Let7d_TNFa_vs_Scramble_TNFa PCA plot using MsigDB Hallmark with Process_Category 1.0 -E2F_TARGETS **Process Category** MYC_TARGETS_V2 cellular_component INTERFERON_ALPHA_RESPONSE development **G2M CHECKPOINT** INTERFERON_GAMMA_RESPONSE DNA_damage immune 0.5 other MYC TARGETS V1 pathway PC2: 15% variance proliferation ALLOGRAFT_REJECTION signaling MITOTIC SPINDLE INFLAMMATORY_RESPONSE UNFOLDED_PROTEIN_RESPONSE IL6_JAK_STAT3_SIGNALING UV_RESPONSE_UP group_drug Let7d_Ator COMPLEMENT 0.0 -Let7d_Lova EPITHELIAL_MESENCHYMAL_TRANSITION TNFA_SIGNALING_VIA_NFKB Let7d_None ESTROGEN_RESPONSE_EARLY Let7d_TNFa APICAL_JUNCTION Scramble_Ator X **HYPOXIA** Scramble_Lova COAGULATION Scramble_None **MYOGENESIS** Scramble TNFa P53_PATHWAY -0.5 **-**KRAS_SIGNALING_UP -1.00.5 -0.50.0 PC1: 83% variance