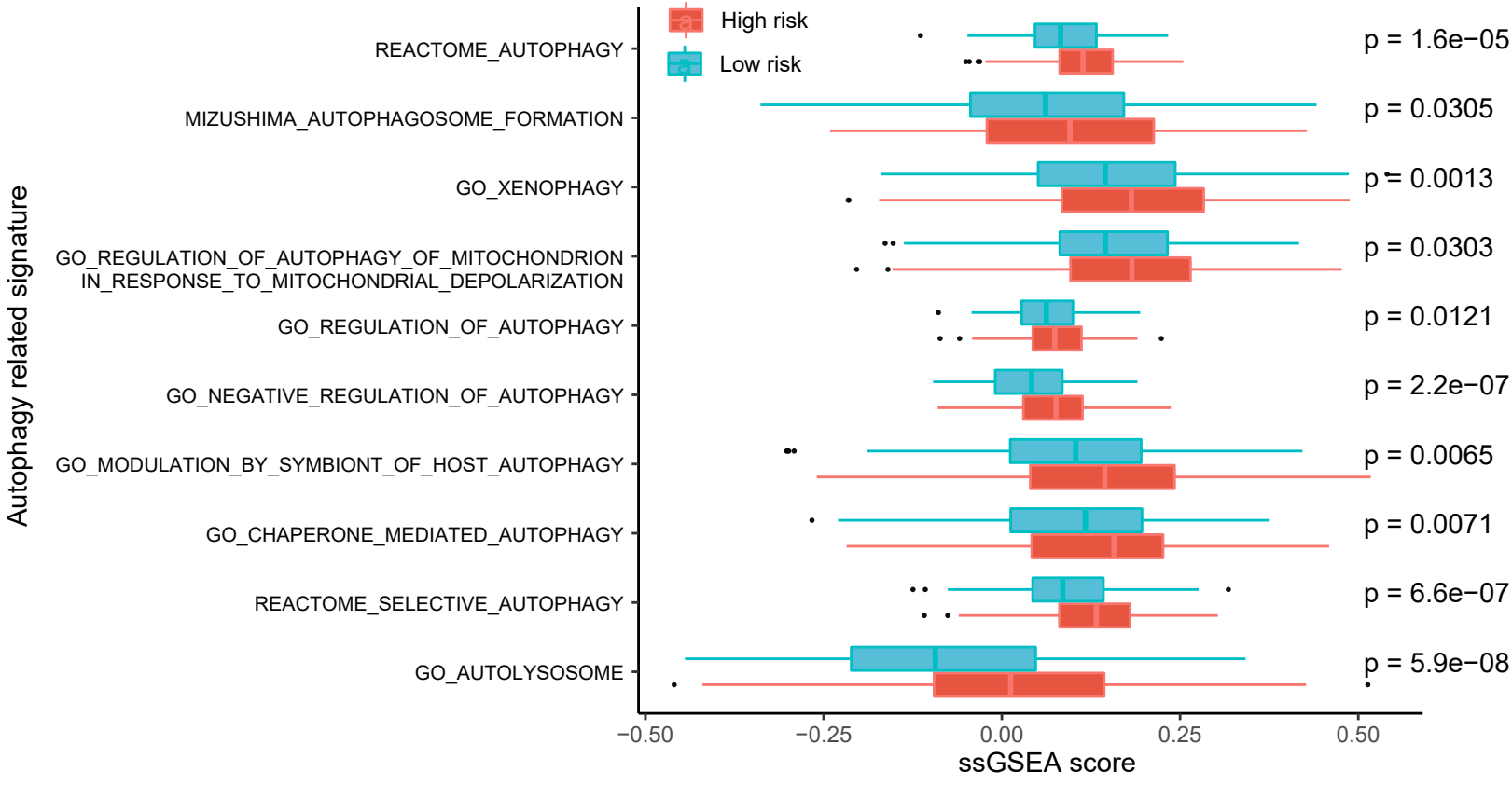
High risk
Low risk
Autophagy related signature
REACTOME_AUTOPHAGY
MIZUSHIMA_AUTOPHAGOSOME_FORMATION
GO_XENOPHAGY
GO_REGULATION_OF_AUTOPHAGY_OF_MITOCHONDRION
IN_RESPONSE_TO_MITOCHONDRIAL_DEPOLARIZATION
GO_REGULATION_OF_AUTOPHAGY
GO_NEGATIVE_REGULATION_OF_AUTOPHAGY
GO_MODULATION_BY_SYMBIONT_OF_HOST_AUTOPHAGY
GO_CHAPERONE_MEDIATED_AUTOPHAGY
REACTOME_SELECTIVE_AUTOPHAGY
GO_AUTOLYSOSOME
p = 1.6e−05
p = 0.0305
p = 0.0013
p = 0.0303
p = 0.0121
p = 2.2e−07
p = 0.0065
p = 0.0071
p = 6.6e−07
p = 5.9e−08
−0.50
−0.25
0.00
0.25
0.50
ssGSEA score