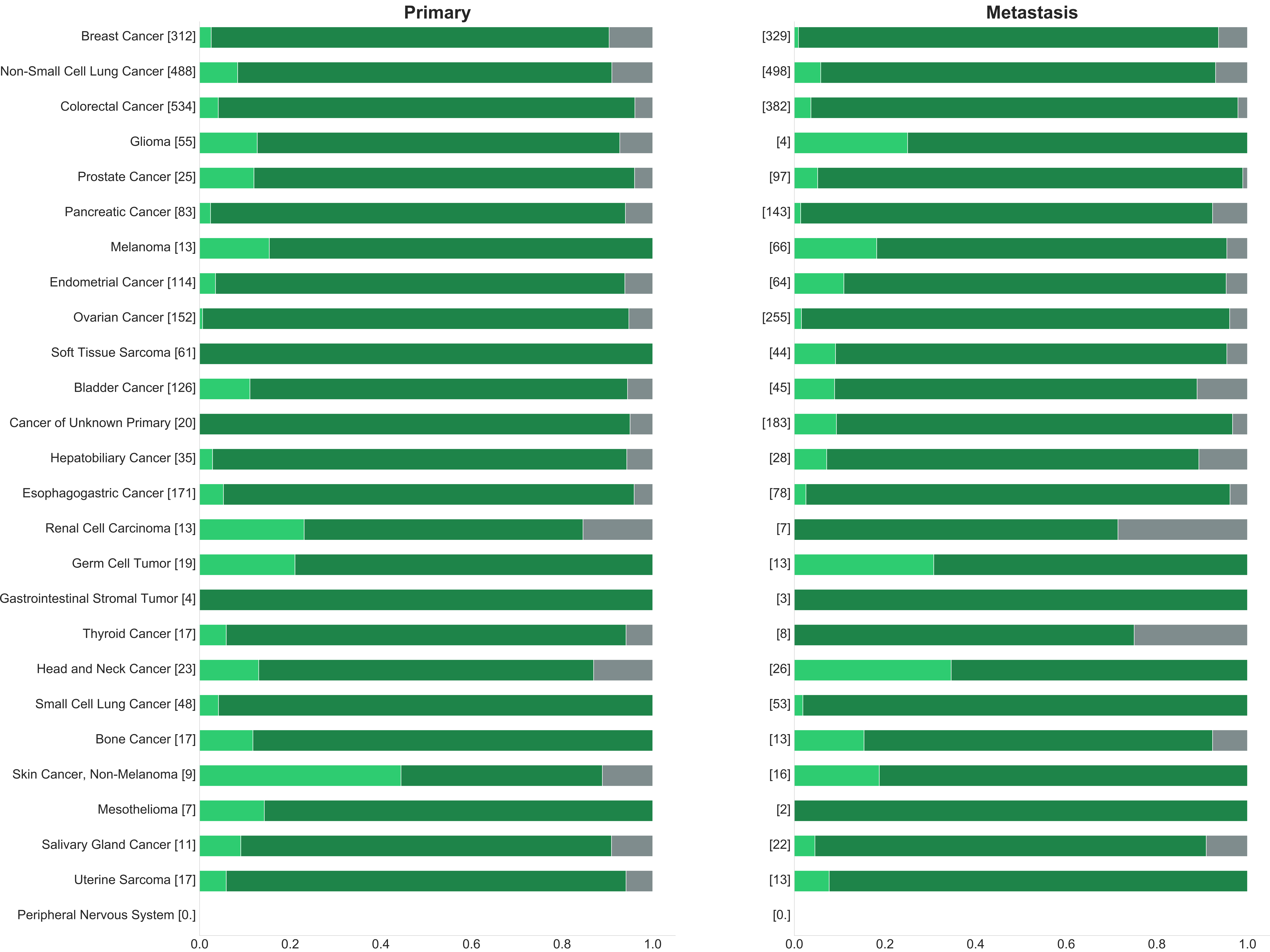
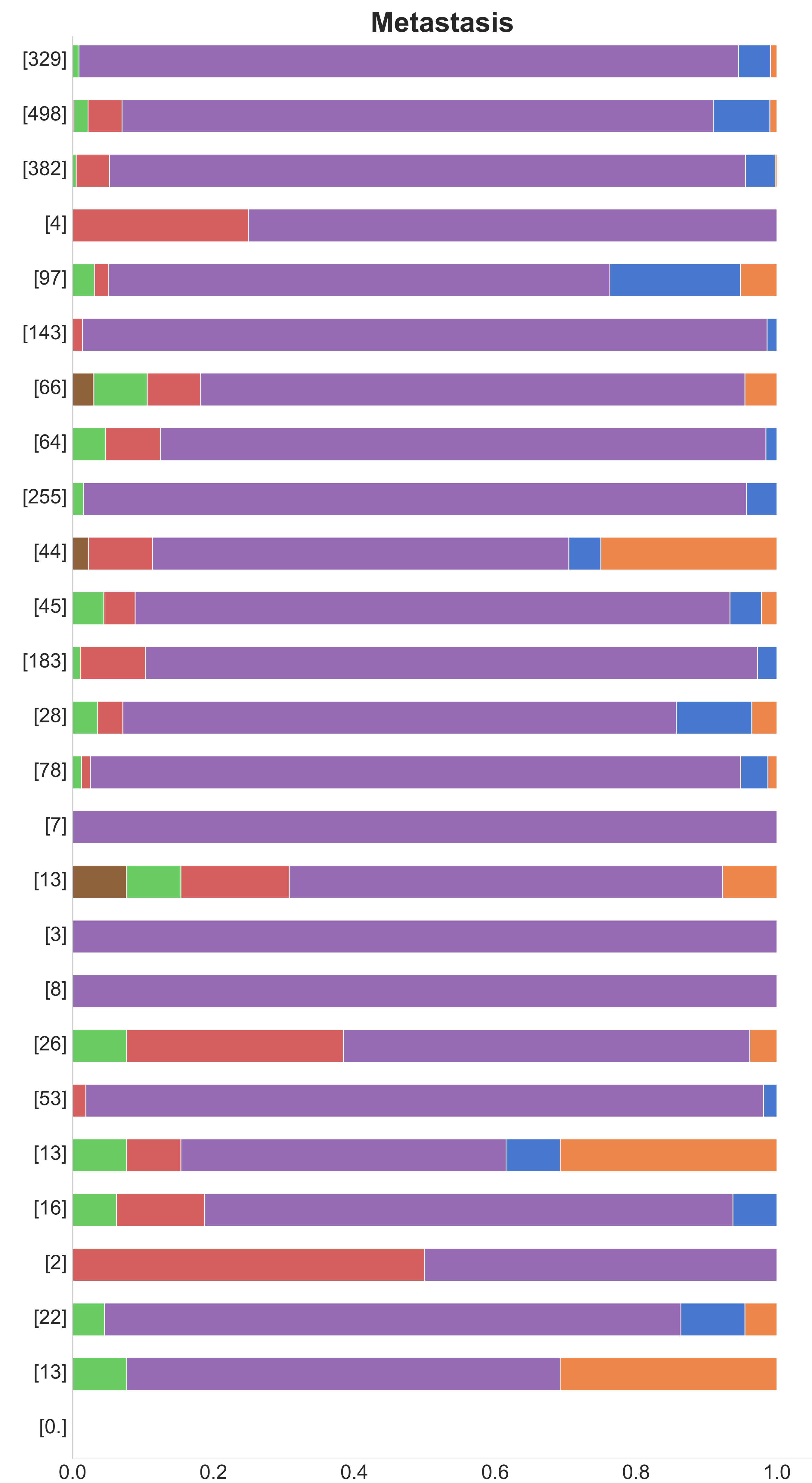
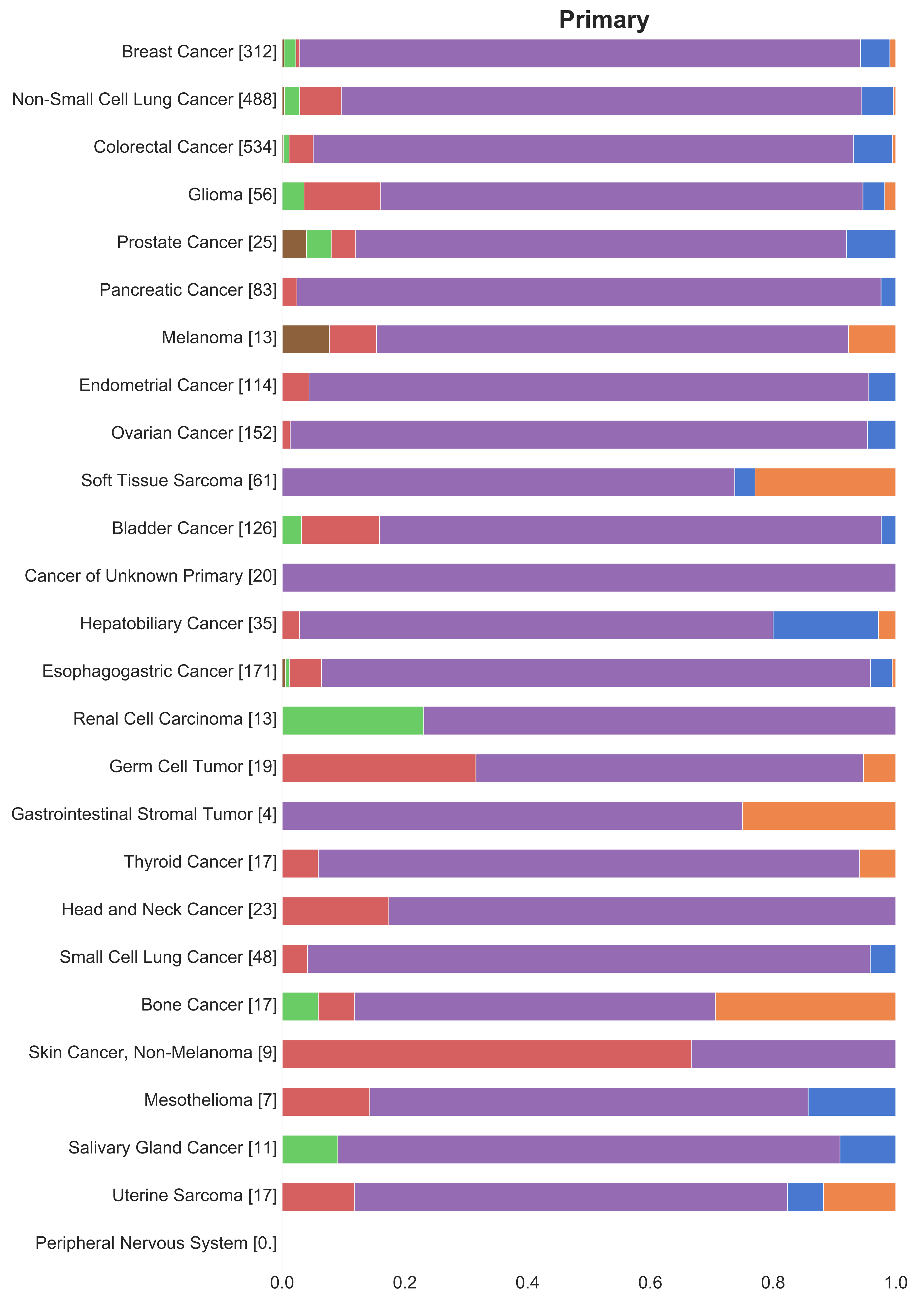


