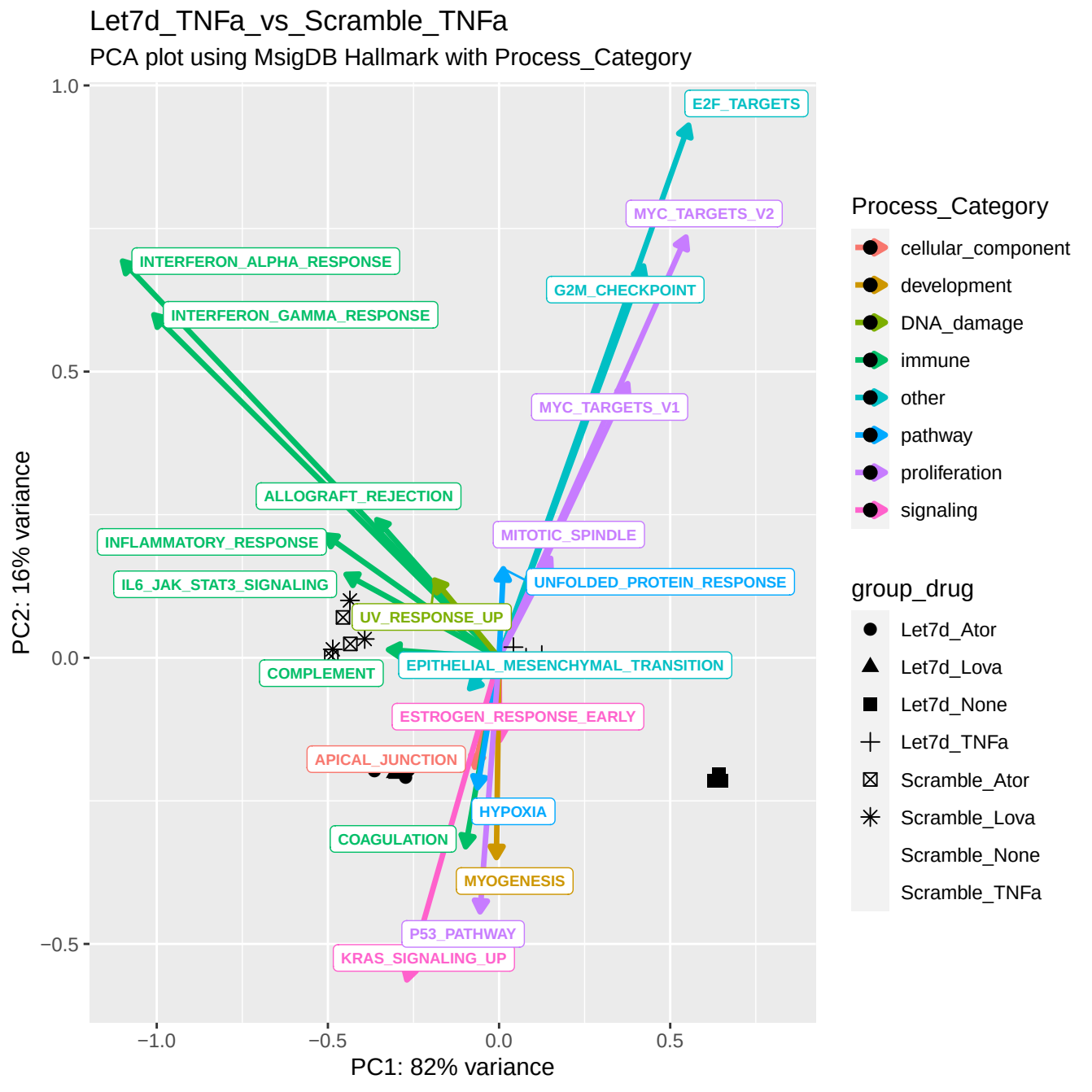
Let7d_TNFa_vs_Scramble_TNFa
PCA plot using MsigDB Hallmark with Process_Category
E2F_TARGETS
MYC_TARGETS_V2
INTERFERON_ALPHA_RESPONSE
G2M_CHECKPOINT
INTERFERON_GAMMA_RESPONSE
MYC_TARGETS_V1
ALLOGRAFT_REJECTION
INFLAMMATORY_RESPONSE
MITOTIC_SPINDLE
IL6_JAK_STAT3_SIGNALING
UNFOLDED_PROTEIN_RESPONSE
UV_RESPONSE_UP
COMPLEMENT
EPITHELIAL_MESENCHYMAL_TRANSITION
ESTROGEN_RESPONSE_EARLY
APICAL_JUNCTION
HYPOXIA
COAGULATION
MYOGENESIS
P53_PATHWAY
KRAS_SIGNALING_UP
1.0
0.5
0.0
−0.5
PC2: 16% variance
−1.0
−0.5
0.0
0.5
PC1: 82% variance
Process_Category
cellular_component
development
DNA_damage
immune
other
pathway
proliferation
signaling
group_drug
Let7d_Ator
Let7d_Lova
Let7d_None
Let7d_TNFa
Scramble_Ator
Scramble_Lova
Scramble_None
Scramble_TNFa