

analysis

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2022-08-01

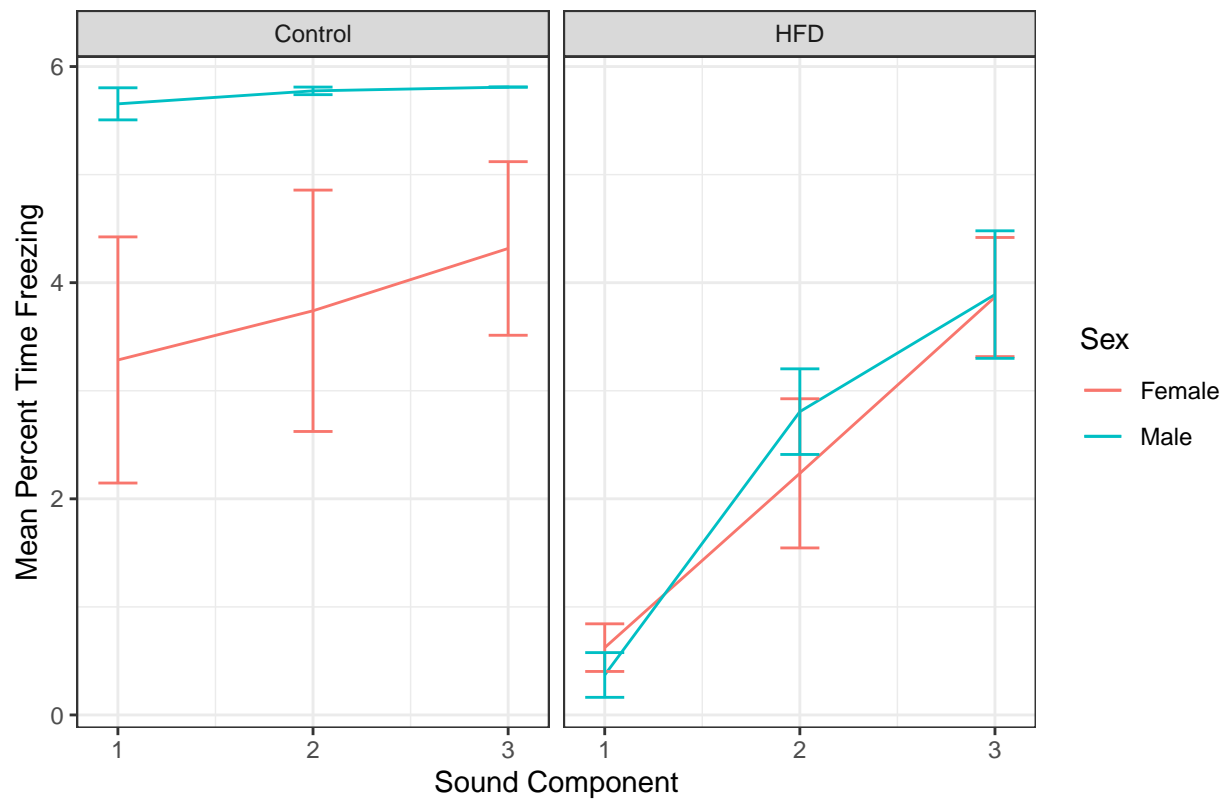
```
## [1] "Mice count"
```

```
## # A tibble: 6 x 7
```

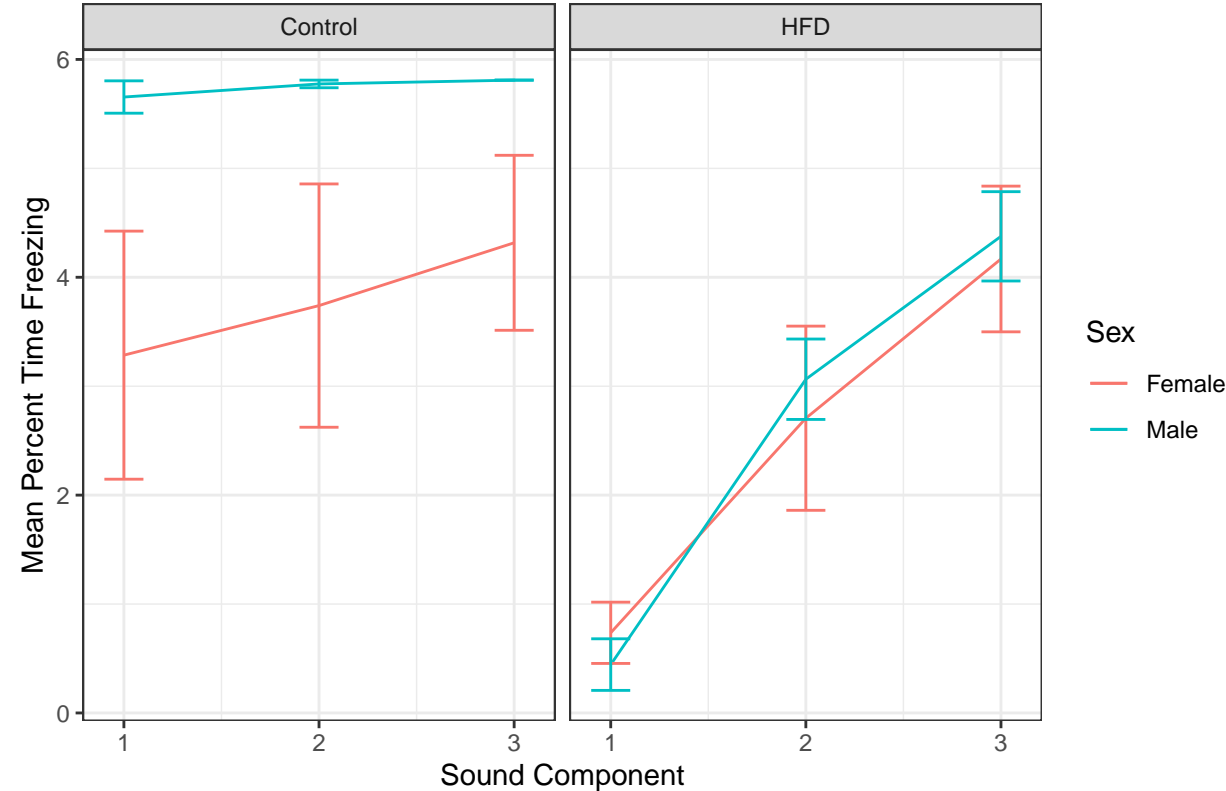
```
## # Groups:   Genotype, Treatment, Sex [4]
```

```
##   Genotype Treatment Sex    Age    Count    sd    len
##   <chr>    <chr>    <chr> <chr>    <int> <dbl> <dbl>
## 1 APOE22   Control   Female 12 months    6  0.493  11.6
## 2 APOE22   Control   Male   12 months    4  0.25   12.0
## 3 APOE22   HFD      Female 12 months    5  1.13   11.9
## 4 APOE22   HFD      Female 18 months    2  0      18.2
## 5 APOE22   HFD      Male   12 months    5  0.219  12.1
## 6 APOE22   HFD      Male   18 months    1 NA      18.2
```

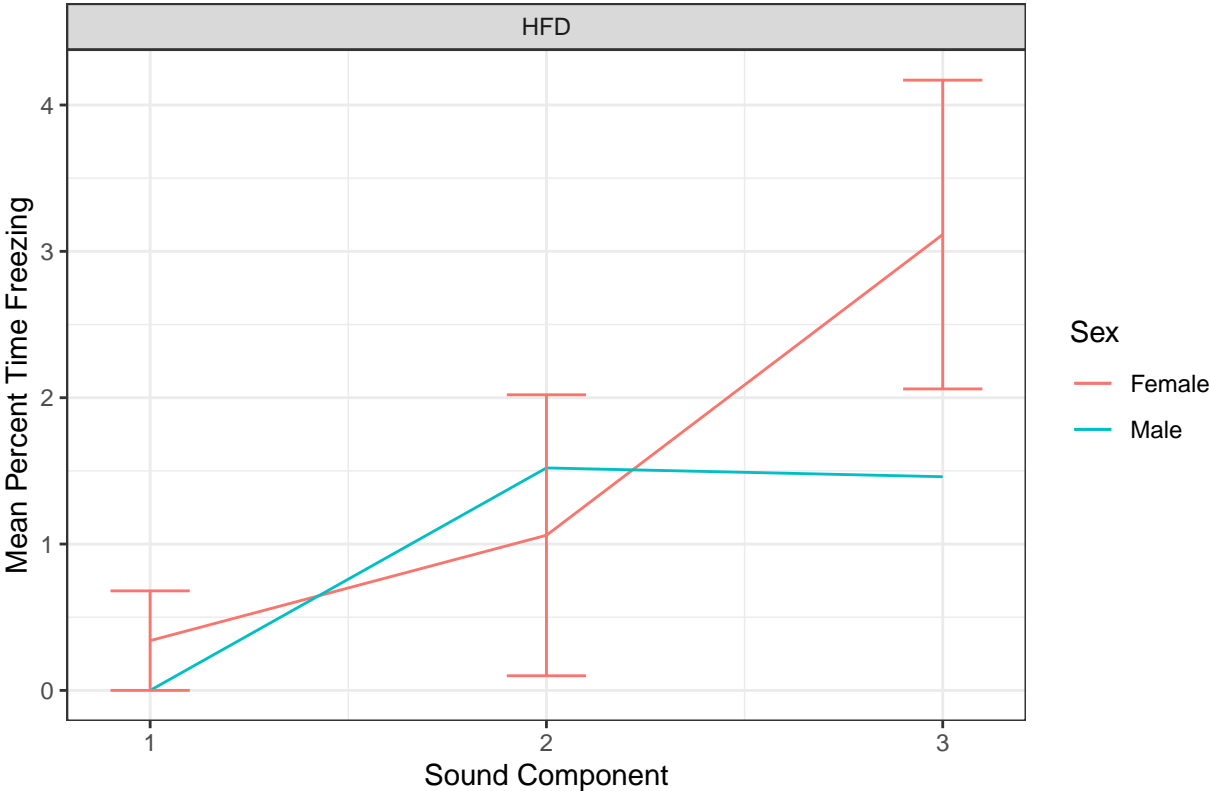
Day 0 Learning Curves by Genotype and Treatment



