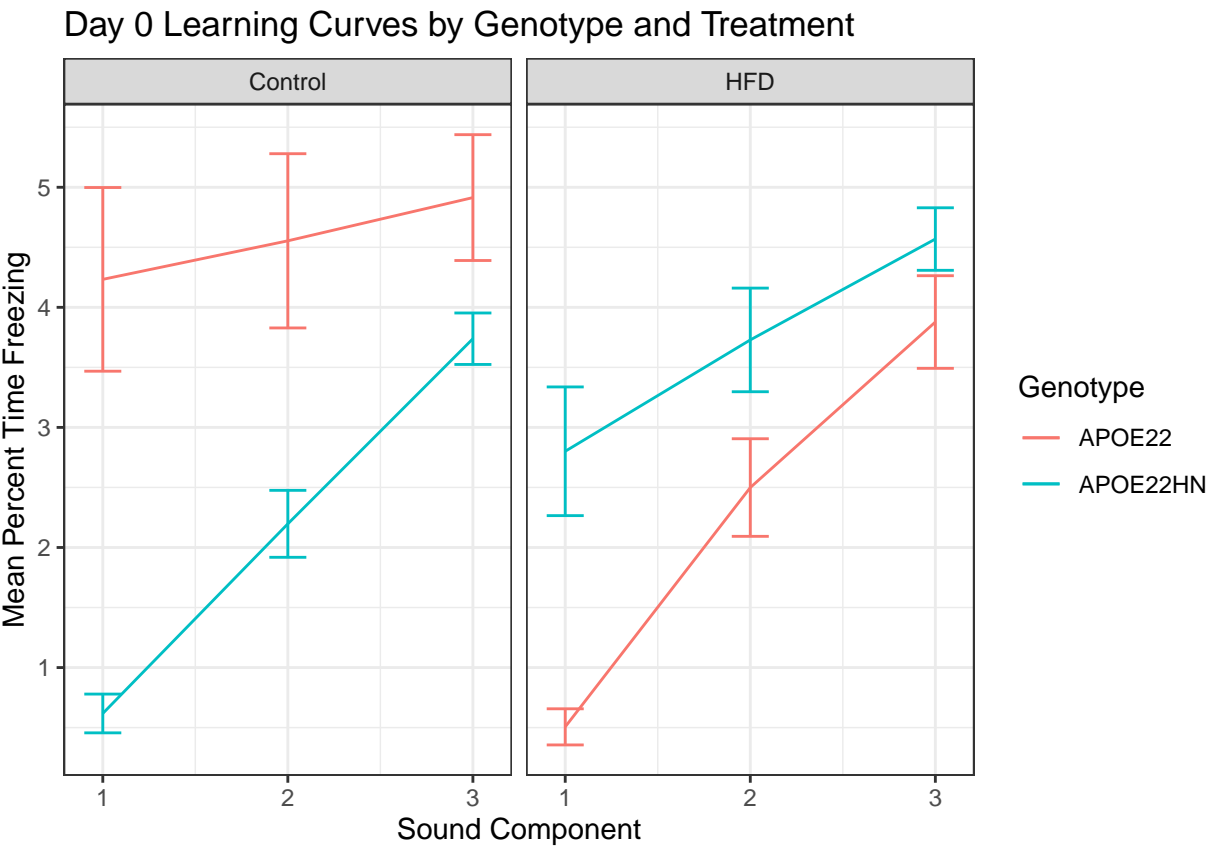


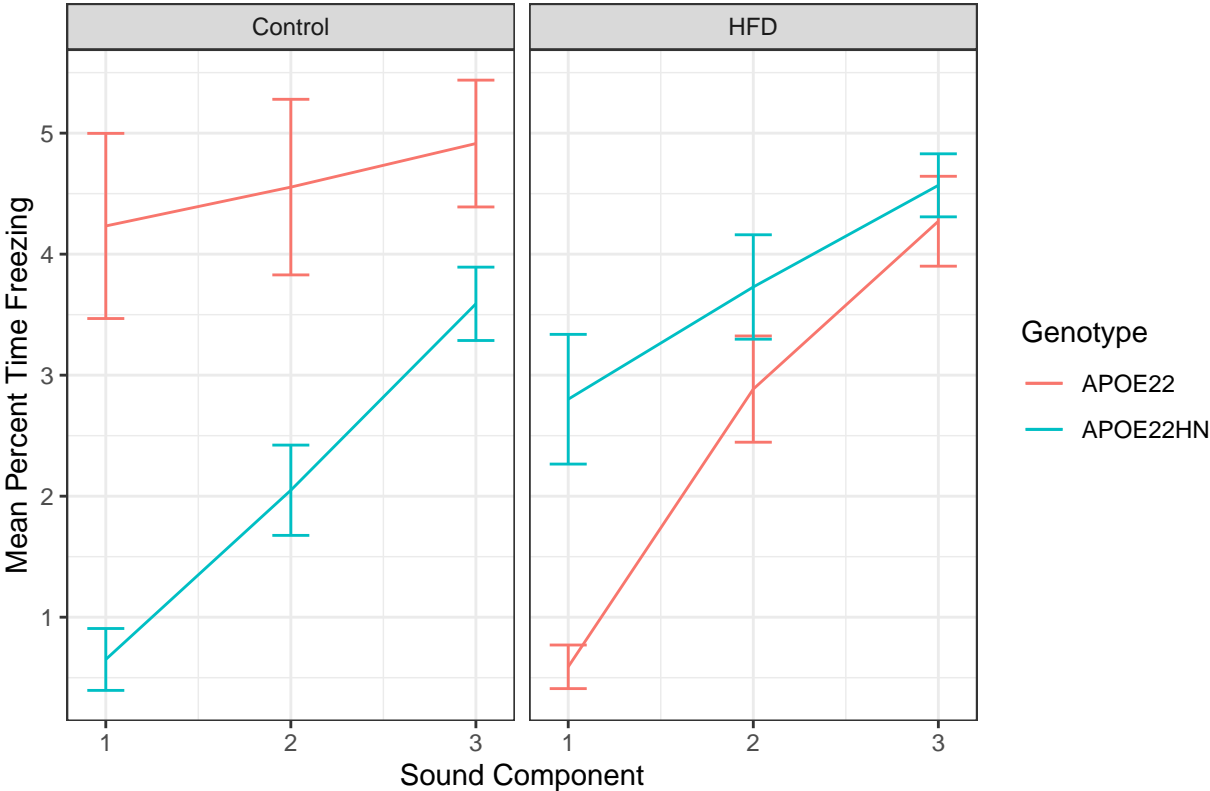
analysis

Daniel Jin

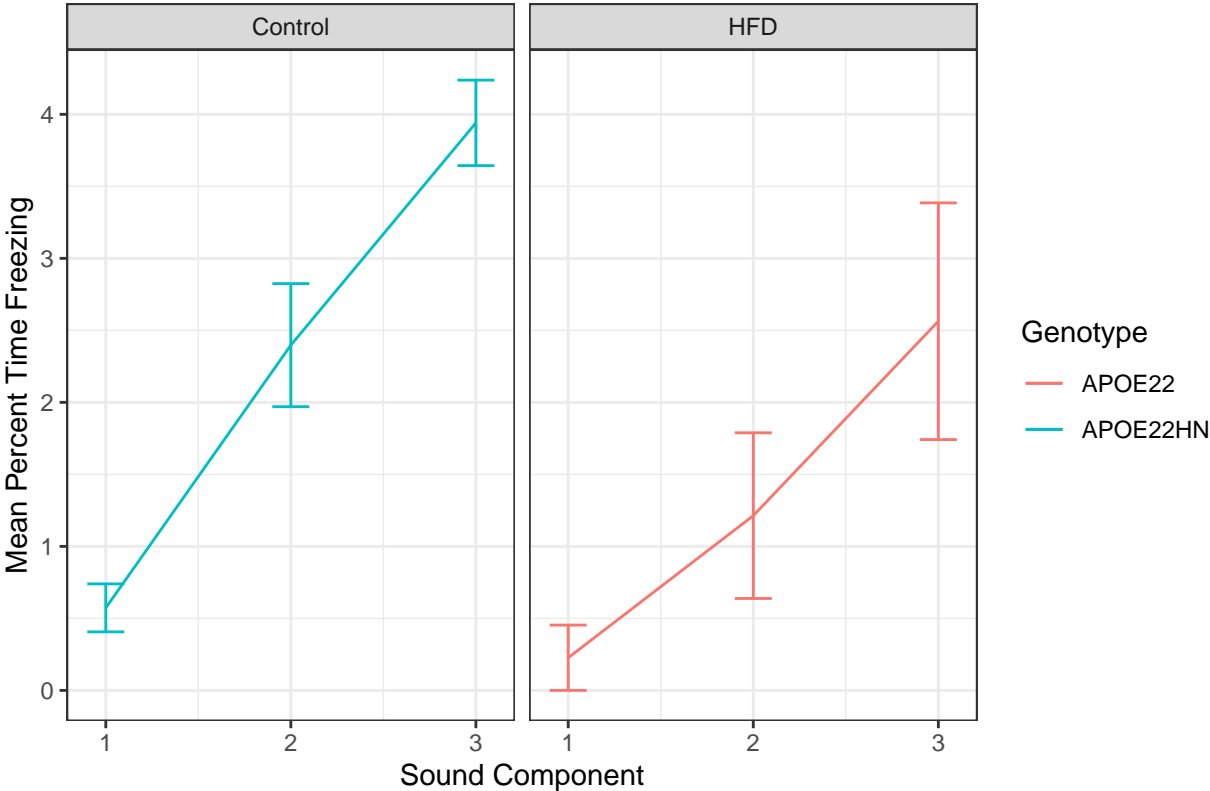
2022-08-01



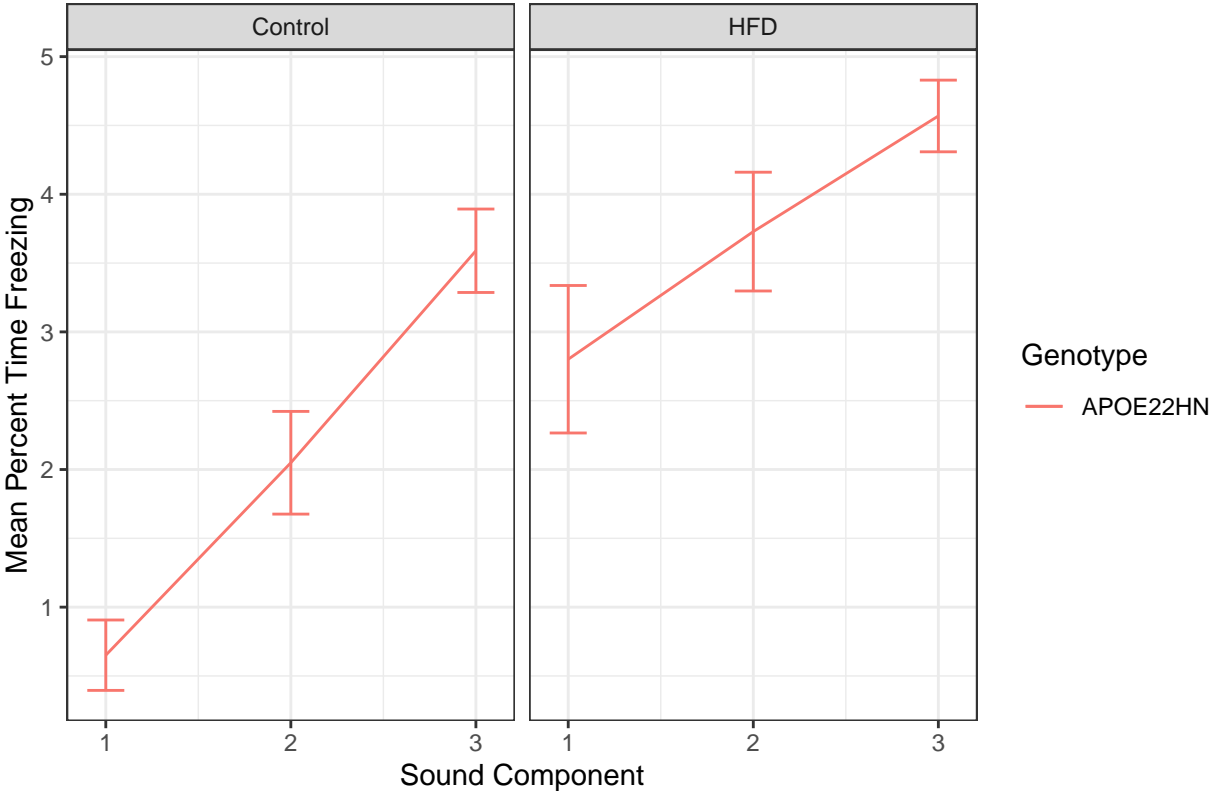
Day 0 12 Months Old Learning Curves by Genotype and Treatment



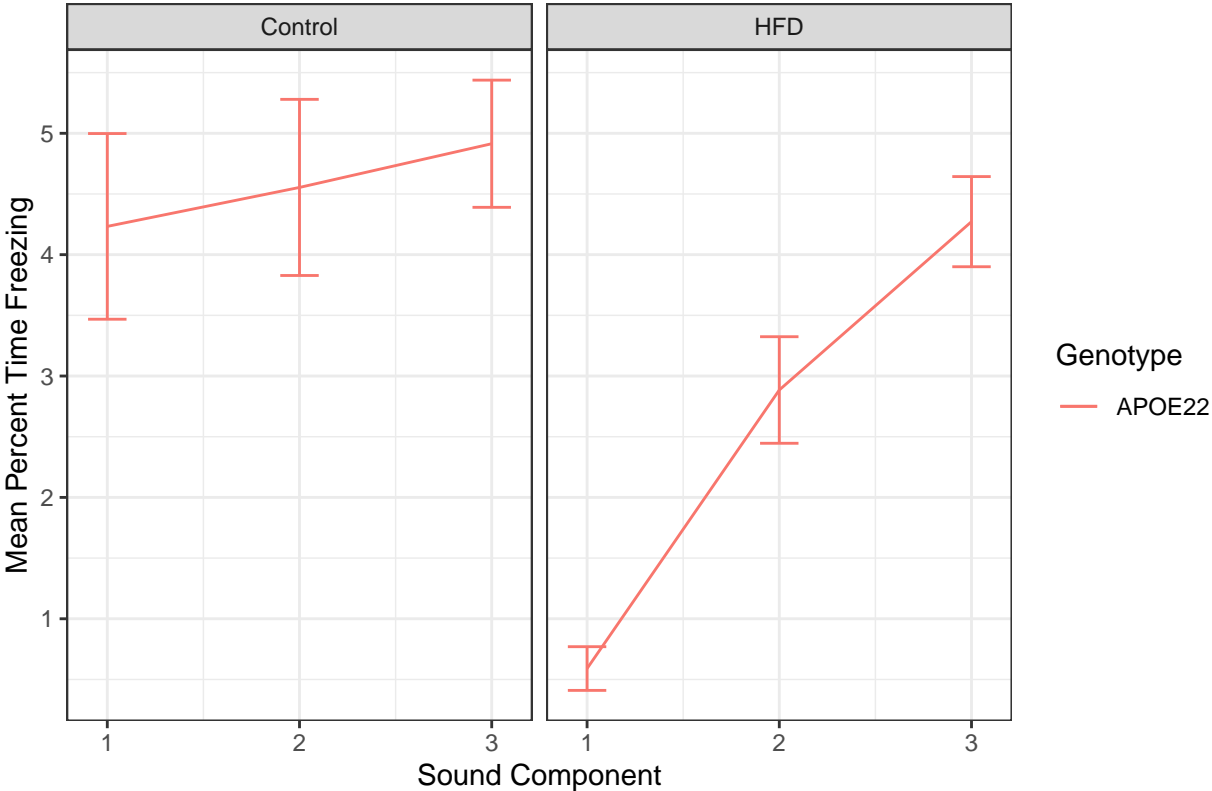
Day 0 18 Months Old Learning Curves by Genotype and Treatment



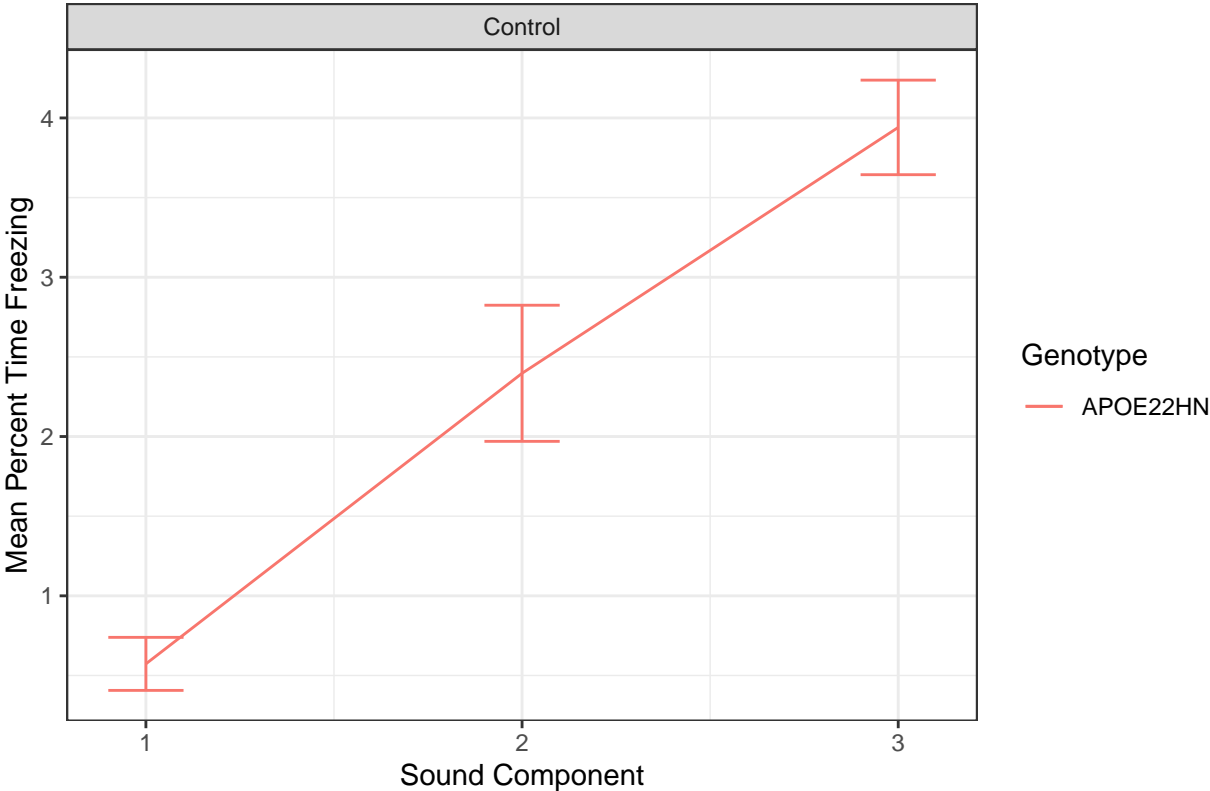
Day 0 12 Months Old Learning Curves by HN Genotype and Treatment



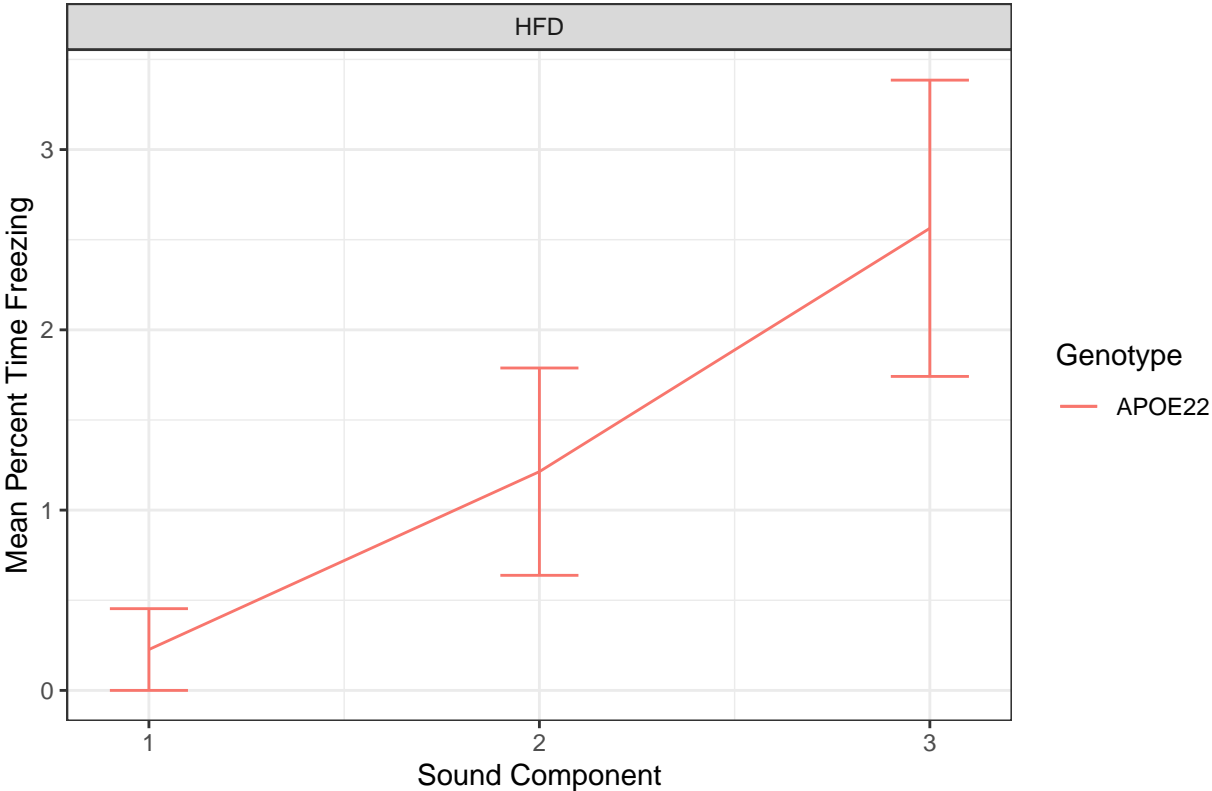
Day 0 12 Months Old Learning Curves by nonHN Genotype and Treatment



