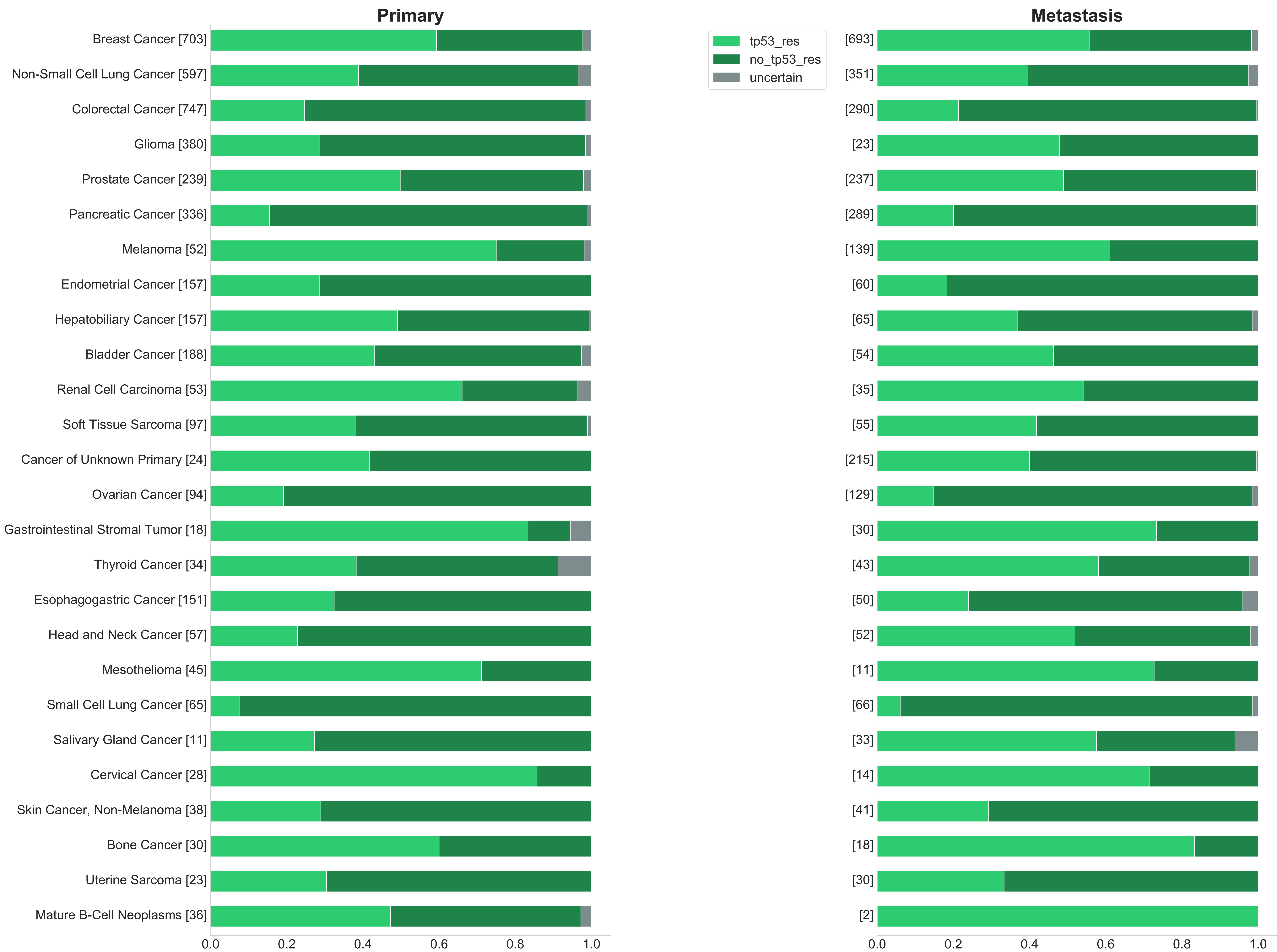
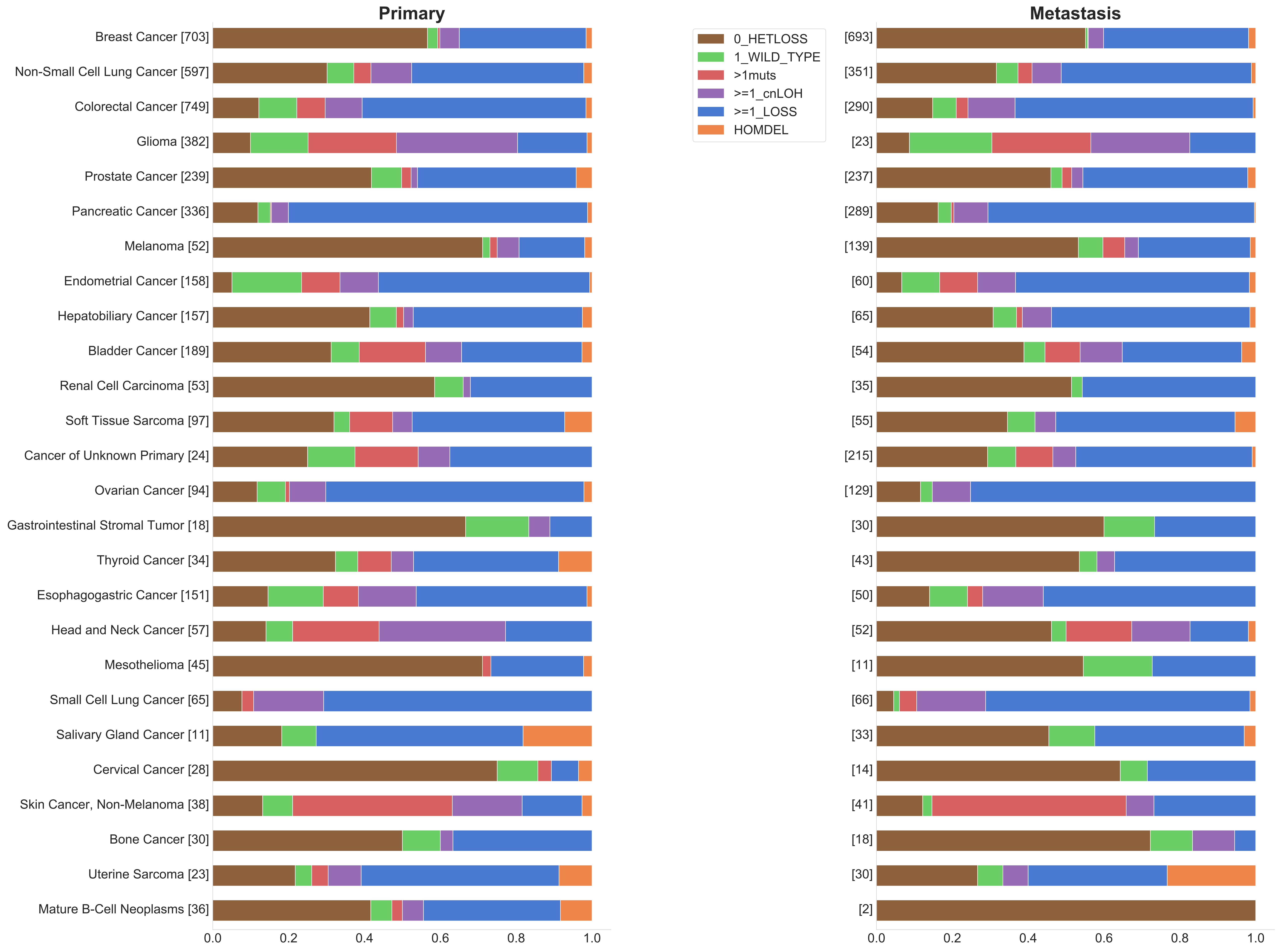


