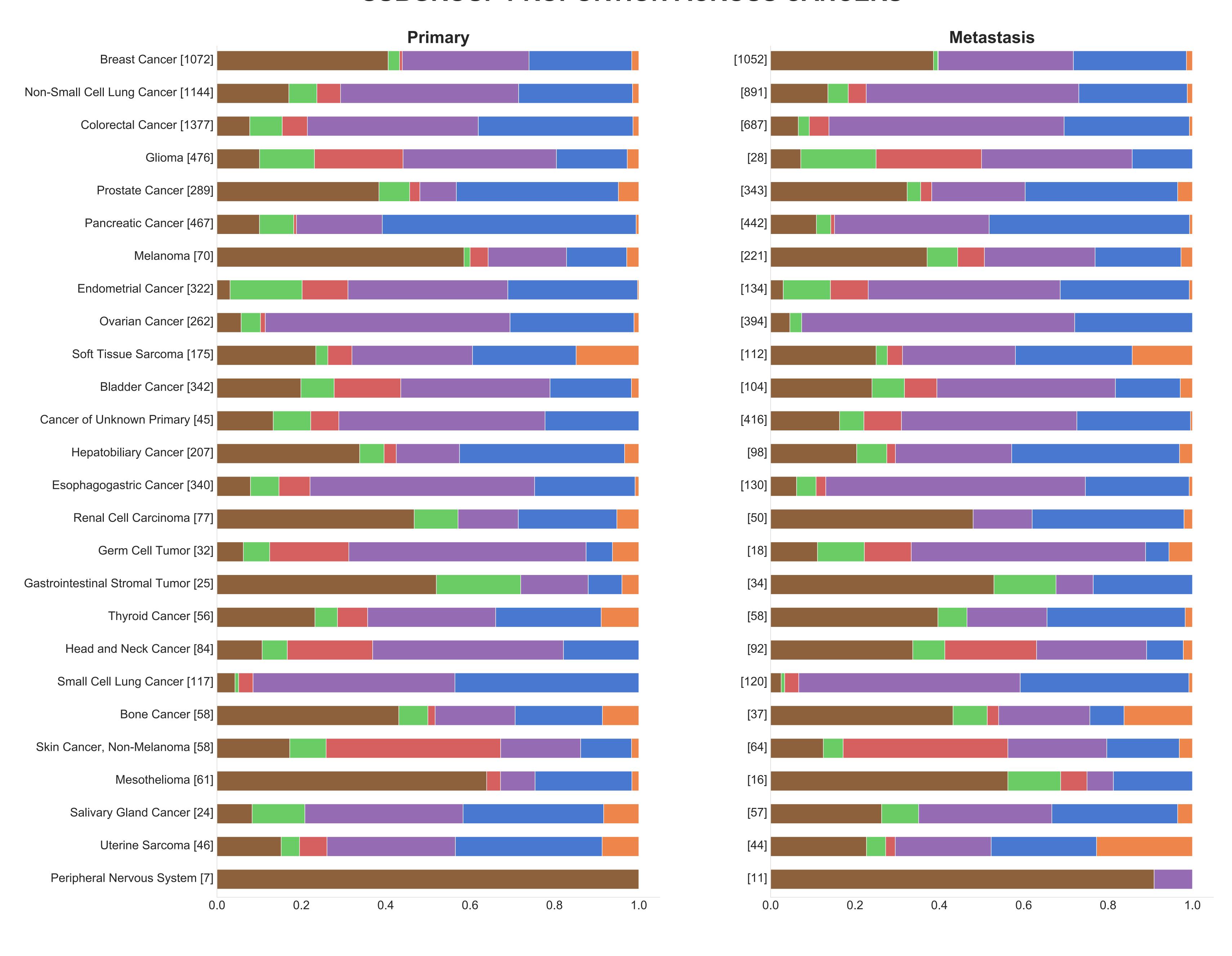
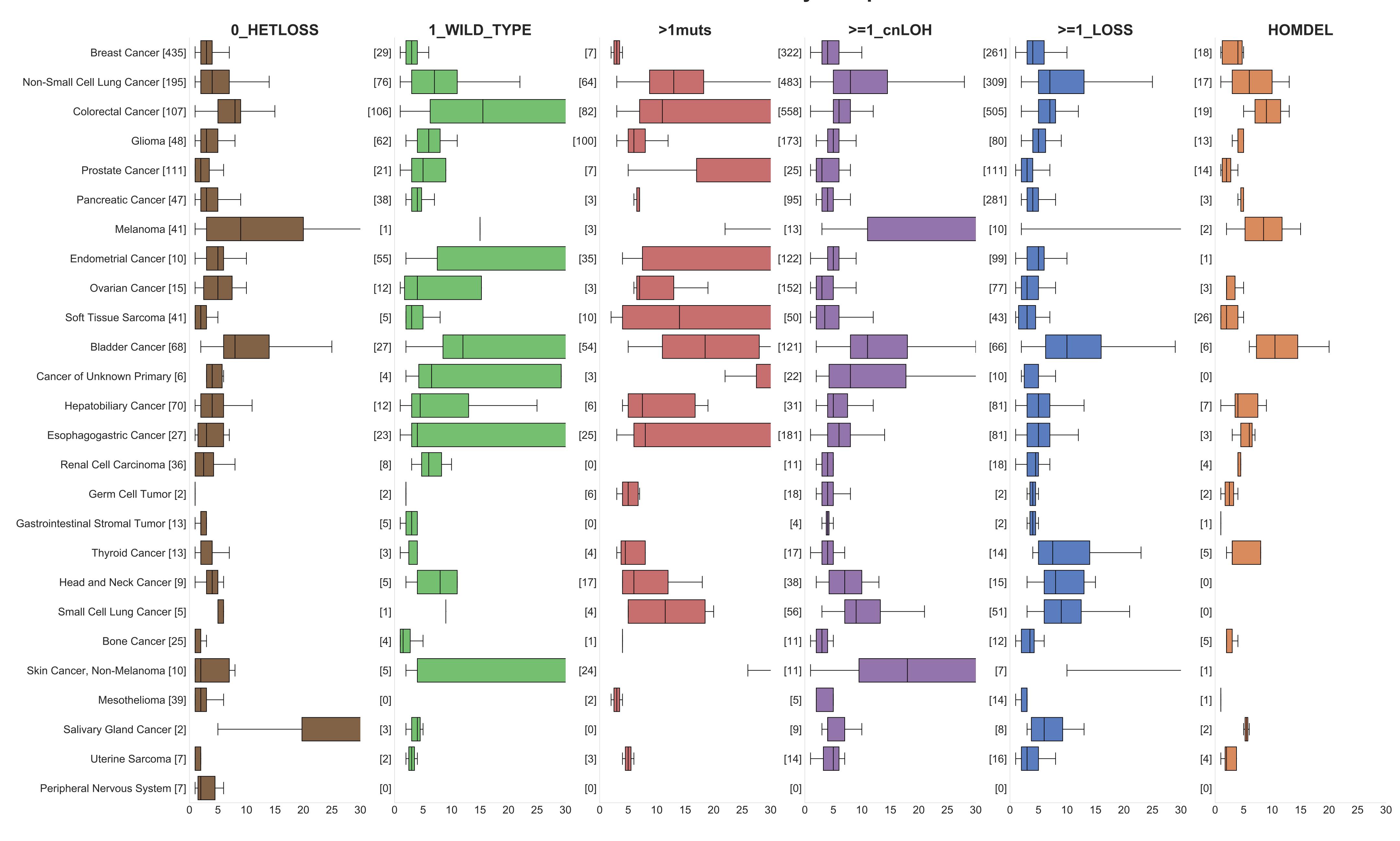
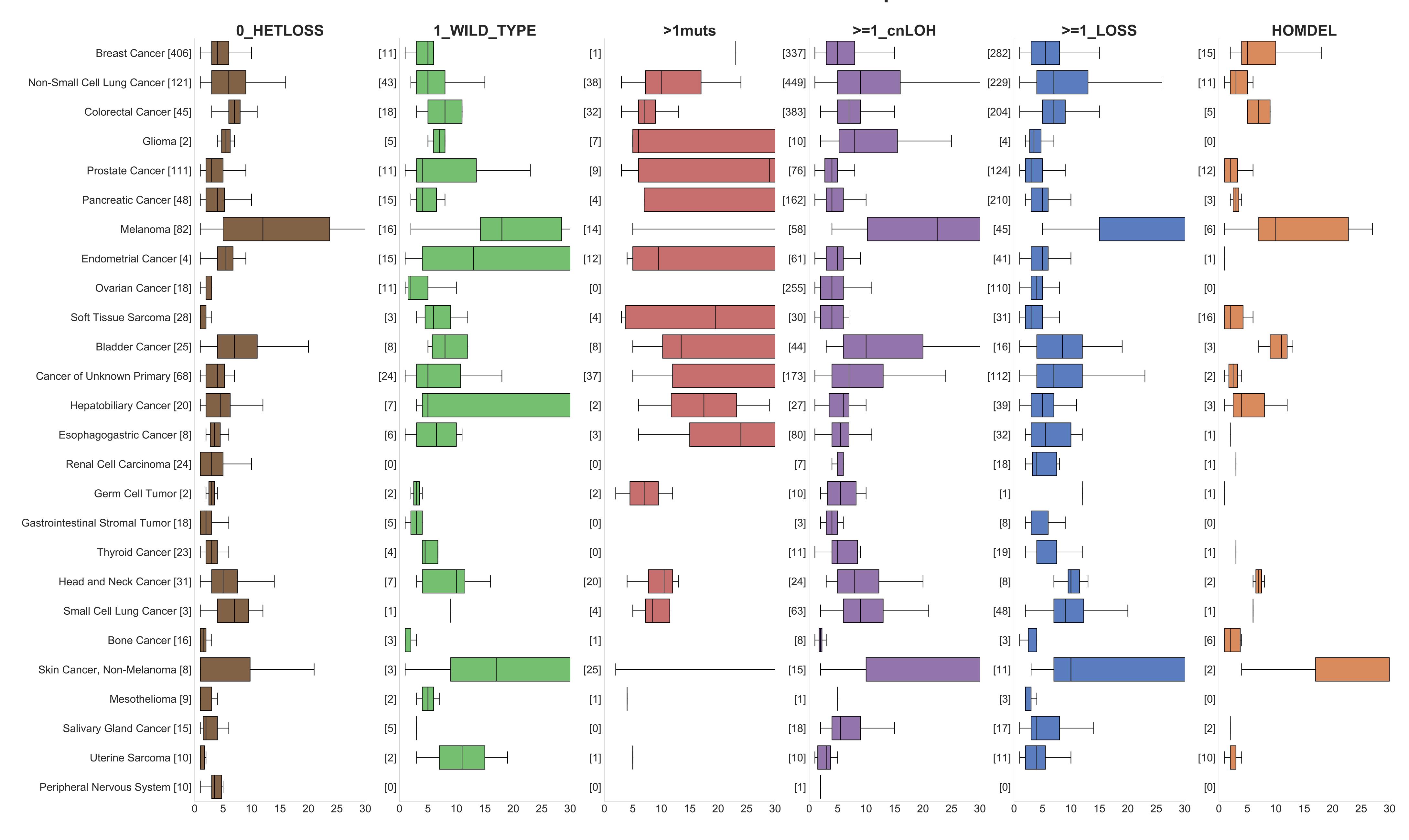
SUBGROUP PROPORTION ACROSS CANCERS