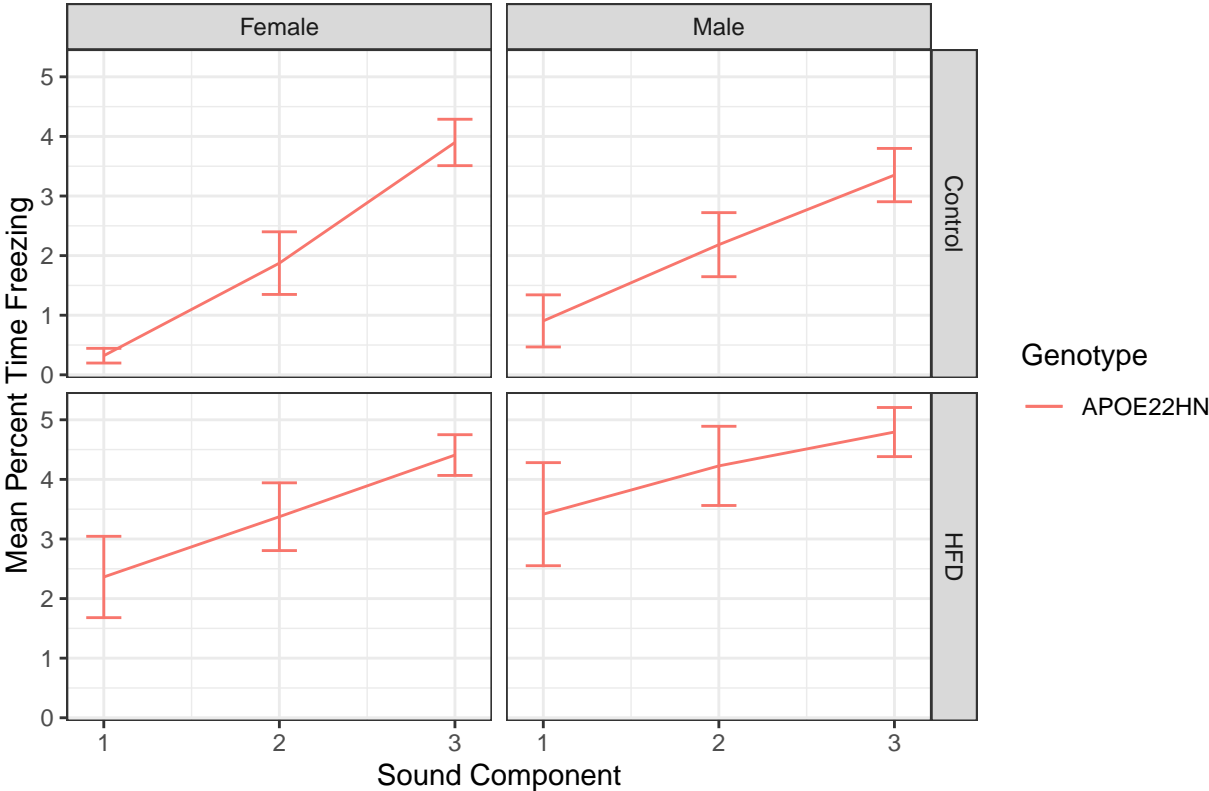
Day 0 18 Months Old Learning Curves by HN Genotype and Treatment



Day 0 18 Months Old Learning Curves by nonHN Genotype and Treatment

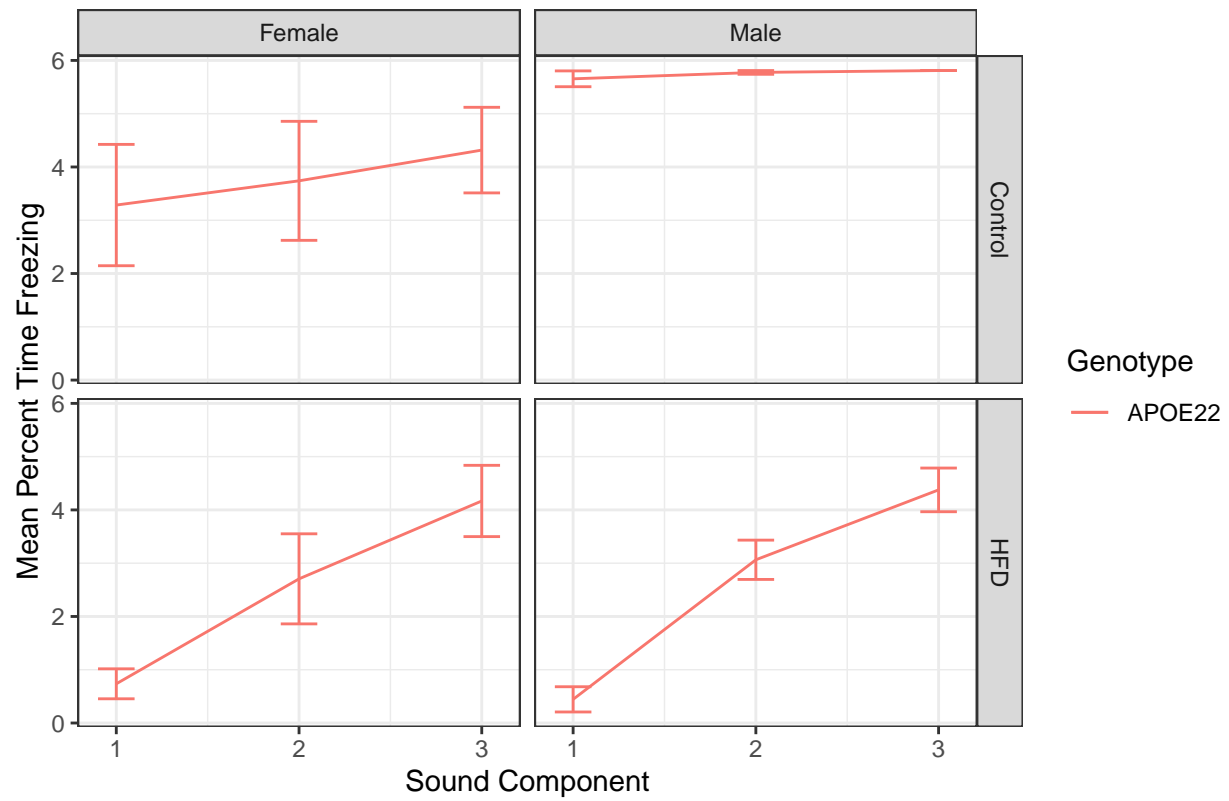


Day 0 12 Months Old HNs by Genotype, Sex, and Treatment

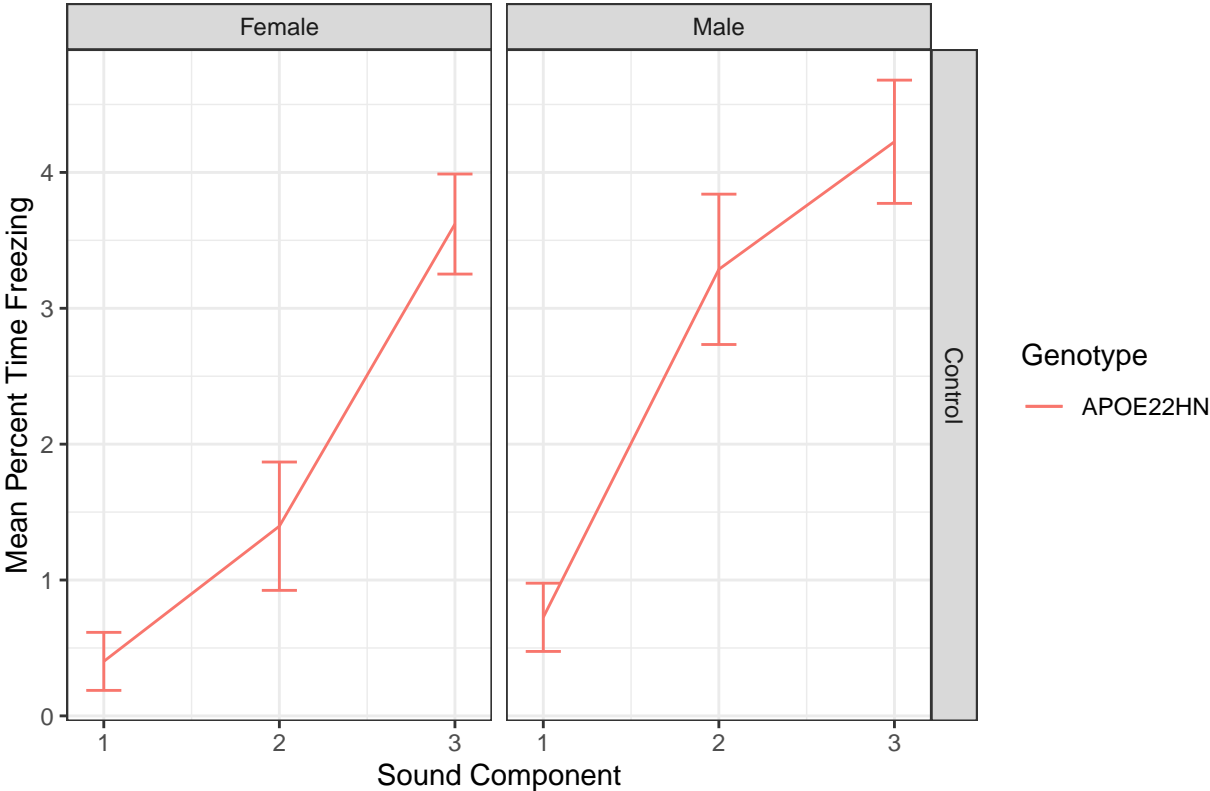




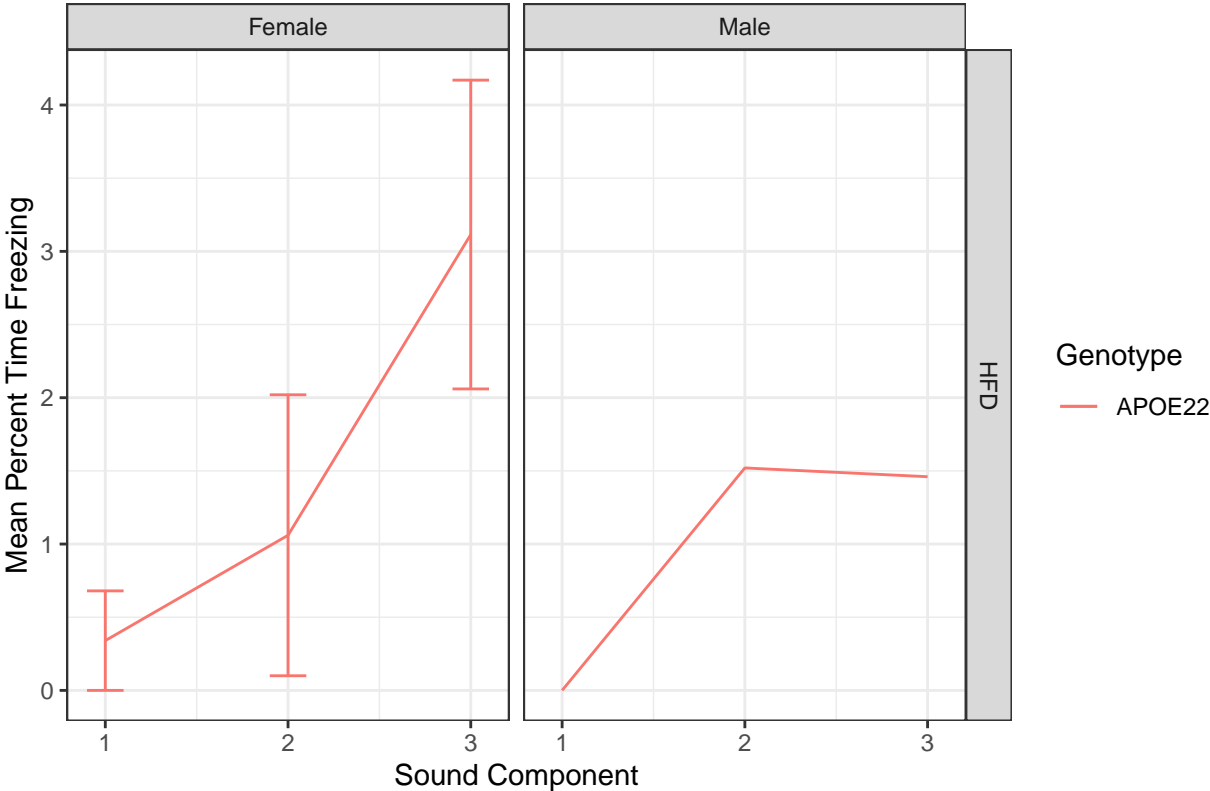
Day 0 12 Months Old nonHNs by Genotype, Sex, and Treatment



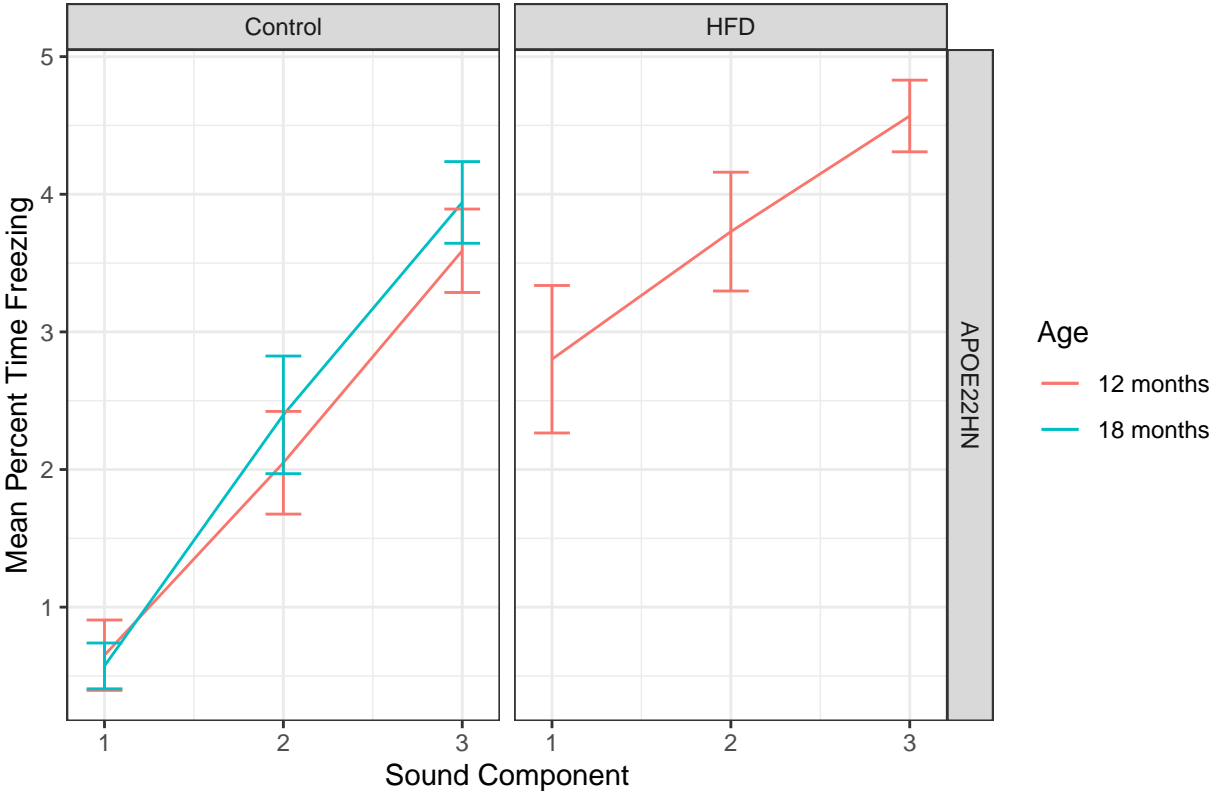
Day 0 18 Months Old HNs by Genotype, Sex, and Treatment



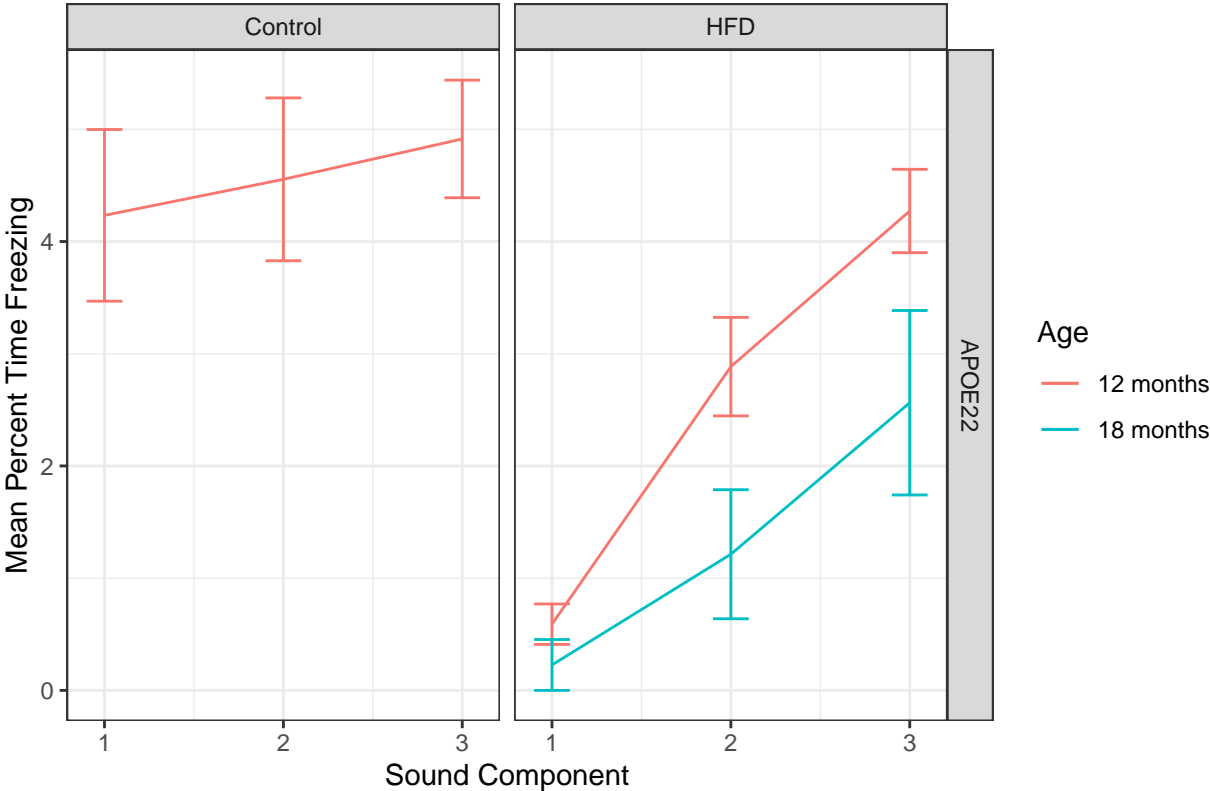
Day 0 18 Months Old nonHNs by Genotype, Sex, and Treatment



