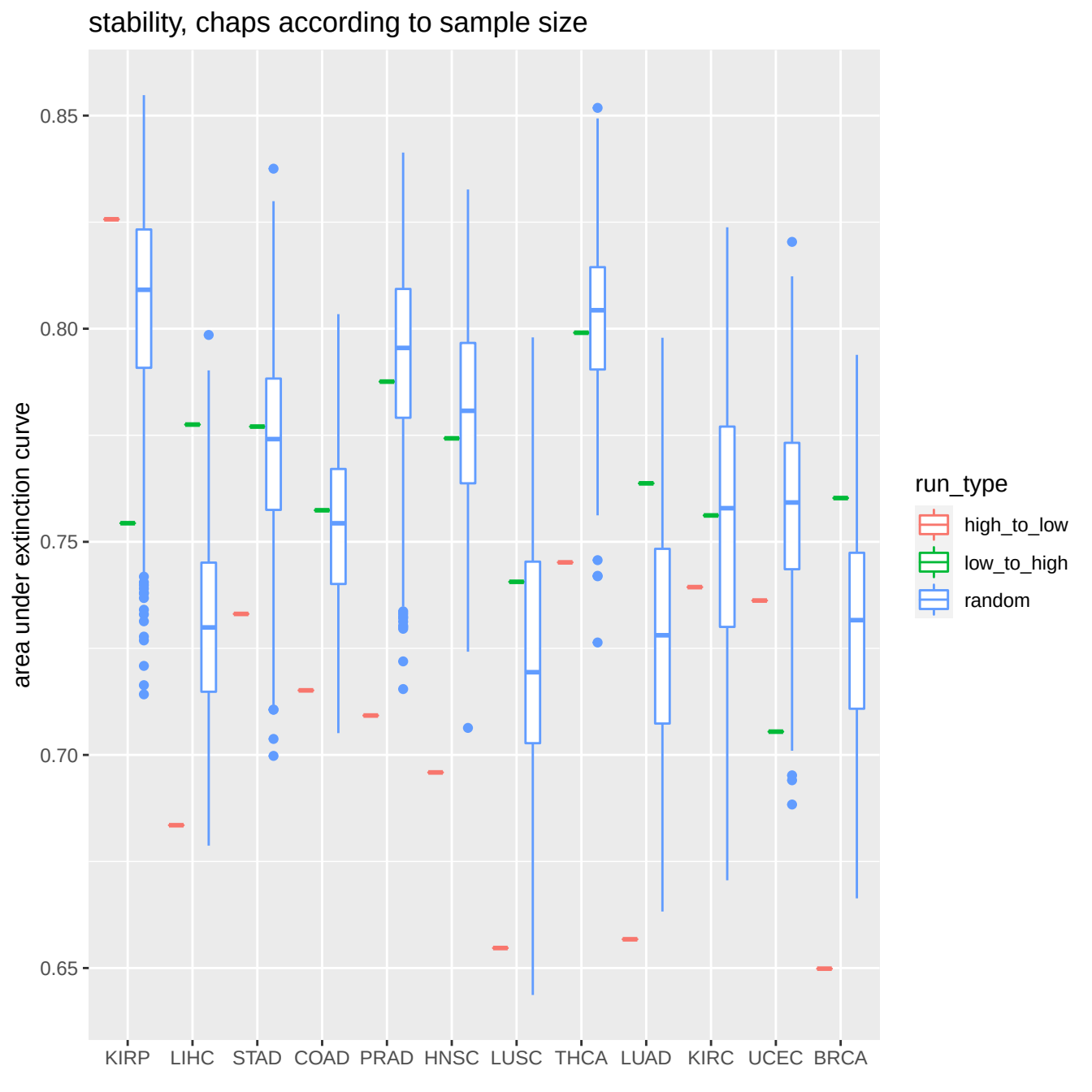
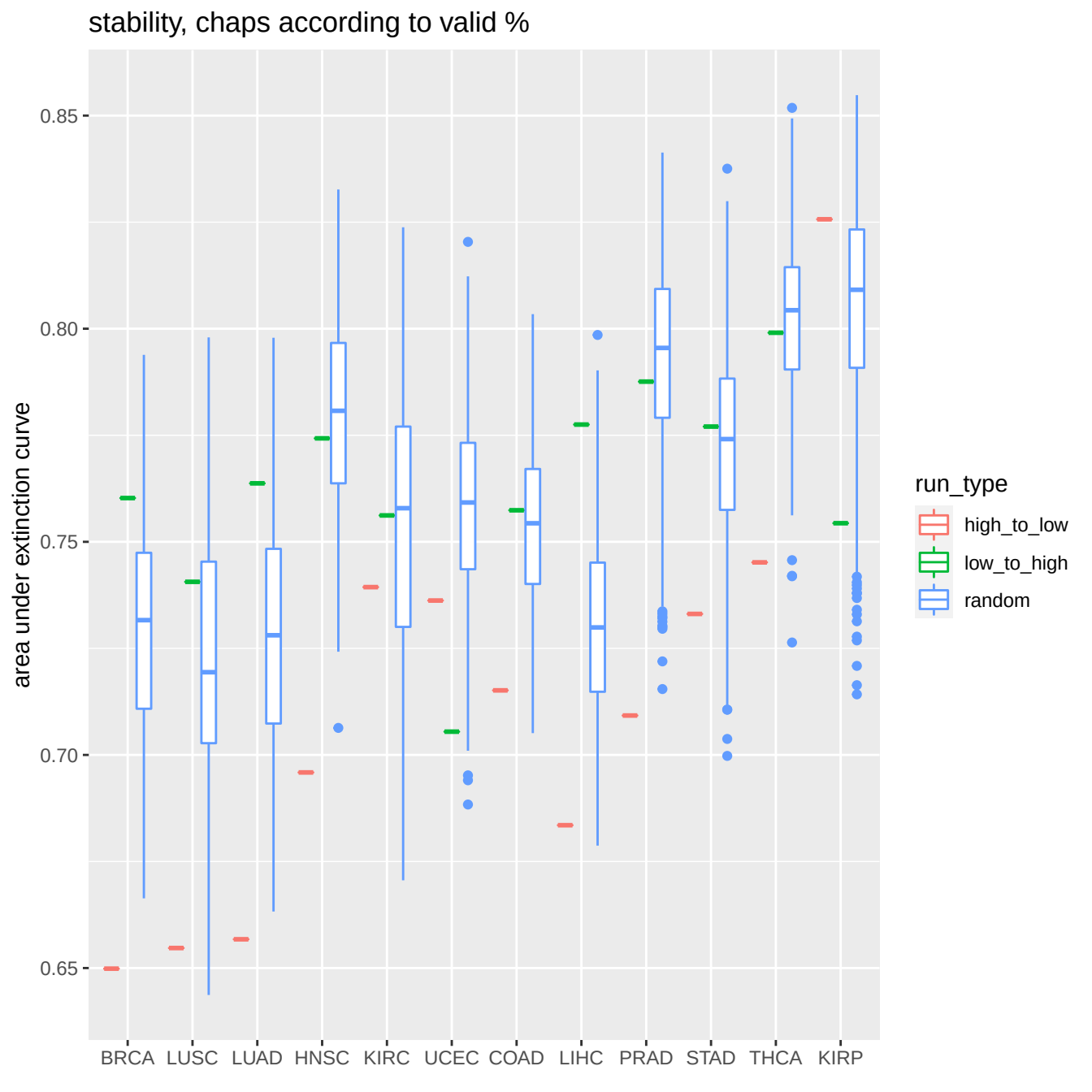
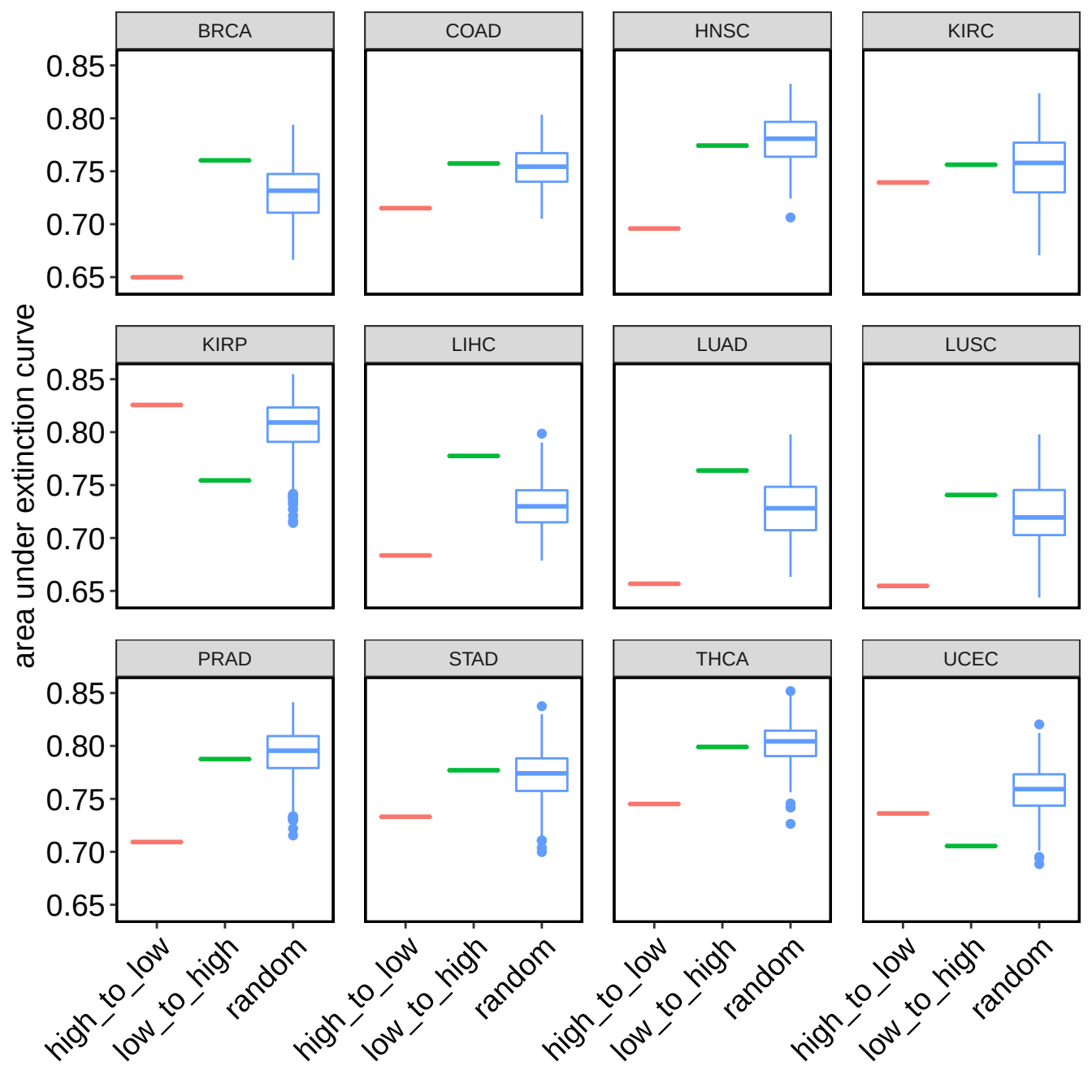
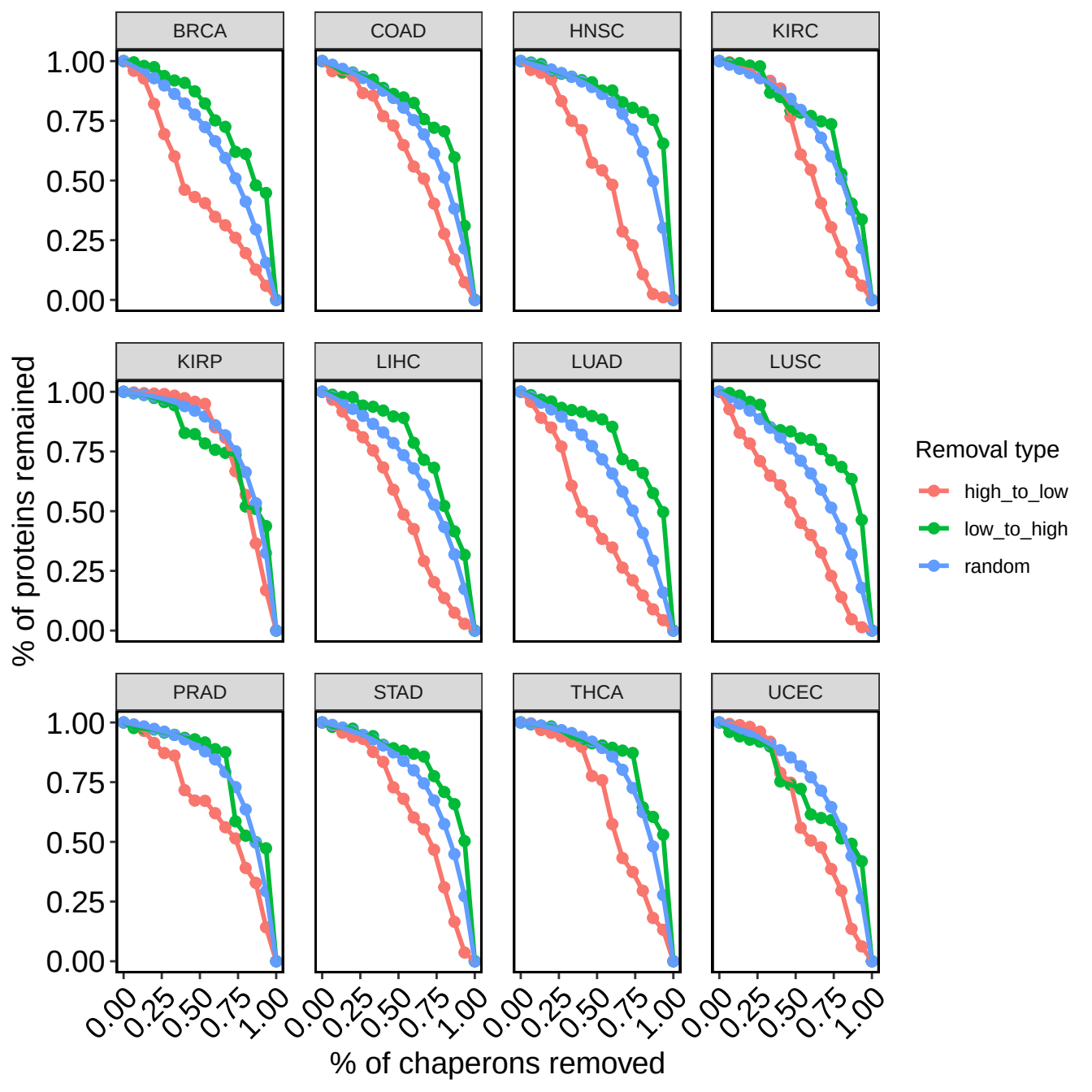

stability, chaps according to sample size
area under extinction curve
0.85
0.80
0.75
0.70
0.65
KIRP
LIHC
STAD
COAD
PRAD
HNSC
LUSC
THCA
LUAD
KIRC
UCEC
BRCA
run_type
high_to_low
low_to_high
random


stability, chaps according to valid %
area under extinction curve
0.85
0.80
0.75
0.70
0.65
BRCA
LUSC
LUAD
HNSC
KIRC
UCEC
COAD
LIHC
PRAD
STAD
THCA
KIRP
run_type
high_to_low
low_to_high
random


BRCA
COAD
HNSC
KIRC
KIRP
LIHC
LUAD
LUSC
PRAD
STAD
THCA
UCEC
area under extinction curve
0.85
0.80
0.75
0.70
0.65
high_to_low
low_to_high
random


BRCA
COAD
HNSC
KIRC
KIRP
LIHC
LUAD
LUSC
PRAD
STAD
THCA
UCEC
% of proteins remained
1.00
0.75
0.50
0.25
0.00
0.00
0.25
0.50
0.75
1.00
% of chaperons removed
Removal type
high_to_low
low_to_high
random