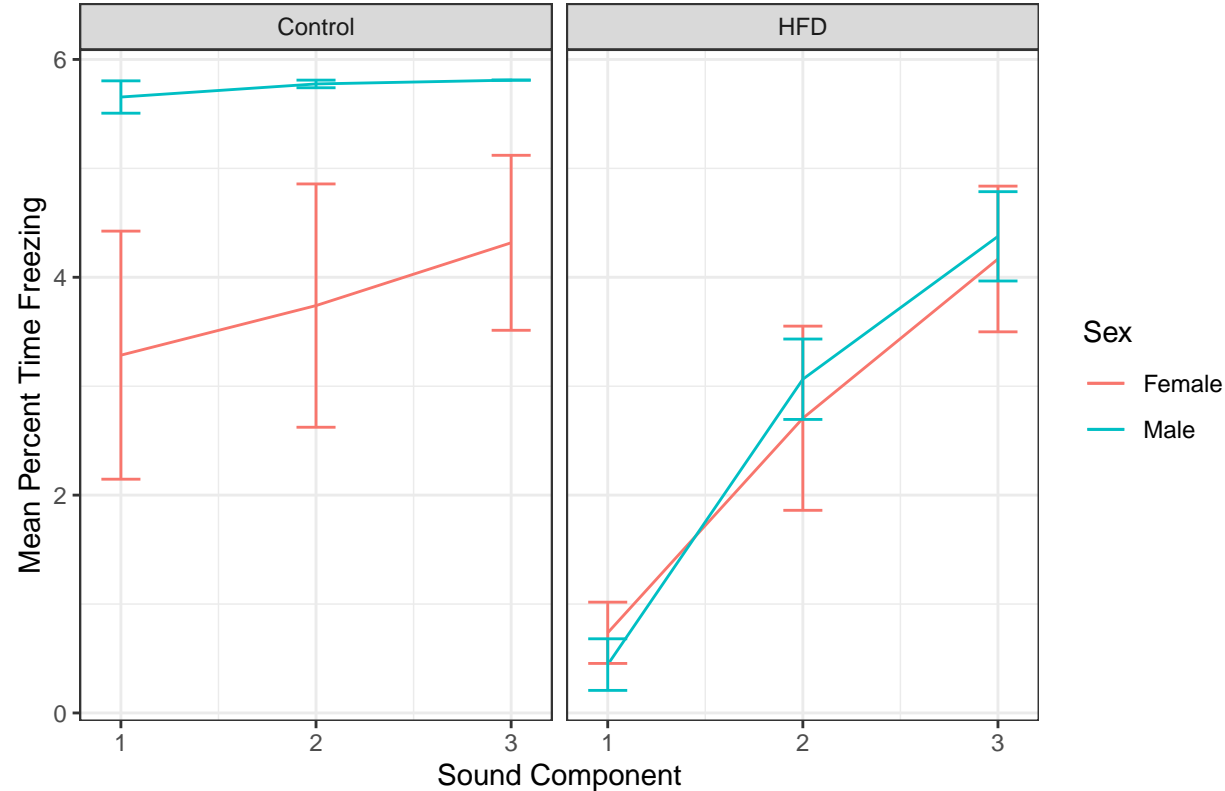
Day 0 12 Months Old Learning Curves by Genotype and Treatment



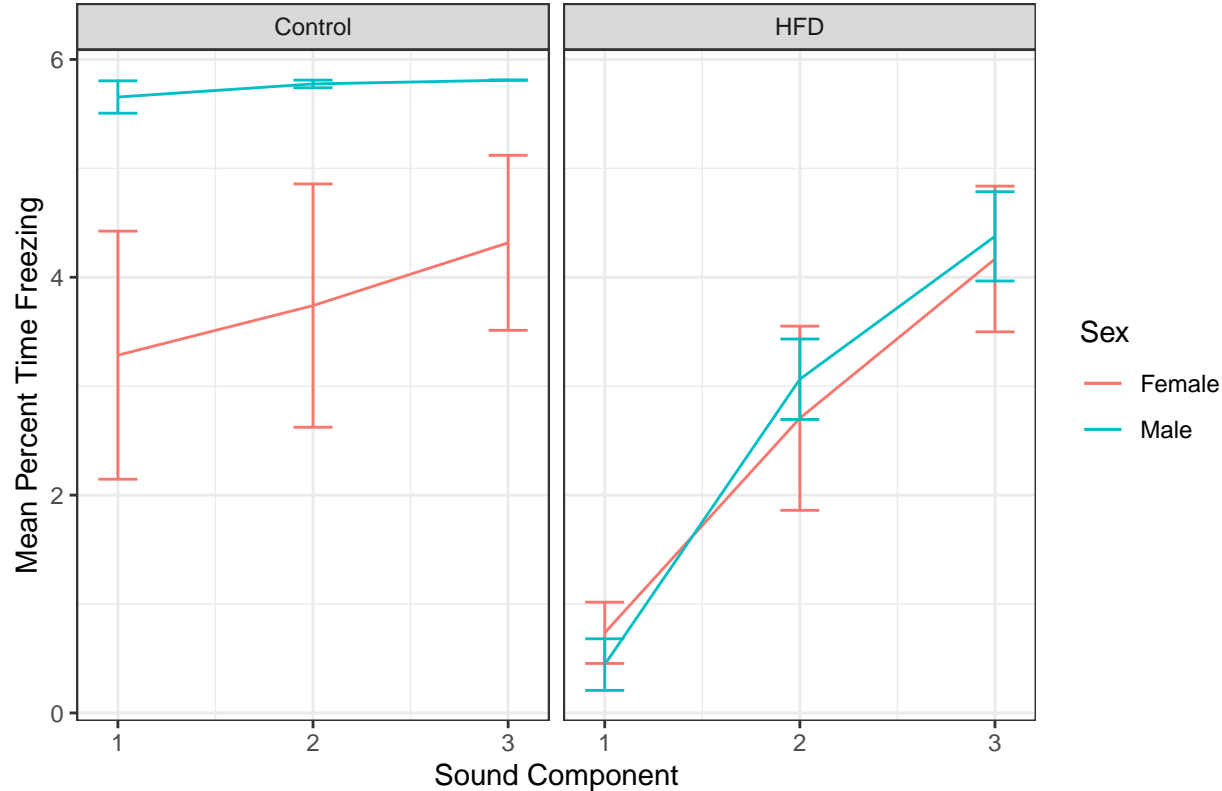
Day 0 18 Months Old Learning Curves by Genotype and Treatment



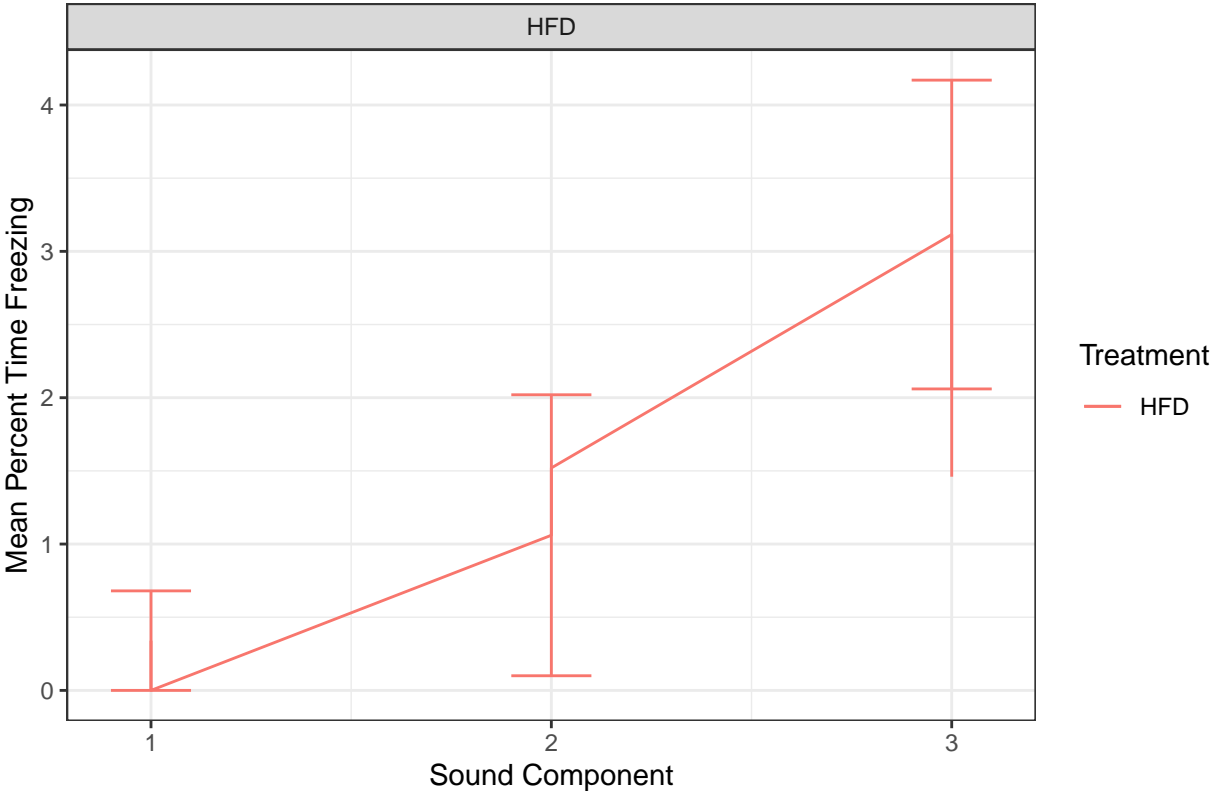
Day 0 12 Months Old Learning Curves by nonHN Genotype and Treatment