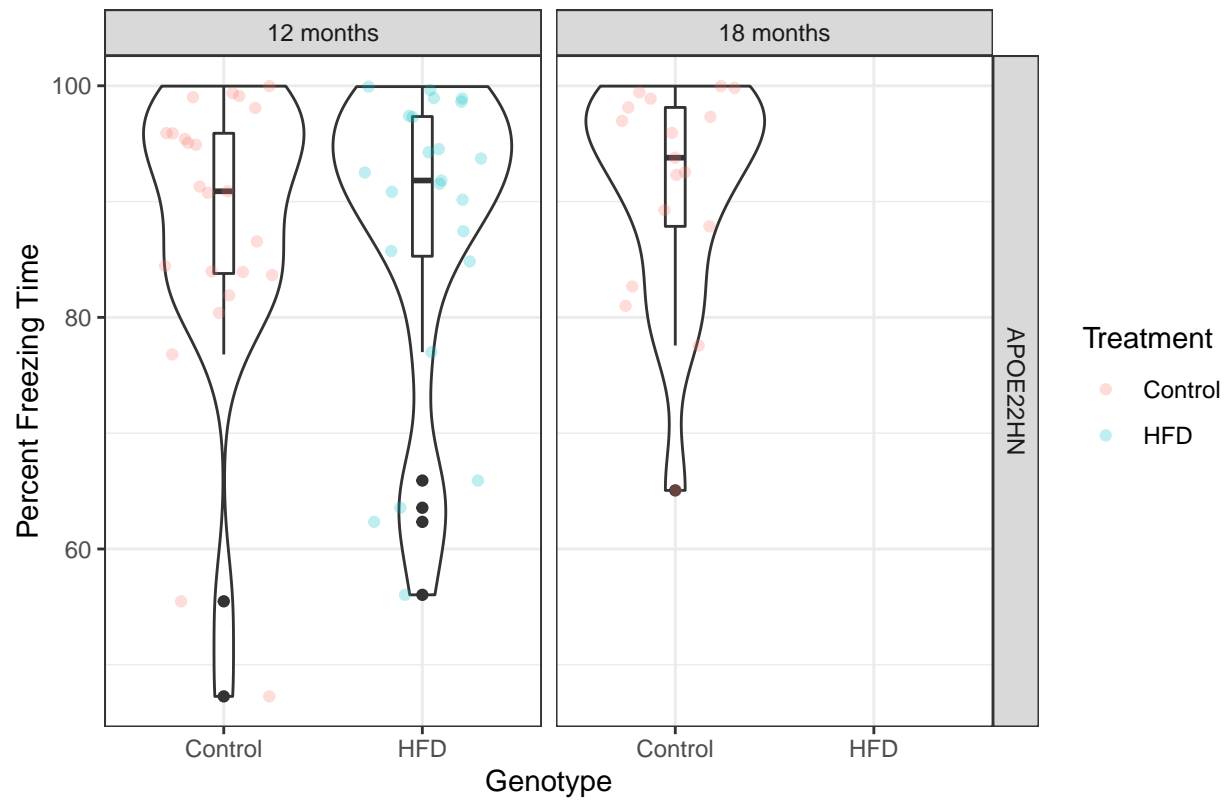
Day 0 Learning Curves by Age, HN Genotype, and Treatment



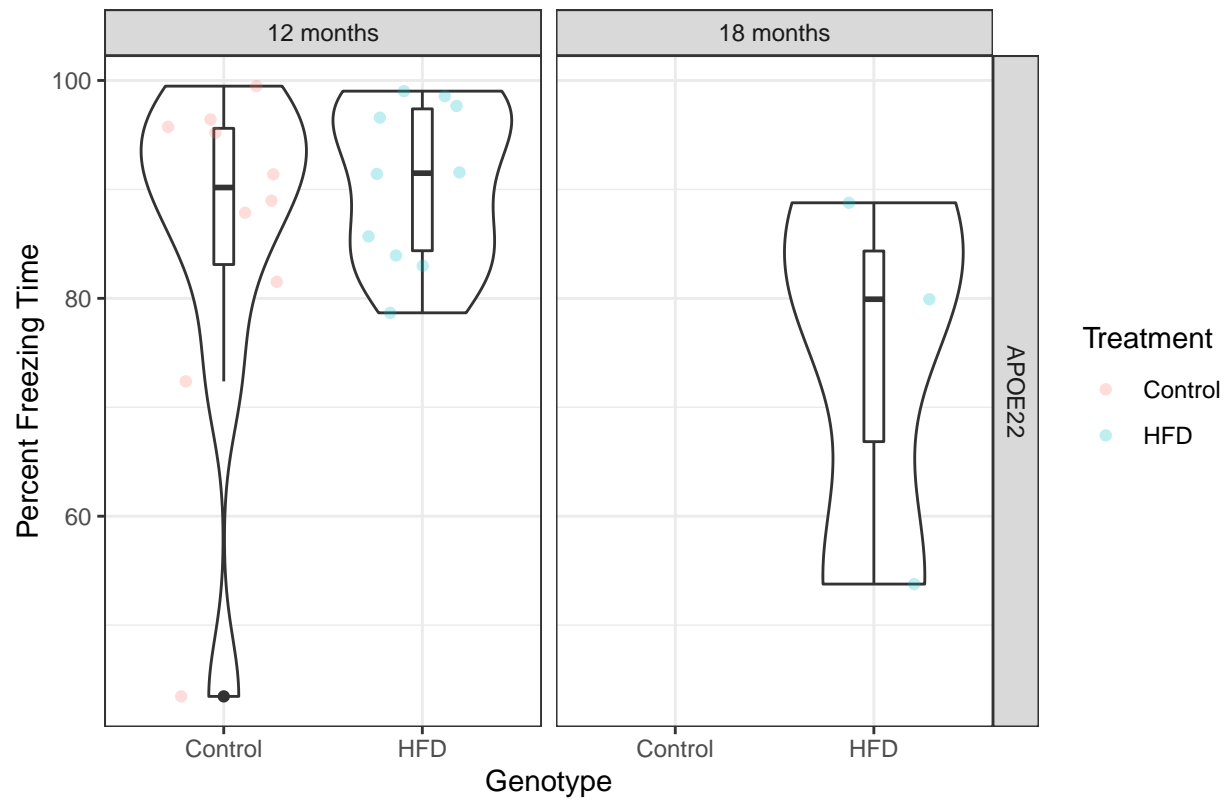
Day 0 Learning Curves by Age, nonHN Genotype, and Treatment



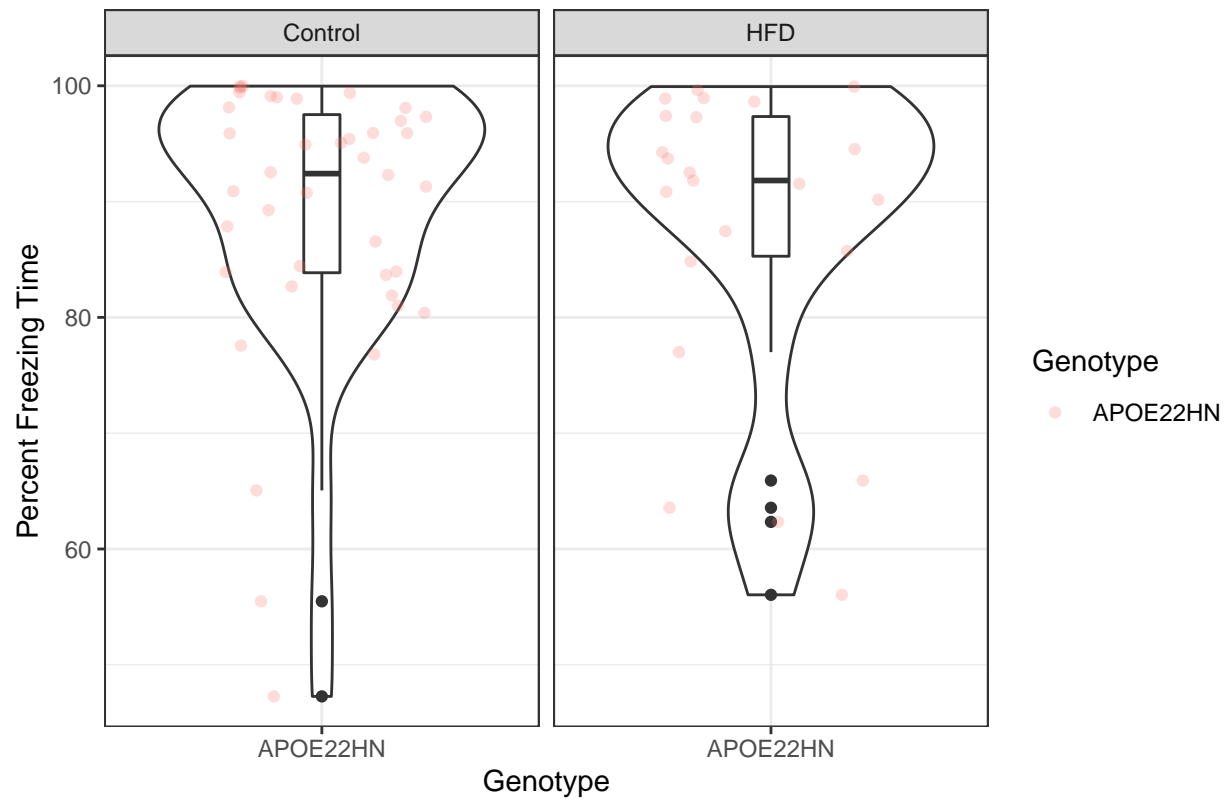
Day 1 Contextual Test by Treatment, Age, and HN Genotype



Day 1 Contextual Test by Treatment, Age, and nonHN Genotype

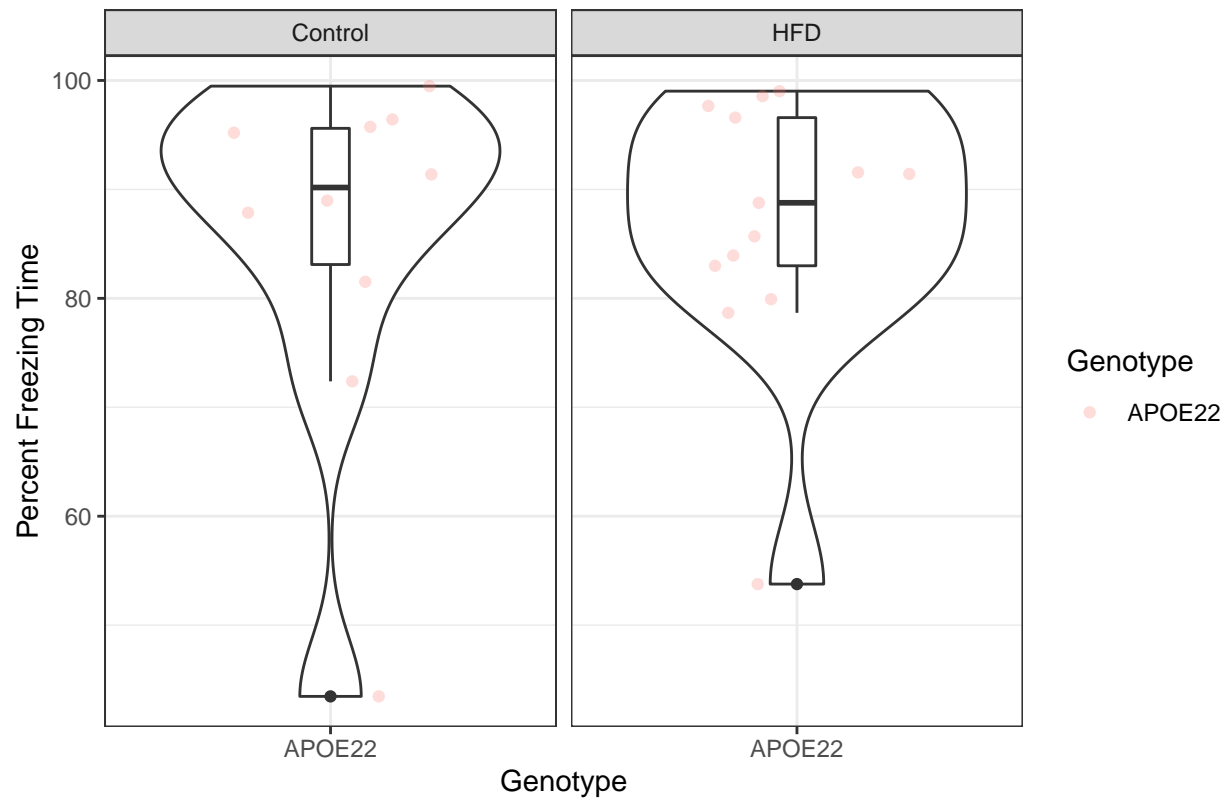


# Day 1 Contextual Test All Ages by Treatment and HN Genotype

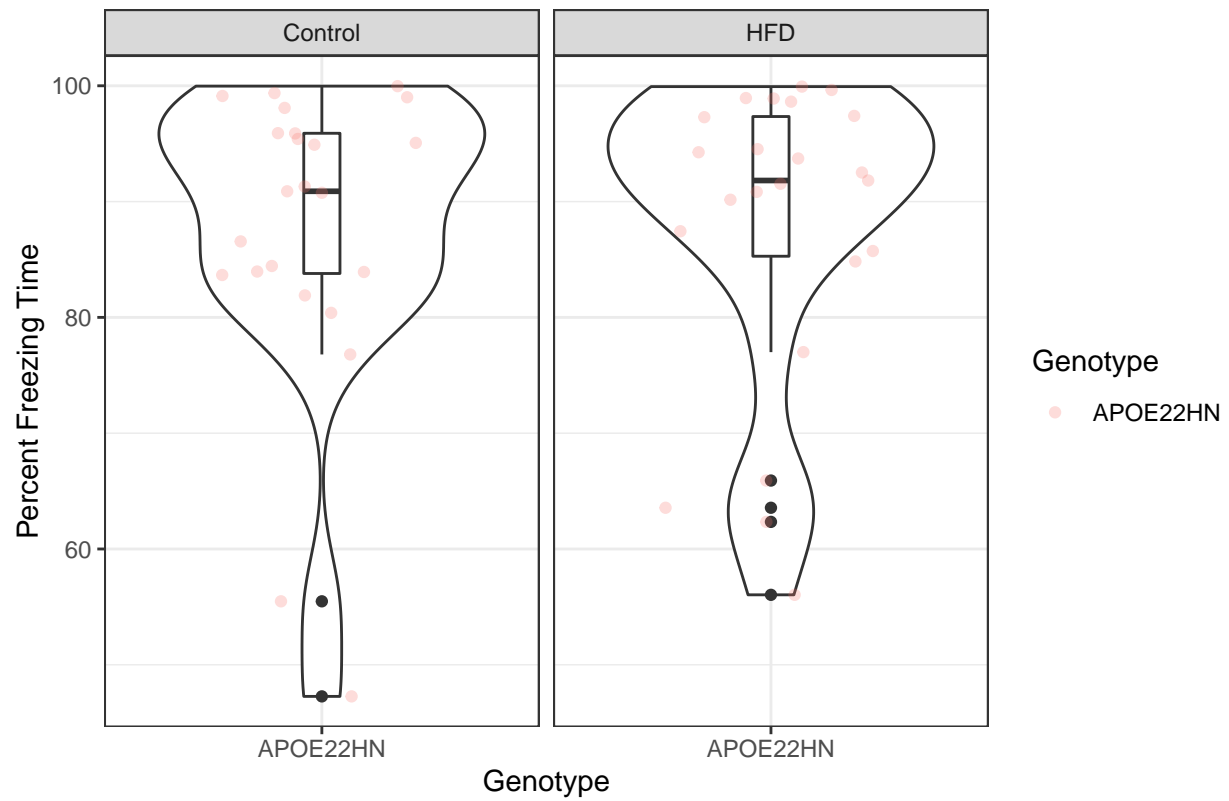




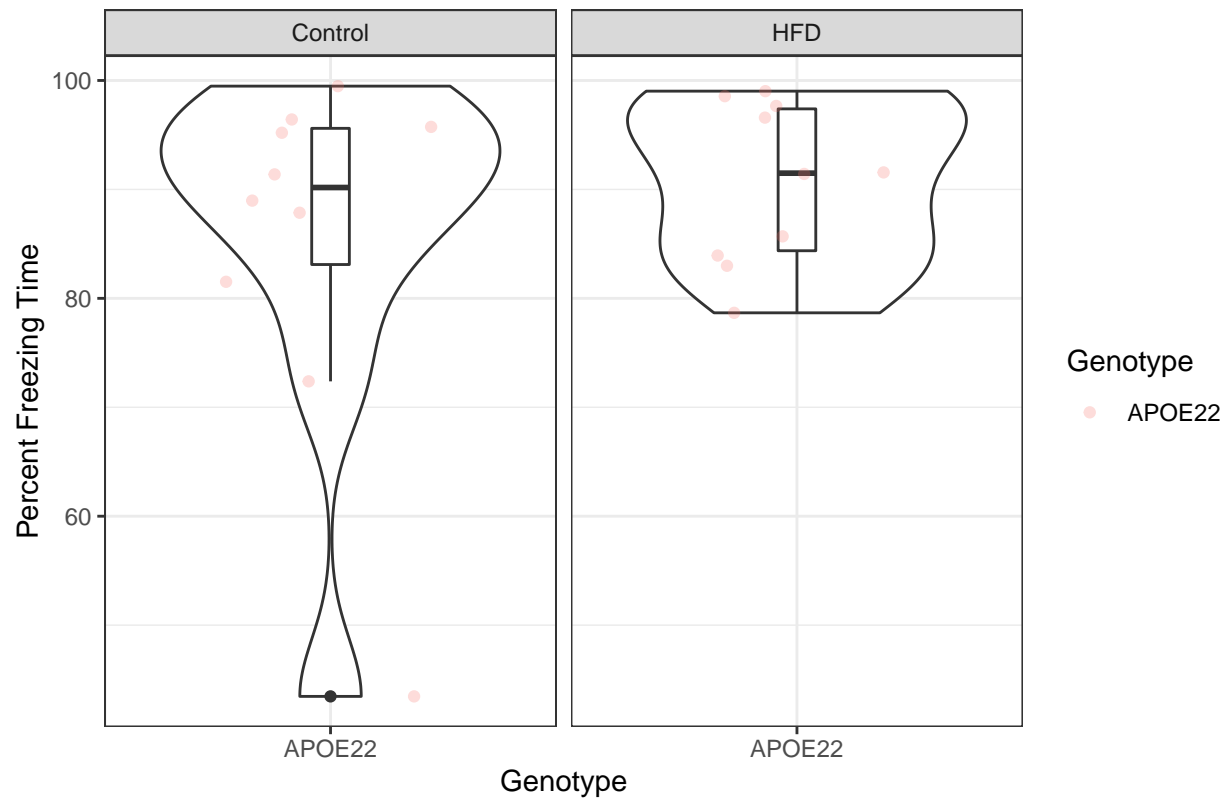
# Day 1 Contextual Test All Ages by Treatment and nonHN Genotype



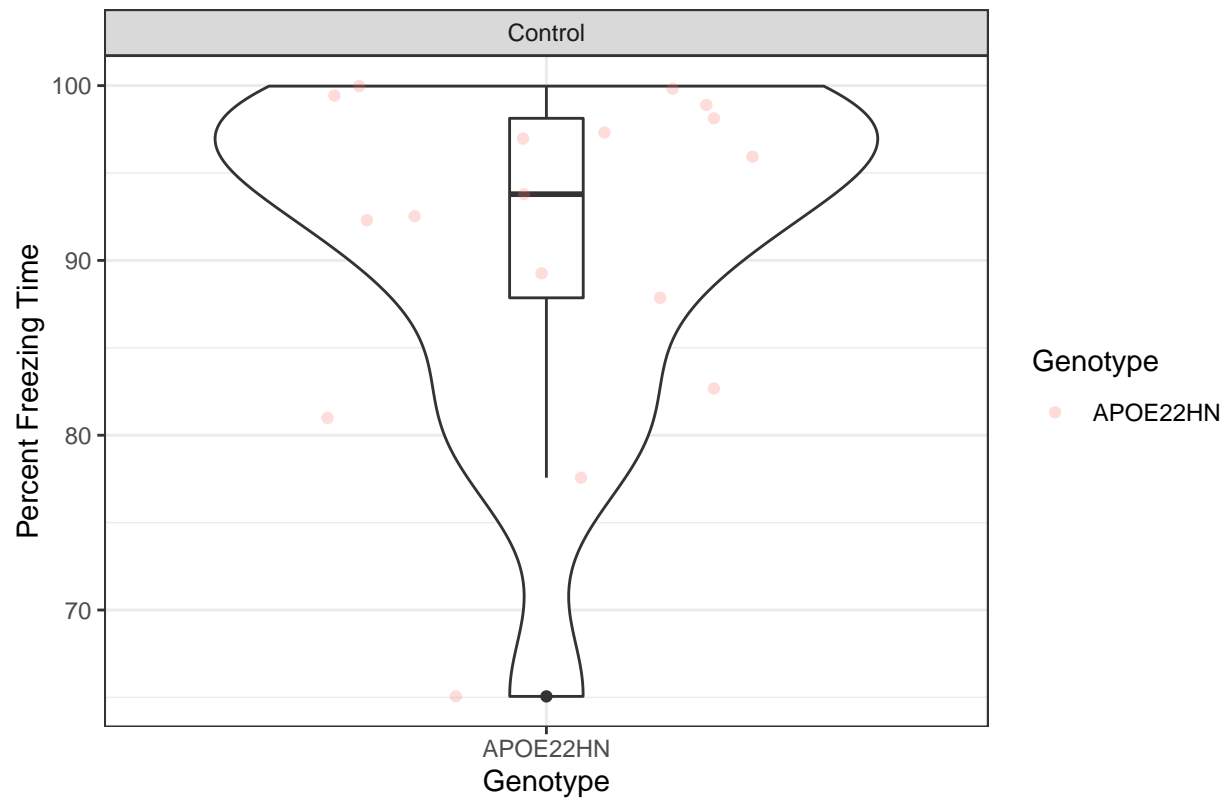
Day 1 Contextual Test 12 Months by Treatment and HN Genotype



