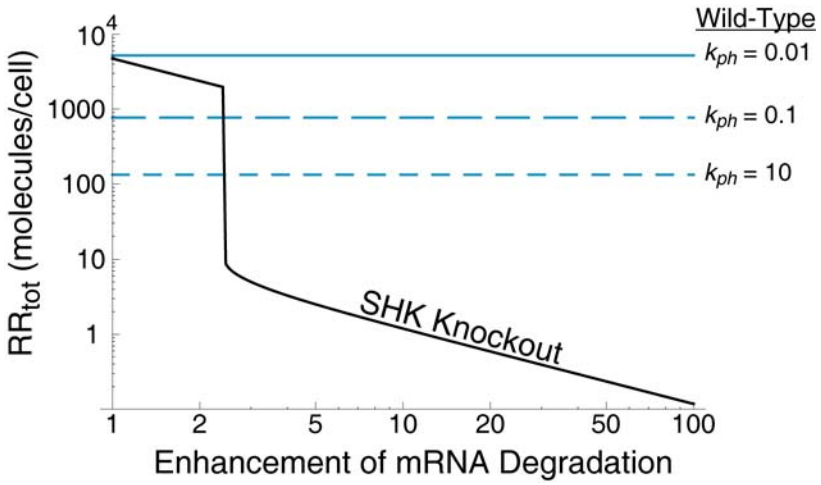

Figure S6. Altered mRNA stability in a simulated SHK knockout changes total RR concentrations. Wild-type concentrations at the default degradation rate for various signal levels are shown for reference. All simulations use the default parameter set (Table S1).