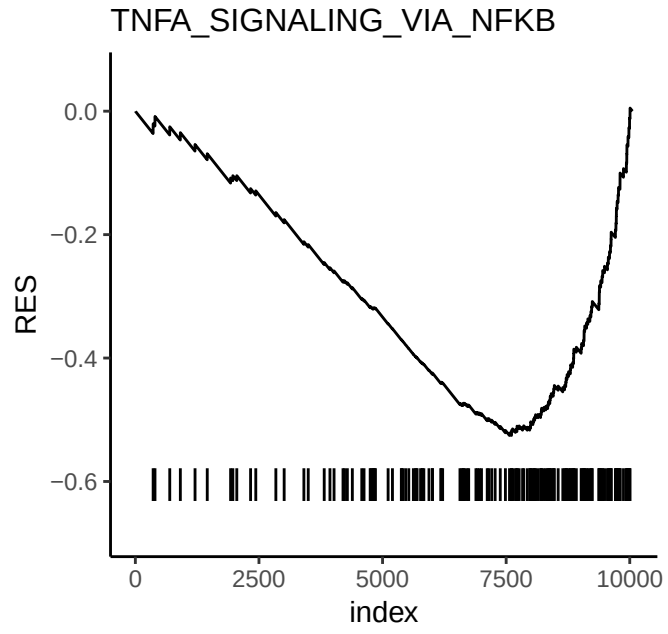
TNFA_SIGNALING_VIA_NFKB
RES
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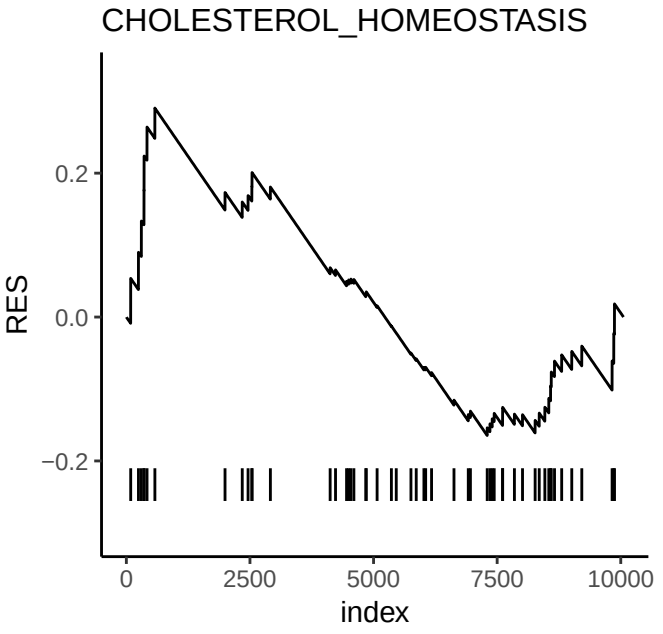
−0.6
0
2500
5000
7500
10000
index


CHOLESTEROL_HOMEOSTASIS
RES
0.2
0.0
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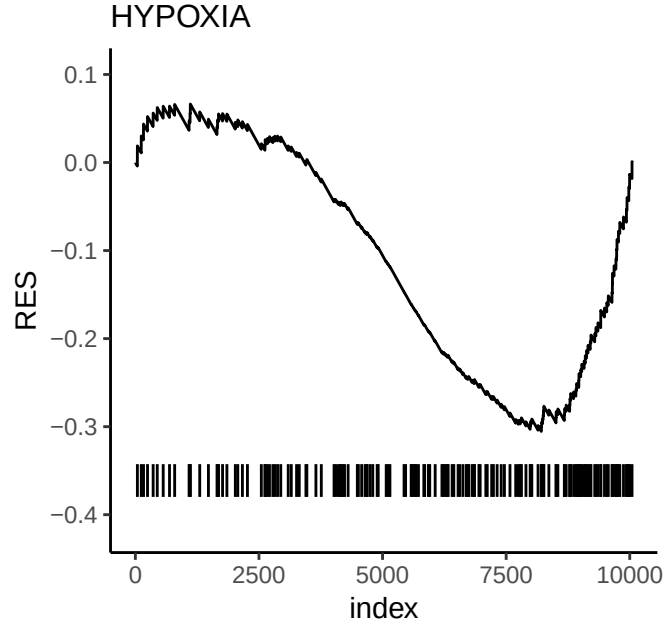
5000
7500
10000
index


HYPOXIA
RES
0.1
0.0
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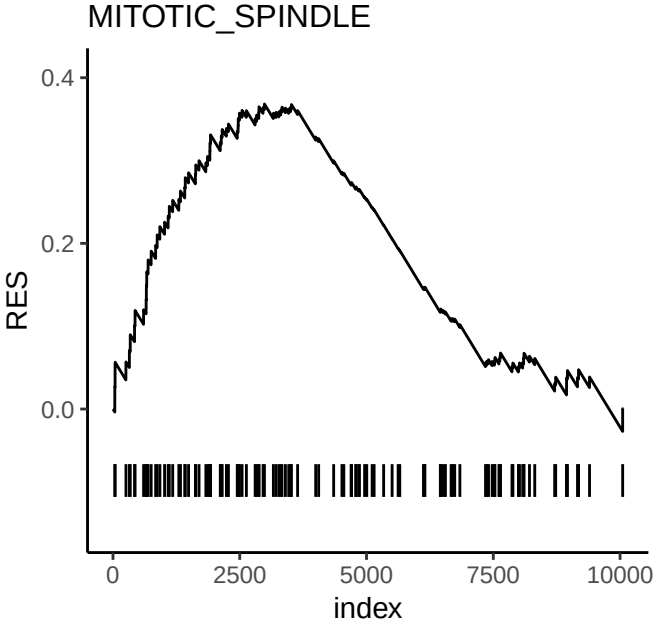
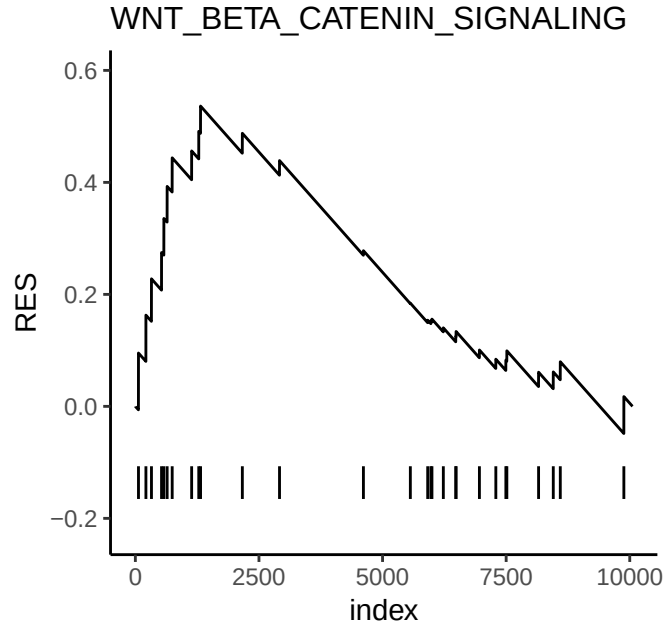
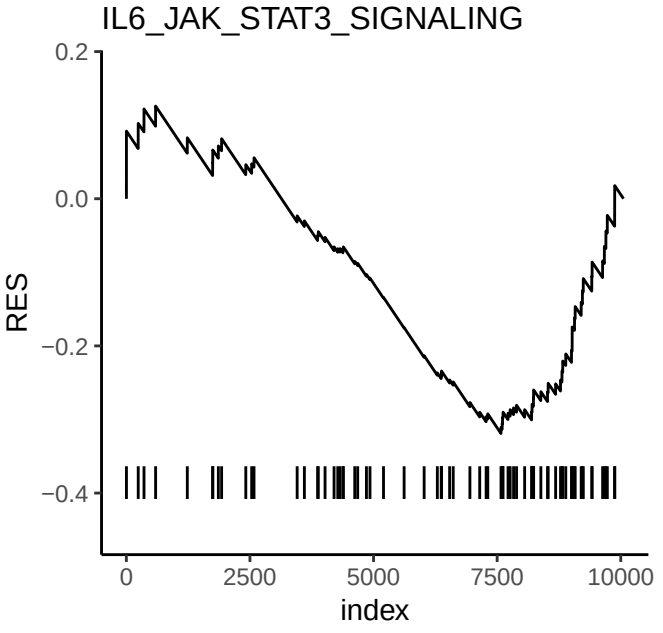
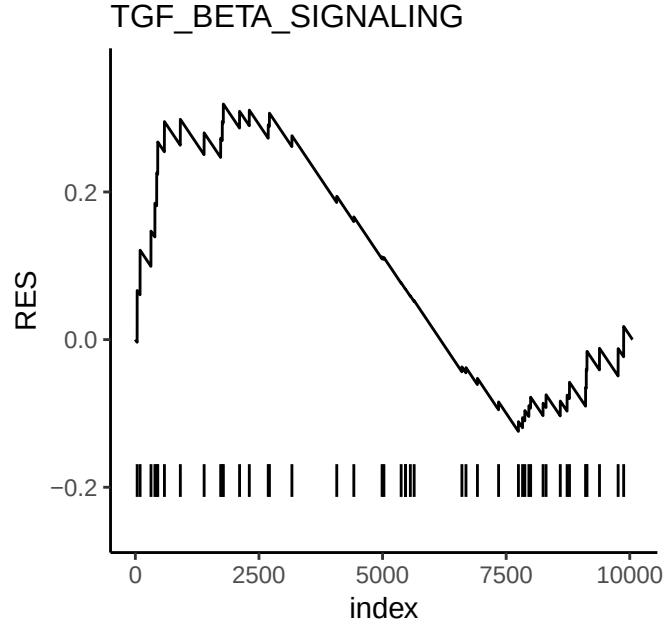
−0.2
−0.3
−0.4
0
2500
5000
7500
10000
index