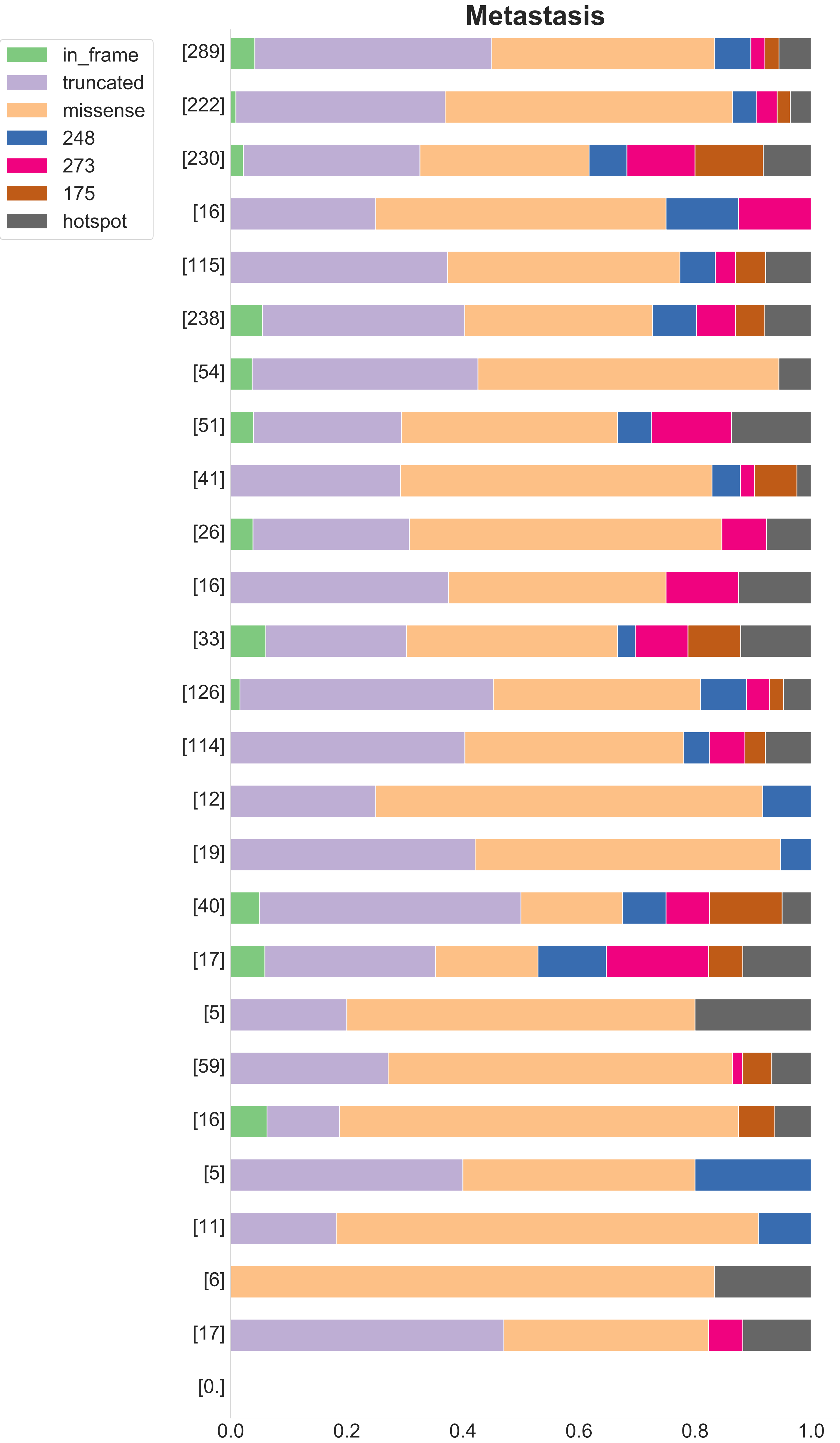
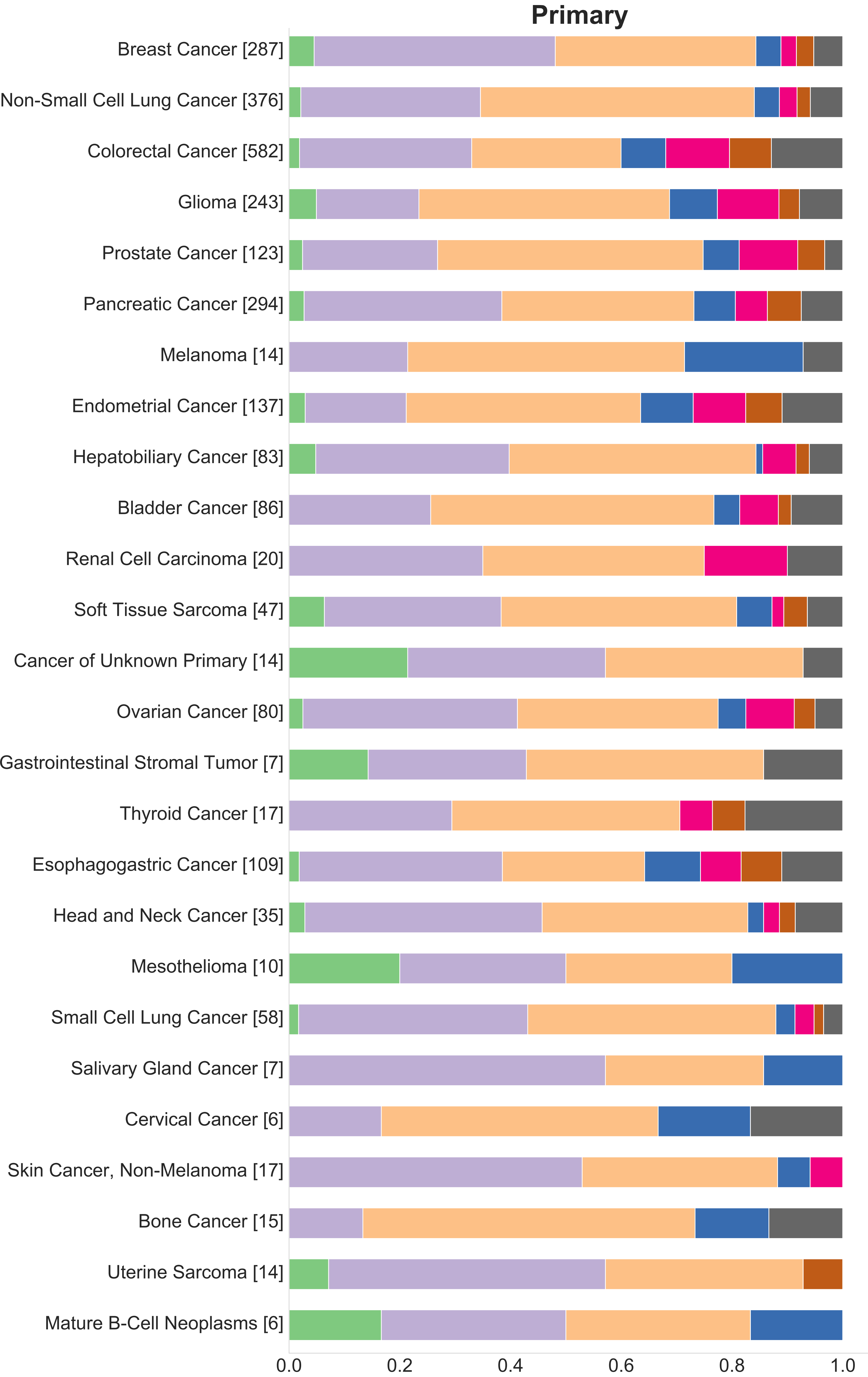
TP53 RESIDUAL SUBGROUP PROPORTION ACROSS CANCERS



TP53 SUBGROUP PROPORTION ACROSS CANCERS



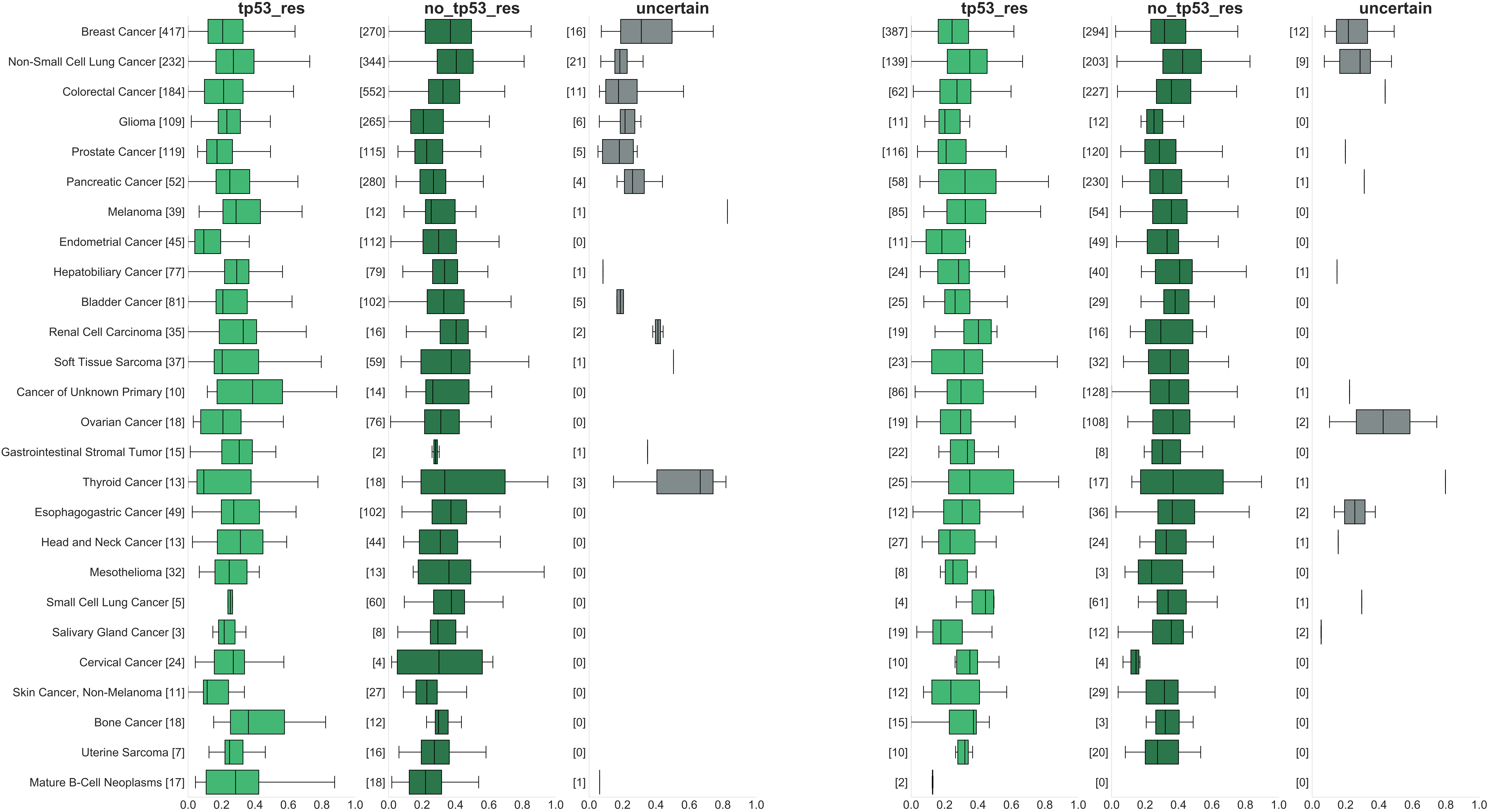
TP53 HOTSPOT DISTRIBUTION ACROSS CANCERS