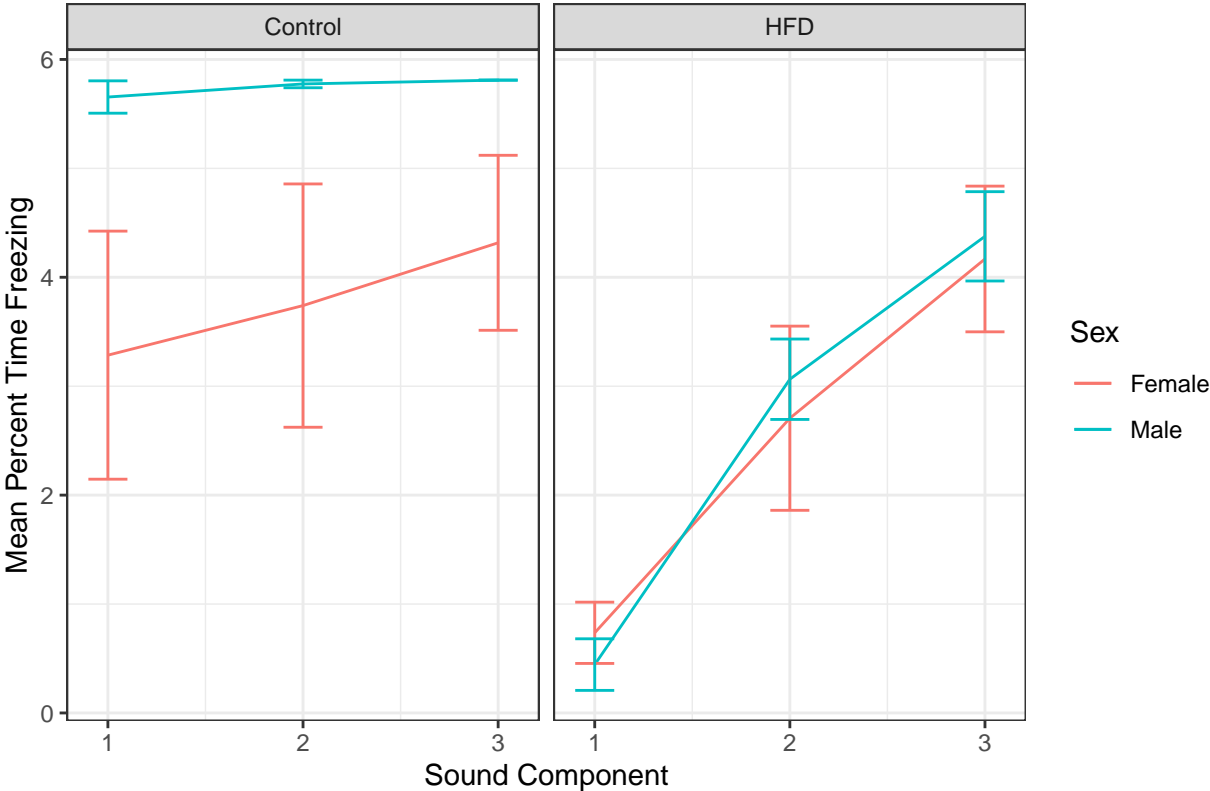
Day 0 12 Months Old Learning Curves by nonHN Genotype and Treatment



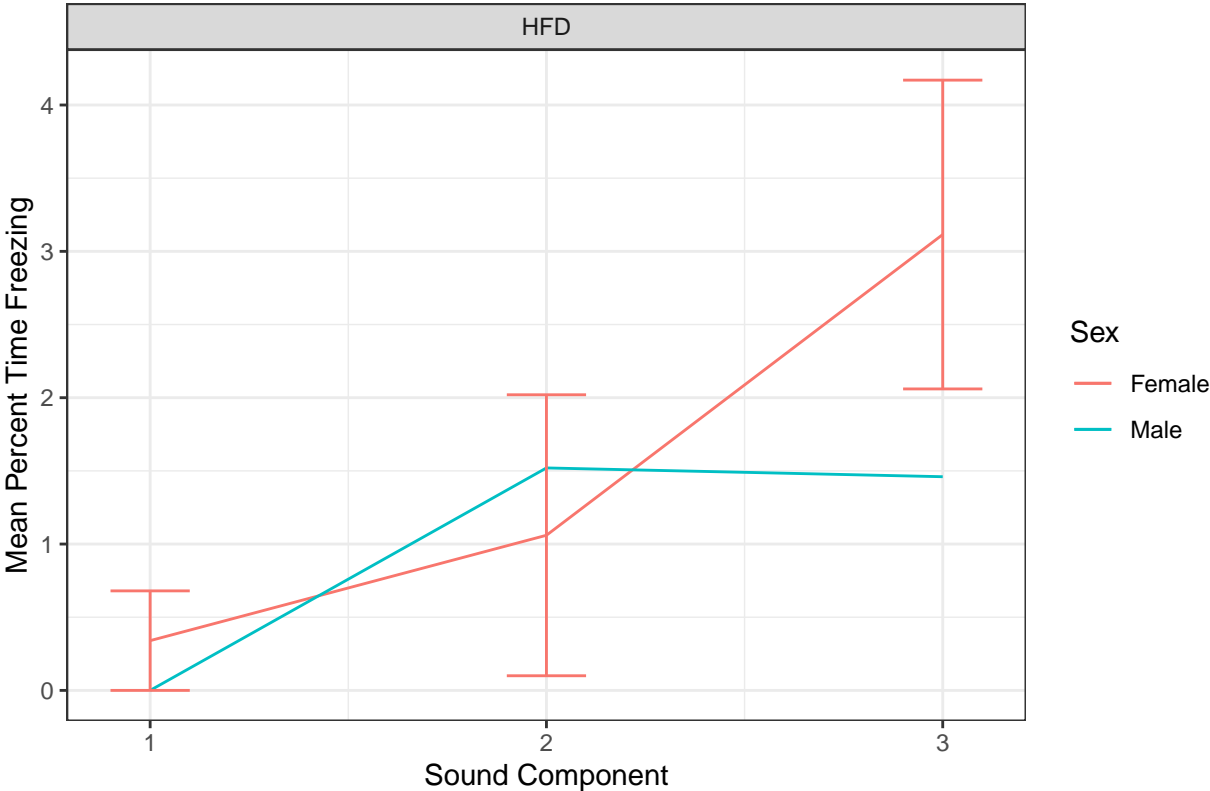
Day 0 18 Months Old Learning Curves by nonHN Genotype and Treatment



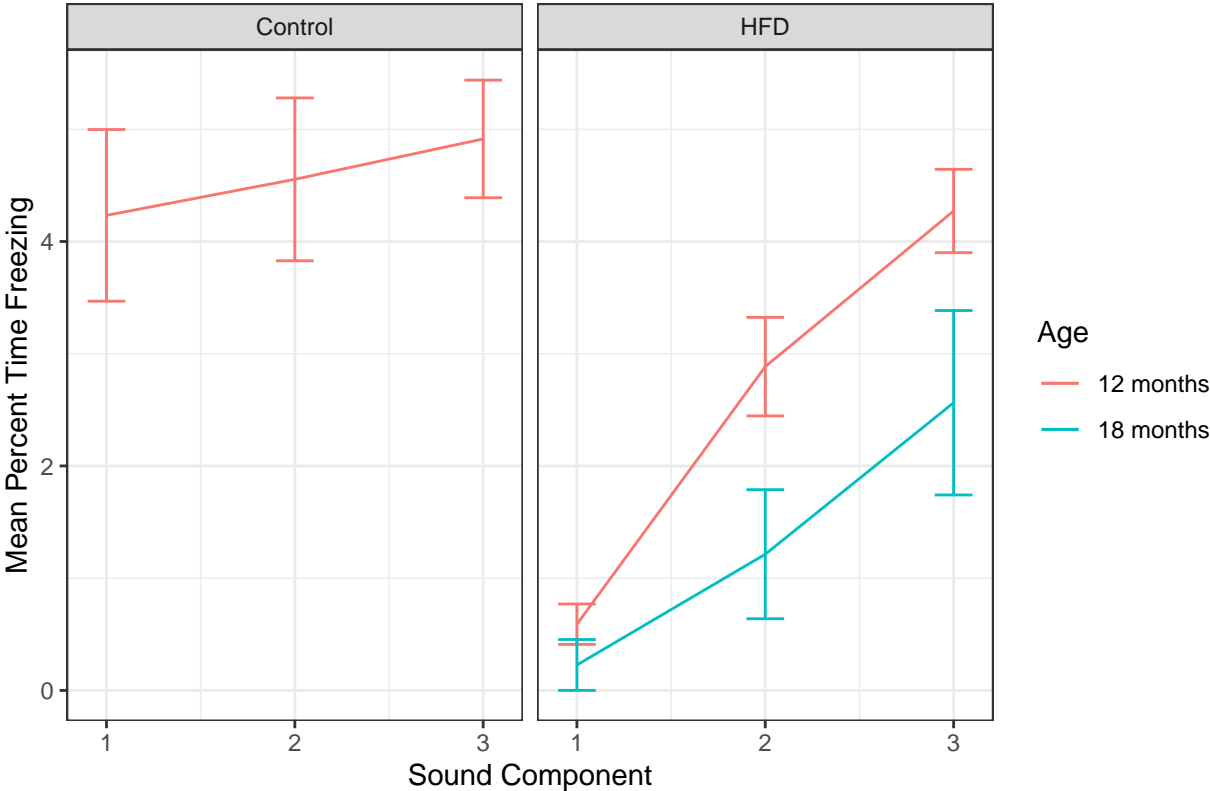
Day 0 12 Months Old nonHNs by Genotype, Sex, and Treatment



