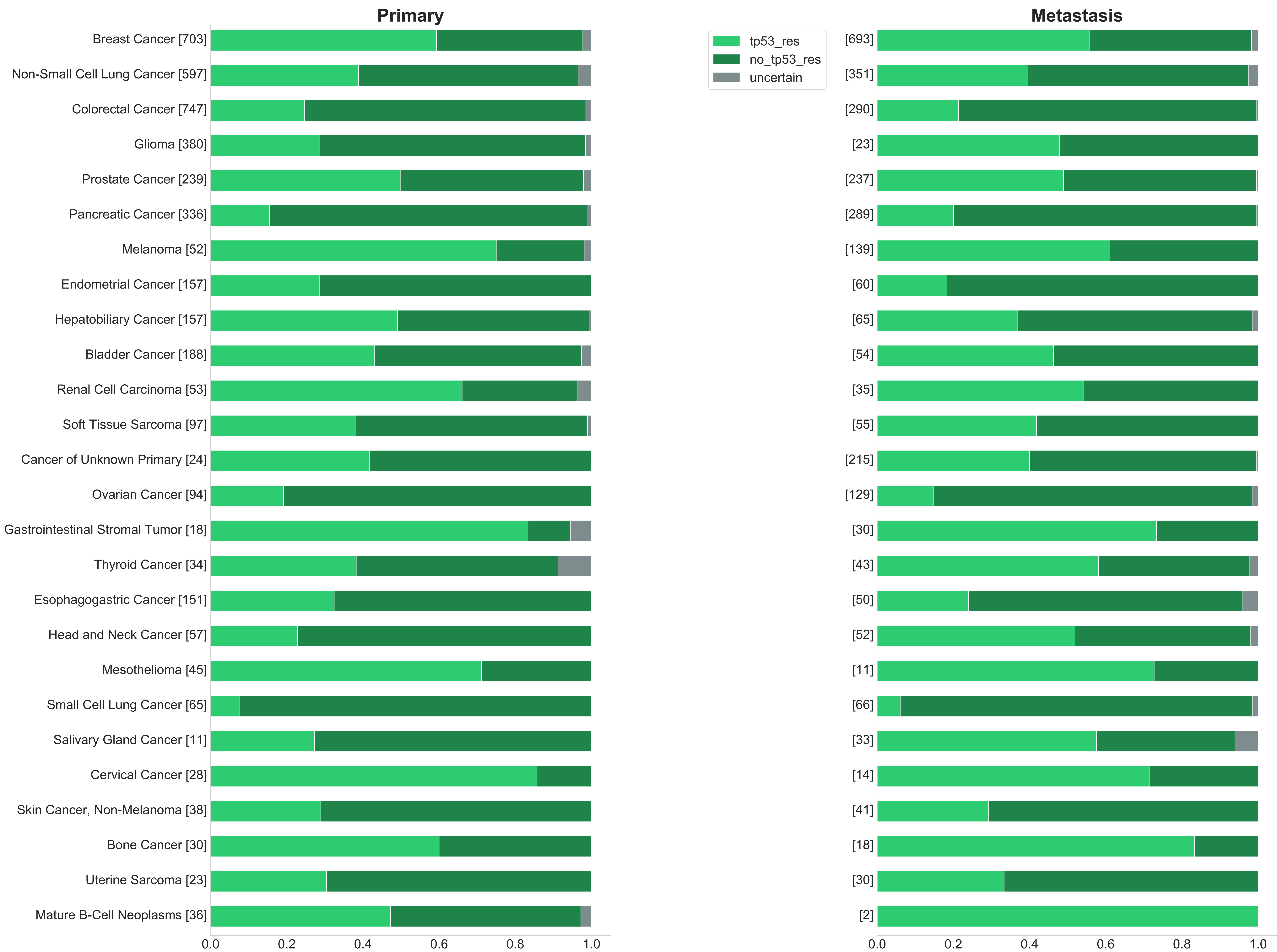
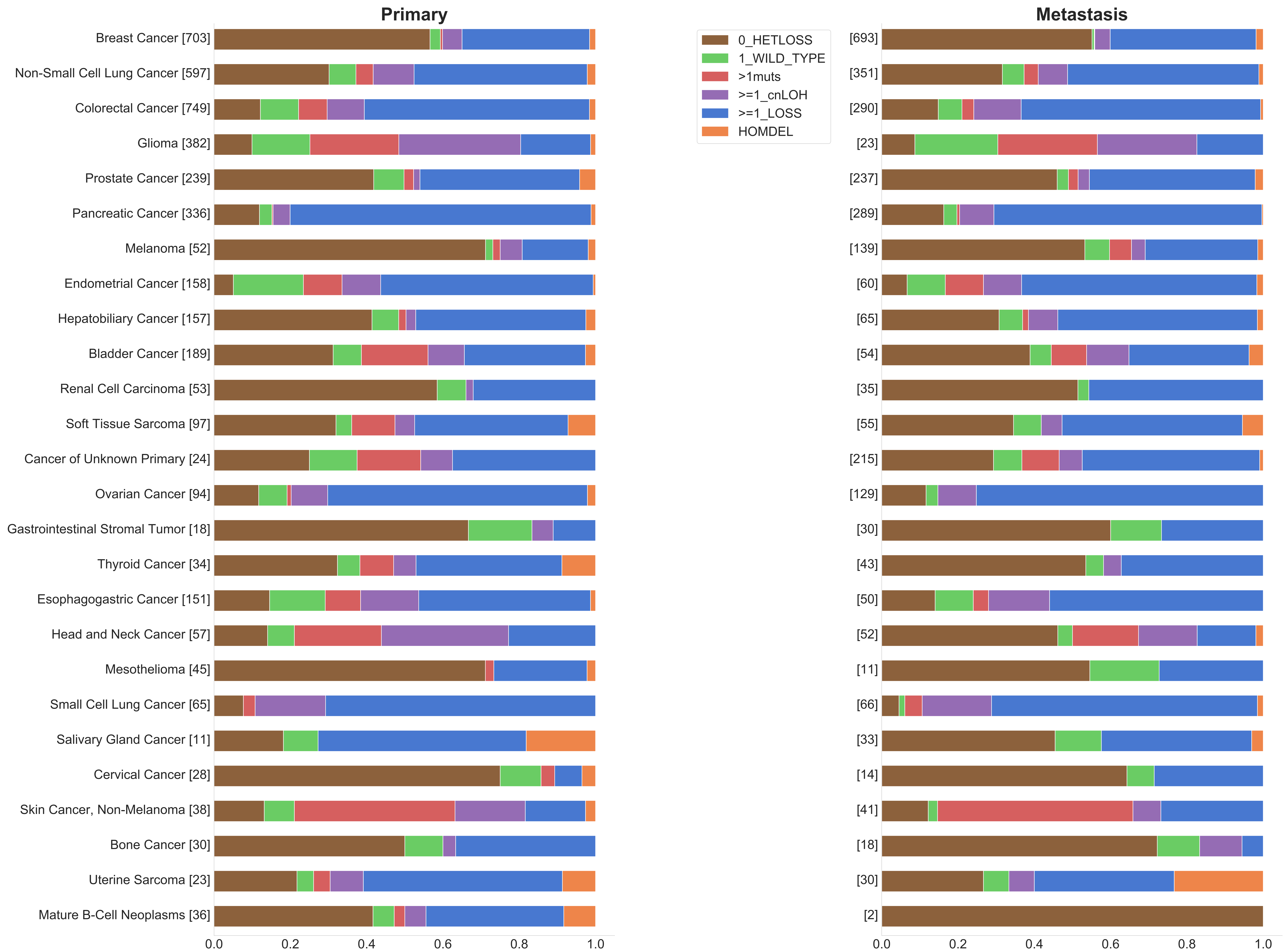


