

MWM Analysis

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```
## Analysis of Variance Table
##               npar    Sum Sq  Mean Sq F value
## Stage          1 0.264193 0.264193 24.8948
## APOE            6 0.139758 0.023293  2.1949
## Sex             1 0.000478 0.000478  0.0450
## Diet            1 0.111274 0.111274 10.4853
## Stage:APOE      6 0.032026 0.005338  0.5030
## Stage:Sex       1 0.031294 0.031294  2.9488
## APOE:Sex        6 0.112192 0.018699  1.7620
## Stage:Diet      1 0.022941 0.022941  2.1617
## APOE:Diet       3 0.101812 0.033937  3.1979
## Sex:Diet        1 0.016718 0.016718  1.5753
## Stage:APOE:Sex  6 0.032211 0.005369  0.5059
## Stage:APOE:Diet 3 0.002704 0.000901  0.0849
## Stage:Sex:Diet  1 0.001843 0.001843  0.1736
## APOE:Sex:Diet   2 0.000171 0.000085  0.0080
## Stage:APOE:Sex:Diet 2 0.025896 0.012948  1.2201

## contrast      estimate      SE df t.ratio p.value
## Female - Male -0.00454 0.0184 292  -0.247  0.8051
##

## Results are averaged over the levels of: APOE

## Linear mixed model fit by REML ['lmerMod']
## Formula: NormSWDist ~ Stage * APOE * Sex * Diet + (1 | Animal)
##   Data: probe_trials
##
## REML criterion at convergence: -227.4
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.05286 -0.49505  0.02678  0.47553  2.57501
