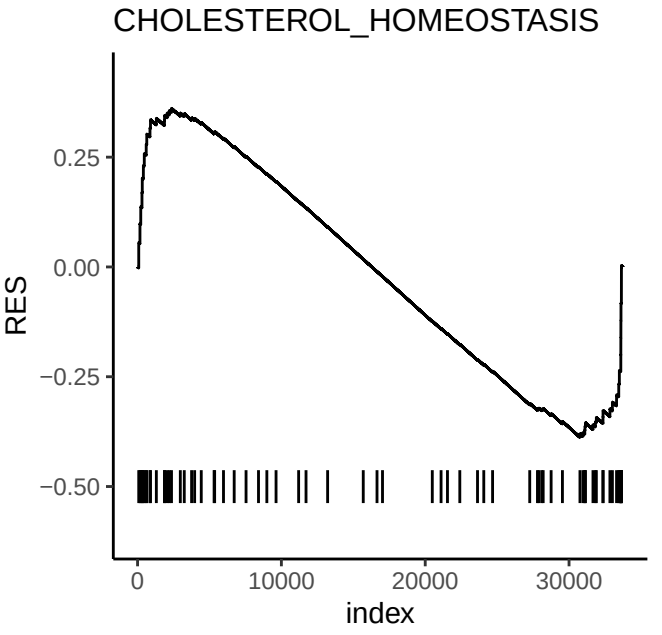
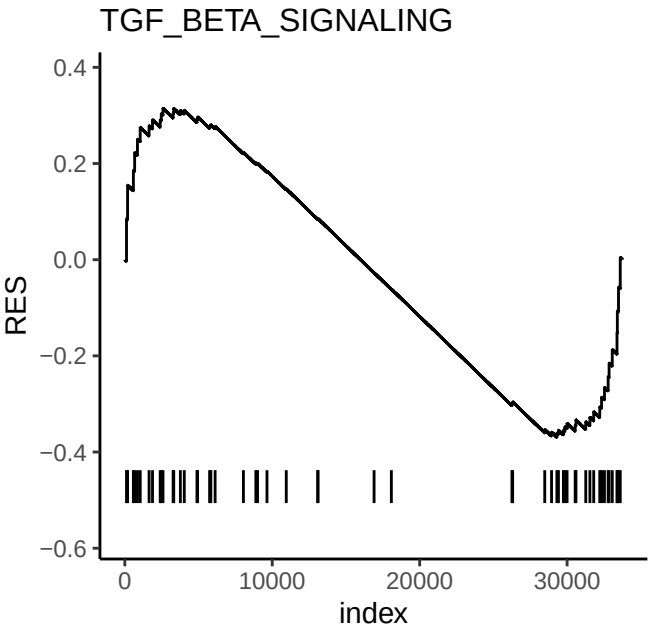

CHOLESTEROL_HOMEOSTASIS
RES
0.25
0.00
−0.25
−0.50
0
10000
20000
30000
index


TGF_BETA_SIGNALING
RES
0.4
0.2
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−0.2
−0.4
−0.6
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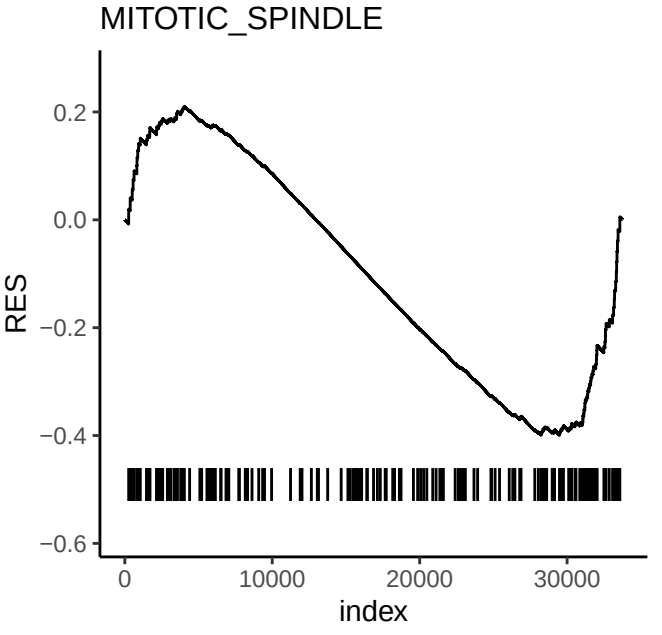
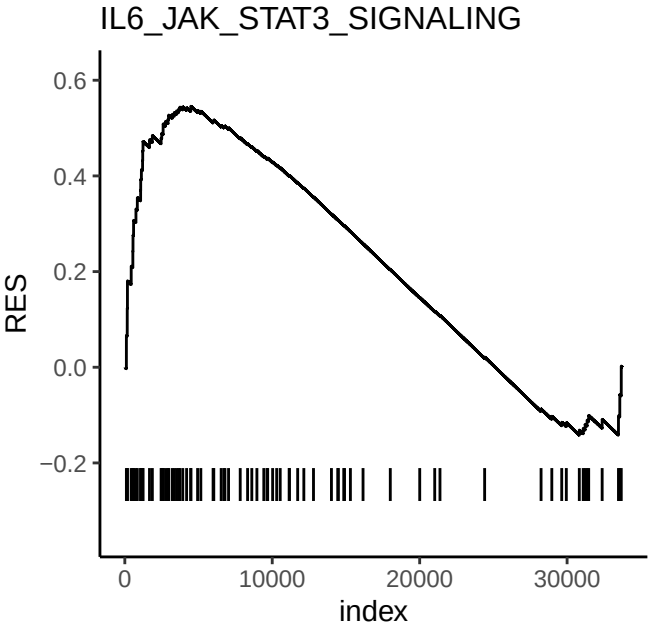
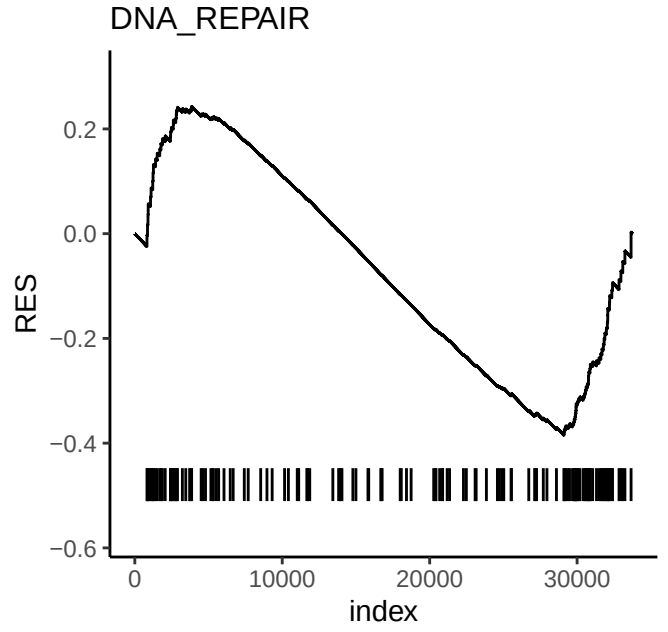
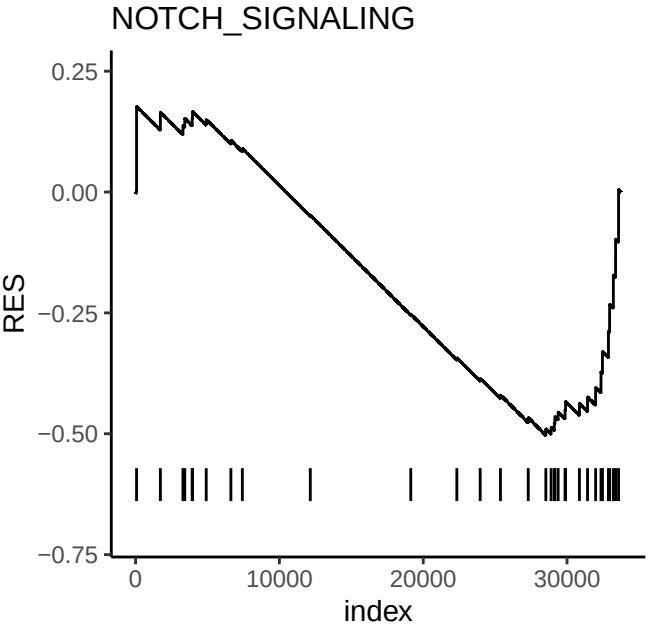
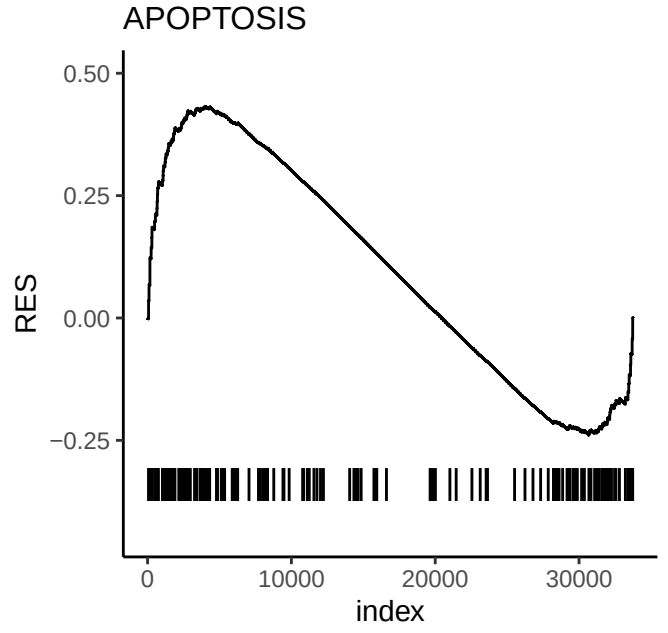
10000
20000
30000
index

