

# Class10R

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## Examine 1000 Genome Data

Q5. What proportion of the Mexican Ancestry in Los Angeles sample population (MXL) are homozygous for the asthma associated SNP (G|G)?

```
# Read genotype file of from Ensembl
mxl <- read.csv("373531-SampleGenotypes-Homo_sapiens_Variation_Sample_rs8067378.csv")
mxl
```

##	Sample..Male.Female.Unknown.	Genotype..forward.strand.	Population.s.	Father
## 1	NA19648 (F)	A A	ALL, AMR, MXL	-
## 2	NA19649 (M)	G G	ALL, AMR, MXL	-
## 3	NA19651 (F)	A A	ALL, AMR, MXL	-
## 4	NA19652 (M)	G G	ALL, AMR, MXL	-
## 5	NA19654 (F)	G G	ALL, AMR, MXL	-
## 6	NA19655 (M)	A G	ALL, AMR, MXL	-
## 7	NA19657 (F)	A G	ALL, AMR, MXL	-
## 8	NA19658 (M)	A A	ALL, AMR, MXL	-
## 9	NA19661 (M)	A G	ALL, AMR, MXL	-
## 10	NA19663 (F)	A A	ALL, AMR, MXL	-
## 11	NA19664 (M)	G A	ALL, AMR, MXL	-
## 12	NA19669 (F)	A A	ALL, AMR, MXL	-
## 13	NA19670 (M)	A A	ALL, AMR, MXL	-
## 14	NA19676 (M)	G G	ALL, AMR, MXL	-
## 15	NA19678 (F)	A A	ALL, AMR, MXL	-
## 16	NA19679 (M)	A G	ALL, AMR, MXL	-
## 17	NA19681 (F)	A G	ALL, AMR, MXL	-
## 18	NA19682 (M)	A G	ALL, AMR, MXL	-
## 19	NA19684 (F)	A G	ALL, AMR, MXL	-
## 20	NA19716 (F)	G A	ALL, AMR, MXL	-
## 21	NA19717 (M)	A G	ALL, AMR, MXL	-
## 22	NA19719 (F)	G G	ALL, AMR, MXL	-
## 23	NA19720 (M)	G G	ALL, AMR, MXL	-
## 24	NA19722 (F)	G A	ALL, AMR, MXL	-
## 25	NA19723 (M)	G G	ALL, AMR, MXL	-
## 26	NA19725 (F)	A G	ALL, AMR, MXL	-
## 27	NA19726 (M)	A A	ALL, AMR, MXL	-
## 28	NA19728 (F)	A A	ALL, AMR, MXL	-
## 29	NA19729 (M)	A G	ALL, AMR, MXL	-
## 30	NA19731 (F)	A A	ALL, AMR, MXL	-
## 31	NA19732 (M)	A G	ALL, AMR, MXL	-