MITOTIC_SPINDLE
RES
0.4
0.2
0.0
0
2500
5000
7500
10000
index


WNT_BETA_CATENIN_SIGNALING
RES
0.6
0.4
0.2
0.0
−0.2
0
2500
5000
7500
10000
index


IL6_JAK_STAT3_SIGNALING
RES
0.2
0.0
−0.2
−0.4
0
2500
5000
7500
10000
index


TGF_BETA_SIGNALING
RES
0.2
0.0
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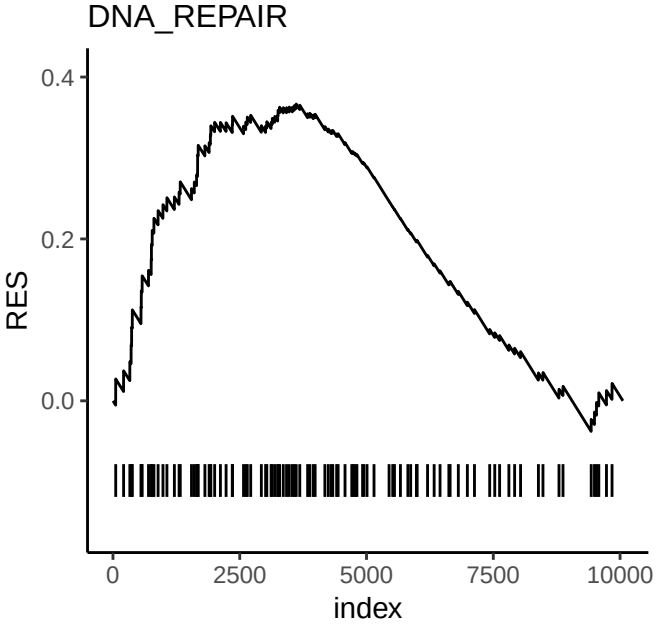
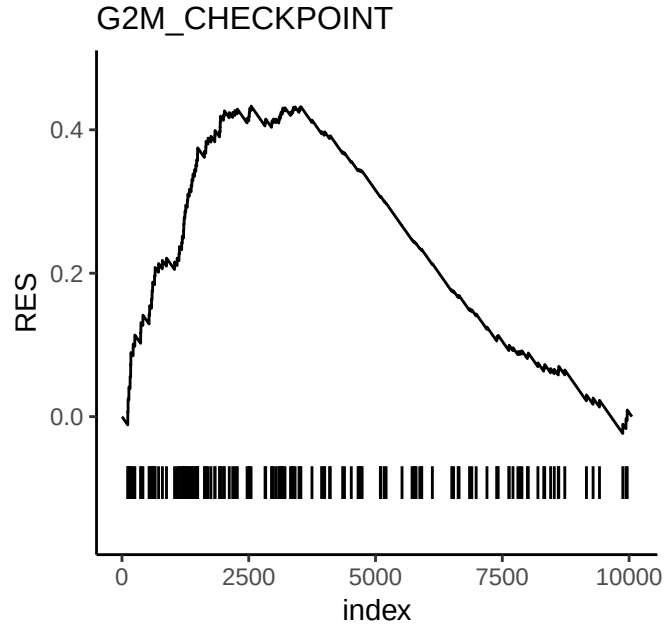
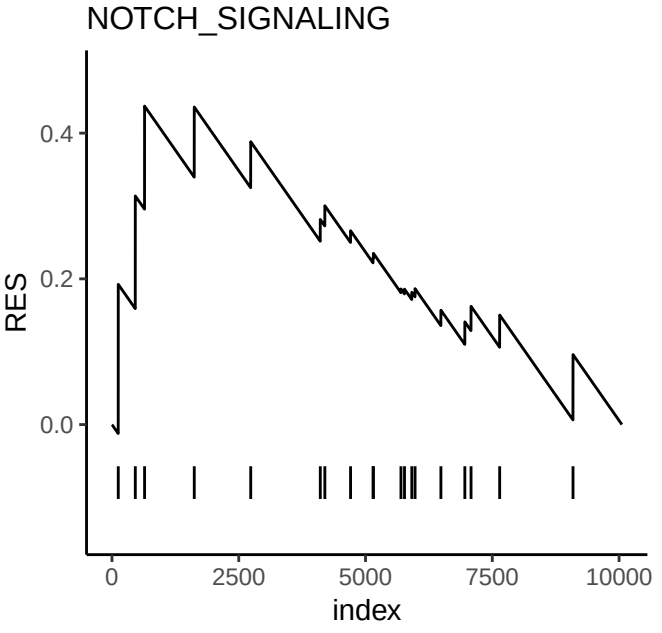
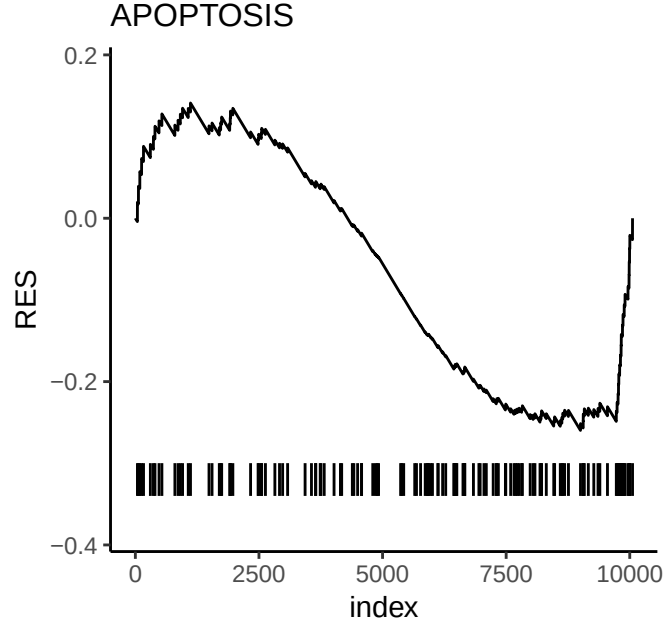
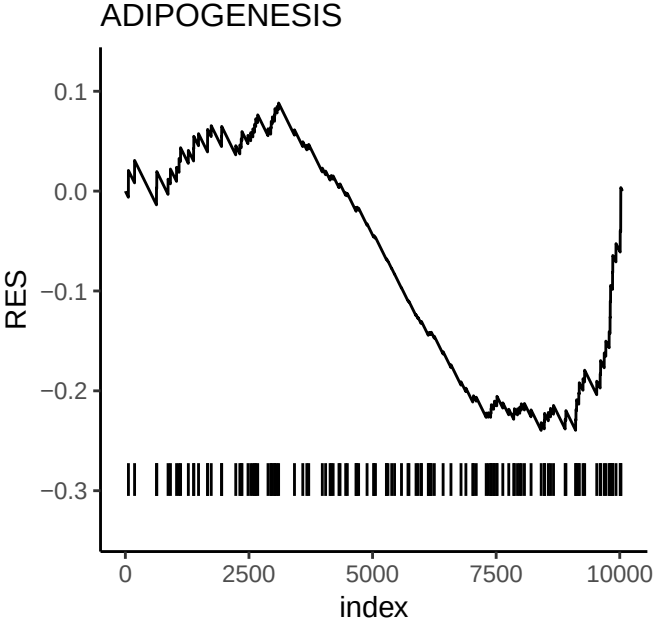
0
2500
5000
7500
10000
index

