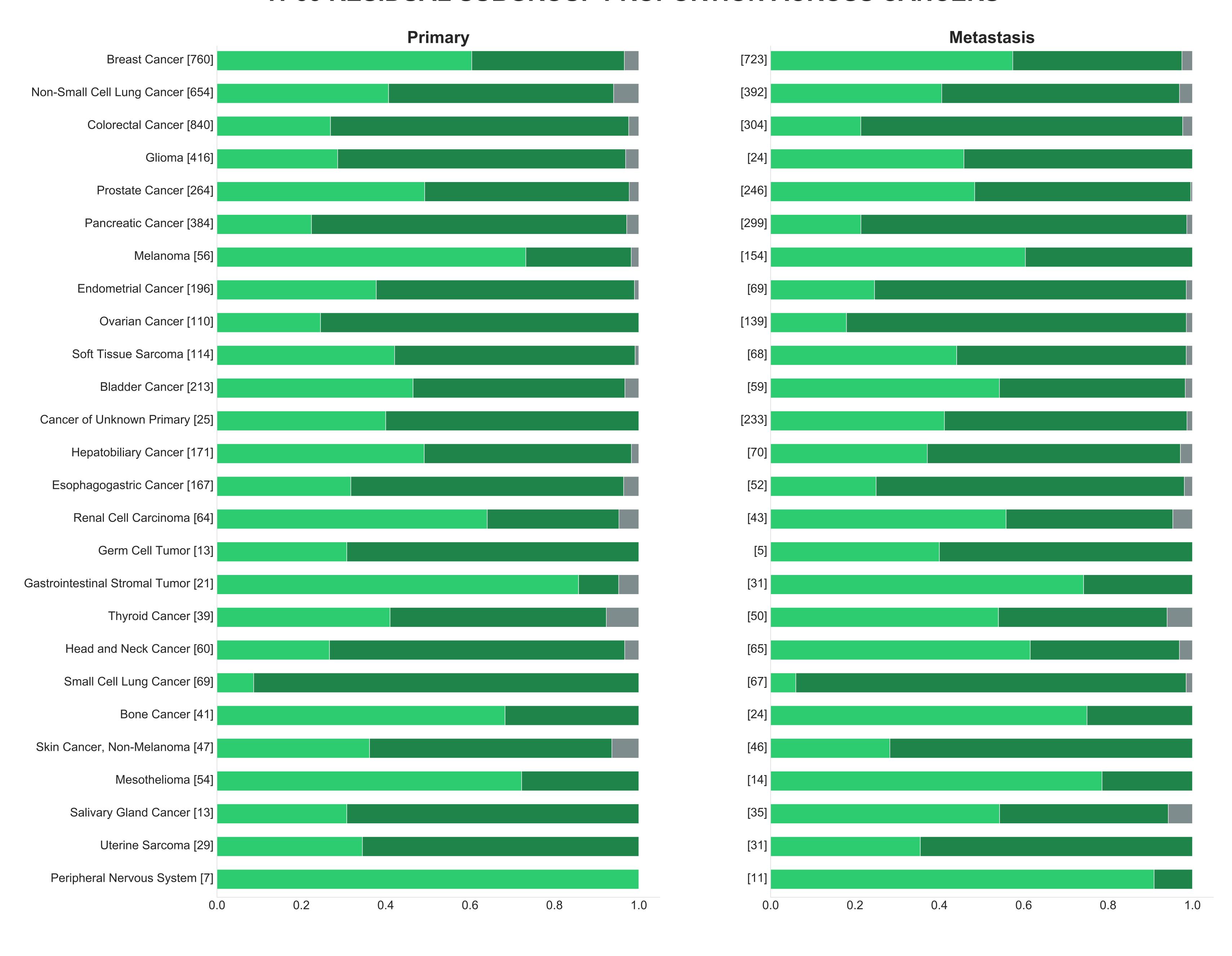
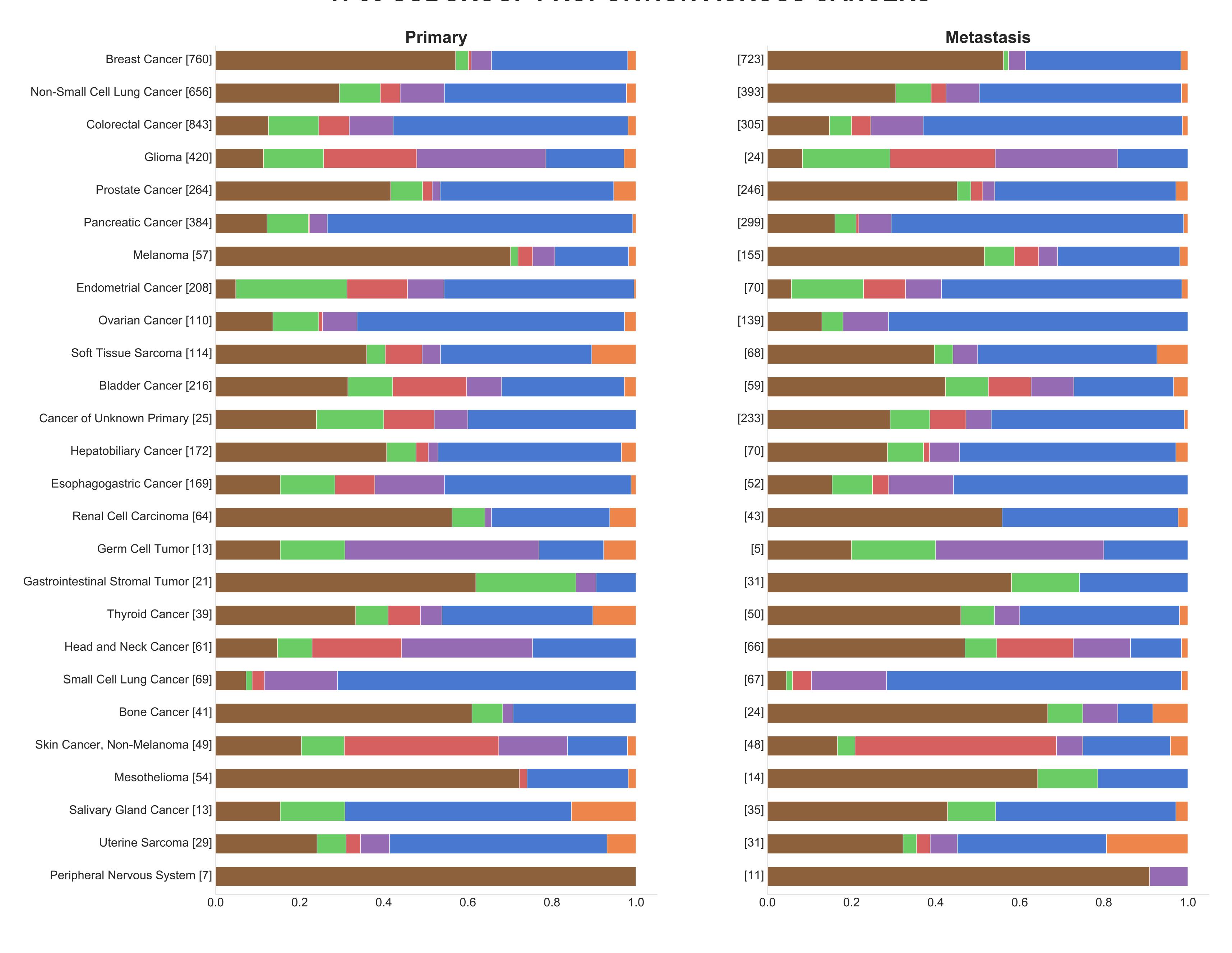
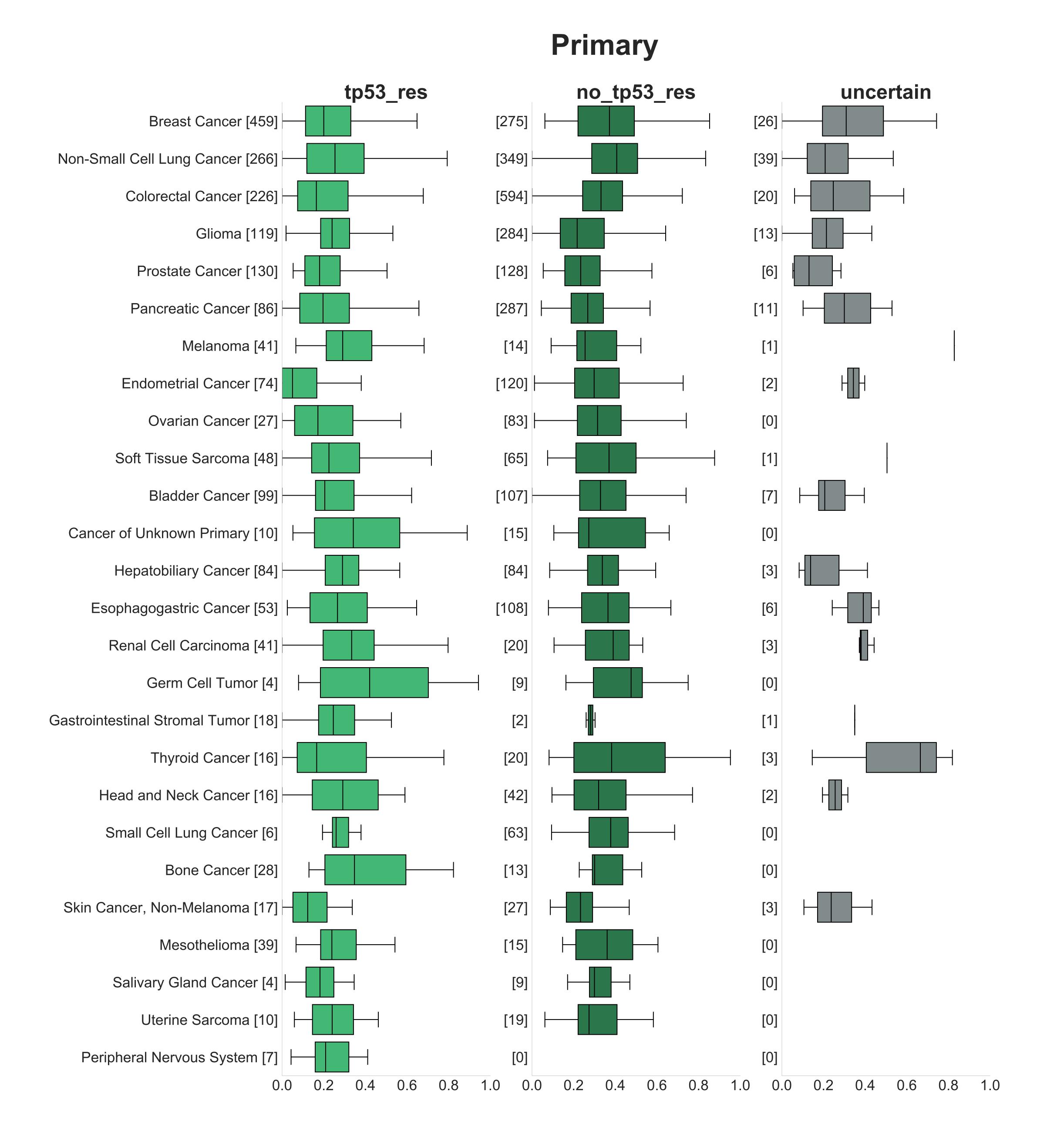
TP53 RESIDUAL SUBGROUP PROPORTION ACROSS CANCERS