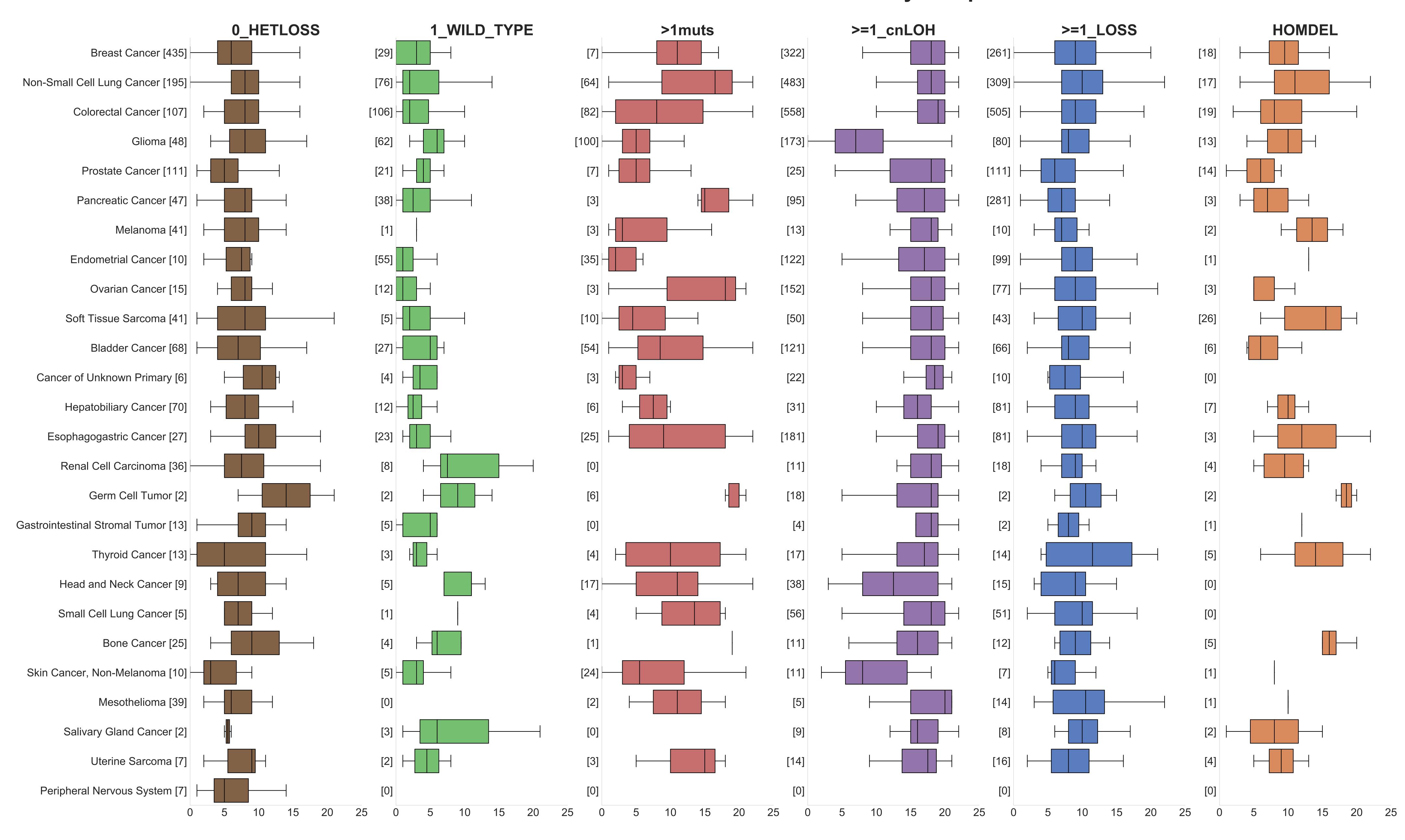
Mutation Count - Primary Samples



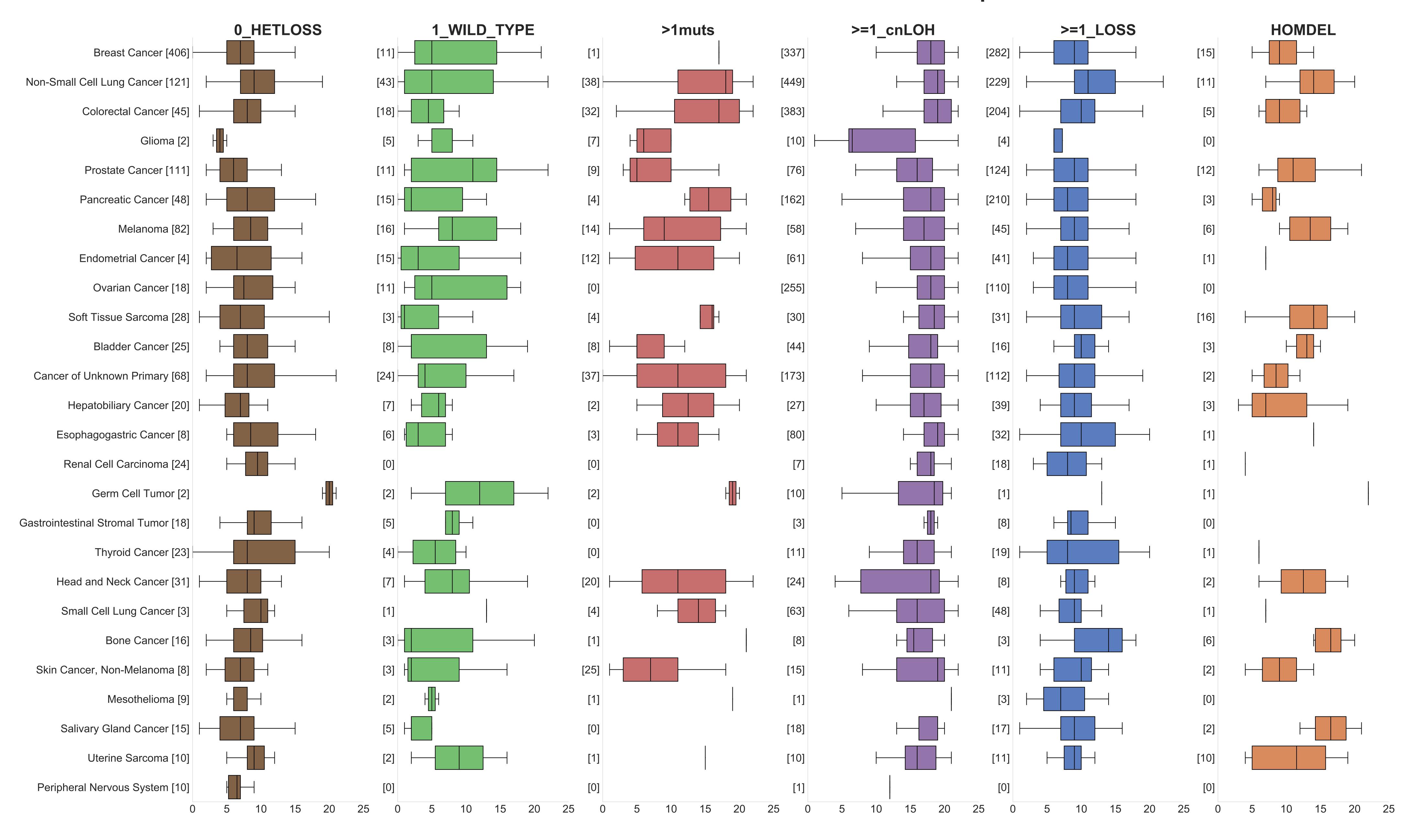
