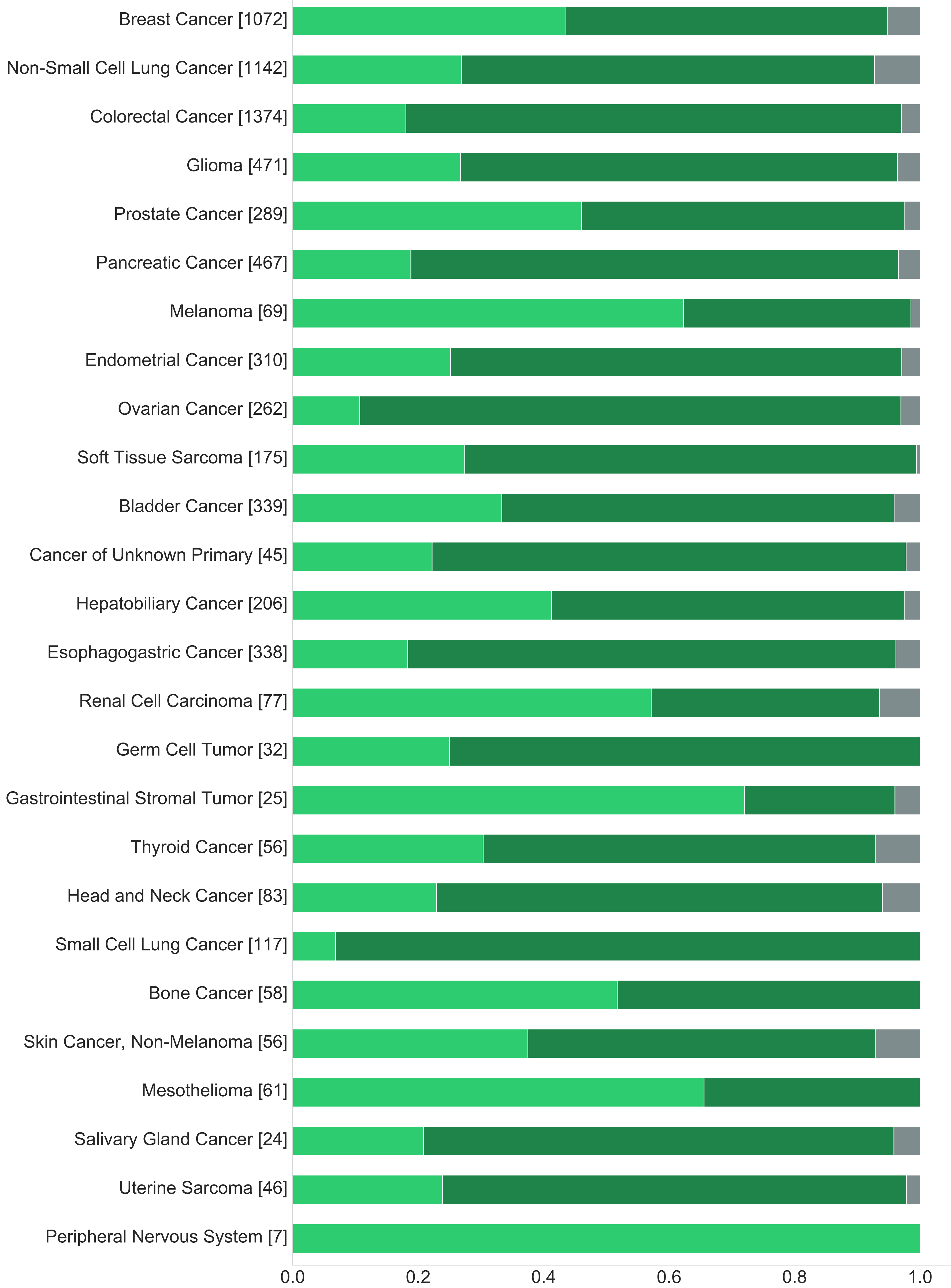
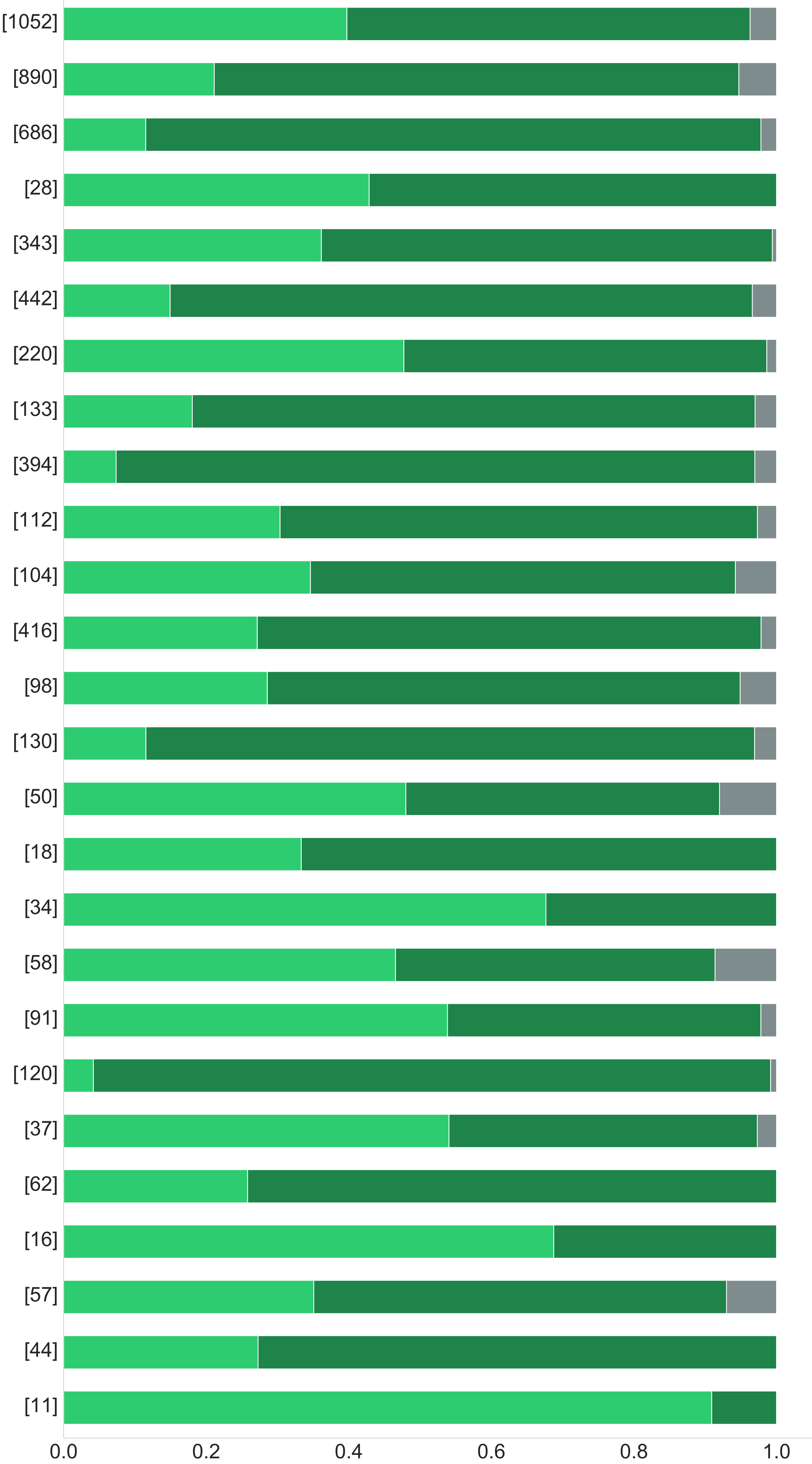