TP53 SUBGROUP PROPORTION ACROSS CANCERS



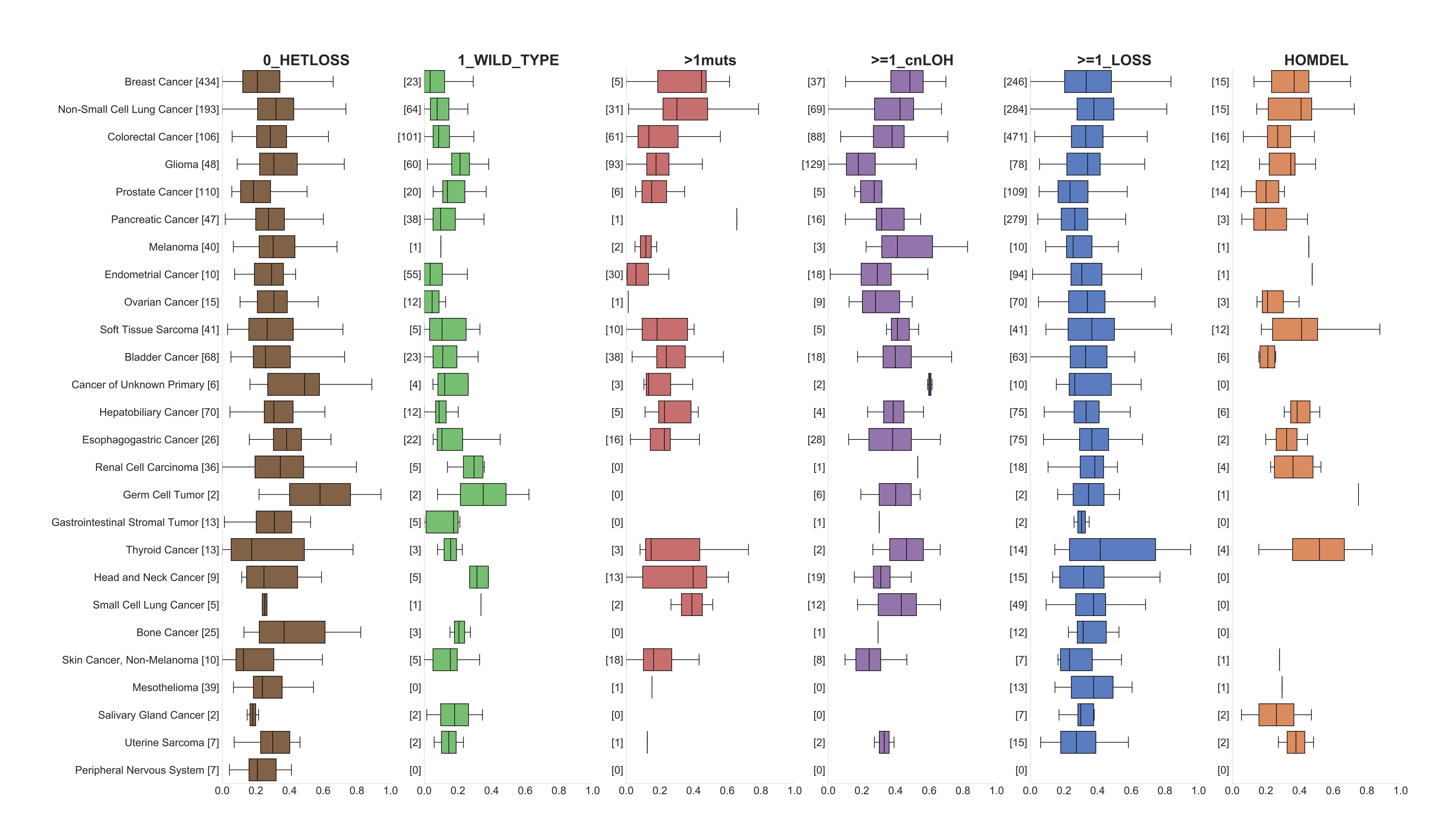
Fraction of genome altered



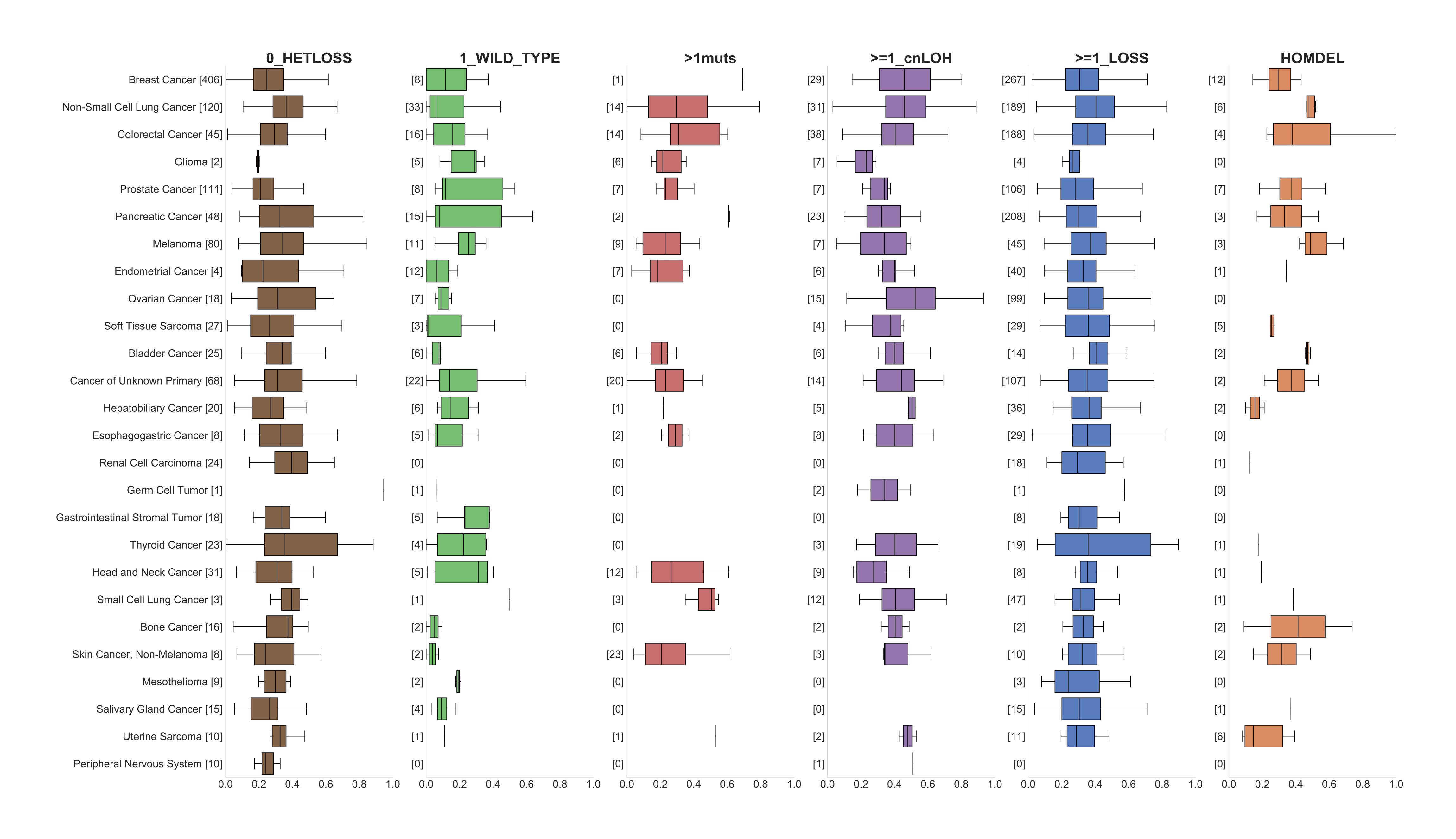
Metastasis



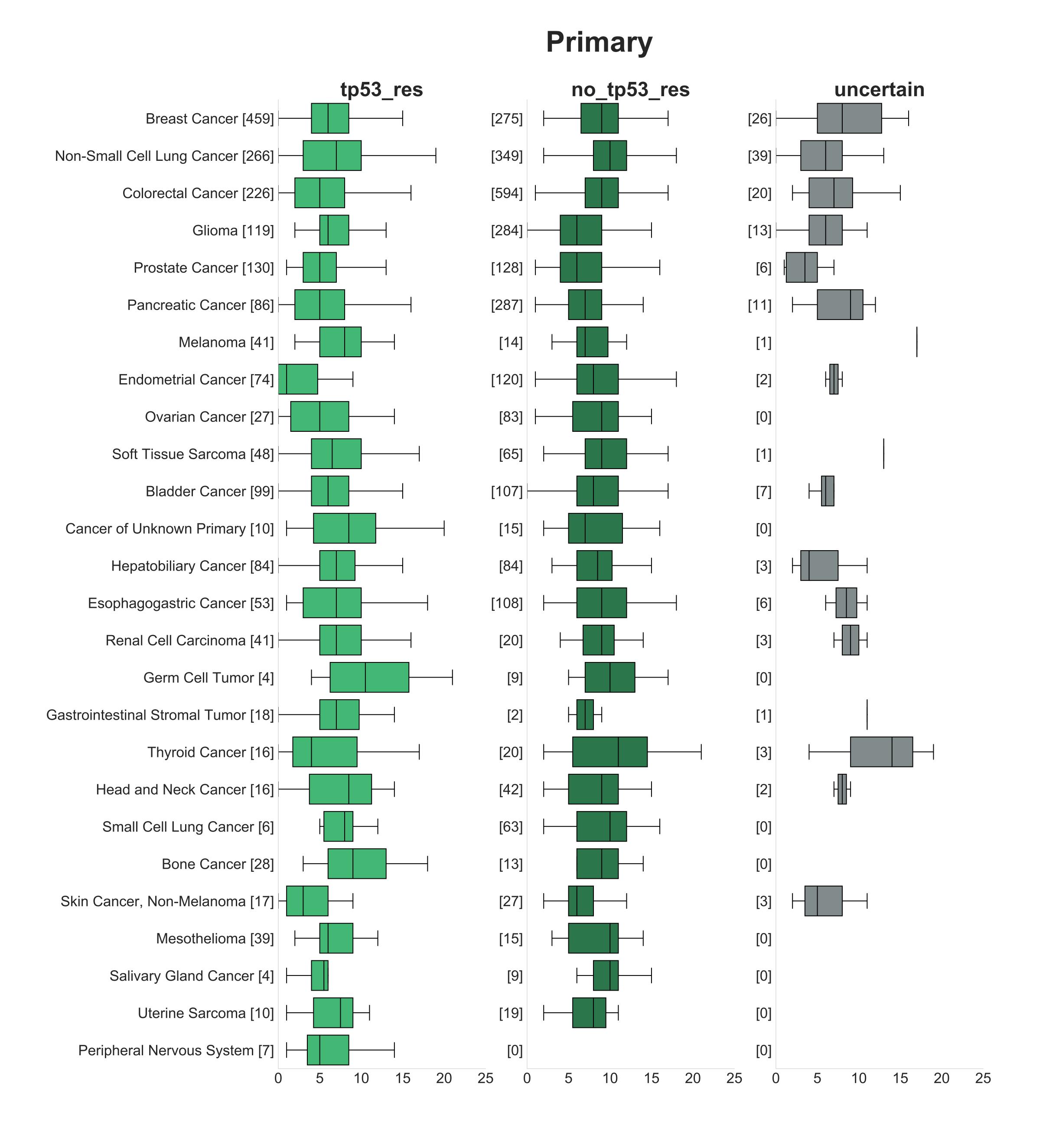
Fraction of genome altered - Primary Samples