TP53 RESIDUAL SUBGROUP PROPORTION ACROSS CANCERS



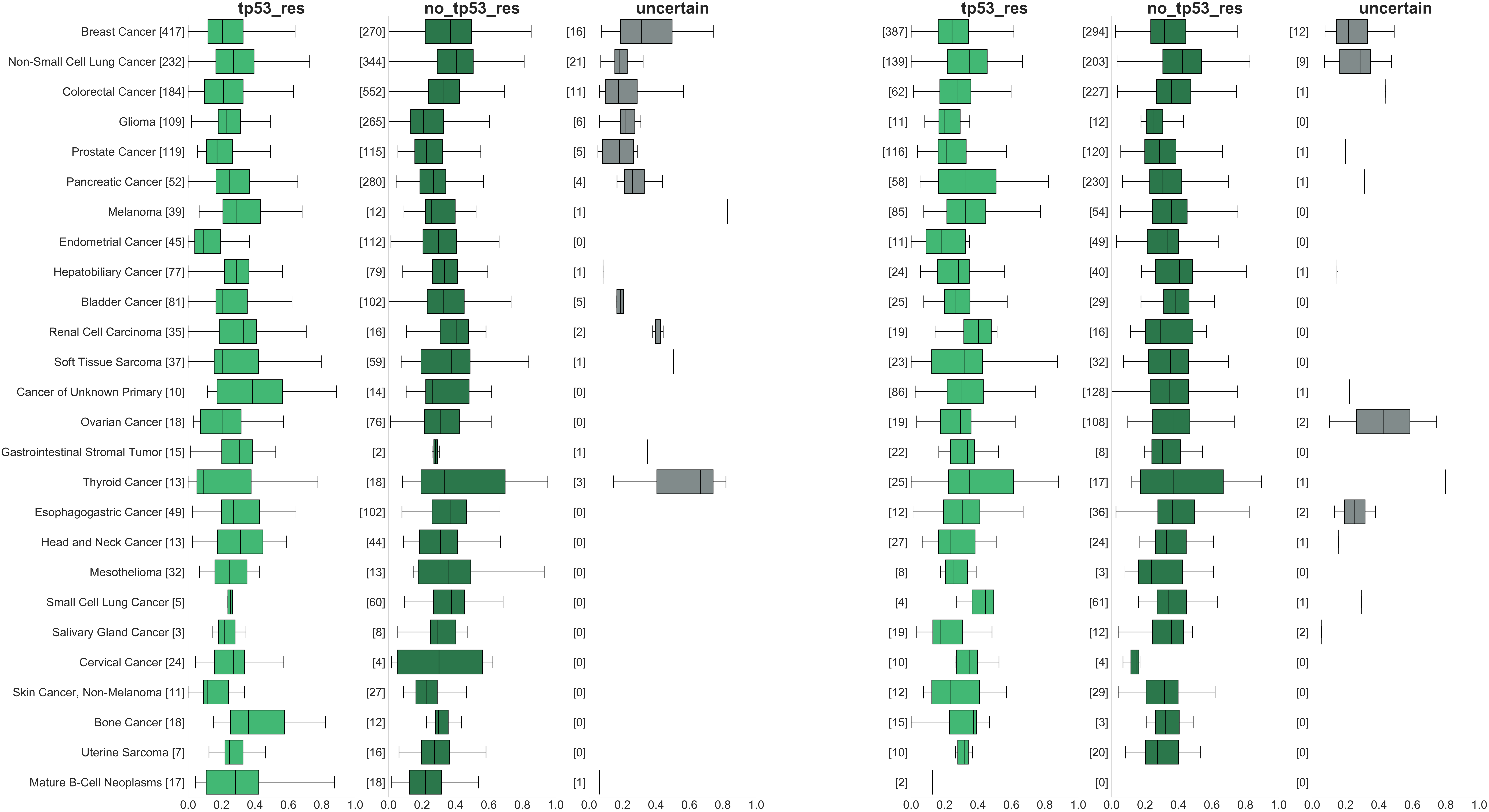
# TP53 SUBGROUP PROPORTION ACROSS CANCERS



