

## Mutations in Cancer

*Positive dataset*



*Ref genome: v38*

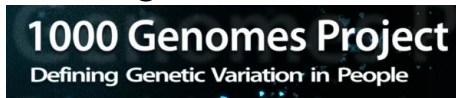
Original:  
127,424 in coding region,  
747,312 in non-coding region



Recurrence Number:  
1,260 in coding region,  
5,260 in non-coding region

## Neutral Mutation

*Negative dataset*



*Ref genome: v38*

Original:  
98,123 variants



MAF Selection:  
66,514 variants



Annotation & Selection:  
432 in coding region,  
7,268 in non-coding region