Mutation Count - Metastatic Samples



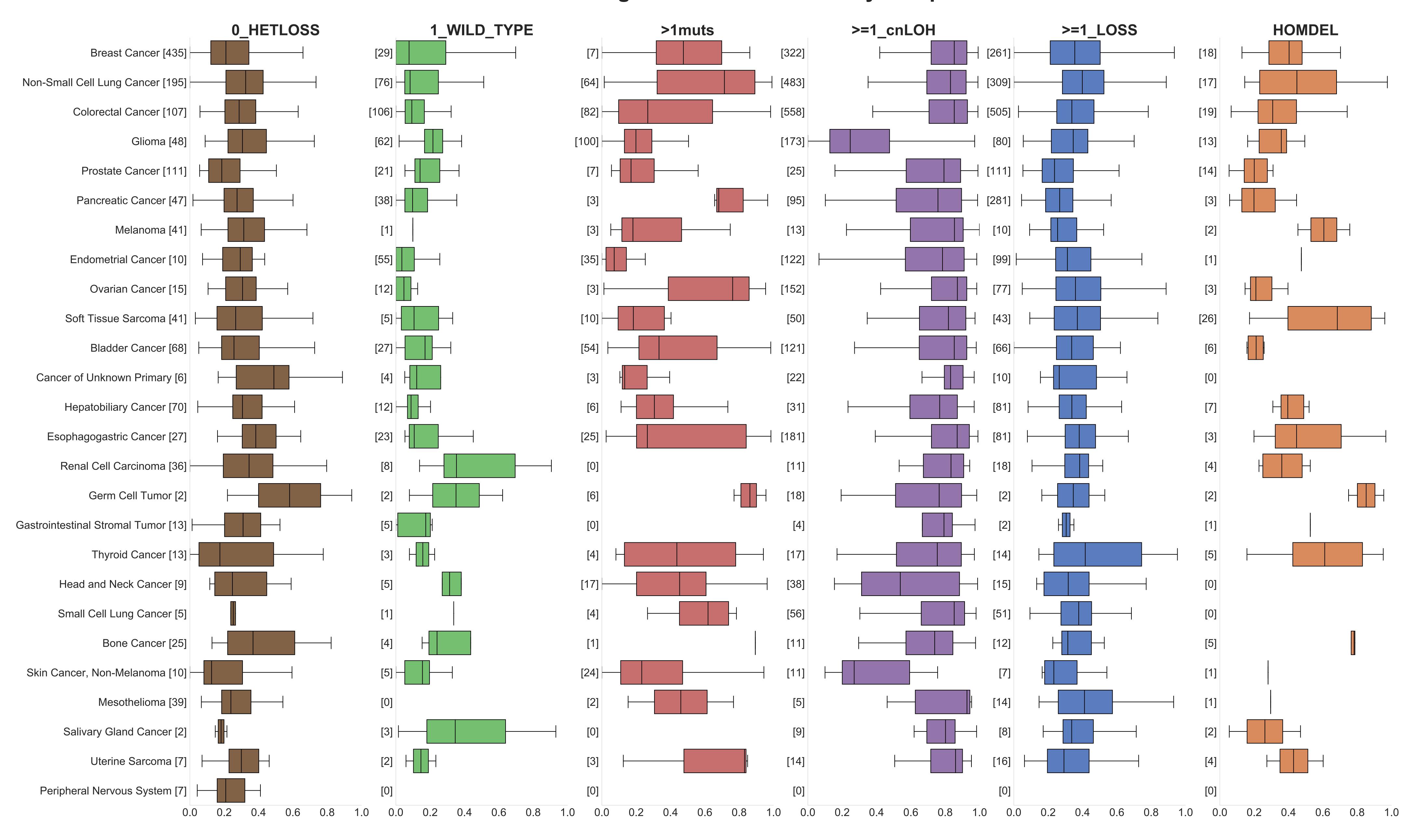
Number of aberrant chromosomes - Primary Samples



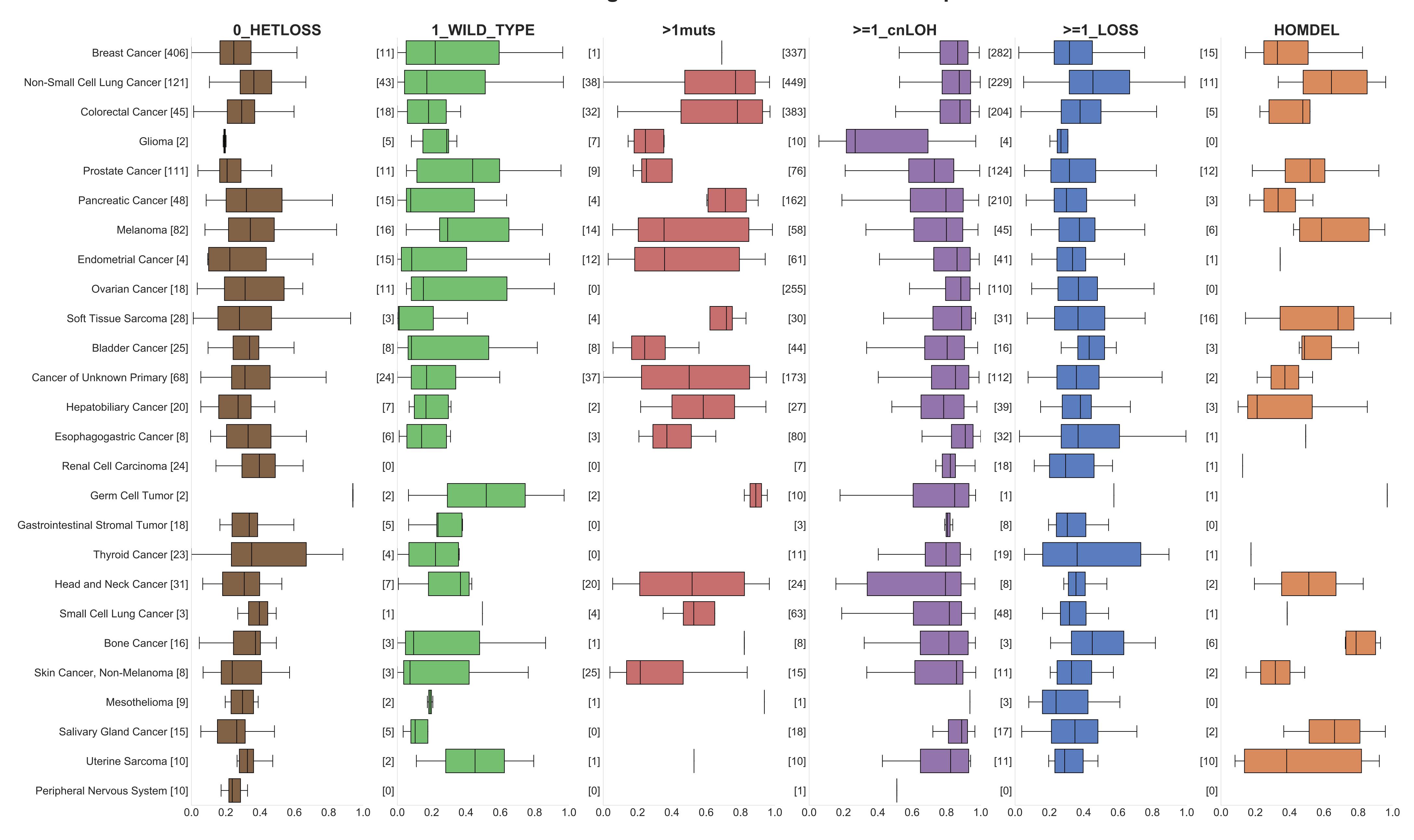