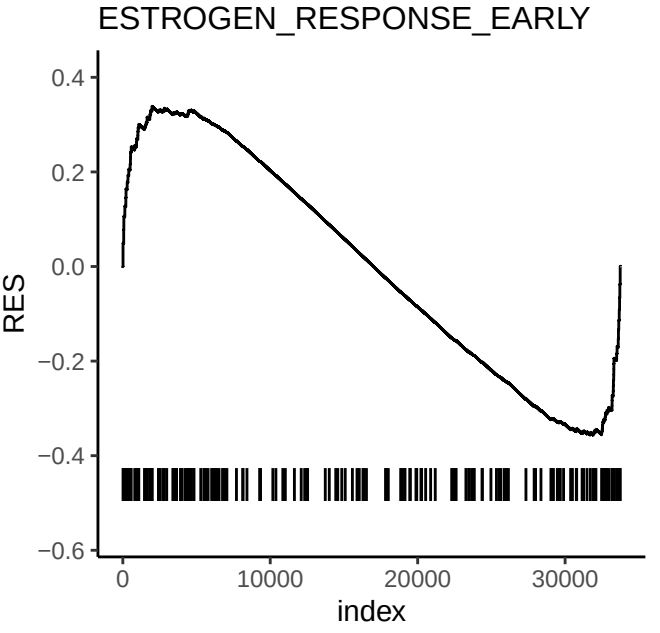

MITOTIC_SPINDLE
RES
0.2
0.0
−0.2
−0.4
−0.6
0
10000
20000
30000
index

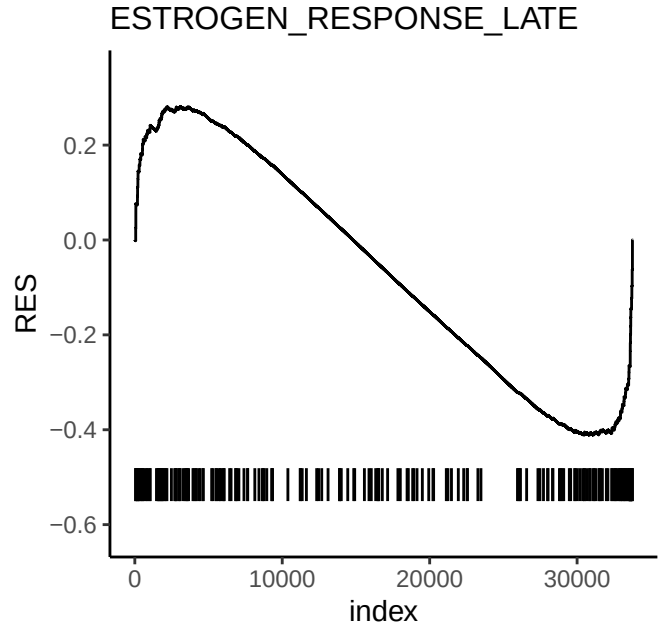

IL6_JAK_STAT3_SIGNALING
RES
0.6
0.4
0.2
0.0
−0.2
0
10000
20000
30000
index

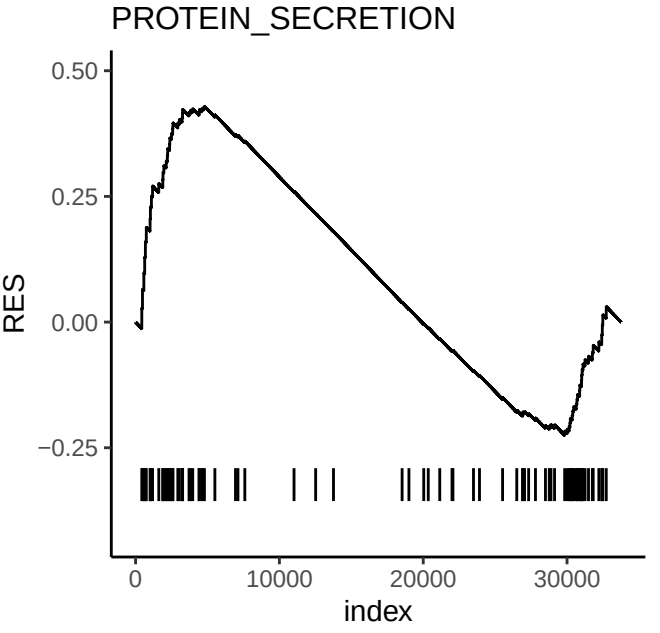

DNA_REPAIR
RES
0.2
0.0
−0.2
−0.4
−0.6
0
10000
20000
30000
index

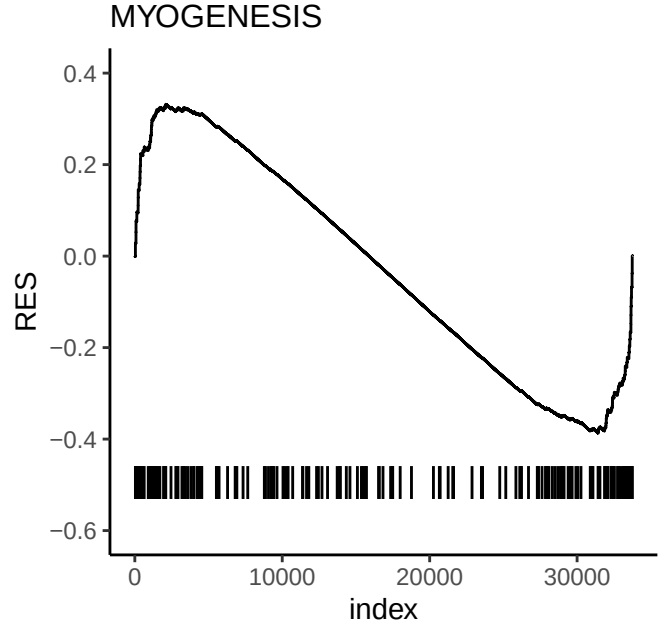

NOTCH_SIGNALING
RES
0.25
0.00
−0.25
−0.50
−0.75
0
10000
20000
30000
index

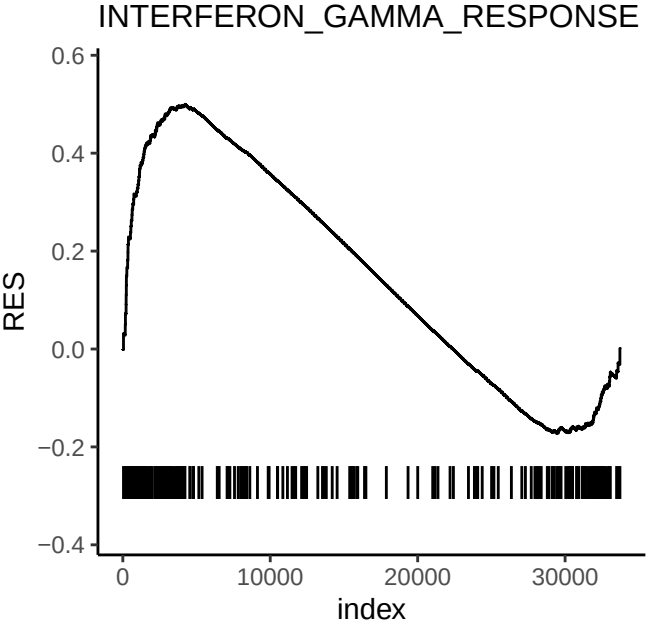

APOPTOSIS
RES
0.50
0.25
0.00
−0.25
0
10000
20000
30000
index

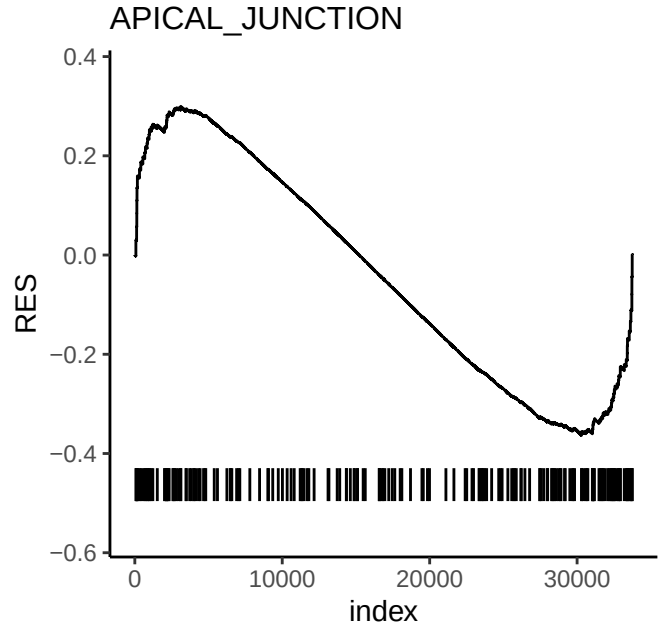