##
## Random effects:
##   Groups   Name      Variance Std.Dev.
##   Animal   (Intercept) 0.01112  0.1054
##   Residual              0.01061  0.1030
## Number of obs: 300, groups:  Animal, 150
##
## Fixed effects:
##
##               Estimate Std. Error t value
## (Intercept)      0.453771   0.052119   8.706
## StageProbe_D8    -0.027854   0.051508  -0.541
## APOEE2HN         -0.042318   0.267839  -0.158
```

```

## APOEE33 -0.018873 0.059880 -0.315
## APOEE3HN -0.052987 0.079613 -0.666
## APOEE44 -0.004948 0.069925 -0.071
## APOEE4HN -0.090720 0.116541 -0.778
## APOEHN 0.210931 0.156356 1.349
## SexMale -0.008881 0.069925 -0.127
## DietHFD 0.194423 0.118194 1.645
## StageProbe_D8:APOEE2HN 0.062442 0.264702 0.236
## StageProbe_D8:APOEE33 -0.029197 0.059178 -0.493
## StageProbe_D8:APOEE3HN -0.033956 0.078680 -0.432
## StageProbe_D8:APOEE44 -0.032199 0.069106 -0.466
## StageProbe_D8:APOEE4HN -0.008777 0.115176 -0.076
## StageProbe_D8:APOEHN 0.068323 0.154525 0.442
## StageProbe_D8:SexMale -0.092360 0.069106 -1.337
## APOEE2HN:SexMale 0.188971 0.228429 0.827
## APOEE33:SexMale 0.104944 0.085504 1.227
## APOEE3HN:SexMale 0.022586 0.100317 0.225
## APOEE44:SexMale 0.050886 0.093457 0.544
## APOEE4HN:SexMale 0.065908 0.163157 0.404
## APOEHN:SexMale -0.464123 0.219889 -2.111
## StageProbe_D8:DietHFD -0.008157 0.116810 -0.070
## APOEE2HN:DietHFD -0.237098 0.225178 -1.053
## APOEE33:DietHFD -0.011052 0.133953 -0.083
## APOEE3HN:DietHFD -0.205805 0.143861 -1.431
## SexMale:DietHFD -0.156062 0.179106 -0.871
## StageProbe_D8:APOEE2HN:SexMale -0.103402 0.225754 -0.458
## StageProbe_D8:APOEE33:SexMale 0.003622 0.084502 0.043
## StageProbe_D8:APOEE3HN:SexMale 0.088046 0.099142 0.888
## StageProbe_D8:APOEE44:SexMale 0.112700 0.092362 1.220
## StageProbe_D8:APOEE4HN:SexMale 0.019104 0.161246 0.118
## StageProbe_D8:APOEHN:SexMale 0.094071 0.217313 0.433
## StageProbe_D8:APOEE2HN:DietHFD -0.058504 0.222541 -0.263
## StageProbe_D8:APOEE33:DietHFD 0.024379 0.132384 0.184
## StageProbe_D8:APOEE3HN:DietHFD 0.098084 0.142176 0.690
## StageProbe_D8:SexMale:DietHFD 0.176732 0.177008 0.998
## APOEE33:SexMale:DietHFD 0.034434 0.203043 0.170
## APOEE3HN:SexMale:DietHFD 0.140895 0.209712 0.672
## StageProbe_D8:APOEE33:SexMale:DietHFD -0.094129 0.200665 -0.469
## StageProbe_D8:APOEE3HN:SexMale:DietHFD -0.276319 0.207256 -1.333
## fit warnings:
## fixed-effect model matrix is rank deficient so dropping 14 columns / coefficients

## 2.5 % 97.5 %
## .sig01 0.08096136 0.11584630
## .sigma 0.08566382 0.10745475
## (Intercept) 0.35865513 0.54888630
## StageProbe_D8 -0.12207747 0.06636994
## APOEE2HN -0.53111933 0.44648242
## APOEE33 -0.12815248 0.09040649
## APOEE3HN -0.19827796 0.09230495
## APOEE44 -0.13255890 0.12266300
## APOEE4HN -0.30340516 0.12196467
## APOEHN -0.07441527 0.49627825
## SexMale -0.13649196 0.11872994

```