TP53 RESIDUAL SUBGROUP PROPORTION ACROSS CANCERS



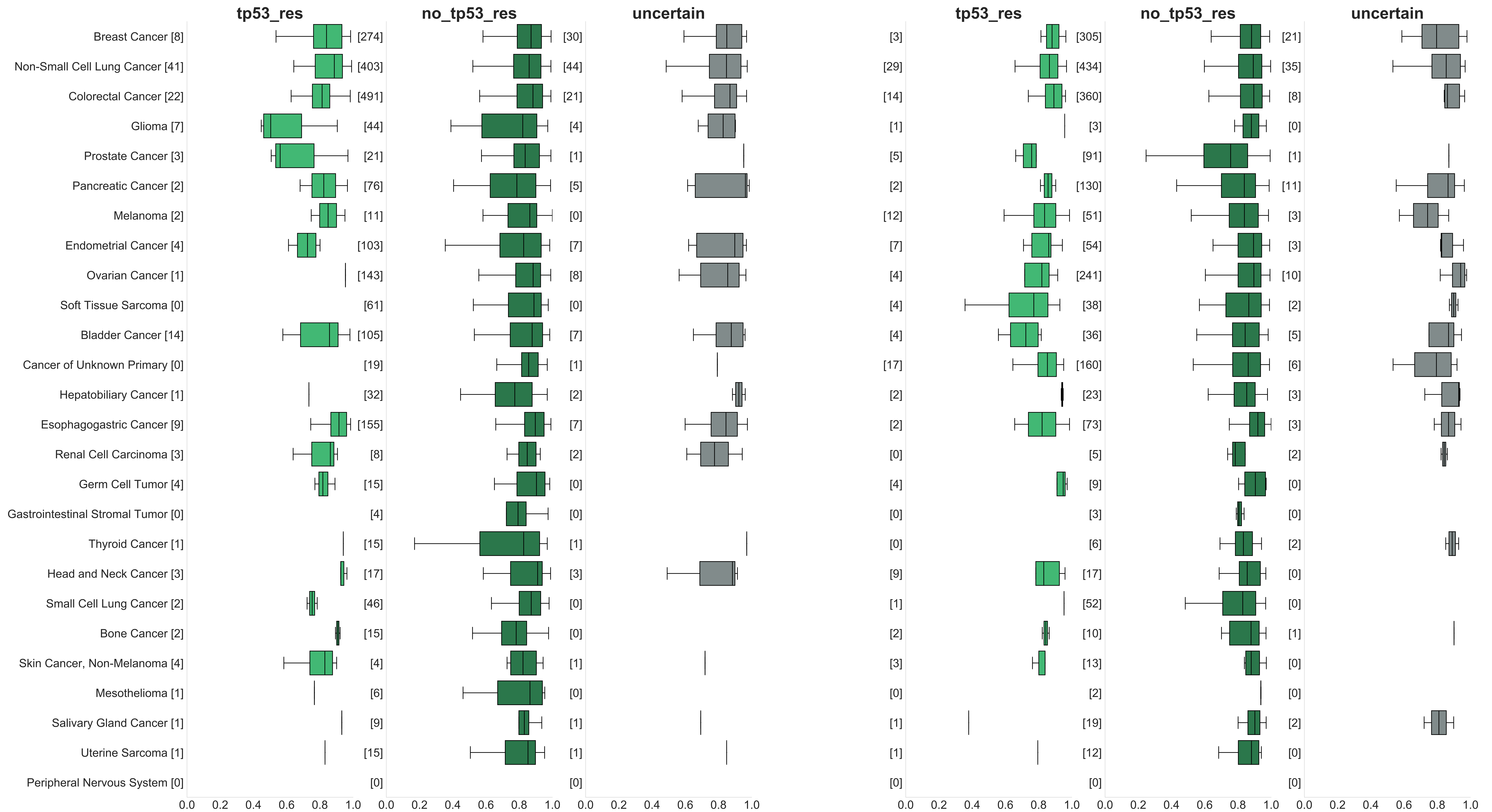
# TP53 SUBGROUP PROPORTION ACROSS CANCERS