ESTROGEN_RESPONSE_EARLY
RES
0.4
0.2
0.0
−0.2
−0.4
−0.6
0
10000
20000
30000
index

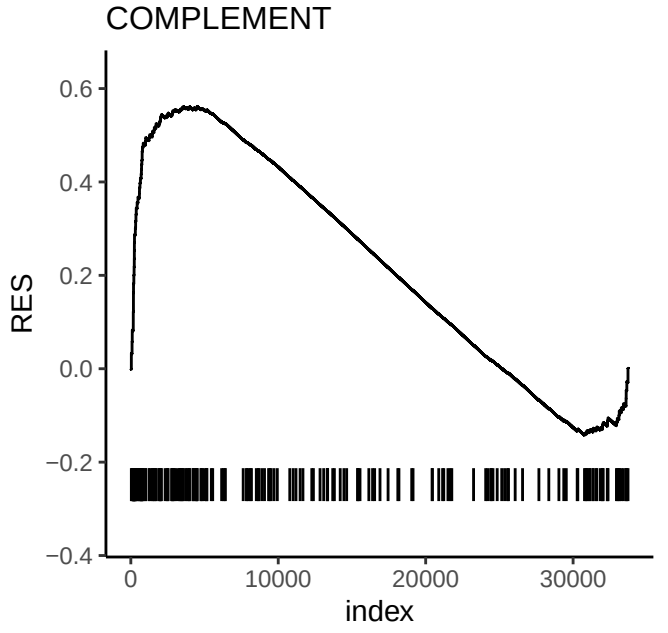

ESTROGEN_RESPONSE_LATE
RES
0.2
0.0
−0.2
−0.4
−0.6
0
10000
20000
30000
index

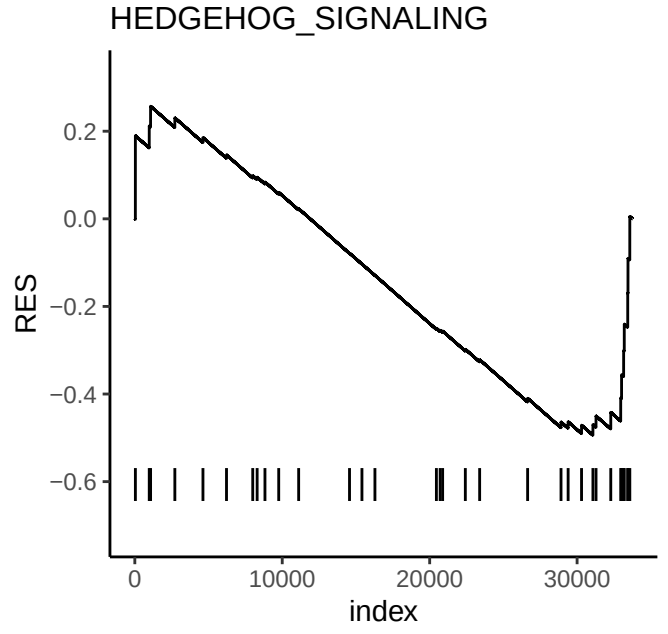

PROTEIN_SECRETION
RES
0.50
0.25
0.00
−0.25
0
10000
20000
30000
index

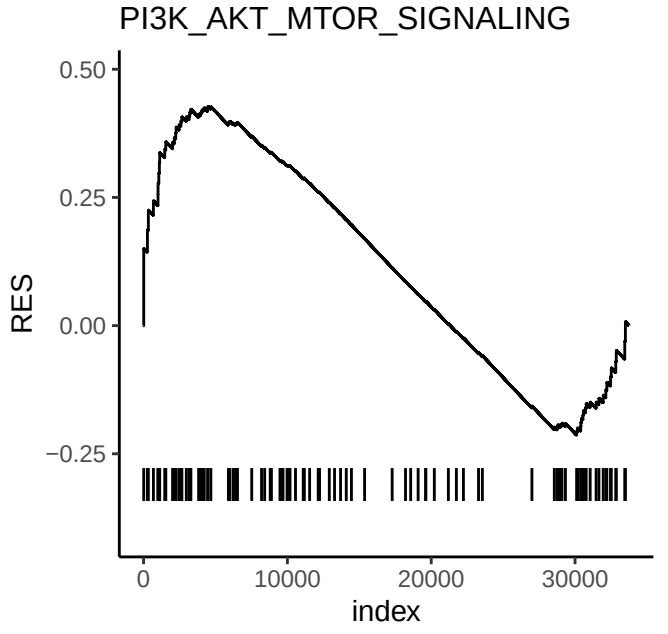

MYOGENESIS
RES
0.4
0.2
0.0
−0.2
−0.4
−0.6
0
10000
20000
30000
index

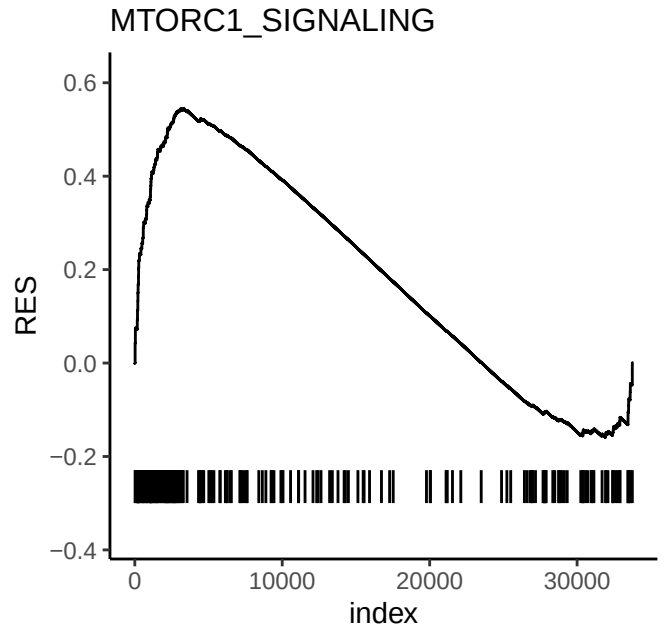

INTERFERON_GAMMA_RESPONSE
RES
0.6
0.4
0.2
0.0
−0.2
−0.4
0
10000
20000
30000
index

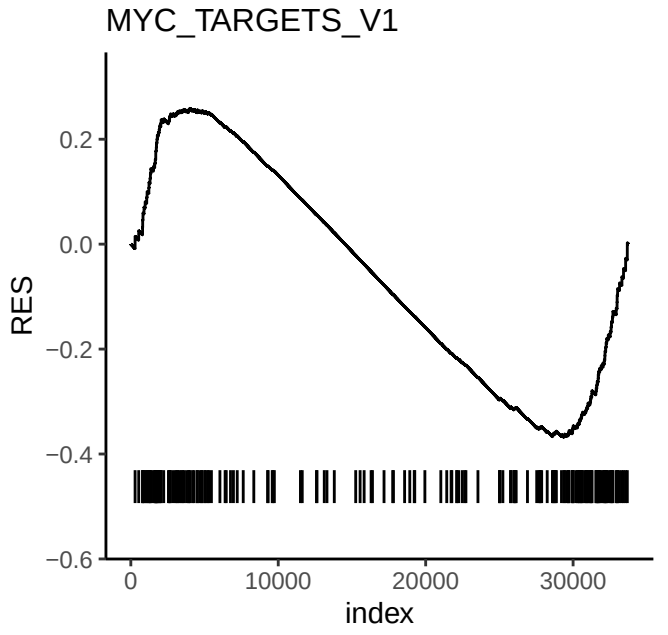

