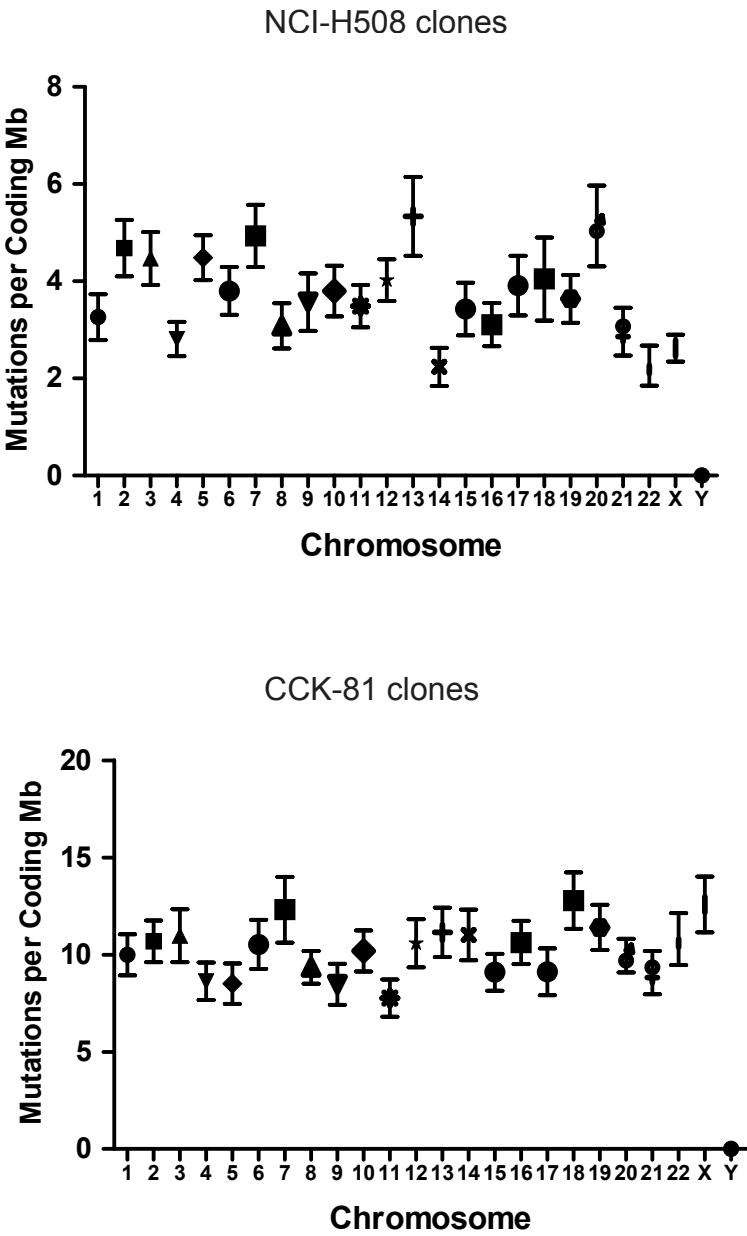


Supplemental Figure S3



Supp Figure S3. Distribution of coding mutations per chromosome. All mutations detected in the NCI-H508 and CCK-81 cetuximab-resistant clones were combined and the mean number of mutations calculated per Mb of coding exon per chromosome.