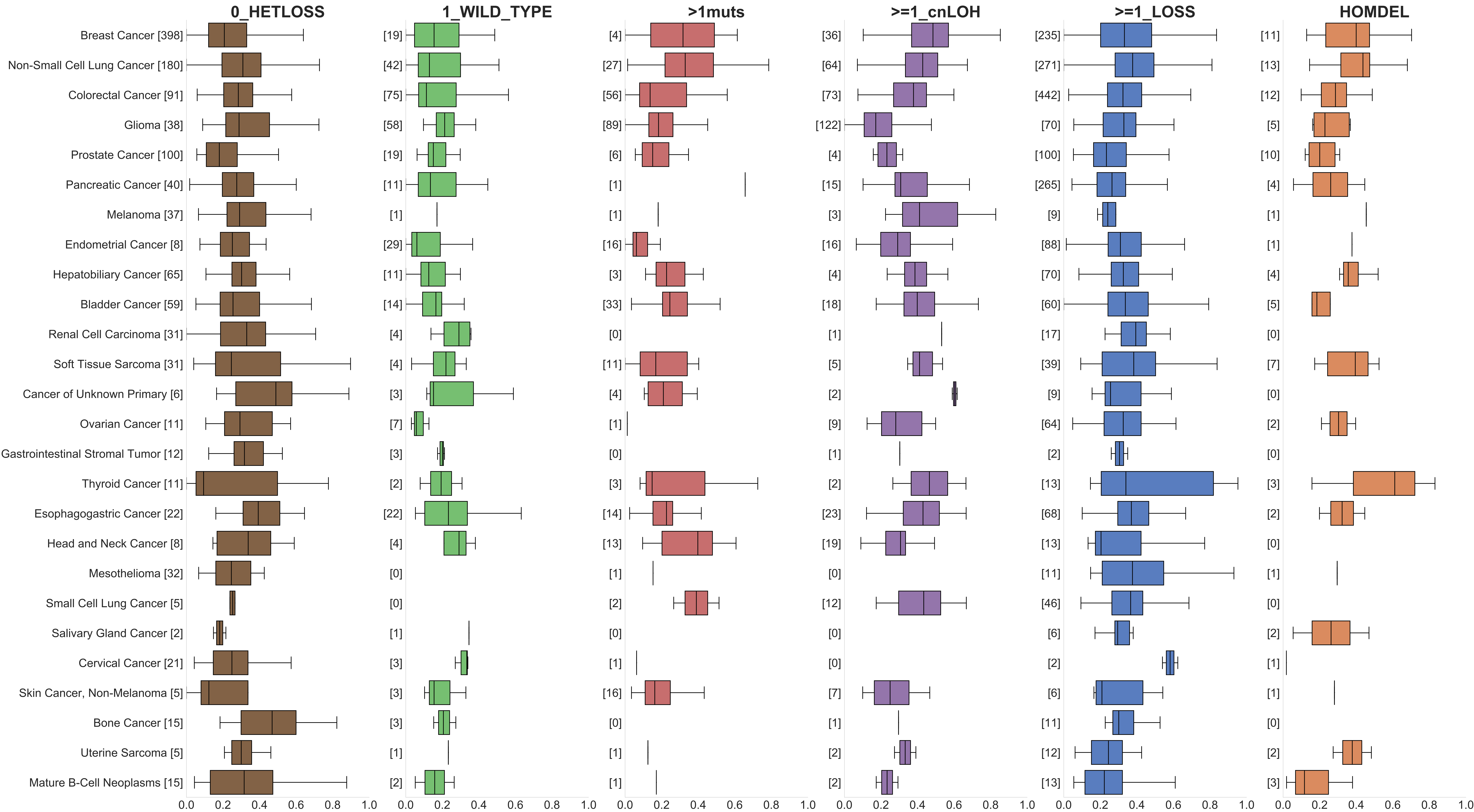
Fraction of genome altered

Primary

Metastasis

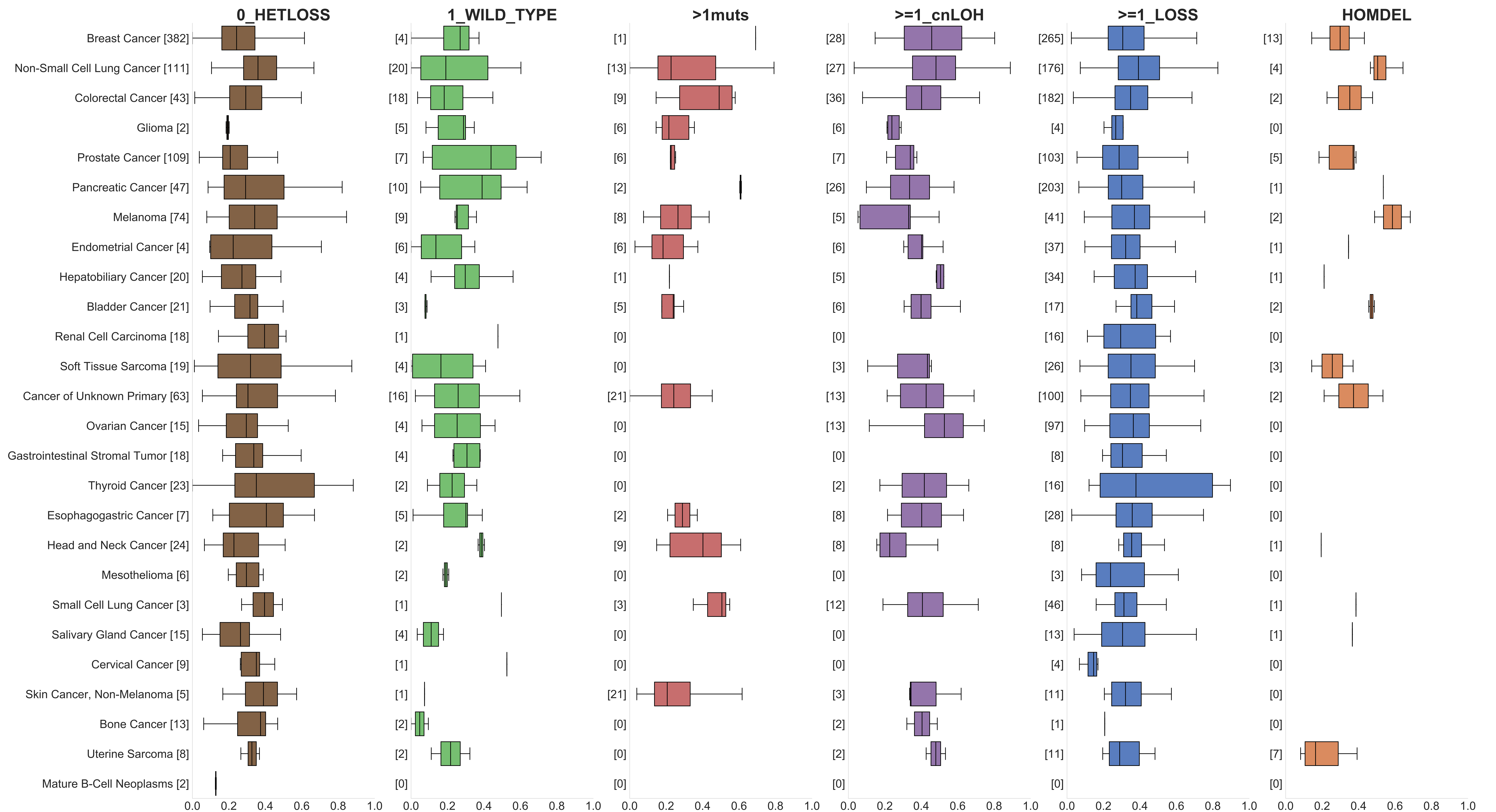


Fraction of genome altered - Primary Samples





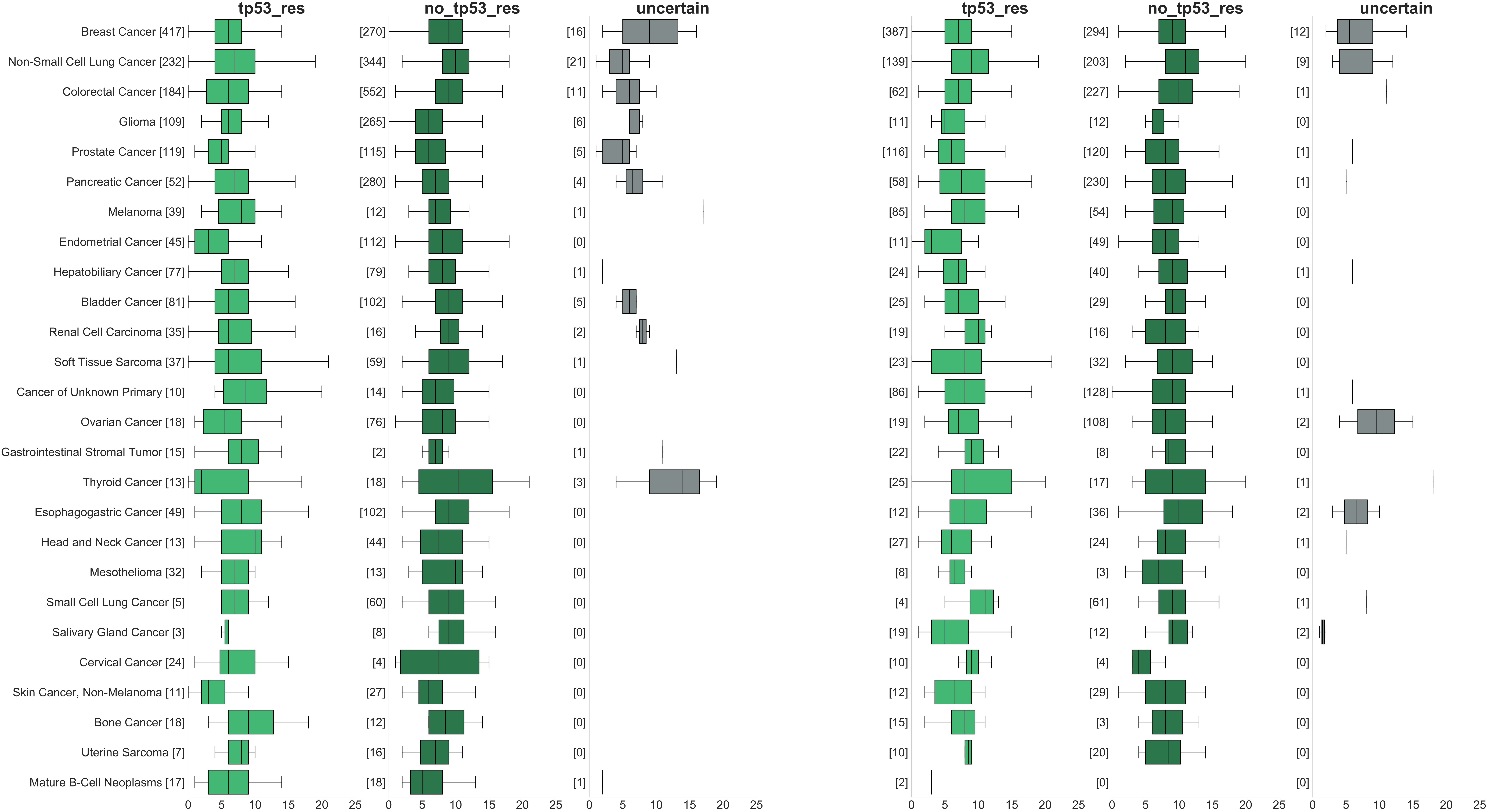
# Fraction of genome altered - Metastatic Samples