APICAL_JUNCTION
RES
index
0.4
0.2
0.0
−0.2
−0.4
−0.6
0
10000
20000
30000

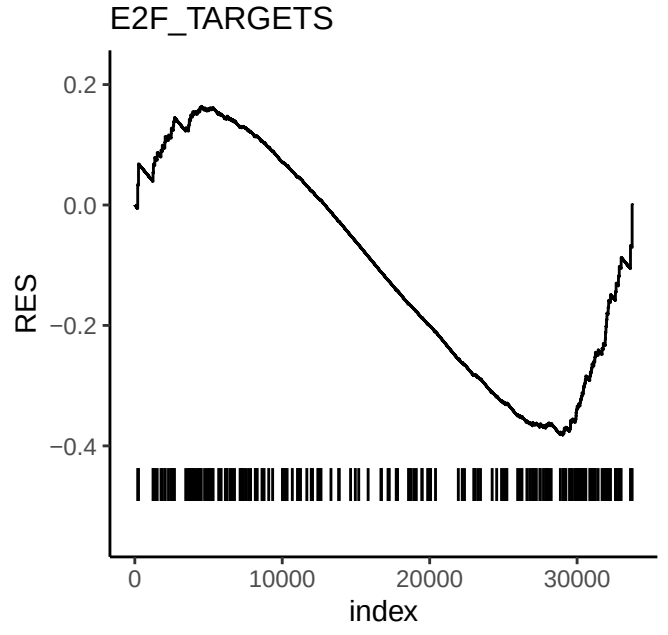

COMPLEMENT
RES
index
0.6
0.4
0.2
0.0
−0.2
−0.4
0
10000
20000
30000

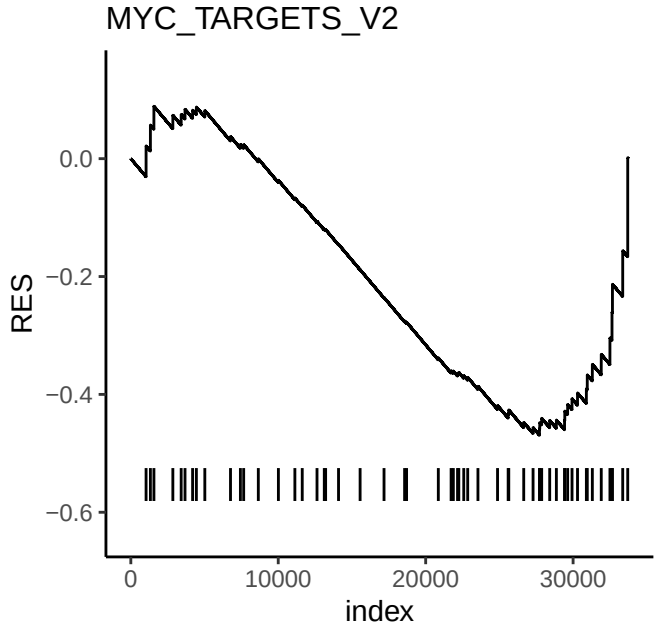

HEDGEHOG_SIGNALING
RES
index
0.2
0.0
−0.2
−0.4
−0.6
0
10000
20000
30000

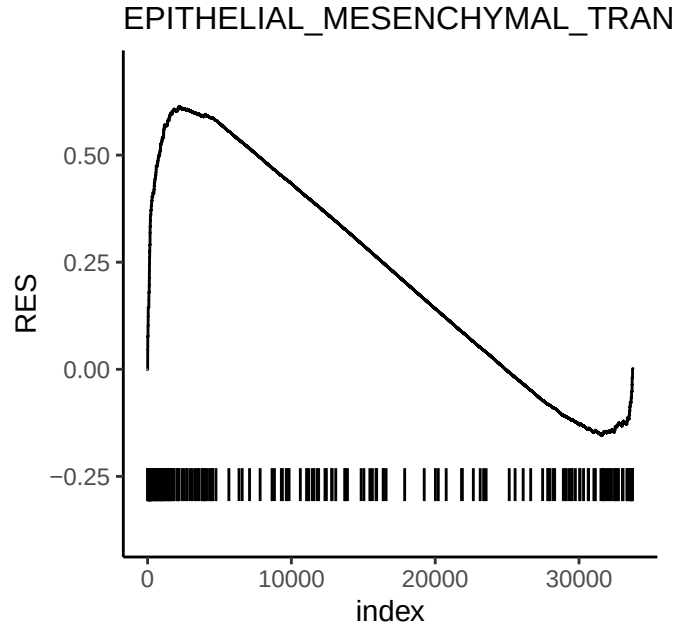

PI3K_AKT_MTOR_SIGNALING
RES
index
0.50
0.25
0.00
−0.25
0
10000
20000
30000

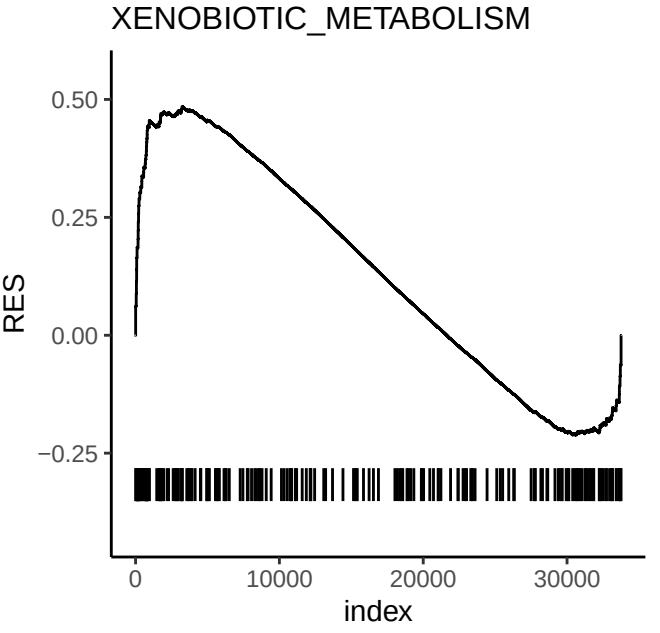

MTORC1_SIGNALING
RES
index
0.6
0.4
0.2
0.0
−0.2
−0.4
0
10000
20000
30000

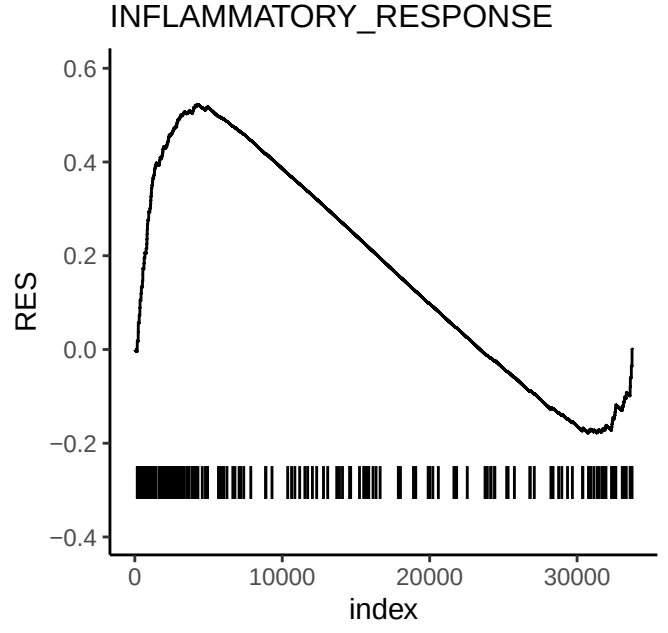

