

## Class\_19

Lizzie (PID: 59010743)

11/28/2021

```
x <- read.csv("MXL.csv", header=TRUE)
head(x)
```

```
##      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      -
##      Mother
## 1      -
## 2      -
## 3      -
## 4      -
## 5      -
## 6      -
```

```
sum(x$Genotype..forward.strand. == "G|G")
```

```
## [1] 9
```

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

```
##      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
## 7  HG00256  A/G 31.48736
## 8  NA18498  A/A 47.64556
## 9  HG00327  G/G 17.67473
## 10 HG00115  A/G 33.85374
## 11 NA20806  A/G 16.29854
## 12 HG00278  A/G 19.73450
## 13 NA20585  A/A 30.71355
## 14 NA19137  A/G 13.96175
```

## 15	HG00235	A/A	25.44983
## 16	NA20798	A/A	34.24915
## 17	NA12546	G/G	18.55622
## 18	NA19116	A/A	35.15014
## 19	HG00381	A/G	18.40351
## 20	NA18488	G/G	23.10383
## 21	HG00259	A/G	34.21985
## 22	HG00177	A/G	23.32404
## 23	NA19214	G/G	30.94554
## 24	NA19247	A/A	24.54684
## 25	NA19098	A/G	23.18606
## 26	NA20589	A/G	18.15997
## 27	NA19207	A/A	49.39612
## 28	HG00112	G/G	21.14387
## 29	NA20518	G/G	18.39547
## 30	HG00335	A/A	28.20755
## 31	NA19119	G/G	12.02809
## 32	HG00247	G/G	17.44761
## 33	NA12155	A/G	28.03580
## 34	NA20771	A/G	30.65270
## 35	NA20758	G/G	29.82254
## 36	HG00121	A/G	20.51327
## 37	NA20759	A/A	28.56199
## 38	NA20816	A/G	29.72309
## 39	NA20542	A/G	22.50789
## 40	NA18511	A/G	31.68959
## 41	NA12249	G/G	23.01983
## 42	NA11830	A/G	28.76435
## 43	NA19159	A/G	35.85543
## 44	NA20778	A/G	37.62403
## 45	NA18908	A/G	20.54885
## 46	HG00320	G/G	13.42470
## 47	NA11843	G/G	22.65437
## 48	HG00105	A/A	51.51787
## 49	NA20588	G/G	11.07445
## 50	NA20510	G/G	28.35841
## 51	NA12342	A/G	31.04941
## 52	HG00249	A/G	18.94583
## 53	NA11894	A/A	38.10956
## 54	HG00240	A/G	32.29483
## 55	HG00132	A/A	31.13741
## 56	HG00118	G/G	28.79371
## 57	NA18520	G/G	27.08956
## 58	NA18508	A/G	27.81775
## 59	HG00353	A/G	19.89903
## 60	NA20792	A/G	48.03410
## 61	NA12234	G/G	16.11138
## 62	HG00377	A/A	39.12999
## 63	NA19143	A/G	27.90313
## 64	NA20787	A/G	36.47949
## 65	NA20513	A/G	20.03116
## 66	HG00243	A/G	29.65063
## 67	NA19172	A/A	32.44173
## 68	NA06994	A/G	34.92257

## 69	NA18510	A/G	16.71385
## 70	HG00337	A/G	16.68151
## 71	NA20503	A/G	25.71008
## 72	NA19152	G/G	26.61928
## 73	NA20761	G/G	30.18323
## 74	NA19235	A/G	11.60808
## 75	HG00382	A/G	19.30953
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## 77	NA18923	G/G	19.40790
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## 84	HG00263	A/G	35.42982
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## 96	HG00174	A/A	26.10355
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## 99	NA20520	A/A	38.77623
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## 101	HG00155	A/G	19.10420
## 102	HG00111	A/A	40.82922
## 103	NA12827	A/G	25.70962
## 104	NA18517	G/G	29.01720
## 105	NA20801	G/G	20.69333
## 106	NA20529	G/G	21.15677
## 107	NA18909	A/G	38.34531
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## 109	HG00349	G/G	18.58691
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## 115	NA20537	G/G	21.12823
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## 119	NA20783	G/G	31.42162
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##	386	HG00339	A/A	34.88439
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##	388	NA12778	A/G	23.27255
##	389	NA18861	A/A	29.29955
##	390	NA20539	A/A	32.87767
##	391	NA11931	G/G	17.91118
##	392	NA20812	A/G	28.69506