Number of aberrant chromosomes - Metastatic Samples



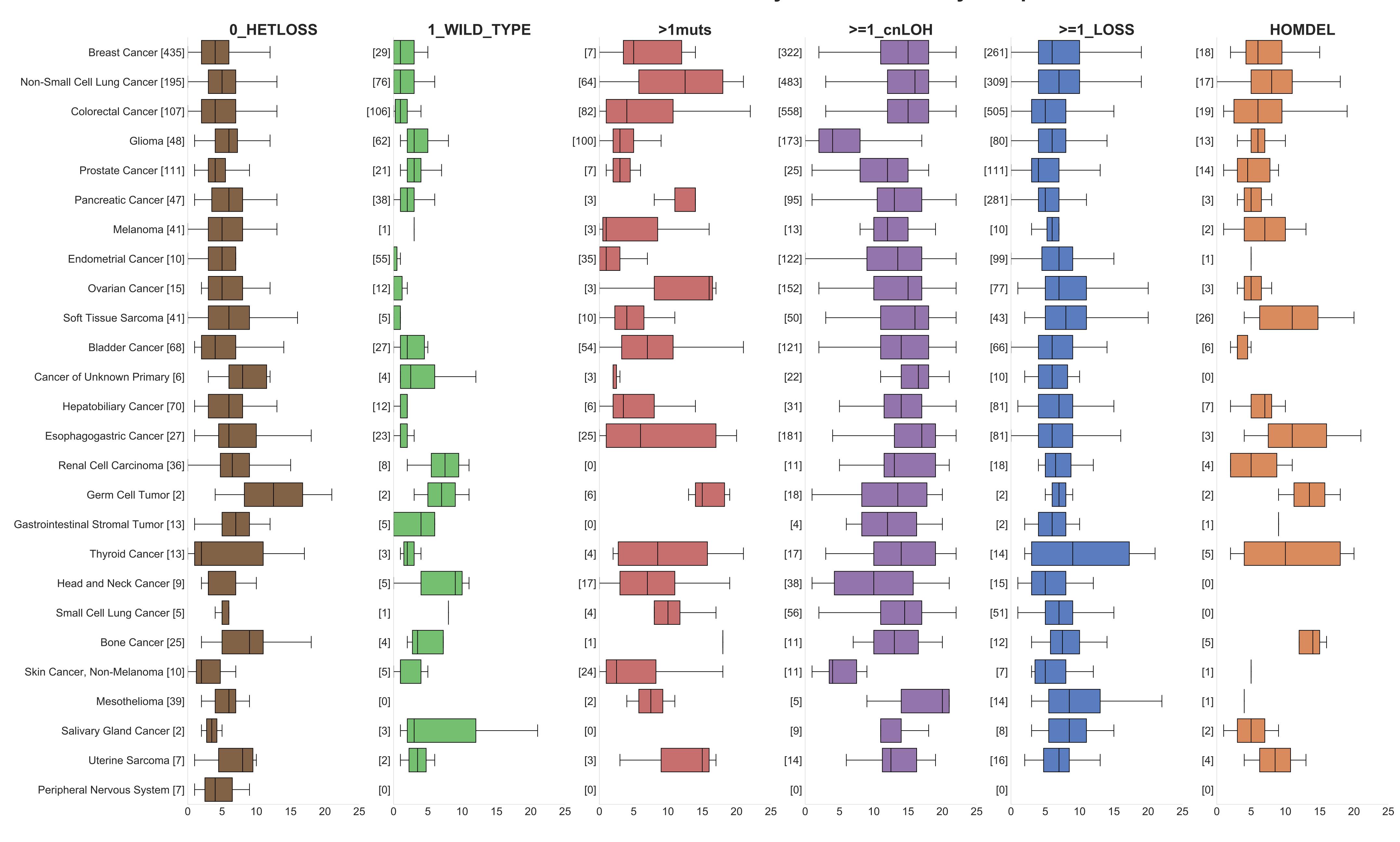
Fraction of genome altered - Primary Samples



