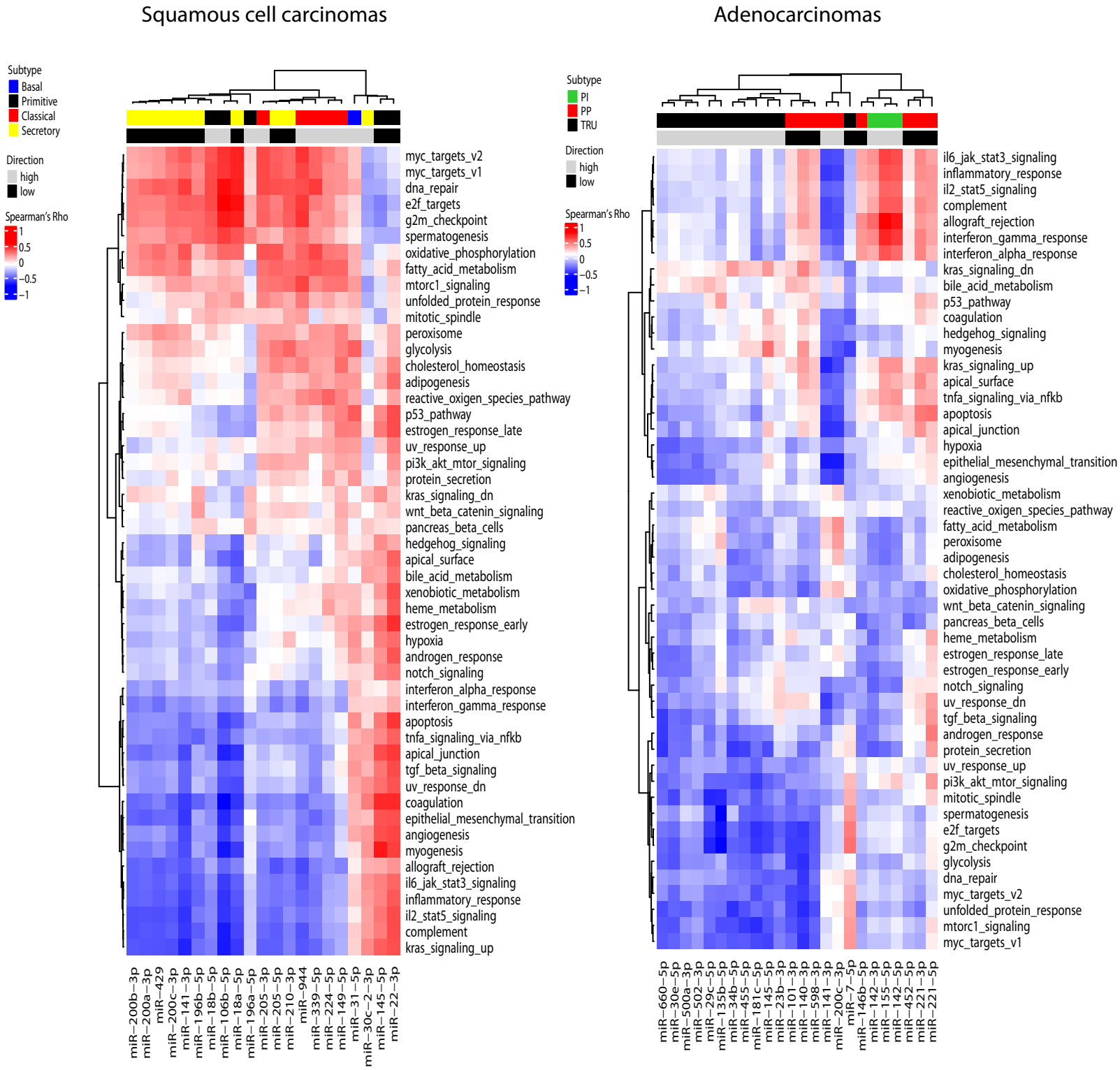
Squamous cell carcinomas
Subtype
Basal
Primitive
Classical
Secretory
Direction
high
low
Spearman's Rho
1
0.5
0
−0.5
−1
myc_targets_v2
myc_targets_v1
dna_repair
e2f_targets
g2m_checkpoint
spermatogenesis
oxidative_phosphorylation
fatty_acid_metabolism
mtorc1_signaling
unfolded_protein_response
mitotic_spindle
peroxisome
glycolysis
cholesterol_homeostasis
adipogenesis
reactive_oxigen_species_pathway
p53_pathway
estrogen_response_late
uv_response_up
pi3k_akt_mtor_signaling
protein_secretion
kras_signaling_dn
wnt_beta_catenin_signaling
pancreas_beta_cells
hedgehog_signaling
apical_surface
bile_acid_metabolism
xenobiotic_metabolism
heme_metabolism
estrogen_response_early
hypoxia
androgen_response
notch_signaling
interferon_alpha_response
interferon_gamma_response
apoptosis
tnfa_signaling_via_nfkb
apical_junction
tgf_beta_signaling
uv_response_dn
coagulation
epithelial_mesenchymal_transition
angiogenesis
myogenesis
allograft_rejection
il6_jak_stat3_signaling
inflammatory_response
il2_stat5_signaling
complement
kras_signaling_up
miR−200b−3p
miR−200a−3p
miR−429
miR−200c−3p
miR−141−3p
miR−196b−5p
miR−18b−5p
miR−106b−5p
miR−18a−5p
miR−196a−5p
miR−205−3p
miR−205−5p
miR−210−3p
miR−944
miR−339−5p
miR−224−5p
miR−149−5p
miR−31−5p
miR−30c−2−3p
miR−145−5p
miR−22−3p
Adenocarcinomas
Subtype
PI
PP
TRU
Direction
high
low
Spearman's Rho
1
0.5
0
−0.5
−1
il6_jak_stat3_signaling
inflammatory_response
il2_stat5_signaling
complement
allograft_rejection
interferon_gamma_response
interferon_alpha_response
kras_signaling_dn
bile_acid_metabolism
p53_pathway
coagulation
hedgehog_signaling
myogenesis
kras_signaling_up
apical_surface
tnfa_signaling_via_nfkb
apoptosis
apical_junction
hypoxia
epithelial_mesenchymal_transition
angiogenesis
xenobiotic_metabolism
reactive_oxigen_species_pathway
fatty_acid_metabolism
peroxisome
adipogenesis
cholesterol_homeostasis
oxidative_phosphorylation
wnt_beta_catenin_signaling
pancreas_beta_cells
heme_metabolism
estrogen_response_late
estrogen_response_early
notch_signaling
uv_response_dn
tgf_beta_signaling
androgen_response
protein_secretion
uv_response_up
pi3k_akt_mtor_signaling
mitotic_spindle
spermatogenesis
e2f_targets
g2m_checkpoint
glycolysis
dna_repair
myc_targets_v2
unfolded_protein_response
mtorc1_signaling
myc_targets_v1
miR−660−5p
miR−30e−5p
miR−500a−3p
miR−502−3p
miR−29c−5p
miR−135b−5p
miR−34b−5p
miR−455−5p
miR−181c−5p
miR−145−5p
miR−23b−3p
miR−101−3p
miR−140−3p
miR−598−3p
miR−141−3p
miR−200c−3p
miR−7−5p
miR−146b−5p
miR−142−3p
miR−155−5p
miR−142−5p
miR−452−5p
miR−221−3p
miR−221−5p