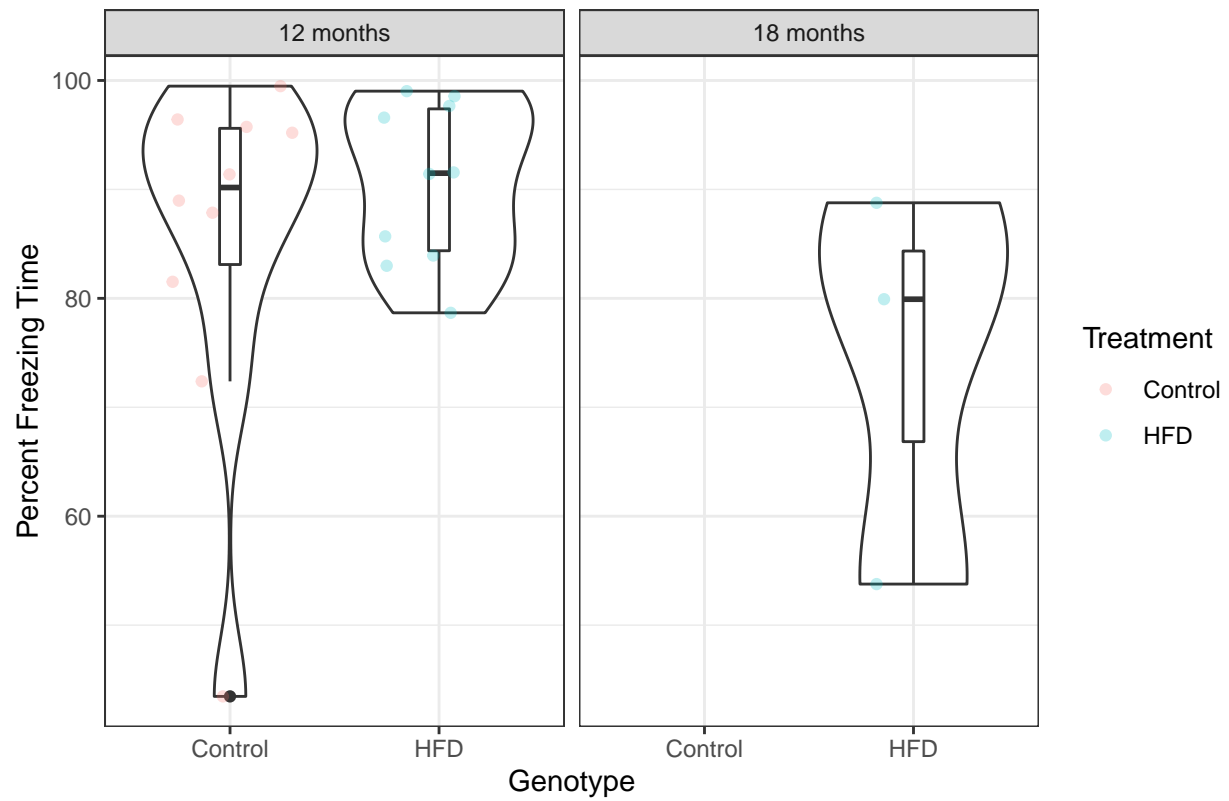
Day 0 18 Months Old nonHNs by Genotype, Sex, and Treatment



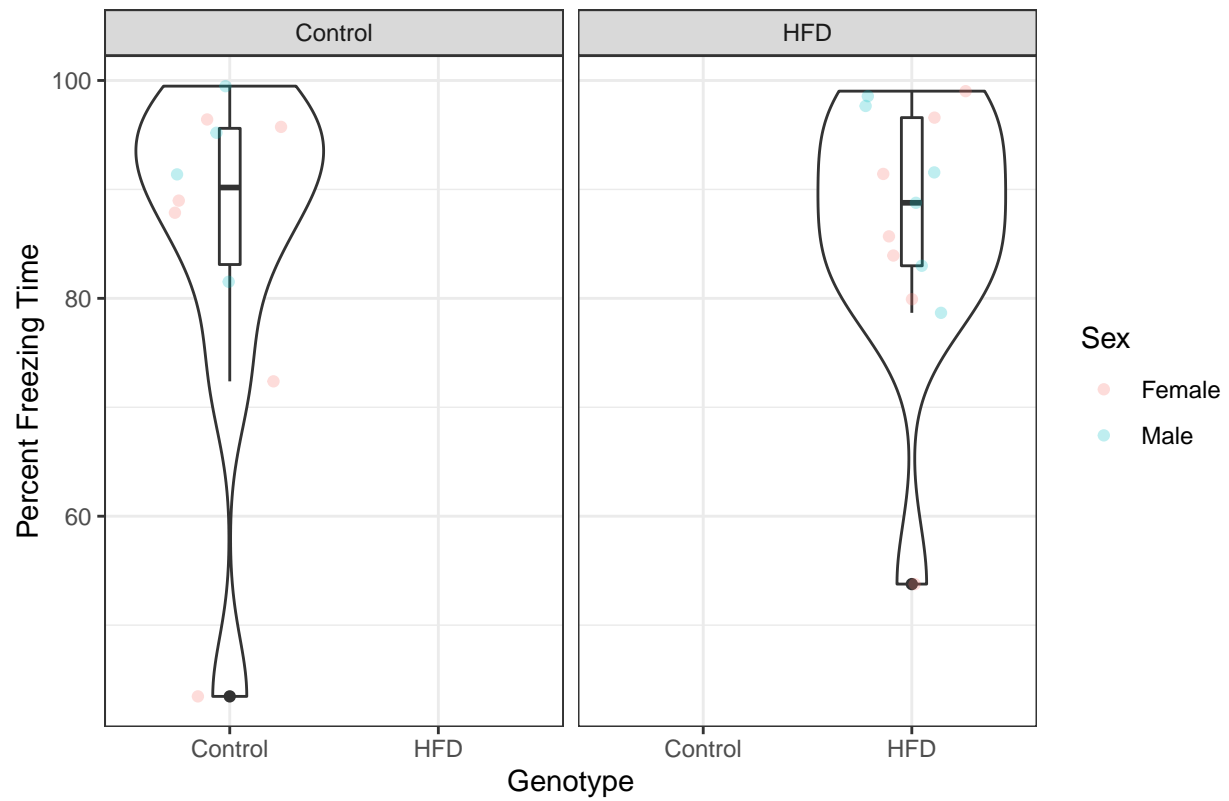
Day 0 Learning Curves by Age, nonHN Genotype, and Treatment