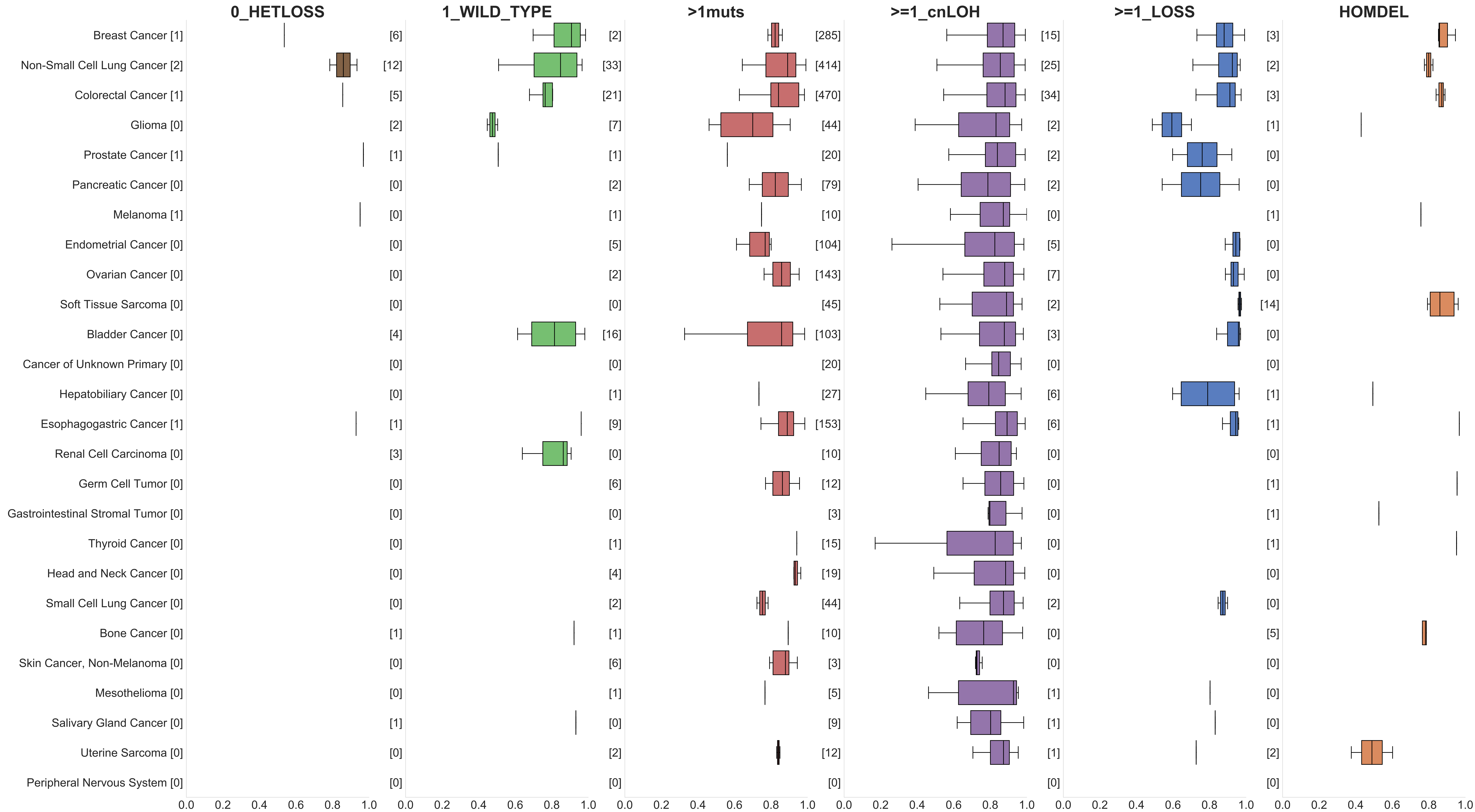
## Fraction of genome altered

## Primary

## Metastasis

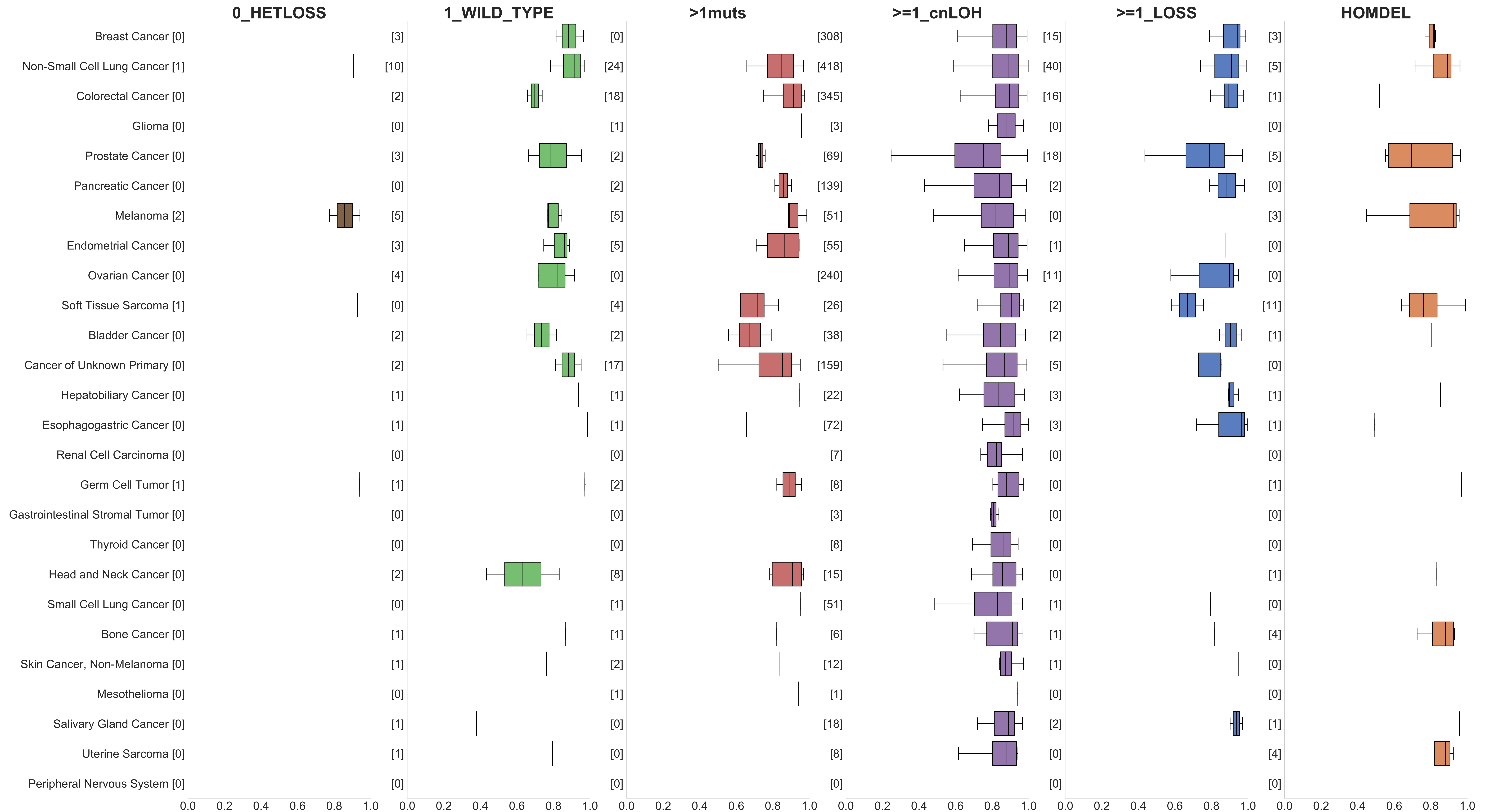


Fraction of genome altered - Primary Samples





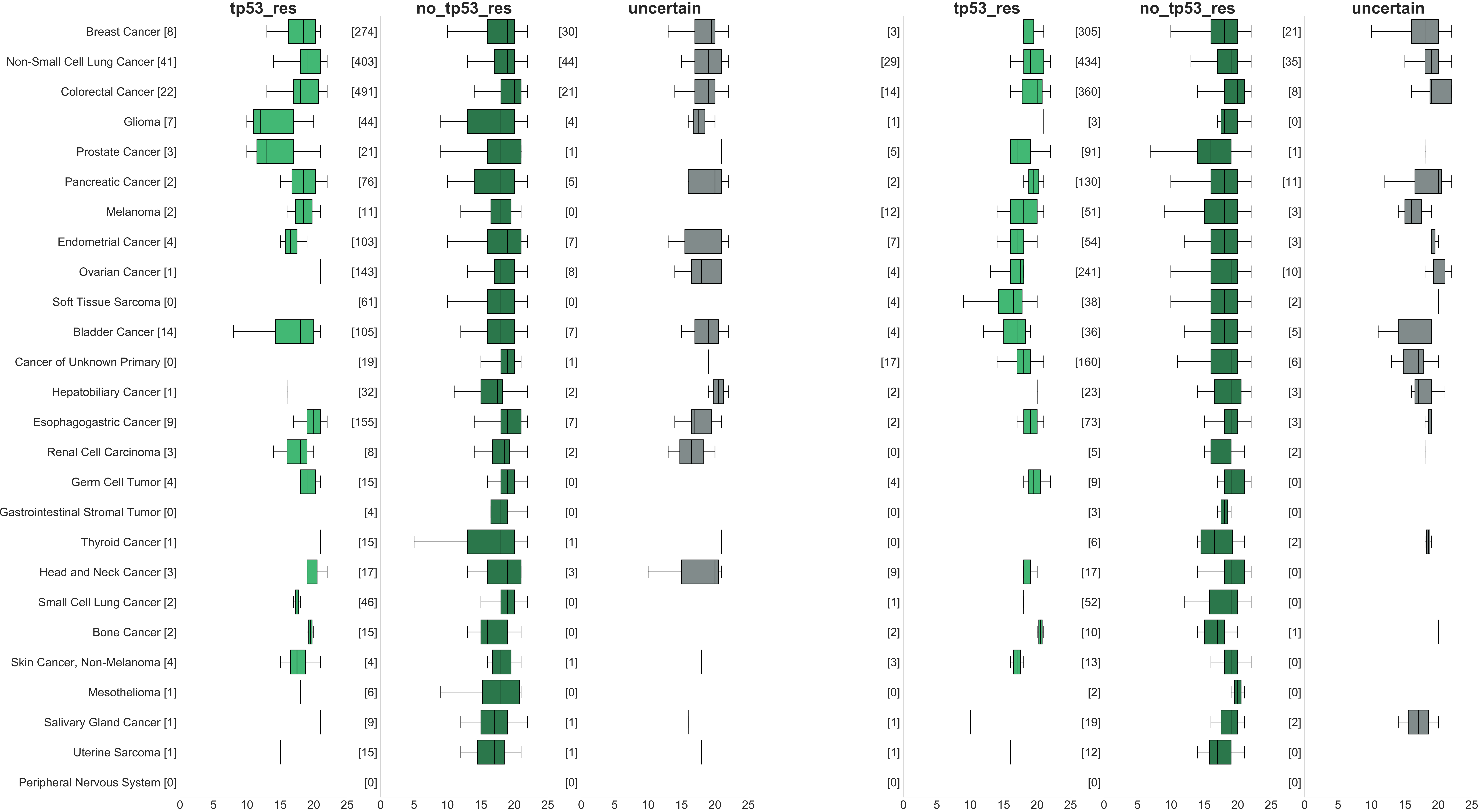
### Fraction of genome altered - Metastatic Samples



