

class10

Phoebe LI

2/18/2022

genotype data from 1000 Genomes

Let us determine the frequency of different alleles MXL population

```
# import the csv file
mxl <- read.csv("373531-SampleGenotypes-Homo_sapiens_Variation_Sample_rs8067378.csv")
head(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 -
## Mother
## 1 -
## 2 -
## 3 -
## 4 -
## 5 -
## 6 -
```

```
mxl$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(mxl$Genotype..forward.strand.)/nrow(mxl)
```

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

Downloaded another data set British in England and Scotland

```
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
```

```
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
```

```
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
```

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

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

```
x[x$geno=="G/G",]
```

```
##      sample geno      exp
## 5   NA18870  G/G 18.25141
## 9   HG00327  G/G 17.67473
## 17  NA12546  G/G 18.55622
## 20  NA18488  G/G 23.10383
## 23  NA19214  G/G 30.94554
## 28  HG00112  G/G 21.14387
## 29  NA20518  G/G 18.39547
## 31  NA19119  G/G 12.02809
## 32  HG00247  G/G 17.44761
## 35  NA20758  G/G 29.82254
## 41  NA12249  G/G 23.01983
## 46  HG00320  G/G 13.42470
## 47  NA11843  G/G 22.65437
## 49  NA20588  G/G 11.07445
## 50  NA20510  G/G 28.35841
## 56  HG00118  G/G 28.79371
## 57  NA18520  G/G 27.08956
## 61  NA12234  G/G 16.11138
## 72  NA19152  G/G 26.61928
## 73  NA20761  G/G 30.18323
## 77  NA18923  G/G 19.40790
## 79  HG00238  G/G 19.52301
## 85  NA12058  G/G 26.56808
## 89  HG00129  G/G 17.34076
## 92  HG00183  G/G 10.74263
## 93  HG00109  G/G 16.66051
## 104 NA18517  G/G 29.01720
## 105 NA20801  G/G 20.69333
## 106 NA20529  G/G 21.15677
## 109 HG00349  G/G 18.58691
## 110 HG00234  G/G 19.04962
## 111 NA19248  G/G 22.81974
## 114 NA12813  G/G 32.01142
## 115 NA20537  G/G 21.12823
## 117 HG00332  G/G 18.61268
## 118 HG00152  G/G 19.37093
## 119 NA20783  G/G 31.42162
## 128 HG00185  G/G 16.67764
## 132 NA20531  G/G 19.08659
## 135 HG00277  G/G 21.55001
## 140 HG00336  G/G  8.29591
## 143 NA20581  G/G 12.58869
## 150 NA20538  G/G 17.34109
## 153 NA20814  G/G 28.23642
## 156 NA19171  G/G 19.99979
## 159 HG00141  G/G 25.55413
## 163 NA19190  G/G 24.45672
```

##	166	NA10851	G/G	23.53572
##	170	HG00116	G/G	22.48273
##	171	NA12272	G/G	14.66862
##	172	NA19096	G/G	33.95602
##	175	NA19236	G/G	18.26466
##	178	HG00345	G/G	16.06661
##	190	HG00156	G/G	17.32504
##	193	HG00282	G/G	19.14766
##	194	HG00343	G/G	12.57599
##	195	HG00139	G/G	22.28749
##	199	HG00232	G/G	17.29261
##	201	HG00122	G/G	24.18141
##	207	NA19149	G/G	16.07627
##	211	HG00189	G/G	14.80495
##	218	HG00126	G/G	23.46573
##	224	HG00265	G/G	28.97074
##	225	HG00378	G/G	27.78837
##	232	NA20796	G/G	23.92355
##	233	NA12399	G/G	9.55902
##	239	HG00099	G/G	12.35836
##	241	NA19114	G/G	22.53910
##	247	NA19210	G/G	21.98118
##	250	HG00276	G/G	16.40569
##	253	HG00181	G/G	25.21931
##	254	HG00346	G/G	24.32857
##	259	HG00142	G/G	19.42882
##	261	HG00315	G/G	26.56993
##	267	HG00250	G/G	13.34557
##	268	NA20769	G/G	16.60507
##	271	NA19144	G/G	24.85165
##	272	NA12815	G/G	21.56943
##	280	NA19175	G/G	23.95528
##	283	NA18519	G/G	16.18962
##	285	NA20535	G/G	22.53720
##	287	HG00260	G/G	26.04123
##	288	HG00372	G/G	6.67482
##	292	HG00261	G/G	20.07363
##	293	HG00273	G/G	19.76527
##	299	HG00358	G/G	18.50772
##	307	NA19121	G/G	20.14146
##	308	NA20515	G/G	18.07151
##	314	NA10847	G/G	6.94390
##	316	NA12400	G/G	22.14277
##	319	HG00342	G/G	14.23742
##	330	HG00136	G/G	19.85388
##	340	NA20765	G/G	27.73467
##	344	NA18502	G/G	19.02064
##	351	NA20772	G/G	14.49816
##	355	HG00257	G/G	26.78940
##	356	NA18486	G/G	20.84709
##	357	HG00188	G/G	10.77316
##	361	HG00280	G/G	12.82128
##	362	HG00308	G/G	16.90256
##	364	NA18910	G/G	29.60045

```
## 369 HG00281 G/G 14.81945
## 373 NA12275 G/G 17.46326
## 375 HG00351 G/G 23.26922
## 376 HG00186 G/G 21.39806
## 378 HG00275 G/G 18.06320
## 379 HG00325 G/G 15.91528
## 380 NA19118 G/G 24.80823
## 381 HG00124 G/G 26.04514
## 383 HG02215 G/G 18.28089
## 385 HG00134 G/G 23.24907
## 391 NA11931 G/G 17.91118
## 393 HG00120 G/G 21.09502
## 421 NA20582 G/G 24.74366
## 428 NA12889 G/G 27.40521
## 435 NA12006 G/G 24.85772
## 436 NA19108 G/G 23.08482
## 446 NA07346 G/G 16.56929
## 454 HG00154 G/G 16.69044
## 457 HG00233 G/G 25.08880
## 458 HG00131 G/G 32.78519
```

```
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$geno=="A/A"],$exp)
```

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

```
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
```

```
library(ggplot2)
ggplot(x)+
  aes(x= geno, y= exp)+
  geom_boxplot()+
  labs(x= "Genotype", y= "Expression")
```