MYC_TARGETS_V1
RES
index
0.2
0.0
−0.2
−0.4
−0.6
0
10000
20000
30000

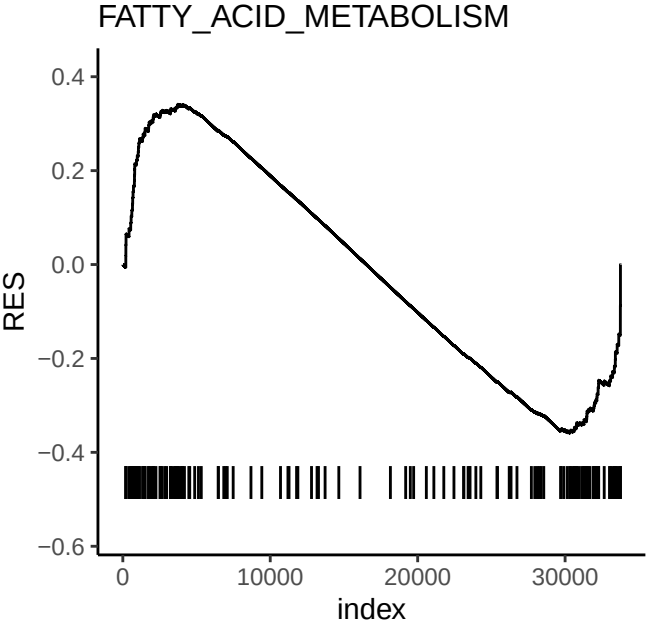

E2F_TARGETS
RES
index
0.2
0.0
−0.2
−0.4
0
10000
20000
30000

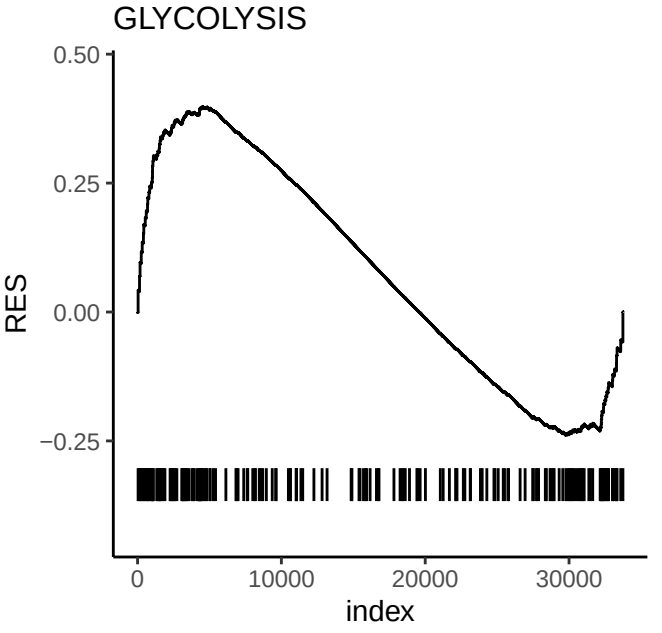

MYC_TARGETS_V2
RES
index
0.0
−0.2
−0.4
−0.6
0
10000
20000
30000

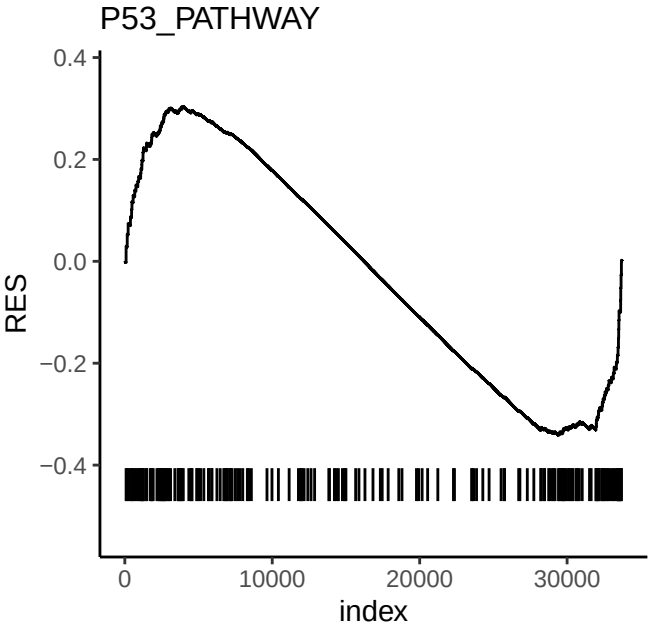

EPITHELIAL_MESENCHYMAL_TRAN
RES
0.50
0.25
0.00
−0.25
0
10000
20000
30000
index

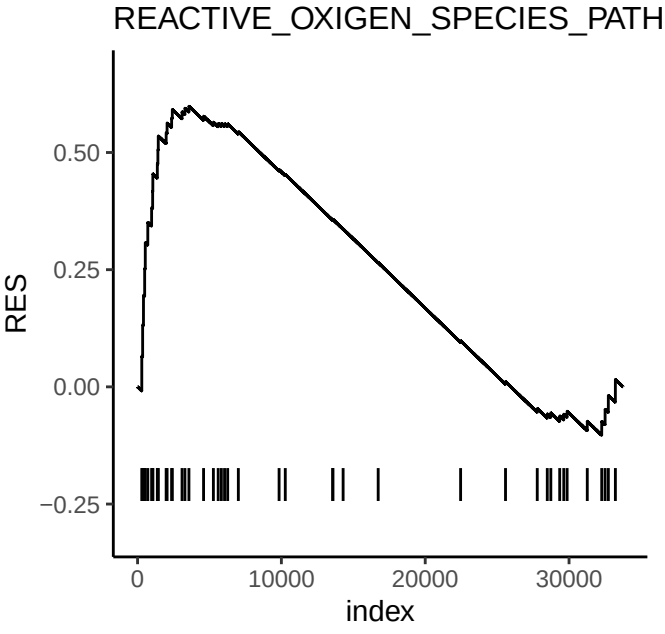

XENOBIOTIC_METABOLISM
RES
0.50
0.25
0.00
−0.25
0
10000
20000
30000
index

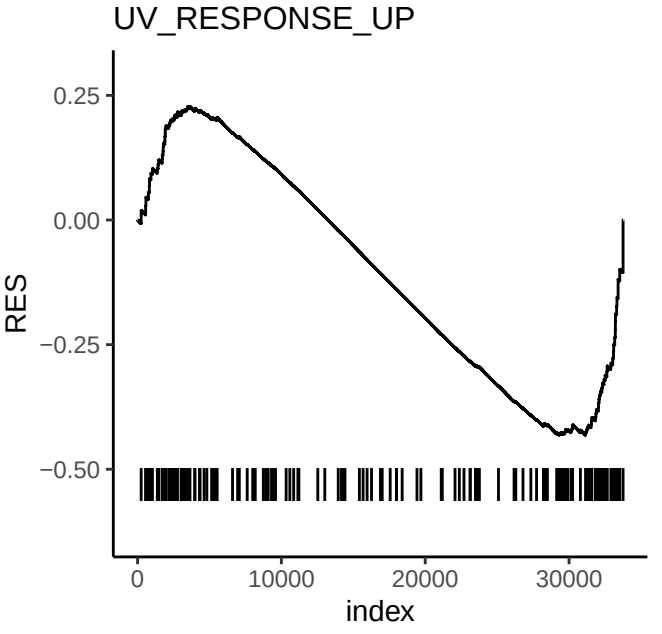

INFLAMMATORY_RESPONSE
RES