Fraction of genome altered - Metastatic Samples



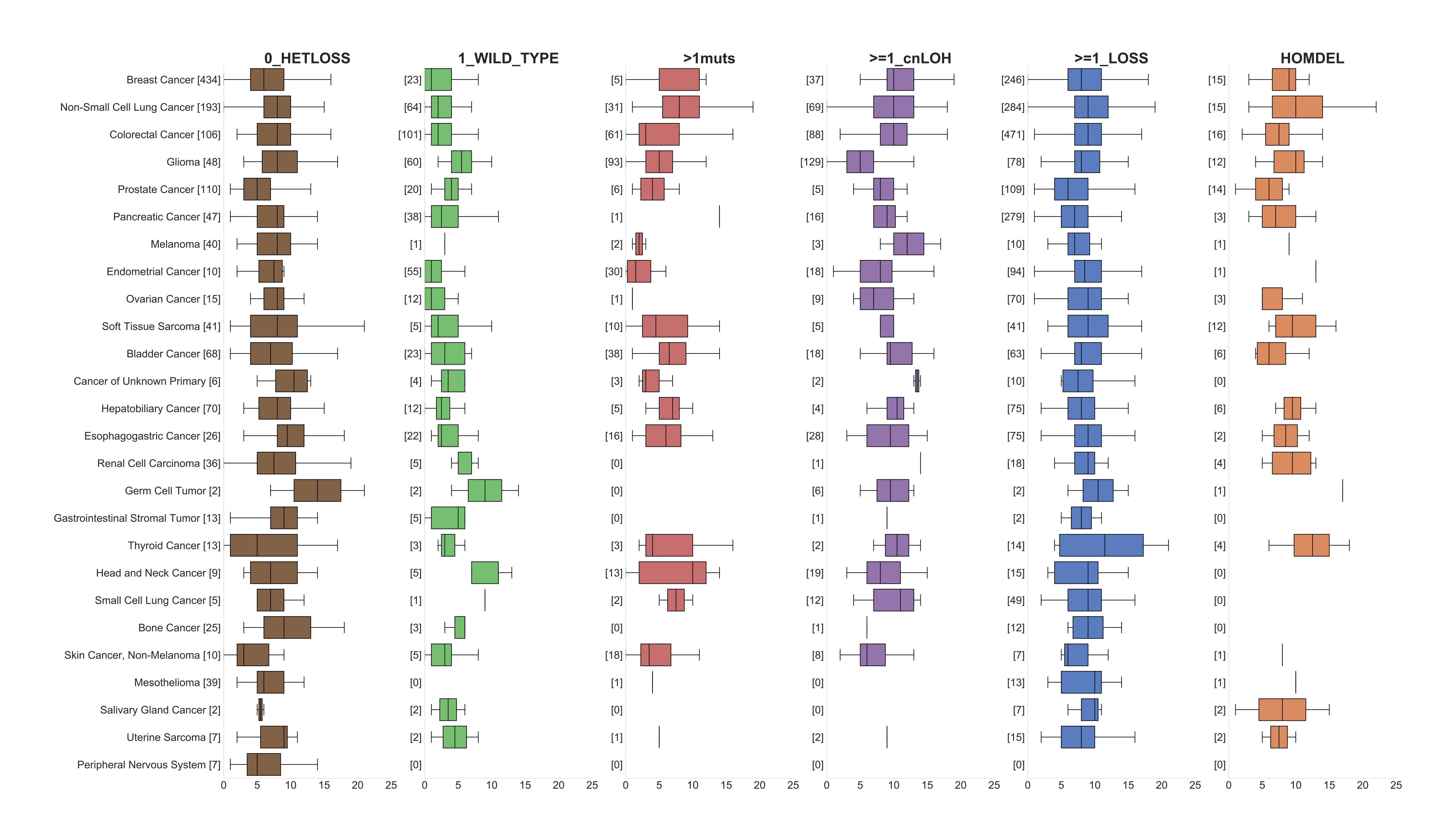
Number of aberrant chromosomes



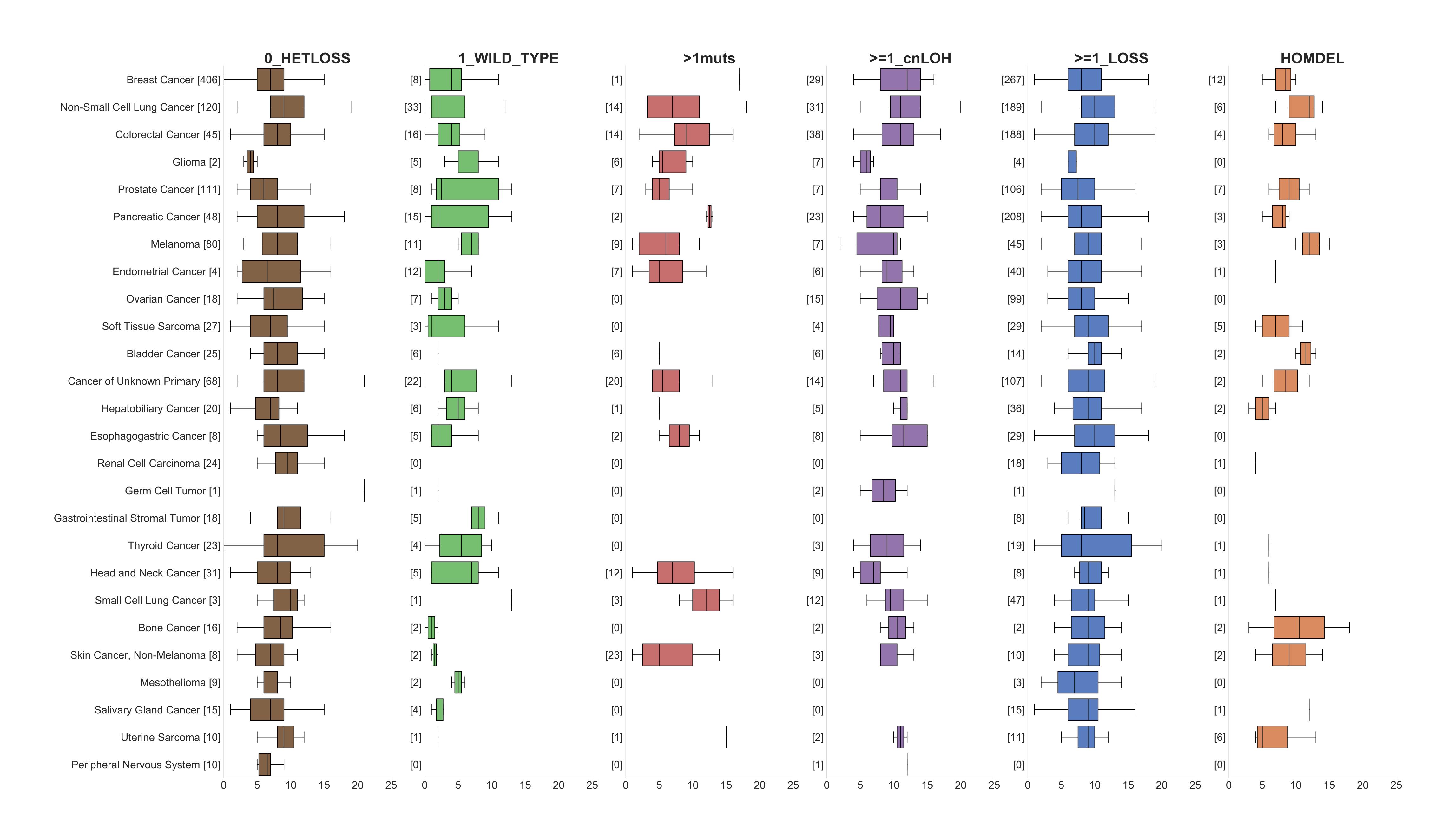
Metastasis



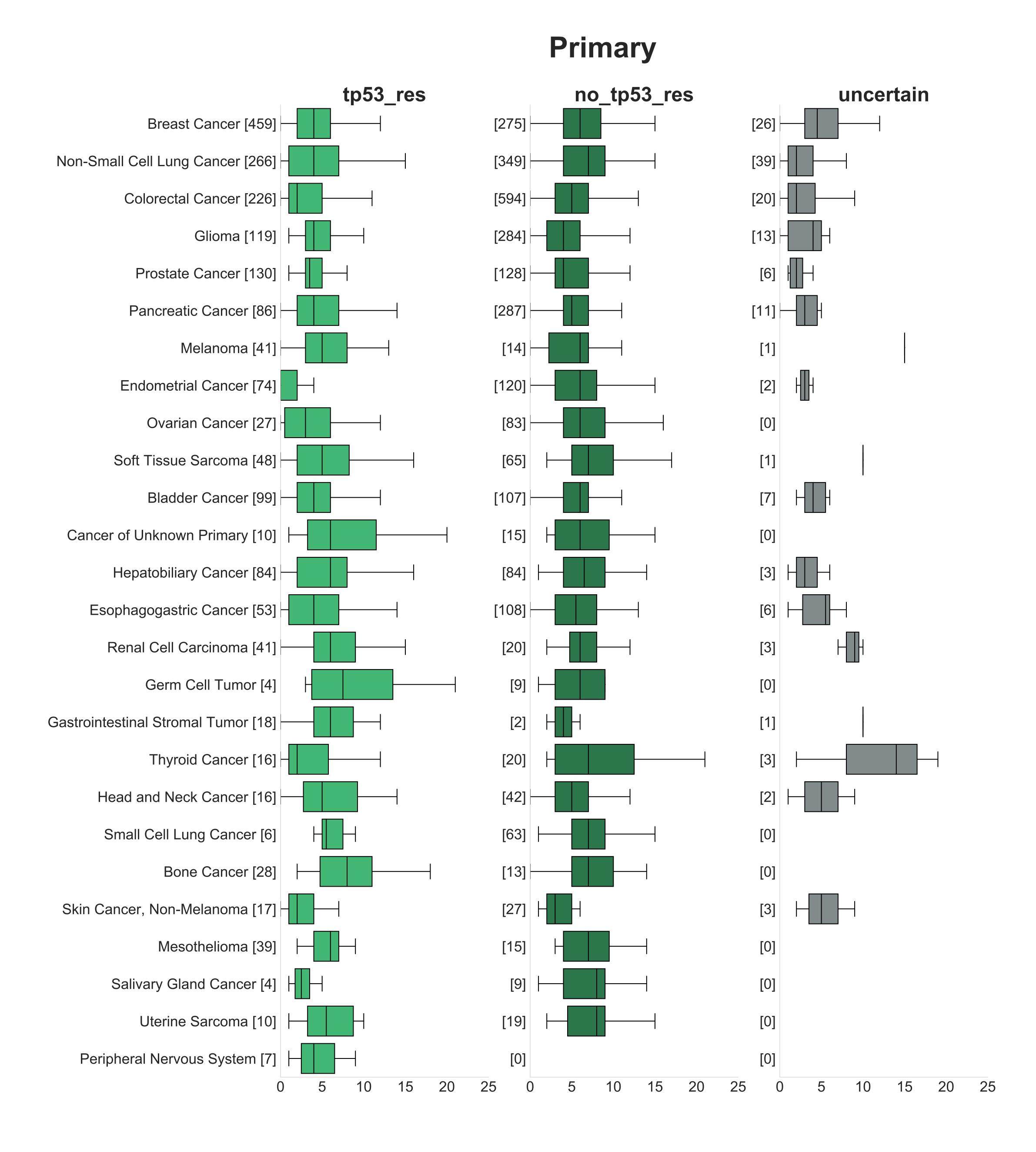
Number of aberrant chromosomes - Primary Samples