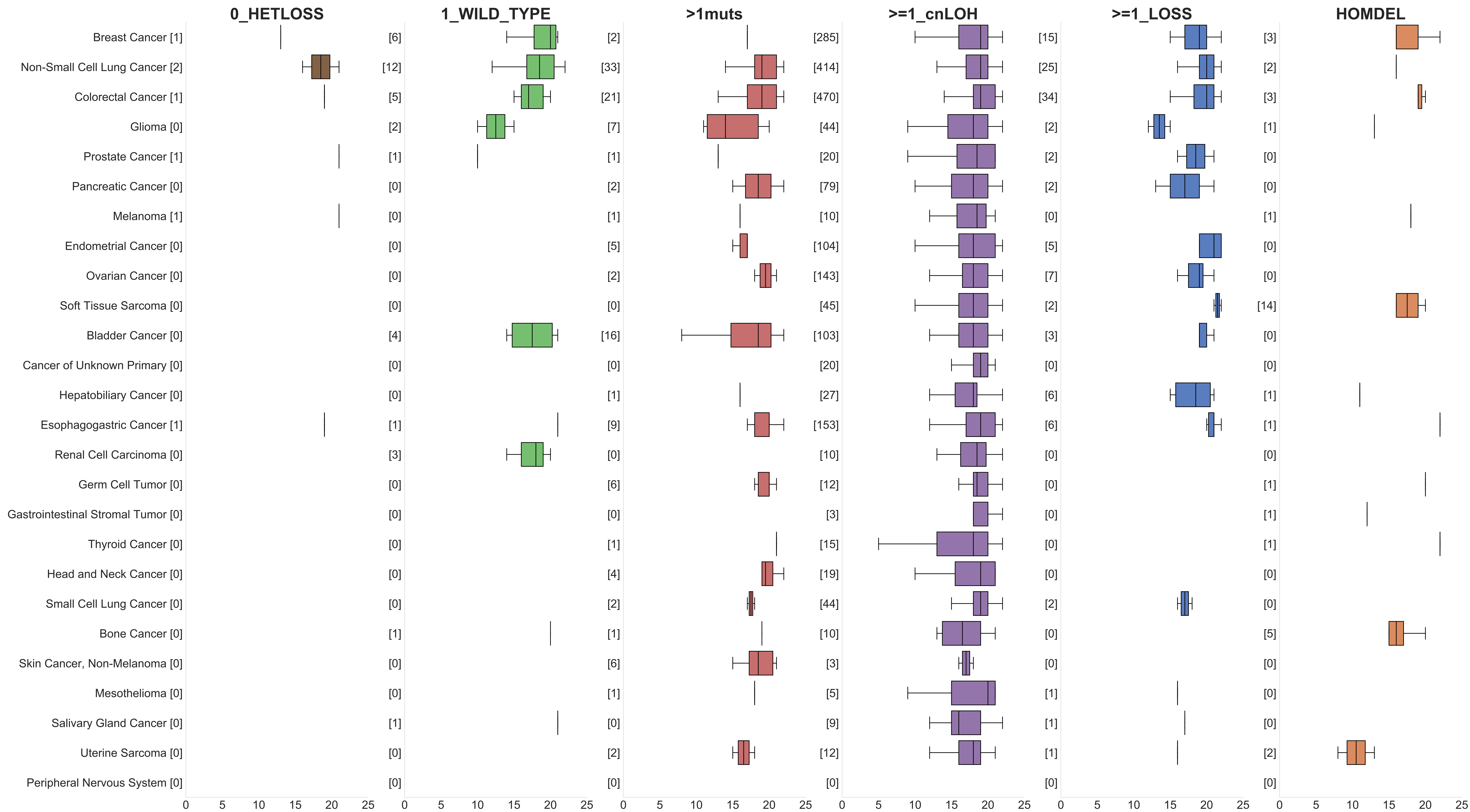
Number of aberrant chromosomes

Primary

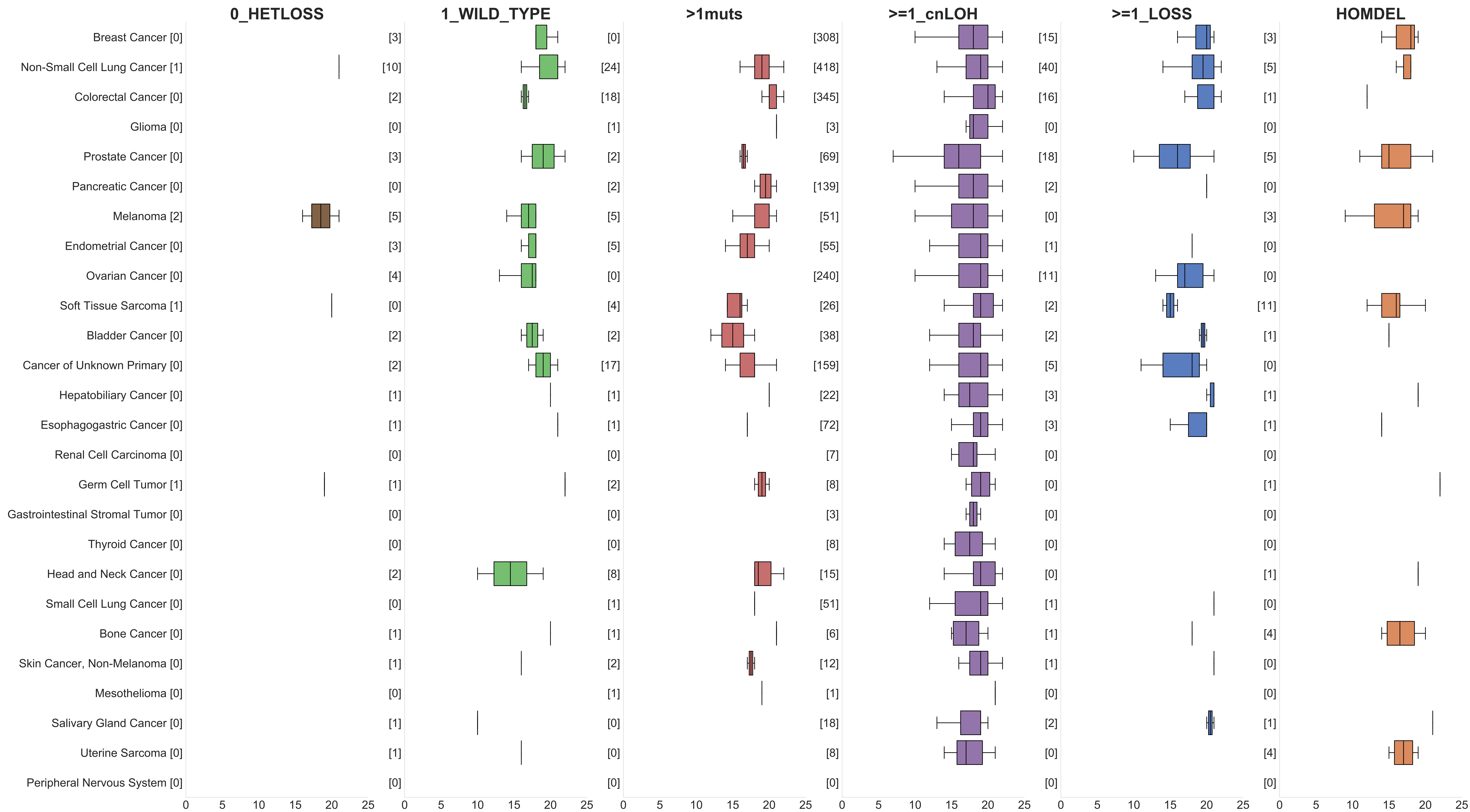
Metastasis



Number of aberrant chromosomes - Primary Samples



Number of aberrant chromosomes - Metastatic Samples

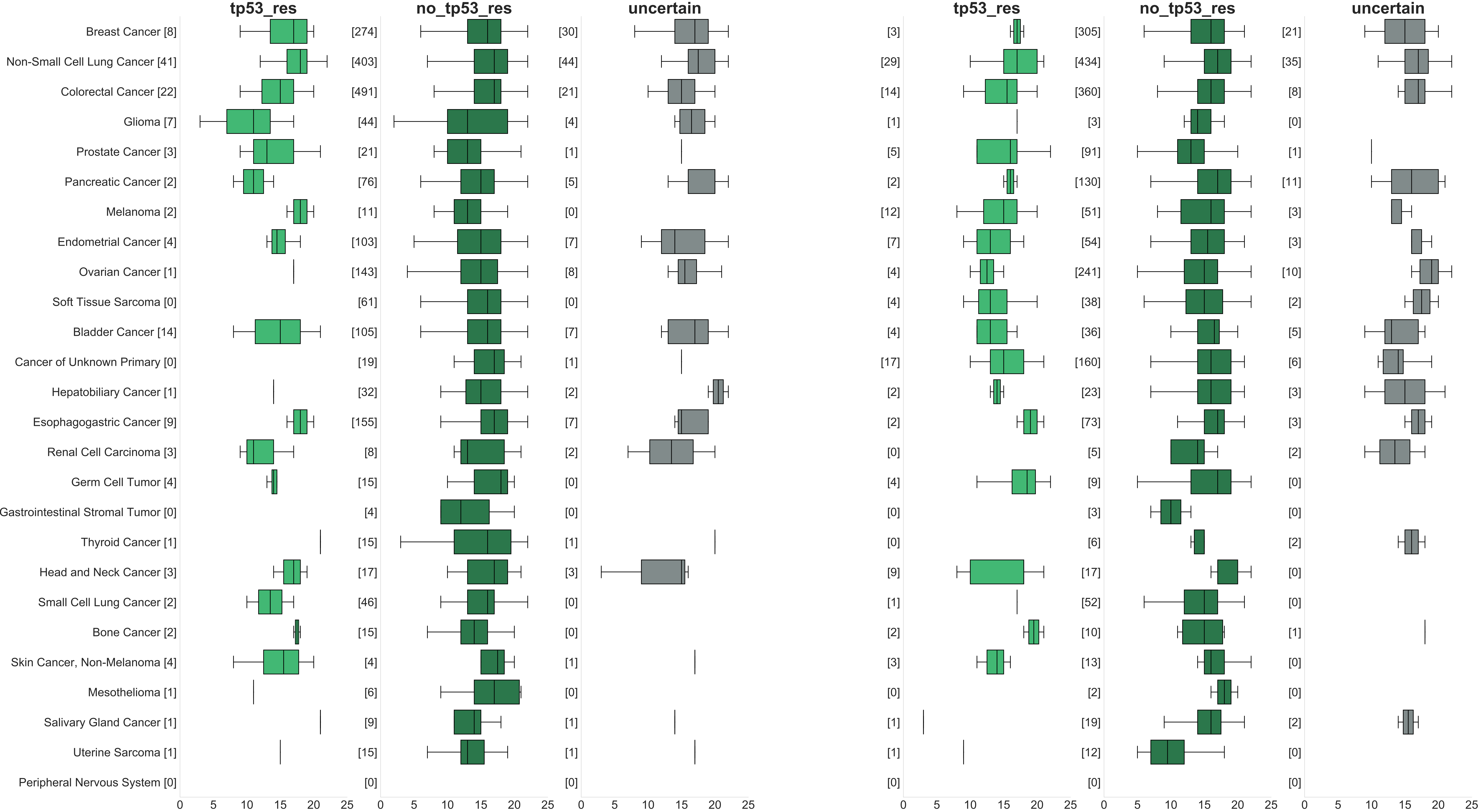




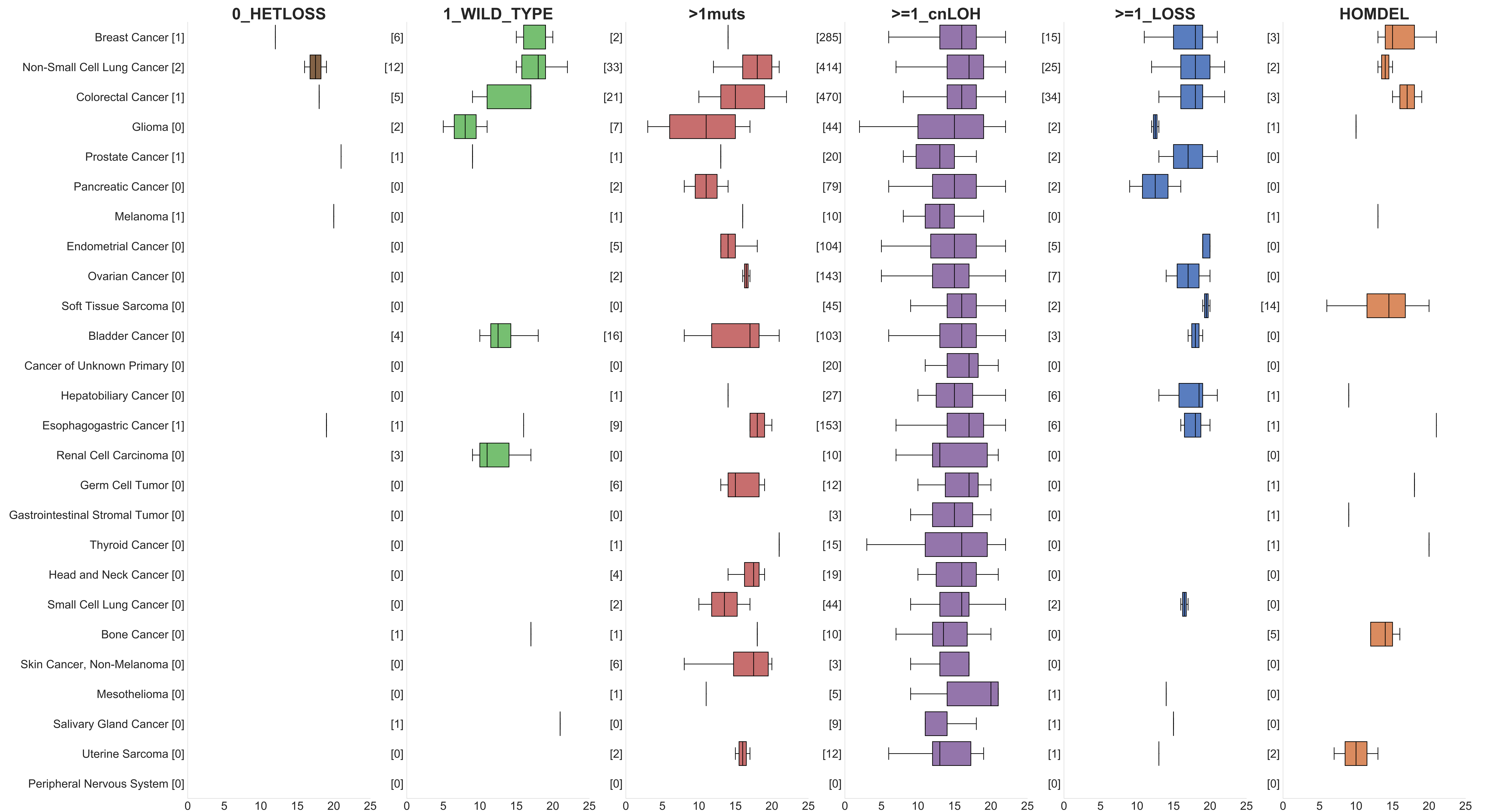
Number of chromosomes affected by a LOSS