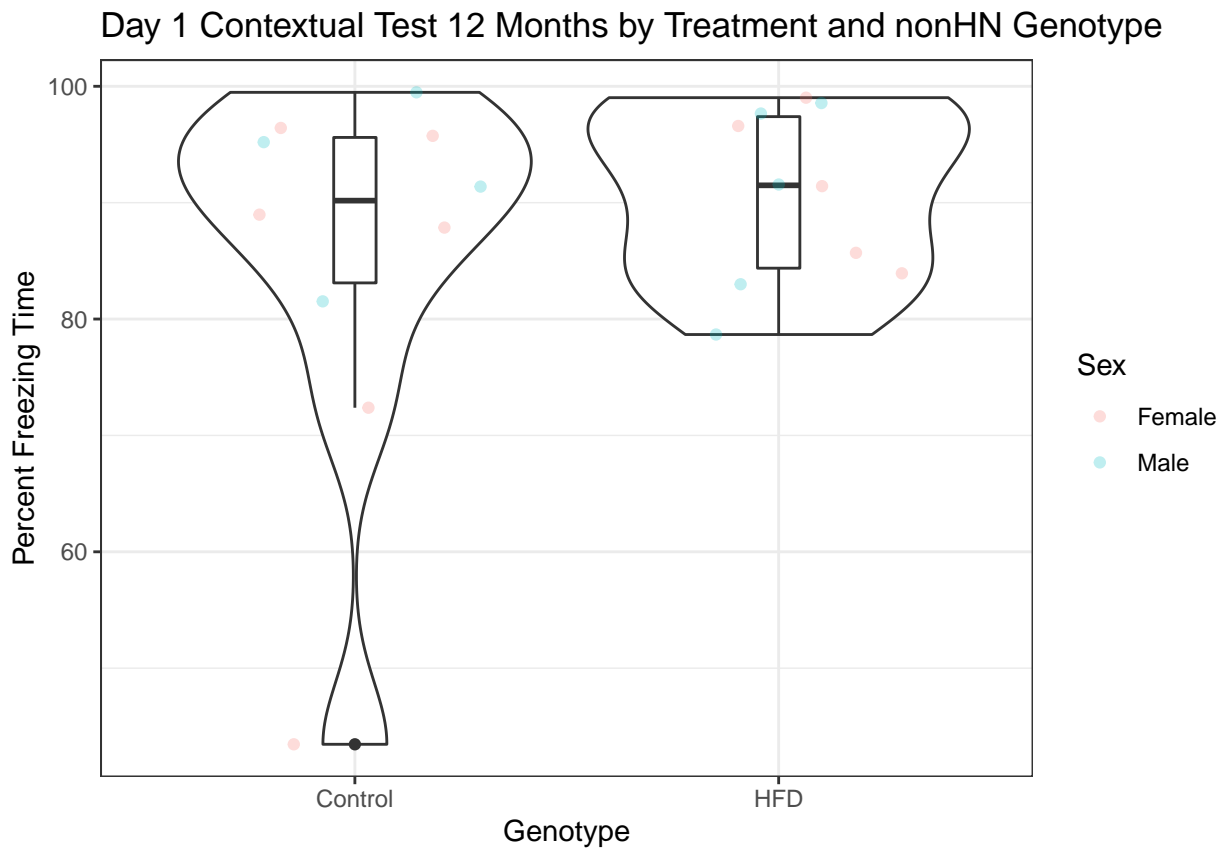


Day 1 Contextual Test by Treatment, Age, and nonHN Genotype



Day 1 Contextual Test All Ages by Treatment and nonHN Genotype

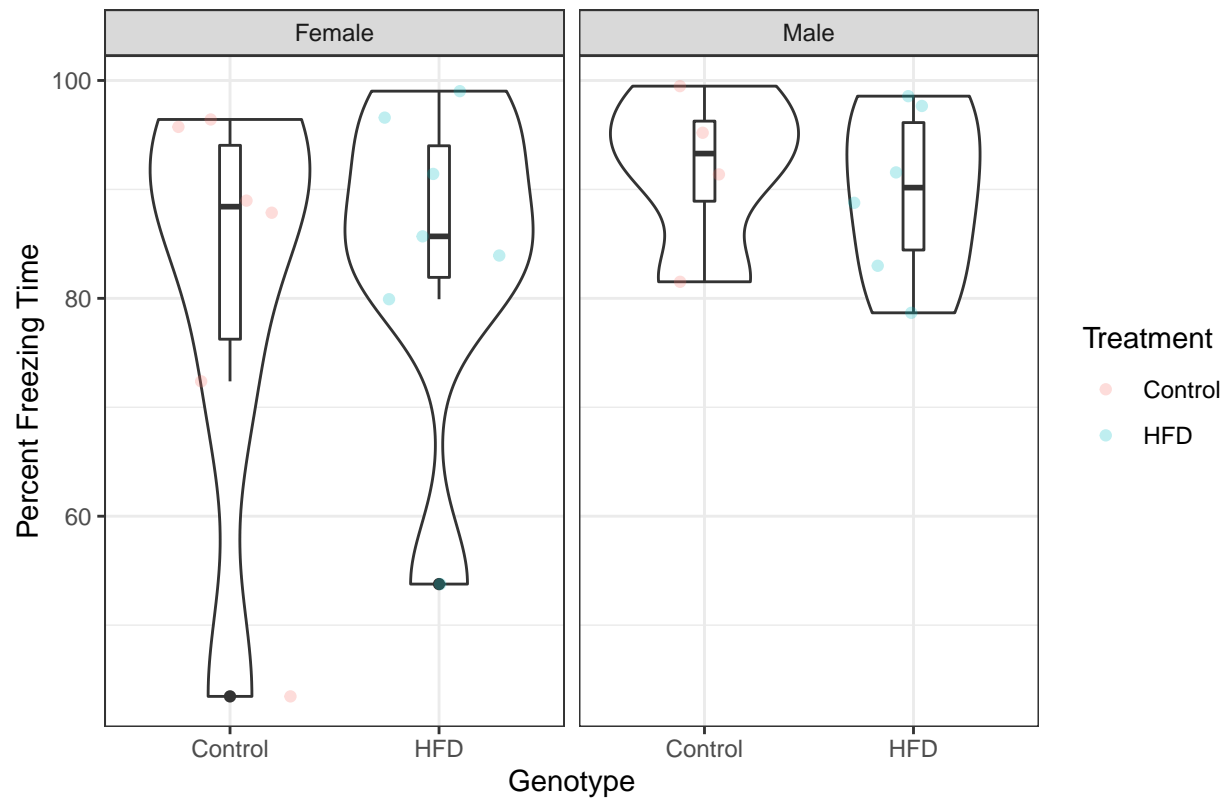




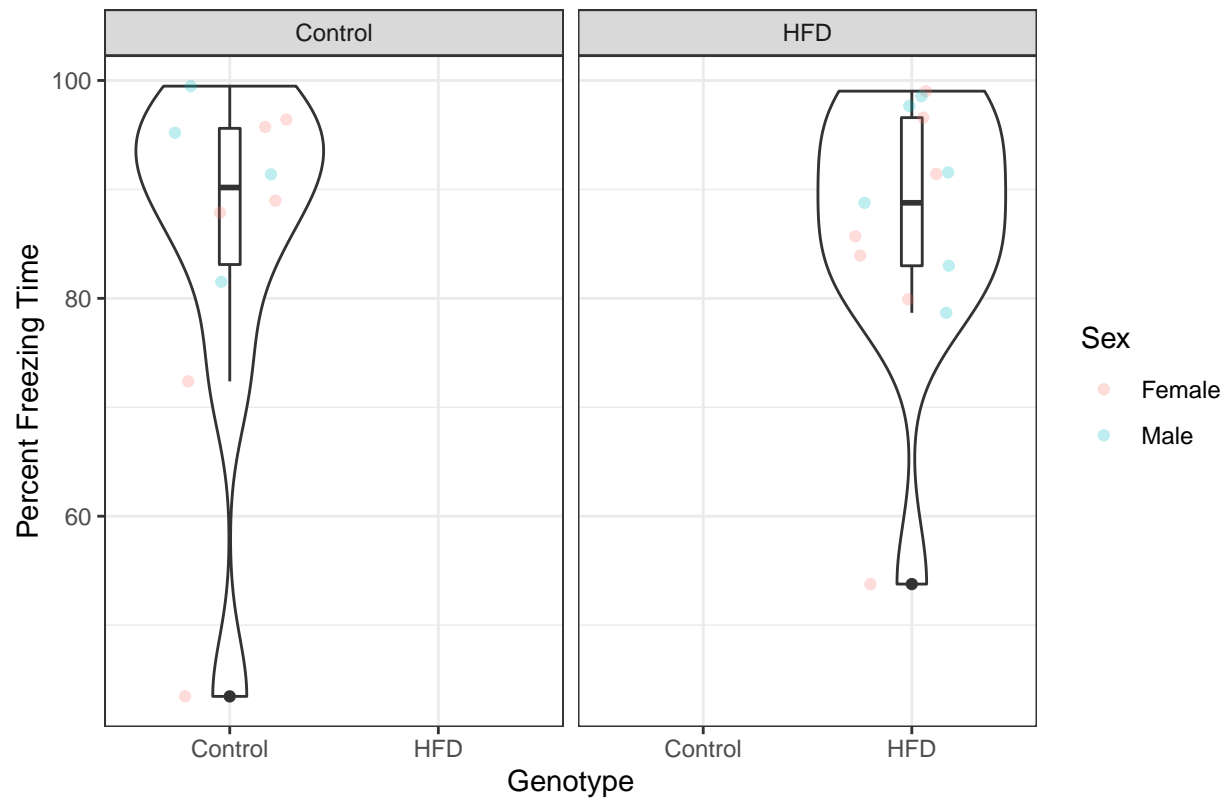
Day 1 Contextual Test 18 Months by Treatment and nonHN Genotype



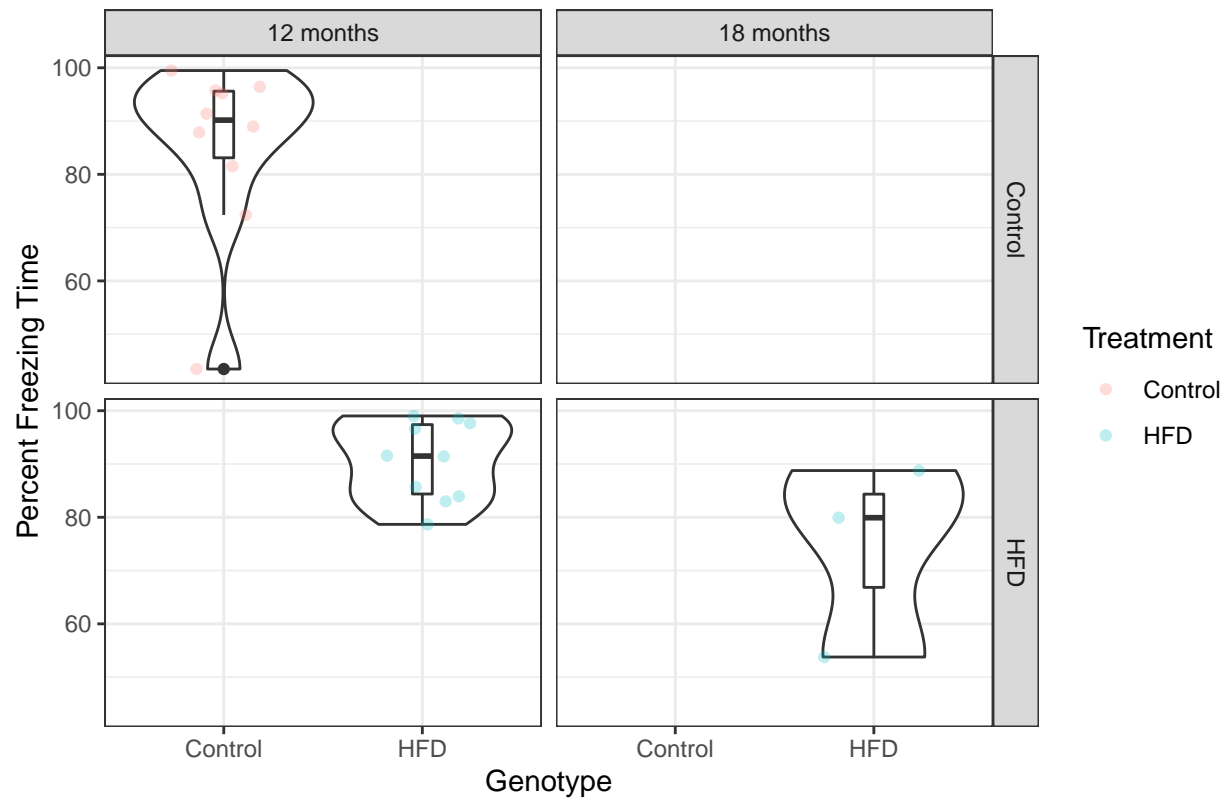
Day 1 Contextual Test All Ages by Sex and nonHN Genotype