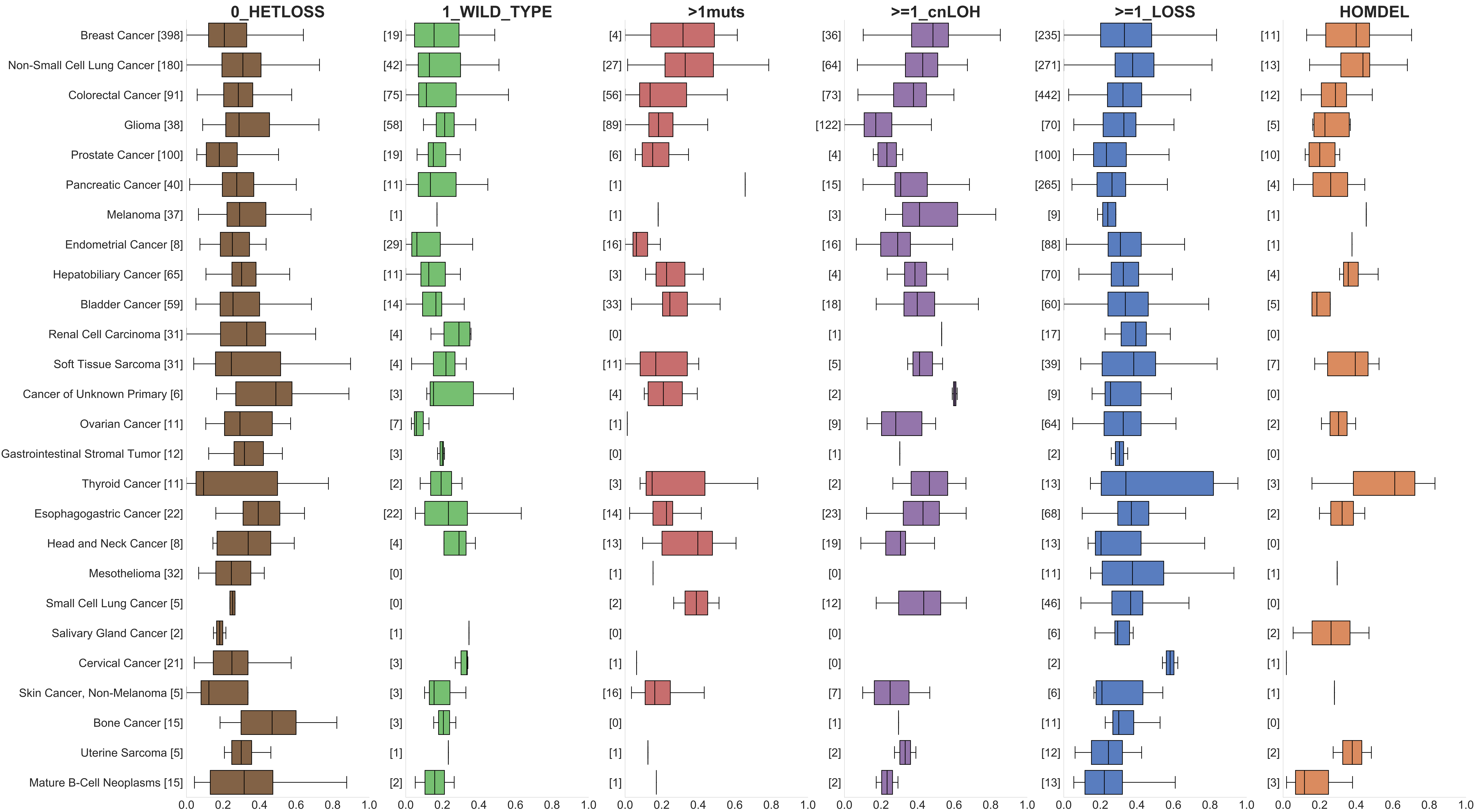
Fraction of genome altered

Primary

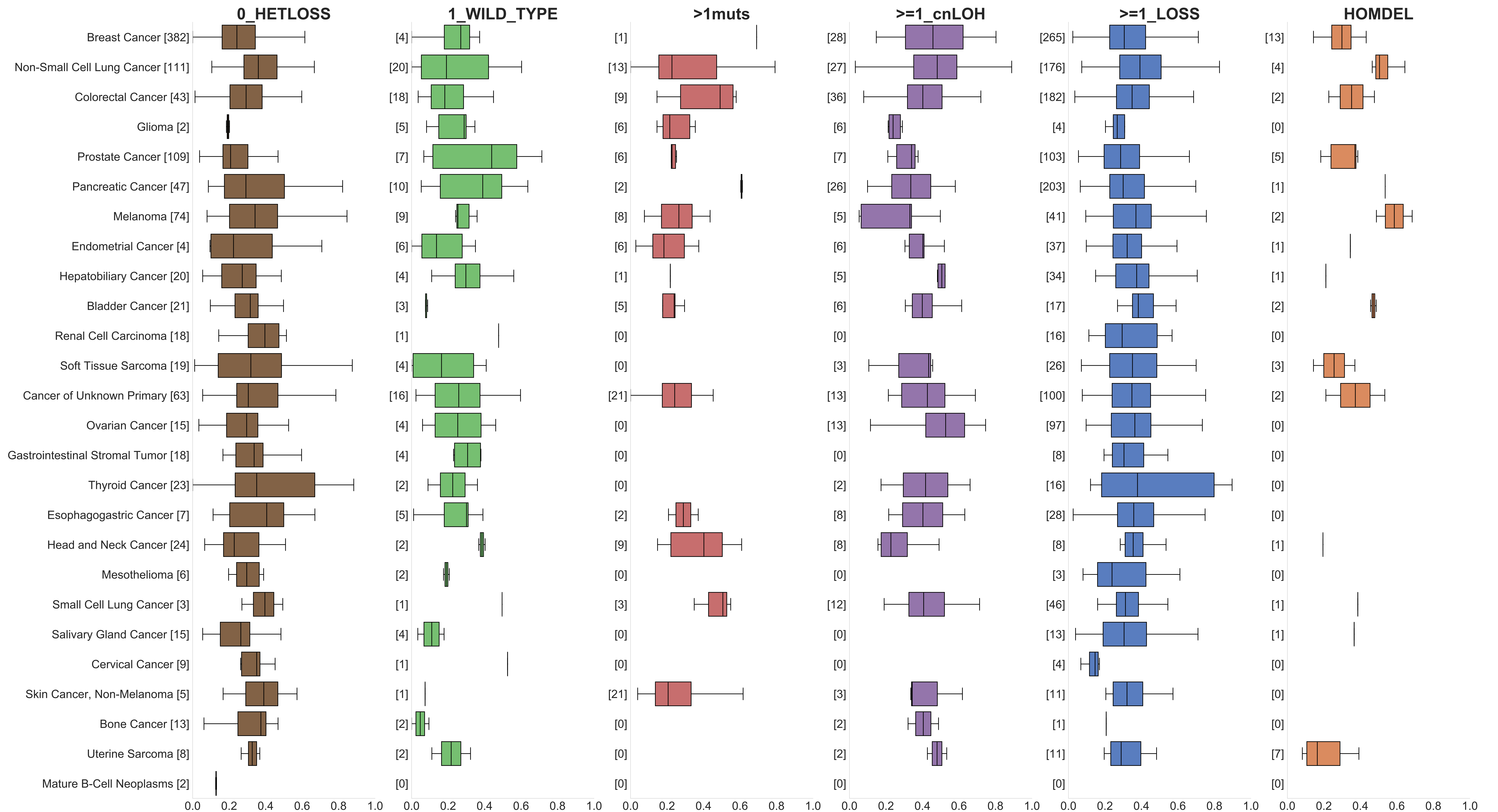
Metastasis



Fraction of genome altered - Primary Samples



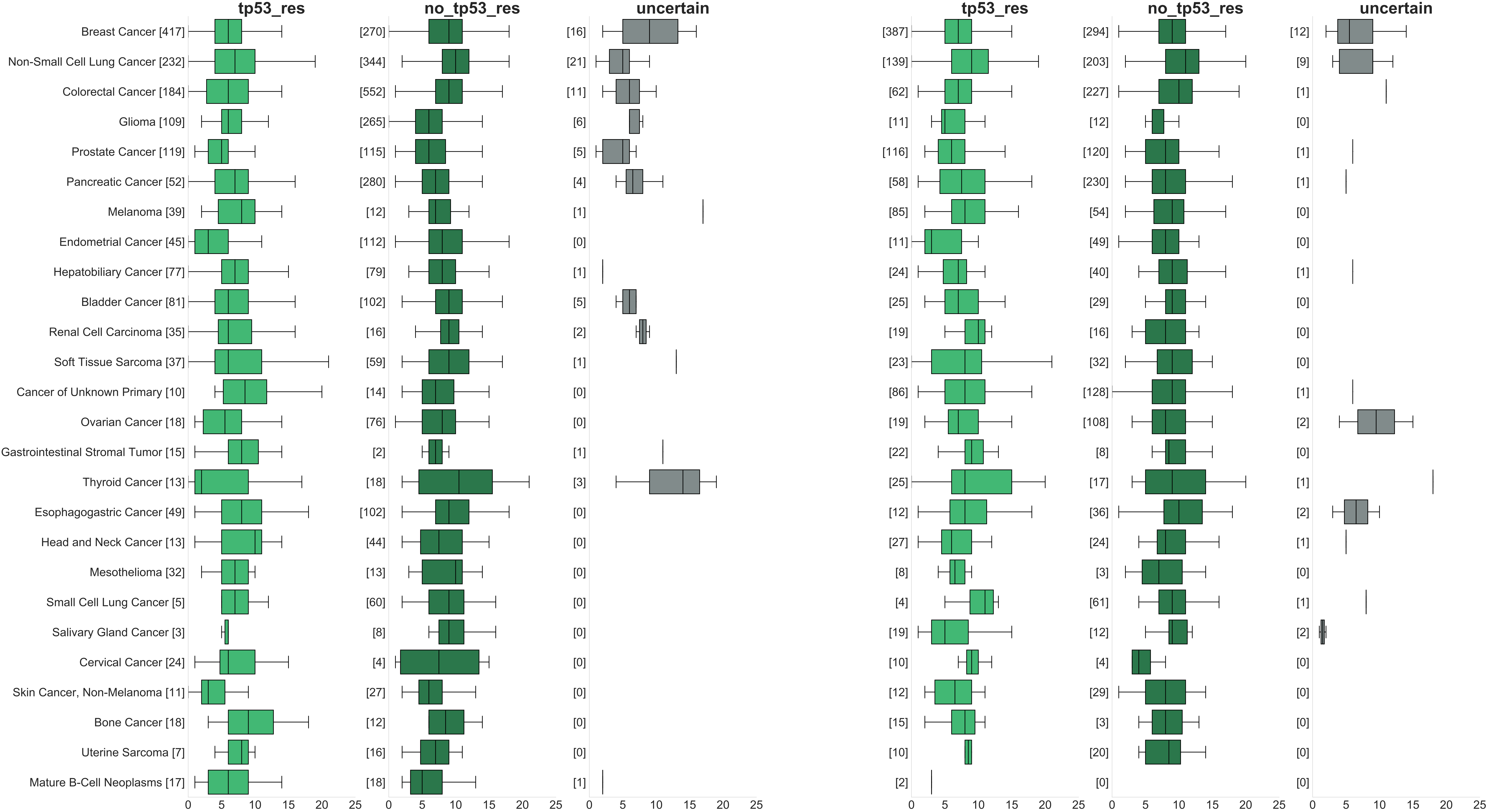
Fraction of genome altered - Metastatic Samples