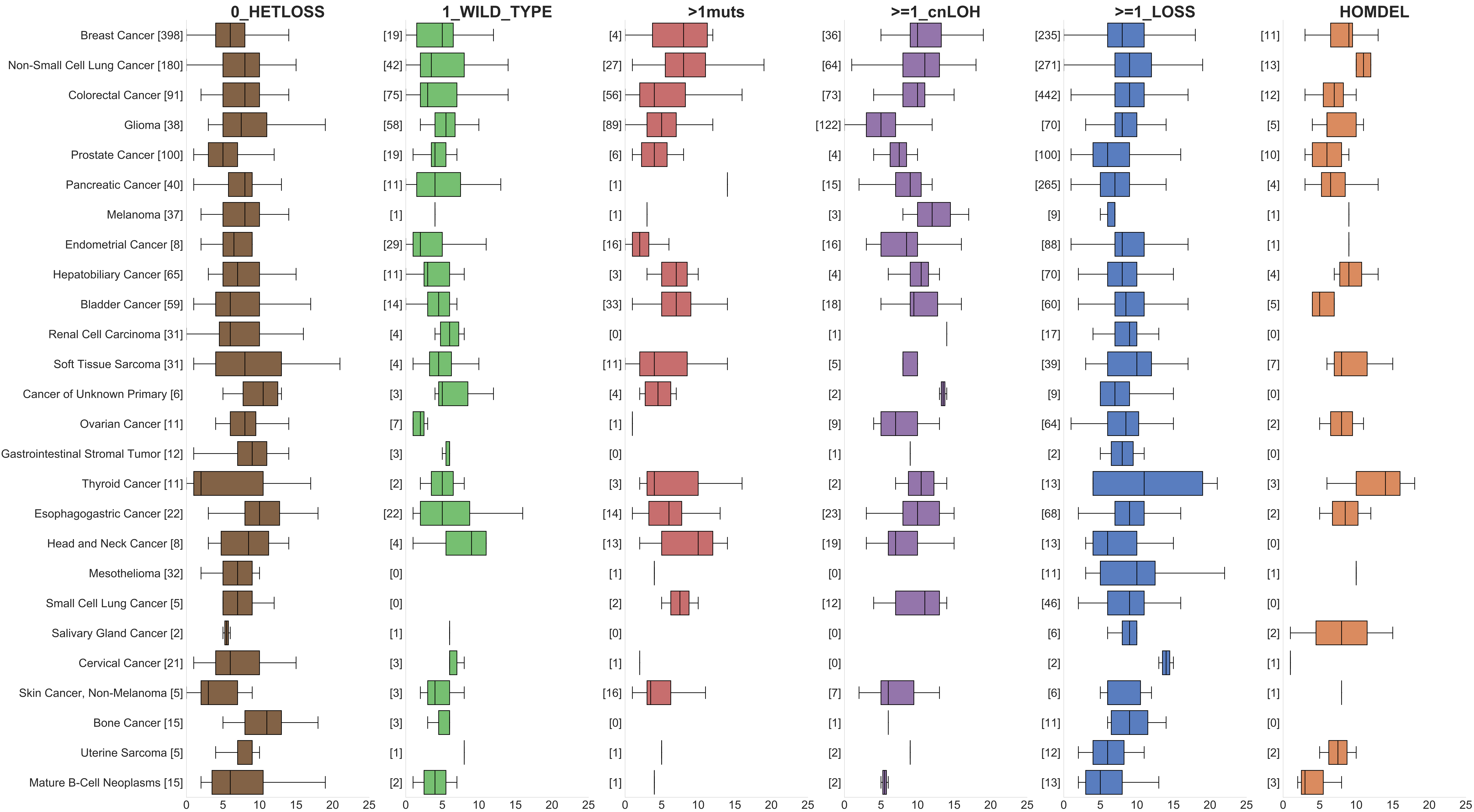
Number of aberrant chromosomes

Primary

Metastasis

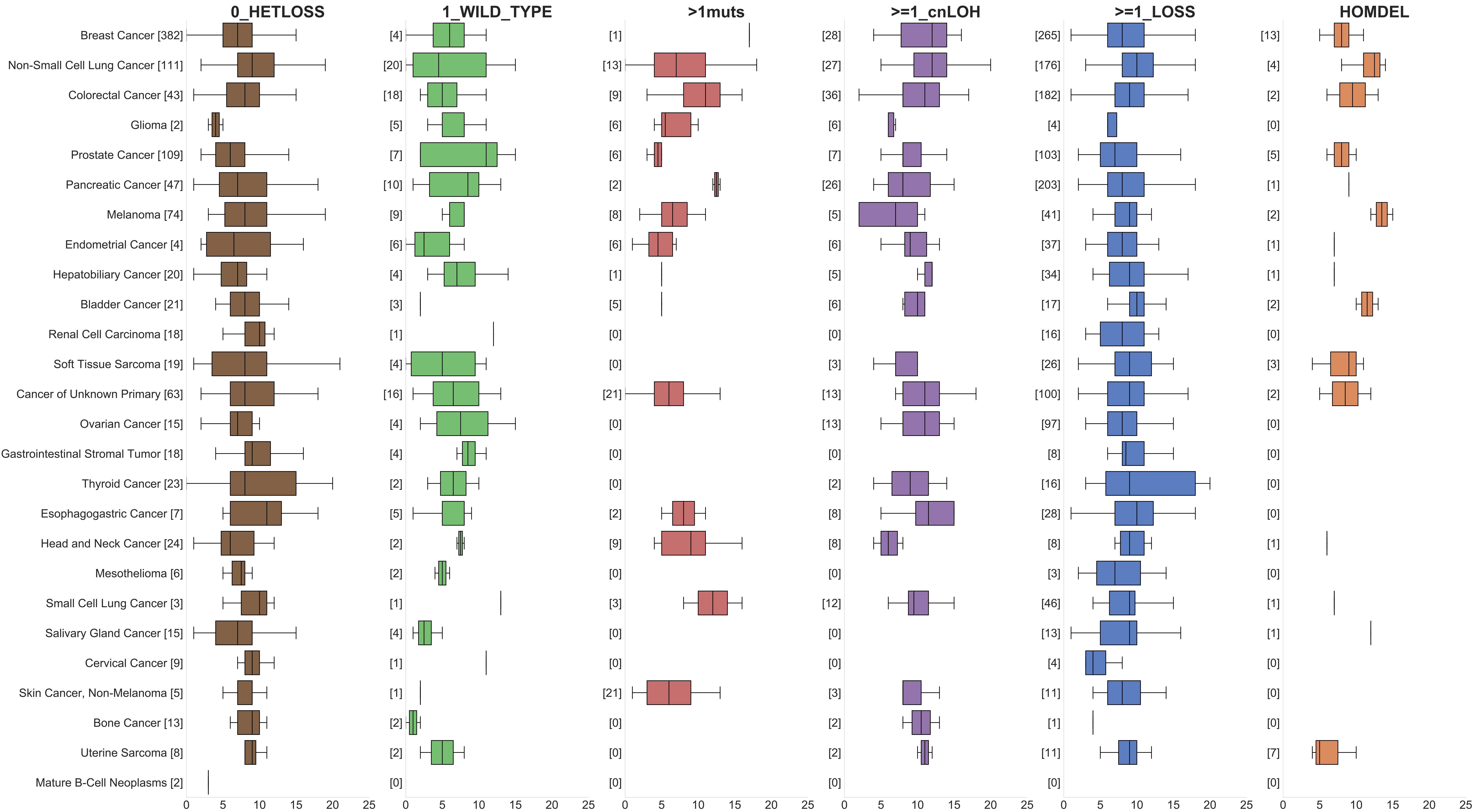


Number of aberrant chromosomes - Primary Samples