Primary

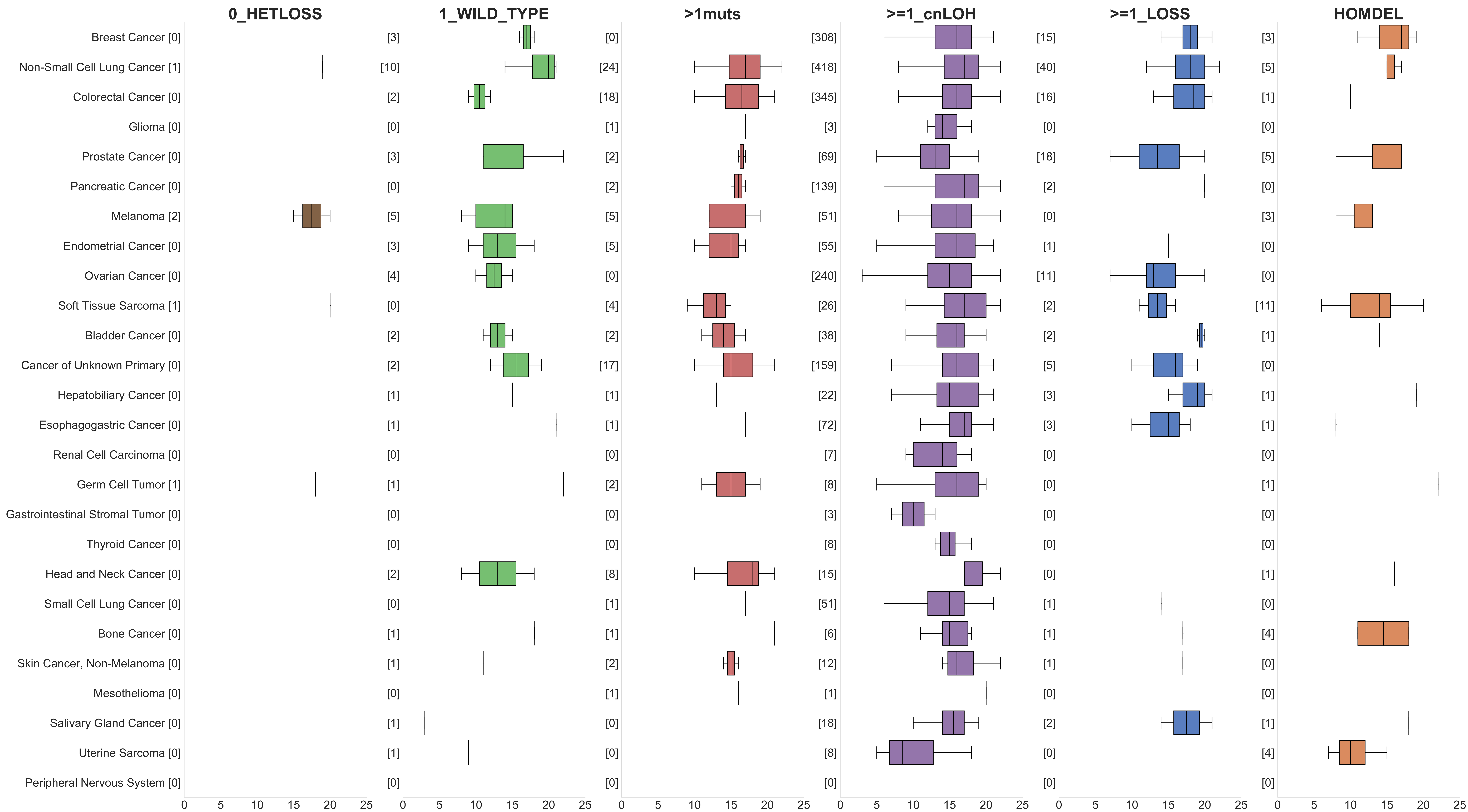
Metastasis



## Number of chromosomes affected by a LOSS - Primary Samples



Number of chromosomes affected by a LOSS - Metastatic Samples

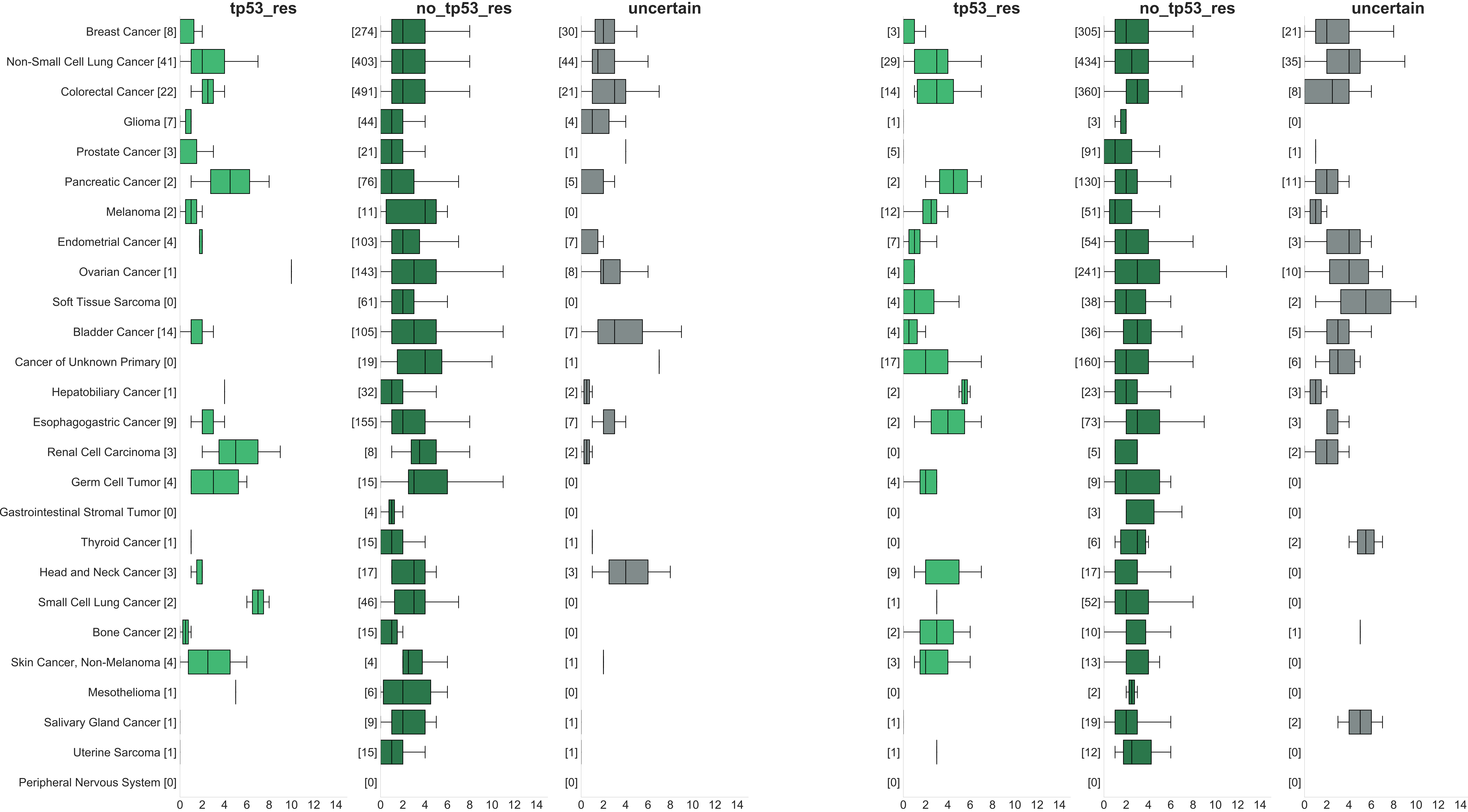




Number of chromosomes affected by a cnLOH

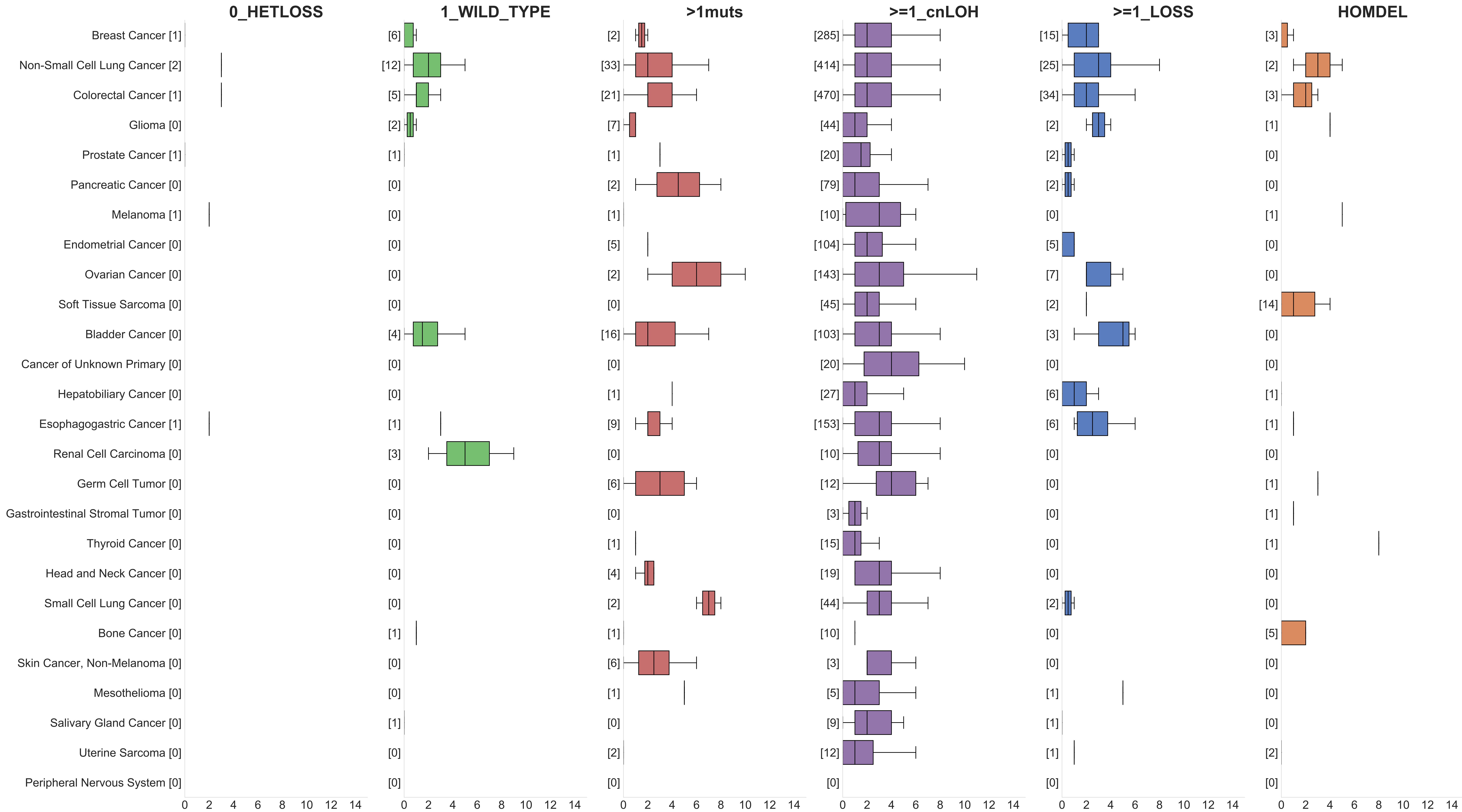