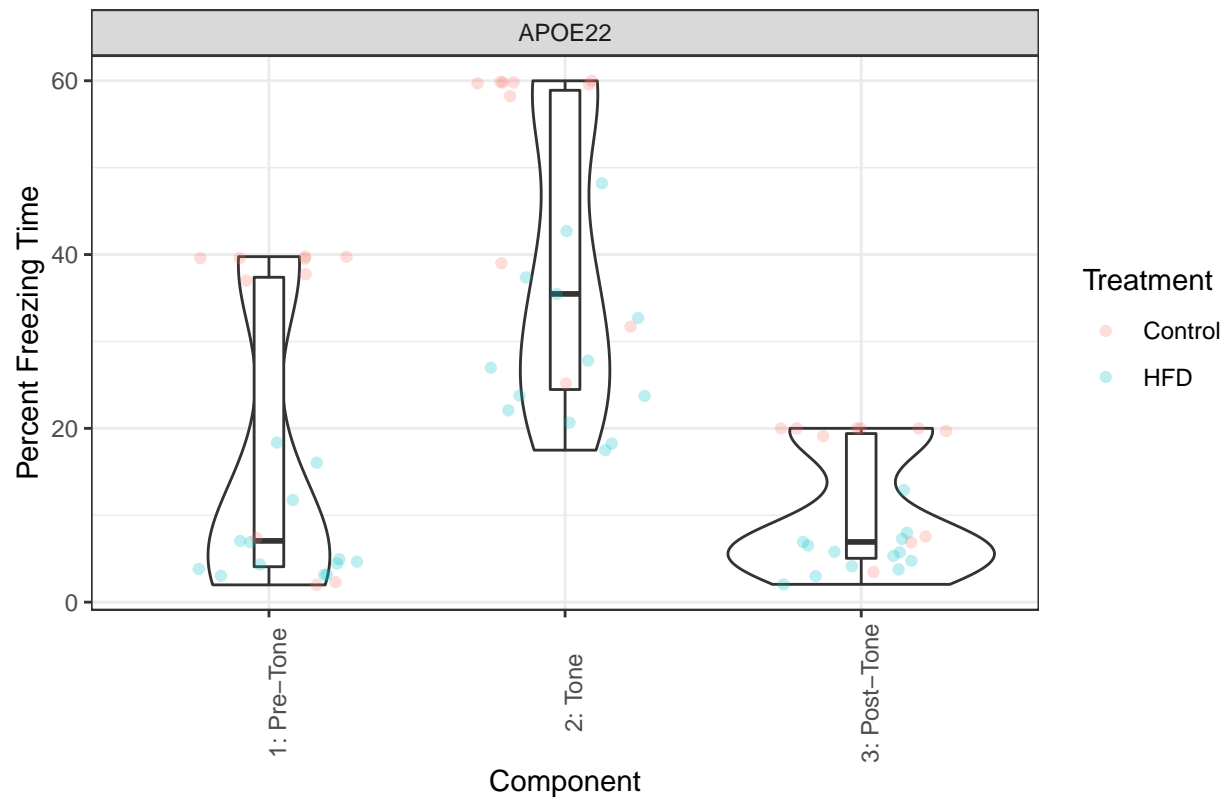
Day 1 Contextual Test All Ages by Treatment, Sex and nonHN Genotype



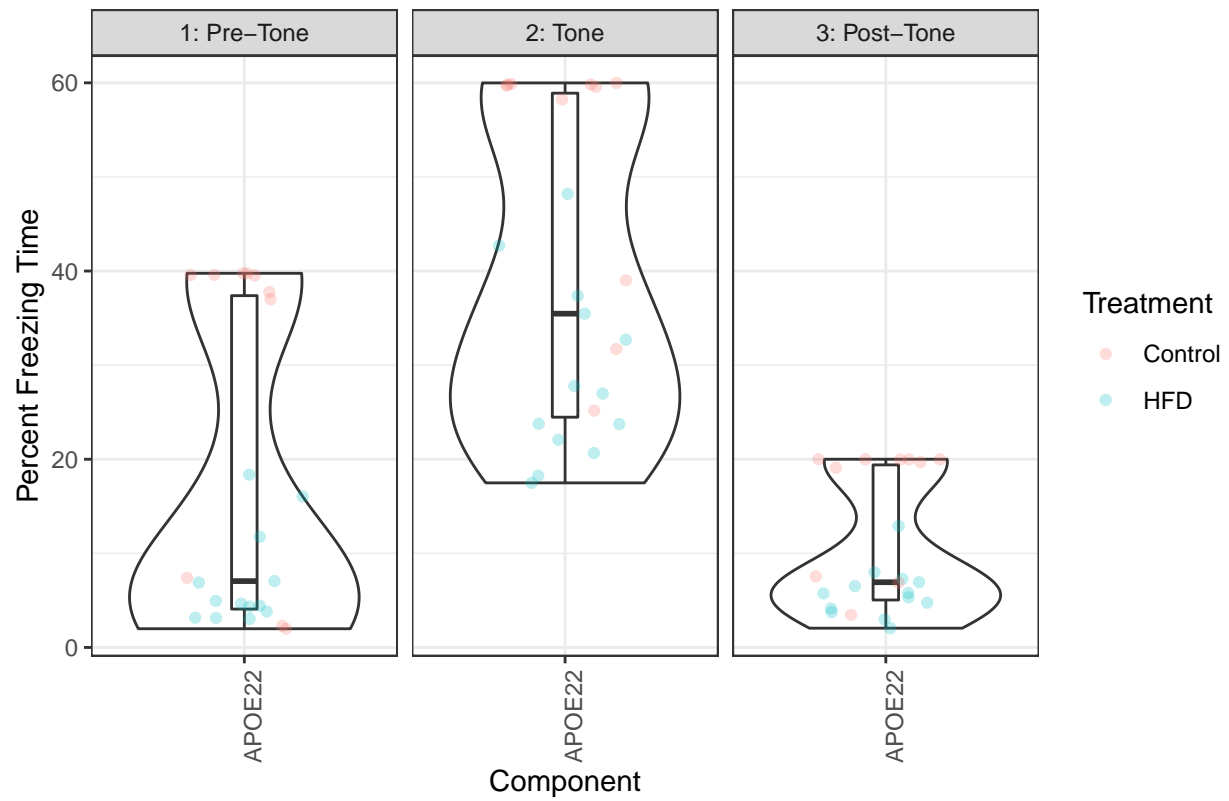
Day 1 Contextual Test Treatment, Age and nonHN Genotype



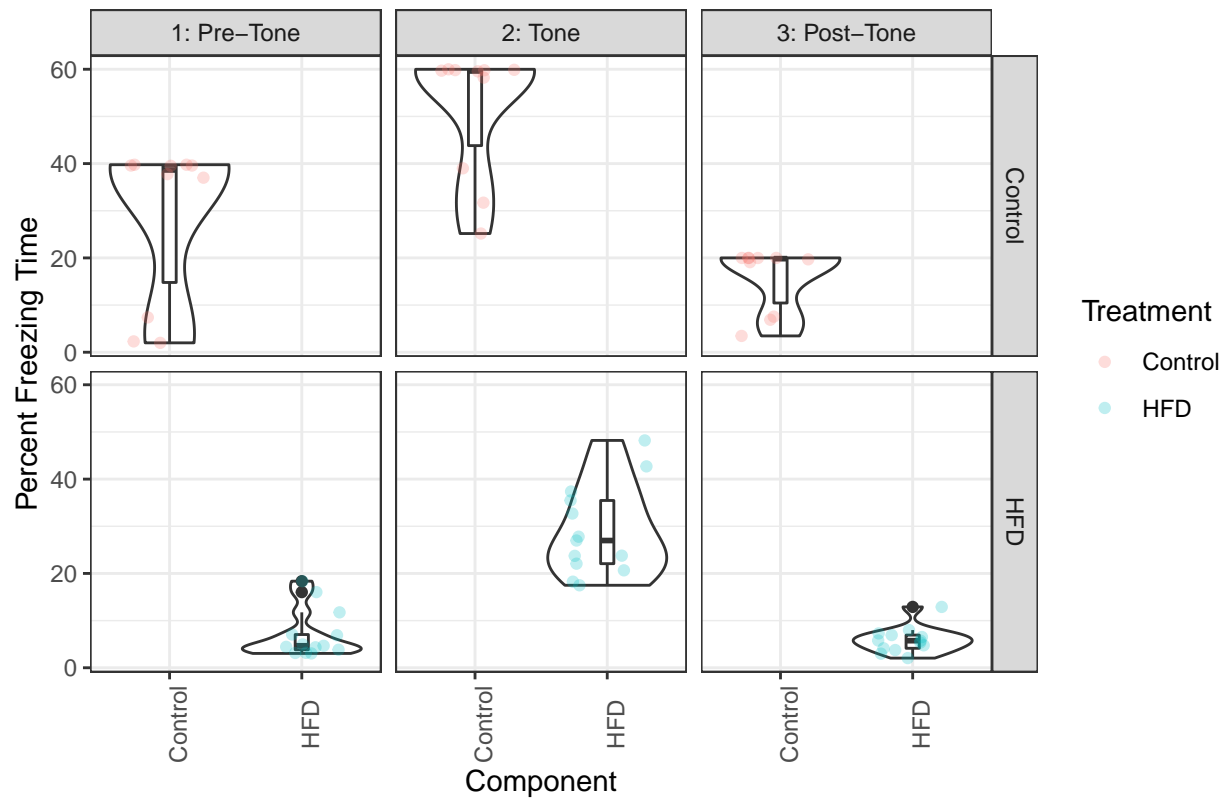
Day 2 All Ages Percent Freezing by Component and nonHN Genotype