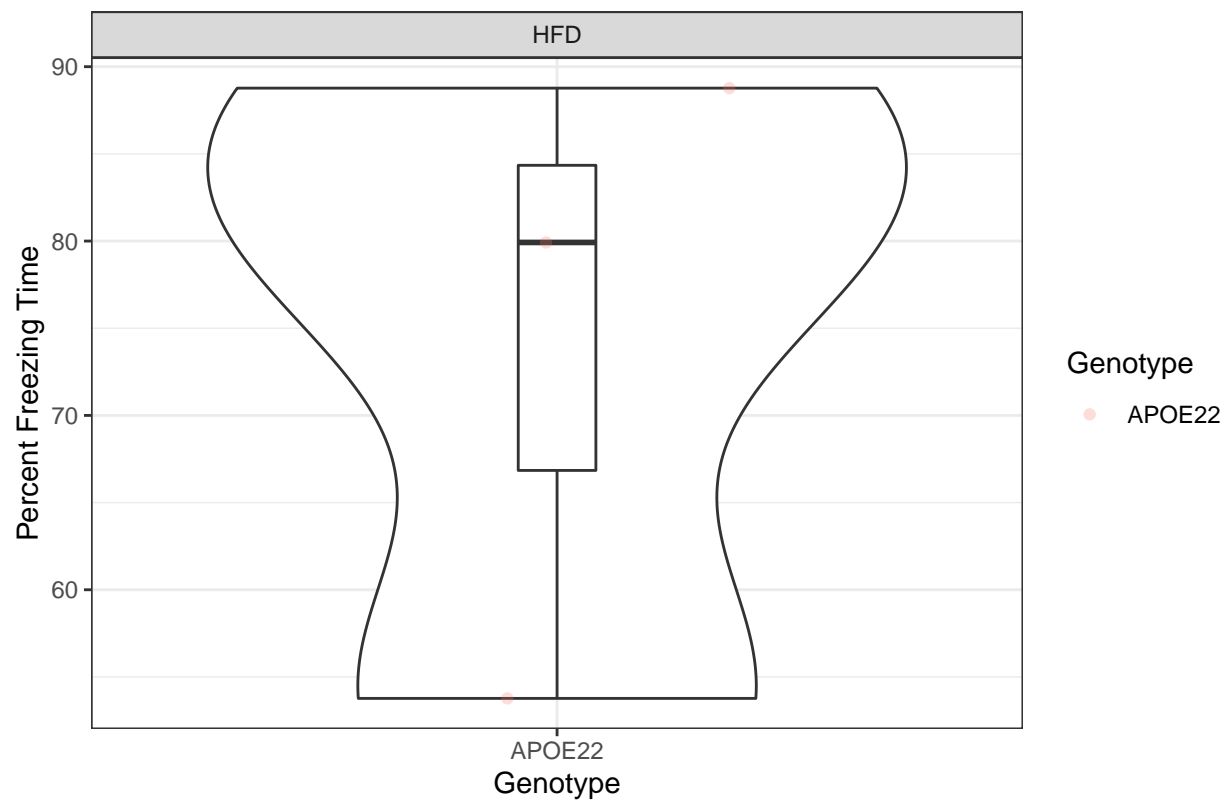
Day 1 Contextual Test 12 Months by Treatment and nonHN Genotype



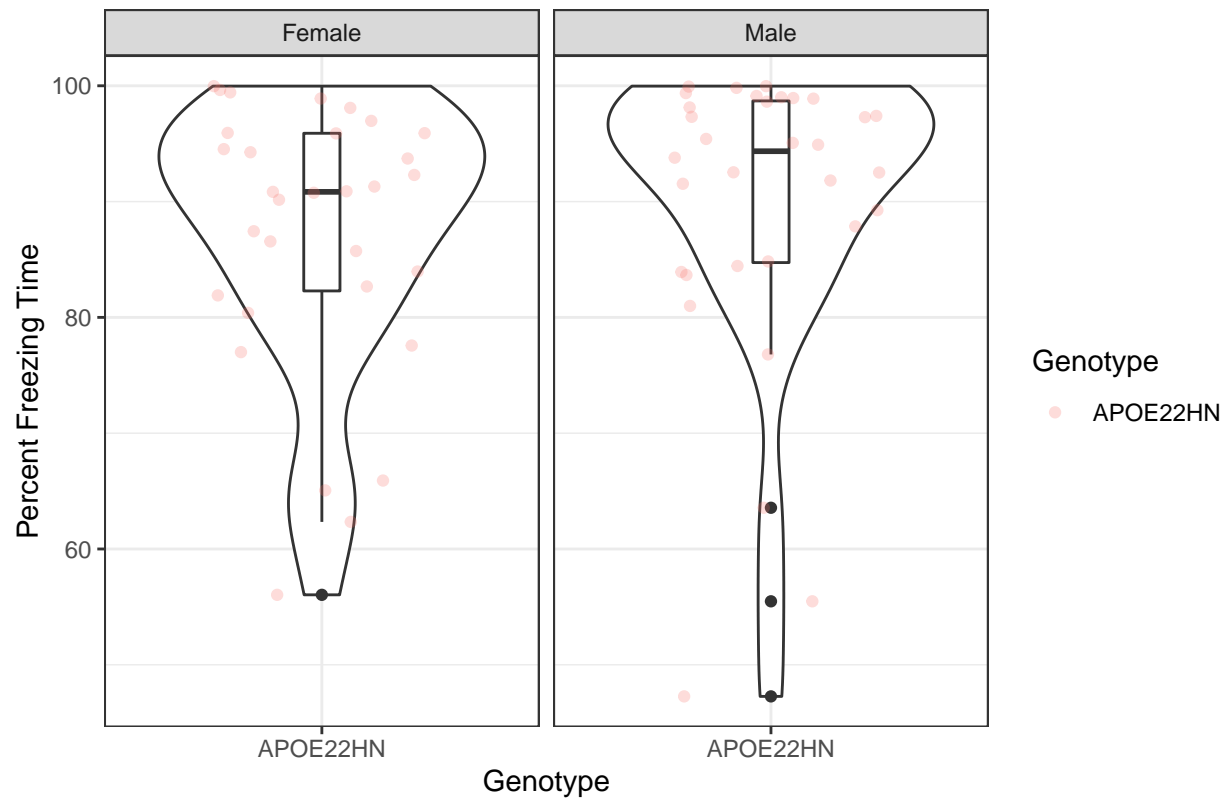
# Day 1 Contextual Test 18 Months by Treatment and HN Genotype



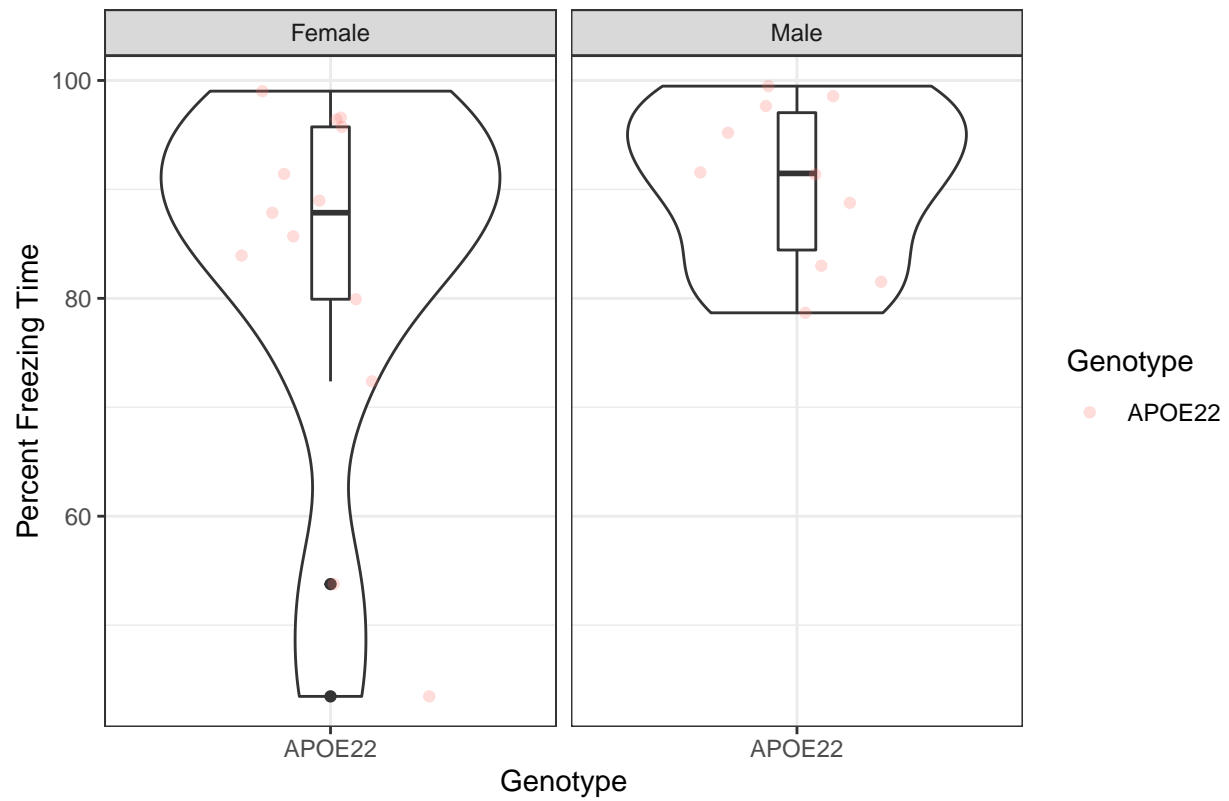
# Day 1 Contextual Test 18 Months by Treatment and nonHN Genotype



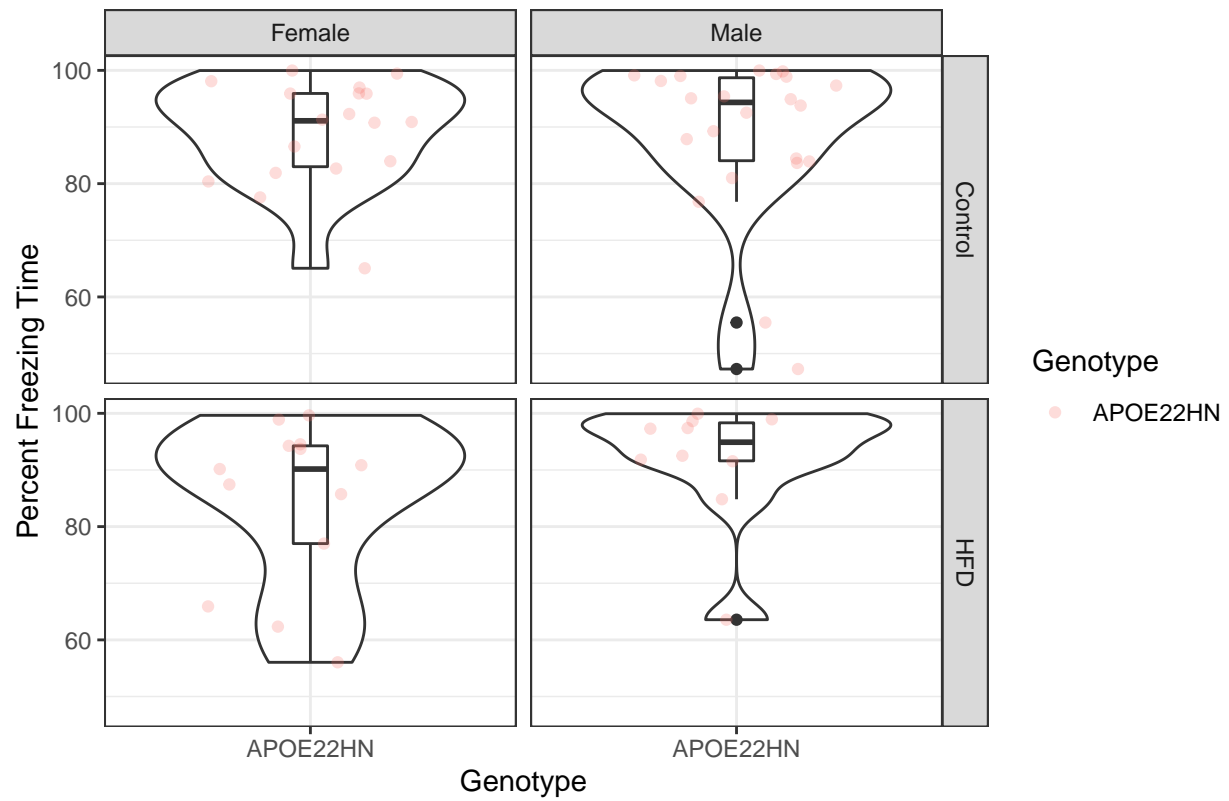
# Day 1 Contextual Test All Ages by Sex and HN Genotype



# Day 1 Contextual Test All Ages by Sex and nonHN Genotype

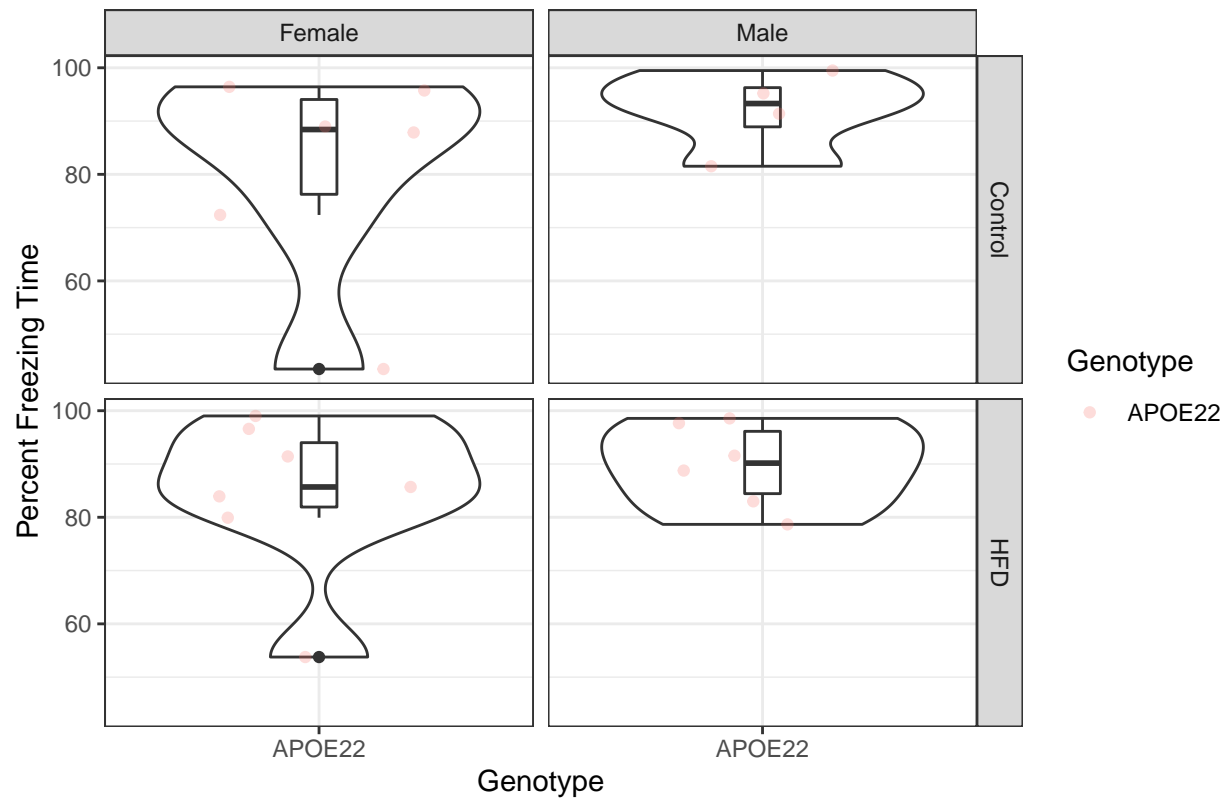


# Day 1 Contextual Test All Ages by Treatment, Sex and HN Genotype

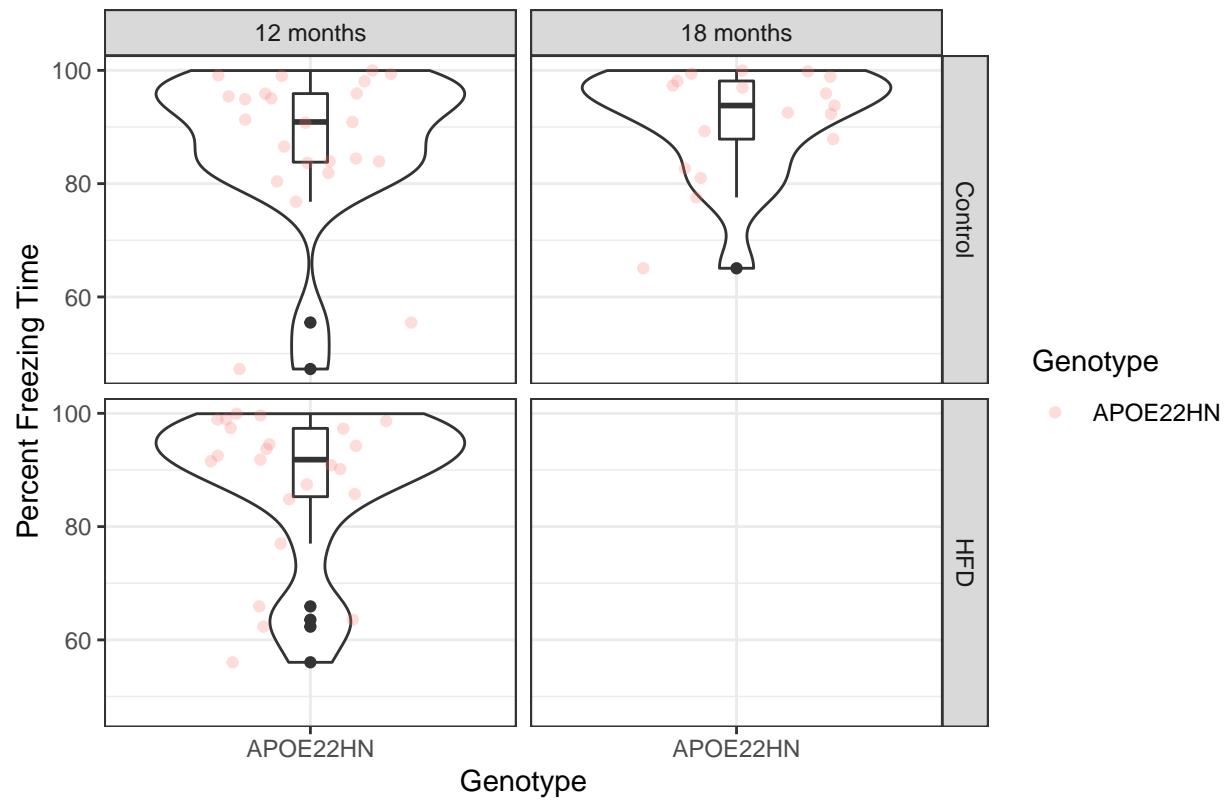




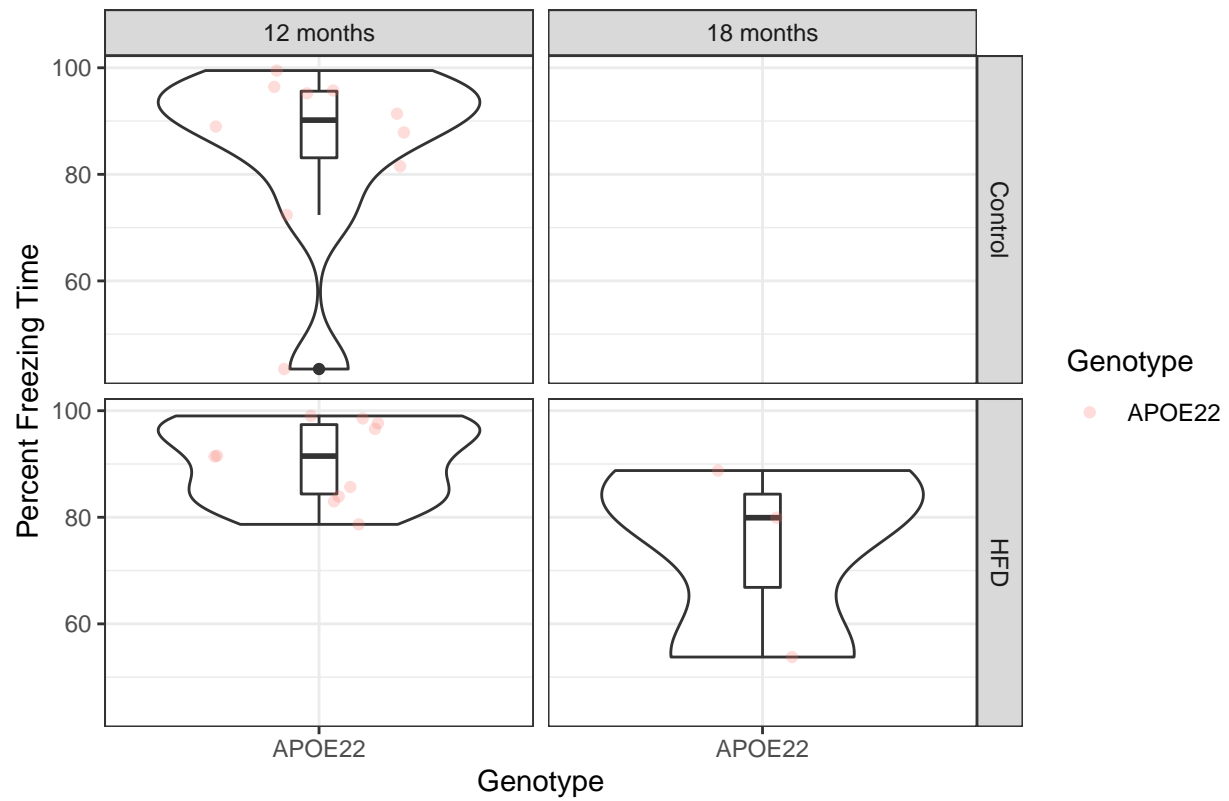
# Day 1 Contextual Test All Ages by Treatment, Sex and nonHN Genotype