TP53 RESIDUAL SUBGROUP PROPORTION ACROSS CANCERS

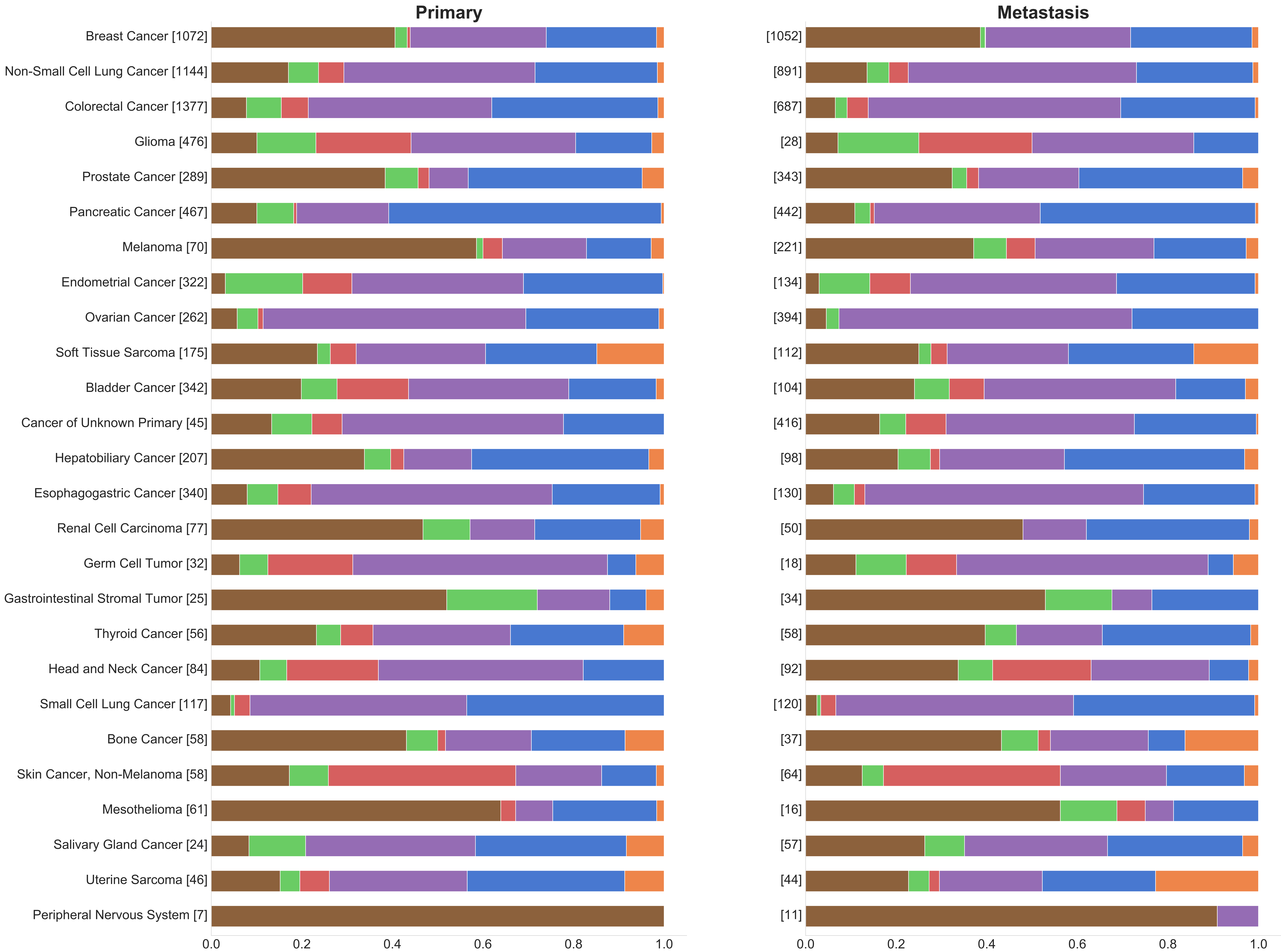
Primary



Metastasis



SUBGROUP PROPORTION ACROSS CANCERS



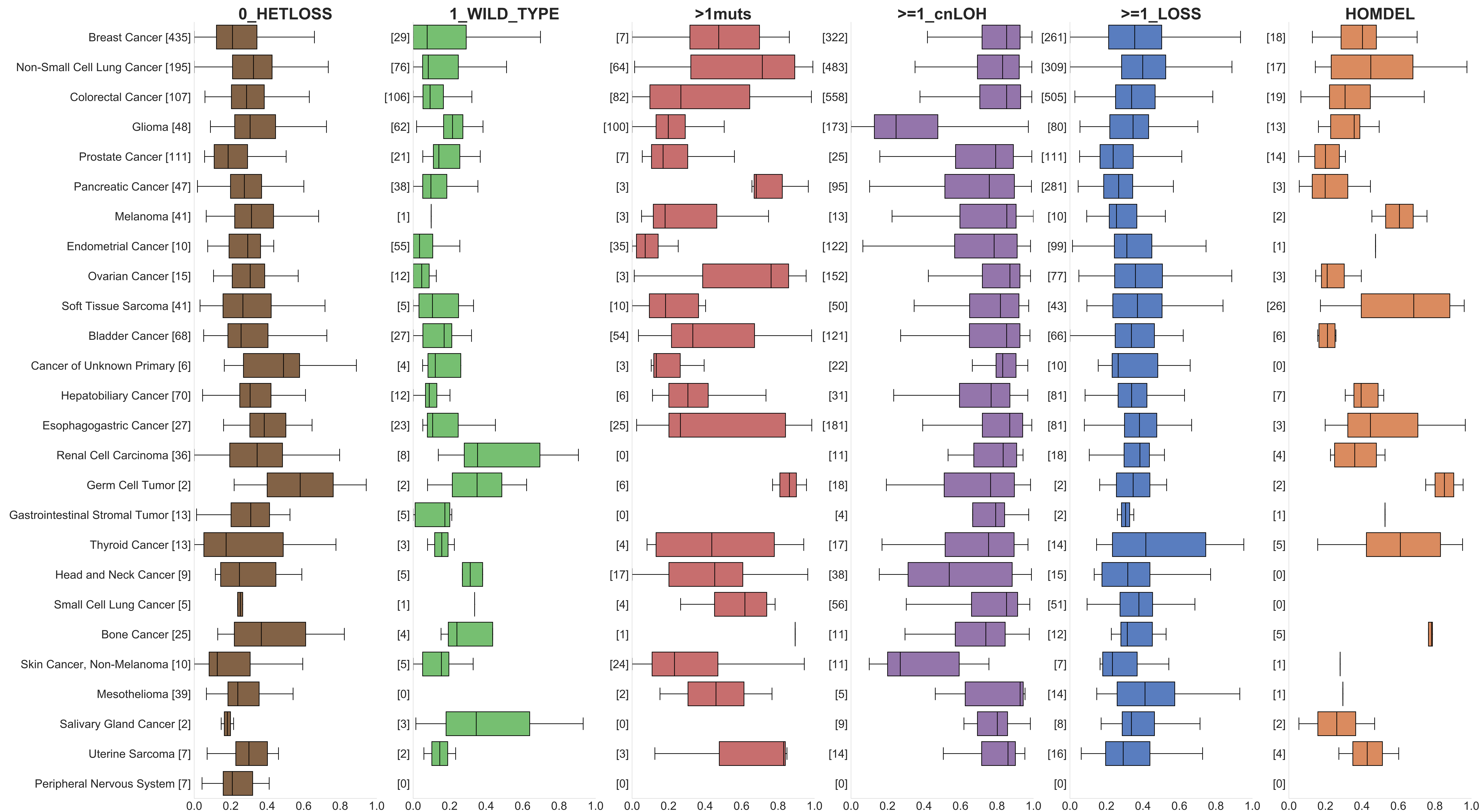
Fraction of genome altered