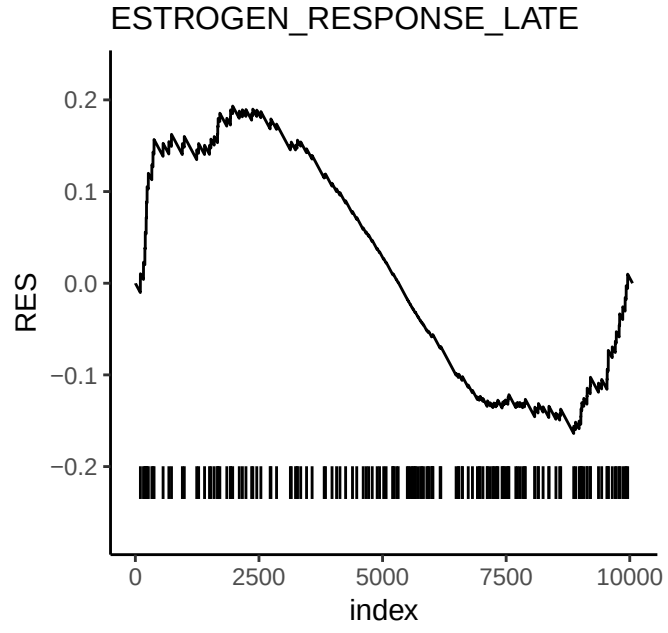

DNA_REPAIR
RES
0.4
0.2
0.0
0
2500
5000
7500
10000
index

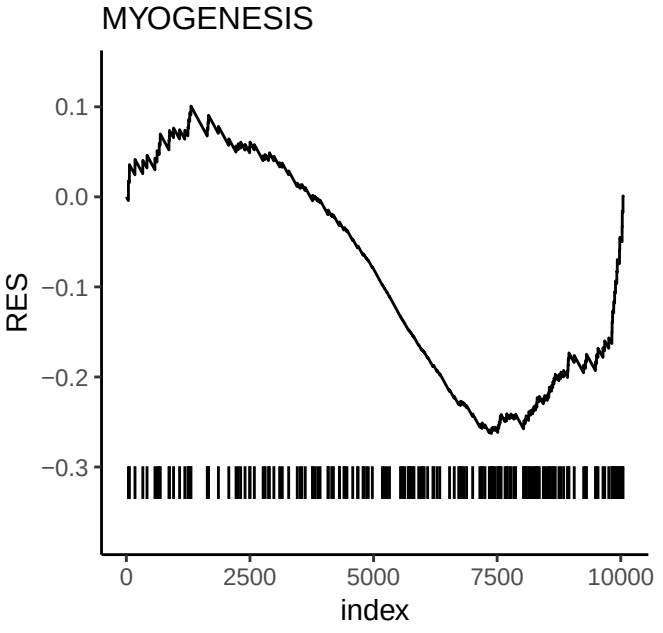

G2M_CHECKPOINT
RES
0.4
0.2
0.0
0
2500
5000
7500
10000
index

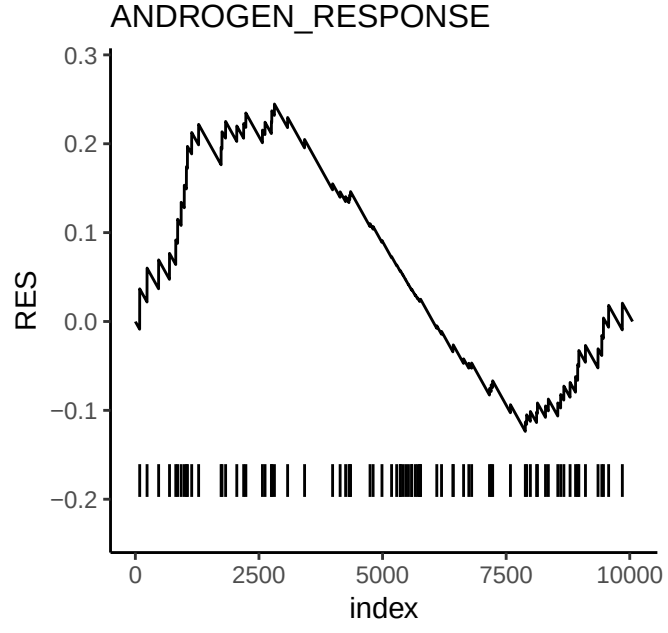

NOTCH_SIGNALING
RES
0.4
0.2
0.0
0
2500
5000
7500
10000
index

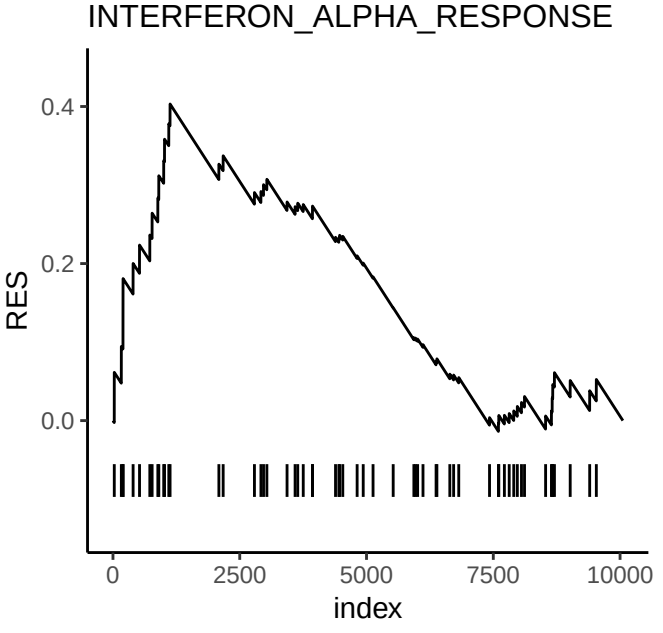

APOPTOSIS
RES
0.2
0.0
−0.2
−0.4
0
2500
5000
7500
10000
index

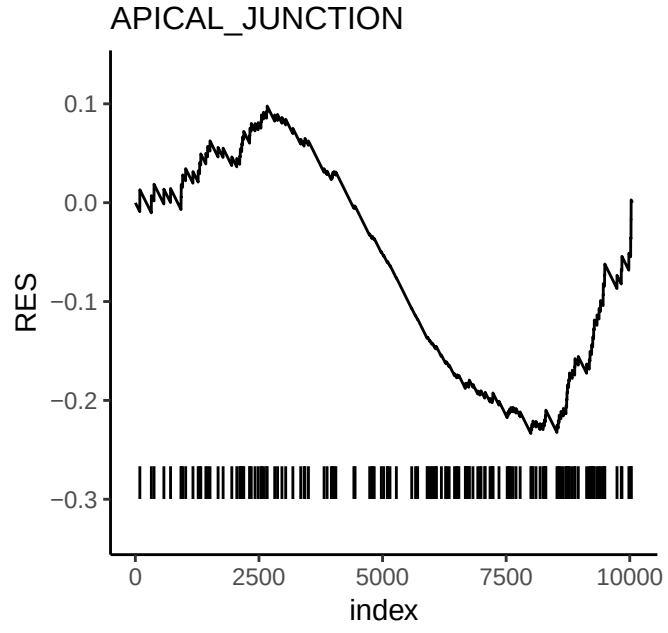

ADIPOGENESIS
RES
0.1
0.0
−0.1
−0.2
−0.3
0
2500
5000
7500
10000
index

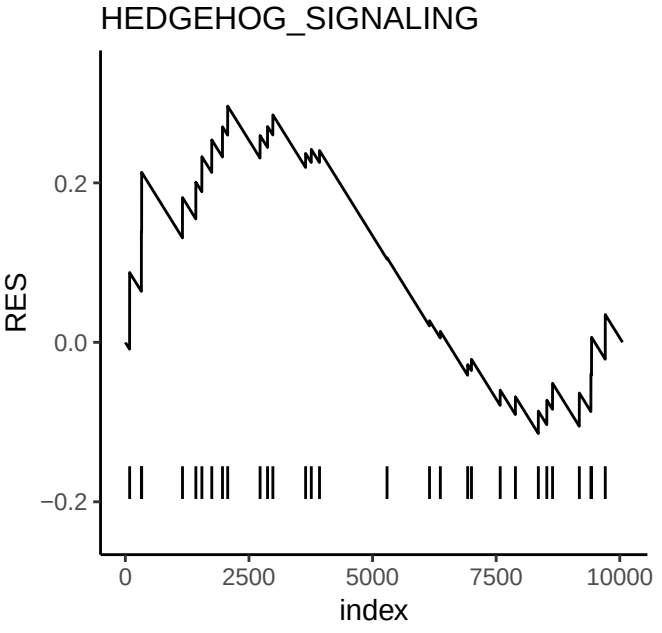

ESTROGEN_RESPONSE_LATE
RES
0.2
0.1
0.0
−0.1