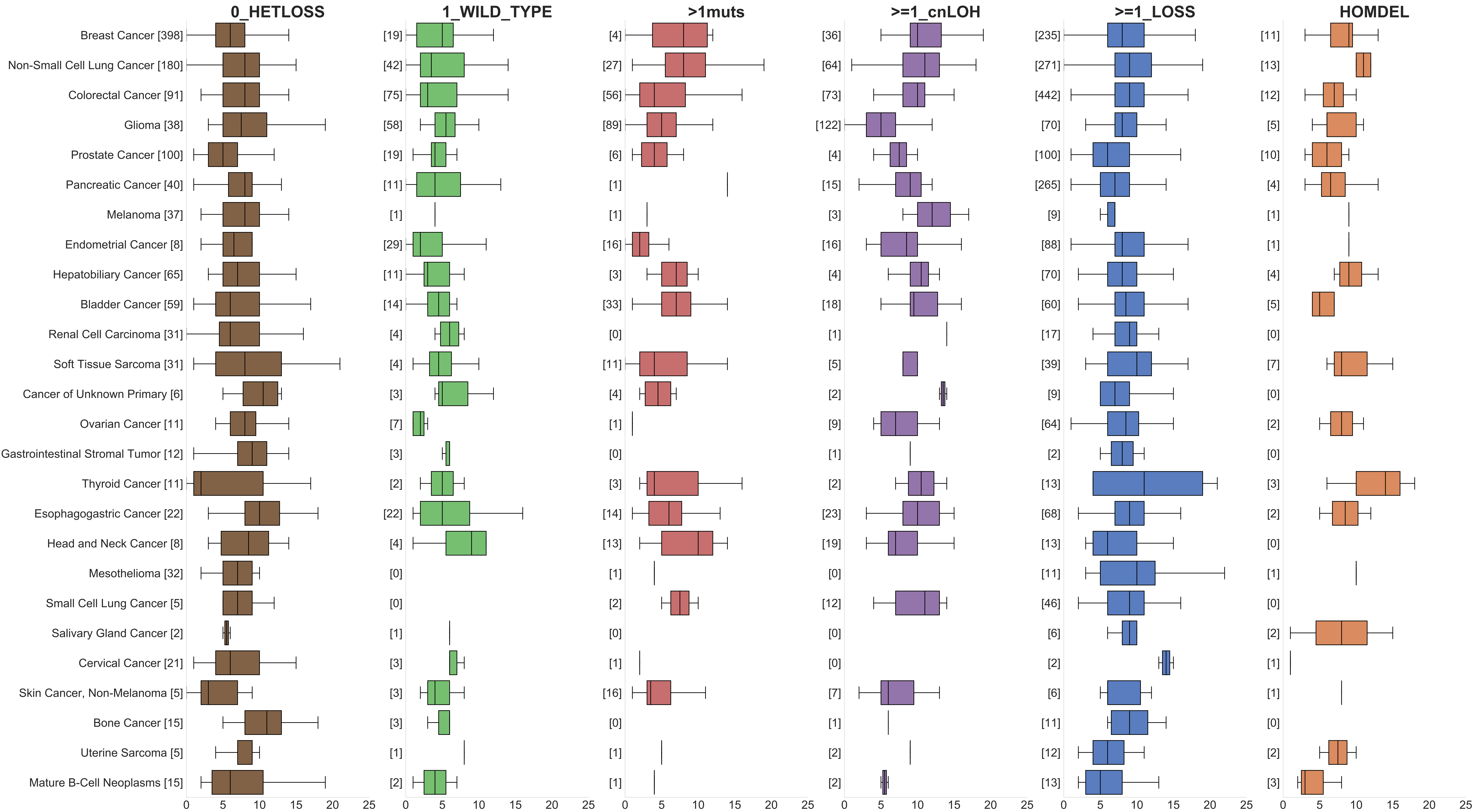
Number of aberrant chromosomes

Primary

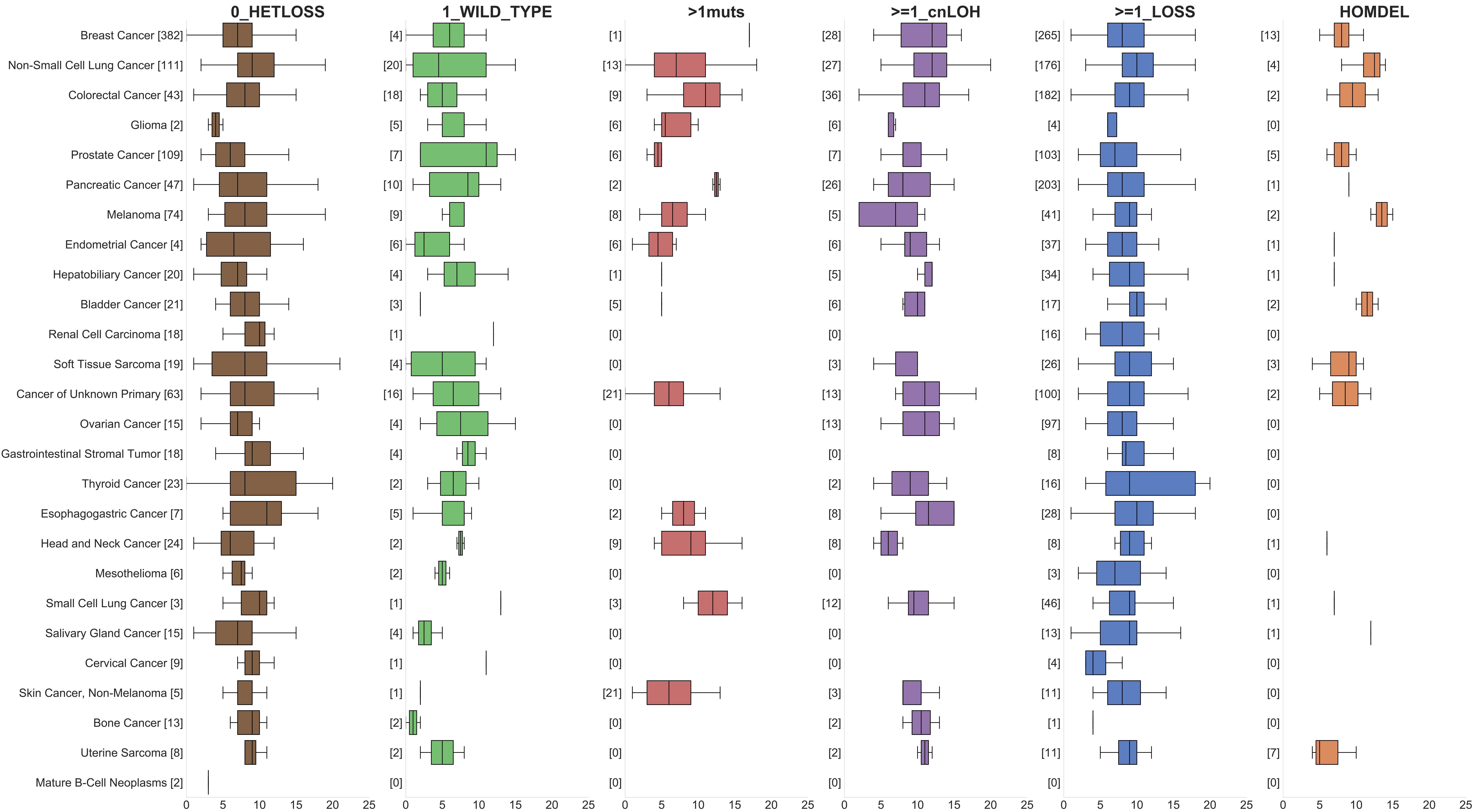
Metastasis



Number of aberrant chromosomes - Primary Samples