0.6
0.4
0.2
0.0
−0.2
−0.4
0
10000
20000
30000
index

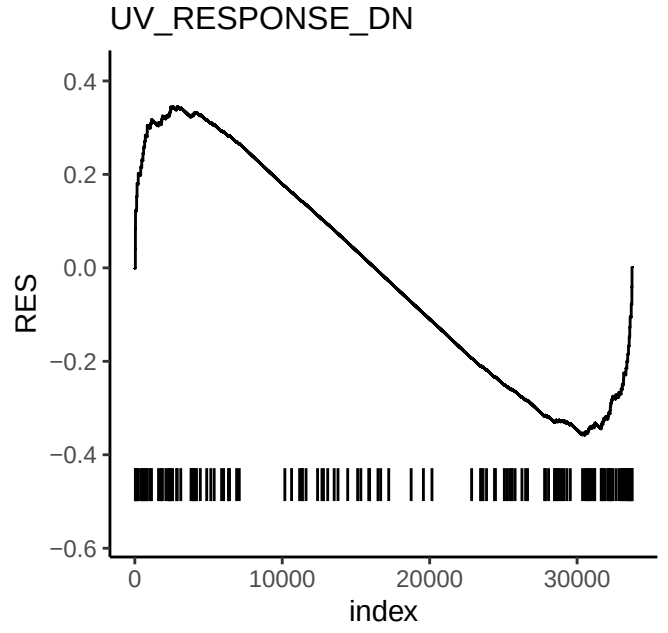

FATTY_ACID_METABOLISM
RES
0.4
0.2
0.0
−0.2
−0.4
−0.6
0
10000
20000
30000
index

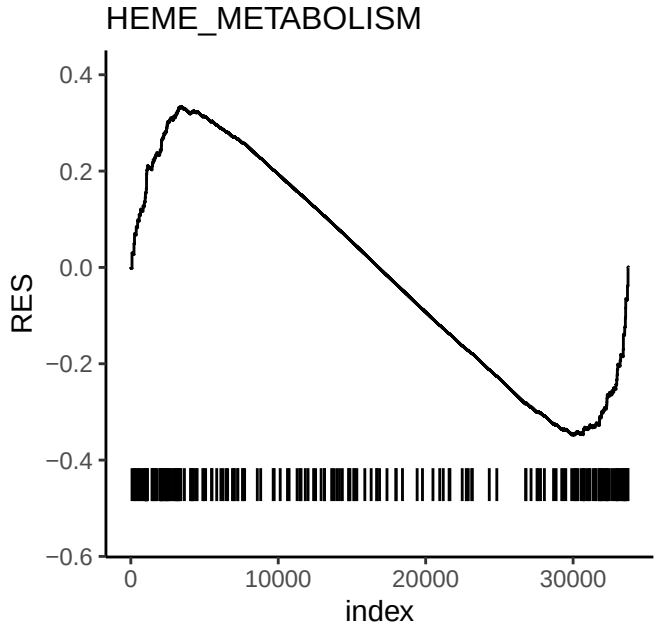

GLYCOLYSIS
RES
0.50
0.25
0.00
−0.25
0
10000
20000
30000
index

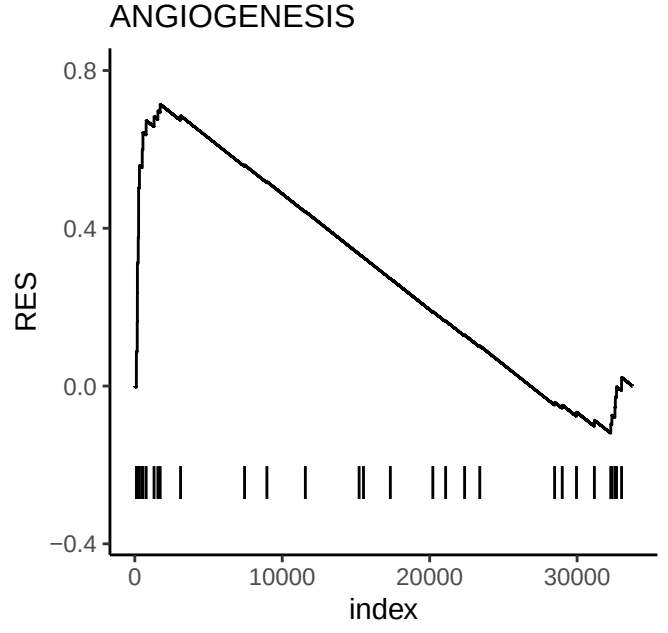

P53_PATHWAY
RES
0.4
0.2
0.0
−0.2
−0.4
0
10000
20000
30000
index

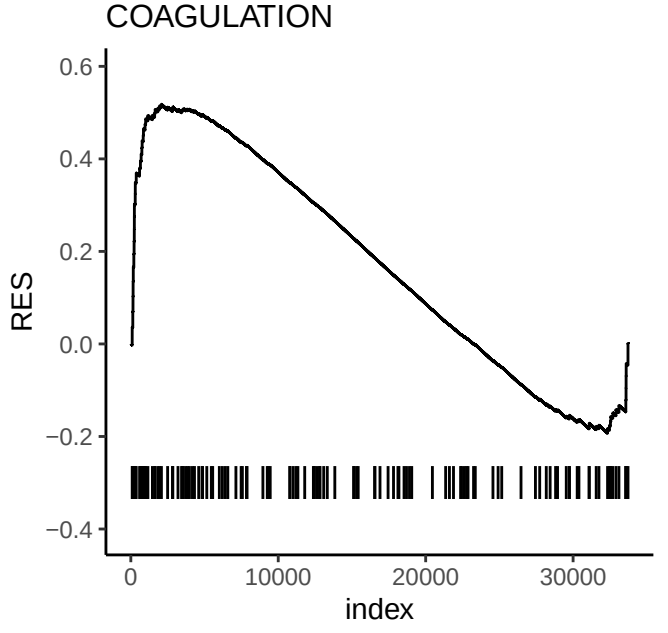

REACTIVE_OXIGEN_SPECIES_PATH
RES
0.50
0.25
0.00
−0.25
0
10000
20000
30000
index

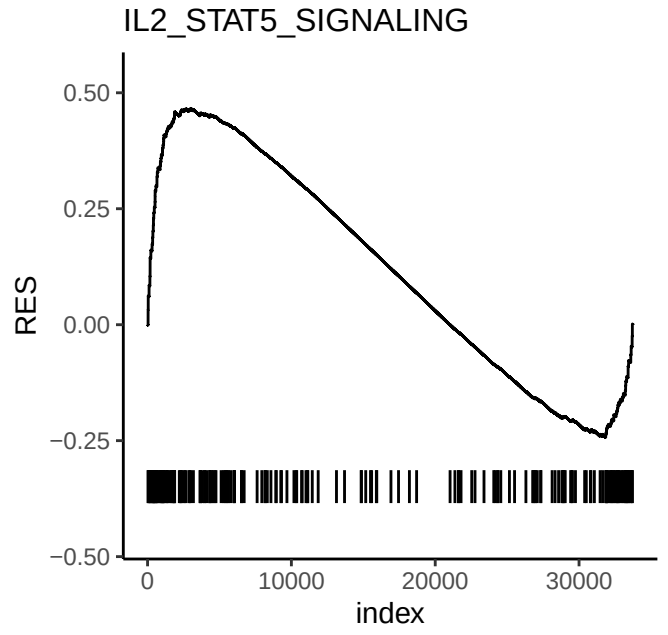

UV_RESPONSE_UP
RES
0.25
0.00
−0.25
