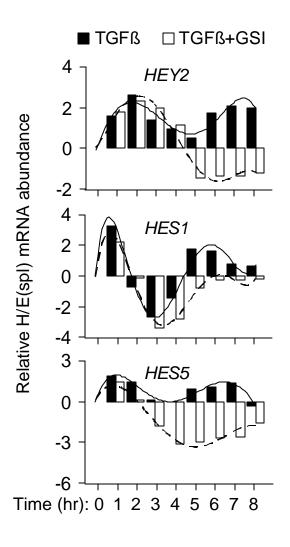


Fig. S1 H/E(spl) repressors HEY2, HES1 and HES5 are targets of coordinated signalling by TGF-ß /Smad and Notch. Bar graphs show averages of relative H/E(spl) mRNA abundance (log2 transformed), determined by quantitative real-time analysis (N=3) in HaCaT PCR keratinocytes treated with TGF-ß in the absence (black bars, solid line) or presence (open bars, broken line) of ?secretase inhibitor (GSI).