Primary

Metastasis

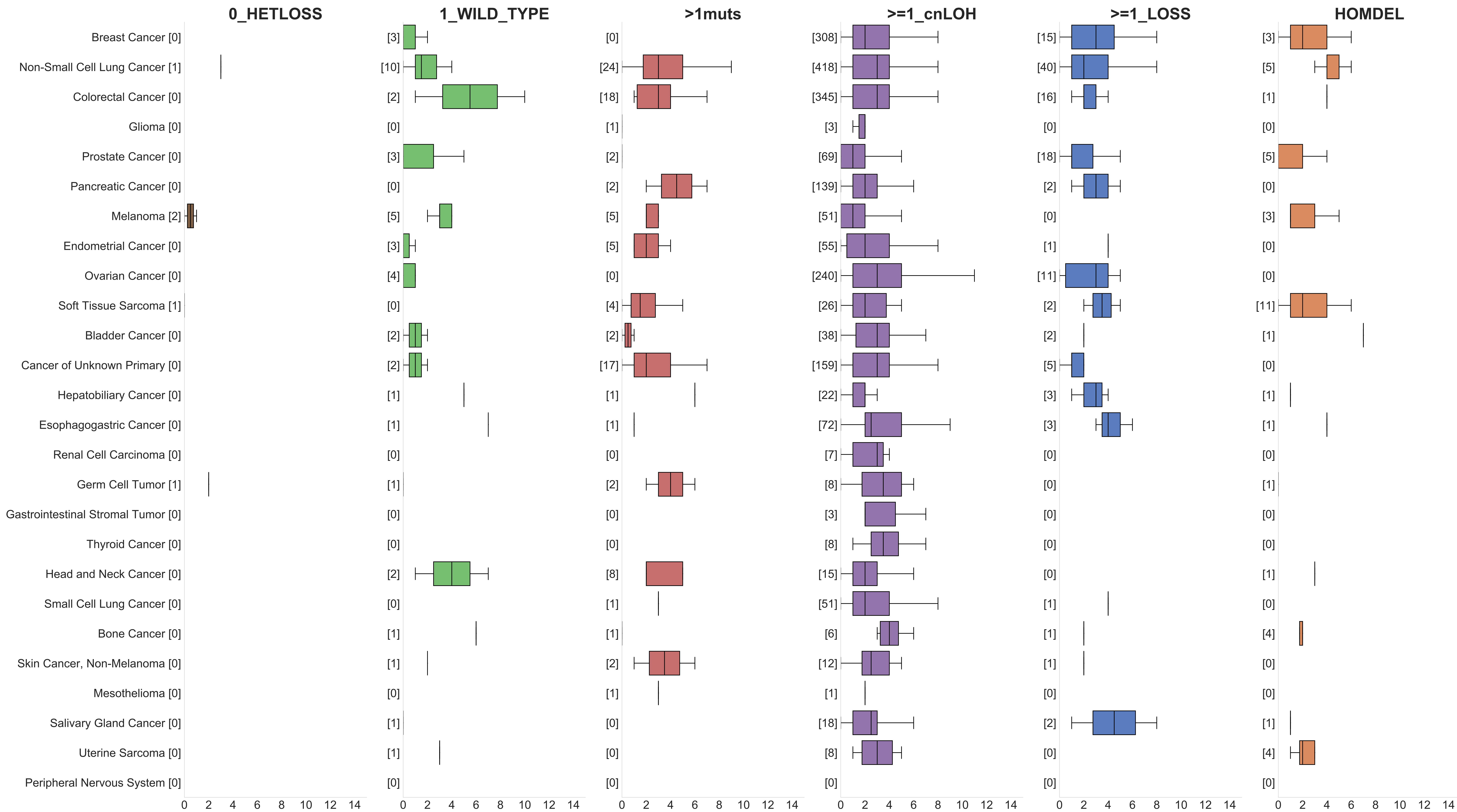




Number of chromosomes affected by a cnLOH - Primary Samples



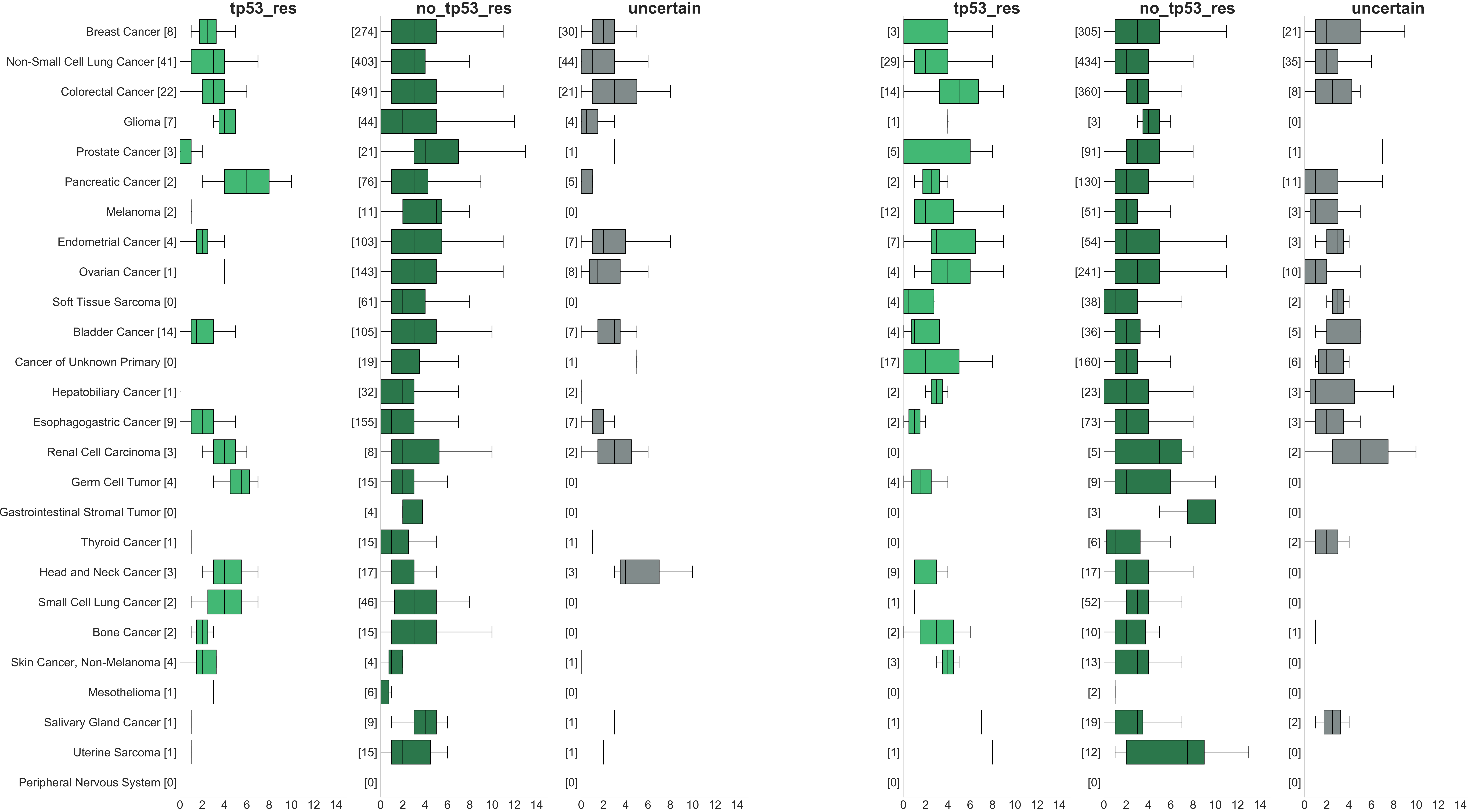
Number of chromosomes affected by a cnLOH - Metastatic Samples



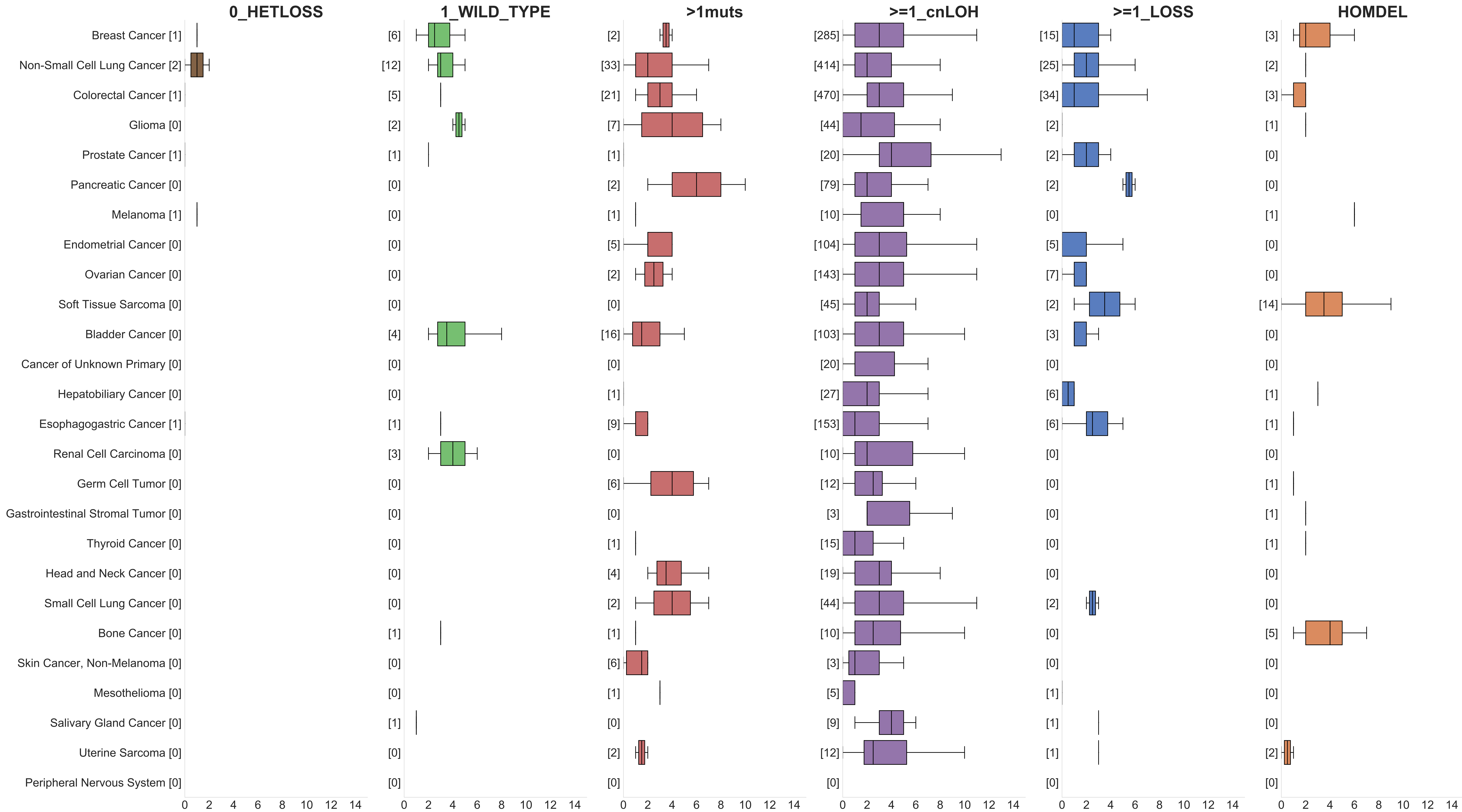
Number of chromosomes affected by a GAIN