−0.50
0
10000
20000
30000
index

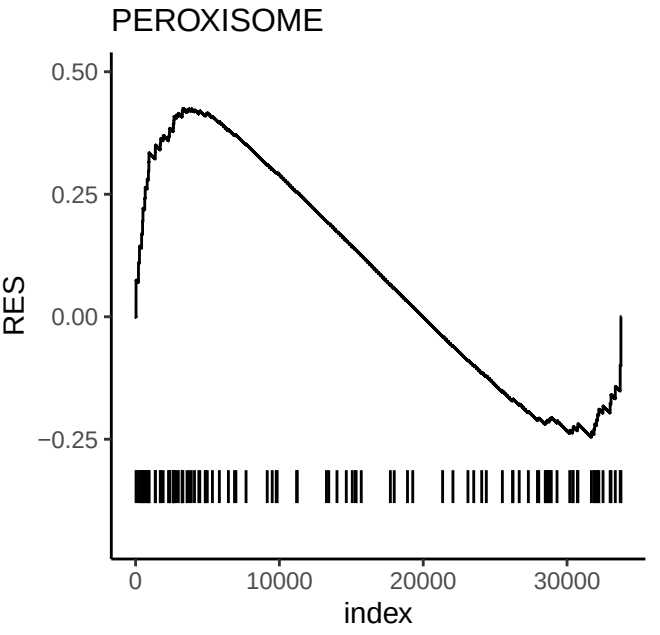

UV_RESPONSE_DN
RES
0.4
0.2
0.0
−0.2
−0.4
−0.6
0
10000
20000
30000
index

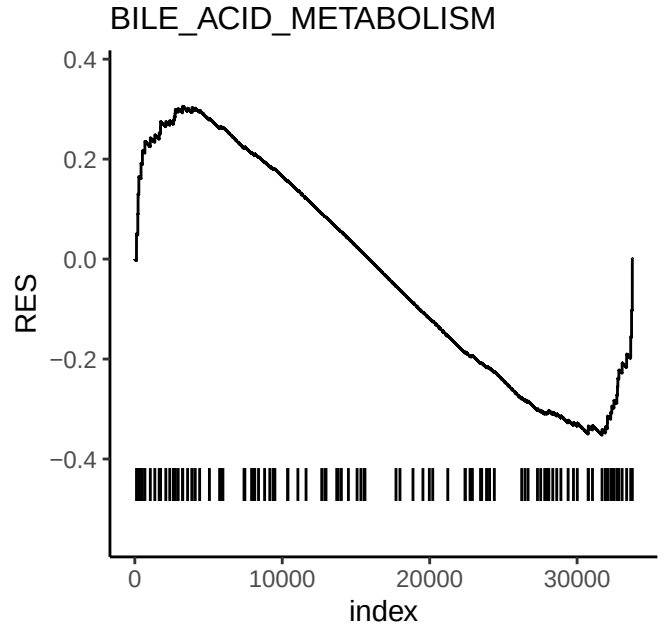

HEME_METABOLISM
RES
0.4
0.2
0.0
−0.2
−0.4
−0.6
0
10000
20000
30000
index

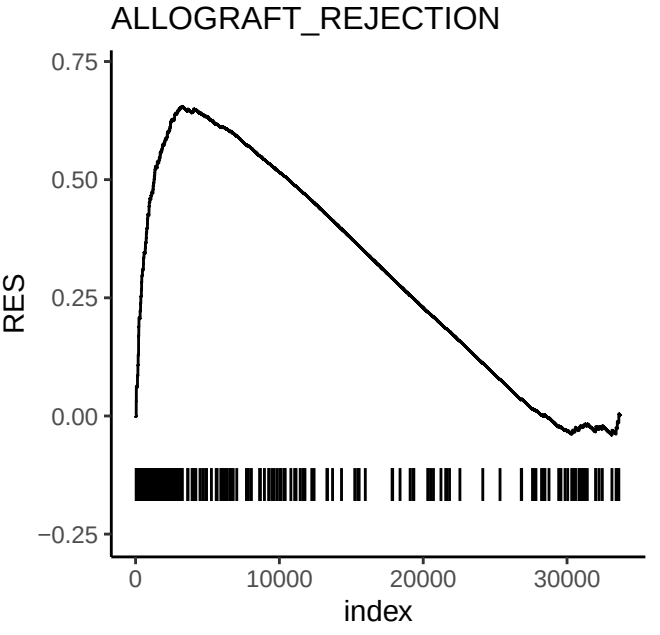

ANGIOGENESIS
RES
0.8
0.4
0.0
−0.4
0
10000
20000
30000
index

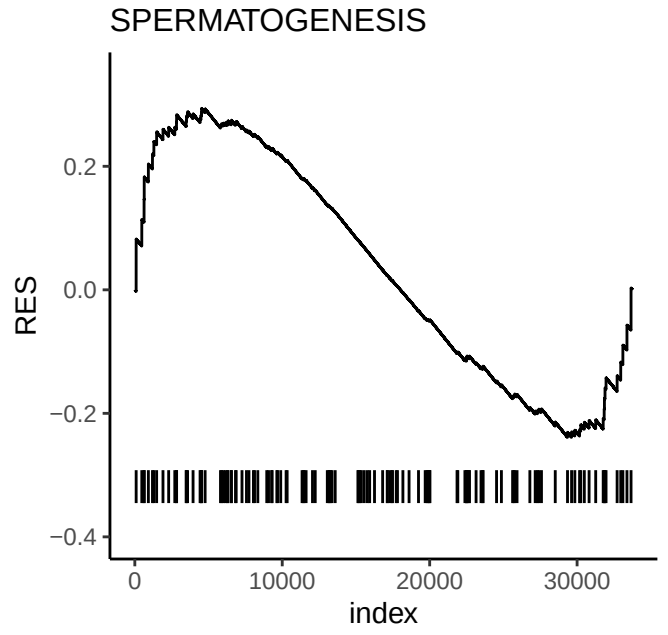

COAGULATION
RES
0.6
0.4
0.2
0.0
−0.2
−0.4
0
10000
20000
30000
index

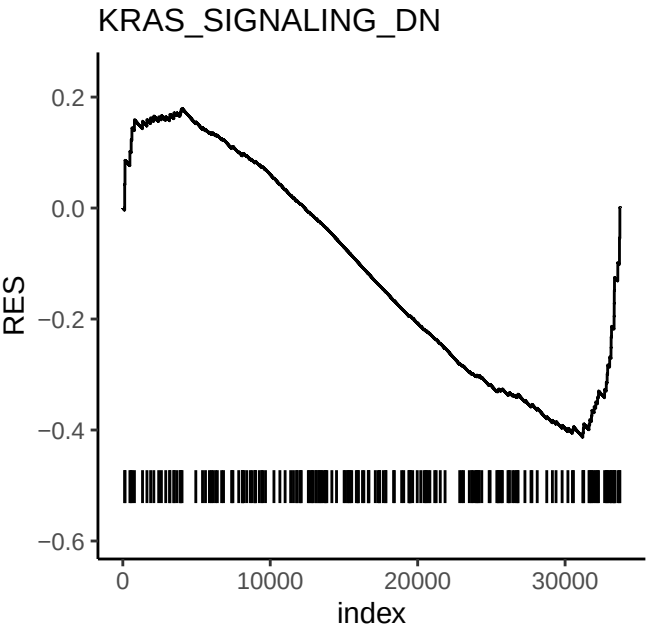

IL2_STAT5_SIGNALING
RES
0.50
0.25
0.00
−0.25
−0.50
0
10000
20000