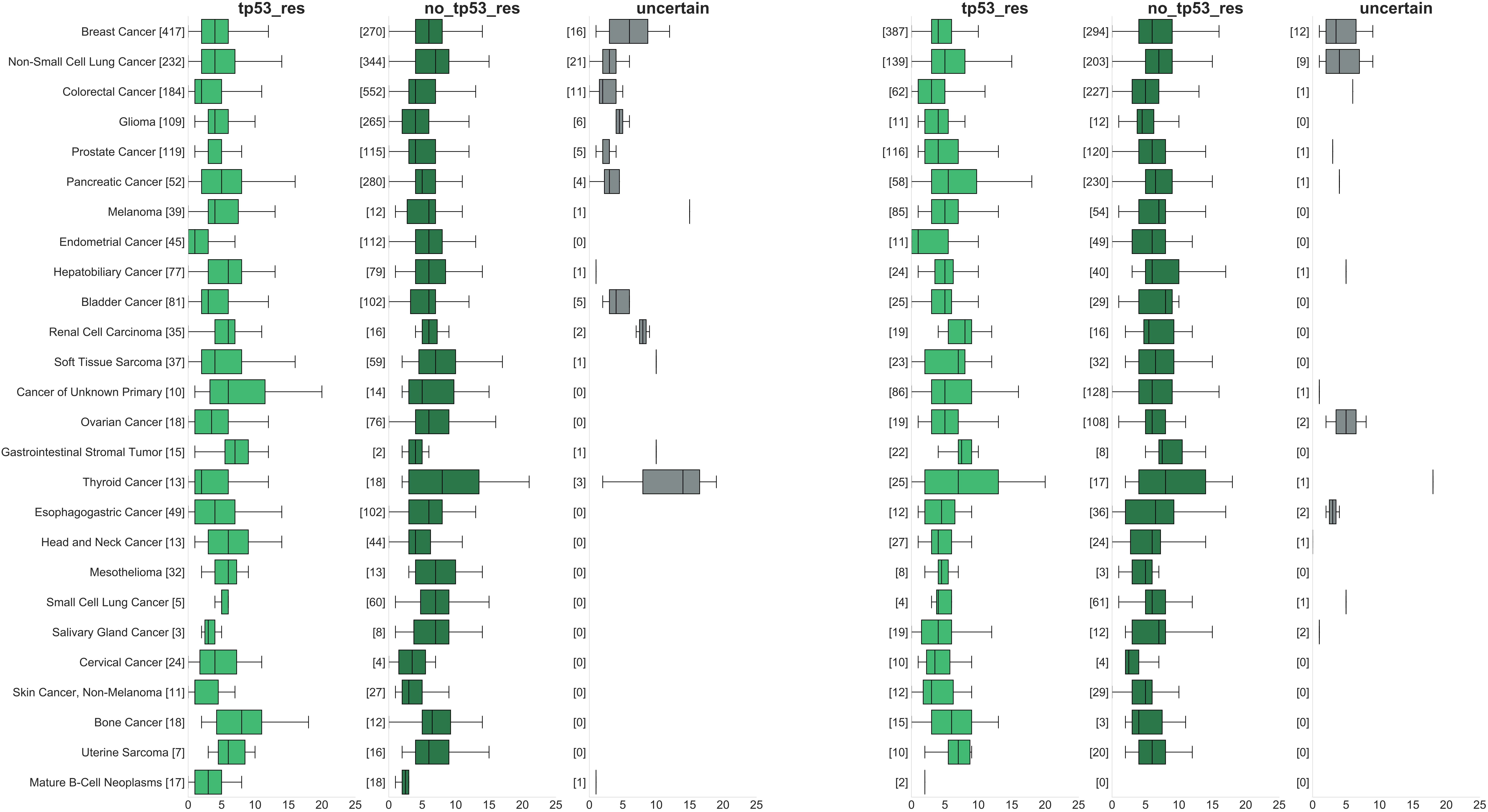
Number of aberrant chromosomes - Metastatic Samples



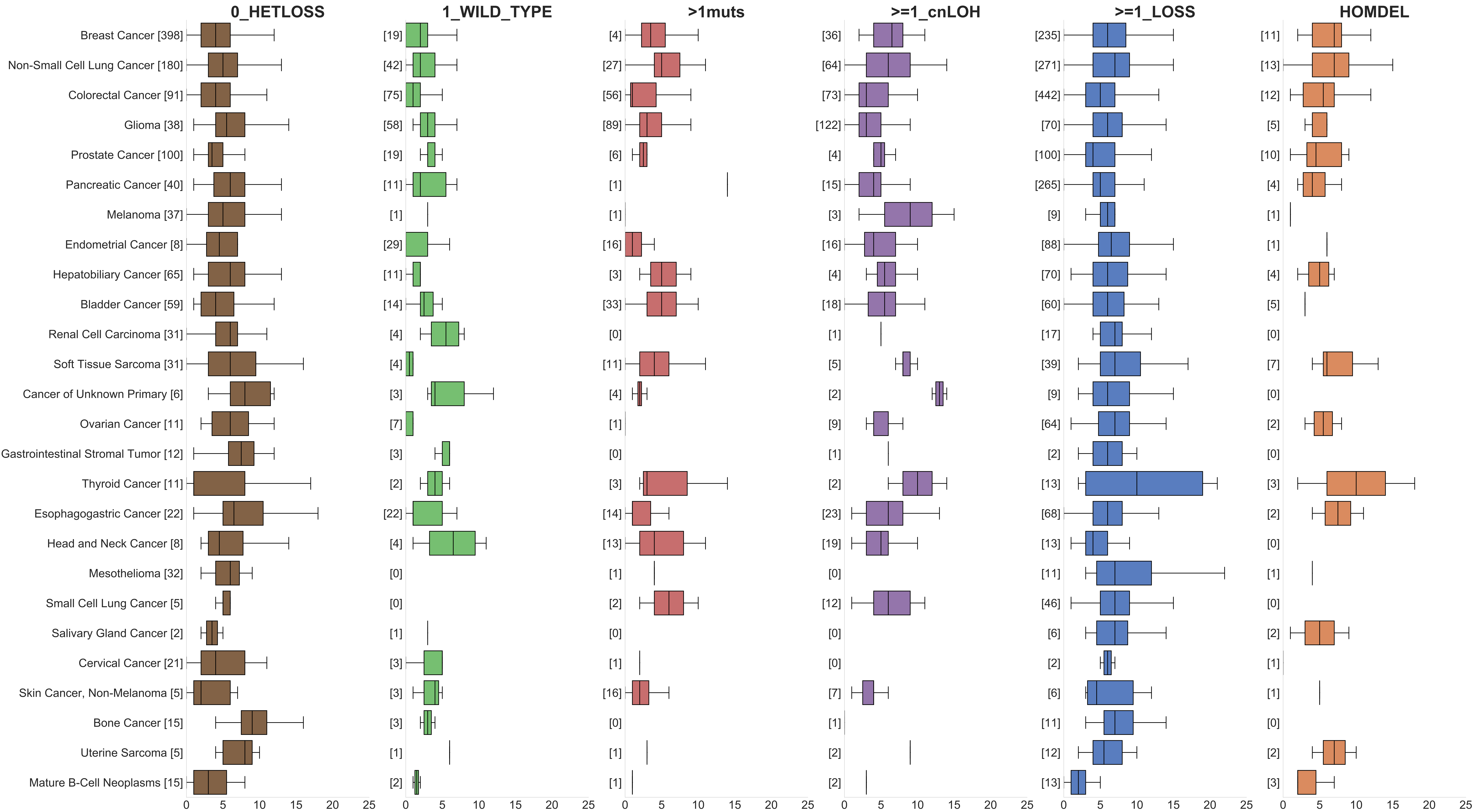
Number of chromosomes affected by a LOSS