−0.2
0
2500
5000
7500
10000
index

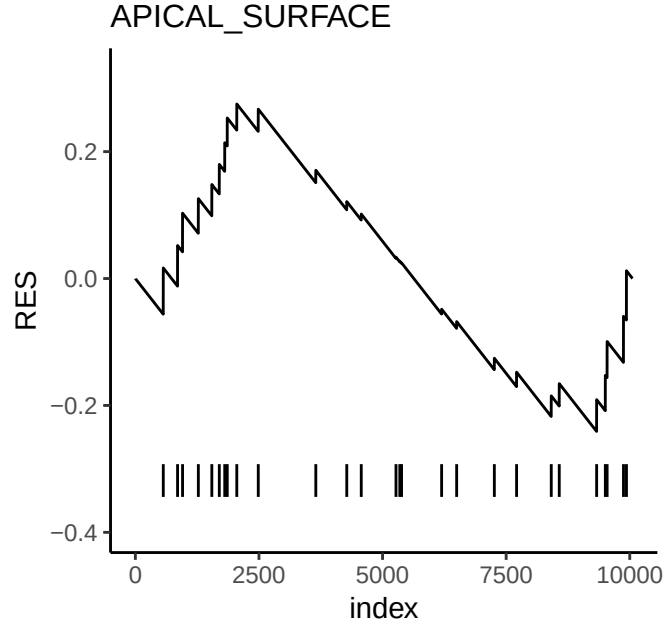

MYOGENESIS
RES
0.1
0.0
−0.1
−0.2
−0.3
0
2500
5000
7500
10000
index

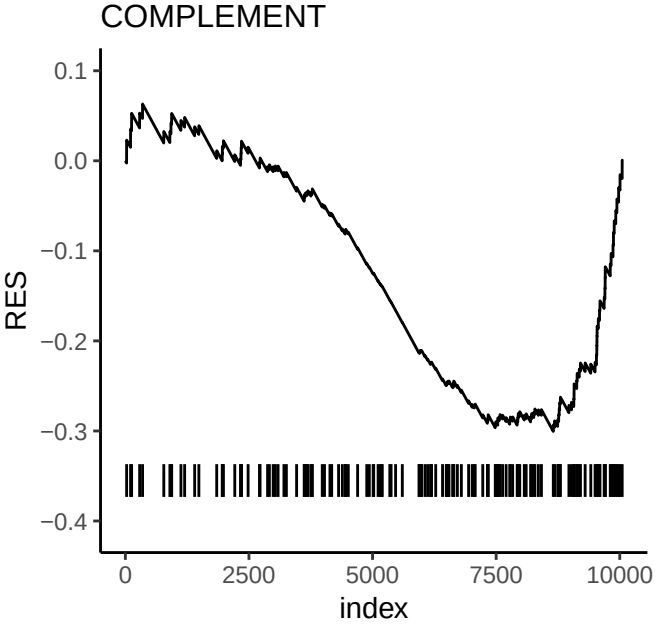

ANDROGEN_RESPONSE
RES
0.3
0.2
0.1
0.0
−0.1
−0.2
0
2500
5000
7500
10000
index


INTERFERON_ALPHA_RESPONSE
RES
0.4
0.2
0.0
0
2500
5000
7500
10000
index


APICAL_JUNCTION
RES
0.1
0.0
−0.1
−0.2
−0.3
0
2500
5000
7500
10000
index


HEDGEHOG_SIGNALING
RES
0.2
0.0
−0.2
0
2500
5000
7500
10000
index


APICAL_SURFACE
RES
0.2
0.0
−0.2
−0.4
0
2500
5000
7500
10000
index


COMPLEMENT
RES