Primary

Metastasis

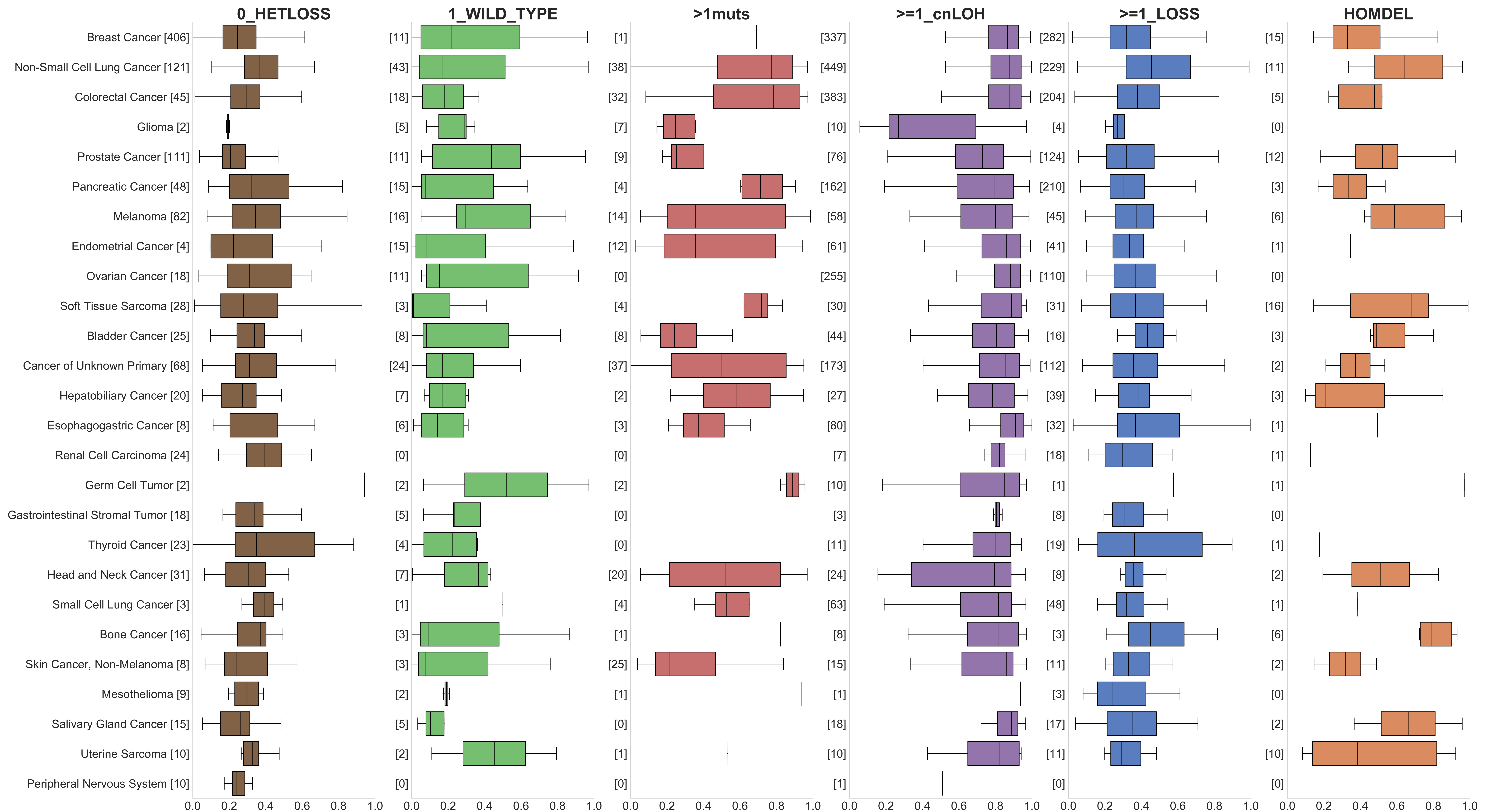


Fraction of genome altered - Primary Samples





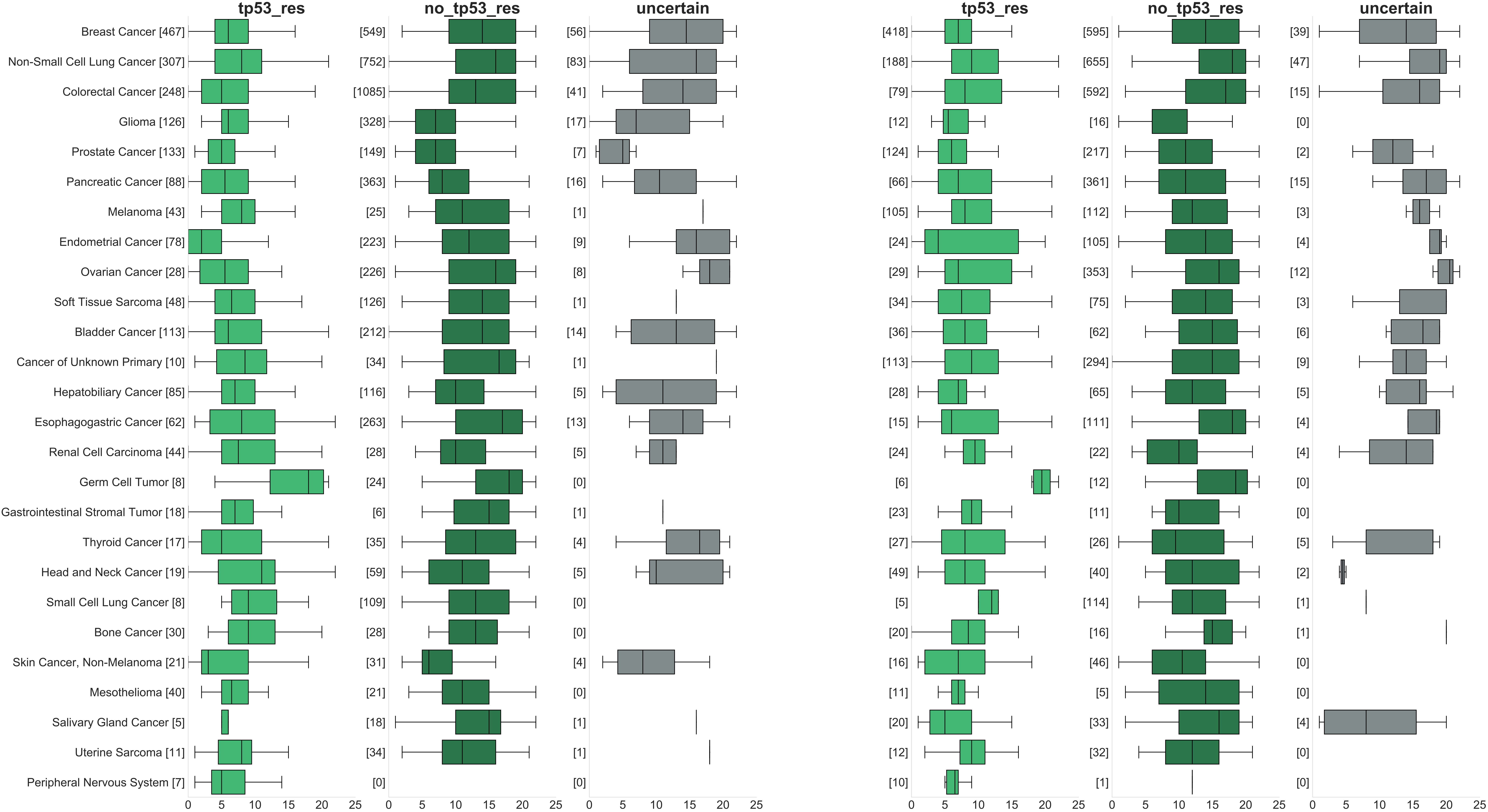
# Fraction of genome altered - Metastatic Samples