Number of aberrant chromosomes - Metastatic Samples

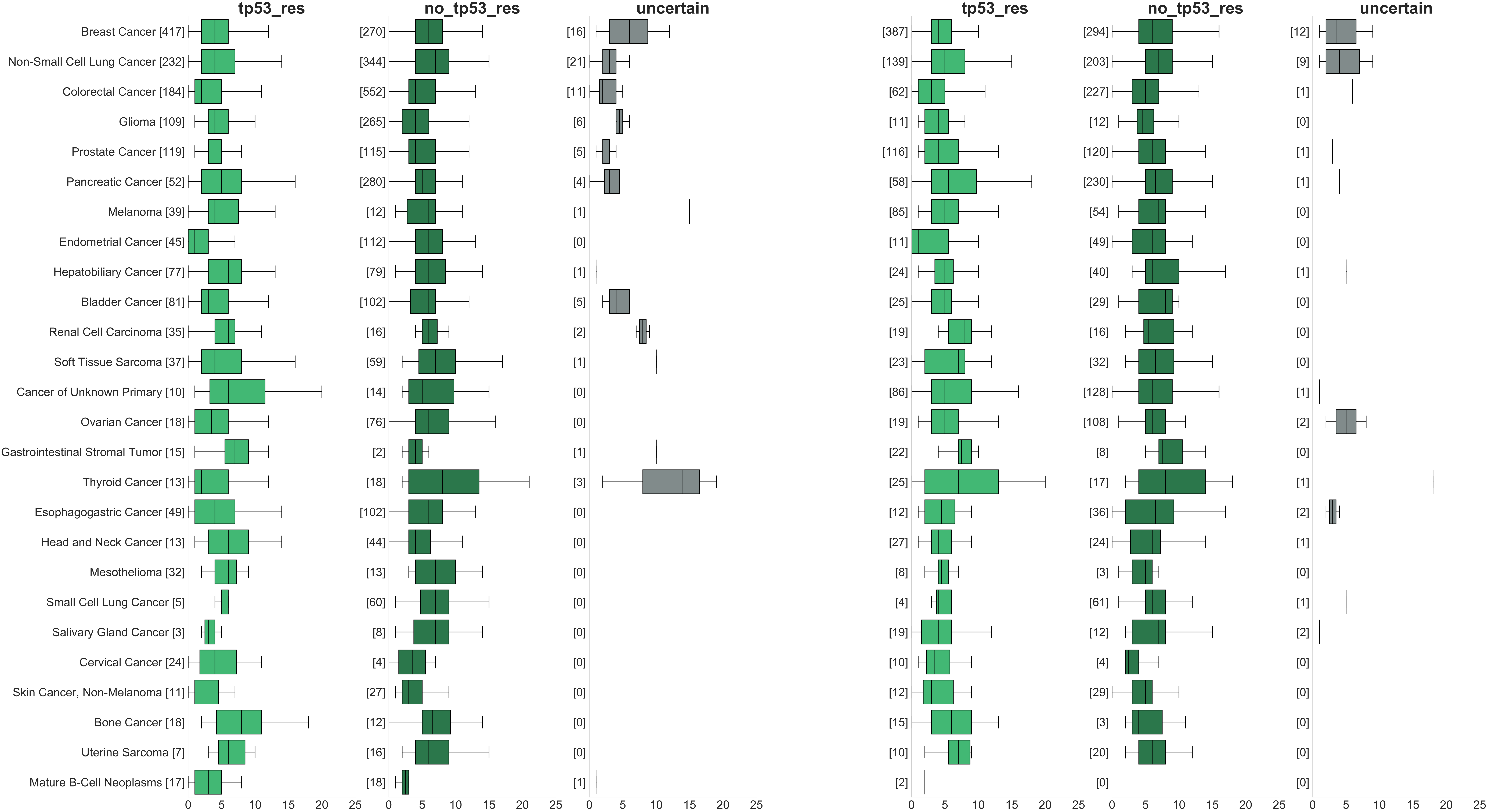




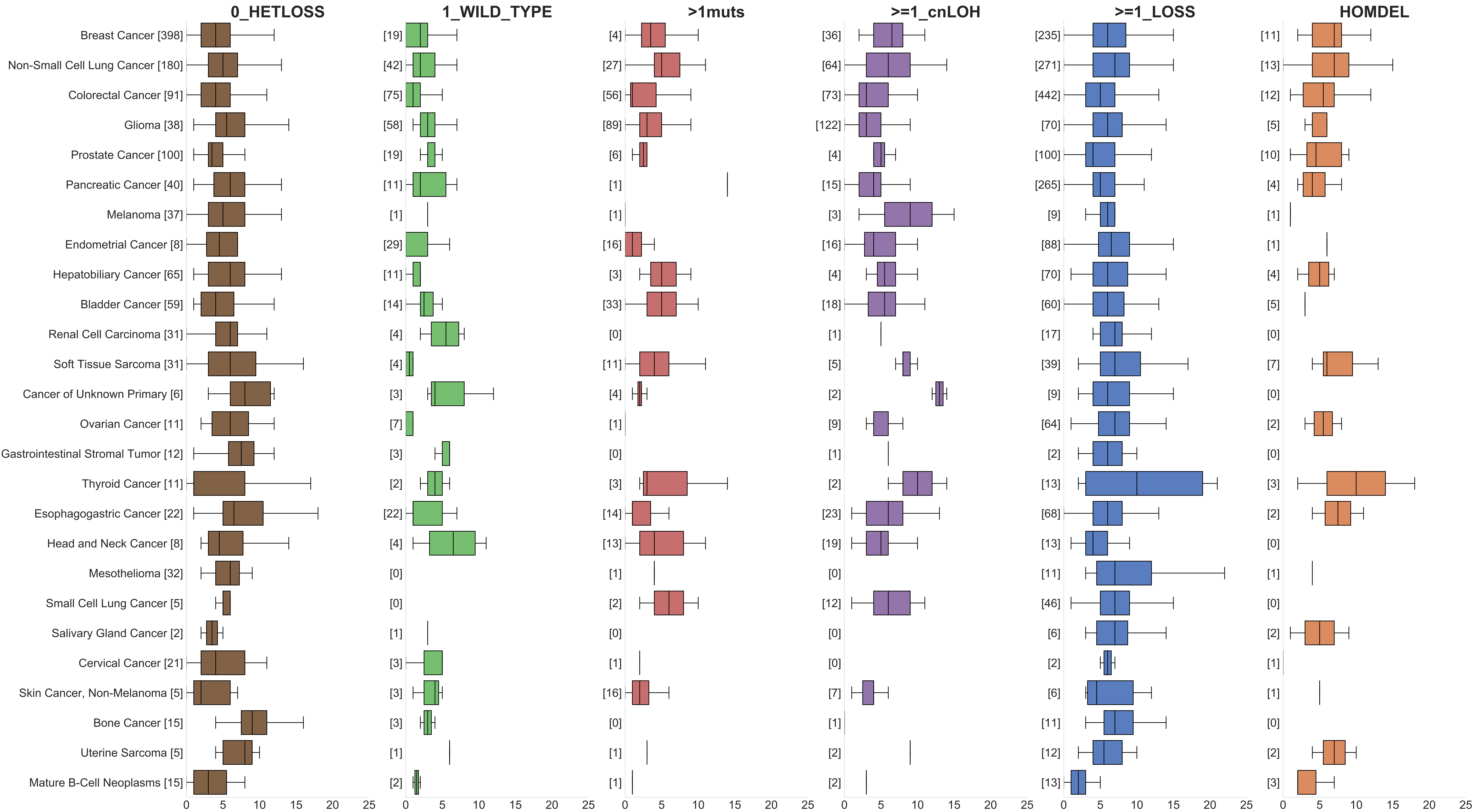
Number of chromosomes affected by a LOSS