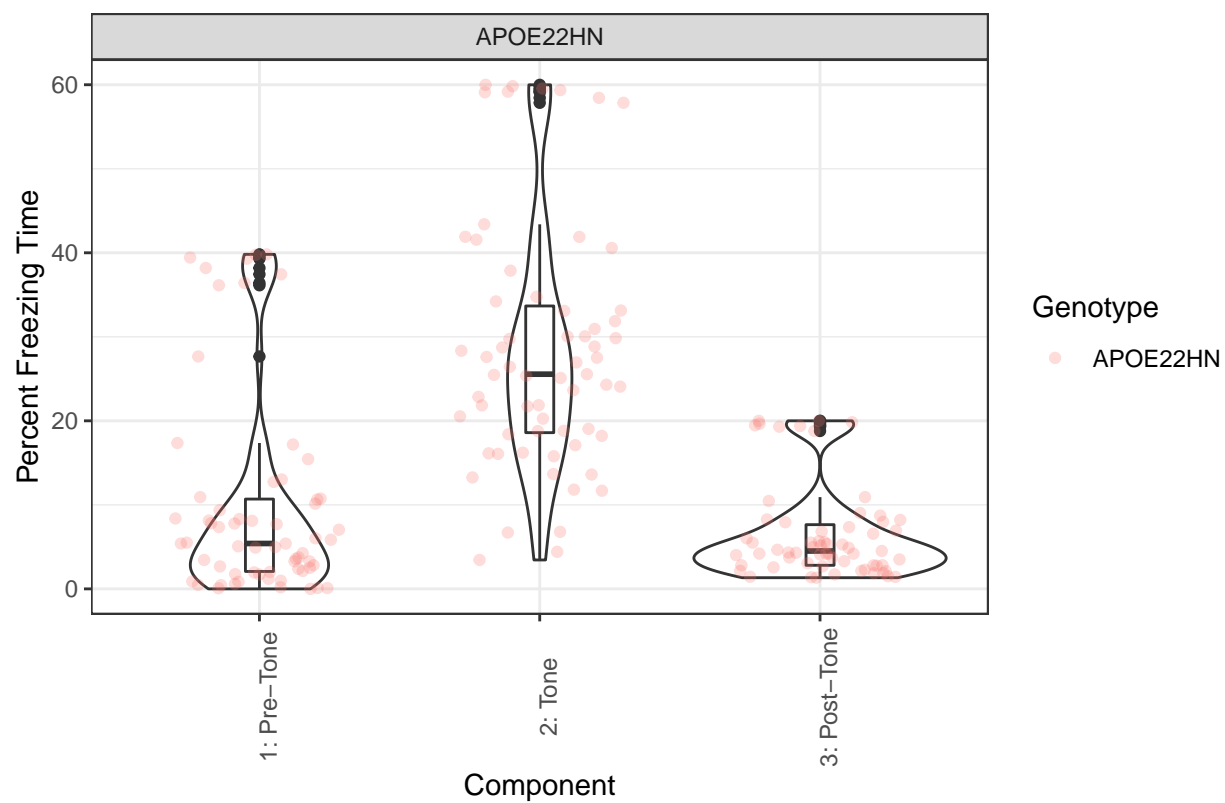
# Day 1 Contextual Test Treatment, Age and HN Genotype



# Day 1 Contextual Test Treatment, Age and nonHN Genotype



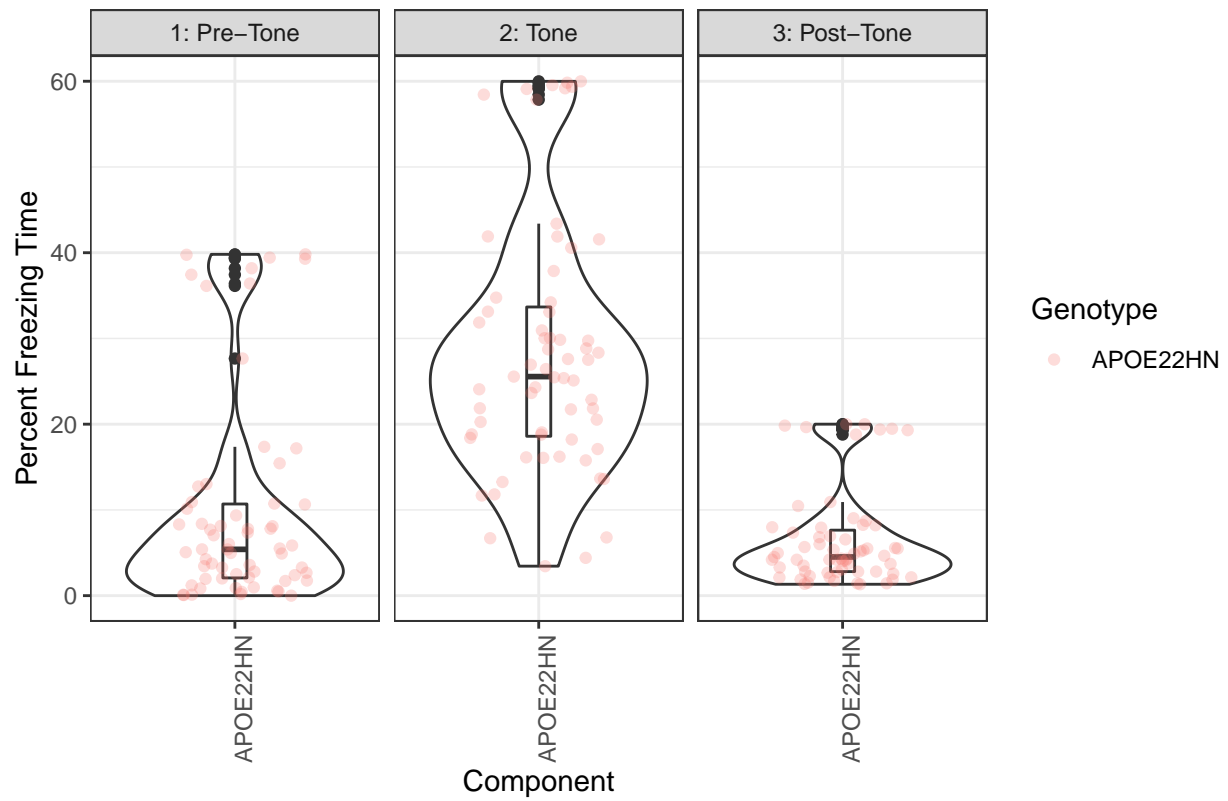
Day 2 All Ages Percent Freezing by Component and HN Genotype



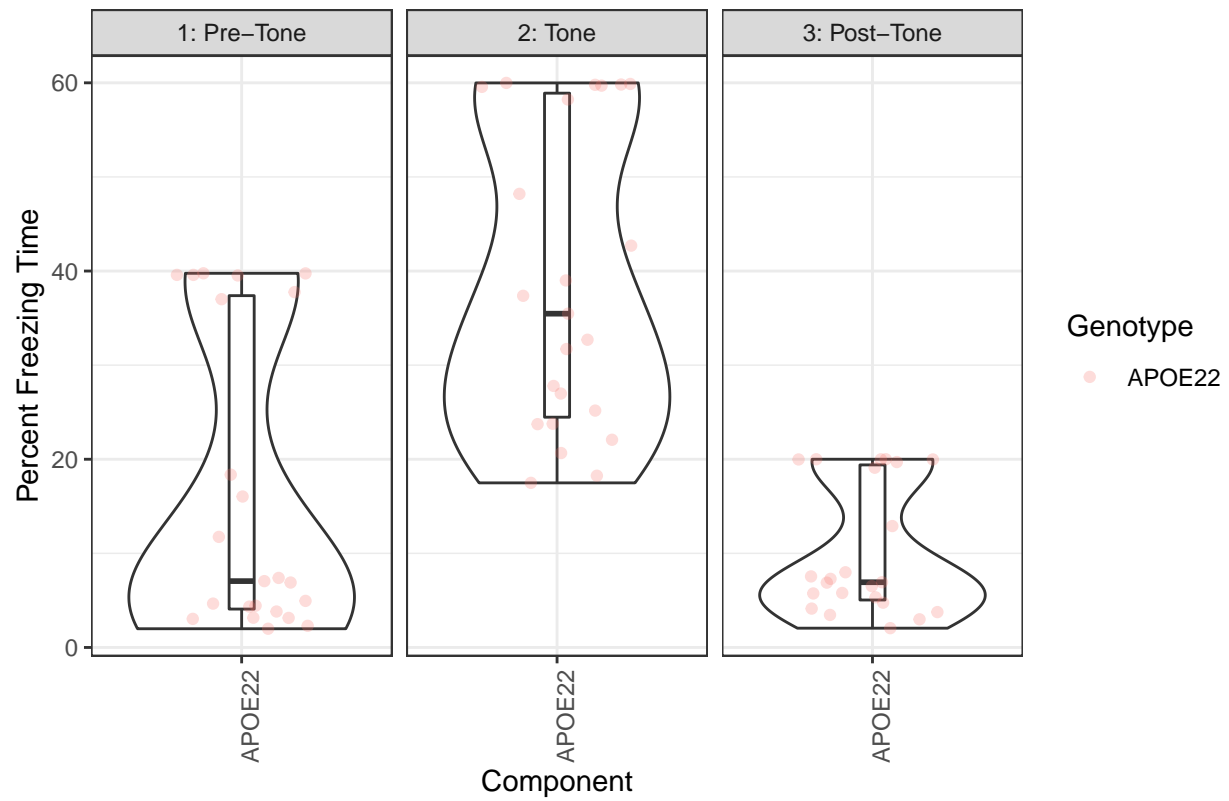
Day 2 All Ages Percent Freezing by Component and nonHN Genotype



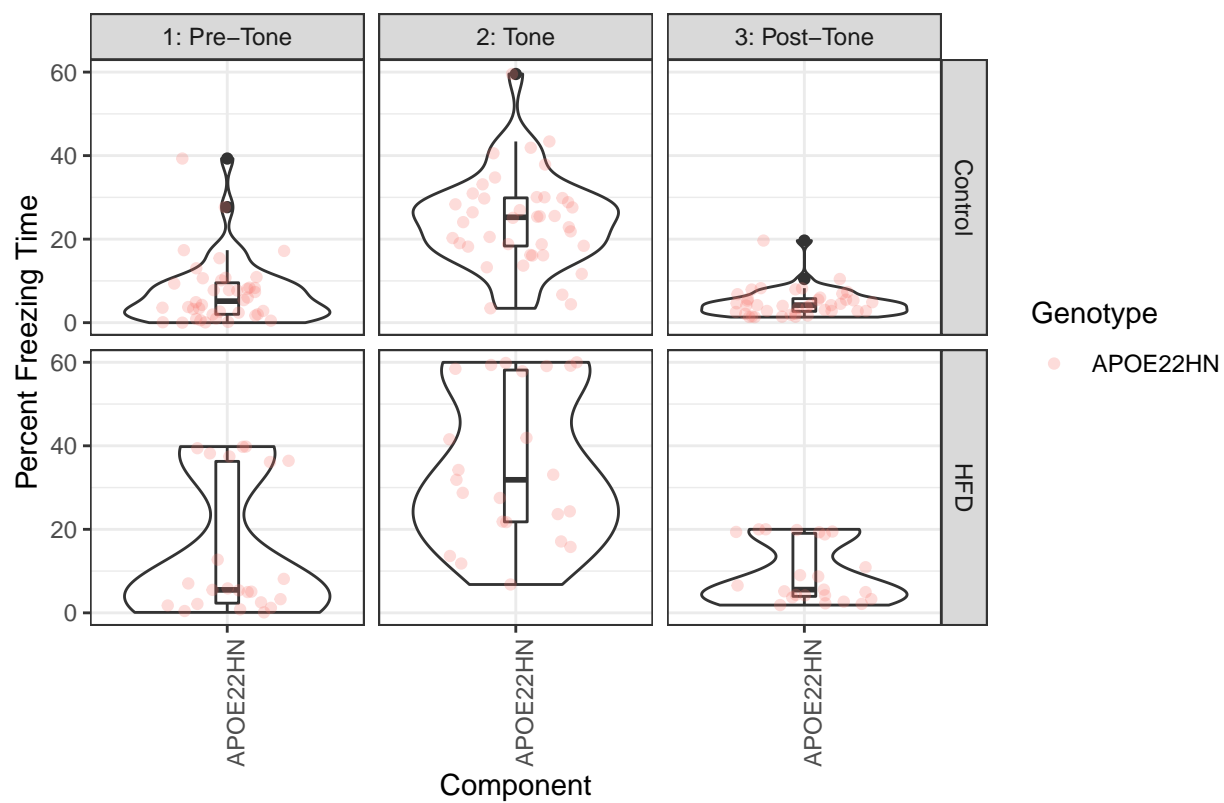
Day 2 All Ages Percent Freezing by Component and HN Genotype



