



Figure S3. SW620, cross-condition comparison. Aggregated display of interventions and timepoints showing directional transcriptional changes relative to control (red for up, blue for down). Track length and endpoint size give a visual proxy for response breadth, supporting like-for-like comparison of effect sizes across all SW620 conditions. Cutoffs: $|\log_2 \text{FC}| > 1$, gene adjusted $p < 0.05$, gene-set FDR < 0.05 .