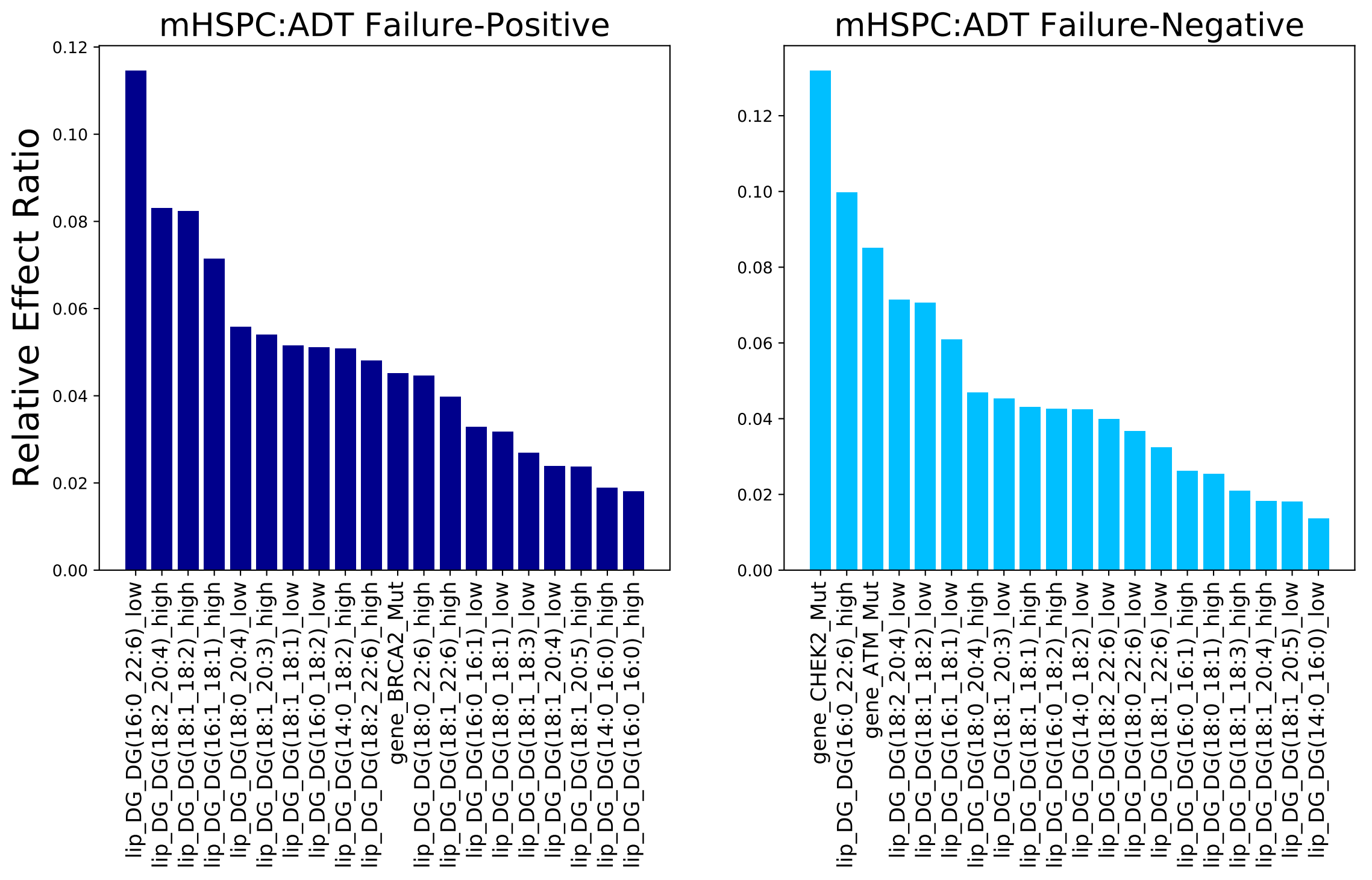
mHSPC:ADT Failure-Positive
Relative Effect Ratio
0.12
0.10
0.08
0.06
0.04
0.02
0.00
lip_DG_DG(16:0_22:6)_low
lip_DG_DG(18:2_20:4)_high
lip_DG_DG(18:1_18:2)_high
lip_DG_DG(16:1_18:1)_high
lip_DG_DG(18:0_20:4)_low
lip_DG_DG(18:1_20:3)_high
lip_DG_DG(18:1_18:1)_low
lip_DG_DG(16:0_18:2)_low
lip_DG_DG(14:0_18:2)_high
lip_DG_DG(18:2_22:6)_high
gene_BRCA2_Mut
lip_DG_DG(18:0_22:6)_high
lip_DG_DG(18:1_22:6)_high
lip_DG_DG(16:0_16:1)_low
lip_DG_DG(18:0_18:1)_low
lip_DG_DG(18:1_18:3)_low
lip_DG_DG(18:1_20:4)_low
lip_DG_DG(18:1_20:5)_high
lip_DG_DG(14:0_16:0)_high
lip_DG_DG(16:0_16:0)_high
mHSPC:ADT Failure-Negative
0.12
0.10
0.08
0.06
0.04
0.02
0.00
gene_CHEK2_Mut
lip_DG_DG(16:0_22:6)_high
gene_ATM_Mut
lip_DG_DG(18:2_20:4)_low
lip_DG_DG(18:1_18:2)_low
lip_DG_DG(16:1_18:1)_low
lip_DG_DG(18:0_20:4)_high
lip_DG_DG(18:1_20:3)_low
lip_DG_DG(18:1_18:1)_high
lip_DG_DG(16:0_18:2)_high
lip_DG_DG(14:0_18:2)_low
lip_DG_DG(18:2_22:6)_low
lip_DG_DG(18:0_22:6)_low
lip_DG_DG(18:1_22:6)_low
lip_DG_DG(16:0_16:1)_high
lip_DG_DG(18:0_18:1)_high
lip_DG_DG(18:1_18:3)_high
lip_DG_DG(18:1_20:4)_high
lip_DG_DG(18:1_20:5)_low
lip_DG_DG(14:0_16:0)_low