Day 2 All Ages Percent Freezing by Component and nonHN Genotype

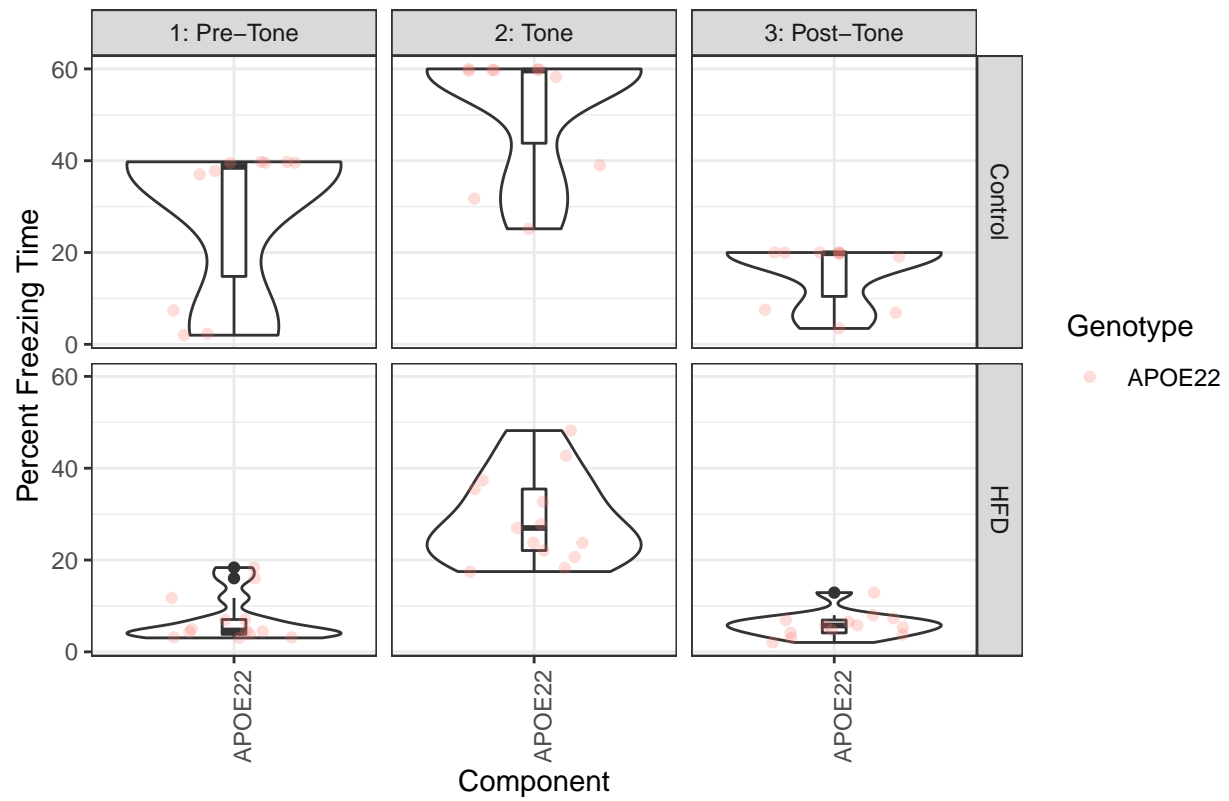


Day 2 All Ages Percent Freezing by Treatment, Component and HN Genotype

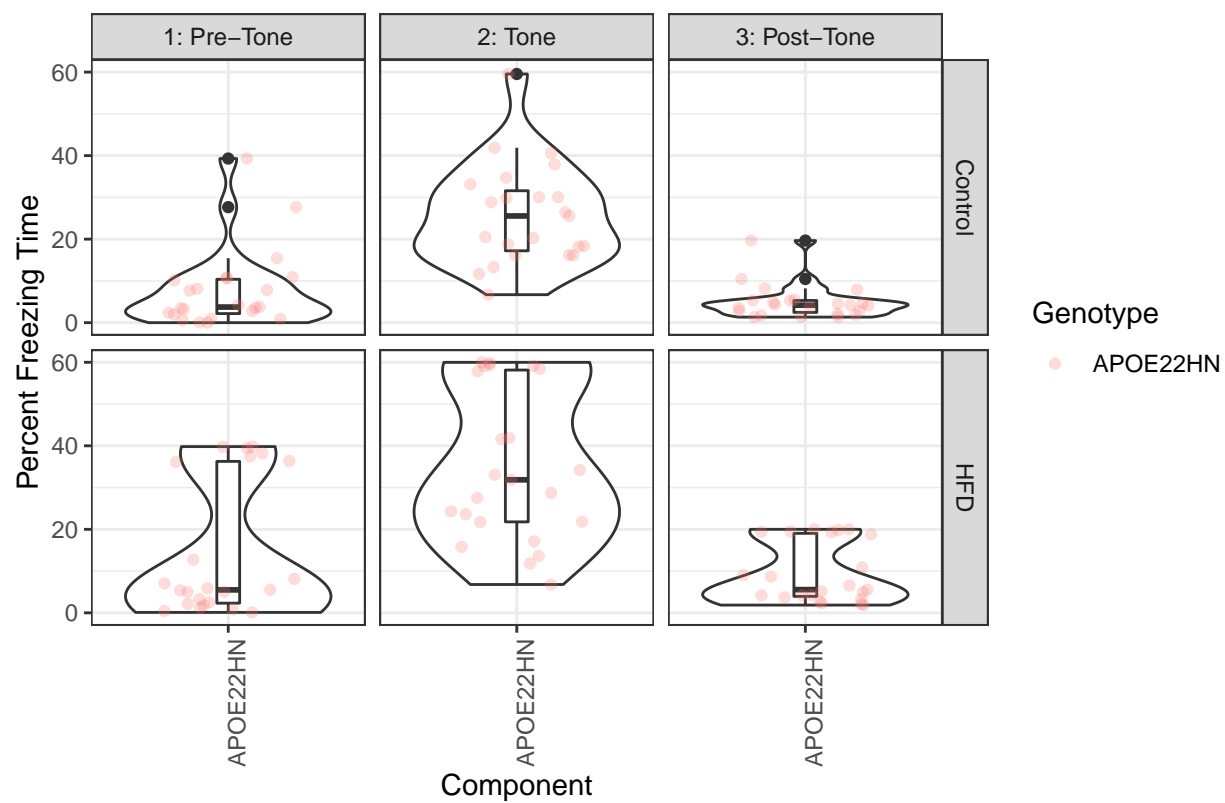




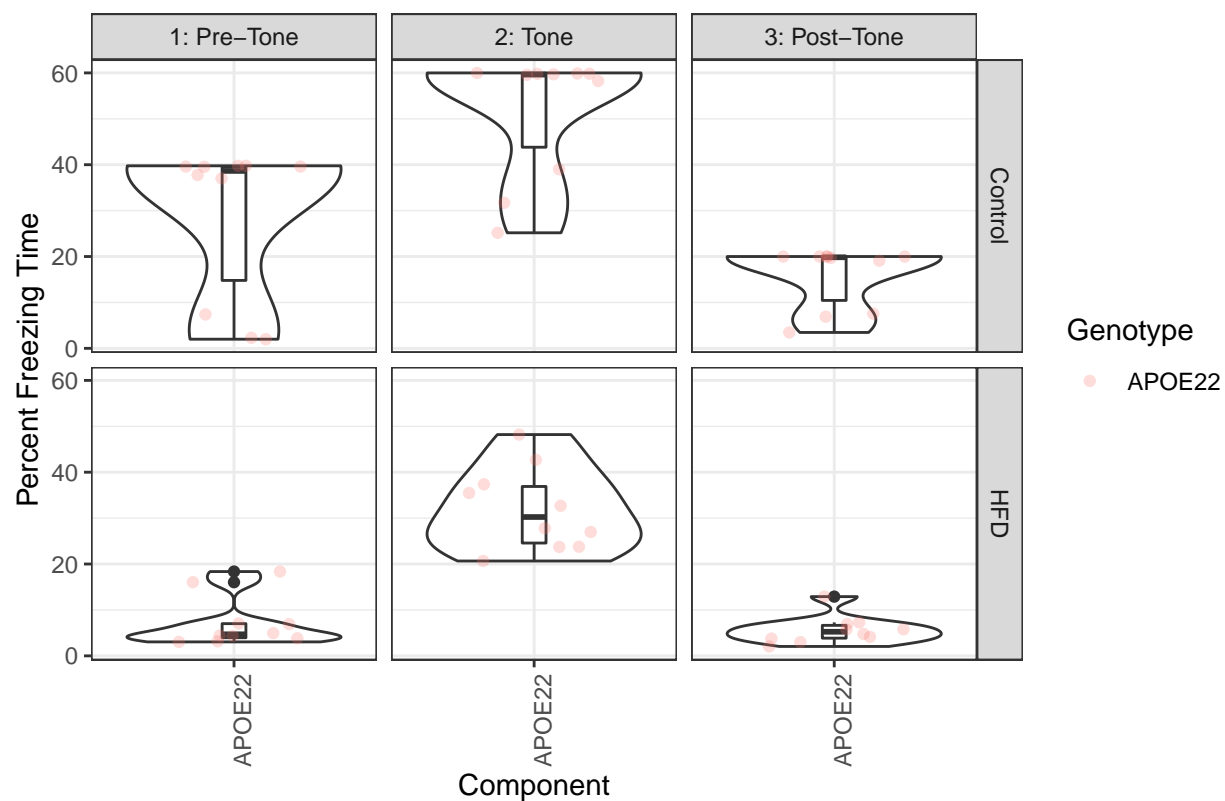
# Day 2 All Ages Percent Freezing by Treatment, Component and nonHN Ger



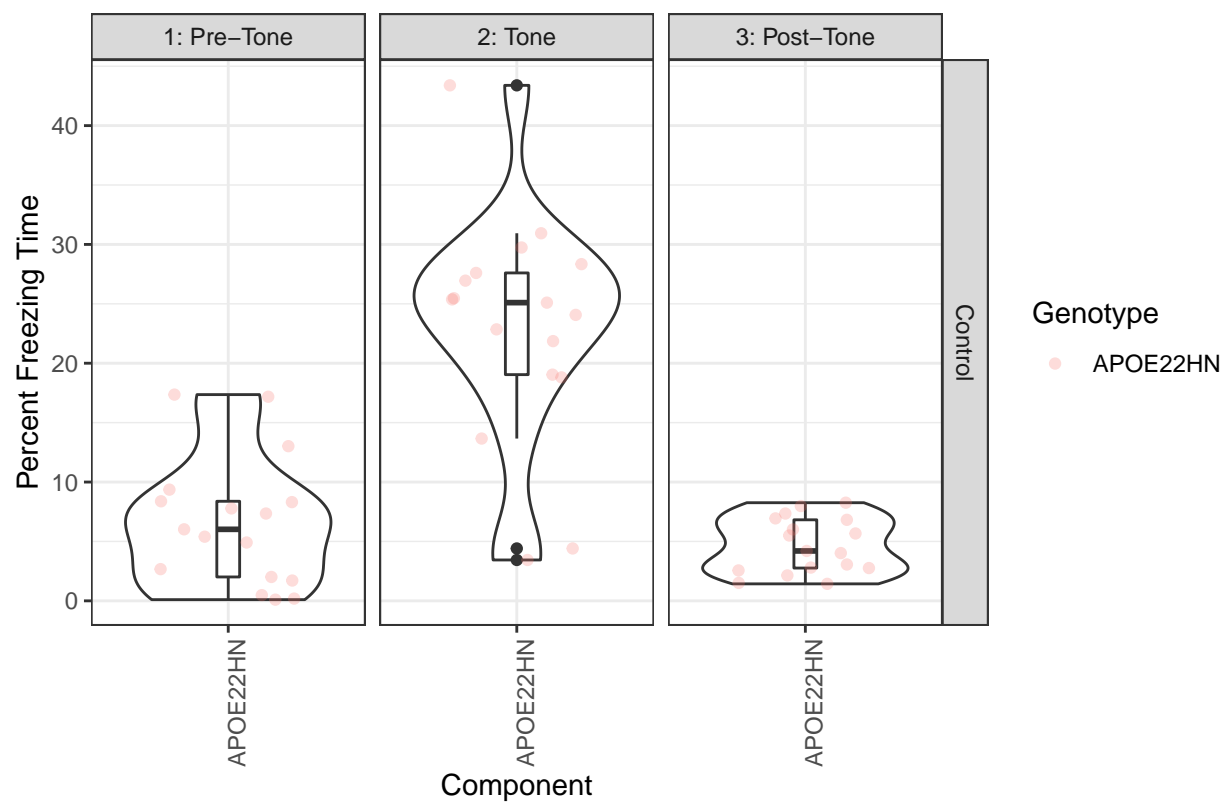
Day 2 12 Months Percent Freezing by Treatment, Component and HN Geno



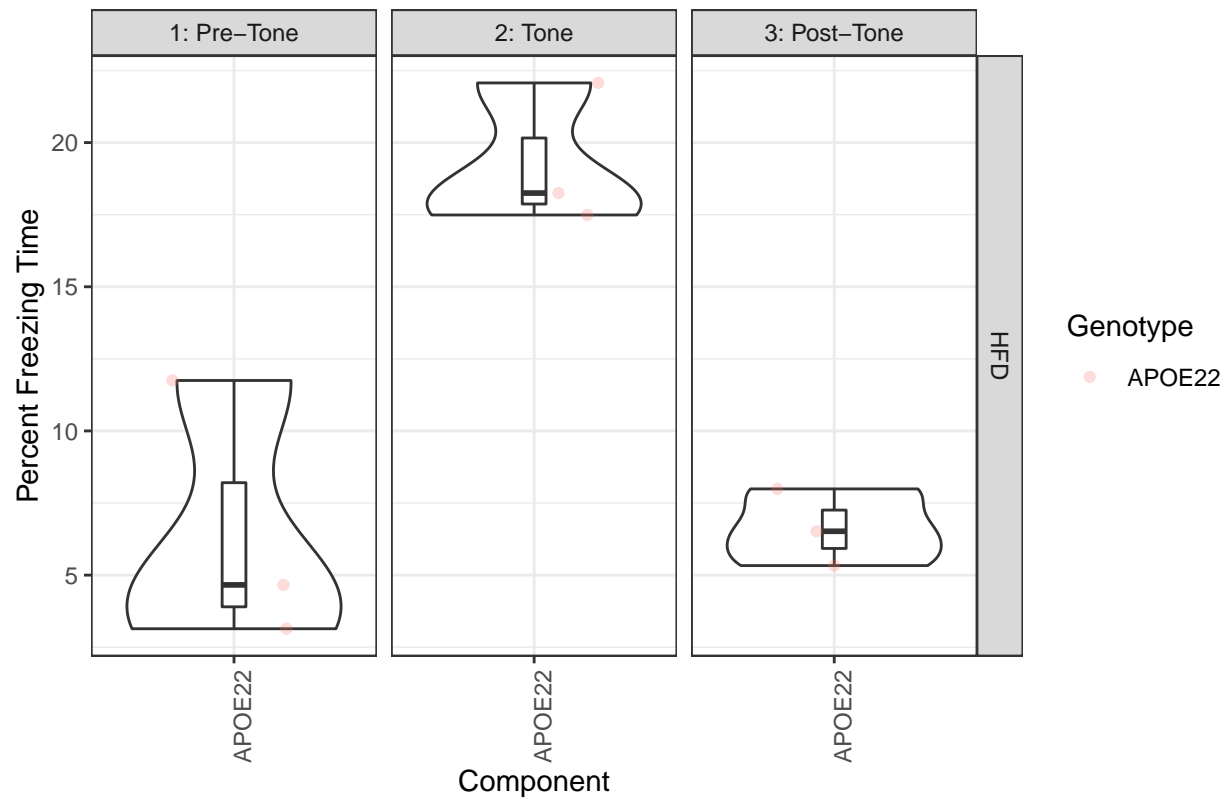
# Day 2 12 Months Percent Freezing by Treatment, Component and nonHN G



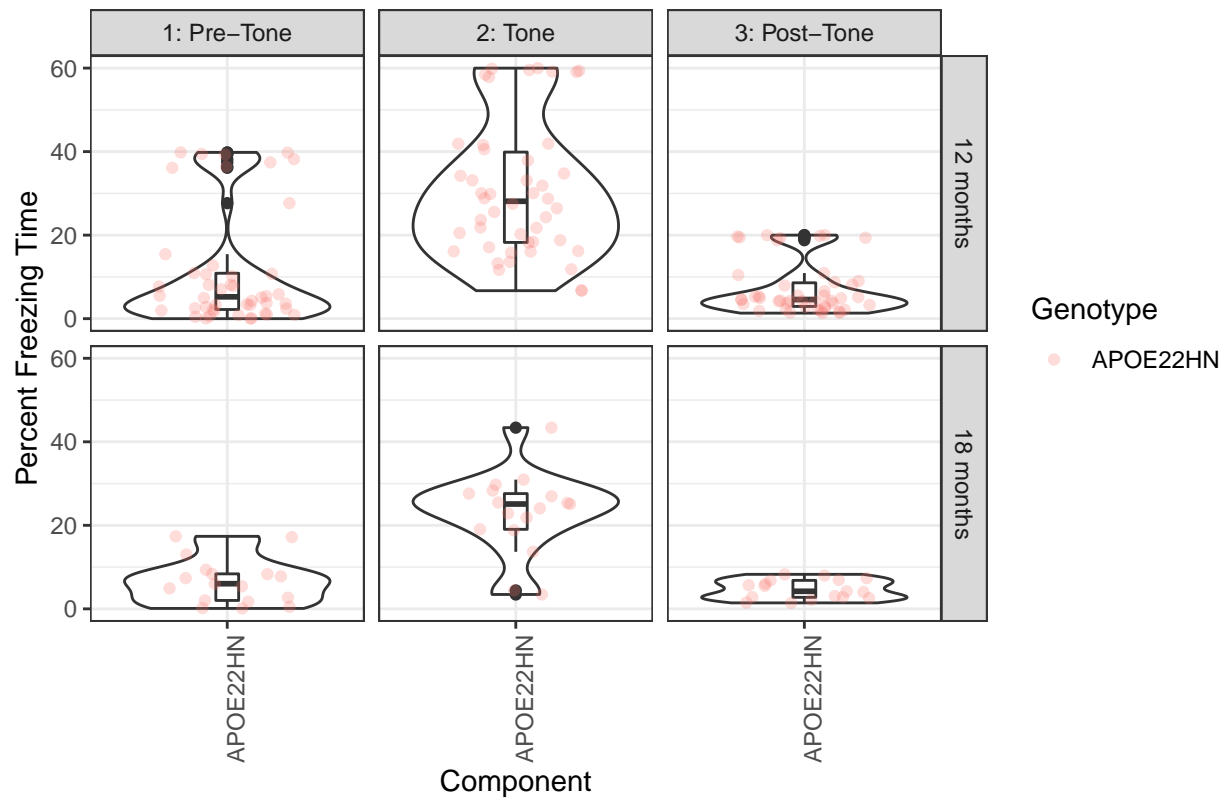
Day 2 18 Months Percent Freezing by Treatment, Component and HN Geno



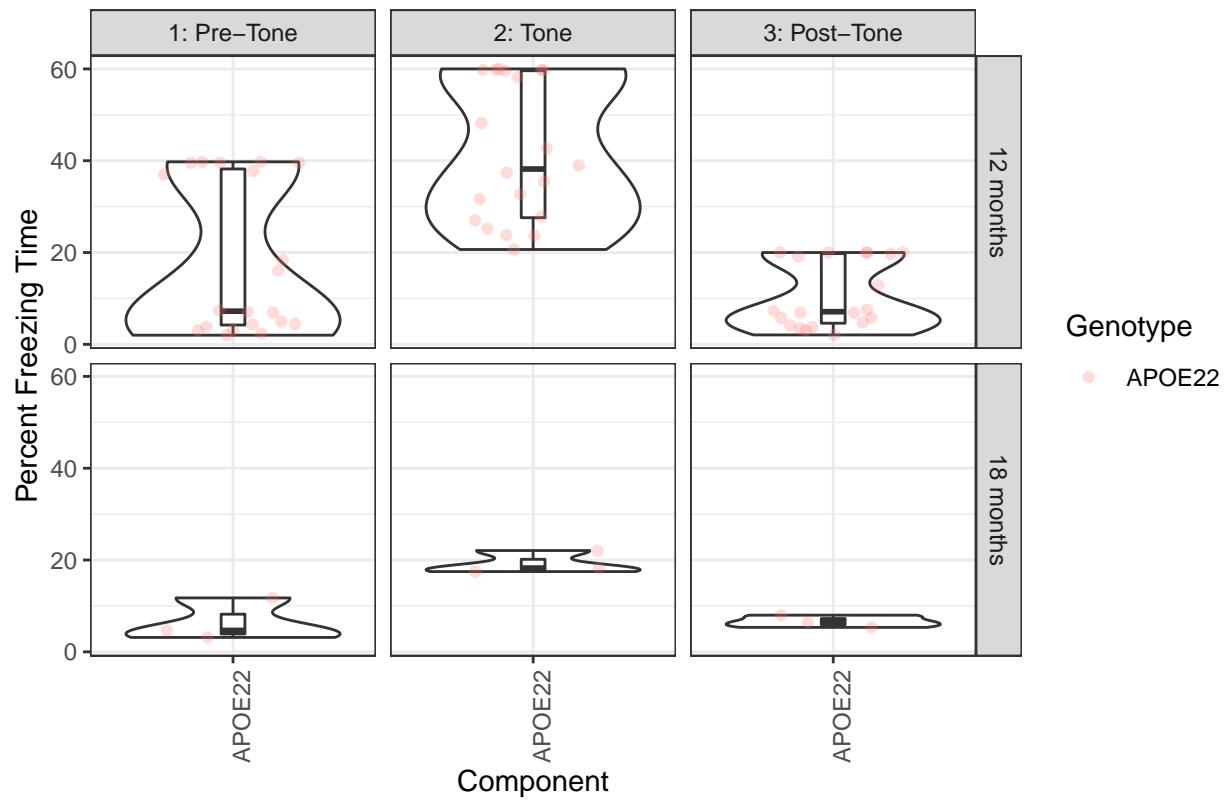
# Day 2 18 Months Percent Freezing by Treatment, Component and nonHN G



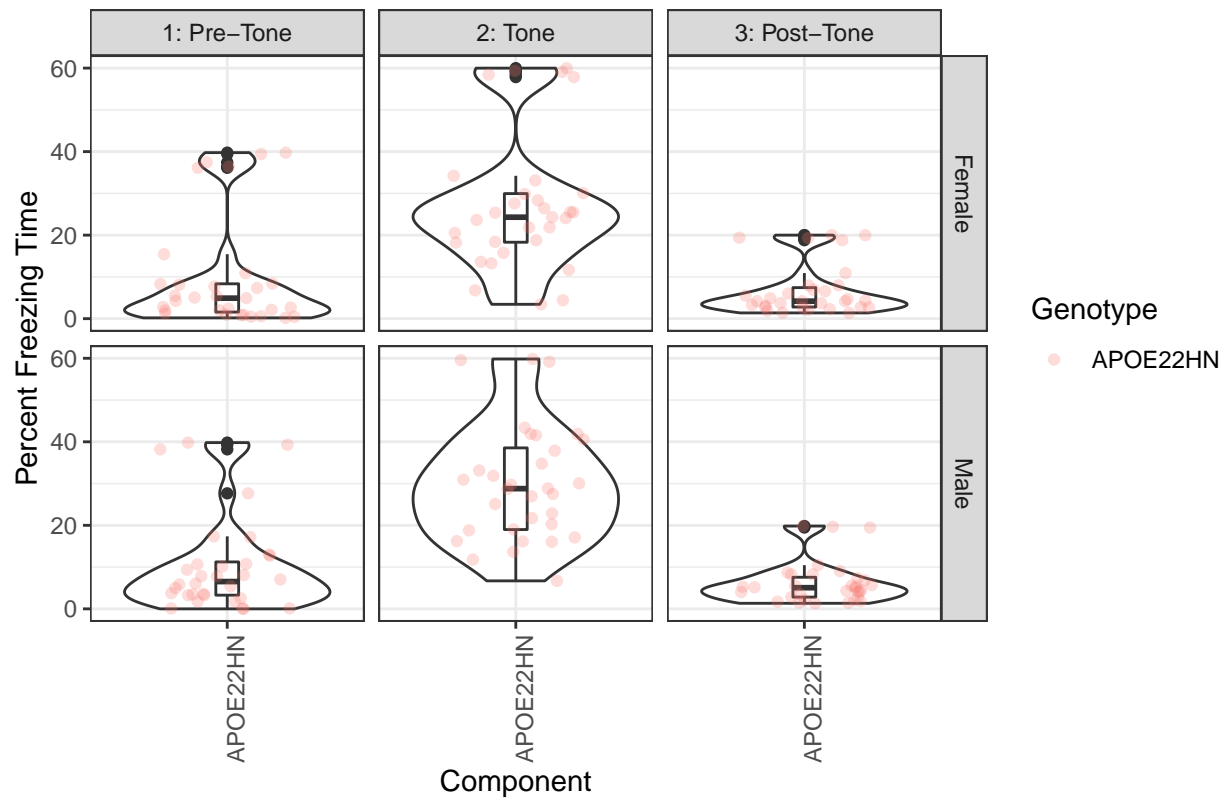
Day 2 Percent Freezing by Age, Component and HN Genotype