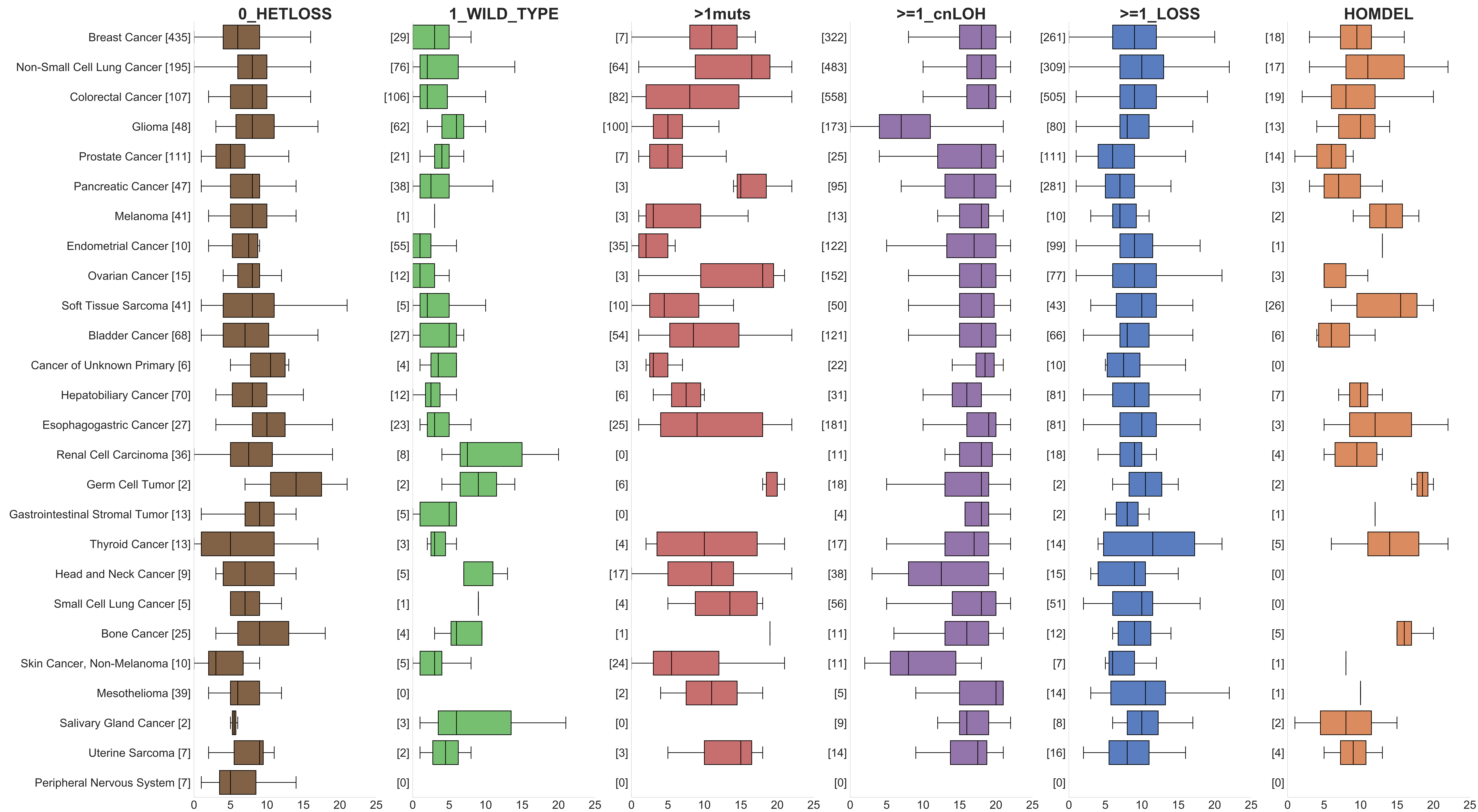
Number of aberrant chromosomes

Primary

Metastasis

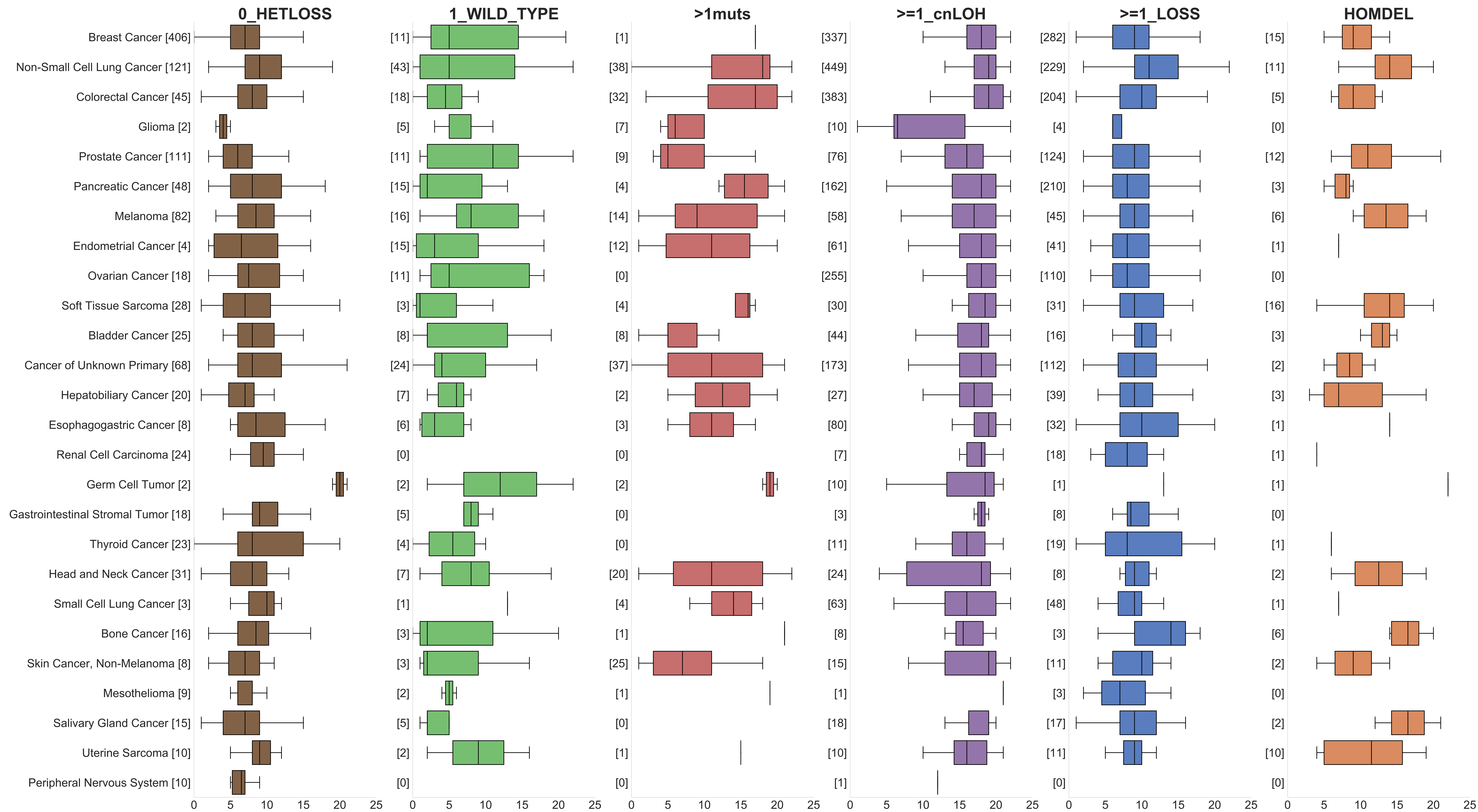


Number of aberrant chromosomes - Primary Samples





Number of aberrant chromosomes - Metastatic Samples





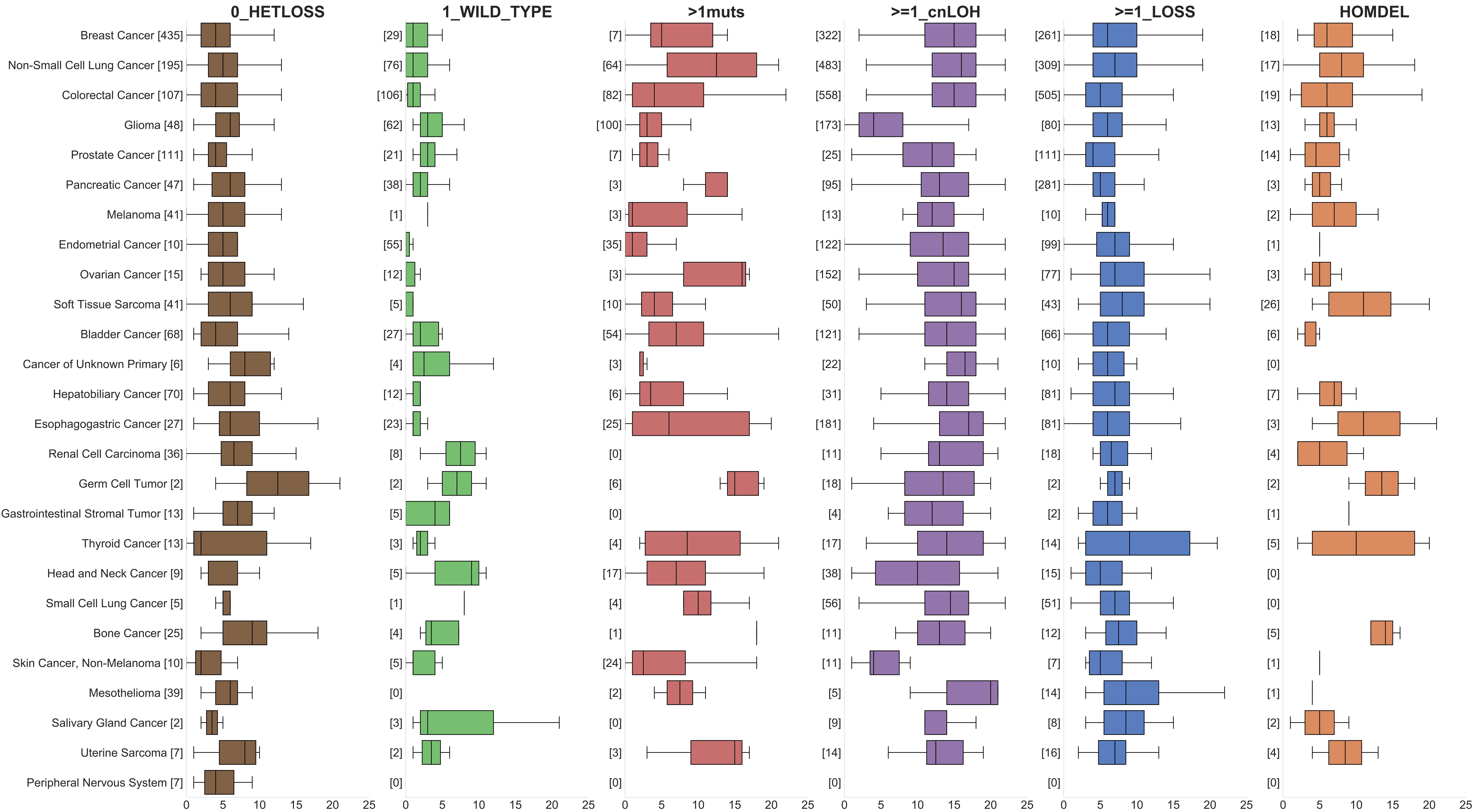
Number of chromosomes affected by a LOSS