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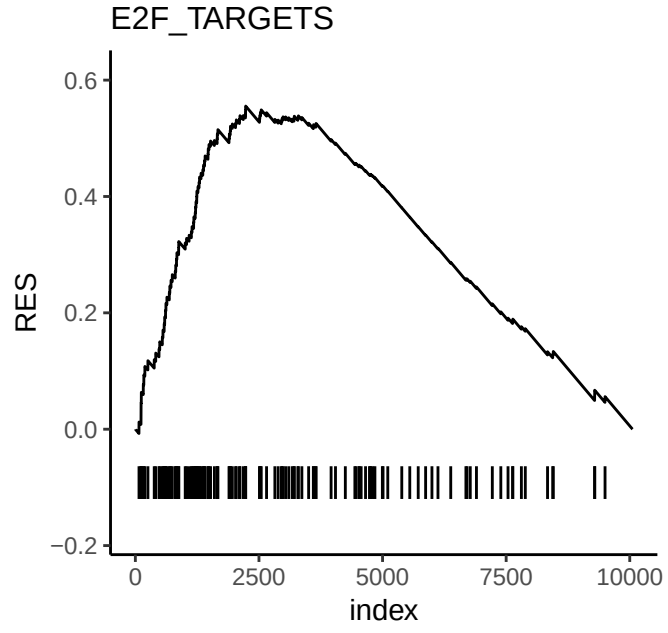
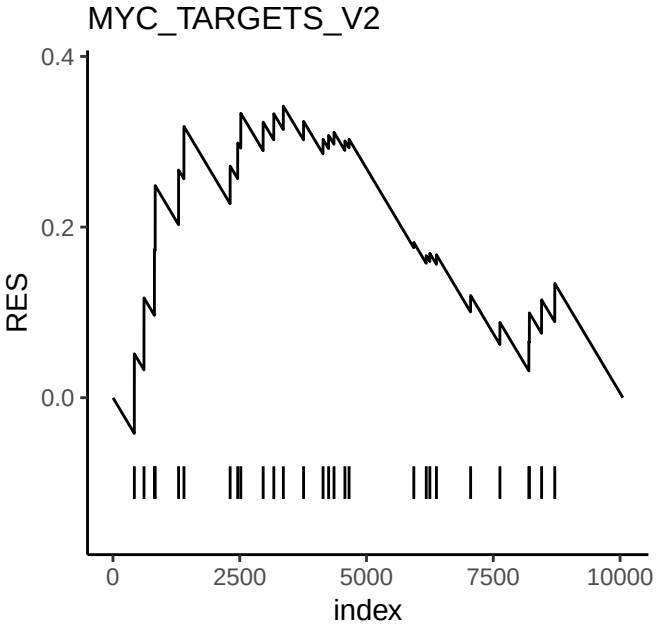
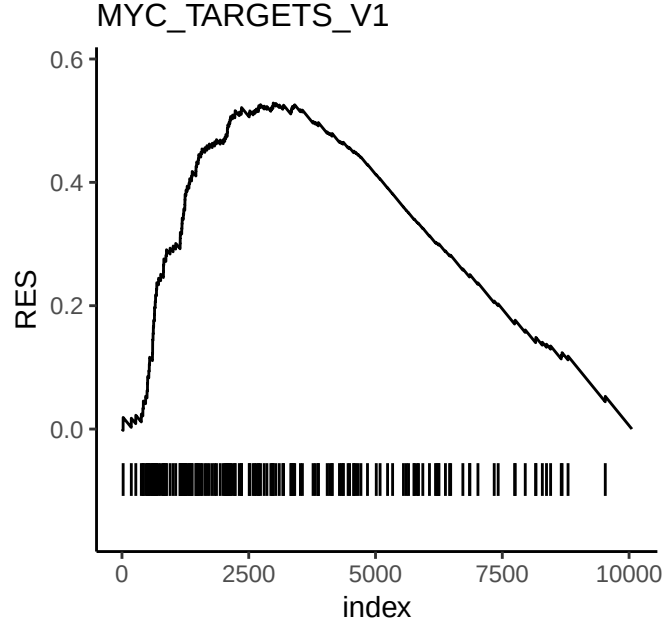
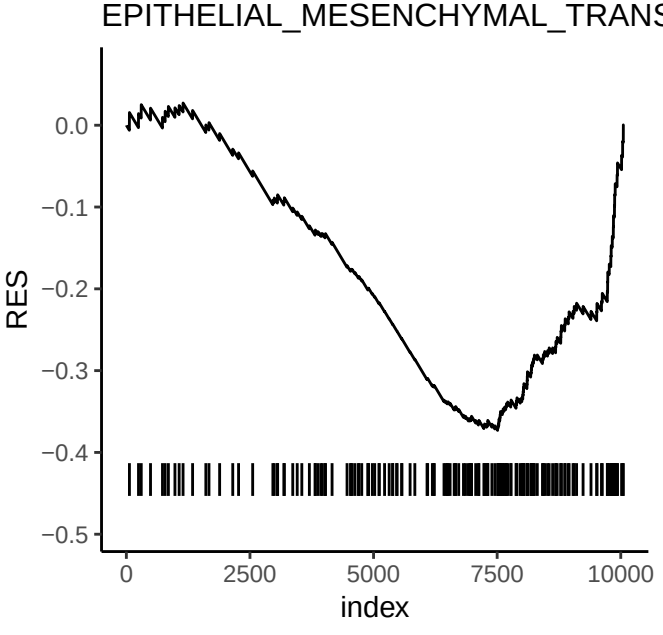
0.0
−0.1
−0.2
−0.3
−0.4
0
2500
5000
7500
10000
index


E2F_TARGETS
RES
0.6
0.4
0.2
0.0
−0.2
0
2500
5000
7500
10000
index


MYC_TARGETS_V2
RES
0.4
0.2
0.0
0
2500
5000
7500
10000
index


MYC_TARGETS_V1
RES
0.6
0.4
0.2
0.0
0
2500
5000
7500
10000
index


EPITHELIAL_MESENCHYMAL_TRANS
RES
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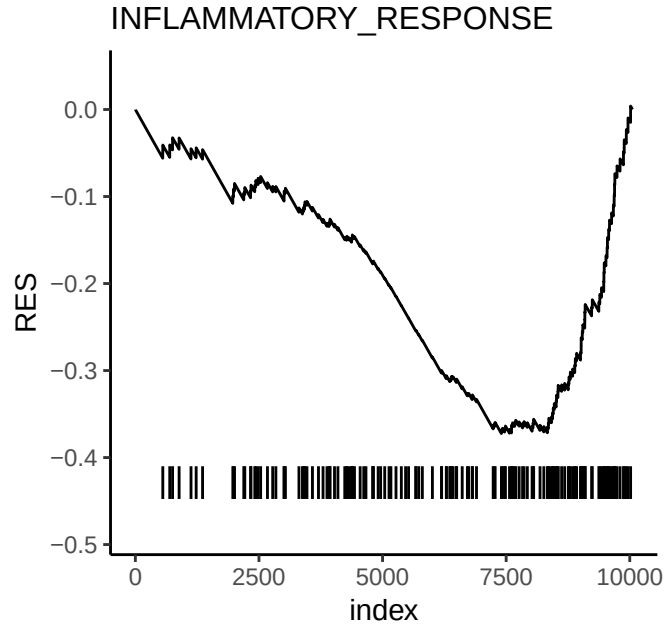
−0.1
−0.2
−0.3
−0.4
−0.5
0
2500
5000
7500
10000
index