##	393	HG00120	G/G	21.09502
##	394	HG00103	A/G	26.52036
##	395	HG00328	A/G	27.49975
##	396	NA20774	A/G	24.66196
##	397	NA18873	A/G	25.81562
##	398	NA20502	A/G	22.49429
##	399	HG00143	A/G	26.88264
##	400	HG00145	A/A	43.43665
##	401	NA19225	A/A	26.56050
##	402	NA12829	A/G	28.98200
##	403	HG00137	A/G	34.31875
##	404	NA20524	A/G	26.40231
##	405	HG00379	A/A	21.87746
##	406	NA18505	A/G	21.67621
##	407	HG01334	A/G	27.56805
##	408	NA18907	A/A	33.42582
##	409	NA19204	A/A	25.38406
##	410	NA12874	A/G	16.16277
##	411	NA20506	A/G	18.28963
##	412	NA20770	A/A	18.20442
##	413	NA12776	A/G	30.55183
##	414	NA18934	A/G	20.70871
##	415	NA19153	A/G	17.66476
##	416	HG00356	A/G	22.79543
##	417	NA12283	A/G	24.03419
##	418	HG00284	A/G	18.02351
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##	420	HG00104	A/A	21.62336
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##	422	NA11840	A/G	27.54976
##	423	HG00383	A/G	14.79717
##	424	NA20786	A/A	35.80093
##	425	NA20802	A/G	25.34921
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##	431	NA12287	A/G	22.43773
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##	439	NA12045	A/G	30.80067
##	440	NA19257	A/G	33.95134
##	441	NA12413	A/G	39.43243
##	442	HG00159	A/A	23.99631
##	443	NA20811	A/A	11.39643
##	444	HG00149	A/G	23.91465
##	445	NA19223	A/G	20.97560
##	446	NA07346	G/G	16.56929

```
## 447 NA20536 A/G 20.02507
## 448 HG01791 A/A 35.24632
## 449 HG00271 A/G 33.44170
## 450 HG00373 A/G 17.32813
## 451 HG00182 A/A 23.38376
## 452 HG00110 A/G 32.61856
## 453 NA20819 A/G 36.77906
## 454 HG00154 G/G 16.69044
## 455 HG00330 A/G 16.84776
## 456 NA12750 A/A 34.94395
## 457 HG00233 G/G 25.08880
## 458 HG00131 G/G 32.78519
## 459 HG00108 A/A 31.92036
## 460 HG00119 A/G 31.53069
## 461 NA19130 A/A 44.27738
## 462 HG00239 A/G 23.18250
```

```
summary(dat)
```

```
##      sample          geno          exp
## Length:462      Length:462      Min.   : 6.675
## Class :character Class :character 1st Qu.:20.004
## Mode  :character Mode  :character Median :25.116
##                                     Mean  :25.640
##                                     3rd Qu.:30.779
##                                     Max.   :51.518
```

```
unique(dat$geno)
```

```
## [1] "A/G" "A/A" "G/G"
```

Q13

```
sum(dat$geno == "A/G") # there are 233 A/G
```

```
## [1] 233
```

```
sum(dat$geno == "G/G") # there are 121 G/G
```

```
## [1] 121
```

```
sum(dat$geno == "A/A") # there are 108 A/A
```

```
## [1] 108
```

```
A_G <- dat$exp[dat$geno == "A/G"]
A_A <- dat$exp[dat$geno == "A/A"]
G_G <- dat$exp[dat$geno == "G/G"]
```

```
summary(A_G) #median level of expression for A/G is 25.065
```

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

```
summary(A_A) #median level of expression for A/A is 31.25
```

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

```
summary(G_G) #median level of expression for G/G is 20.074
```

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

Q14

```
mycol <- c("blueviolet", "cyan3", "aquamarine2")
boxplot(A_G,A_A,G_G,notch= TRUE, col = mycol , ylab= "Expression", xlab= "Genotype")
```