Primary

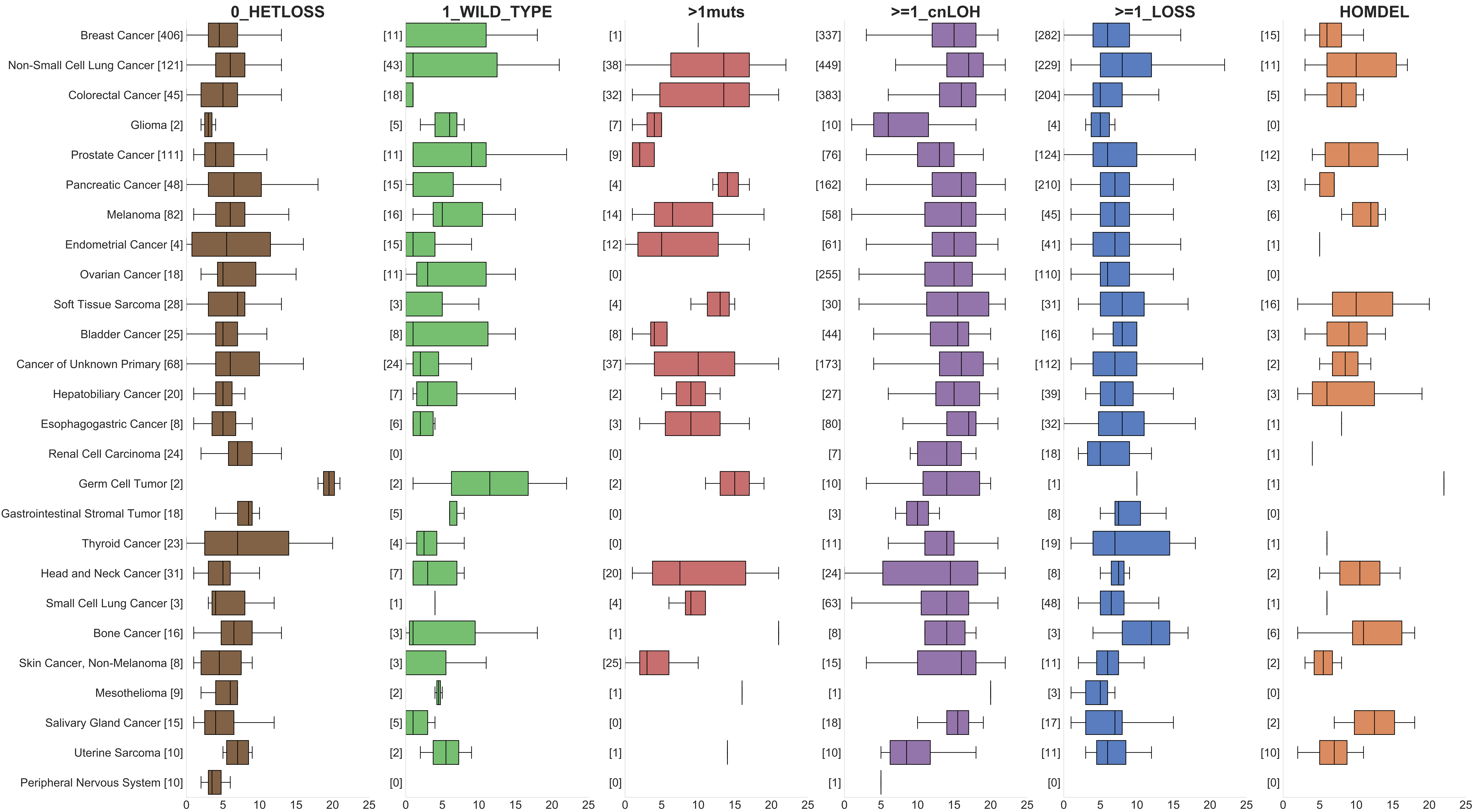
Metastasis



Number of chromosomes affected by a LOSS - Primary Samples

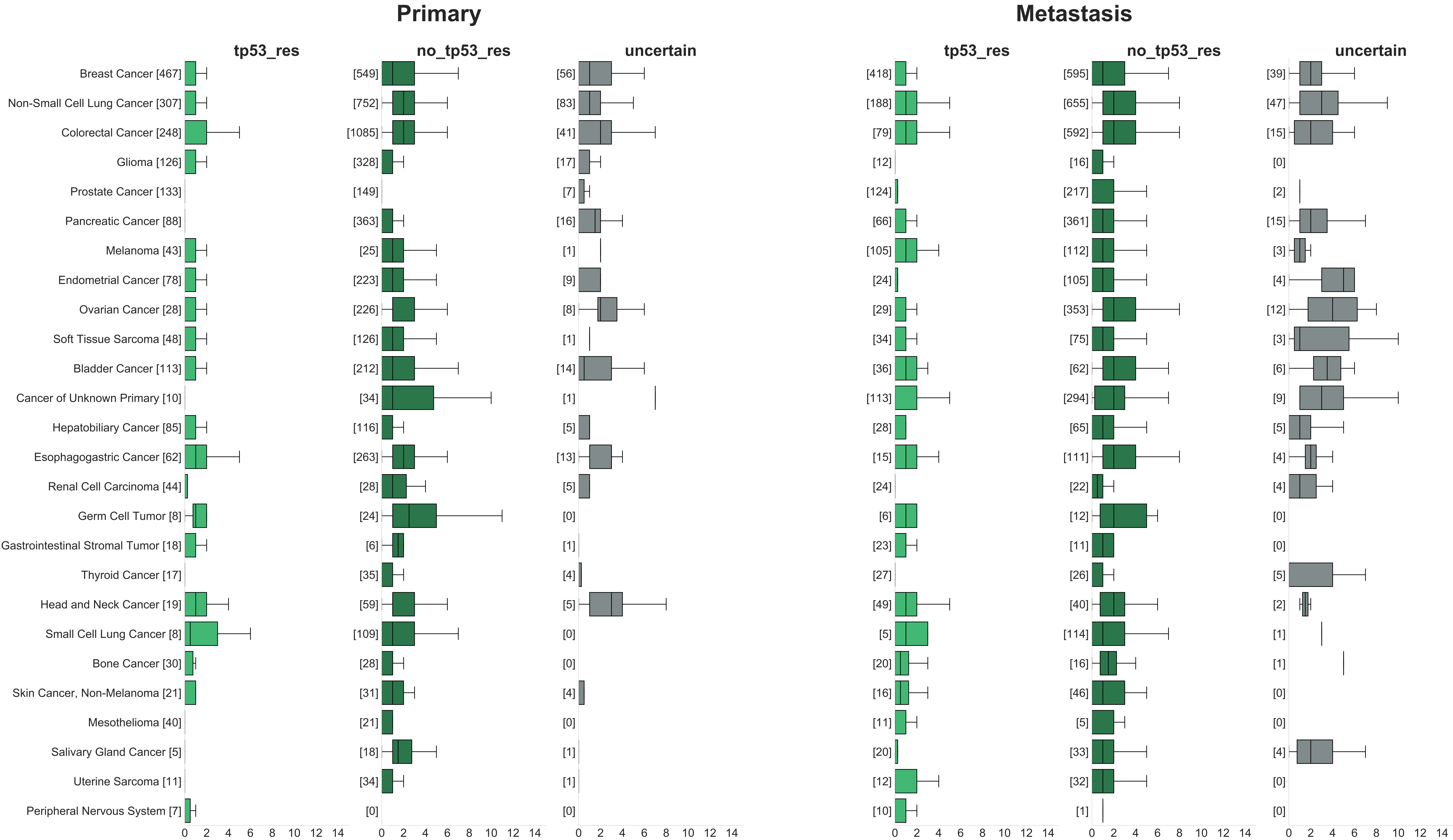


Number of chromosomes affected by a LOSS - Metastatic Samples

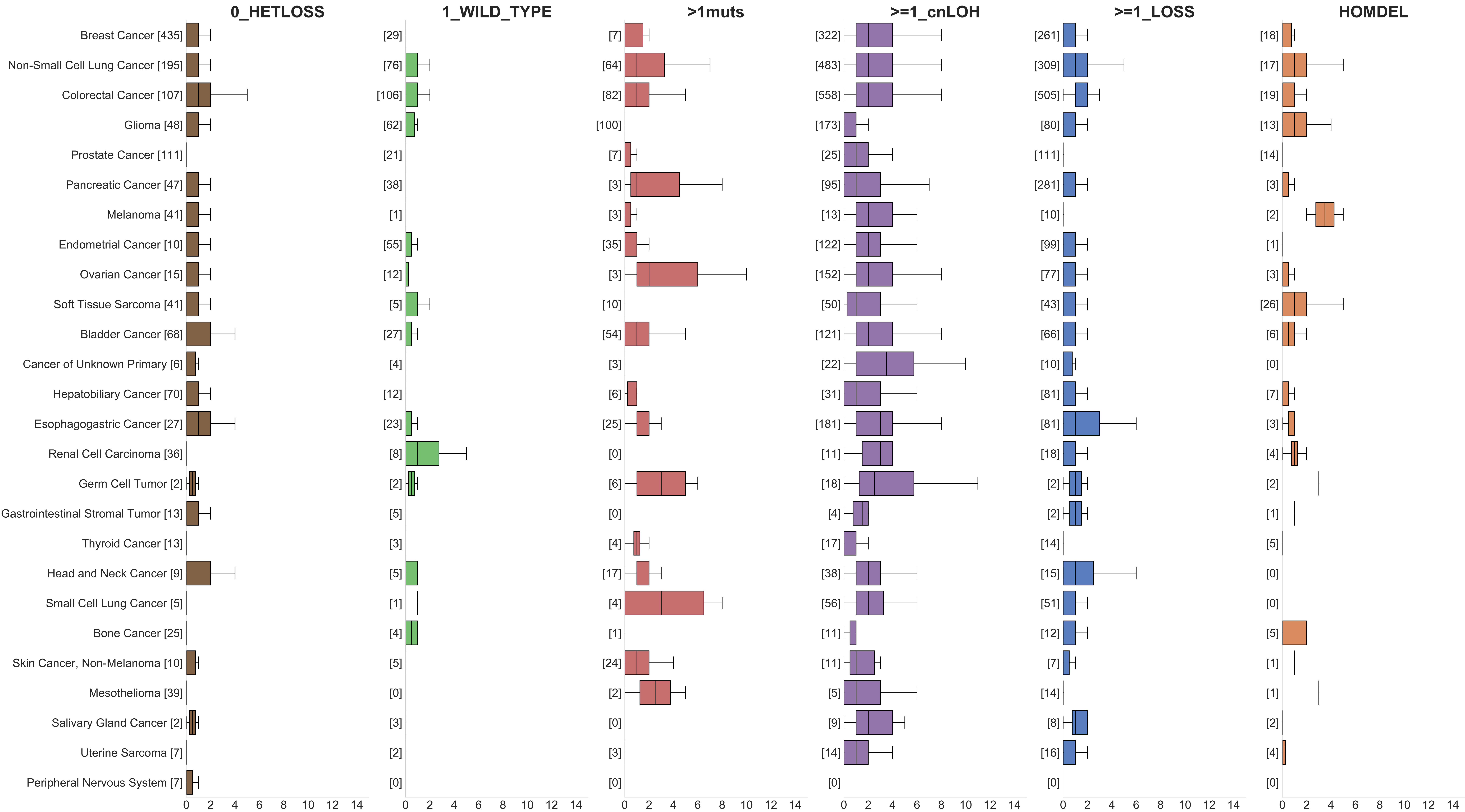




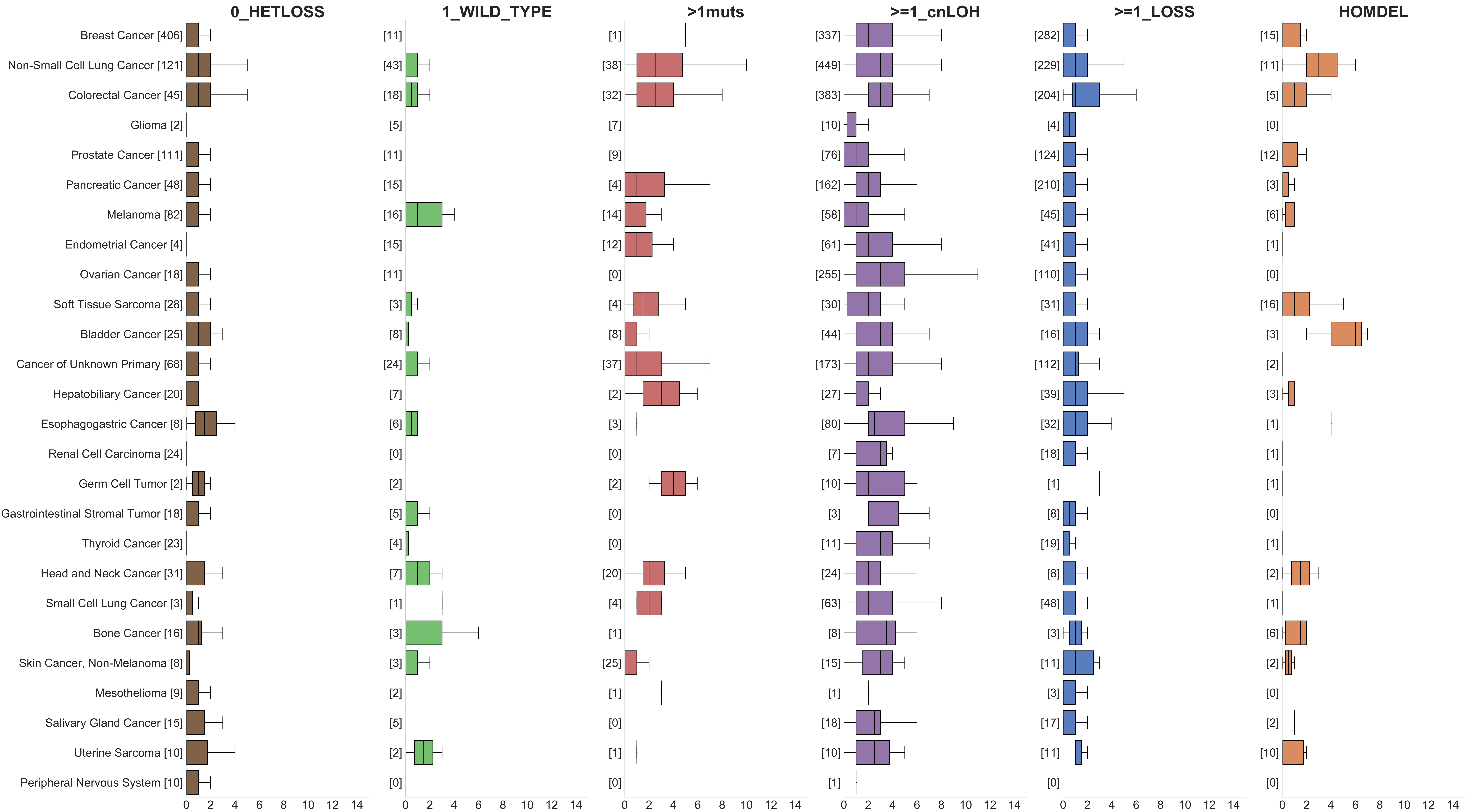
Number of chromosomes affected by a cnLOH