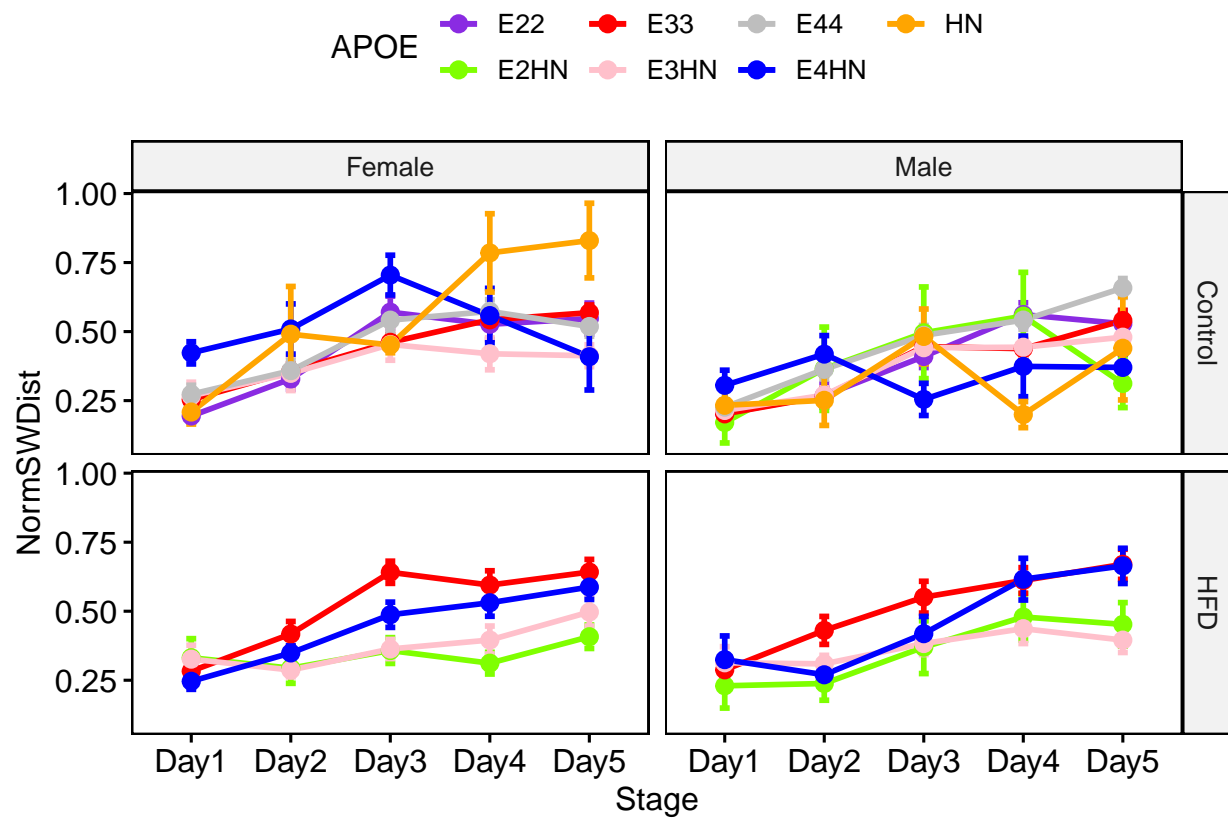
## DietHFD	-0.02127937	0.41012438
## StageProbe_D8:APOEE2HN	-0.42177555	0.54665941
## StageProbe_D8:APOEE33	-0.13745194	0.07905765
## StageProbe_D8:APOEE3HN	-0.17788476	0.10997341
## StageProbe_D8:APOEE44	-0.15861355	0.09421518
## StageProbe_D8:APOEE4HN	-0.21946761	0.20191360
## StageProbe_D8:APOEHN	-0.21434817	0.35099406
## StageProbe_D8:SexMale	-0.21877460	0.03405413
## APOEE2HN:SexMale	-0.22790758	0.60585036
## APOEE33:SexMale	-0.05109861	0.26098654
## APOEE3HN:SexMale	-0.16049002	0.20566234
## APOEE44:SexMale	-0.11967095	0.22144196
## APOEE4HN:SexMale	-0.23185125	0.36366651
## APOEHN:SexMale	-0.86541658	-0.06283038
## StageProbe_D8:DietHFD	-0.22183581	0.20552275
## APOEE2HN:DietHFD	-0.64804456	0.17384804
## APOEE33:DietHFD	-0.25551458	0.23340967
## APOEE3HN:DietHFD	-0.46834919	0.05673873
## SexMale:DietHFD	-0.48292695	0.17080224
## StageProbe_D8:APOEE2HN:SexMale	-0.51637207	0.30956787
## StageProbe_D8:APOEE33:SexMale	-0.15095715	0.15820164
## StageProbe_D8:APOEE3HN:SexMale	-0.09331343	0.26940559
## StageProbe_D8:APOEE44:SexMale	-0.05625739	0.28165696
## StageProbe_D8:APOEE4HN:SexMale	-0.27586311	0.31407060
## StageProbe_D8:APOEHN:SexMale	-0.30345915	0.49160135
## StageProbe_D8:APOEE2HN:DietHFD	-0.46559644	0.34858943
## StageProbe_D8:APOEE33:DietHFD	