Primary

Metastasis

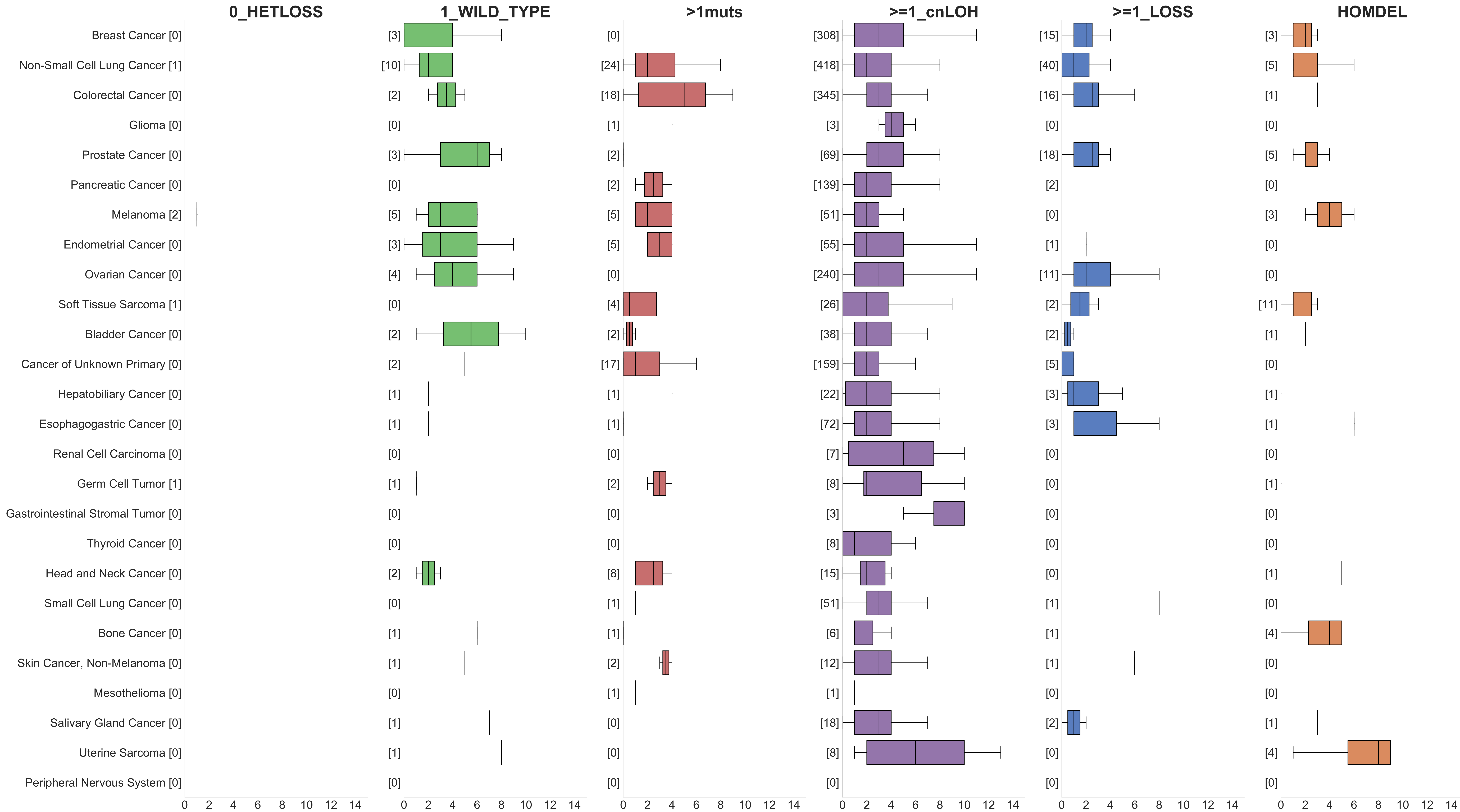


Number of chromosomes affected by a GAIN - Primary Samples





Number of chromosomes affected by a GAIN - Metastatic Samples



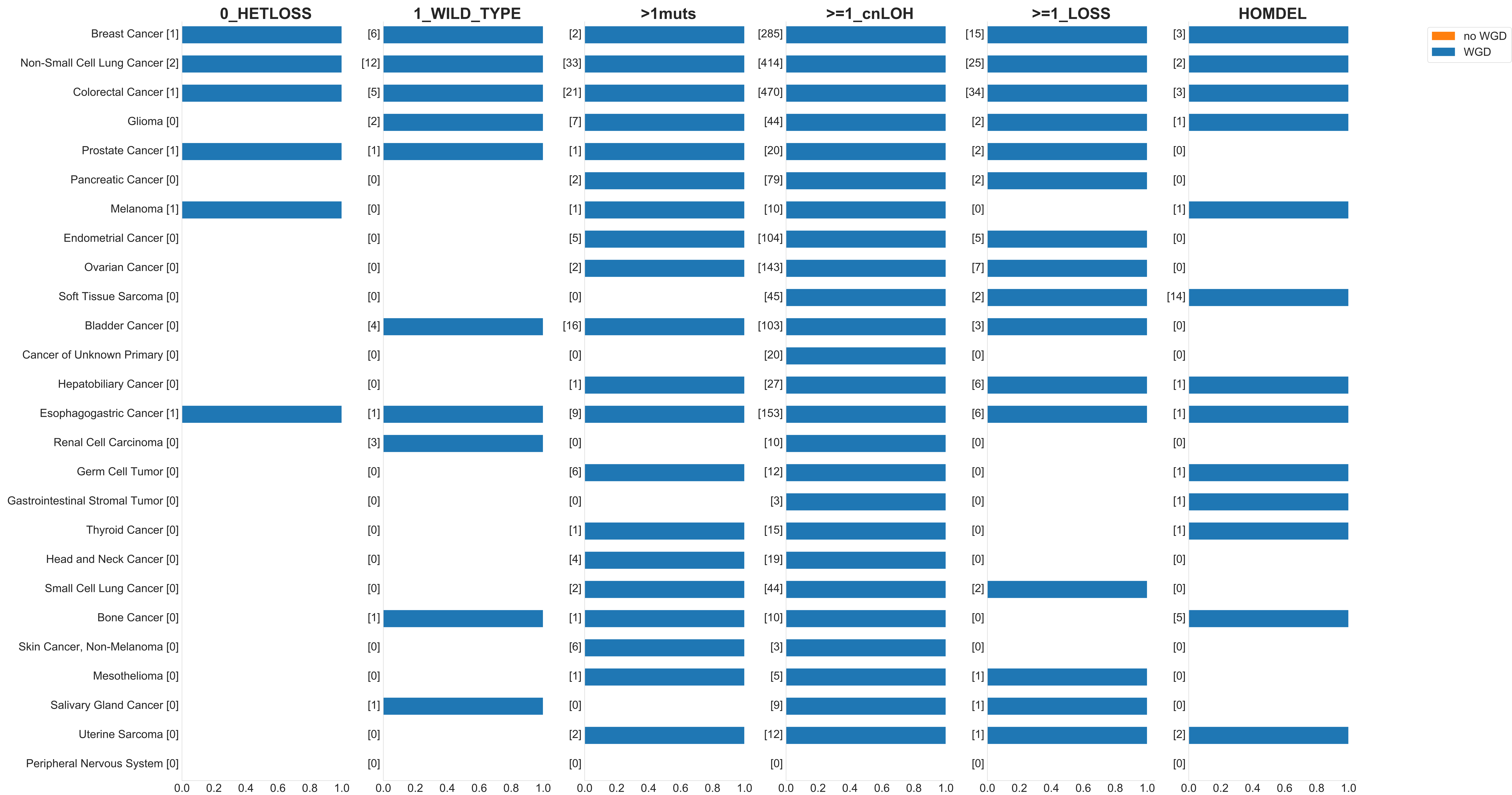
Proportion of WGD

Primary

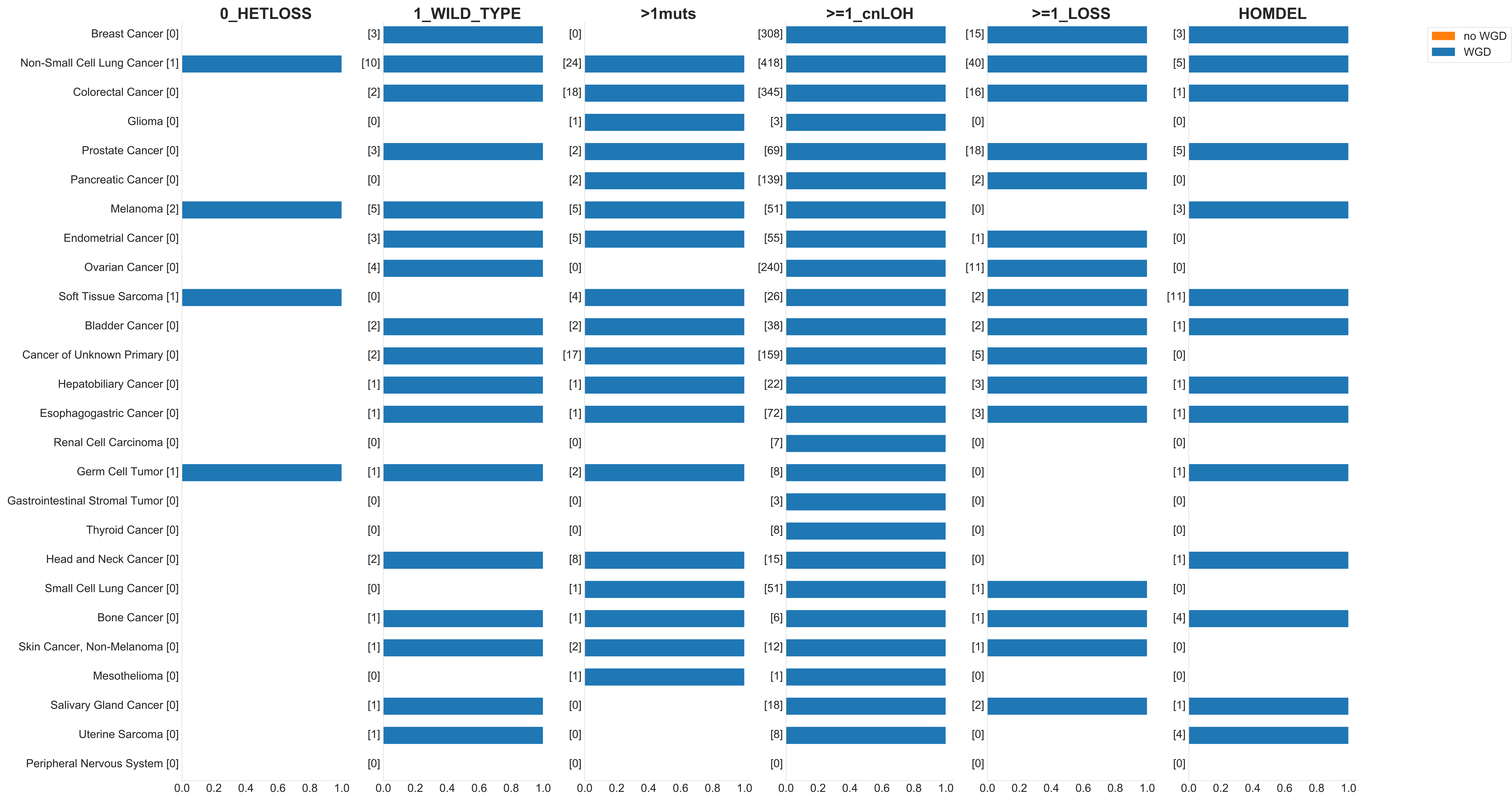
Metastasis