Number of chromosomes affected by a cnLOH - Primary Samples



Number of chromosomes affected by a cnLOH - Metastatic Samples

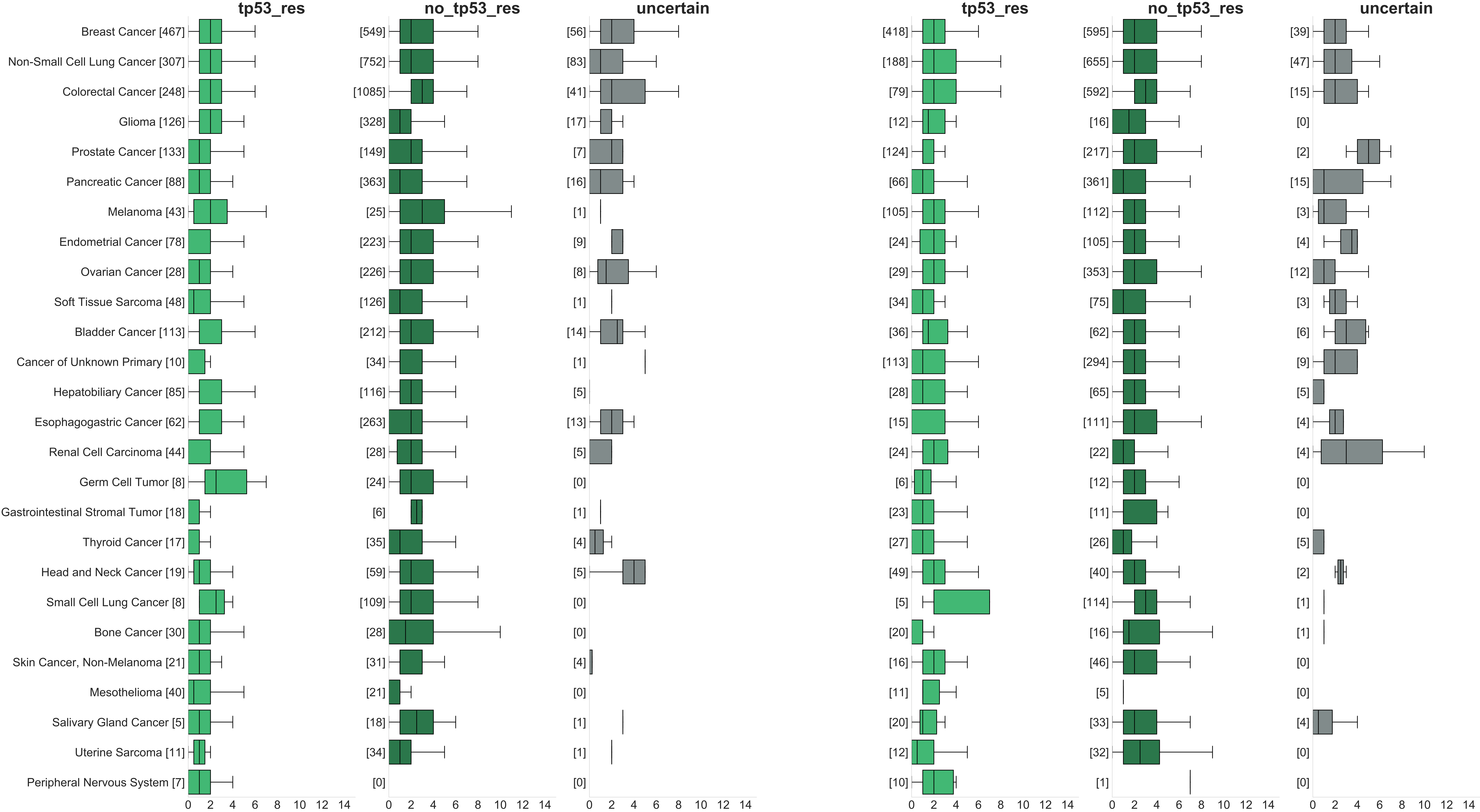




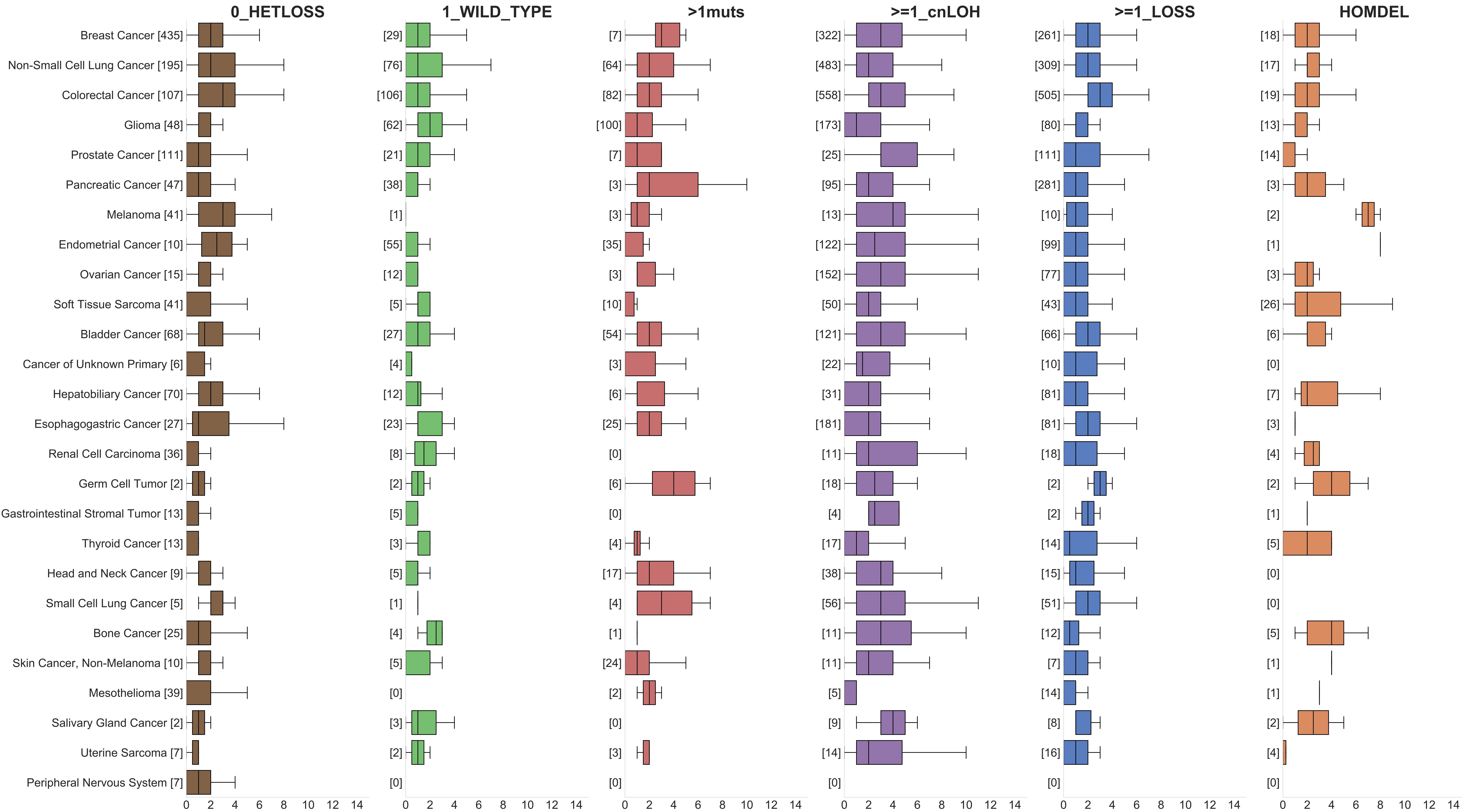
Number of chromosomes affected by a GAIN

