Let7d_Ator_vs_Scramble_Ator PCA plot using MsigDB Hallmark with Process Category 1.2 -E2F_TARGETS group_drug G2M_CHECKPOINT Let7d_Ator Let7d_Lova 0.8 -Let7d_None Let7d_TNFa INTERFERON_ALPHA_RESPONSE Scramble Ator Scramble_Lova PC2: 13% variance INTERFERON GAMMA RESPONSE Scramble_None 0.4 -Scramble_TNFa MITOTIC_SPINDLE ALLOGRAFT_REJECTION UV RESPONSE UP Process_Category IL6_JAK_STAT3_SIGNALING PROTEIN_SECRETION development 0.0 -DNA_damage **INFLAMMATORY RESPONSE** COMPLEMENT ╁┼ immune **EPITHELIAL MESENCHYMAL TRANSITION** metabolic **HYPOXIA** other TNFA_SIGNALING_VIA_NFKB pathway **MYOGENESIS** -0.4 proliferation signaling CHOLESTEROL_HOMEOSTASIS KRAS_SIGNALING_UP -1.00.0 -0.50.5 PC1: 85% variance