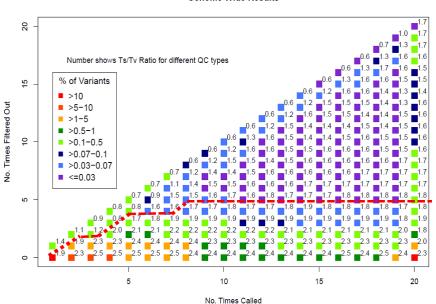
## Genome Wide Results



265 duplicates within and between cohorts 100 200 300 500 number of differing genotypes between sample pairs (out of 1000)