INFLAMMATORY_RESPONSE
RES
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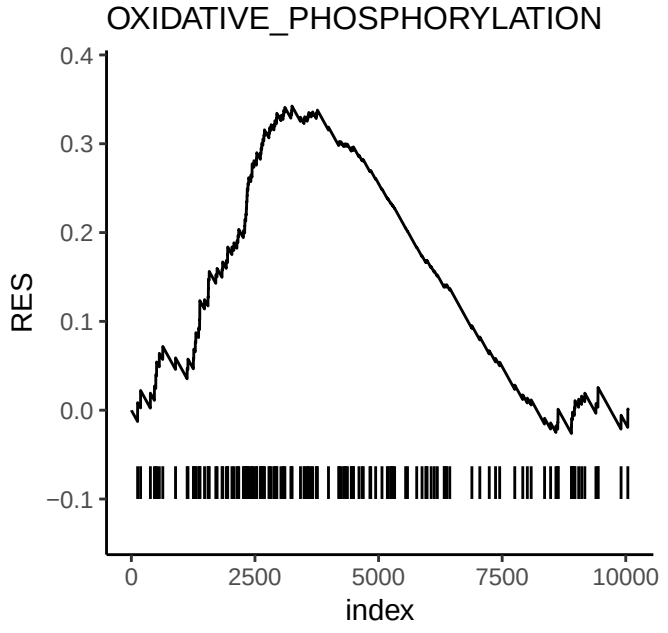
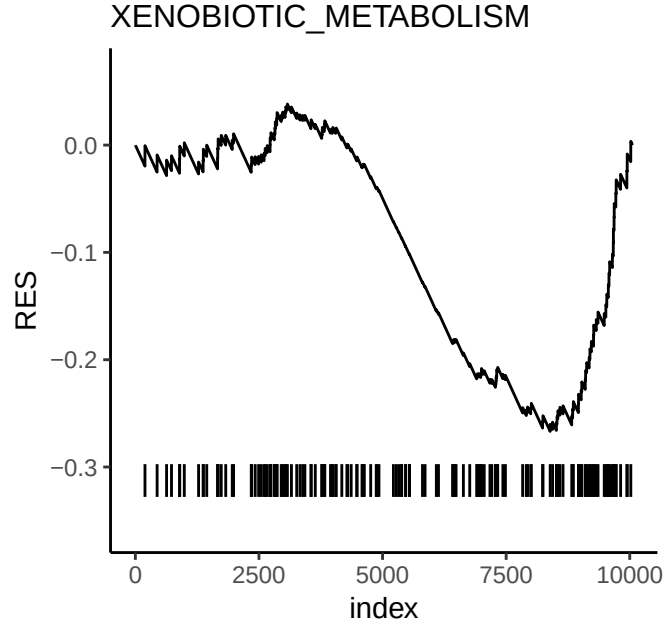
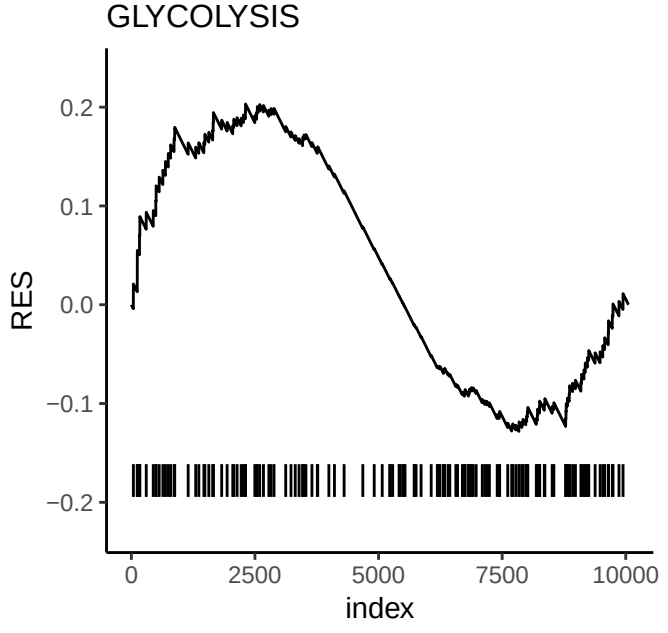
−0.1
−0.2
−0.3
−0.4
−0.5
0
2500
5000
7500
10000
index