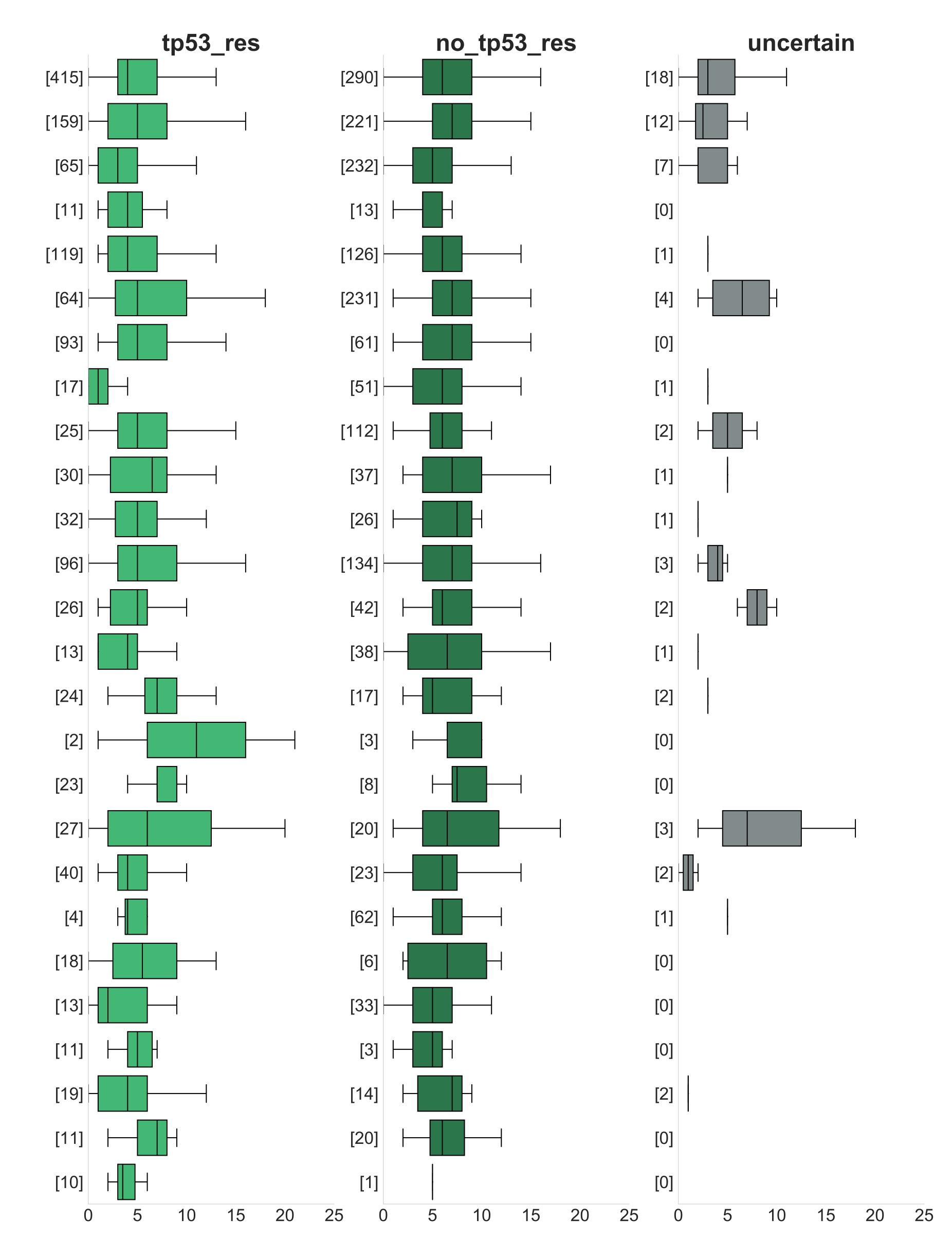
Number of aberrant chromosomes - Metastatic Samples



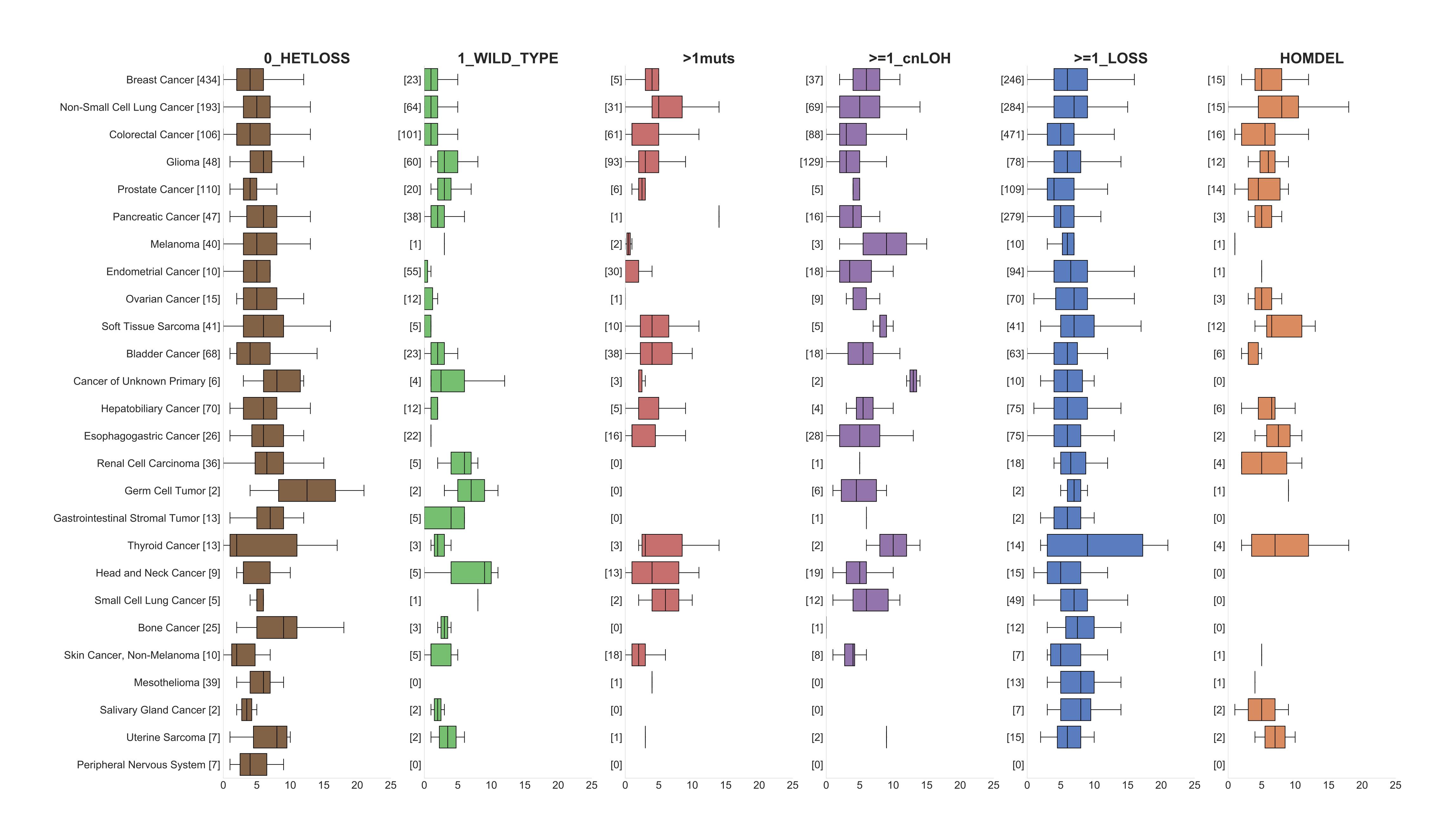
Number of chromosomes affected by a LOSS