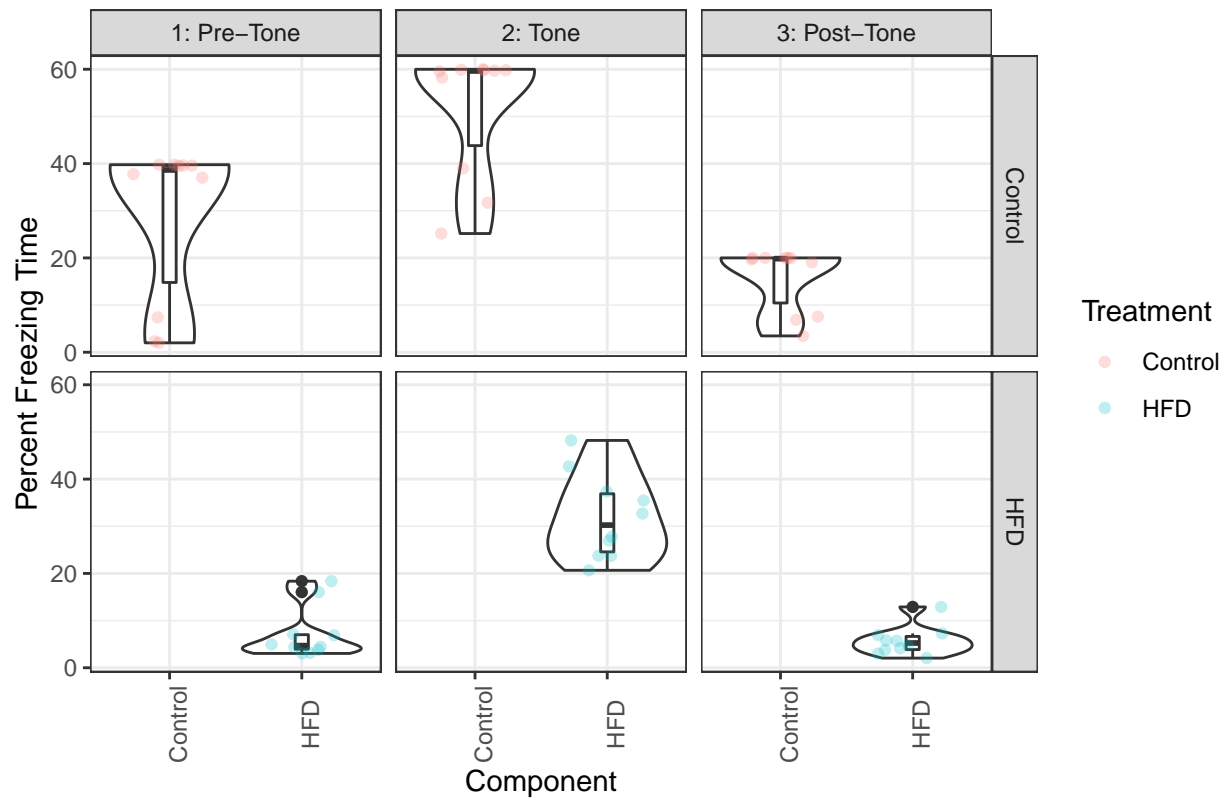
Day 2 All Ages Percent Freezing by Component and nonHN Genotype



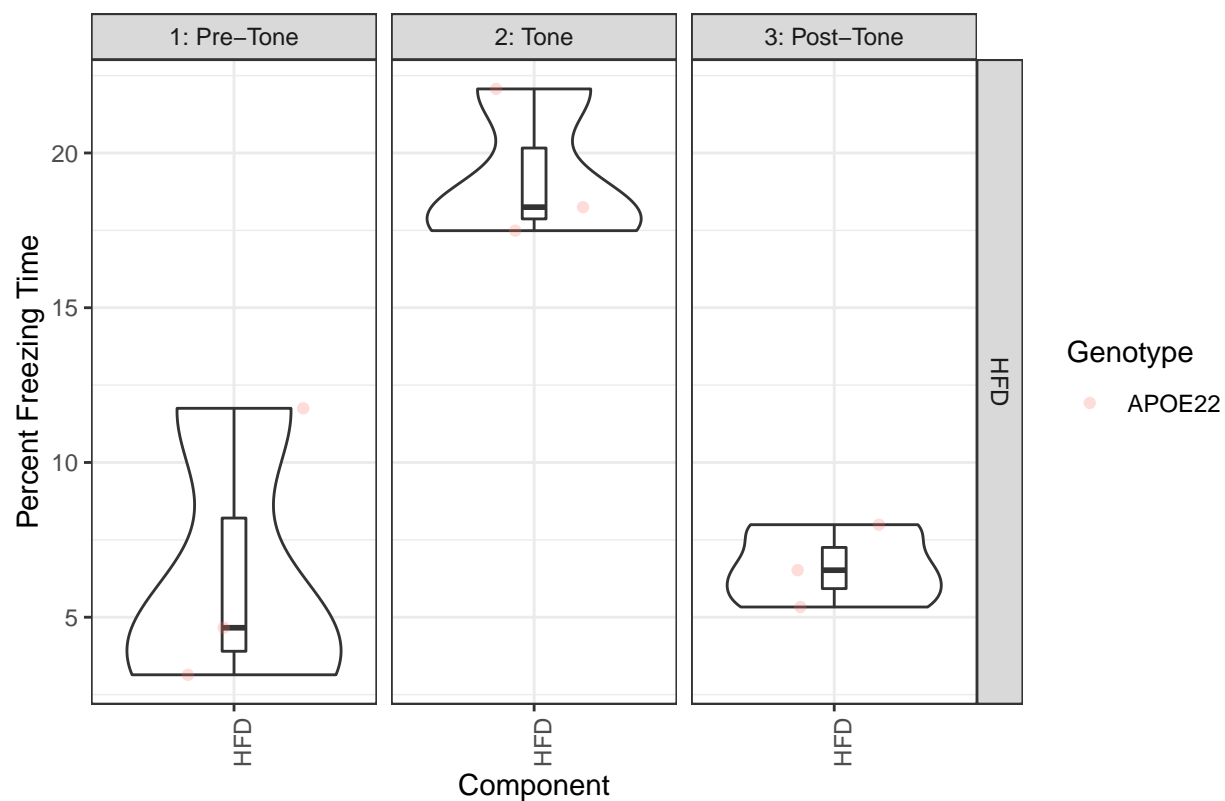
Day 2 All Ages Percent Freezing by Treatment, Component and nonHN Ger



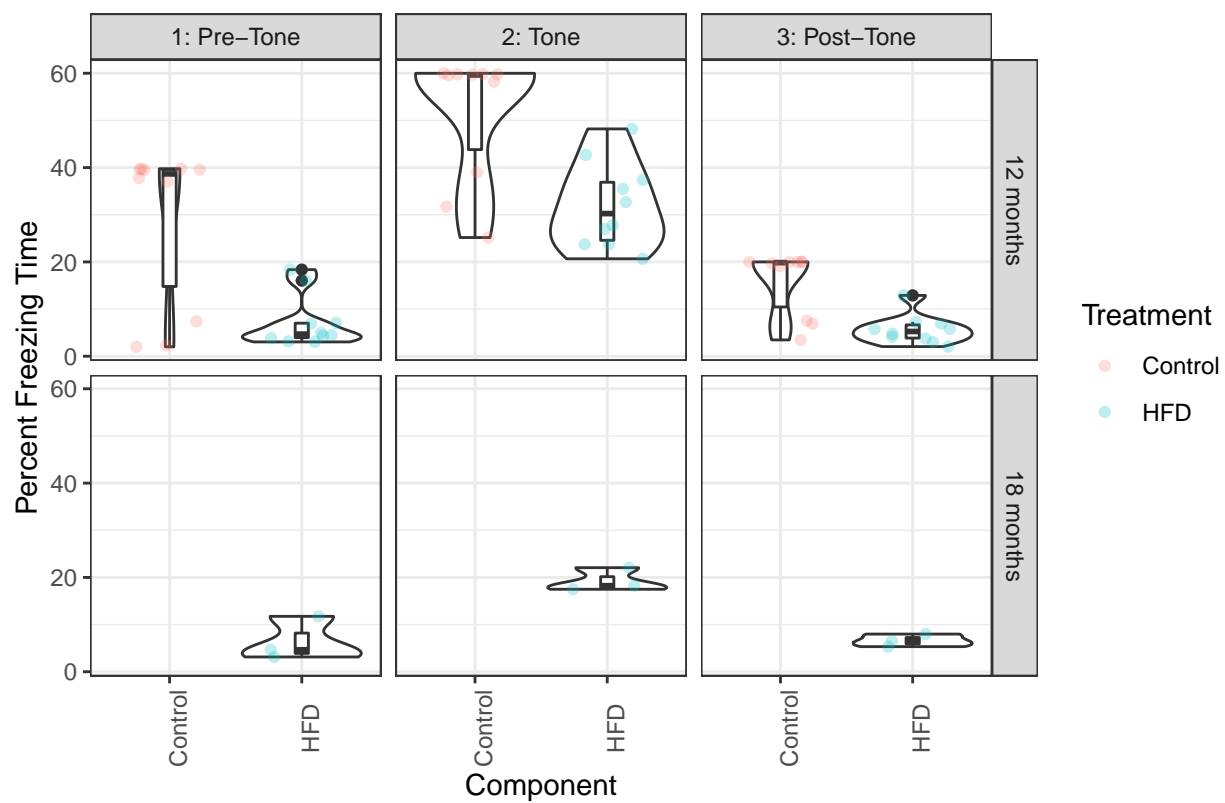
Day 2 12 Months Percent Freezing by Treatment, Component and nonHN G