## 32	NA19734 (F)	G A ALL, AMR, MXL	-
## 33	NA19735 (M)	G G ALL, AMR, MXL	-
## 34	NA19740 (F)	A A ALL, AMR, MXL	-
## 35	NA19741 (M)	A A ALL, AMR, MXL	-
## 36	NA19746 (F)	A A ALL, AMR, MXL	-
## 37	NA19747 (M)	G A ALL, AMR, MXL	-
## 38	NA19749 (F)	A G ALL, AMR, MXL	-
## 39	NA19750 (M)	A G ALL, AMR, MXL	-
## 40	NA19752 (F)	A G ALL, AMR, MXL	-
## 41	NA19755 (F)	A A ALL, AMR, MXL	-
## 42	NA19756 (M)	G A ALL, AMR, MXL	-
## 43	NA19758 (F)	A G ALL, AMR, MXL	-
## 44	NA19759 (M)	G A ALL, AMR, MXL	-
## 45	NA19761 (F)	G A ALL, AMR, MXL	-
## 46	NA19762 (M)	A A ALL, AMR, MXL	-
## 47	NA19764 (F)	A A ALL, AMR, MXL	-
## 48	NA19770 (F)	A G ALL, AMR, MXL	-
## 49	NA19771 (M)	A A ALL, AMR, MXL	-
## 50	NA19773 (F)	A A ALL, AMR, MXL	-
## 51	NA19774 (M)	A G ALL, AMR, MXL	-
## 52	NA19776 (F)	A G ALL, AMR, MXL	-
## 53	NA19777 (M)	A A ALL, AMR, MXL	-
## 54	NA19779 (F)	G A ALL, AMR, MXL	-
## 55	NA19780 (M)	A A ALL, AMR, MXL	-
## 56	NA19782 (F)	G A ALL, AMR, MXL	-
## 57	NA19783 (M)	A G ALL, AMR, MXL	-
## 58	NA19785 (F)	A A ALL, AMR, MXL	-
## 59	NA19786 (M)	G A ALL, AMR, MXL	-
## 60	NA19788 (F)	A G ALL, AMR, MXL	-
## 61	NA19789 (M)	G G ALL, AMR, MXL	-
## 62	NA19792 (M)	A A ALL, AMR, MXL	-
## 63	NA19794 (F)	G A ALL, AMR, MXL	-
## 64	NA19795 (M)	A G ALL, AMR, MXL	-
##	Mother		
## 1	-		
## 2	-		
## 3	-		
## 4	-		
## 5	-		
## 6	-		
## 7	-		
## 8	-		
## 9	-		
## 10	-		
## 11	-		
## 12	-		
## 13	-		
## 14	-		
## 15	-		
## 16	-		
## 17	-		
## 18	-		
## 19	-		
## 20	-		

```
## 21      -
## 22      -
## 23      -
## 24      -
## 25      -
## 26      -
## 27      -
## 28      -
## 29      -
## 30      -
## 31      -
## 32      -
## 33      -
## 34      -
## 35      -
## 36      -
## 37      -
## 38      -
## 39      -
## 40      -
## 41      -
## 42      -
## 43      -
## 44      -
## 45      -
## 46      -
## 47      -
## 48      -
## 49      -
## 50      -
## 51      -
## 52      -
## 53      -
## 54      -
## 55      -
## 56      -
## 57      -
## 58      -
## 59      -
## 60      -
## 61      -
## 62      -
## 63      -
## 64      -
```

```
# Use table() to answer question
mx1$Genotype..forward.strand.
```

```
## [1] "A|A" "G|G" "A|A" "G|G" "G|G" "A|G" "A|G" "A|A" "A|G" "A|A" "G|A" "A|A"
## [13] "A|A" "G|G" "A|A" "A|G" "A|G" "A|G" "A|G" "G|A" "A|G" "G|G" "G|G" "G|A"
## [25] "G|G" "A|G" "A|A" "A|A" "A|G" "A|A" "A|G" "G|A" "G|G" "A|A" "A|A" "A|A"
## [37] "G|A" "A|G" "A|G" "A|G" "A|A" "G|A" "A|G" "G|A" "G|A" "A|A" "A|A" "A|G"
## [49] "A|A" "A|A" "A|G" "A|G" "A|A" "G|A" "A|A" "G|A" "A|G" "A|A" "G|A" "A|G"
## [61] "G|G" "A|A" "G|A" "A|G"
```

```
table(mx1$Genotype..forward.strand.)/nrow(mx1)
```

```
##
##      A|A      A|G      G|A      G|G
## 0.343750 0.328125 0.187500 0.140625
```

14% of mx1 are homozygous for the asthma associated SNP (G|G).

