



**Figure S5.** U138, RF/AMRF-HT  $\pm$  RT. Intervention- and timepoint-resolved counts of up-regulated (red) and down-regulated (blue) transcripts relative to control. Visual encodings (track length, endpoint size) reflect the number of genes per direction, enabling comparison of RF/AMRF-HT alone and in combination with RT in U138 cells. Cutoffs:  $|\log_2 \text{FC}| > 1$ , gene adjusted  $p < 0.05$ , gene-set FDR  $< 0.05$ .