stability. chaps according to sample size 0.75 -0.70 under_curve 0.60 -0.55 **-**

PRAD

COAD

HNSC

fct_relevel(cancer, cancer_sample_size_order)

LUSC

THCA

LUAD

KIRC

UCEC

BRCA

KIRP

LIHC

STAD

stability. chaps according to valid % 0.75 -0.70 under_curve 0.60 -0.55 -BRCA LUSC KIRC UCEC LUAD HNSC COAD LIHC PRAD STAD THCA KIRP fct_relevel(cancer, cancer_percent_of_valid_interactions)

stability. by removal type 0.75 -0.70 nnder_curve 0.60 -0.55 high_to_low low_to_high random

run_type