Primary

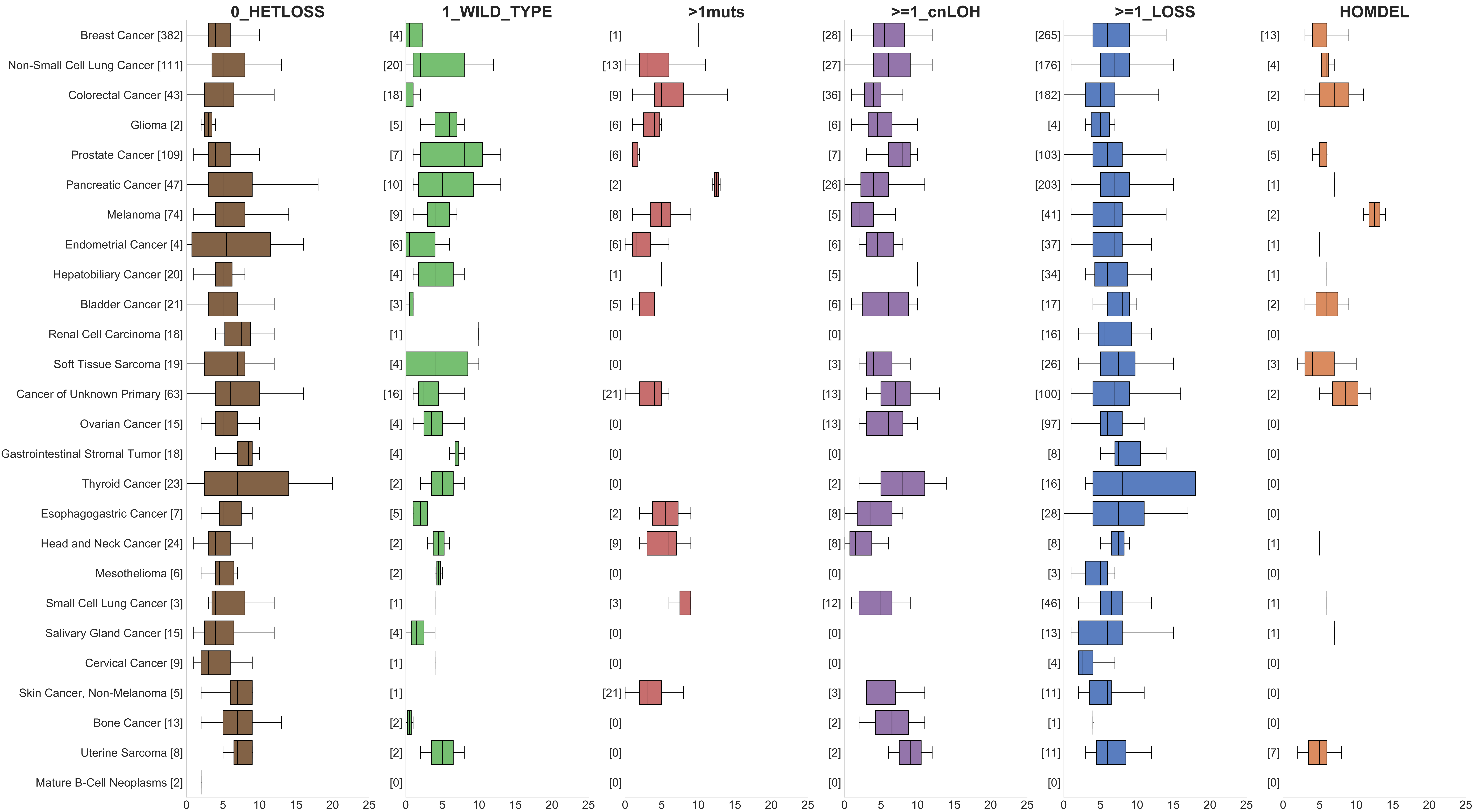
Metastasis



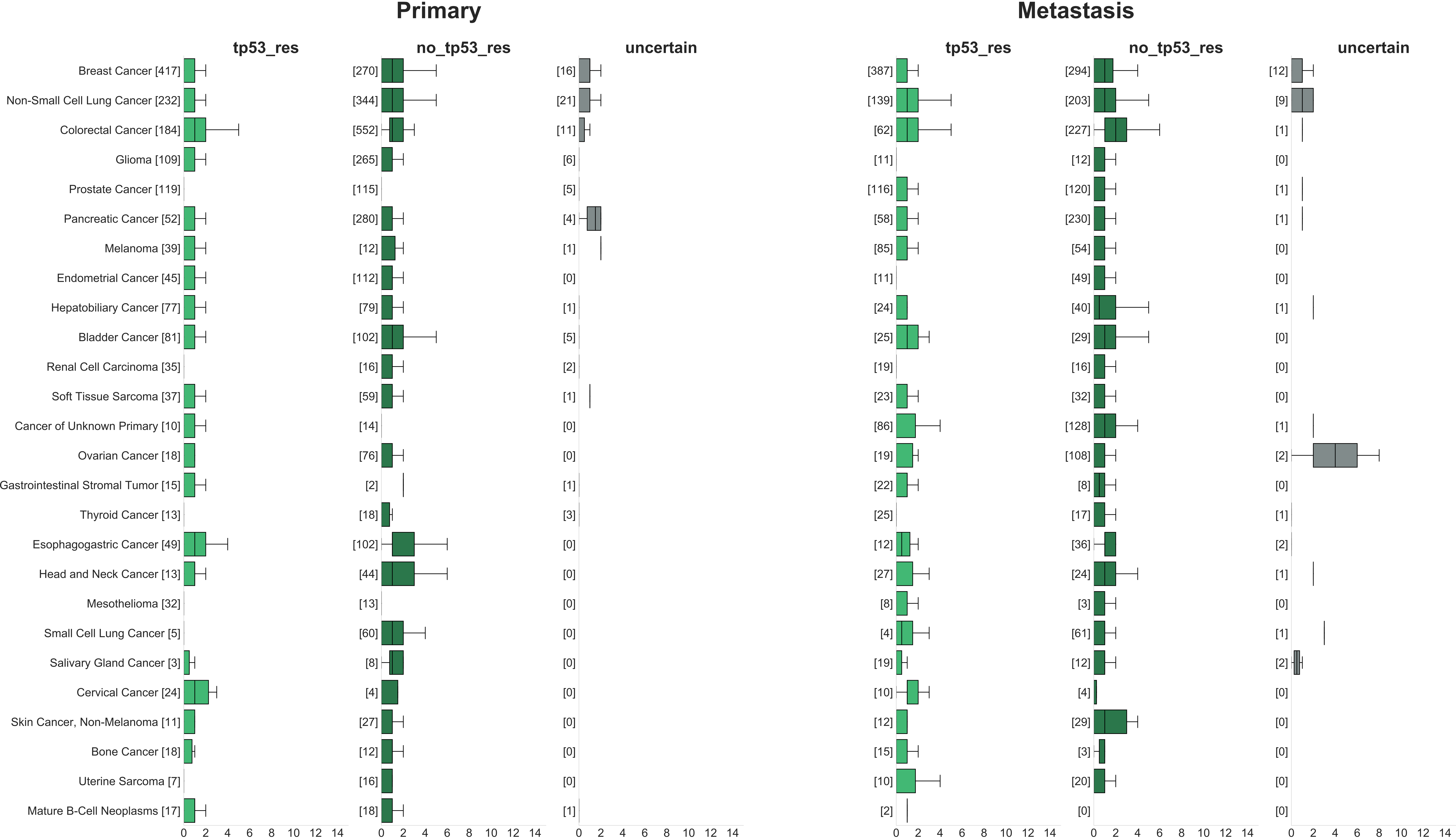
Number of chromosomes affected by a LOSS - Primary Samples



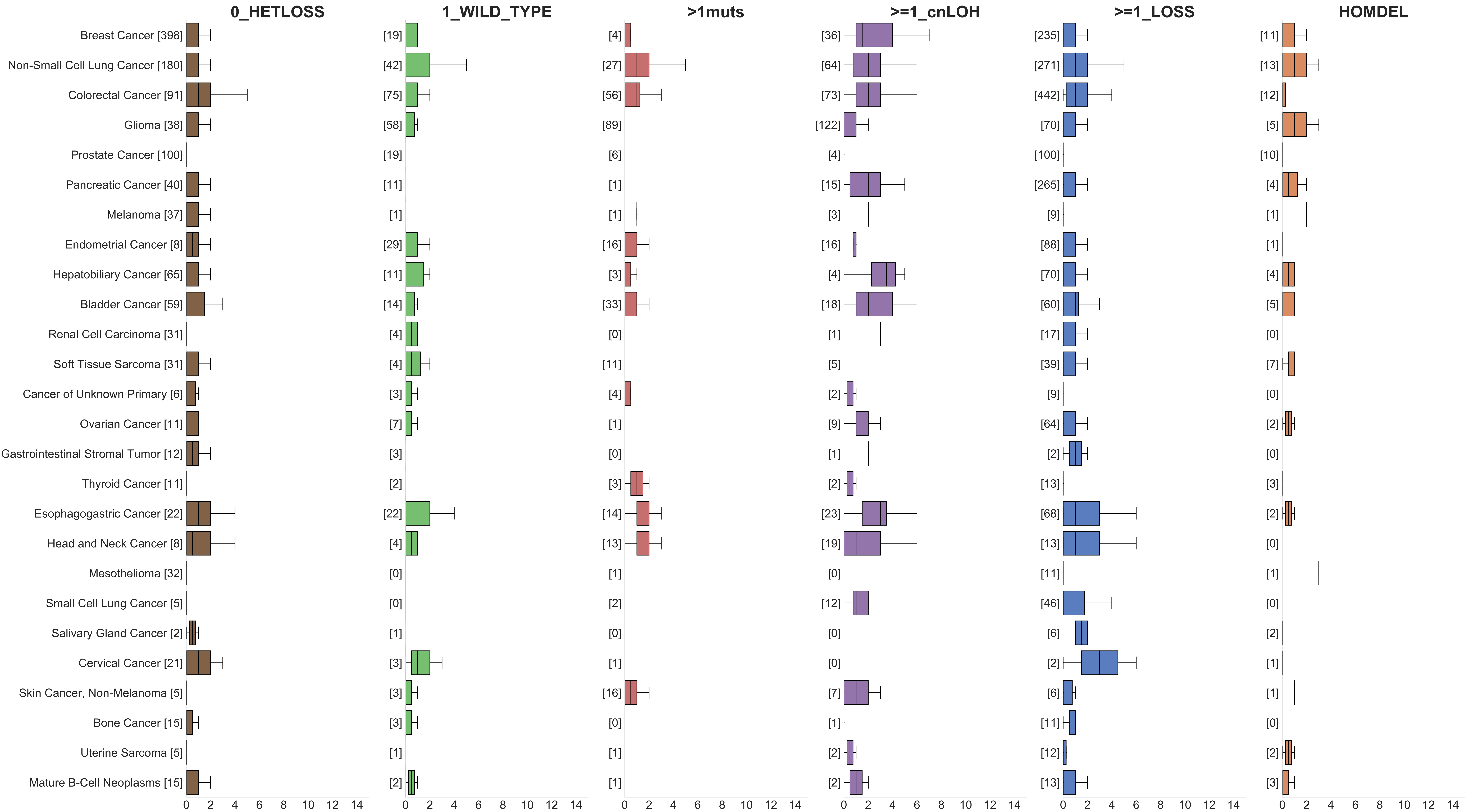
Number of chromosomes affected by a LOSS - Metastatic Samples



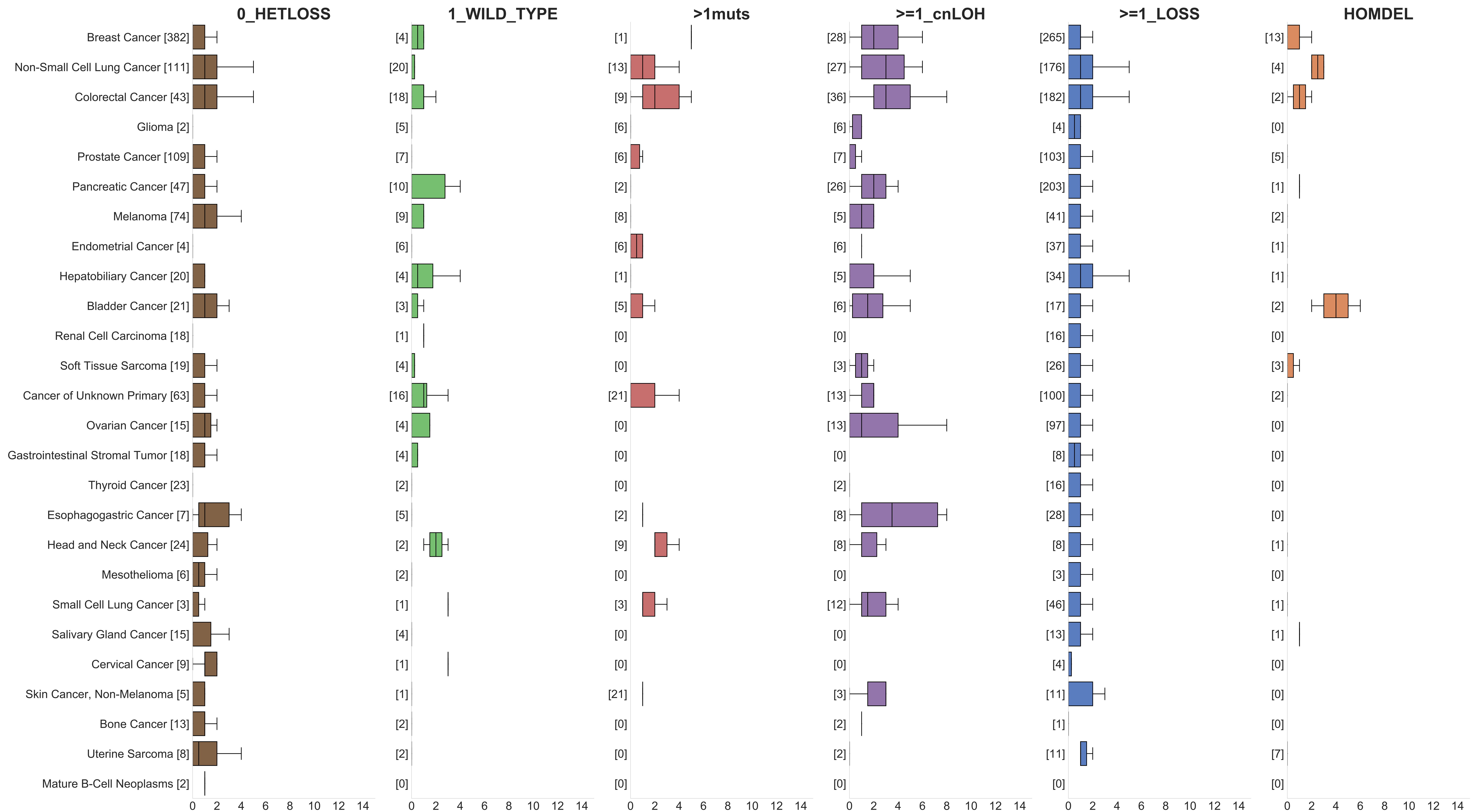
Number of chromosomes affected by a cnLOH