Primary

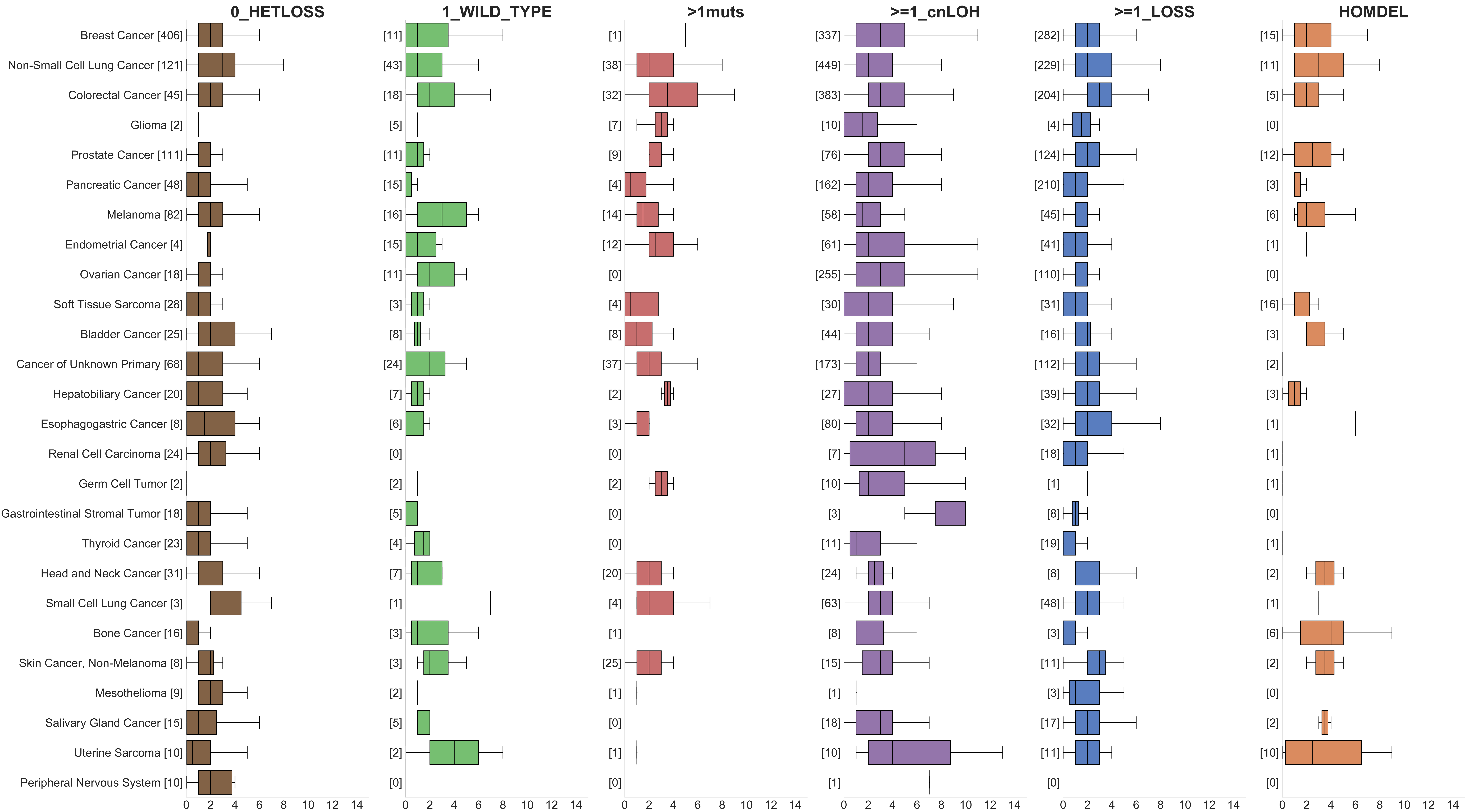
Metastasis



Number of chromosomes affected by a GAIN - Primary Samples

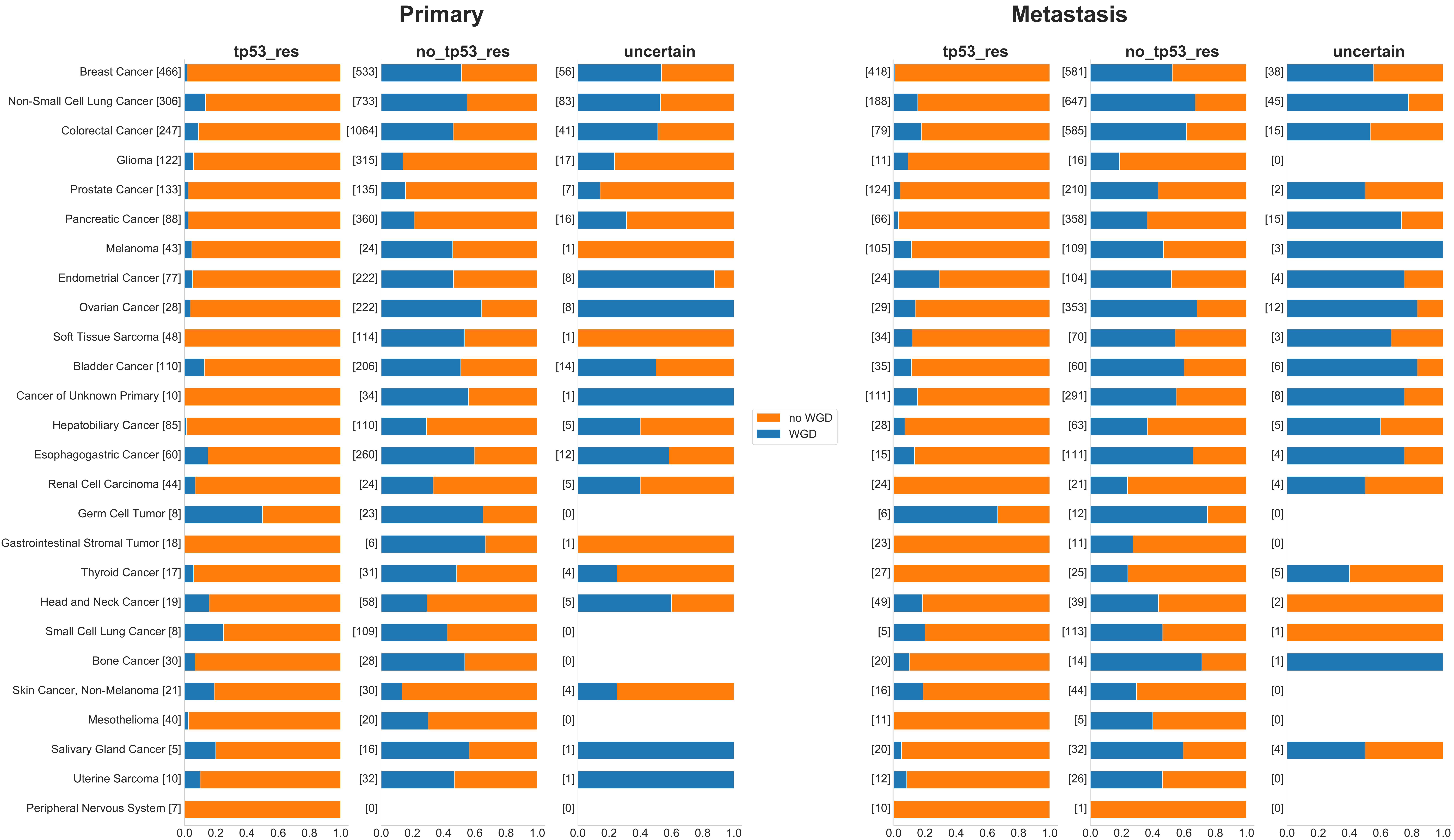


Number of chromosomes affected by a GAIN - Metastatic Samples





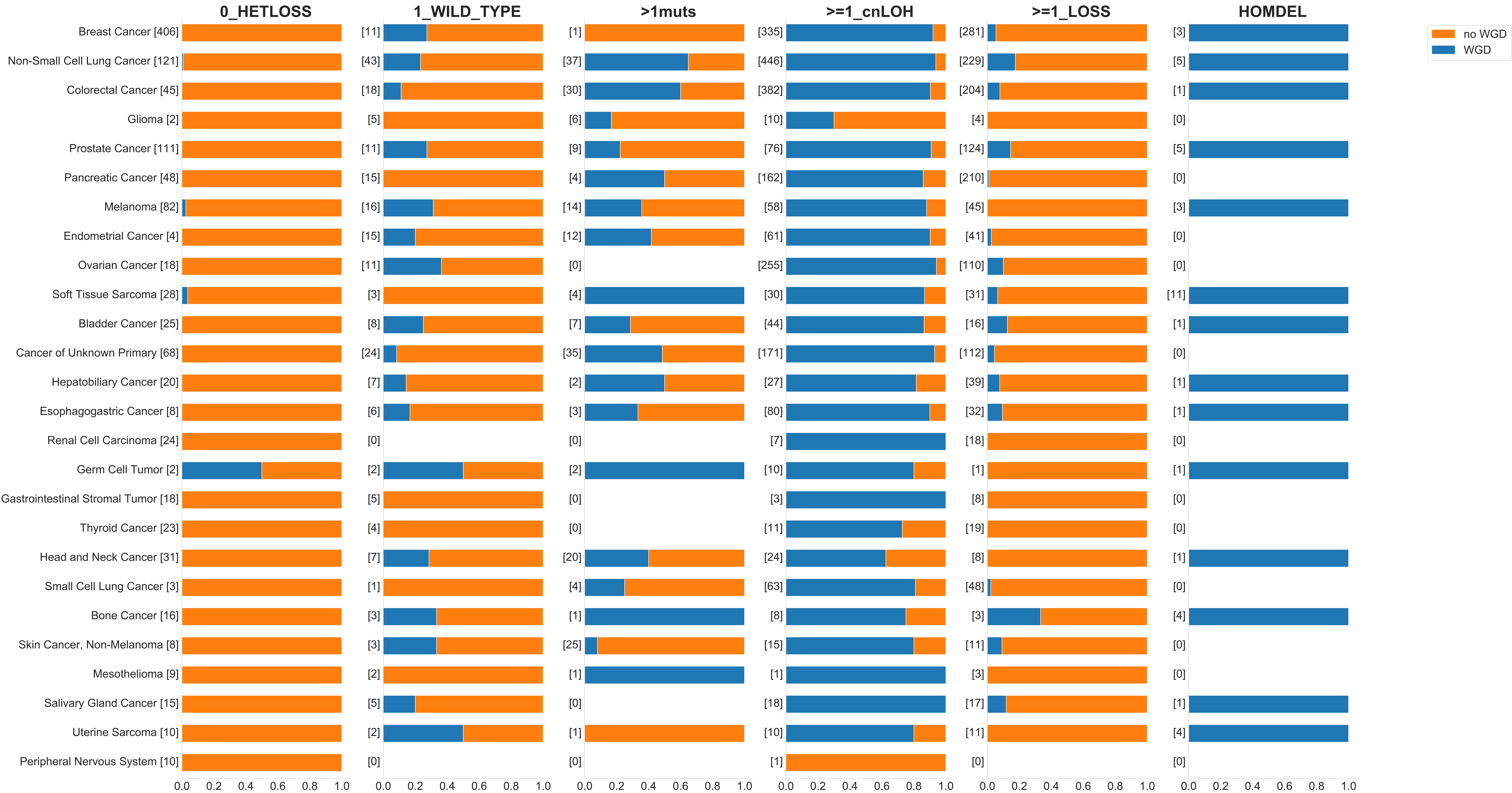