OXIDATIVE_PHOSPHORYLATION
RES
0.4
0.3
0.2
0.1
0.0
−0.1
0
2500
5000
7500
10000
index


XENOBIOTIC_METABOLISM
RES
0.0
−0.1
−0.2
−0.3
0
2500
5000
7500
10000
index


GLYCOLYSIS
RES
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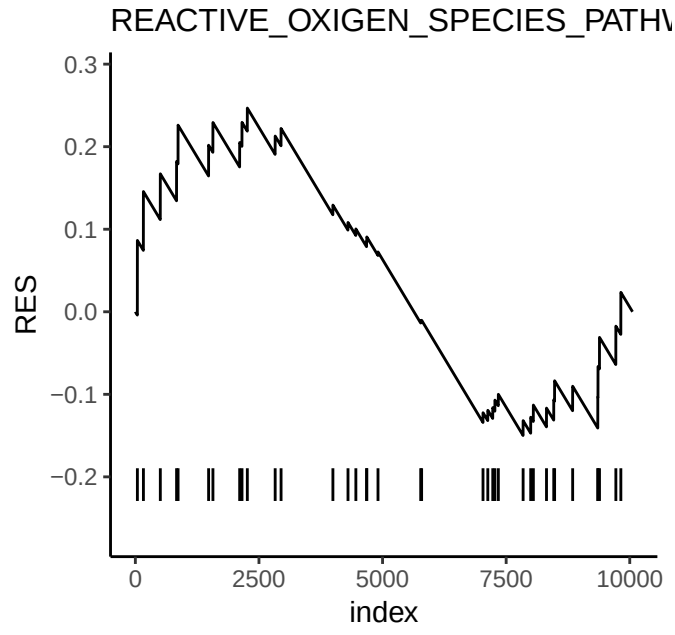
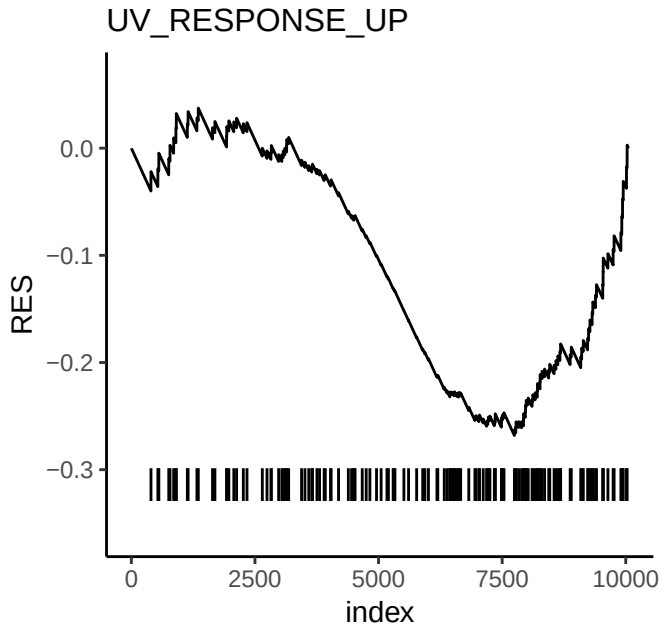
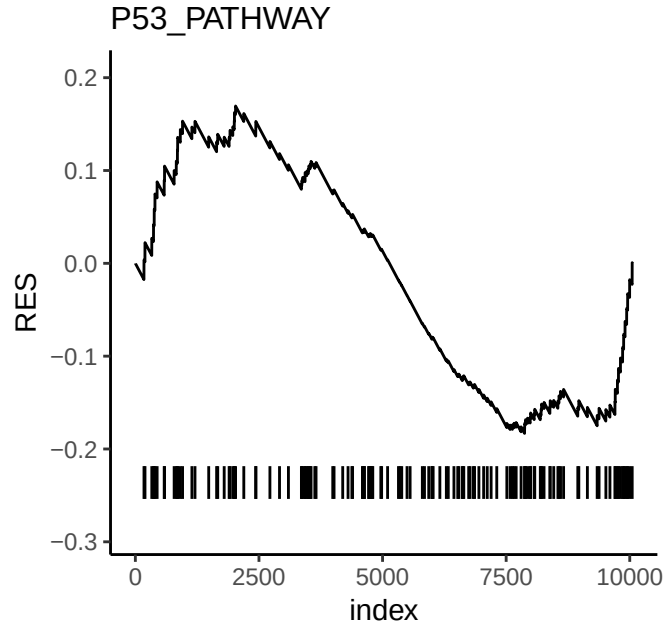
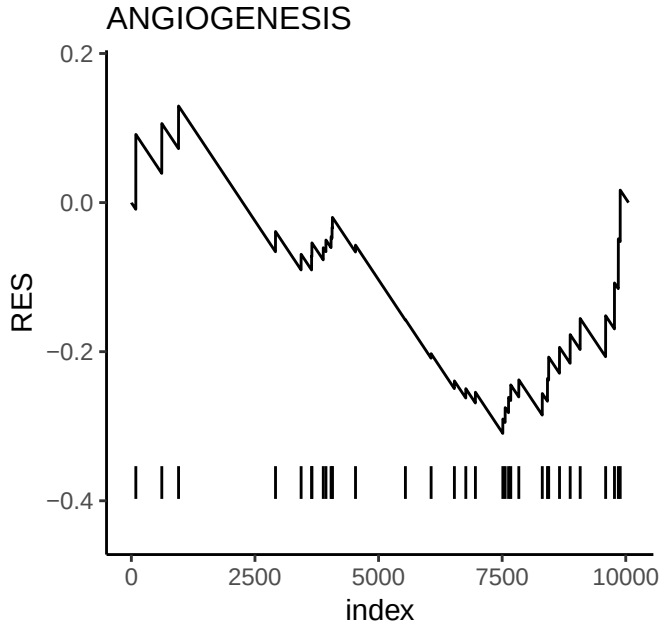
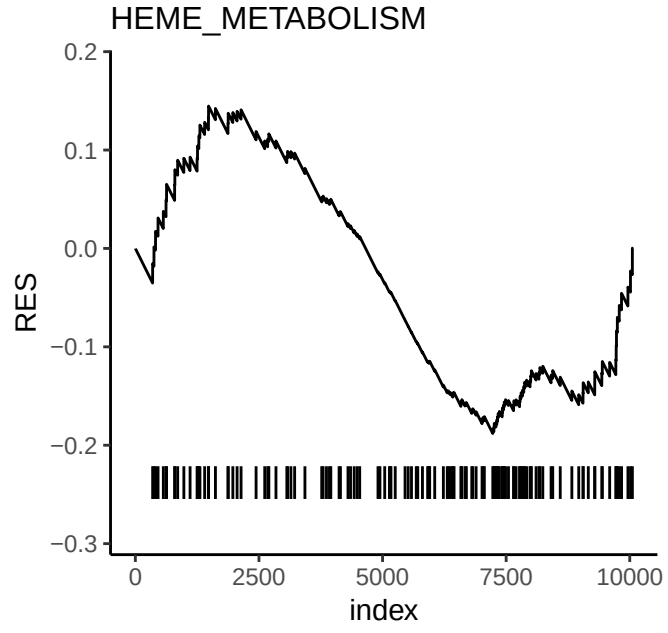
0.1
0.0
−0.1
−0.2
0
2500
5000
7500
10000
index