Number of chromosomes affected by a cnLOH - Primary Samples



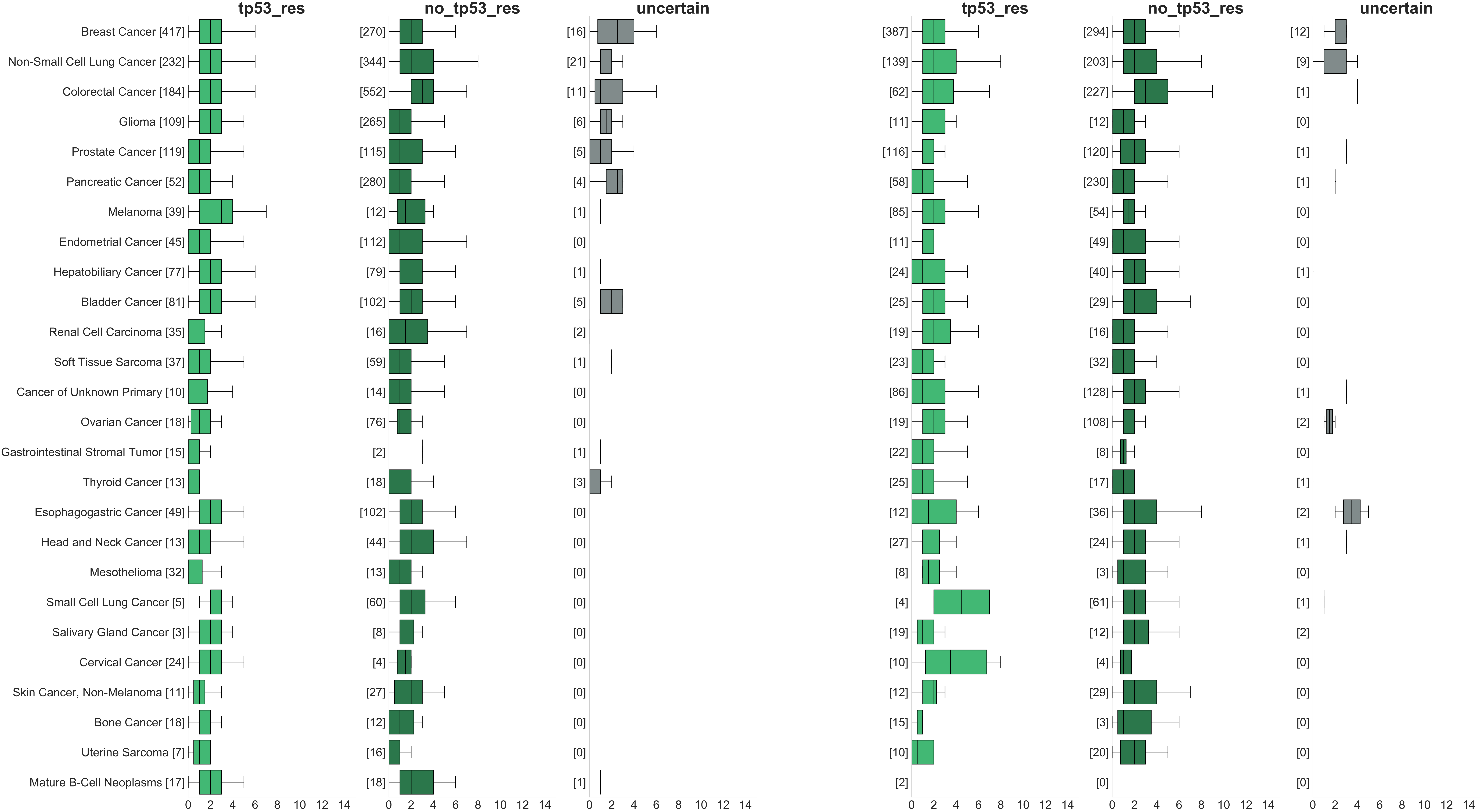
Number of chromosomes affected by a cnLOH - Metastatic Samples



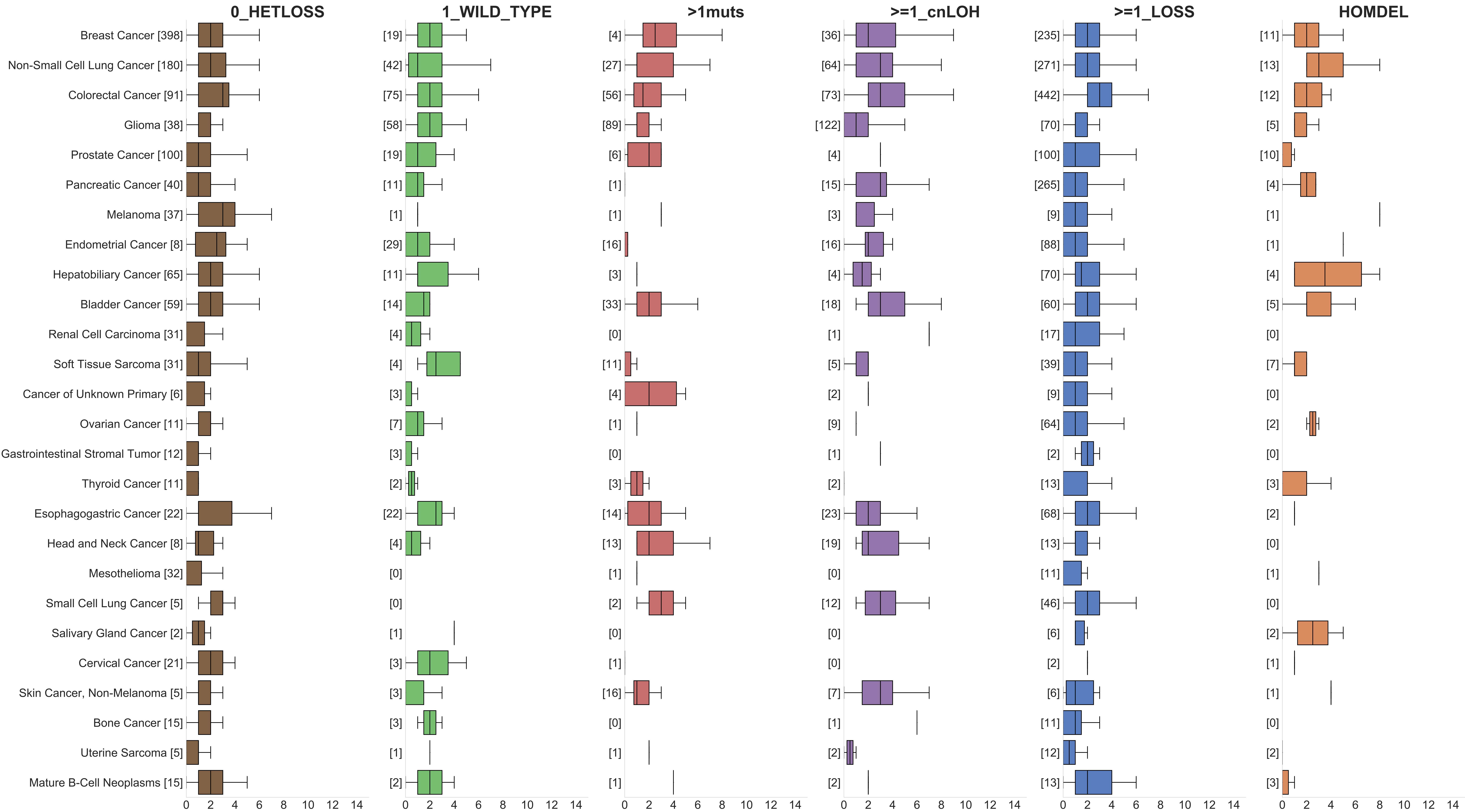
Number of chromosomes affected by a GAIN