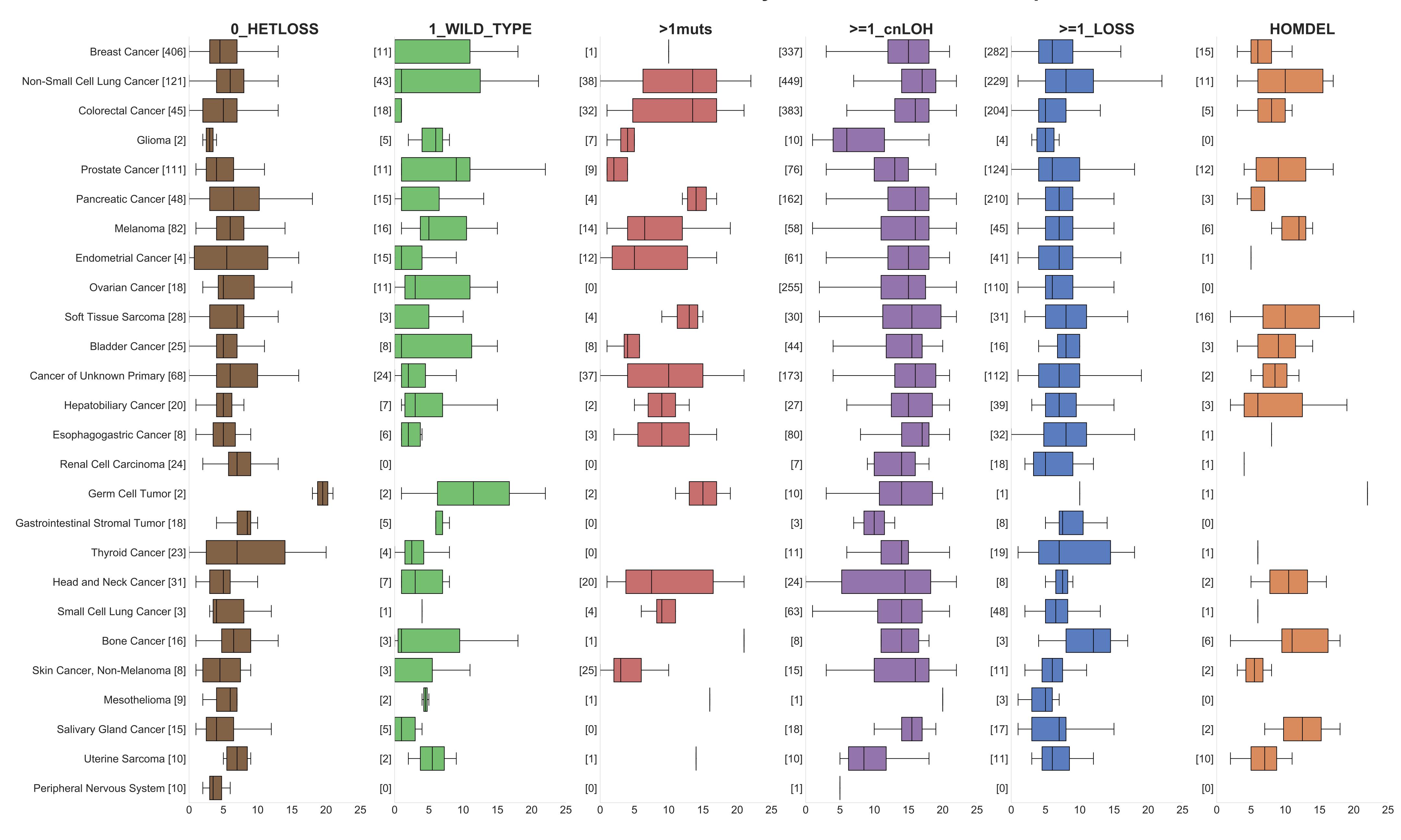
Fraction of genome altered - Metastatic Samples



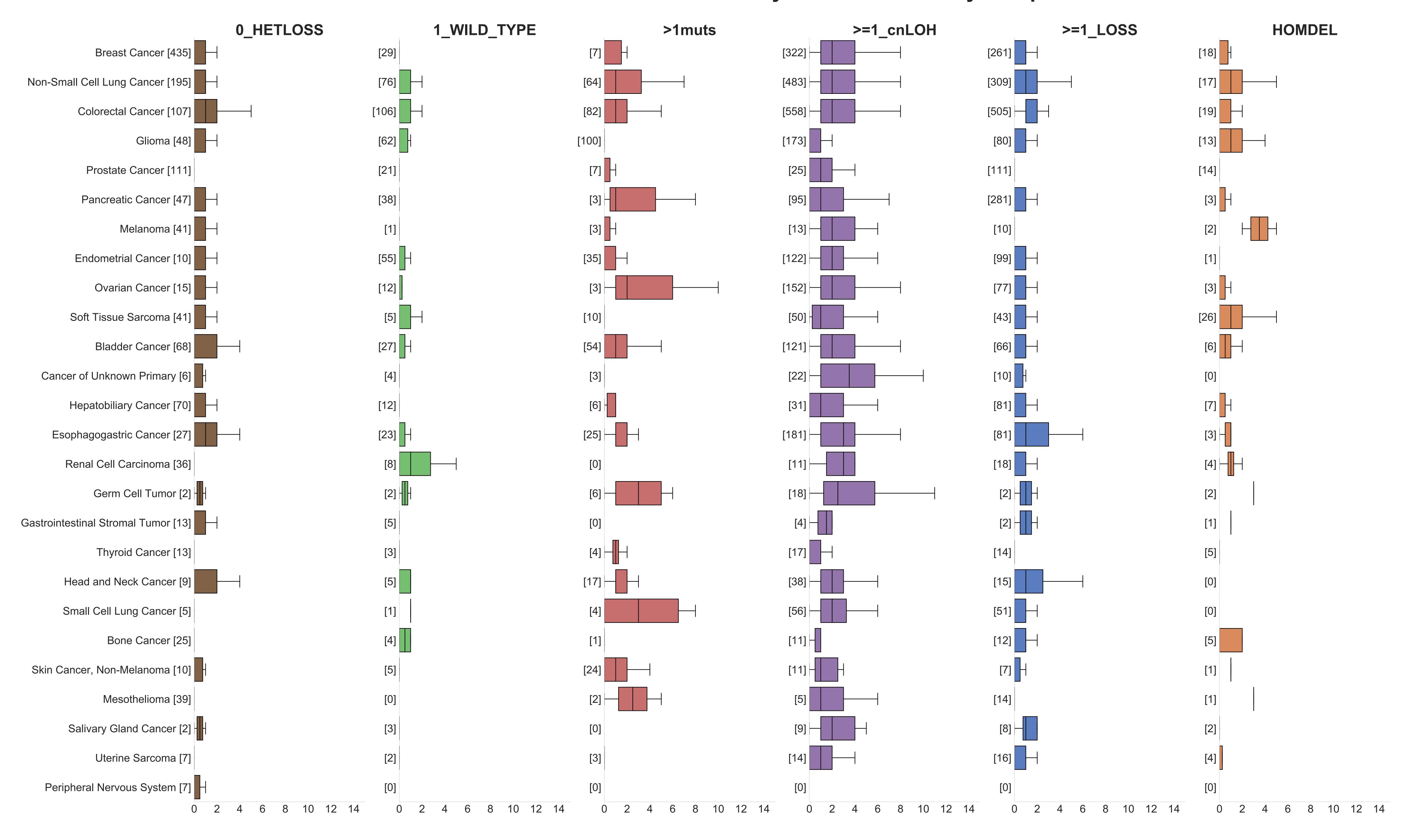