Proportion of WGD



Proportion of WGD - Primary Samples

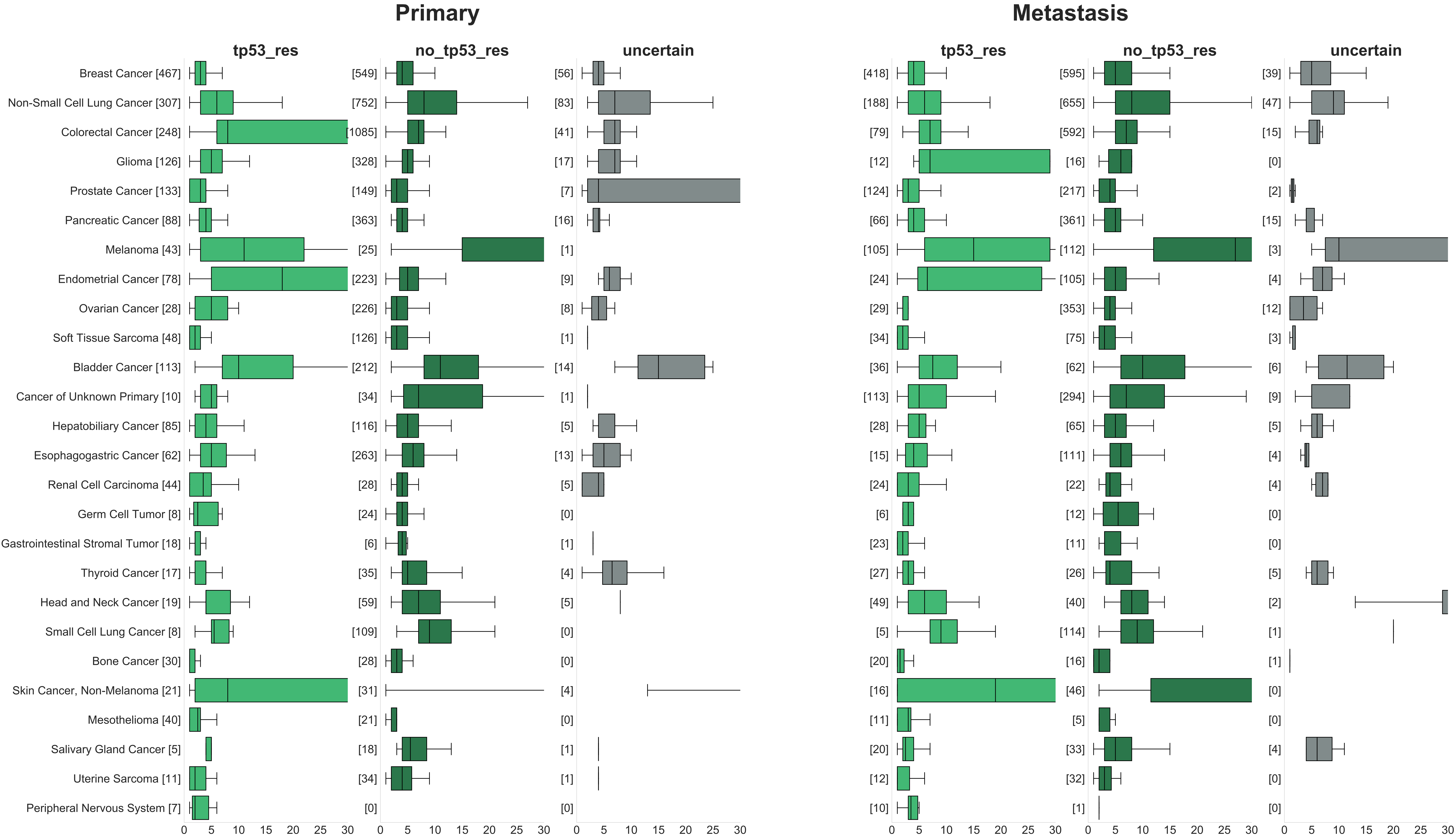


Proportion of WGD - Metastatic Samples

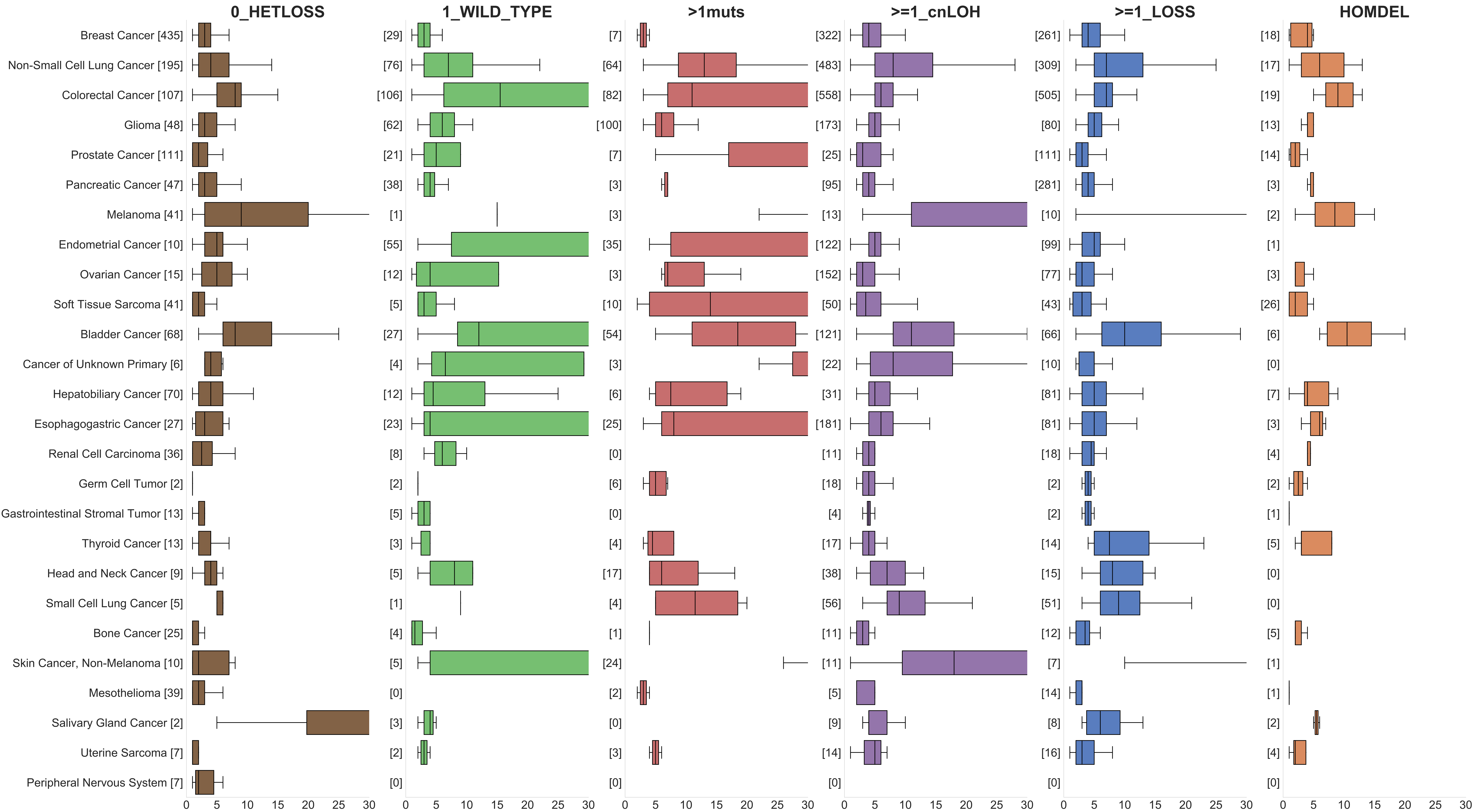




Mutation Count



Mutation Count - Primary Samples



Mutation Count - Metastatic Samples