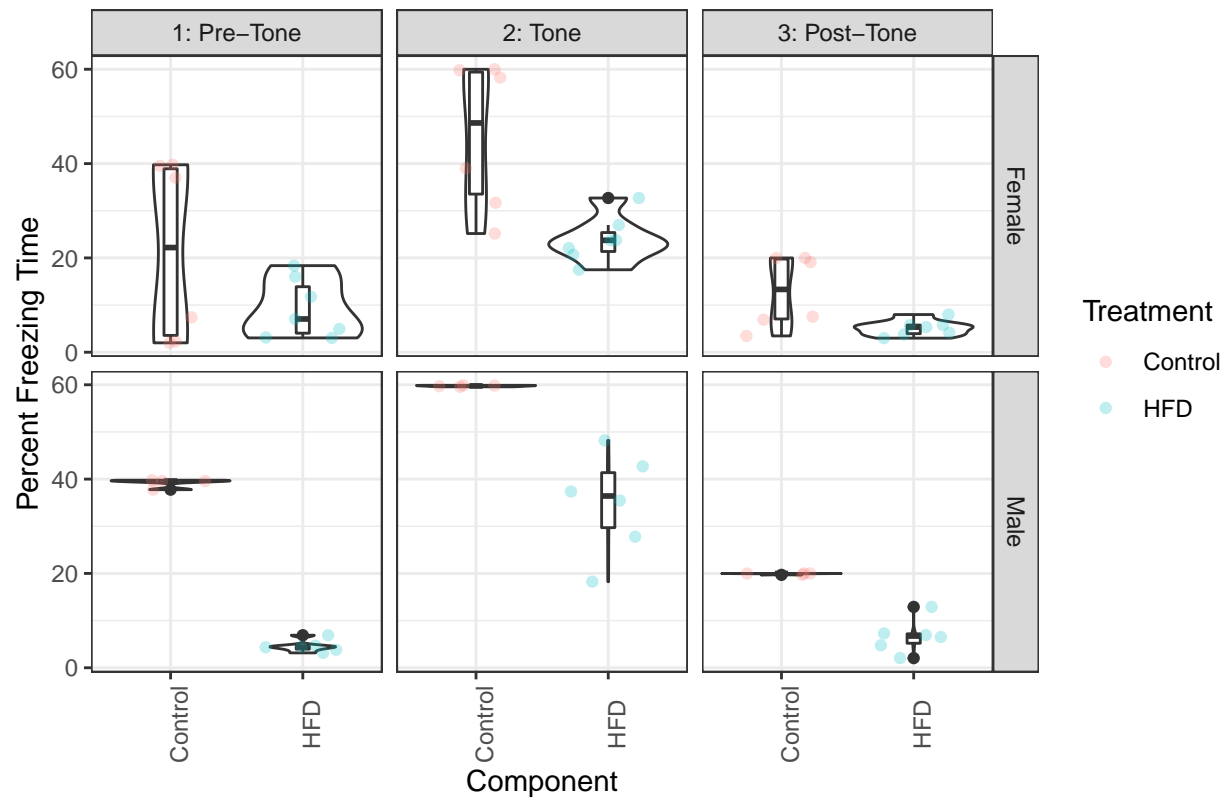
Day 2 18 Months Percent Freezing by Treatment, Component and nonHN G



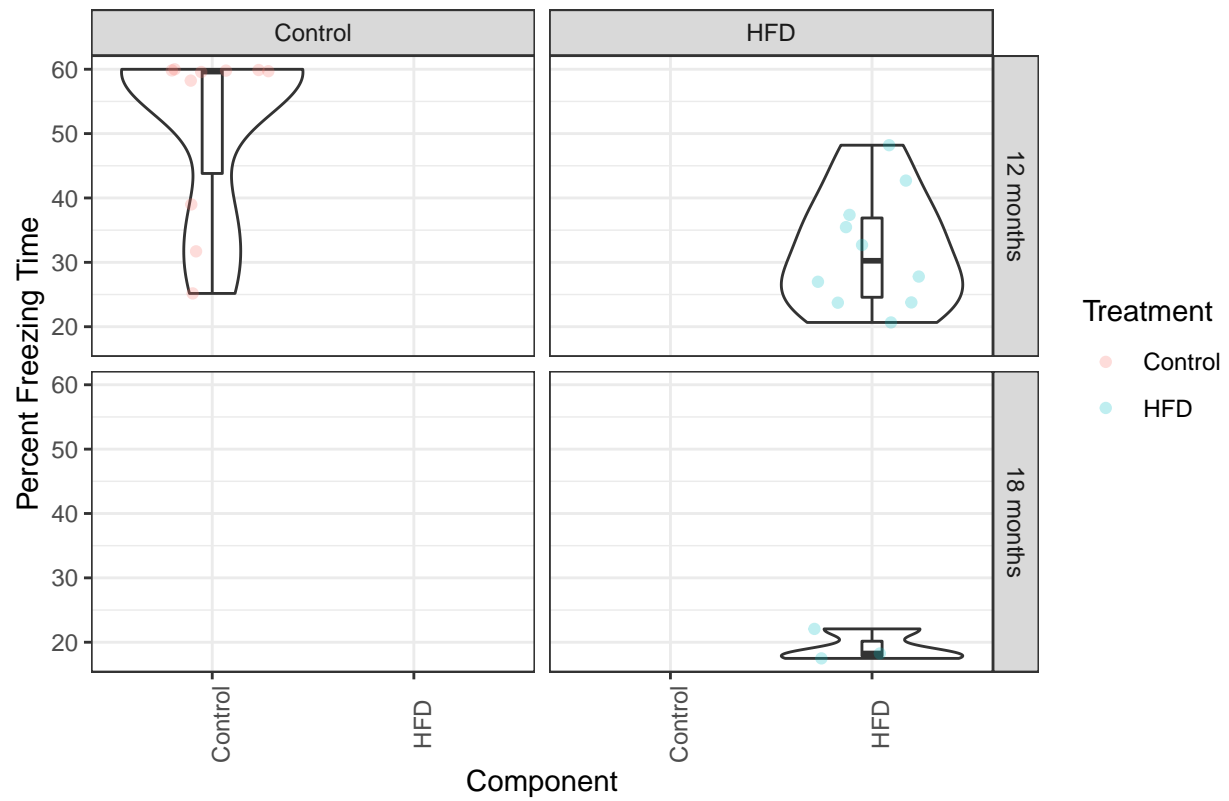
Day 2 Percent Freezing by Age, Component and nonHN Genotype



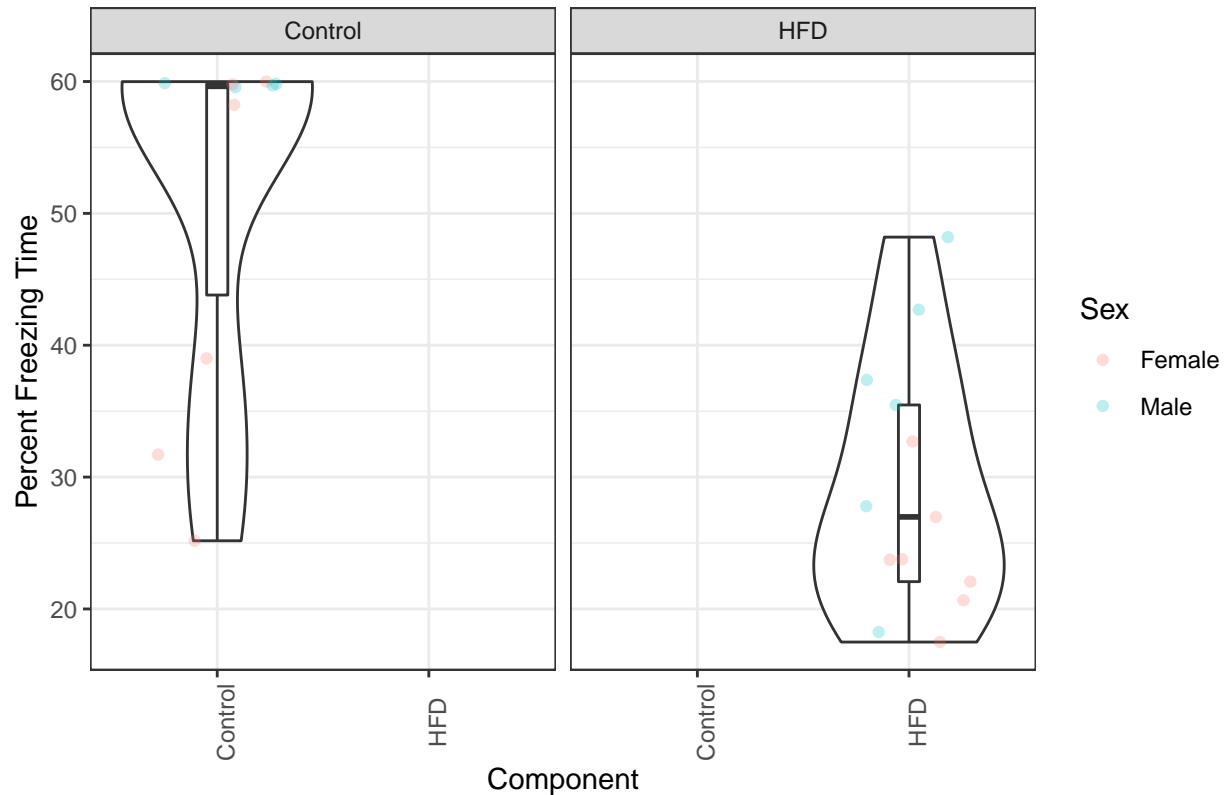
Day 2 All Ages Percent Freezing by Sex, Component and nonHN Genotype



Day 2 Percent Freezing Within Tone by Age and nonHN 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

```