Number of chromosomes affected by a LOSS - Primary Samples



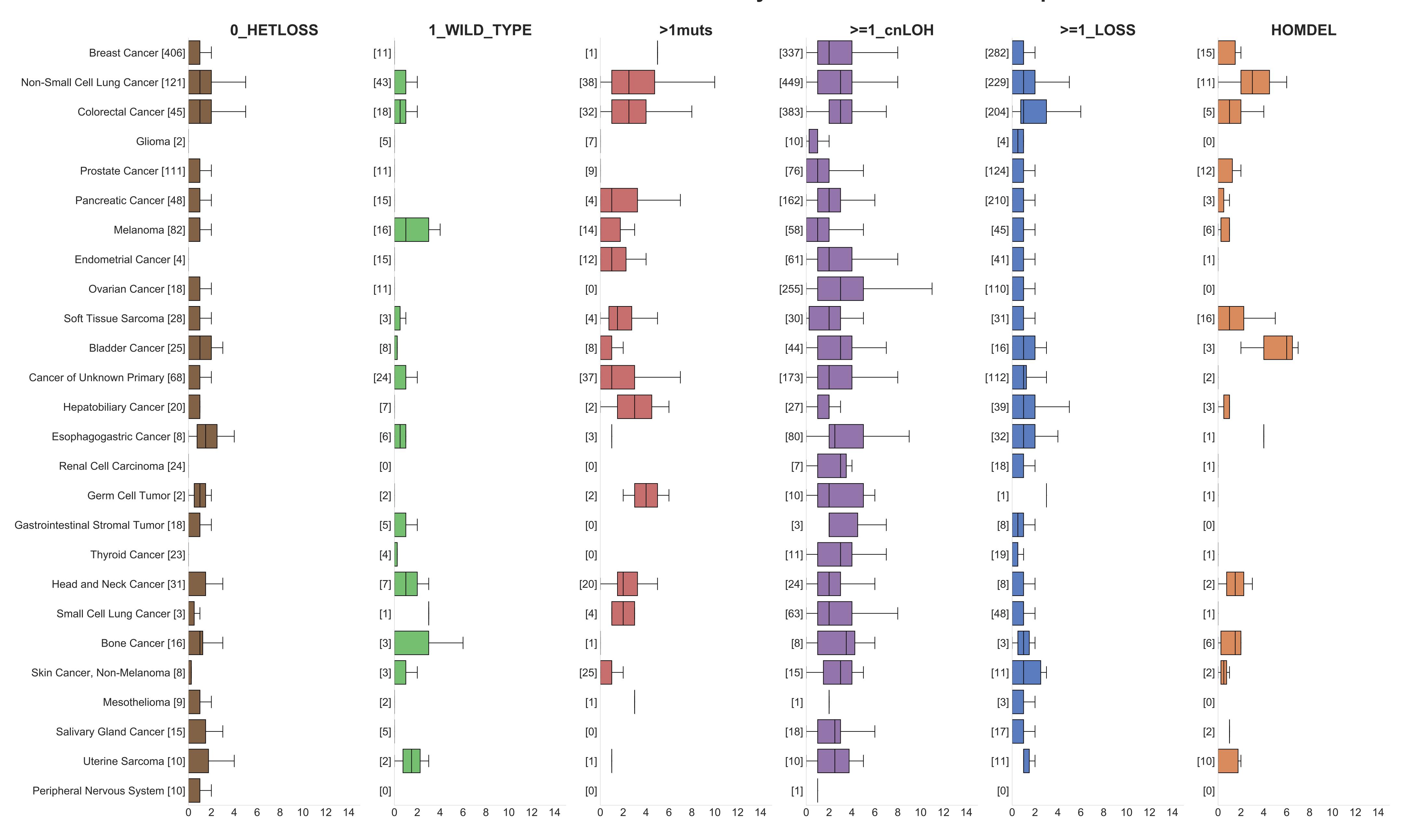
Number of chromosomes affected by a LOSS - Metastatic Samples