Primary

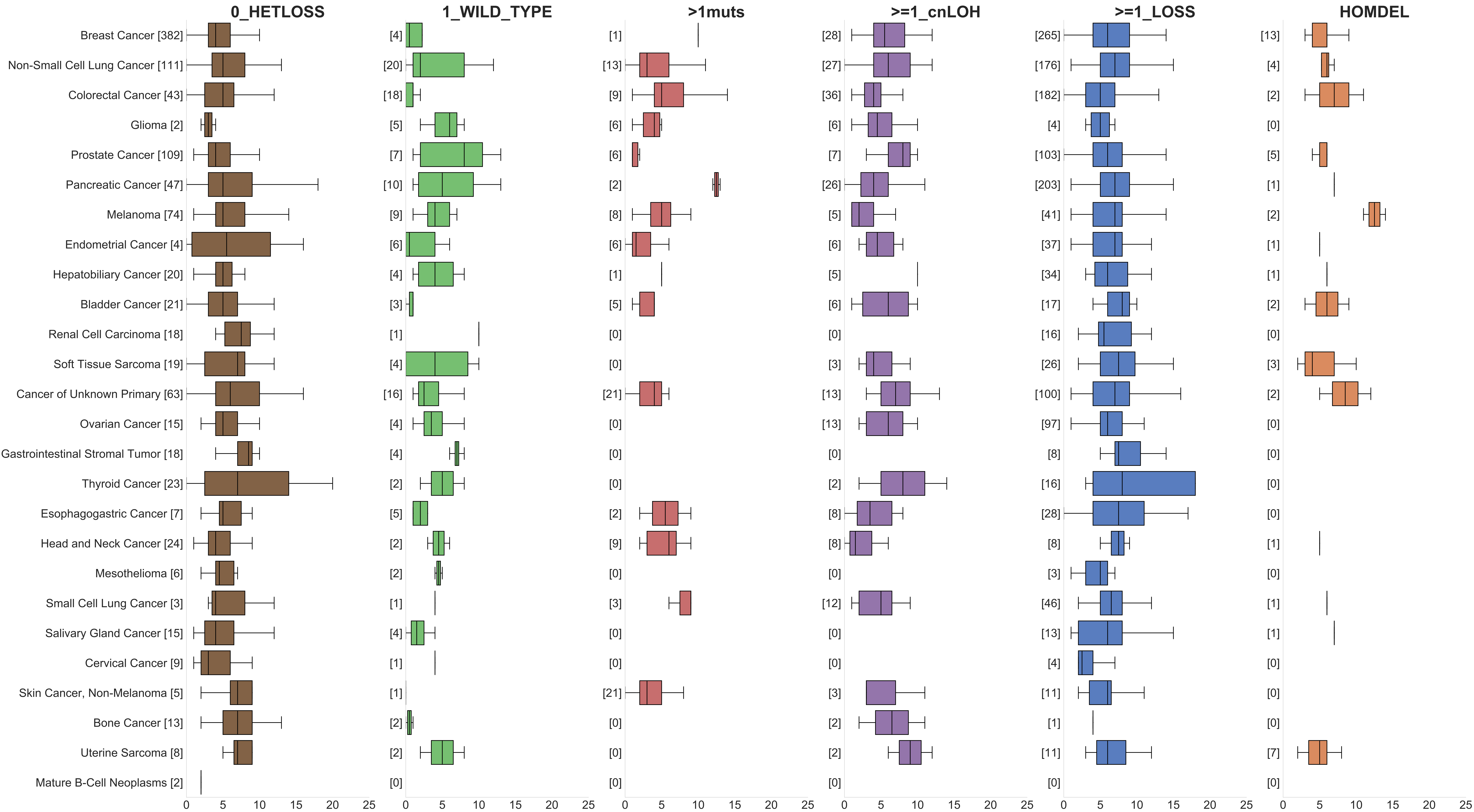
Metastasis



Number of chromosomes affected by a LOSS - Primary Samples

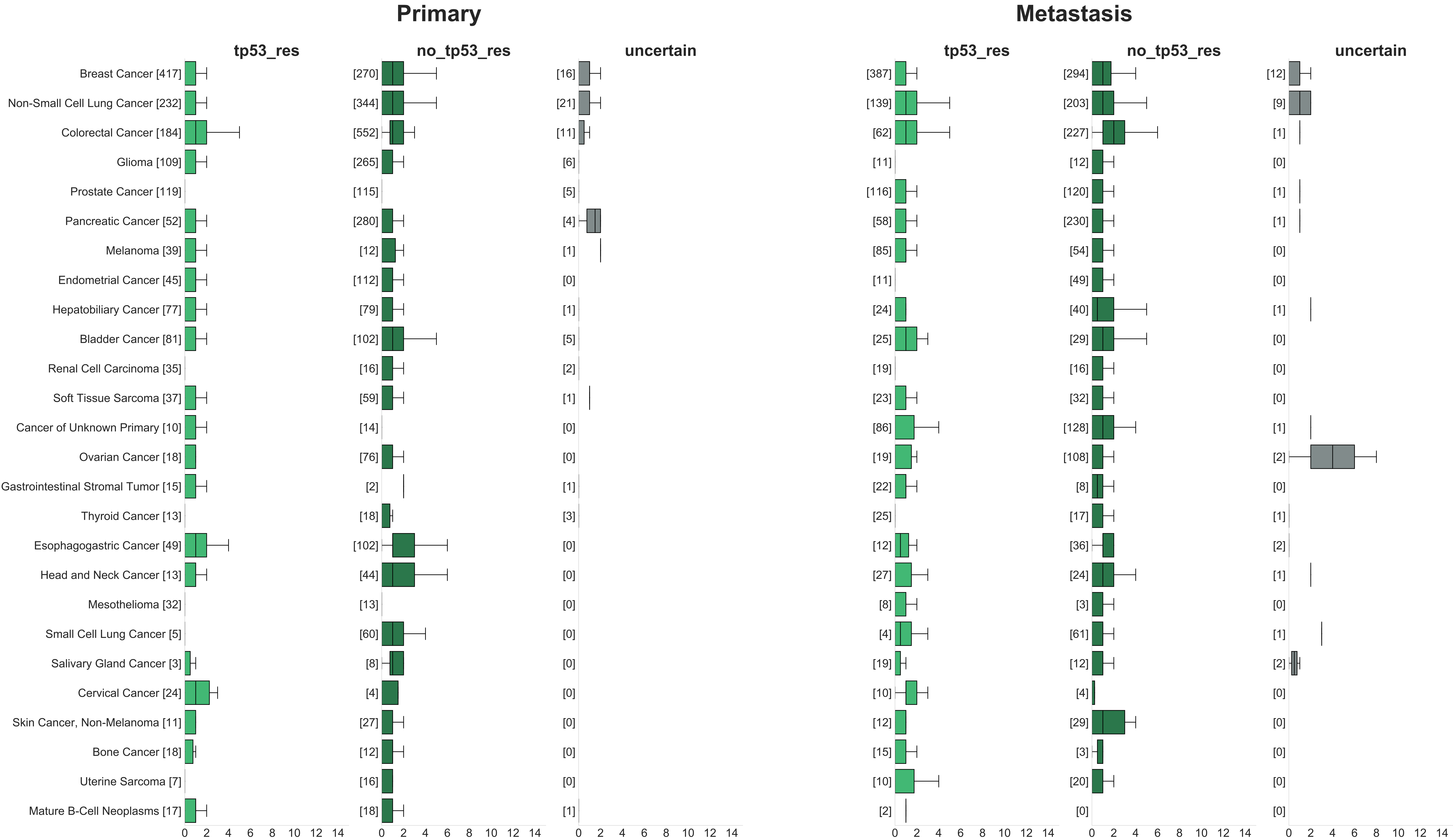


Number of chromosomes affected by a LOSS - Metastatic Samples



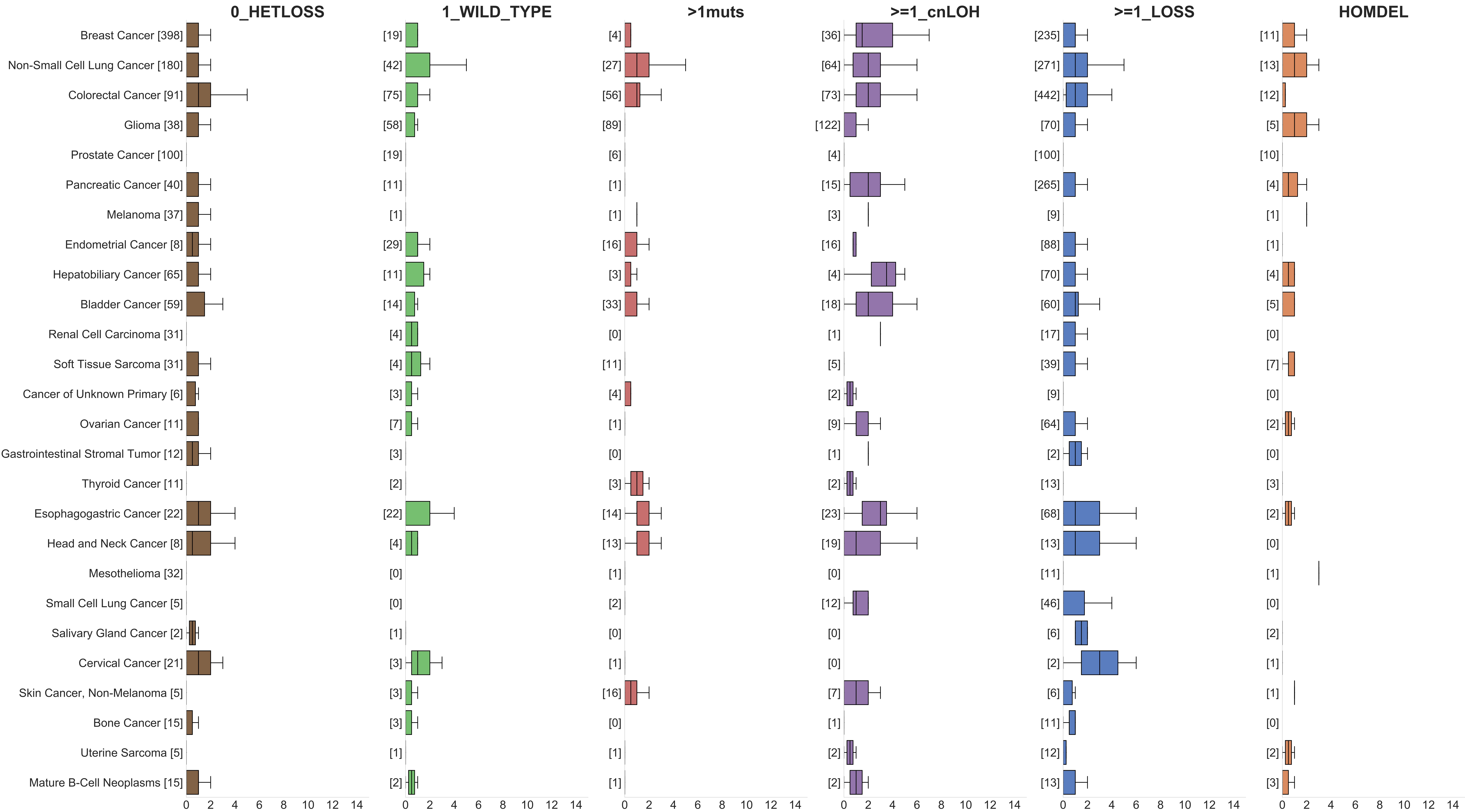


Number of chromosomes affected by a cnLOH

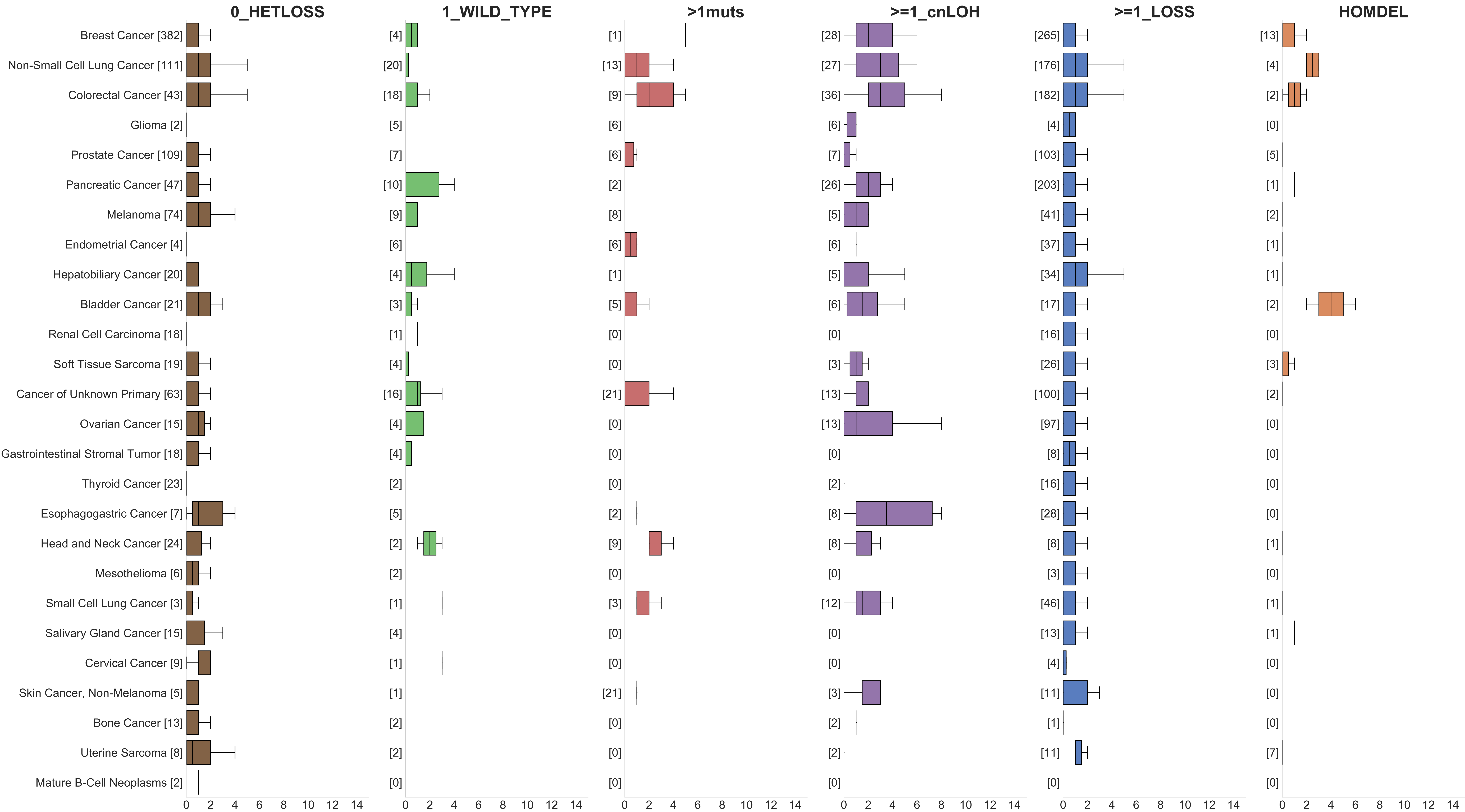