Day 2 Percent Freezing by Age, Component and nonHN Genotype

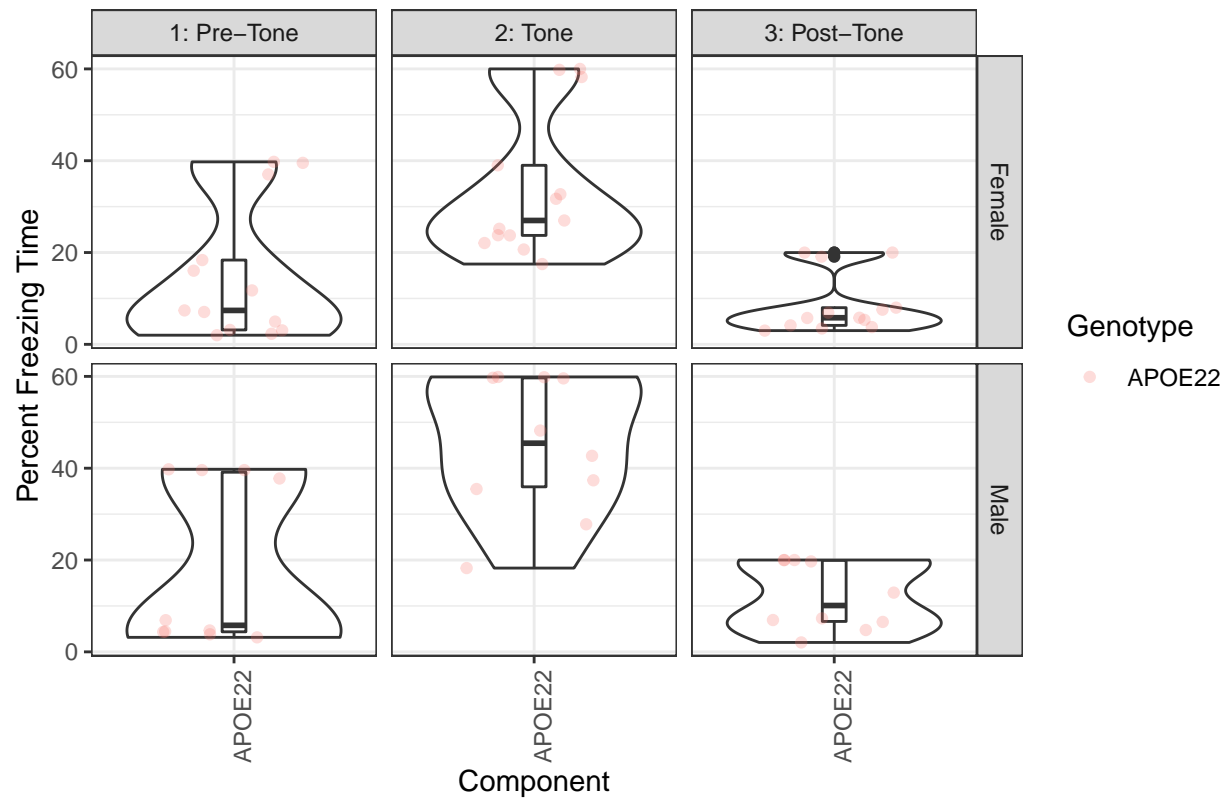


Day 2 All Ages Percent Freezing by Sex, Component and HN Genotype

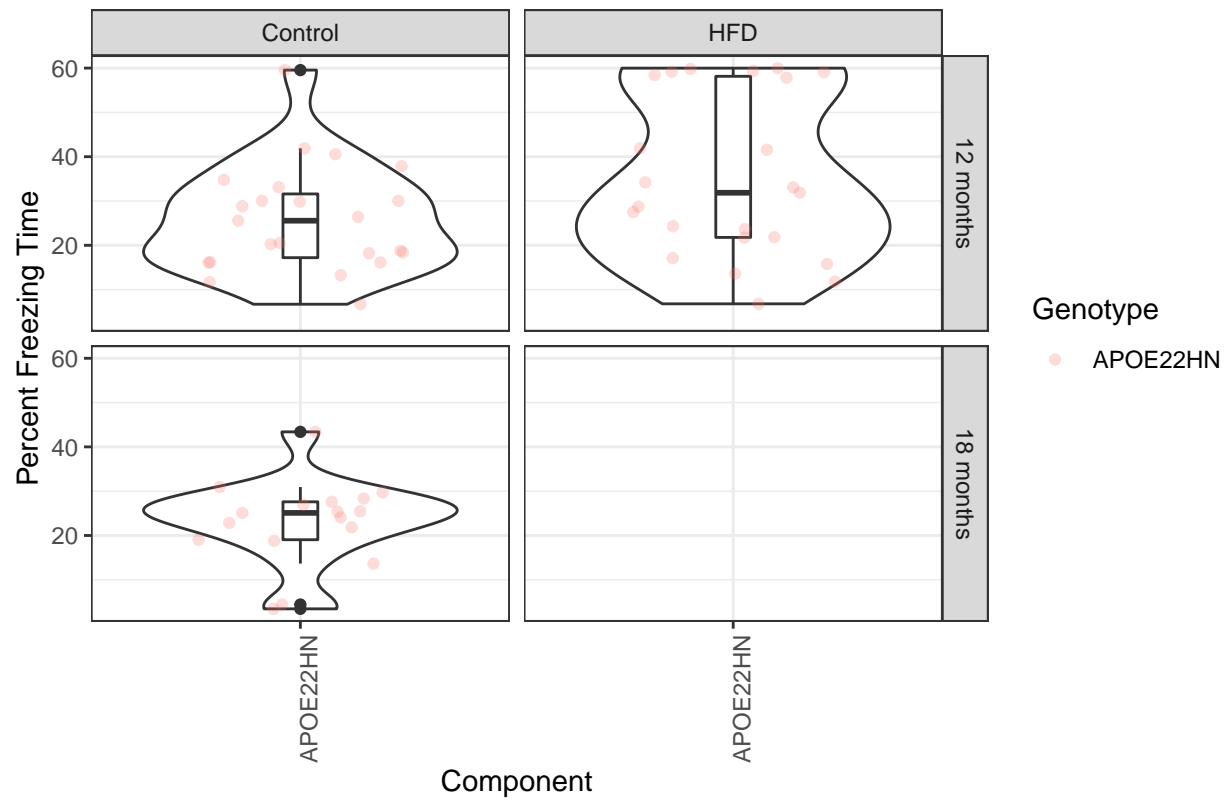




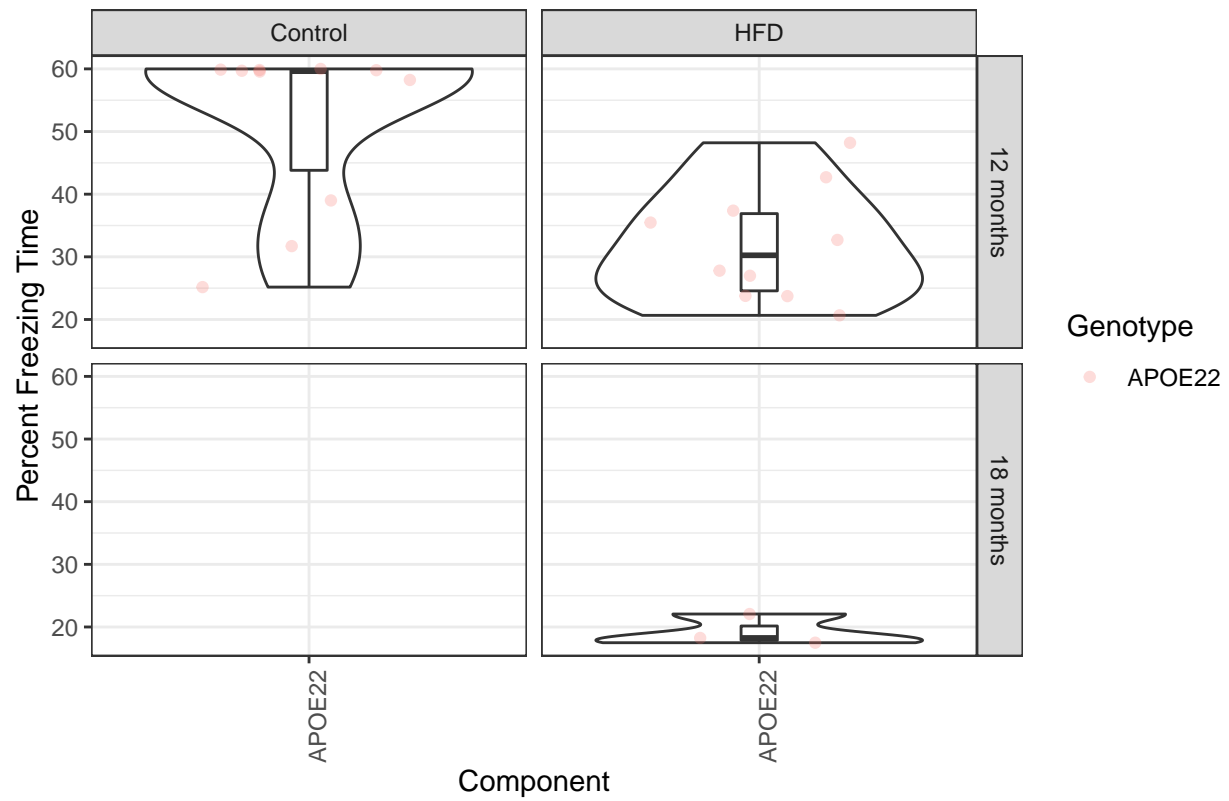
Day 2 All Ages Percent Freezing by Sex, Component and nonHN Genotype



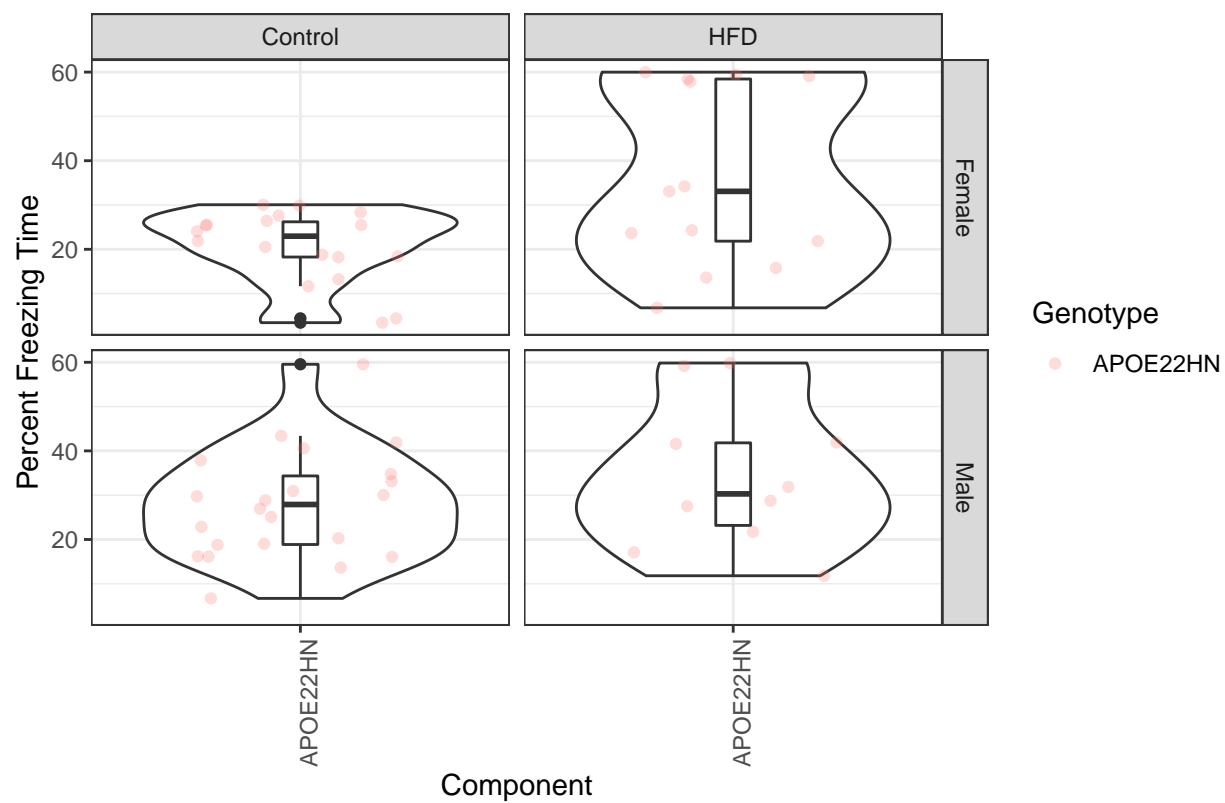
Day 2 Percent Freezing Within Tone by Age and HN Genotype



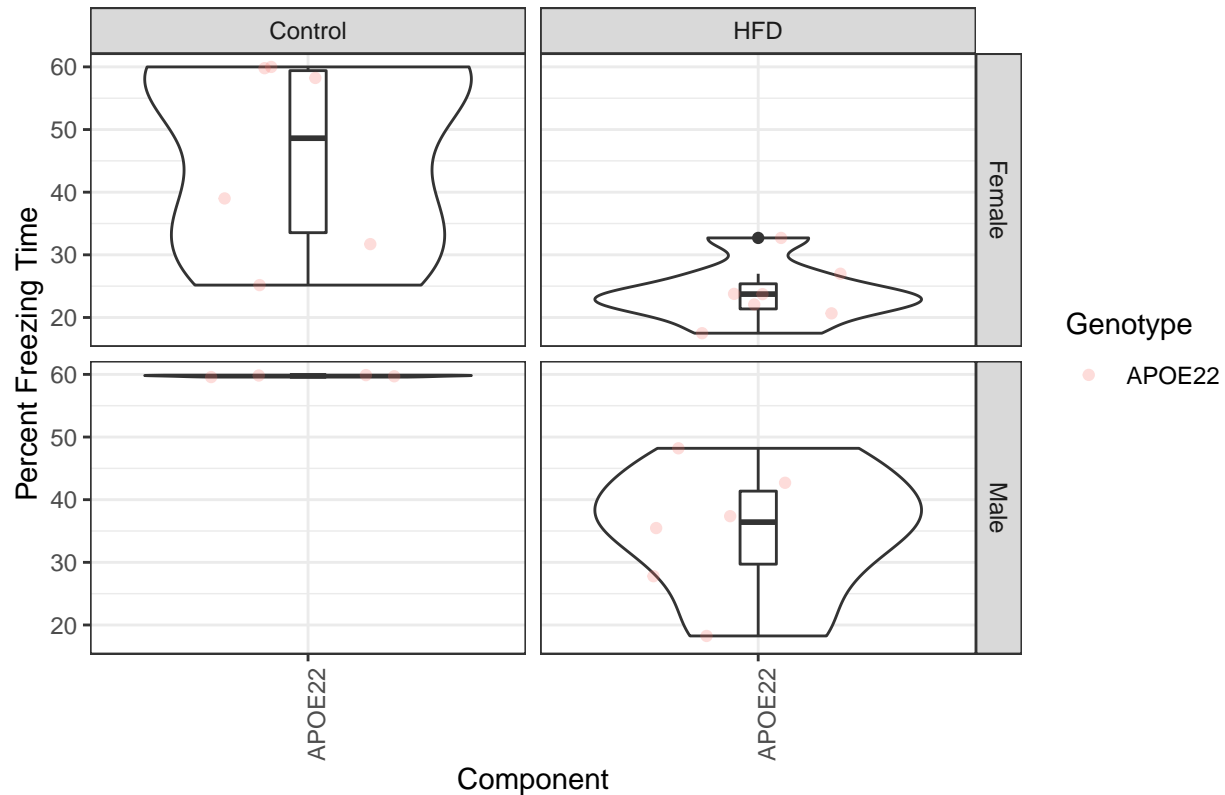
Day 2 Percent Freezing Within Tone by Age and nonHN Genotype



Day 2 Percent Freezing Within Tone by Sex and HN Genotype



## Day 2 Percent Freezing Within Tone by Sex and nonHN Genotype



```
## Type III Analysis of Variance Table with Satterthwaite's method
##               Sum Sq Mean Sq NumDF   DenDF F value    Pr(>F)
## Age              0.0153   0.0153     1 30.211   0.0356   0.85169
## Sex              0.7958   0.7958     1 30.211   1.8539   0.18340
## Treatment       18.4282  18.4282     1 30.211  42.9312 2.913e-07 ***
## Time           20.5480  20.5480     1 40.000  47.8696 2.434e-08 ***
## Age:Sex          0.1047   0.1047     1 30.211   0.2440   0.62490
## Sex:Treatment    1.7537   1.7537     1 30.211   4.0856   0.05219 .
## Age:Time         2.5765   2.5765     1 40.000   6.0023   0.01876 *
## Sex:Time         0.9347   0.9347     1 40.000   2.1776   0.14786
## Treatment:Time   23.3629  23.3629     1 40.000  54.4275 5.620e-09 ***
## Age:Sex:Time     0.8669   0.8669     1 40.000   2.0196   0.16303
## Sex:Treatment:Time 1.1603   1.1603     1 40.000   2.7032   0.10799
## Age:Treatment
## Age:Sex:Treatment
## Age:Treatment:Time
## Age:Sex:Treatment:Time
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
## Type III Analysis of Variance Table with Satterthwaite's method
##               Sum Sq Mean Sq NumDF   DenDF F value    Pr(>F)
## Age           48.552  48.552     1 584.67  88.6462 < 2.2e-16
## Genotype      72.330  14.466     5 583.78  26.4116 < 2.2e-16
## Sex            1.445   1.445     1 584.50   2.6378 0.1048852
```

## Treatment	12.952	12.952	1	583.63	23.6481	1.489e-06
## Time	248.766	248.766	1	572.09	454.1938	< 2.2e-16
## Age:Genotype	44.694	8.939	5	584.27	16.3204	4.357e-15
## Age:Sex	0.255	0.255	1	584.69	0.4658	0.4951958
## Genotype:Sex	4.240	0.848	5	583.65	1.5483	0.1729463
## Age:Treatment	6.937	6.937	1	583.80	12.6658	0.0004027
## Genotype:Treatment	53.511	10.702	5	583.80	19.5399	< 2.2e-16
## Sex:Treatment	1.311	1.311	1	583.27	2.3933	0.1223962
## Age:Time	14.497	14.497	1	572.09	26.4680	3.687e-07
## Genotype:Time	47.698	9.540	5	572.09	17.4171	4.548e-16
## Sex:Time	0.340	0.340	1	572.09	0.6214	0.4308598
## Treatment:Time	11.753	11.753	1	572.09	21.4580	4.481e-06
## Age:Genotype:Sex	2.599	0.520	5	584.12	0.9492	0.4485815
## Age:Genotype:Treatment	19.690	9.845	2	584.12	17.9746	2.658e-08
## Age:Sex:Treatment	1.593	1.593	1	583.51	2.9090	0.0886176
## Genotype:Sex:Treatment	18.036	3.607	5	583.81	6.5860	5.633e-06
## Age:Genotype:Time	24.794	4.959	5	572.09	9.0537	2.690e-08
## Age:Sex:Time	0.230	0.230	1	572.09	0.4204	0.5170184
## Genotype:Sex:Time	5.219	1.044	5	572.09	1.9056	0.0915716
## Age:Treatment:Time	6.095	6.095	1	572.09	11.1285	0.0009052
## Genotype:Treatment:Time	40.366	8.073	5	572.09	14.7400	1.299e-13
## Sex:Treatment:Time	1.661	1.661	1	572.09	3.0333	0.0821113
## Age:Genotype:Sex:Treatment	1.504	1.504	1	583.51	2.7468	0.0979885
## Age:Genotype:Sex:Time	6.634	1.327	5	572.09	2.4224	0.0345539
## Age:Genotype:Treatment:Time	14.443	7.222	2	572.09	13.1853	2.522e-06
## Age:Sex:Treatment:Time	2.039	2.039	1	572.09	3.7232	0.0541540
## Genotype:Sex:Treatment:Time	10.008	2.002	5	572.09	3.6545	0.0029044
## Age:Genotype:Sex:Treatment:Time	3.948	3.948	1	572.09	7.2076	0.0074701
##						
## Age	***					
## Genotype	***					
## Sex						
## Treatment	***					
## Time	***					
## Age:Genotype	***					
## Age:Sex						
## Genotype:Sex						
## Age:Treatment	***					
## Genotype:Treatment	***					
## Sex:Treatment						
## Age:Time	***					
## Genotype:Time	***					
## Sex:Time						
## Treatment:Time	***					
## Age:Genotype:Sex						
## Age:Genotype:Treatment	***					
## Age:Sex:Treatment	.					
## Genotype:Sex:Treatment	***					
## Age:Genotype:Time	***					
## Age:Sex:Time						
## Genotype:Sex:Time	.					
## Age:Treatment:Time	***					
## Genotype:Treatment:Time	***					
## Sex:Treatment:Time	.					

```

## Age:Genotype:Sex:Treatment      .
## Age:Genotype:Sex:Time           *
## Age:Genotype:Treatment:Time     ***
## Age:Sex:Treatment:Time          .
## Genotype:Sex:Treatment:Time     **
## Age:Genotype:Sex:Treatment:Time **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

## Type III Analysis of Variance Table with Satterthwaite's method
##              Sum Sq Mean Sq NumDF  DenDF  F value    Pr(>F)
## HN              0.82    0.82      1 577.86   1.0590    0.3039
## APOE            61.53   30.77      2 577.67  39.6953 < 2.2e-16 ***
## Time           367.04  367.04      1 607.00 473.5769 < 2.2e-16 ***
## HN:APOE         27.63   13.81      2 577.67  17.8245 3.079e-08 ***
## HN:Time          0.01    0.01      1 607.00   0.0112    0.9159
## APOE:Time        41.01   20.50      2 607.00  26.4545 9.647e-12 ***
## HN:APOE:Time     21.07   10.53      2 607.00  13.5899 1.684e-06 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

## Type III Analysis of Variance Table with Satterthwaite's method
##              Sum Sq Mean Sq NumDF  DenDF  F value    Pr(>F)
## Age            194.08  194.08      1 831.99 14.6623 0.0001382
## Genotype       84.17   16.83      5 832.08  1.2718 0.2739922
## Sex             9.59    9.59      1 832.01  0.7242 0.3950035
## Treatment       9.76    9.76      1 832.10  0.7375 0.3907051
## Time          624.09  624.09      1 571.70 47.1476 1.721e-11
## Age:Genotype   148.27   29.65      5 832.03  2.2402 0.0485525
## Age:Sex        27.01   27.01      1 831.99  2.0409 0.1534971
## Genotype:Sex    5.65    1.13      5 832.09  0.0854 0.9945399
## Age:Treatment   6.44    6.44      1 832.08  0.4866 0.4856403
## Genotype:Treatment 63.05   12.61      5 832.08  0.9527 0.4460844
## Sex:Treatment   0.03    0.03      1 832.13  0.0022 0.9622443
## Age:Time       517.49  517.49      1 571.70 39.0941 7.909e-10
## Genotype:Time   364.04   72.81      5 571.70  5.5003 5.884e-05
## Sex:Time        53.24   53.24      1 571.70  4.0224 0.0453697
## Treatment:Time  50.74   50.74      1 571.70  3.8334 0.0507265
## Age:Genotype:Sex 11.51    2.30      5 832.05  0.1739 0.9723016
## Age:Genotype:Treatment 45.95   22.98      2 832.05  1.7357 0.1769124
## Age:Sex:Treatment 8.39    8.39      1 832.11  0.6335 0.4262868
## Genotype:Sex:Treatment 19.93    3.99      5 832.08  0.3011 0.9122806
## Age:Genotype:Time 432.91   86.58      5 571.70  6.5409 6.258e-06
## Age:Sex:Time     91.56   91.56      1 571.70  6.9168 0.0087688
## Genotype:Sex:Time 24.50    4.90      5 571.70  0.3702 0.8691022
## Age:Treatment:Time 27.45   27.45      1 571.70  2.0735 0.1504249
## Genotype:Treatment:Time 252.96   50.59      5 571.70  3.8220 0.0020526
## Sex:Treatment:Time 0.57    0.57      1 571.70  0.0428 0.8362204
## Age:Genotype:Sex:Treatment 0.43    0.43      1 832.11  0.0328 0.8564128
## Age:Genotype:Sex:Time 43.89    8.78      5 571.70  0.6632 0.6515412
## Age:Genotype:Treatment:Time 125.29   62.64      2 571.70  4.7324 0.0091530
## Age:Sex:Treatment:Time 5.80    5.80      1 571.70  0.4382 0.5082508
## Genotype:Sex:Treatment:Time 153.12   30.62      5 571.70  2.3135 0.0426173
## Age:Genotype:Sex:Treatment:Time 6.82    6.82      1 571.70  0.5156 0.4730371