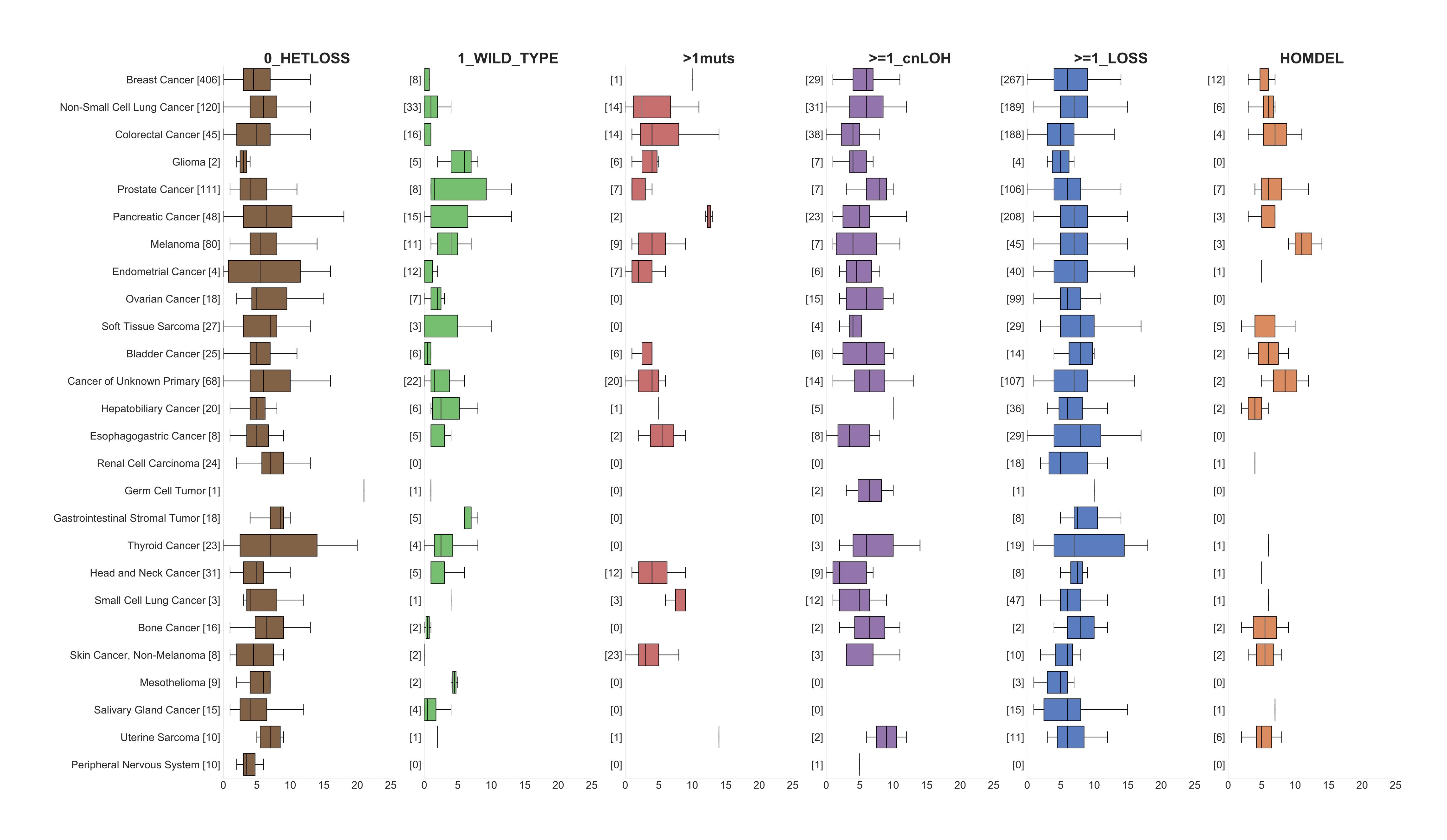
Metastasis



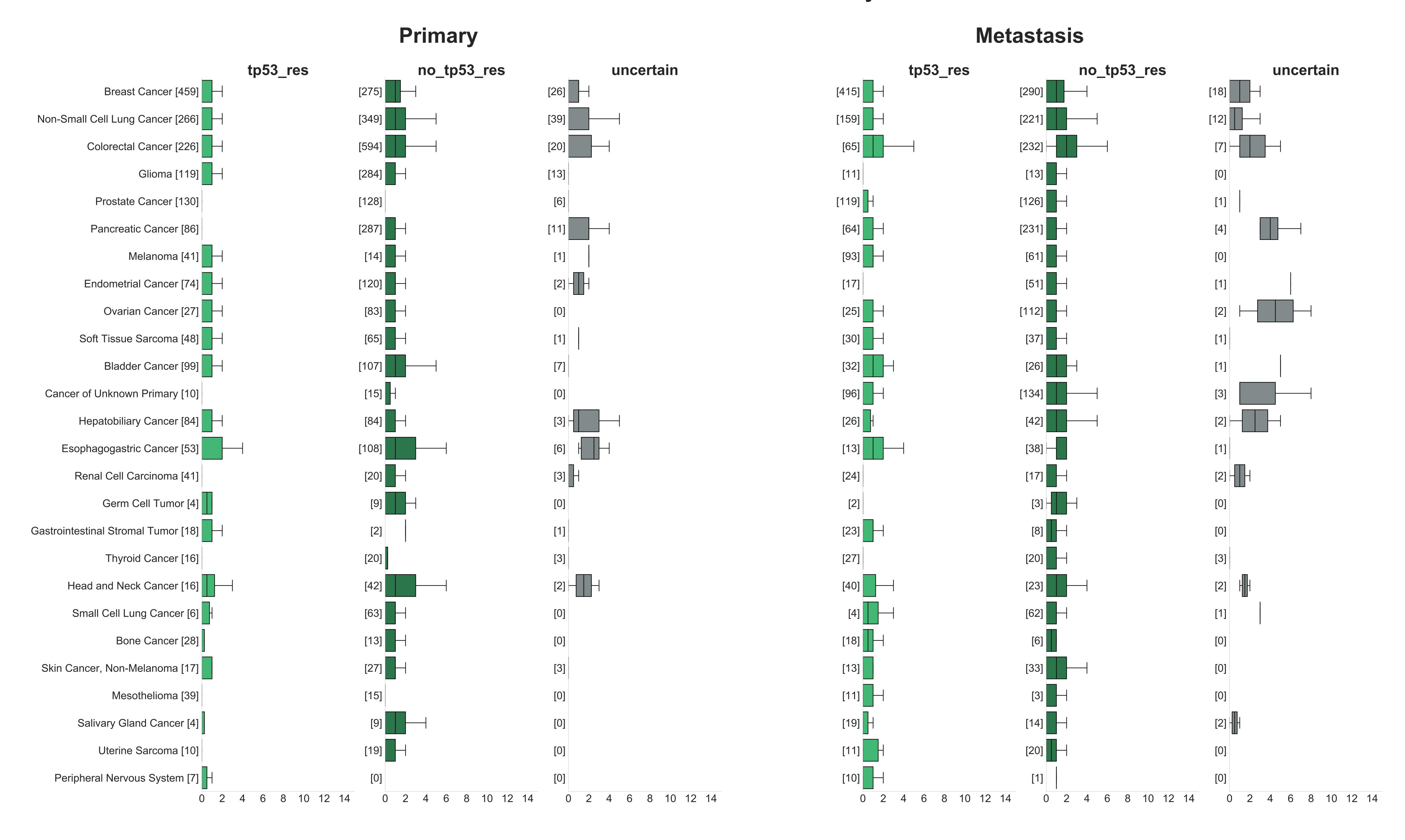
Number of chromosomes affected by a LOSS - Primary Samples



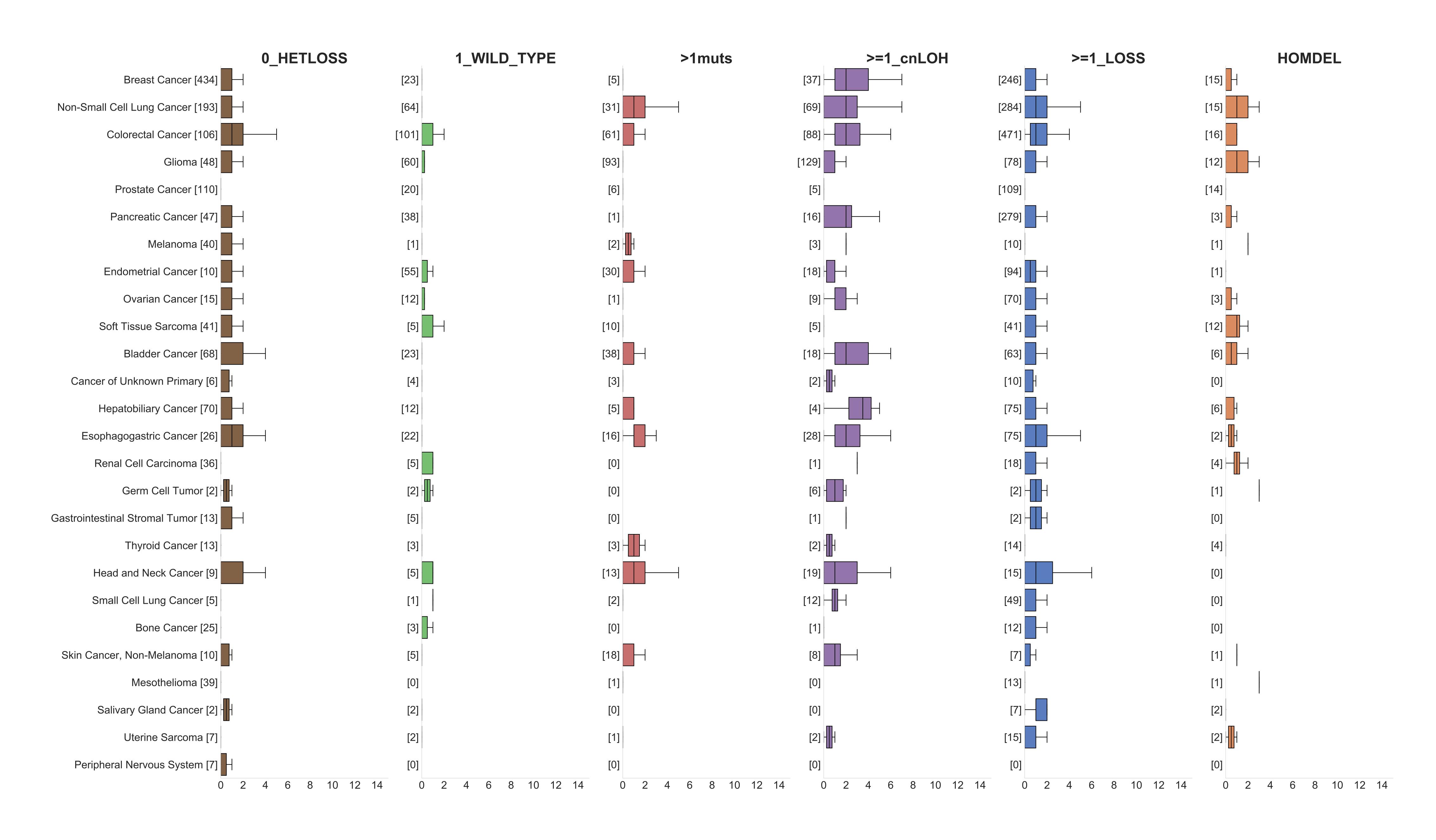
Number of chromosomes affected by a LOSS - Metastatic Samples