Number of chromosomes affected by a cnLOH - Primary Samples



Number of chromosomes affected by a cnLOH - Metastatic Samples



Number of chromosomes affected by a GAIN

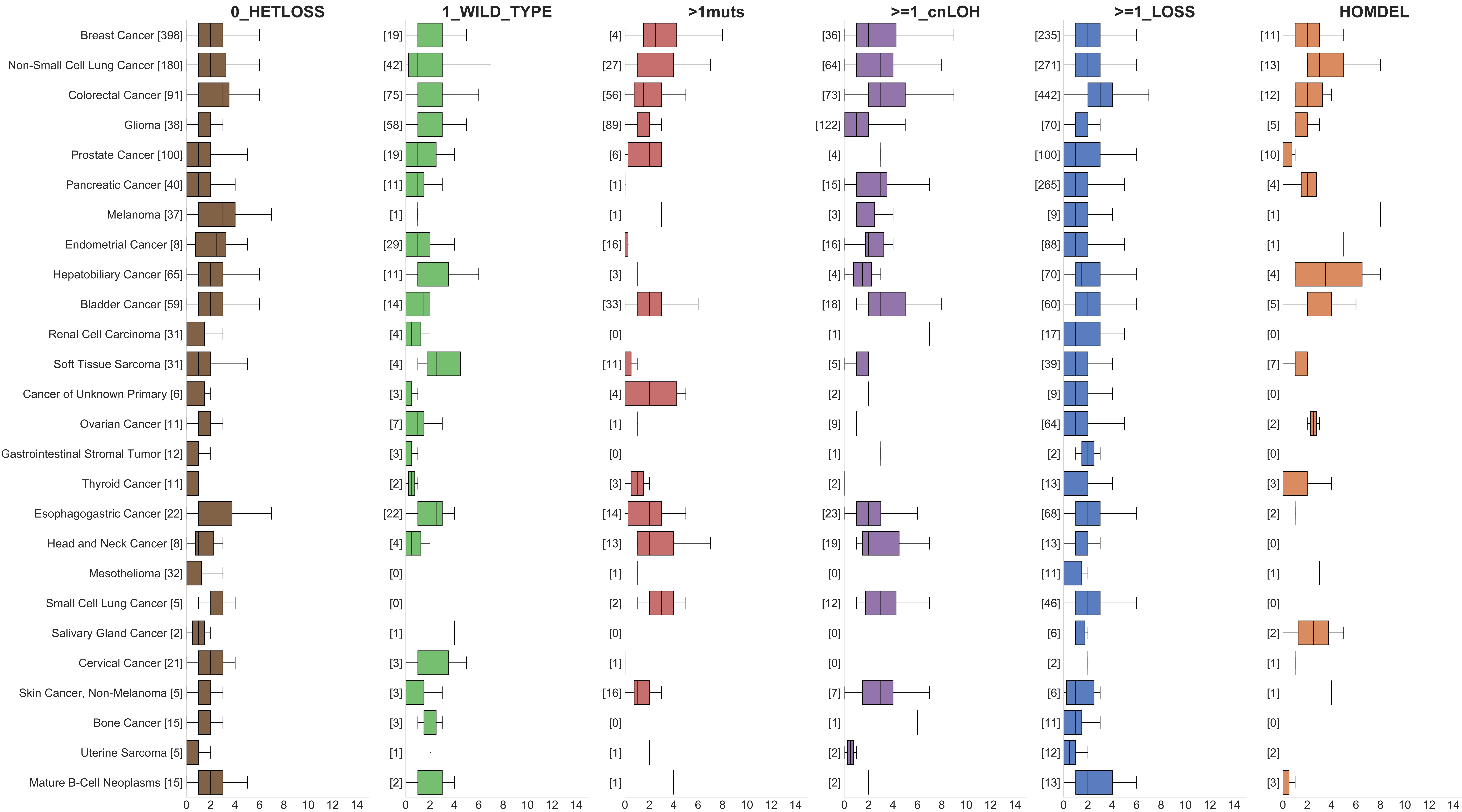
Primary

Metastasis



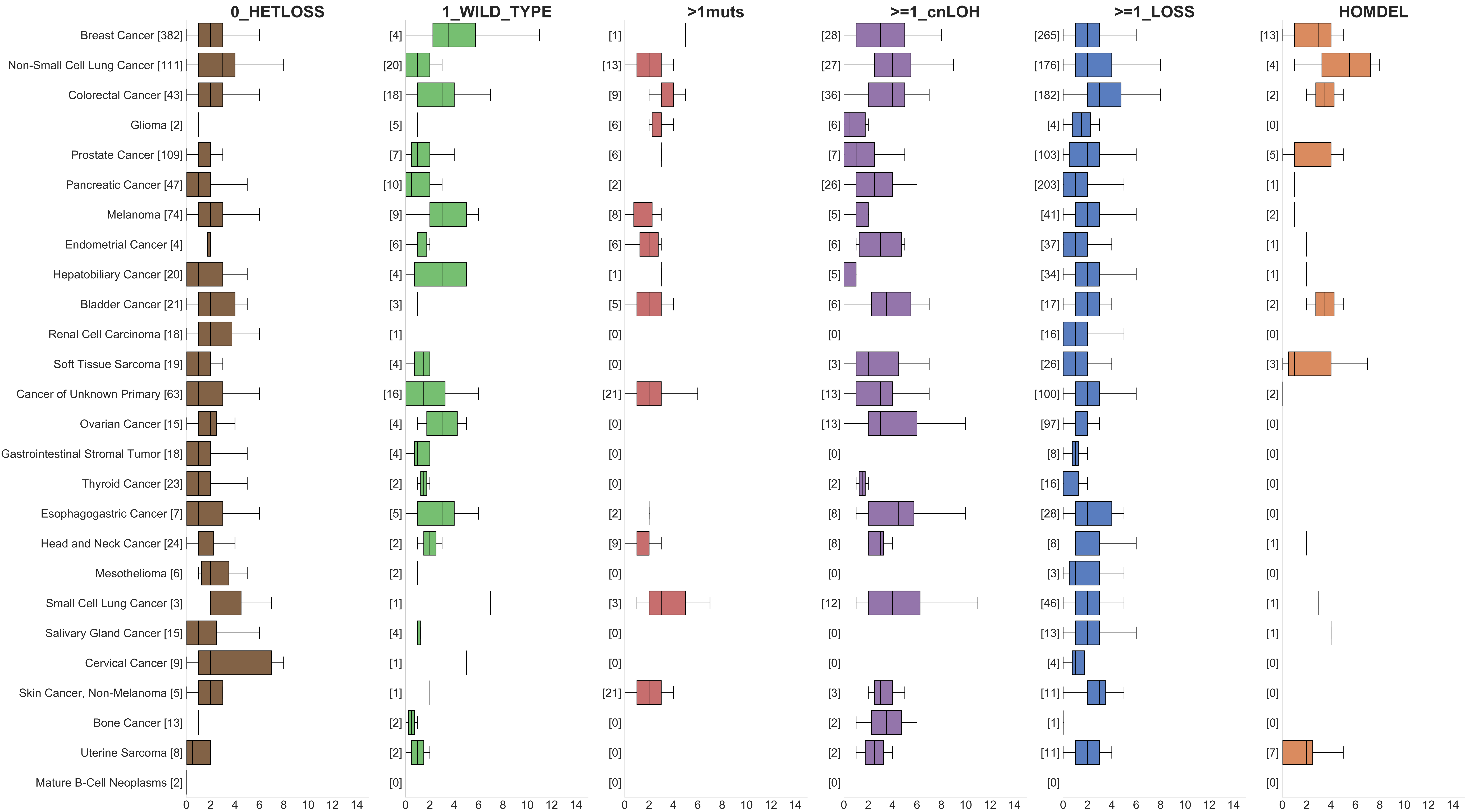


Number of chromosomes affected by a GAIN - Primary Samples





