



Figure S5 Comparison of qPCR results from all single and double *her/hes* perturbations with fit from predicted GRN. qPCR values are shown as normalized log fold change. For each knockdown condition, results for all six genes: *her1*, *her7*, *her11*, *her12*, *hes6*, and *her15* are given. Knockdowns are labeled above each graph, and the order of genes shown is constant. For simplicity, each gene is labeled only once per plot. The order is consistent throughout. Experimental results are shown with standard deviation in green. Five replicates were performed for all single knockdowns, and three replicates for all double knockdowns. Computed qPCR values from the predicted GRN are shown as pink circles.