Examine GBR data set for the same SNP to compare populations.

```
gbr <- read.csv("373522-SampleGenotypes-Homo_sapiens_Variation_Sample_rs8067378.csv")
gbr
```

```
##      Sample..Male.Female.Unknown. Genotype..forward.strand. Population.s. Father
## 1          HG00096 (M)          A|A ALL, EUR, GBR      -
## 2          HG00097 (F)          G|A ALL, EUR, GBR      -
## 3          HG00099 (F)          G|G ALL, EUR, GBR      -
## 4          HG00100 (F)          A|A ALL, EUR, GBR      -
## 5          HG00101 (M)          A|A ALL, EUR, GBR      -
## 6          HG00102 (F)          A|A ALL, EUR, GBR      -
## 7          HG00103 (M)          A|G ALL, EUR, GBR      -
## 8          HG00105 (M)          A|A ALL, EUR, GBR      -
## 9          HG00106 (F)          G|A ALL, EUR, GBR      -
## 10         HG00107 (M)          G|G ALL, EUR, GBR      -
## 11         HG00108 (M)          A|A ALL, EUR, GBR      -
## 12         HG00109 (M)          G|G ALL, EUR, GBR      -
## 13         HG00110 (F)          A|G ALL, EUR, GBR      -
## 14         HG00111 (F)          A|A ALL, EUR, GBR      -
## 15         HG00112 (M)          G|G ALL, EUR, GBR      -
## 16         HG00113 (M)          G|G ALL, EUR, GBR      -
## 17         HG00114 (M)          G|A ALL, EUR, GBR      -
## 18         HG00115 (M)          A|G ALL, EUR, GBR      -
## 19         HG00116 (M)          G|G ALL, EUR, GBR      -
## 20         HG00117 (M)          A|A ALL, EUR, GBR      -
## 21         HG00118 (F)          G|G ALL, EUR, GBR      -
## 22         HG00119 (M)          G|A ALL, EUR, GBR      -
## 23         HG00120 (F)          G|G ALL, EUR, GBR      -
## 24         HG00121 (F)          A|G ALL, EUR, GBR      -
## 25         HG00122 (F)          G|G ALL, EUR, GBR      -
## 26         HG00123 (F)          G|A ALL, EUR, GBR      -
## 27         HG00125 (F)          A|G ALL, EUR, GBR      -
## 28         HG00126 (M)          G|G ALL, EUR, GBR      -
## 29         HG00127 (F)          G|A ALL, EUR, GBR      -
## 30         HG00128 (F)          A|G ALL, EUR, GBR      -
## 31         HG00129 (M)          G|G ALL, EUR, GBR      -
## 32         HG00130 (F)          A|G ALL, EUR, GBR      -
## 33         HG00131 (M)          G|G ALL, EUR, GBR      -
## 34         HG00132 (F)          A|A ALL, EUR, GBR      -
## 35         HG00133 (F)          G|A ALL, EUR, GBR      -
## 36         HG00136 (M)          G|G ALL, EUR, GBR      -
## 37         HG00137 (F)          G|A ALL, EUR, GBR      -
## 38         HG00138 (M)          A|A ALL, EUR, GBR      -
## 39         HG00139 (M)          G|G ALL, EUR, GBR      -
```

## 40	HG00140 (M)	G A ALL, EUR, GBR	-
## 41	HG00141 (M)	G G ALL, EUR, GBR	-
## 42	HG00142 (M)	G G ALL, EUR, GBR	-
## 43	HG00143 (M)	G A ALL, EUR, GBR	-
## 44	HG00145 (M)	A A ALL, EUR, GBR	-
## 45	HG00146 (F)	A A ALL, EUR, GBR	-
## 46	HG00148 (M)	G A ALL, EUR, GBR	-
## 47	HG00149 (M)	G A ALL, EUR, GBR	-
## 48	HG00150 (F)	G A ALL, EUR, GBR	-
## 49	HG00151 (M)	G A ALL, EUR, GBR	-
## 50	HG00154 (F)	G G ALL, EUR, GBR	-
## 51	HG00155 (M)	A G ALL, EUR, GBR	-
## 52	HG00157 (M)	A A ALL, EUR, GBR	-
## 53	HG00158 (F)	A A ALL, EUR, GBR	-
## 54	HG00159 (M)	A A ALL, EUR, GBR	-
## 55	HG00160 (M)	A A ALL, EUR, GBR	-
## 56	HG00231 (F)	A G ALL, EUR, GBR	-
## 57	HG00232 (F)	G G ALL, EUR, GBR	-
## 58	HG00233 (F)	G G ALL, EUR, GBR	-
## 59	HG00234 (M)	G G ALL, EUR, GBR	-
## 60	HG00235 (F)	A A ALL, EUR, GBR	-
## 61	HG00236 (F)	A A ALL, EUR, GBR	-
## 62	HG00237 (F)	A A ALL, EUR, GBR	-
## 63	HG00238 (F)	G G ALL, EUR, GBR	-
## 64	HG00239 (F)	G A ALL, EUR, GBR	-
## 65	HG00240 (F)	G A ALL, EUR, GBR	-
## 66	HG00242 (M)	G A ALL, EUR, GBR	-
## 67	HG00243 (M)	A G ALL, EUR, GBR	-
## 68	HG00244 (M)	G A ALL, EUR, GBR	-
## 69	HG00245 (F)	A G ALL, EUR, GBR	-
## 70	HG00246 (M)	A G ALL, EUR, GBR	-
## 71	HG00250 (F)	G G ALL, EUR, GBR	-
## 72	HG00251 (M)	G A ALL, EUR, GBR	-
## 73	HG00252 (M)	G A ALL, EUR, GBR	-
## 74	HG00253 (F)	A A ALL, EUR, GBR	-
## 75	HG00254 (F)	A G ALL, EUR, GBR	-
## 76	HG00255 (F)	A G ALL, EUR, GBR	-
## 77	HG00256 (M)	A G ALL, EUR, GBR	-
## 78	HG00257 (F)	G G ALL, EUR, GBR	-
## 79	HG00258 (F)	A A ALL, EUR, GBR	-
## 80	HG00259 (F)	G A ALL, EUR, GBR	-
## 81	HG00260 (M)	G G ALL, EUR, GBR	-
## 82	HG00261 (F)	G G ALL, EUR, GBR	-
## 83	HG00262 (F)	A A ALL, EUR, GBR	-
## 84	HG00263 (F)	G A ALL, EUR, GBR	-
## 85	HG00264 (M)	A G ALL, EUR, GBR	-
## 86	HG00265 (M)	G G ALL, EUR, GBR	-
## 87	HG01334 (M)	A G ALL, EUR, GBR	-
## 88	HG01789 (M)	G A ALL, EUR, GBR	-
## 89	HG01790 (F)	G A ALL, EUR, GBR	-
## 90	HG01791 (M)	A A ALL, EUR, GBR	-
## 91	HG02215 (F)	G G ALL, EUR, GBR	-
##	Mother		
## 1	-		