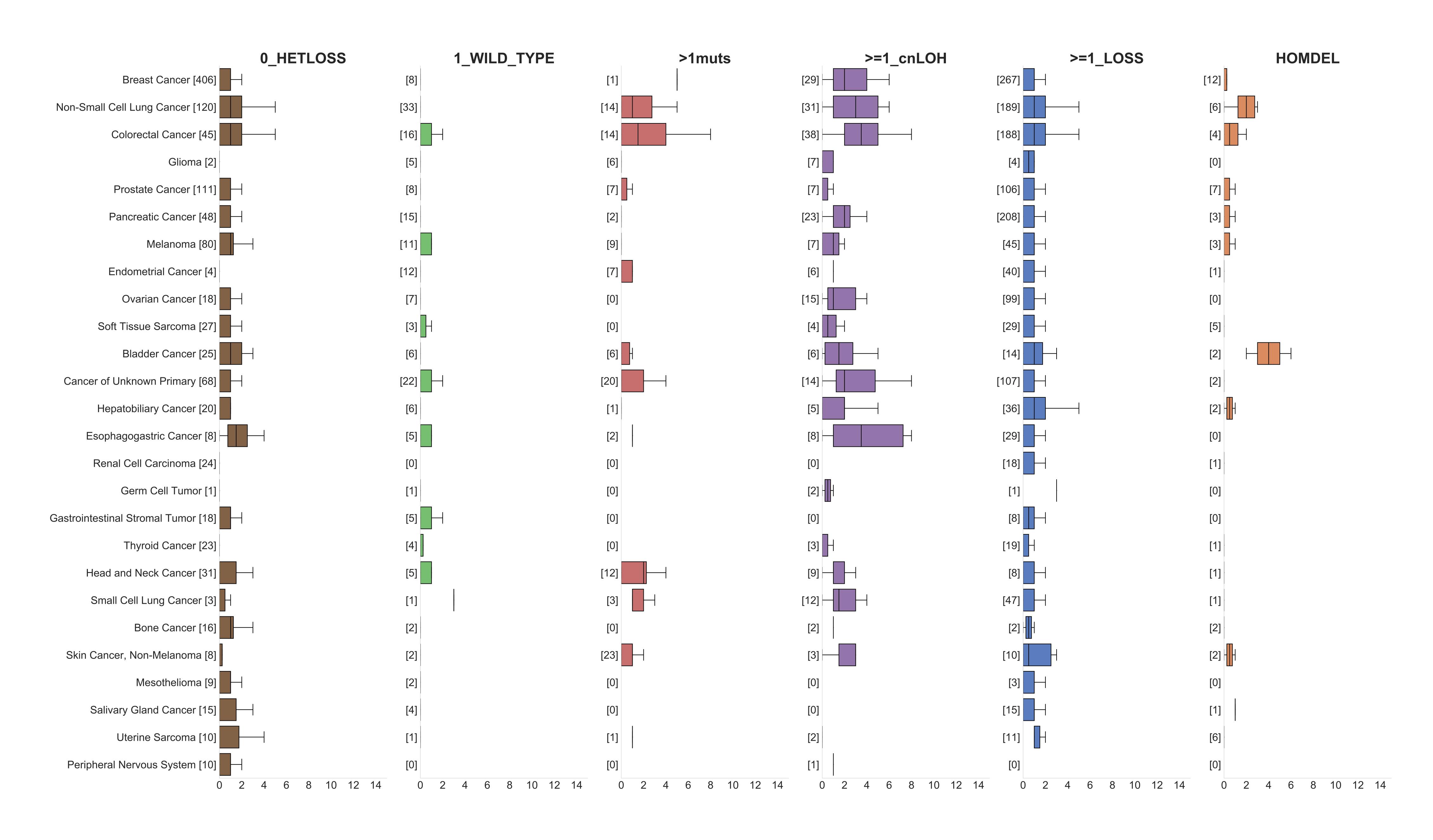
Number of chromosomes affected by a cnLOH



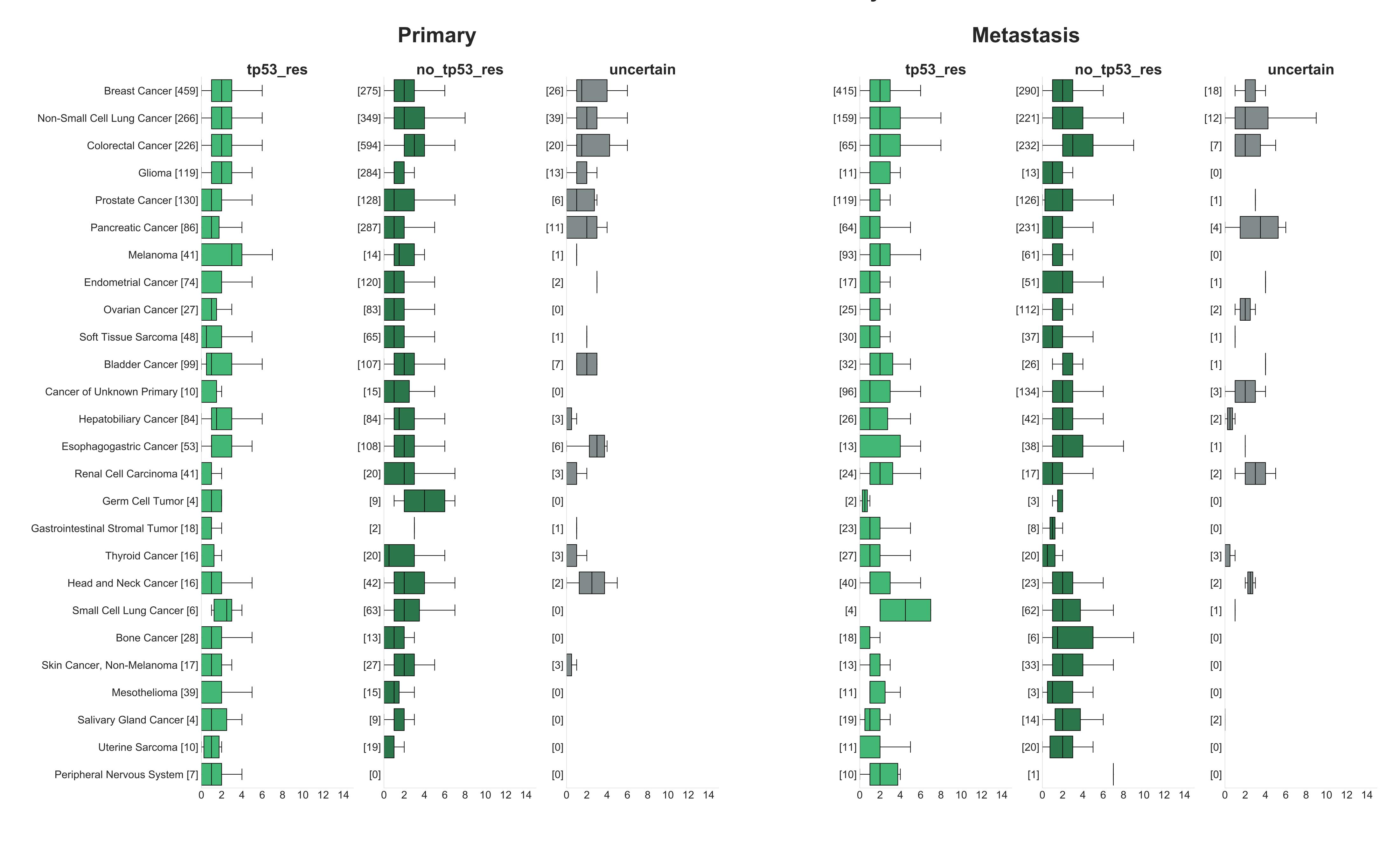
Number of chromosomes affected by a cnLOH - Primary Samples



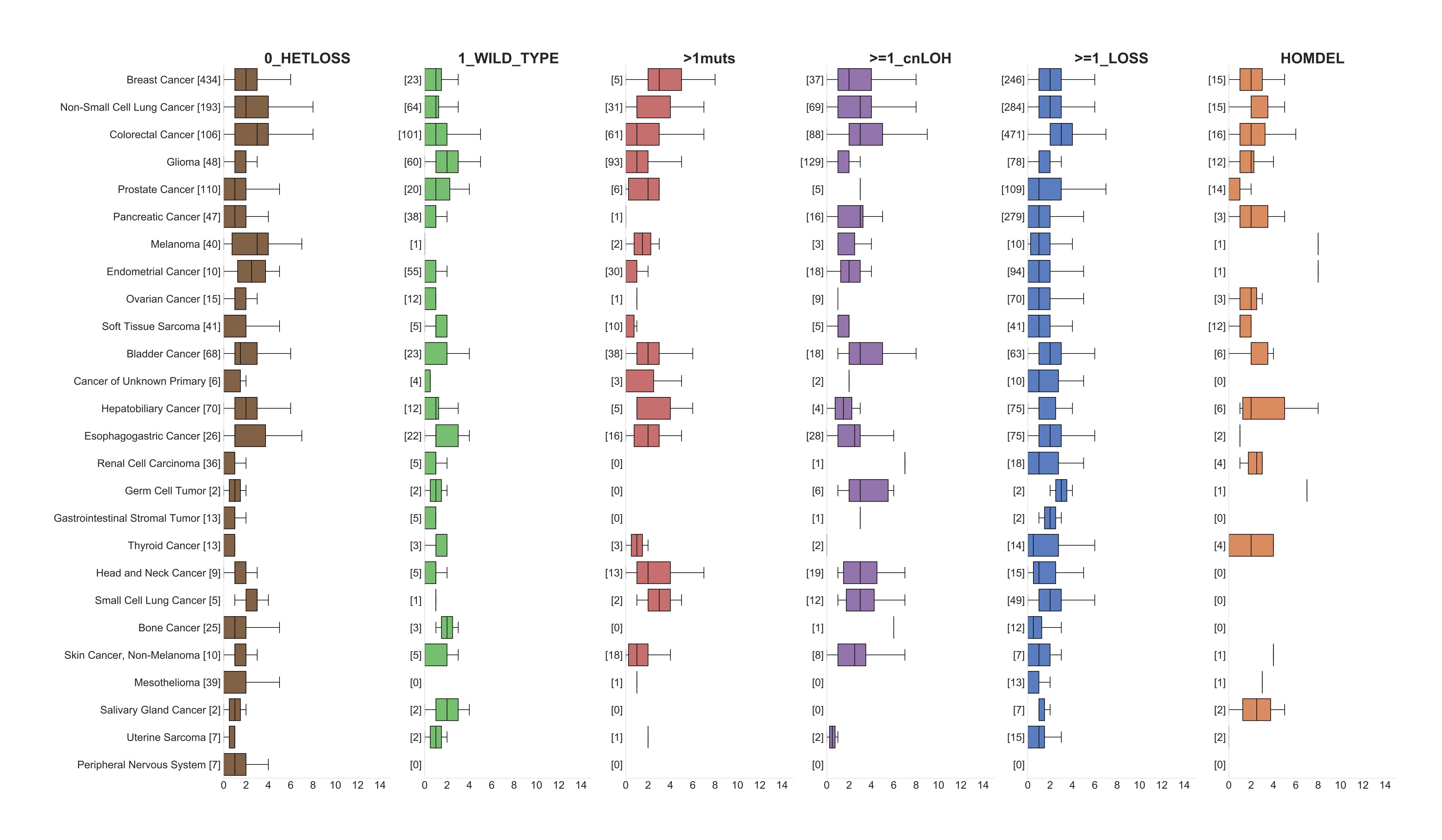
Number of chromosomes affected by a cnLOH - Metastatic Samples