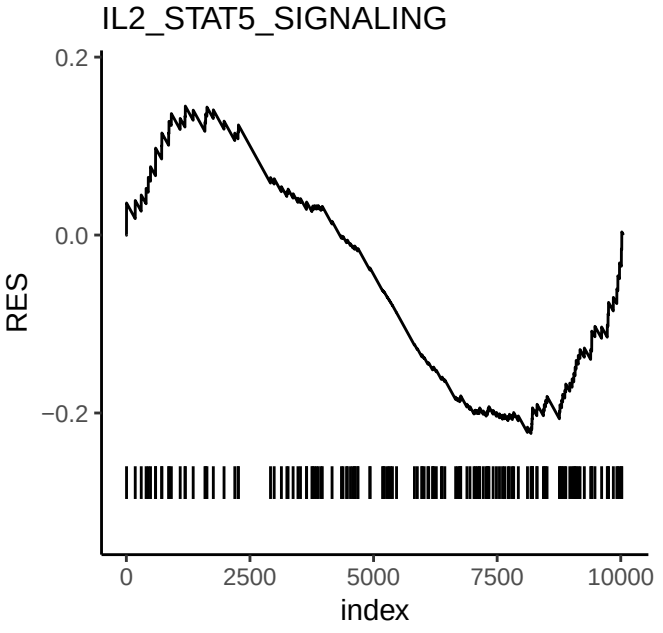

REACTIVE_OXIGEN_SPECIES_PATH
RES
index
0.3
0.2
0.1
0.0
−0.1
−0.2
0
2500
5000
7500
10000

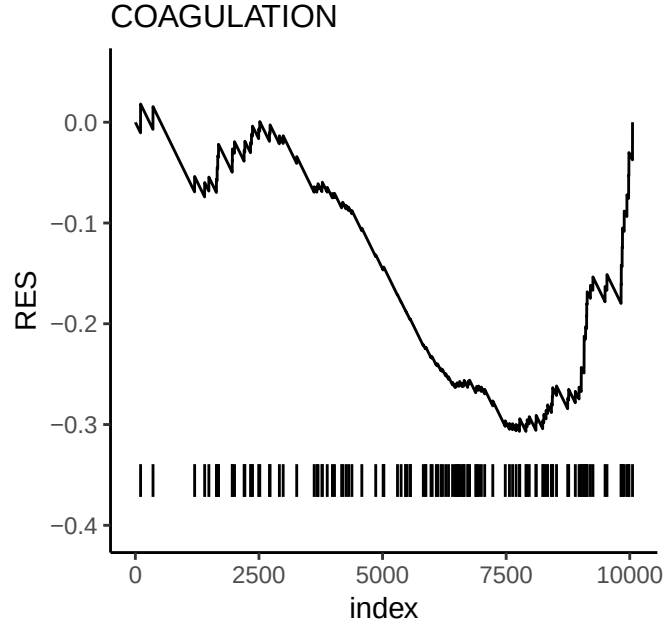

UV_RESPONSE_UP
RES
index
0.0
−0.1
−0.2
−0.3
0
2500
5000
7500
10000

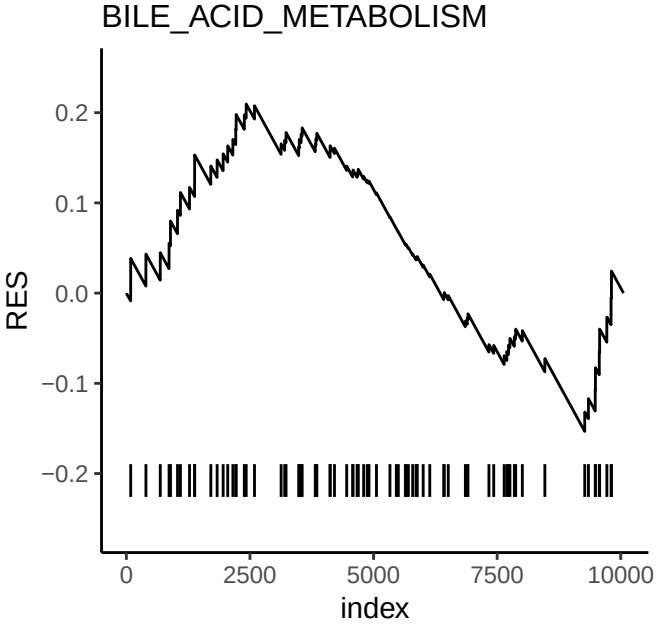

P53_PATHWAY
RES
index
0.2
0.1
0.0
−0.1
−0.2
−0.3
0
2500
5000
7500
10000

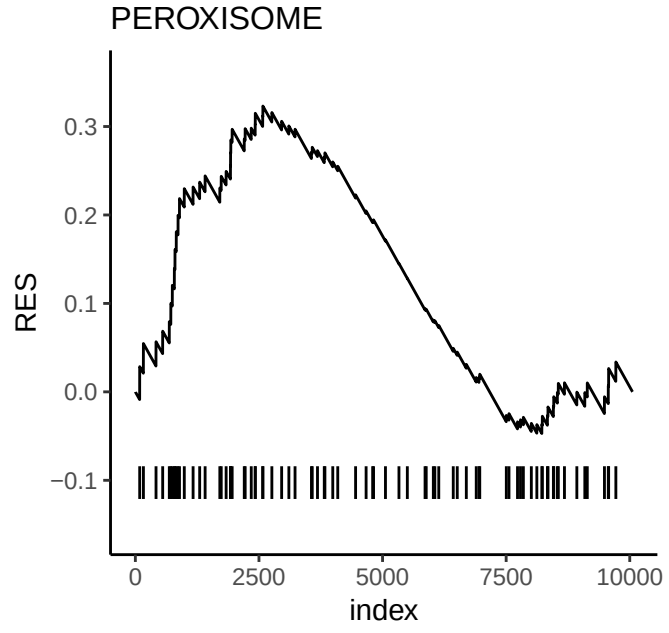

ANGIOGENESIS
RES
index
0.2
0.0
−0.2
−0.4
0
2500
5000
7500
10000

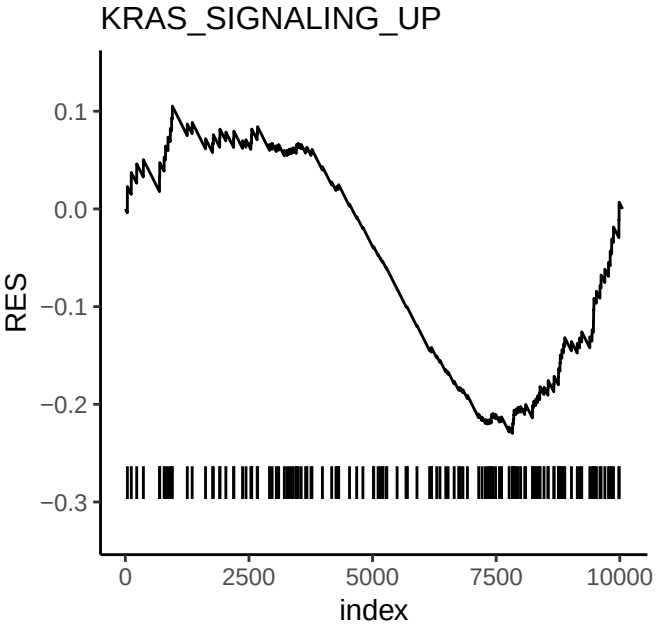

HEME_METABOLISM
RES
index
0.2
0.1
0.0
−0.1
−0.2
−0.3
0
2500
5000
7500
10000

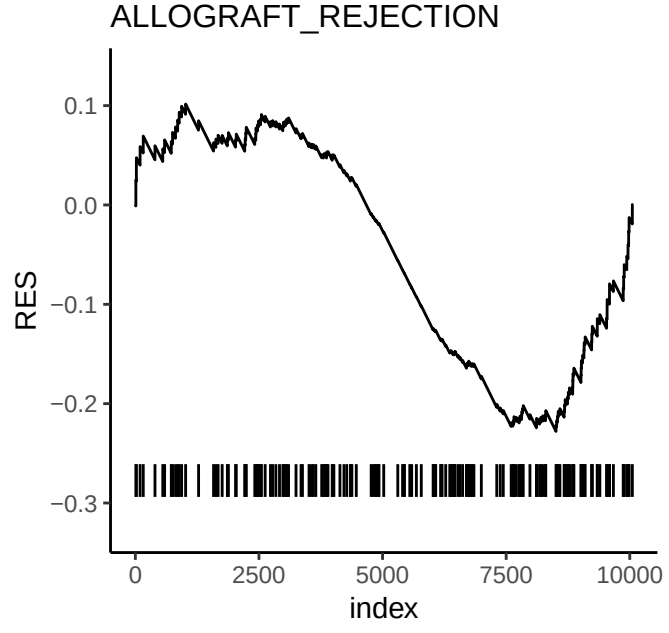

IL2_STAT5_SIGNALING
RES
index
0.2
0.0
−0.2
0
2500
5000
7500
10000

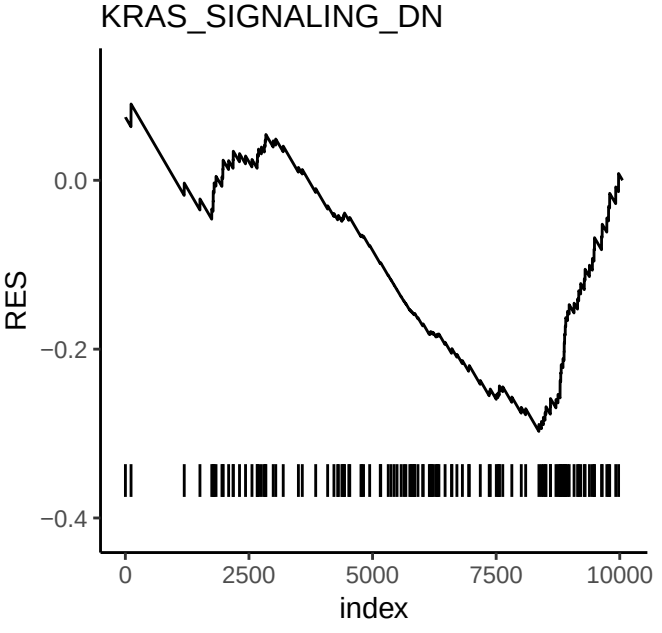

COAGULATION
RES
index
0.0
−0.1
−0.2
−0.3
−0.4
0
2500
5000
7500
10000


BILE_ACID_METABOLISM
RES
index
0.2
0.1
0.0
−0.1
−0.2
0
2500
5000
7500
10000


PEROXISOME
RES
0.3
0.2
0.1
0.0
−0.1
0
2500
5000
7500
10000
index


KRAS_SIGNALING_UP
RES
0.1
0.0
−0.1
−0.2
−0.3
0
2500
5000
7500
10000
index


ALLOGRAFT_REJECTION
RES
0.1
0.0
−0.1
−0.2
−0.3
0
2500
5000
7500
10000
index


KRAS_SIGNALING_DN
RES
0.0
−0.2
−0.4
0
2500
5000
7500
10000
index