## 2	-
## 3	-
## 4	-
## 5	-
## 6	-
## 7	-
## 8	-
## 9	-
## 10	-
## 11	-
## 12	-
## 13	-
## 14	-
## 15	-
## 16	-
## 17	-
## 18	-
## 19	-
## 20	-
## 21	-
## 22	-
## 23	-
## 24	-
## 25	-
## 26	-
## 27	-
## 28	-
## 29	-
## 30	-
## 31	-
## 32	-
## 33	-
## 34	-
## 35	-
## 36	-
## 37	-
## 38	-
## 39	-
## 40	-
## 41	-
## 42	-
## 43	-
## 44	-
## 45	-
## 46	-
## 47	-
## 48	-
## 49	-
## 50	-
## 51	-
## 52	-
## 53	-
## 54	-
## 55	-

```
## 56      -
## 57      -
## 58      -
## 59      -
## 60      -
## 61      -
## 62      -
## 63      -
## 64      -
## 65      -
## 66      -
## 67      -
## 68      -
## 69      -
## 70      -
## 71      -
## 72      -
## 73      -
## 74      -
## 75      -
## 76      -
## 77      -
## 78      -
## 79      -
## 80      -
## 81      -
## 82      -
## 83      -
## 84      -
## 85      -
## 86      -
## 87      -
## 88      -
## 89      -
## 90      -
## 91      -
```

```
table(gbr$Genotype..forward.strand.)/nrow(gbr)
```

```
##
##      A|A      A|G      G|A      G|G
## 0.2527473 0.1868132 0.2637363 0.2967033
```

About 29% of gbr are homozygous for the same SNP.

## Expression by Genotype Analysis

I want to read my RNA-Seq expression results into R. This file is not a CSV but rather has fields separated by space. Because of this, use `read.table()`

```
x <- read.table("rs8067378_ENSG00000172057.6.txt")
head(x)
```

```
##      sample geno      exp
## 1 HG00367   A/G 28.96038
## 2 NA20768   A/G 20.24449
## 3 HG00361   A/A 31.32628
## 4 HG00135   A/A 34.11169
## 5 NA18870   G/G 18.25141
## 6 NA11993   A/A 32.89721
```

Q13. Read this file into R and determine the sample size for each genotype and their corresponding median expression levels for each of these genotypes.

Is the mean expression different based on genotype?

```
table(x$geno)
```

```
##
## A/A A/G G/G
## 108 233 121
```

```
summary(x[x$geno == "A/A",]$exp)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##      11.40   27.02   31.25   31.82   35.92   51.52
```

```
summary(x[x$geno == "A/G",]$exp)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##       7.075  20.626  25.065  25.397  30.552  48.034
```

```
summary(x[x$geno == "G/G",]$exp)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##       6.675  16.903  20.074  20.594  24.457  33.956
```

Based on the summary statistics of each genotype, the median expression levels are 31.25 for A/A, 25.065 for A/G, and 20.074 for G/G.