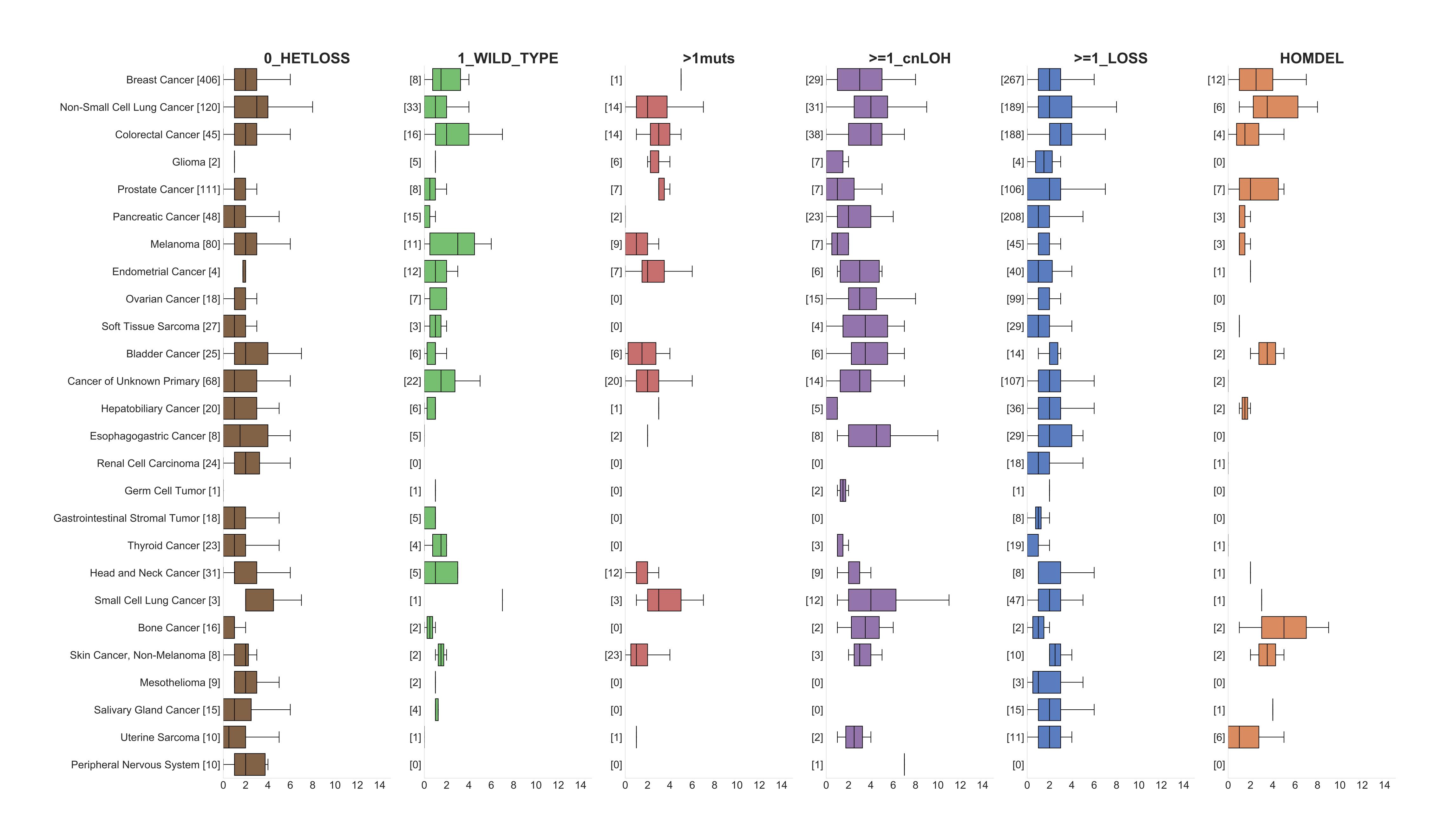
Number of chromosomes affected by a GAIN



Number of chromosomes affected by a GAIN - Primary Samples



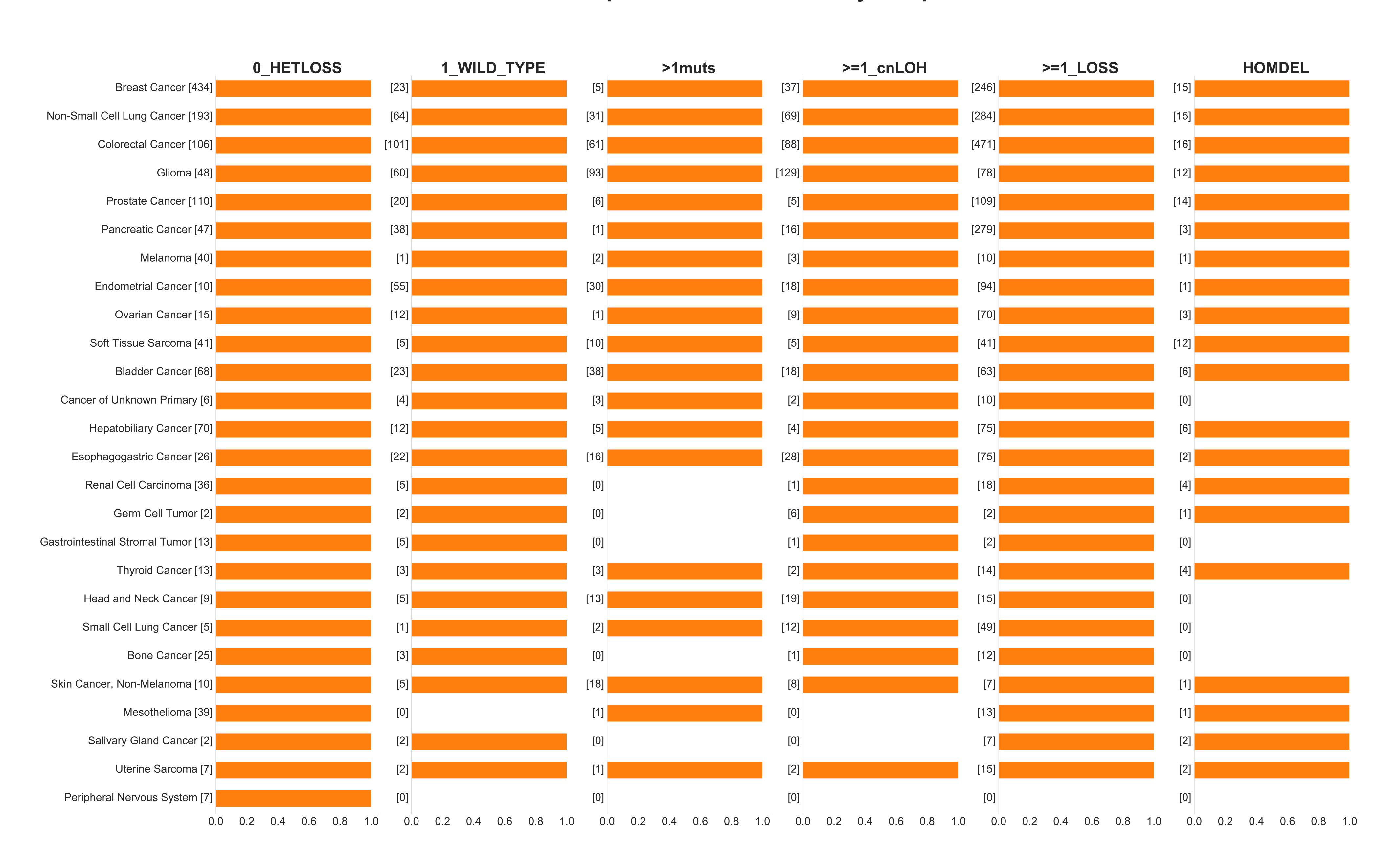
Number of chromosomes affected by a GAIN - Metastatic Samples