30000
index

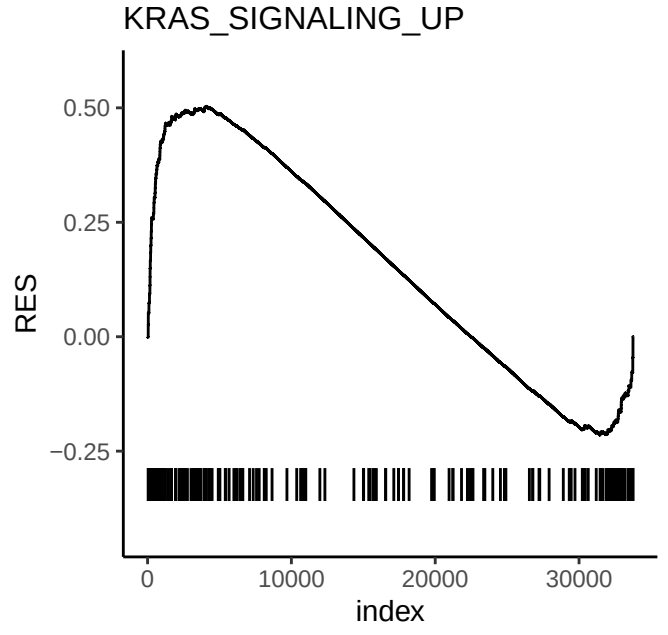

PEROXISOME
RES
0.50
0.25
0.00
−0.25
0
10000
20000
30000
index

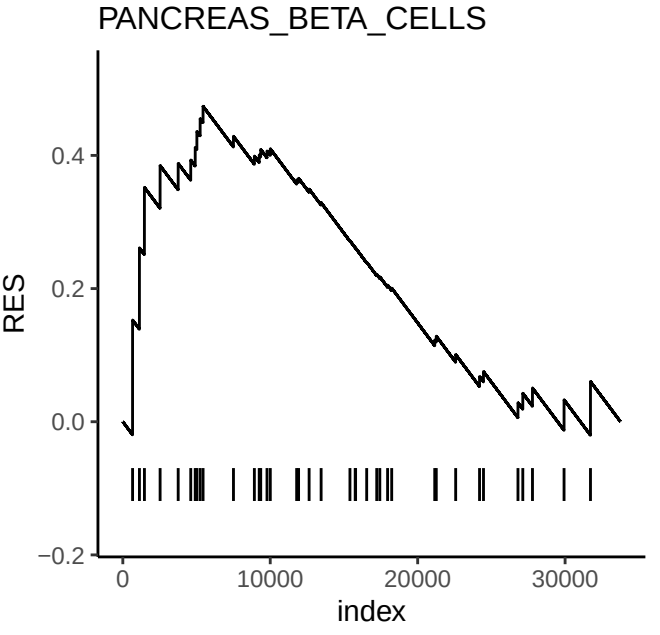

BILE_ACID_METABOLISM
RES
0.4
0.2
0.0
−0.2
−0.4
0
10000
20000
30000
index


ALLOGRAFT_REJECTION
RES
0.75
0.50
0.25
0.00
−0.25
0
10000
20000
30000
index


SPERMATOGENESIS
RES
0.2
0.0
−0.2
−0.4
0
10000
20000
30000
index


KRAS_SIGNALING_DN
RES
0.2
0.0
−0.2
−0.4
−0.6
0
10000
20000
30000
index


KRAS_SIGNALING_UP
RES
0.50
0.25
0.00
−0.25
0
10000
20000
30000
index


PANCREAS_BETA_CELLS
RES
0.4
0.2
0.0
−0.2
0
10000
20000
30000
index