Number of chromosomes affected by a GAIN - Metastatic Samples



Proportion of WGD



Proportion of WGD - Primary Samples





Proportion of WGD - Metastatic Samples

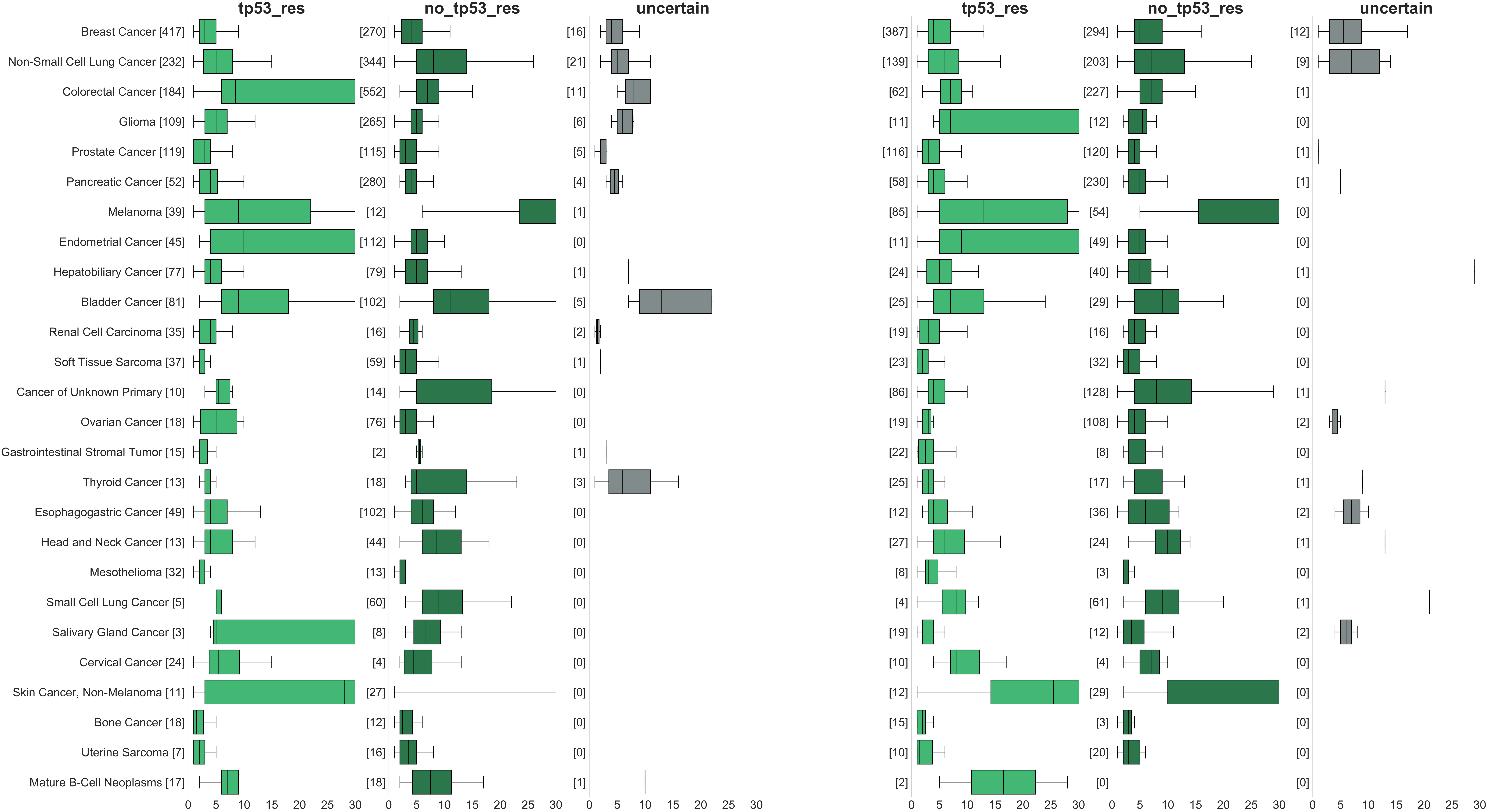




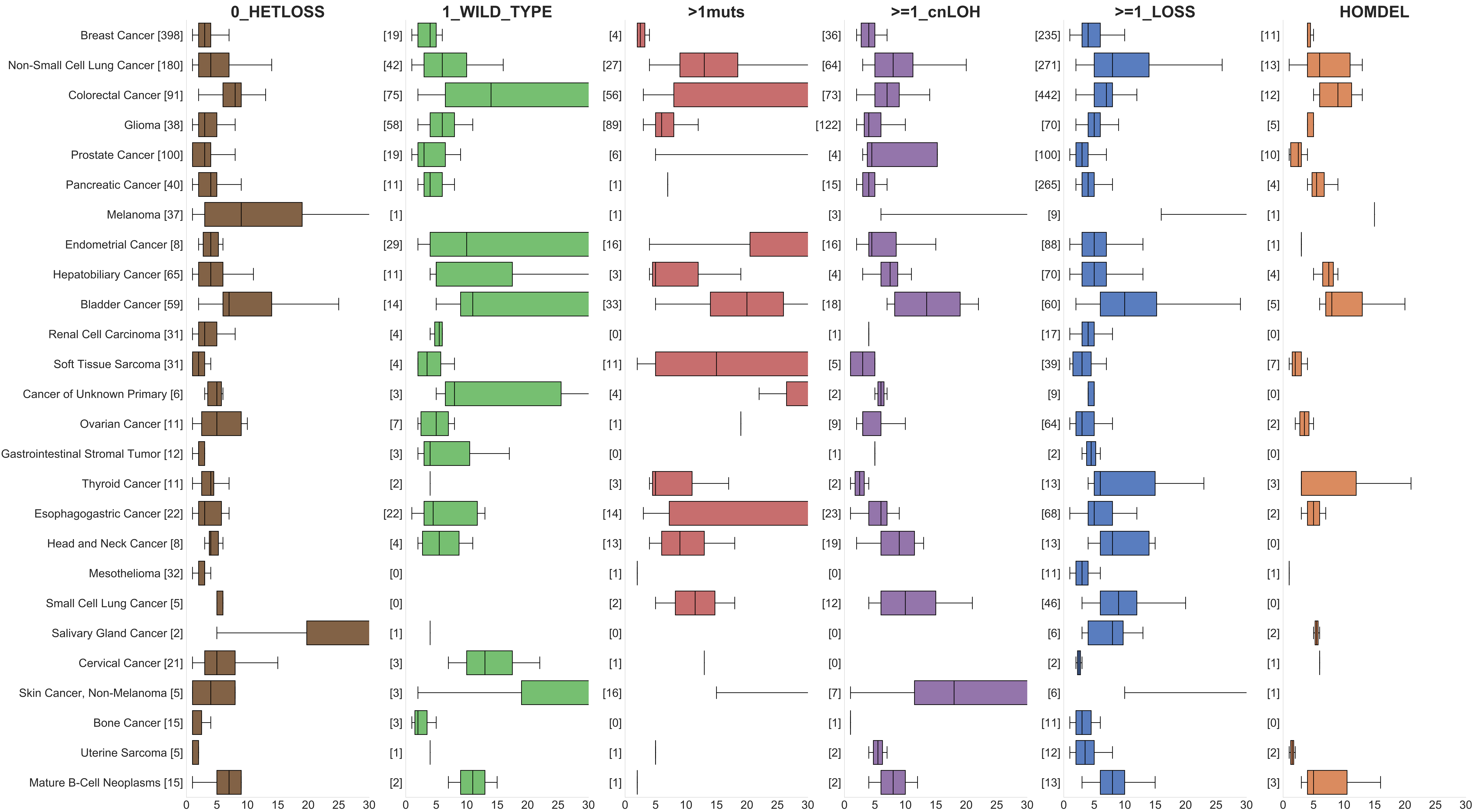
Mutation Count

Primary

Metastasis



Mutation Count - Primary Samples



Mutation Count - Metastatic Samples