Proportion of WGD

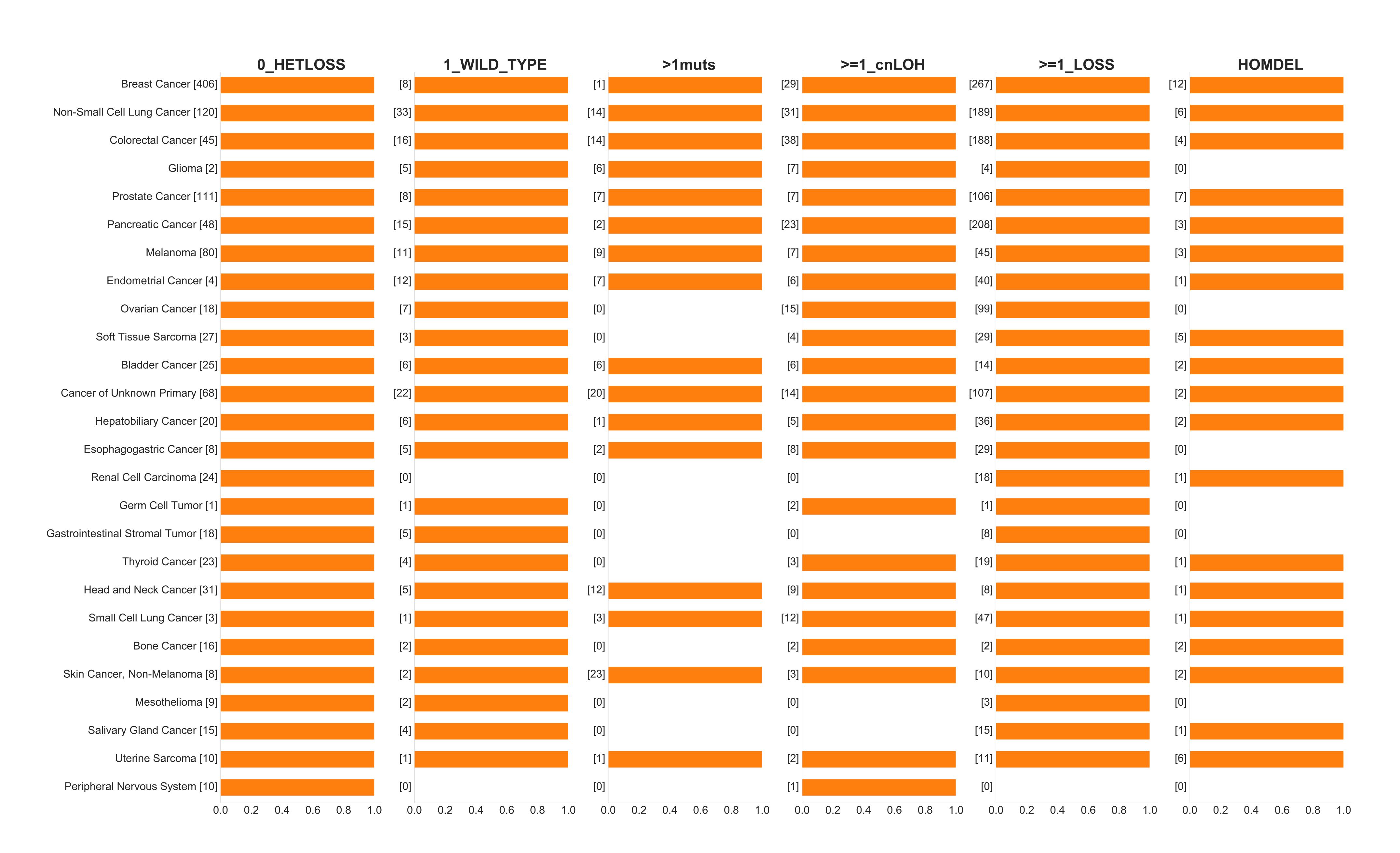


Proportion of WGD - Primary Samples



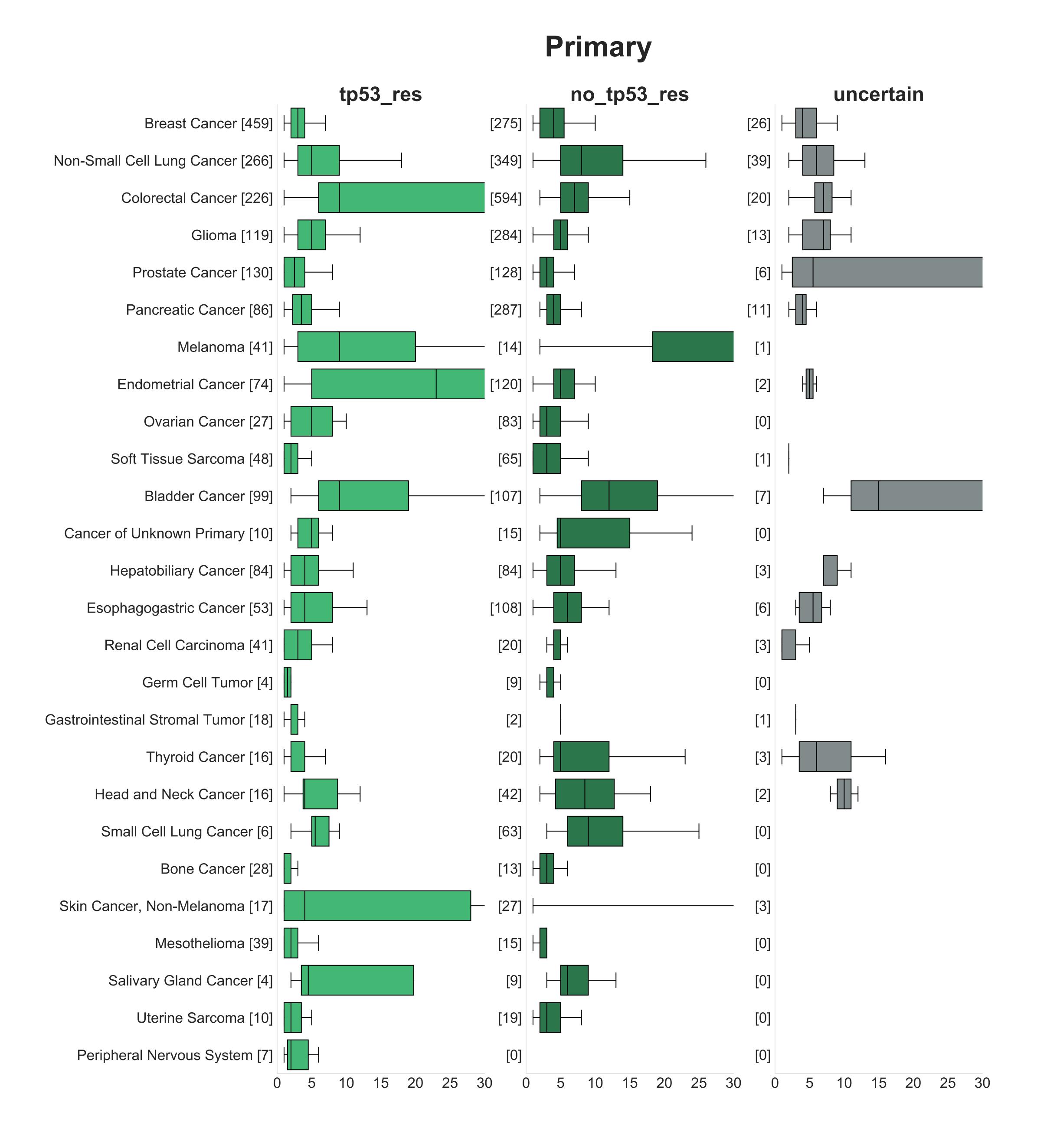


Proportion of WGD - Metastatic Samples

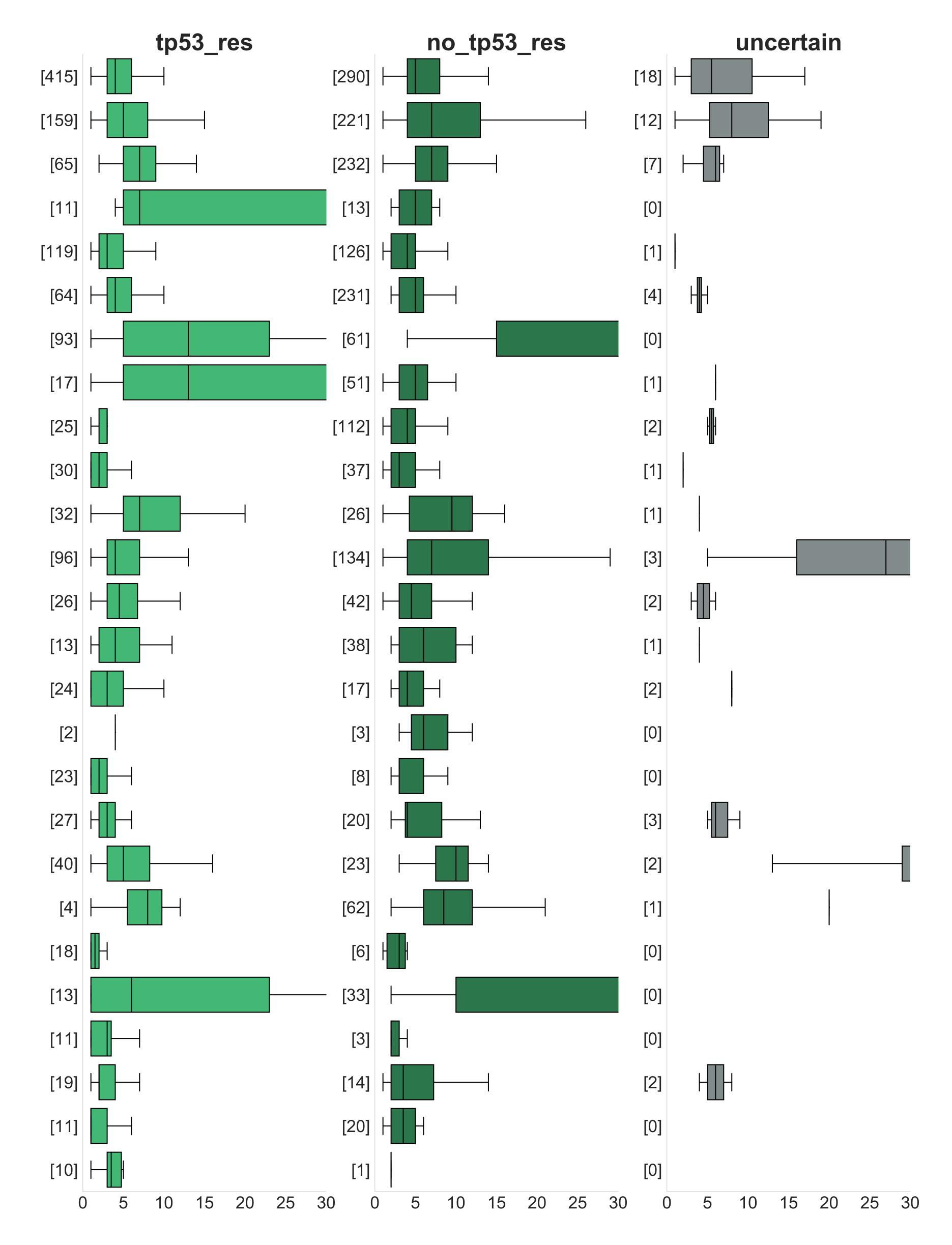




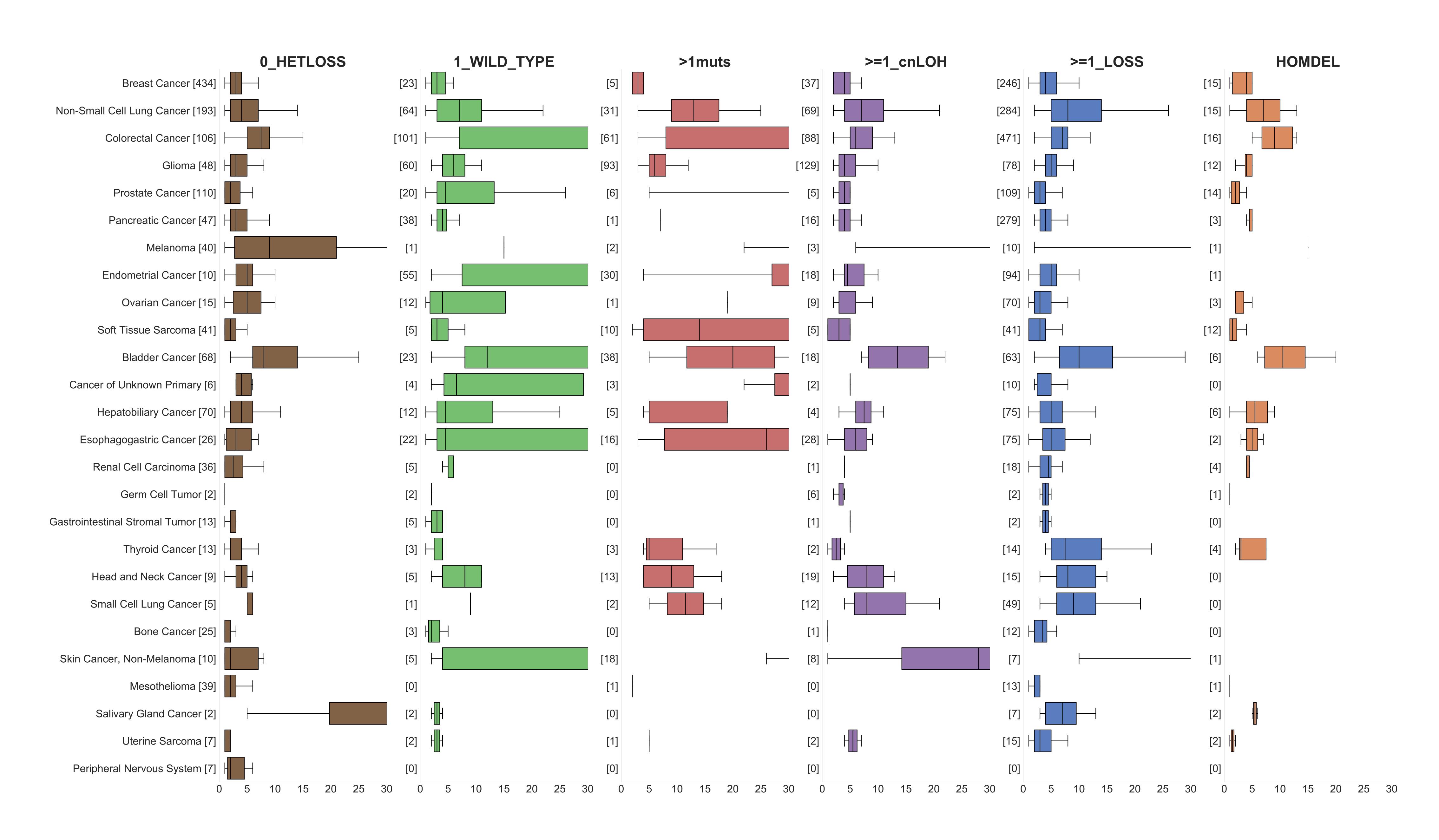
Mutation Count



Metastasis



Mutation Count - Primary Samples



Mutation Count - Metastatic Samples