```

```

##
## Age ***
## Genotype
## Sex
## Treatment
## Time ***
## Age:Genotype *
## Age:Sex
## Genotype:Sex
## Age:Treatment
## Genotype:Treatment
## Sex:Treatment
## Age:Time ***
## Genotype:Time ***
## Sex:Time *
## Treatment:Time .
## Age:Genotype:Sex
## Age:Genotype:Treatment
## Age:Sex:Treatment
## Genotype:Sex:Treatment
## Age:Genotype:Time ***
## Age:Sex:Time **
## Genotype:Sex:Time
## Age:Treatment:Time
## Genotype:Treatment:Time **
## Sex:Treatment:Time
## Age:Genotype:Sex:Treatment
## Age:Genotype:Sex:Time
## Age:Genotype:Treatment:Time **
## Age:Sex:Treatment:Time
## Genotype:Sex:Treatment:Time *
## Age:Genotype:Sex:Treatment:Time
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

## Type III Analysis of Variance Table with Satterthwaite's method
##              Sum Sq Mean Sq NumDF  DenDF F value    Pr(>F)
## HN           30.81   30.81     1 897.26  1.8342 0.1759685
## APOE          37.78   18.89     2 897.31  1.1248 0.3251857
## Time         326.45  326.45     1 606.75 19.4370 1.23e-05 ***
## HN:APOE       10.55    5.28     2 897.31  0.3141 0.7304985
## HN:Time       77.78   77.78     1 606.75  4.6312 0.0317891 *
## APOE:Time     287.00 143.50     2 606.75  8.5441 0.0002191 ***
## HN:APOE:Time  91.99   45.99     2 606.75  2.7385 0.0654676 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

## Analysis of Variance Table
##
## Response: Pct.Total.Time.Freezing
##              Df Sum Sq Mean Sq F value    Pr(>F)
## Age           1    297   296.99  2.1877 0.140323
## Genotype      5   2633   526.69  3.8797 0.002078 **
## Sex           1    425   424.69  3.1284 0.078110 .

```



```
## Treatment      1      237  237.25  1.7477 0.187329
## Age:Genotype   5     1661  332.14  2.4466 0.034469 *
## Age:Sex        1       16   15.58  0.1147 0.735087
## Genotype:Sex    5     921  184.20  1.3568 0.241036
## Age:Treatment  1     309  308.70  2.2740 0.132773
## Genotype:Treatment  5     881  176.30  1.2986 0.264819
## Sex:Treatment   1       33   32.59  0.2401 0.624565
## Age:Genotype:Sex  5     408   81.57  0.6008 0.699340
## Age:Genotype:Treatment  2     502  250.92  1.8483 0.159562
## Age:Sex:Treatment  1        8    7.55  0.0556 0.813774
## Genotype:Sex:Treatment  5     618  123.50  0.9097 0.475175
## Age:Genotype:Sex:Treatment  1      17   17.01  0.1253 0.723666
## Residuals      261   35432  135.76
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
## Analysis of Variance Table
##
## Response: Pct.Total.Time.Freezing
##           Df Sum Sq Mean Sq F value    Pr(>F)
## Genotype    5   2893   578.67  4.1271 0.001224 **
## Residuals  296  41503   140.21
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
## Analysis of Variance Table
##
## Response: Pct.Total.Time.Freezing
##           Df Sum Sq Mean Sq F value    Pr(>F)
## HN           1    178   178.29  1.2716 0.2603803
## APOE          2   2171  1085.37  7.7409 0.0005286 ***
## HN:APOE       2    544   272.16  1.9410 0.1453794
## Residuals  296  41503   140.21
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
## Analysis of Variance Table
##
## Response: Freeze.Cnt
##           Df Sum Sq Mean Sq F value    Pr(>F)
## Age          1   1397   1397.0  1.0728 0.301267
## Genotype      5  24053   4810.5  3.6941 0.003012 **
## Sex           1   5983   5983.4  4.5948 0.032993 *
## Treatment     1   3046   3045.7  2.3389 0.127390
## Age:Genotype   5  13069   2613.7  2.0072 0.078003 .
## Age:Sex        1   1164   1164.4  0.8941 0.345234
## Genotype:Sex    5   7465   1493.0  1.1465 0.336203
## Age:Treatment  1   1329   1328.8  1.0204 0.313363
## Genotype:Treatment  5  11659   2331.8  1.7906 0.115061
## Sex:Treatment   1    568    568.0  0.4362 0.509543
## Age:Genotype:Sex  5   5087   1017.4  0.7813 0.563920
## Age:Genotype:Treatment  2   4047   2023.3  1.5537 0.213408
## Age:Sex:Treatment  1   1166   1166.4  0.8957 0.344807
```

```
## Genotype:Sex:Treatment      5   3251   650.2  0.4993 0.776662
## Age:Genotype:Sex:Treatment  1    228   228.3  0.1753 0.675784
## Residuals                   261 339876 1302.2
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
## Analysis of Variance Table
##
```

```
## Response: Freeze.Cnt
##           Df Sum Sq Mean Sq F value    Pr(>F)
## HN          1     94    94.4  0.0702 0.7912505
## APOE         2  21853 10926.7  8.1273 0.0003663 ***
## HN:APOE      2   3482  1741.0  1.2949 0.2754596
## Residuals 296 397958  1344.5
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
## Type III Analysis of Variance Table with Satterthwaite's method
```

```
##           Sum Sq Mean Sq NumDF  DenDF F value
## Age          4741.3   4741.3      1  731.03 19.0036
## Genotype     9547.5   1909.5      5  731.03  7.6535
## Sex          102.3    102.3      1  731.03  0.4099
## Treatment     853.8    853.8      1  731.03  3.4220
## Component    2880.5   2880.5      1  563.00 11.5453
## Age:Genotype 2631.4    526.3      5  731.03  2.1094
## Age:Sex        4.7      4.7      1  731.03  0.0187
## Genotype:Sex   166.5     33.3      5  731.03  0.1335
## Age:Treatment  523.5    523.5      1  731.03  2.0982
## Genotype:Treatment 3477.7   695.5      5  731.03  2.7878
## Sex:Treatment  133.8    133.8      1  731.03  0.5361
## Age:Component  965.6    965.6      1  563.00  3.8704
## Genotype:Component 1920.0   384.0      5  563.00  1.5391
## Sex:Component   17.1     17.1      1  563.00  0.0686
## Treatment:Component 231.5    231.5      1  563.00  0.9281
## Age:Genotype:Sex 332.2     66.4      5  731.03  0.2663
## Age:Genotype:Treatment 1033.0   516.5      2  731.03  2.0702
## Age:Sex:Treatment  33.7     33.7      1  731.03  0.1351
## Genotype:Sex:Treatment 2130.0   426.0      5  731.03  1.7075
## Age:Genotype:Component 470.6     94.1      5  563.00  0.3773
## Age:Sex:Component   3.4      3.4      1  563.00  0.0136
## Genotype:Sex:Component 131.6     26.3      5  563.00  0.1055
## Age:Treatment:Component 121.5    121.5      1  563.00  0.4870
## Genotype:Treatment:Component 670.5    134.1      5  563.00  0.5375
## Sex:Treatment:Component 25.8     25.8      1  563.00  0.1036
## Age:Genotype:Sex:Treatment 194.7    194.7      1  731.03  0.7805
## Age:Genotype:Sex:Component 66.4     13.3      5  563.00  0.0532
## Age:Genotype:Treatment:Component 140.5     70.3      2  563.00  0.2816
## Age:Sex:Treatment:Component 0.1      0.1      1  563.00  0.0005
## Genotype:Sex:Treatment:Component 480.4     96.1      5  563.00  0.3851
## Age:Genotype:Sex:Treatment:Component 44.9     44.9      1  563.00  0.1798
##           Pr(>F)
## Age          1.492e-05 ***
## Genotype      5.035e-07 ***
## Sex           0.5222357
```

```

## Treatment                0.0647375 .
## Component                 0.0007272 ***
## Age:Genotype              0.0624116 .
## Age:Sex                   0.8912927
## Genotype:Sex              0.9846508
## Age:Treatment             0.1479015
## Genotype:Treatment        0.0166839 *
## Sex:Treatment             0.4642709
## Age:Component             0.0496347 *
## Genotype:Component        0.1757663
## Sex:Component             0.7934127
## Treatment:Component       0.3357798
## Age:Genotype:Sex          0.9314894
## Age:Genotype:Treatment    0.1269014
## Age:Sex:Treatment         0.7133060
## Genotype:Sex:Treatment    0.1304937
## Age:Genotype:Component    0.8643798
## Age:Sex:Component         0.9073119
## Genotype:Sex:Component    0.9910321
## Age:Treatment:Component   0.4855358
## Genotype:Treatment:Component 0.7479285
## Sex:Treatment:Component   0.7476664
## Age:Genotype:Sex:Treatment 0.3772733
## Age:Genotype:Sex:Component 0.9982233
## Age:Genotype:Treatment:Component 0.7546626
## Age:Sex:Treatment:Component 0.9817590
## Genotype:Sex:Treatment:Component 0.8590817
## Age:Genotype:Sex:Treatment:Component 0.6717321
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

## Type III Analysis of Variance Table with Satterthwaite's method
##              Sum Sq Mean Sq NumDF  DenDF F value    Pr(>F)
## HN              495.3   495.3      1  852.92  2.0477  0.152801
## APOE            7675.9  3838.0      2  852.92 15.8673 1.714e-07 ***
## Component     10227.5 10227.5      1  598.00 42.2836 1.668e-10 ***
## HN:APOE        2692.2  1346.1      2  852.92  5.5652  0.003969 **
## HN:Component    110.3   110.3      1  598.00  0.4561  0.499733
## APOE:Component  1882.5   941.3      2  598.00  3.8915  0.020933 *
## HN:APOE:Component  331.7   165.9      2  598.00  0.6857  0.504109
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

## Type III Analysis of Variance Table with Satterthwaite's method
##              Sum Sq Mean Sq NumDF  DenDF F value
## Age              542.1   542.1      1 780.51  1.2171
## Genotype        2009.5   401.9      5 780.51  0.9023
## Sex              272.1   272.1      1 780.51  0.6109
## Treatment        768.9   768.9      1 780.51  1.7262
## Component     10147.0 10147.0      1 563.00 22.7812
## Age:Genotype     556.2   111.2      5 780.51  0.2497
## Age:Sex          212.7   212.7      1 780.51  0.4776
## Genotype:Sex     1739.9   348.0      5 780.51  0.7813
## Age:Treatment     58.0    58.0      1 780.51  0.1302

```

## Genotype:Treatment	1052.4	210.5	5 780.51	0.4725
## Sex:Treatment	183.2	183.2	1 780.51	0.4113
## Age:Component	208.8	208.8	1 563.00	0.4689
## Genotype:Component	1744.3	348.9	5 563.00	0.7832
## Sex:Component	39.2	39.2	1 563.00	0.0879
## Treatment:Component	524.8	524.8	1 563.00	1.1782
## Age:Genotype:Sex	1597.9	319.6	5 780.51	0.7175
## Age:Genotype:Treatment	135.7	67.9	2 780.51	0.1524
## Age:Sex:Treatment	366.2	366.2	1 780.51	0.8221
## Genotype:Sex:Treatment	1768.0	353.6	5 780.51	0.7939
## Age:Genotype:Component	504.6	100.9	5 563.00	0.2266
## Age:Sex:Component	1.0	1.0	1 563.00	0.0024
## Genotype:Sex:Component	445.2	89.0	5 563.00	0.1999
## Age:Treatment:Component	0.5	0.5	1 563.00	0.0010
## Genotype:Treatment:Component	347.4	69.5	5 563.00	0.1560
## Sex:Treatment:Component	479.4	479.4	1 563.00	1.0763
## Age:Genotype:Sex:Treatment	118.7	118.7	1 780.51	0.2664
## Age:Genotype:Sex:Component	663.2	132.6	5 563.00	0.2978
## Age:Genotype:Treatment:Component	26.9	13.4	2 563.00	0.0302
## Age:Sex:Treatment:Component	4.0	4.0	1 563.00	0.0089
## Genotype:Sex:Treatment:Component	2001.1	400.2	5 563.00	0.8985
## Age:Genotype:Sex:Treatment:Component	0.0	0.0	1 563.00	0.0000
##	Pr(>F)			
## Age	0.2703			
## Genotype	0.4789			
## Sex	0.4347			
## Treatment	0.1893			
## Component	2.317e-06 ***			
## Age:Genotype	0.9400			
## Age:Sex	0.4897			
## Genotype:Sex	0.5633			
## Age:Treatment	0.7183			
## Genotype:Treatment	0.7969			
## Sex:Treatment	0.5215			
## Age:Component	0.4938			
## Genotype:Component	0.5620			
## Sex:Component	0.7670			
## Treatment:Component	0.2782			
## Age:Genotype:Sex	0.6104			
## Age:Genotype:Treatment	0.8587			
## Age:Sex:Treatment	0.3648			
## Genotype:Sex:Treatment	0.5542			
## Age:Genotype:Component	0.9510			
## Age:Sex:Component	0.9613			
## Genotype:Sex:Component	0.9625			
## Age:Treatment:Component	0.9742			
## Genotype:Treatment:Component	0.9782			
## Sex:Treatment:Component	0.3000			
## Age:Genotype:Sex:Treatment	0.6059			
## Age:Genotype:Sex:Component	0.9141			
## Age:Genotype:Treatment:Component	0.9703			
## Age:Sex:Treatment:Component	0.9249			
## Genotype:Sex:Treatment:Component	0.4817			
## Age:Genotype:Sex:Treatment:Component	0.9945			

```

## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

## Type III Analysis of Variance Table with Satterthwaite's method
##              Sum Sq Mean Sq NumDF  DenDF F value    Pr(>F)
## HN              2312.0  2312.0      1  858.01  5.3355  0.02113 *
## APOE              359.5   179.8      2  858.01  0.4148  0.66058
## Component      24890.6 24890.6      1  598.00 57.4416 1.333e-13 ***
## HN:APOE           70.2    35.1      2  858.01  0.0810  0.92218
## HN:Component      597.5   597.5      1  598.00  1.3789  0.24076
## APOE:Component    1465.2   732.6      2  598.00  1.6907  0.18527
## HN:APOE:Component  532.7   266.3      2  598.00  0.6146  0.54119
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

## Analysis of Variance Table
##
## Response: Pct.Total.Time.Freezing
##              Df Sum Sq Mean Sq F value    Pr(>F)
## Age              1  13539  13538.5 88.6893 < 2.2e-16 ***
## Genotype          5  24295  4859.1 31.8313 < 2.2e-16 ***
## Sex              1    442   441.7  2.8934  0.090133 .
## Treatment         1    207   207.4  1.3586  0.244845
## Age:Genotype      5   7750  1550.0 10.1542 6.682e-09 ***
## Age:Sex           1    140   140.1  0.9176  0.338996
## Genotype:Sex      5    697   139.3  0.9126  0.473304
## Age:Treatment     1    221   221.4  1.4503  0.229577
## Genotype:Treatment 5   4921   984.2  6.4472 1.140e-05 ***
## Sex:Treatment     1    145   144.9  0.9495  0.330740
## Age:Genotype:Sex  5    377    75.5  0.4945  0.780238
## Age:Genotype:Treatment 2  1513   756.3  4.9548 0.007727 **
## Age:Sex:Treatment 1     5     5.5  0.0360  0.849723
## Genotype:Sex:Treatment 5   2844   568.9  3.7267 0.002823 **
## Age:Genotype:Sex:Treatment 1   177   177.1  1.1603 0.282393
## Residuals        261  39842   152.7
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

## Analysis of Variance Table
##
## Response: Pct.Total.Time.Freezing
##              Df Sum Sq Mean Sq F value    Pr(>F)
## HN              1   3692  3691.8 14.289 0.0001896 ***
## APOE              2  11713  5856.5 22.668 6.925e-10 ***
## HN:APOE          2   5235  2617.6 10.132 5.546e-05 ***
## Residuals      296  76476   258.4
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

## Analysis of Variance Table
##
## Response: Freeze.Cnt
##              Df Sum Sq Mean Sq F value    Pr(>F)

```

```

## Age                1  23230 23230.3 39.9439 1.124e-09 ***
## Genotype           5  44242  8848.3 15.2144 4.045e-13 ***
## Sex                1    489   489.0  0.8408  0.36001
## Treatment          1    682   682.4  1.1734  0.27971
## Age:Genotype       5   3390   678.0  1.1658  0.32641
## Age:Sex            1    315   314.6  0.5409  0.46271
## Genotype:Sex       5   5102  1020.5  1.7547  0.12259
## Age:Treatment      1    999   999.1  1.7179  0.19112
## Genotype:Treatment 5   6577  1315.4  2.2617  0.04879 *
## Sex:Treatment      1     0     0.1  0.0002  0.98972
## Age:Genotype:Sex   5   5846  1169.3  2.0105  0.07753 .
## Age:Genotype:Treatment 2  1075   537.4  0.9240  0.39821
## Age:Sex:Treatment  1     1     1.1  0.0019  0.96565
## Genotype:Sex:Treatment 5  7032  1406.5  2.4184  0.03636 *
## Age:Genotype:Sex:Treatment 1   909   908.7  1.5625  0.21242
## Residuals          261 151791   581.6
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

## Analysis of Variance Table
##
## Response: Freeze.Cnt
##              Df Sum Sq Mean Sq F value    Pr(>F)
## HN              1   8606  8606.0 11.9270 0.000634 ***
## APOE            2  16123  8061.5 11.1725  2.1e-05 ***
## HN:APOE         2  13371  6685.6  9.2656 0.000125 ***
## Residuals      296 213580    721.6
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```