**Make a summary overview figure.**

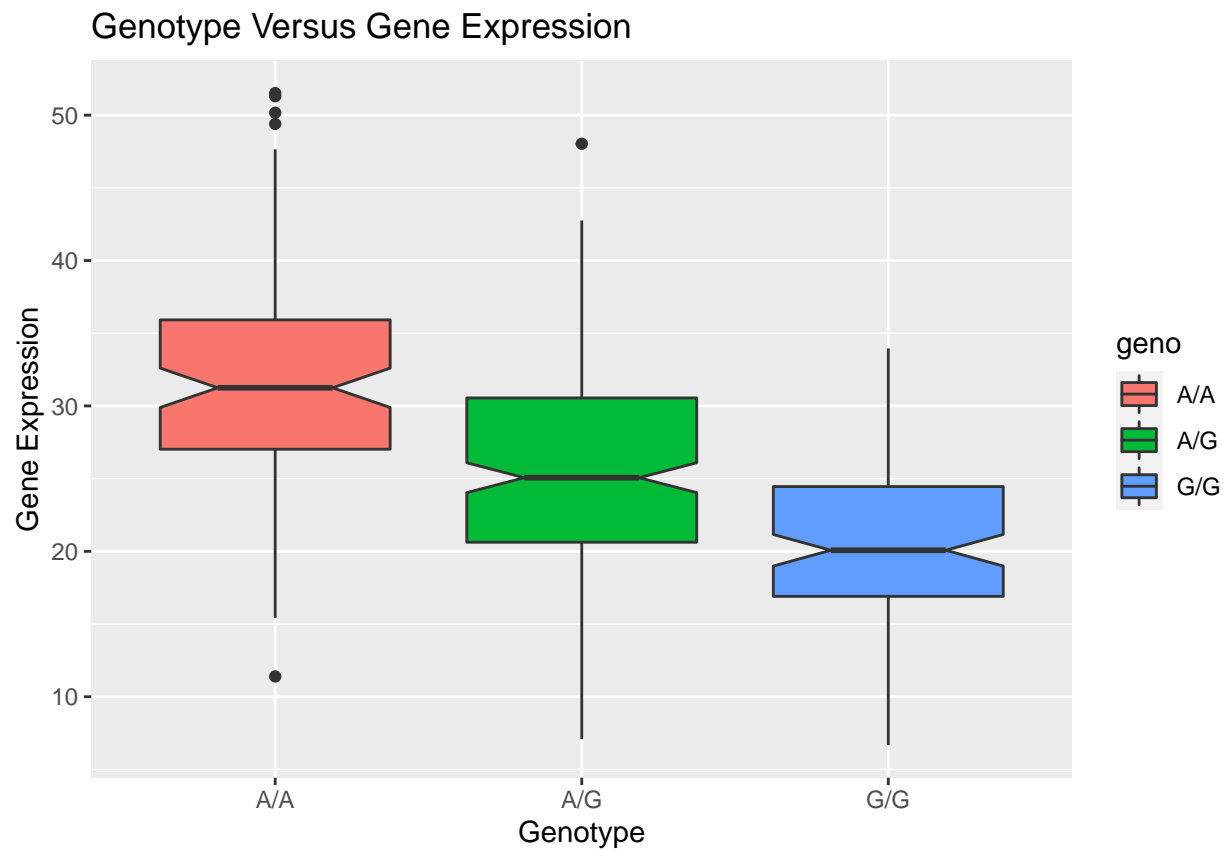
Use ggplot to make a boxplot with data.

Q14. Generate a boxplot with a box per genotype, what could you infer from the relative expression value between A/A and G/G displayed in this plot? Does the SNP effect the expression of ORMDL3?

```
library(ggplot2)
```



```
ggplot(x) + aes(x = geno, y = exp, fill = geno) + geom_boxplot(notch = TRUE) + labs(title = "Genotype Versus Gene Expression")
```



Based on the boxplot, the A/A genotype is correlated with a higher gene expression level compared to both A/G and G/G.