Primary

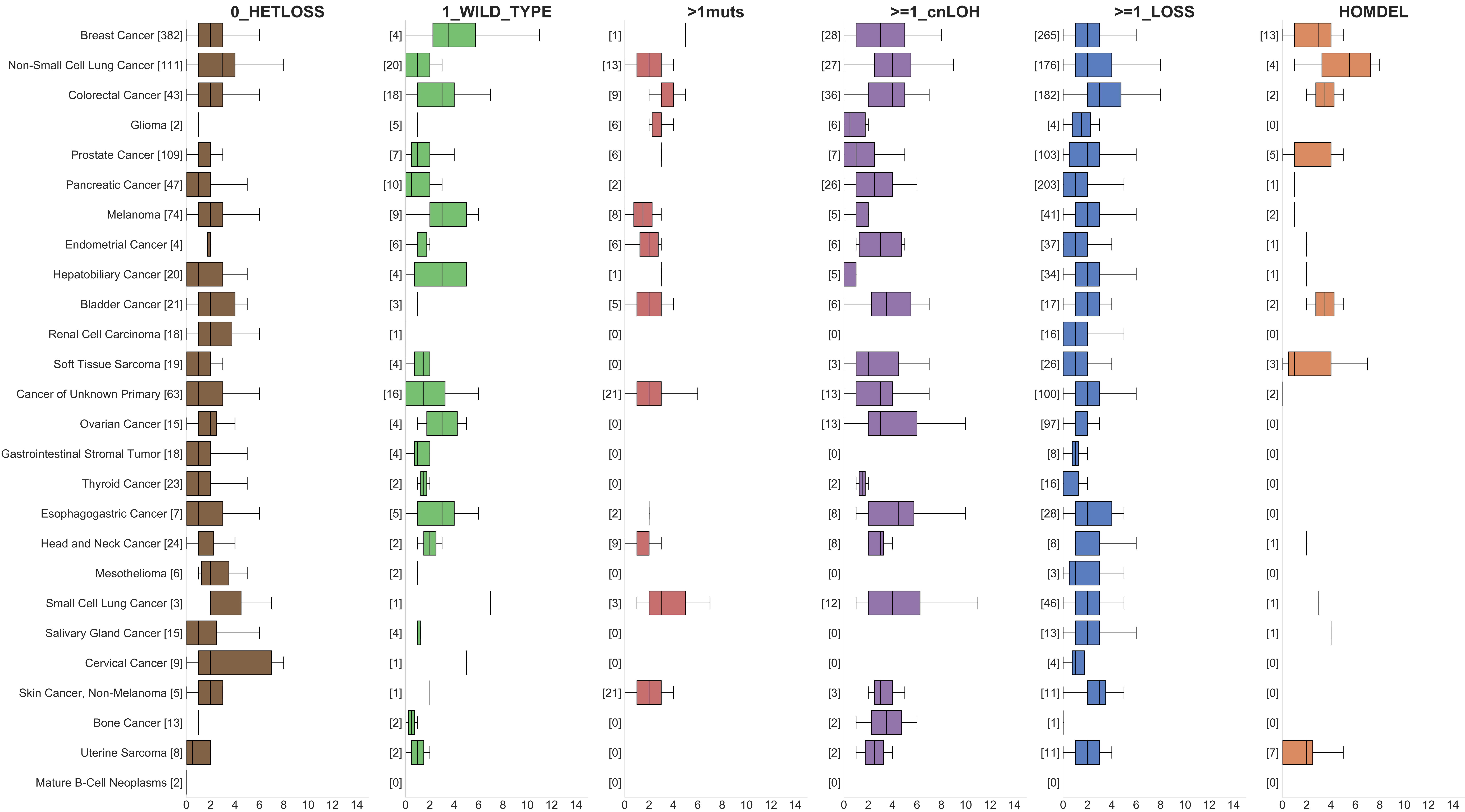
Metastasis



Number of chromosomes affected by a GAIN - Primary Samples



Number of chromosomes affected by a GAIN - Metastatic Samples



Proportion of WGD



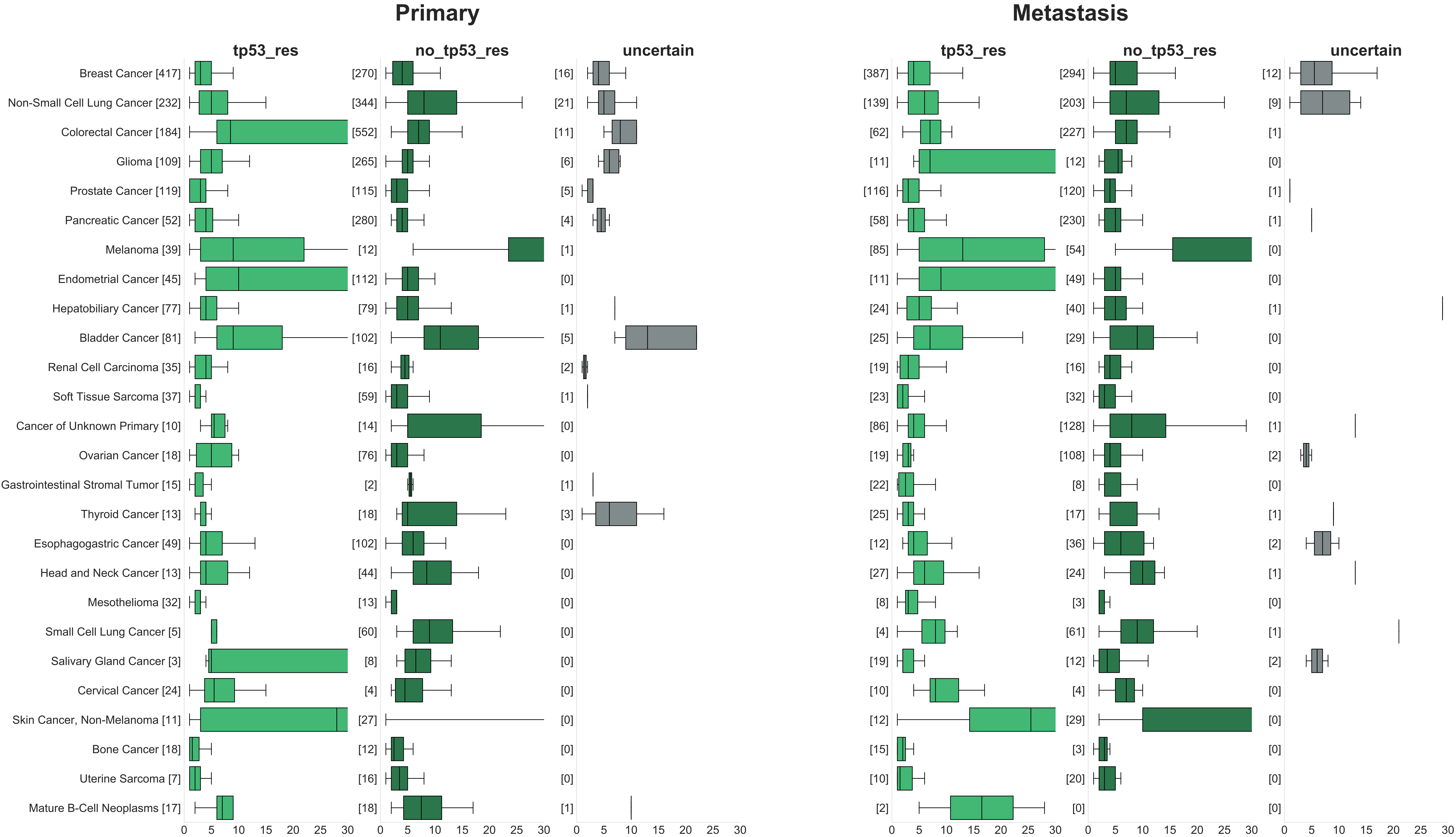
Proportion of WGD - Primary Samples



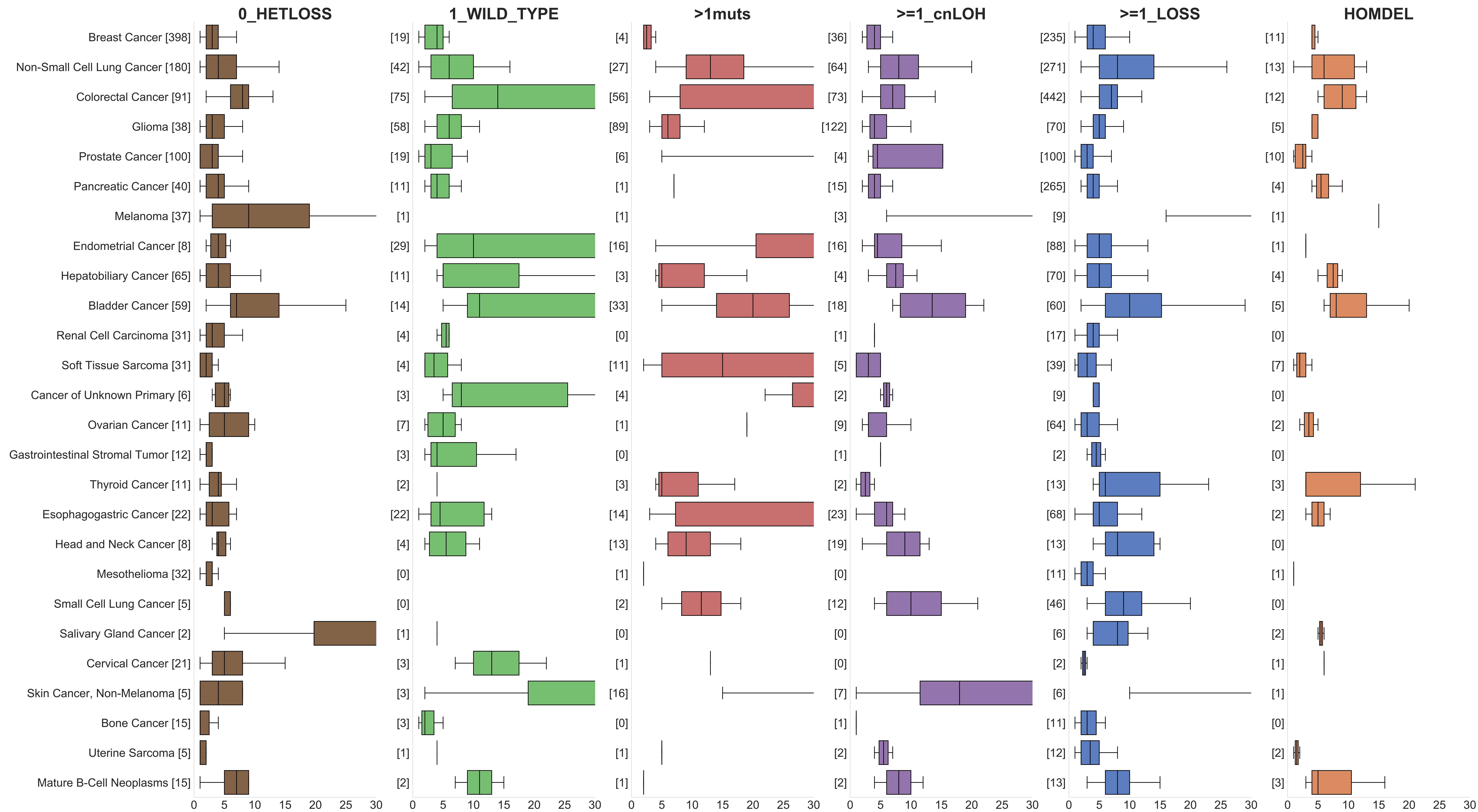
Proportion of WGD - Metastatic Samples



Mutation Count



Mutation Count - Primary Samples



Mutation Count - Metastatic Samples