Proportion of WGD - Primary Samples



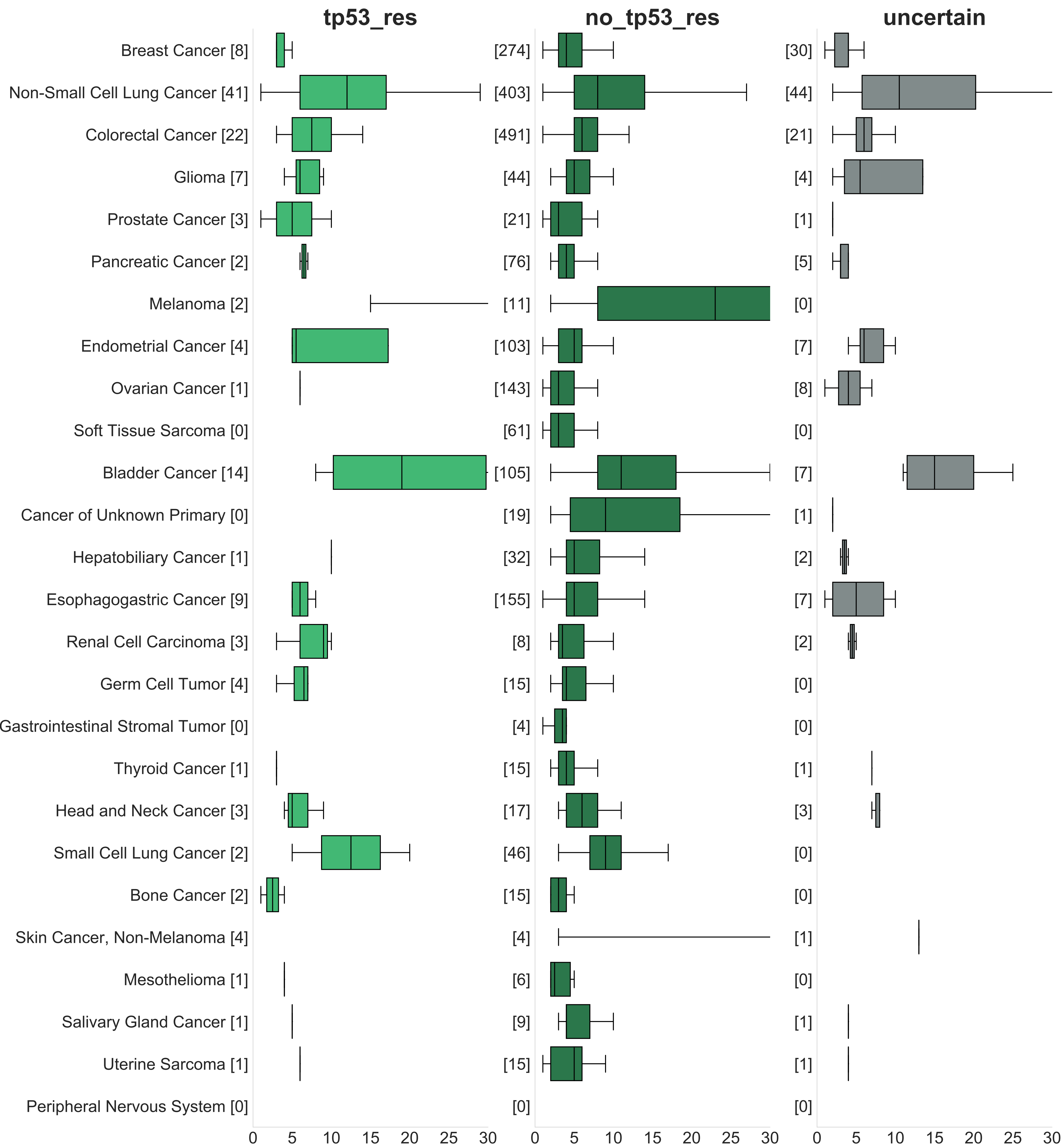
Proportion of WGD - Metastatic Samples



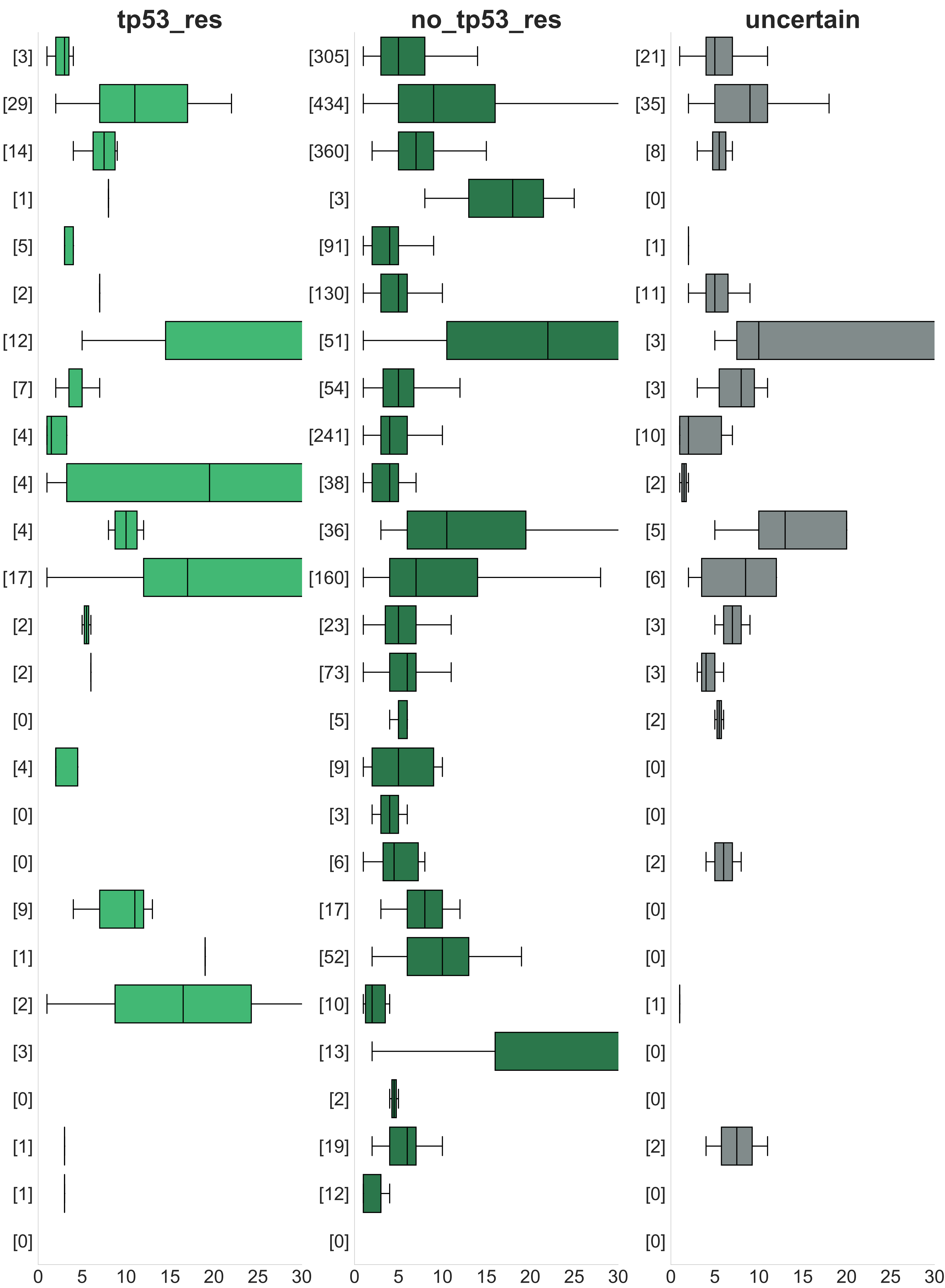


Mutation Count

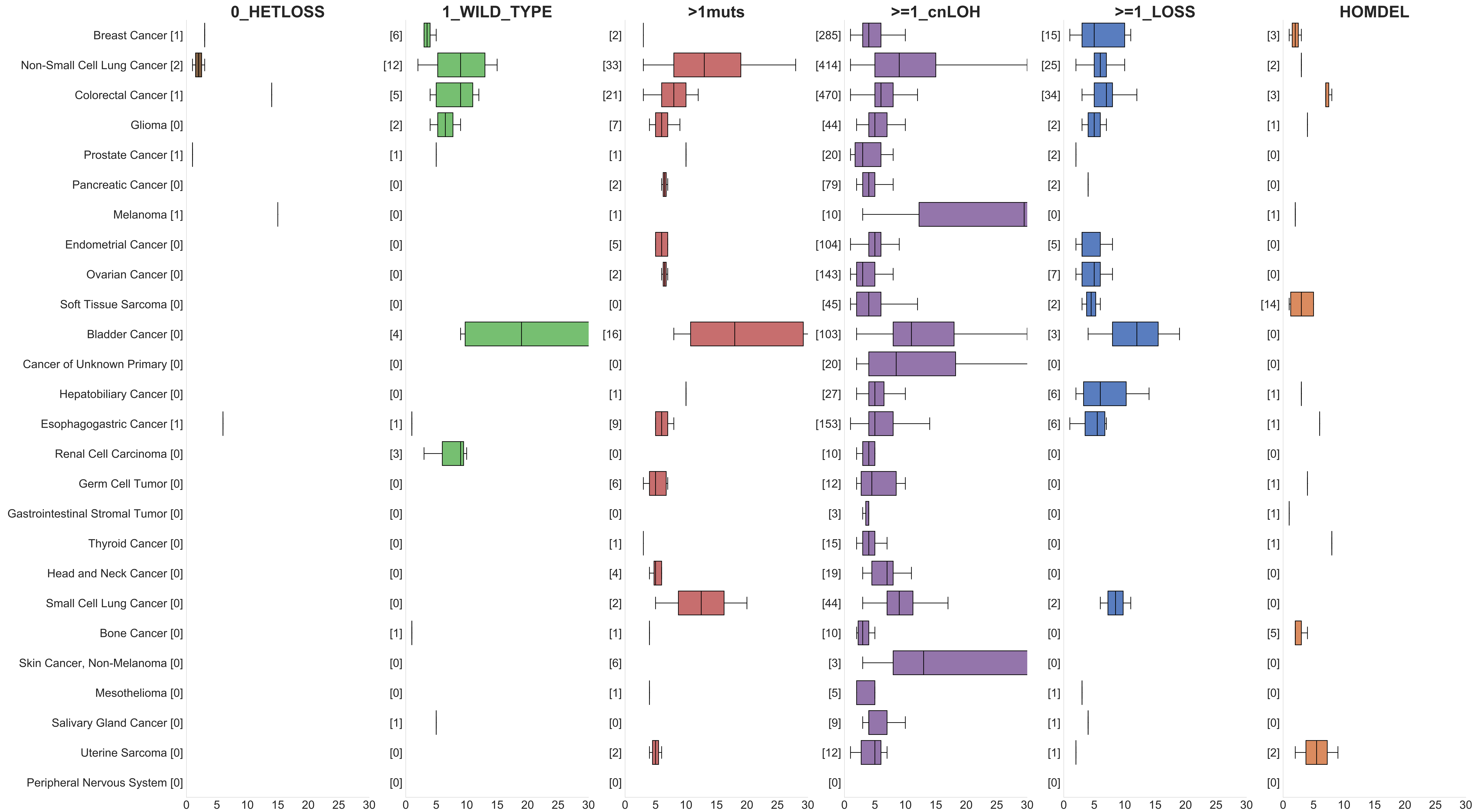
Primary



Metastasis



Mutation Count - Primary Samples



Mutation Count - Metastatic Samples