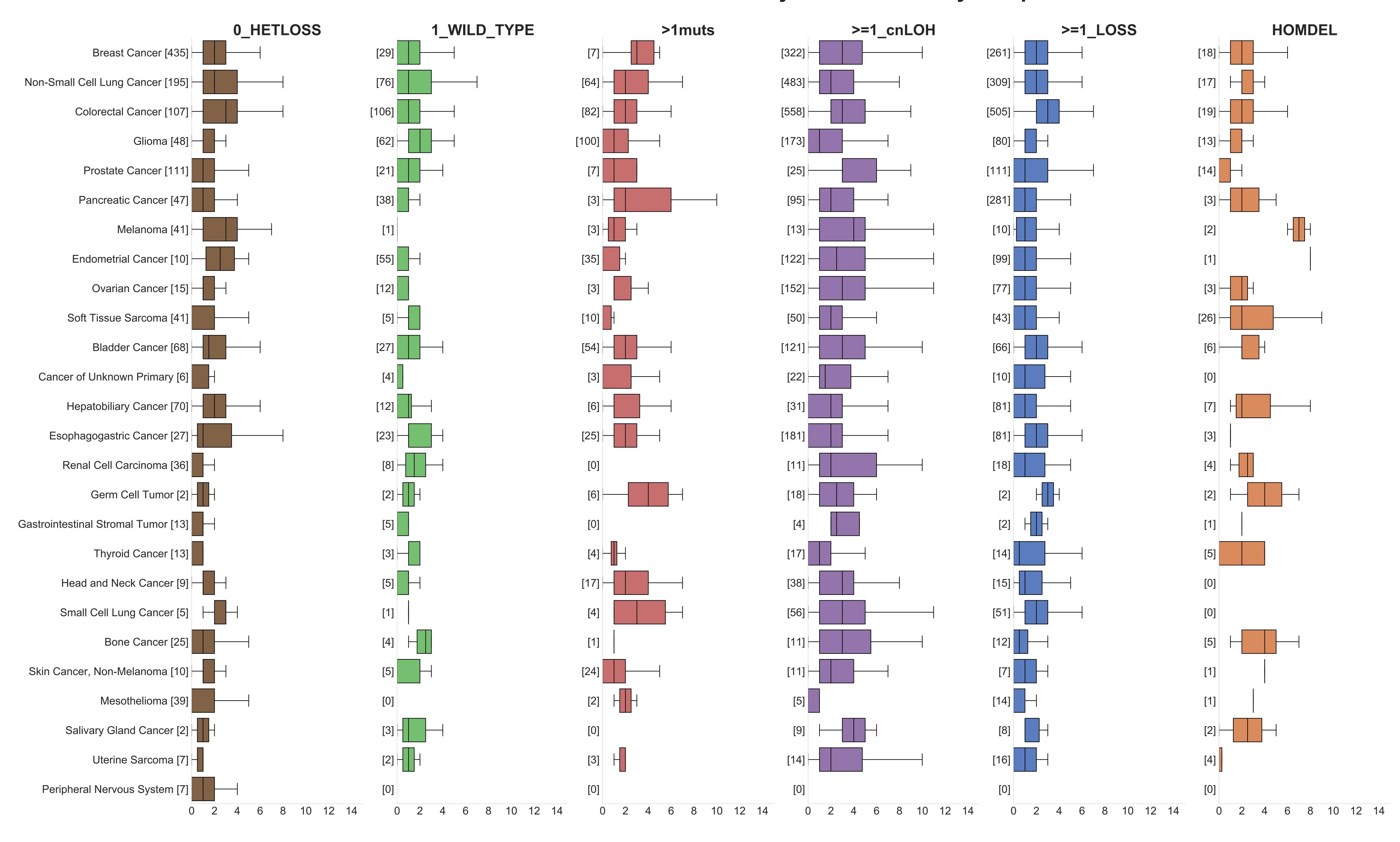
Number of chromosomes affected by a cnLOH - Primary Samples



Number of chromosomes affected by a cnLOH - Metastatic Samples



Number of chromosomes affected by a GAIN - Primary Samples



Number of chromosomes affected by a GAIN - Metastatic Samples