-0.21779039	0.26654931
## StageProbe_D8:APOEE3HN:DietHFD	-0.16199783	0.35816644
## StageProbe_D8:SexMale:DietHFD	-0.14706721	0.50053208
## APOEE33:SexMale:DietHFD	-0.33611551	0.40498429
## APOEE3HN:SexMale:DietHFD	-0.24182538	0.52361522
## StageProbe_D8:APOEE33:SexMale:DietHFD	-0.46120411	0.27294653
## StageProbe_D8:APOEE3HN:SexMale:DietHFD	-0.65545107	0.10281214

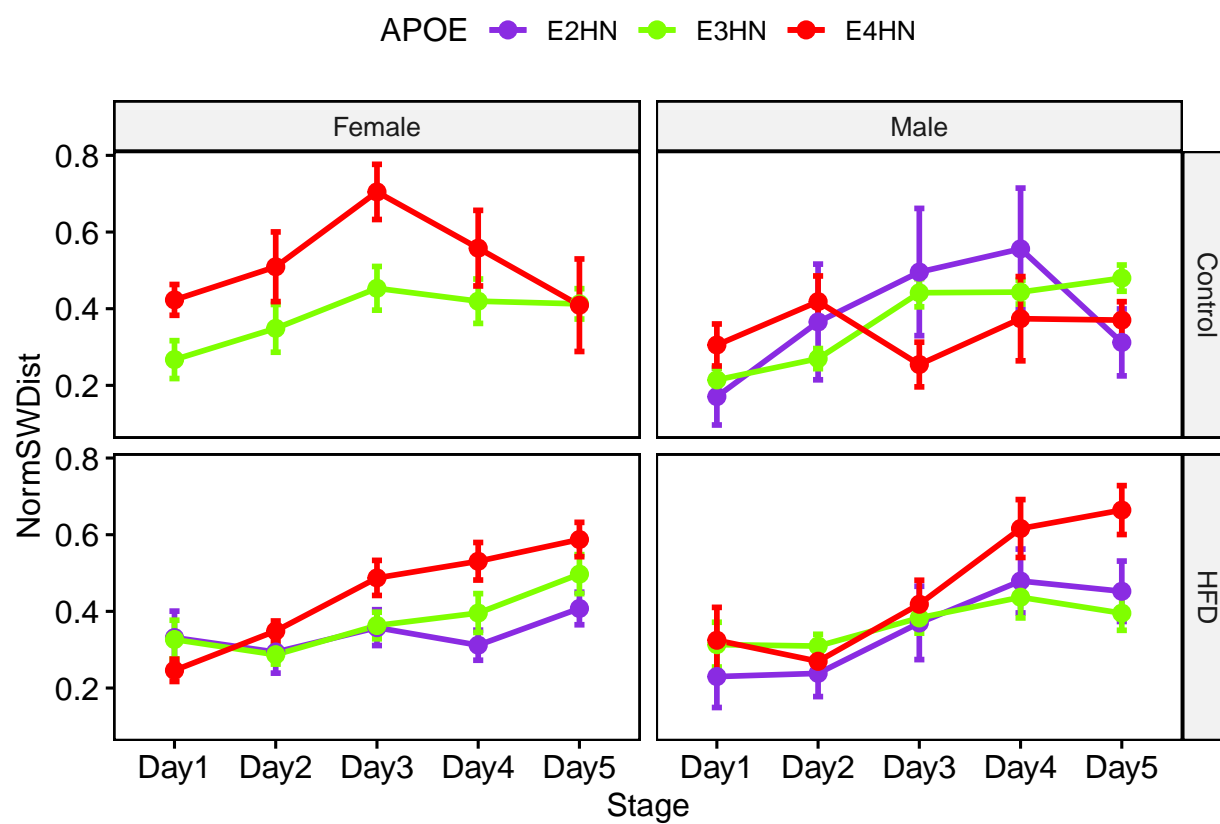
```
## contrast      estimate SE df z.ratio p.value
## Female - Male nonEst NA NA      NA      NA
##
## Results are averaged over the levels of: Stage, APOE, Diet
## Degrees-of-freedom method: kenward-roger
```

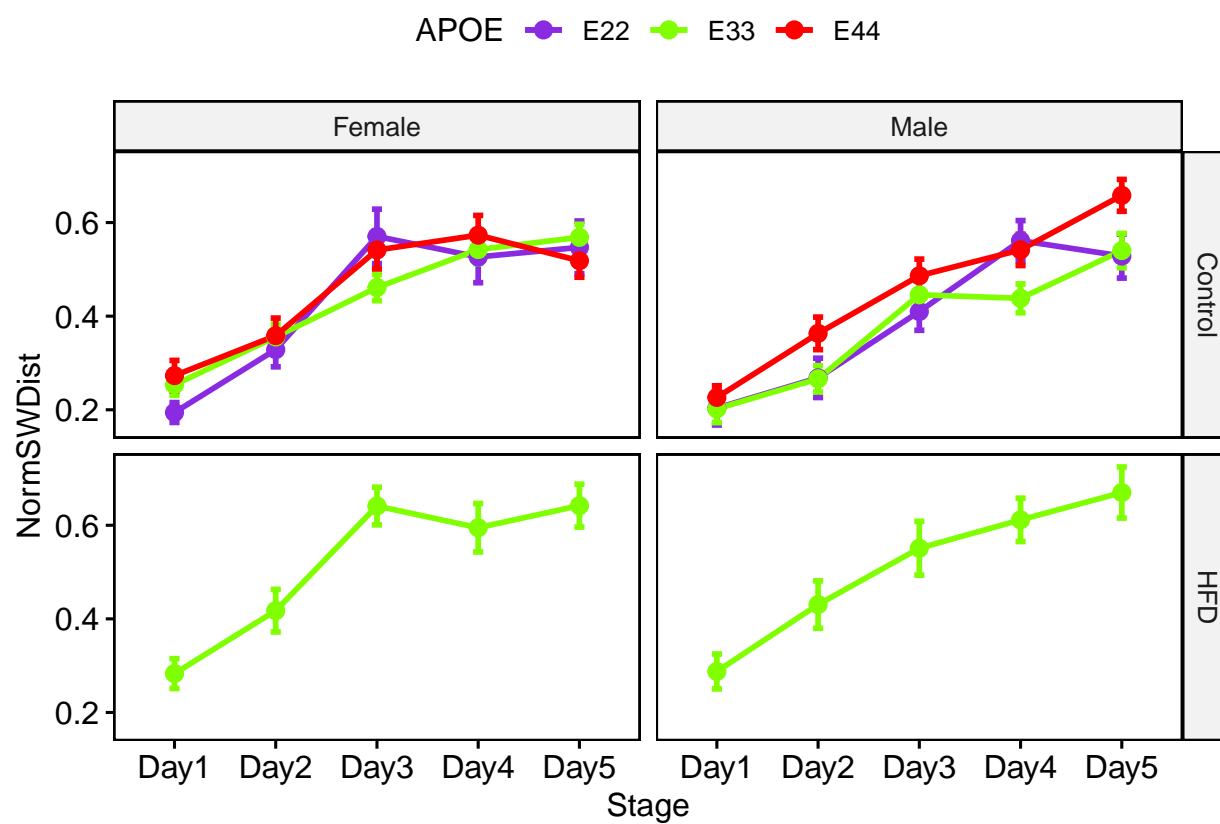
## contrast	estimate	SE	df	t.ratio	p.value
## E22 - E2HN	nonEst	NA	NA	NA	NA
## E22 - E33	nonEst	NA	NA	NA	NA
## E22 - E3HN	nonEst	NA	NA	NA	NA
## E22 - E44	nonEst	NA	NA	NA	NA
## E22 - E4HN	nonEst	NA	NA	NA	NA
## E22 - HN	nonEst	NA	NA	NA	NA
## E2HN - E33	nonEst	NA	NA	NA	NA
## E2HN - E3HN	nonEst	NA	NA	NA	NA
## E2HN - E44	nonEst	NA	NA	NA	NA
## E2HN - E4HN	nonEst	NA	NA	NA	NA
## E2HN - HN	nonEst	NA	NA	NA	NA
## E33 - E3HN	0.1317	0.0315	129	4.176	0.0010
## E33 - E44	nonEst	NA	NA	NA	NA

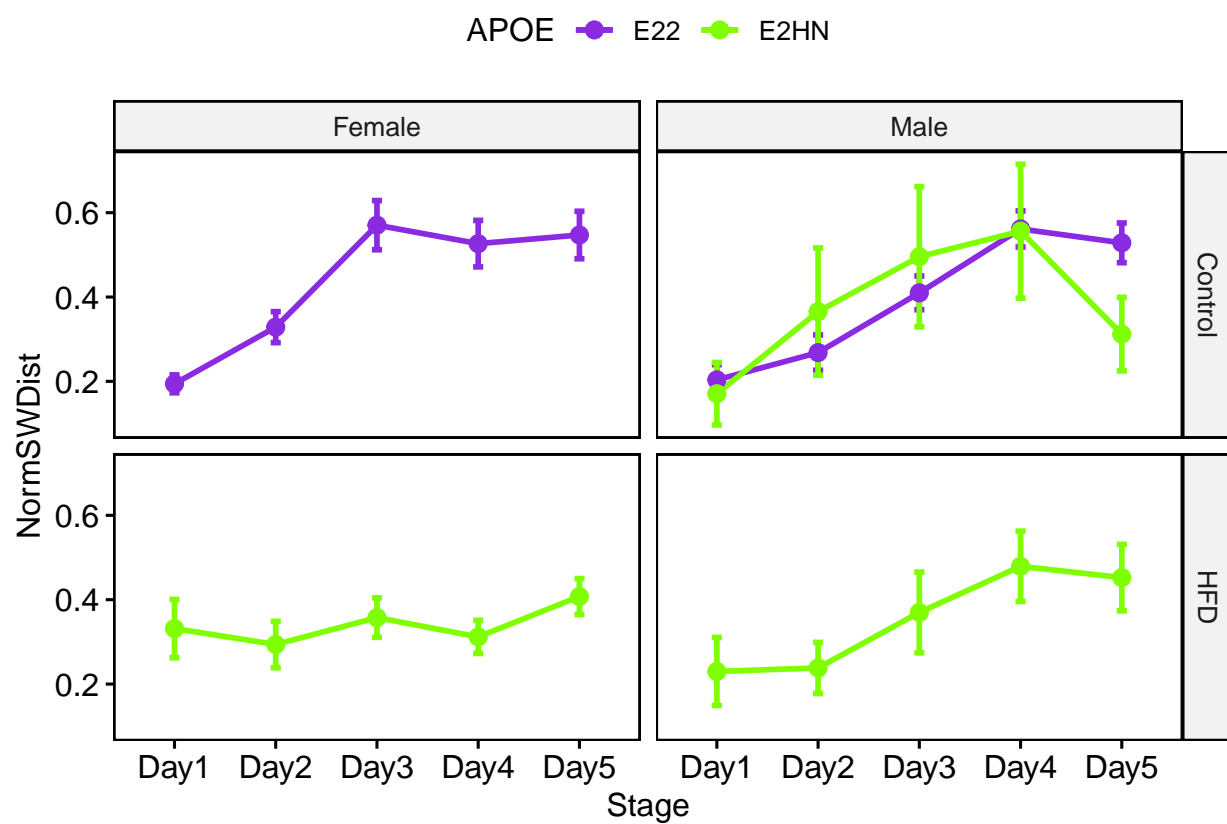
```
## E33 - E4HN      0.0747 0.0441 129    1.693 0.6222
## E33 - HN        nonEst      NA  NA      NA      NA
## E3HN - E44      nonEst      NA  NA      NA      NA
## E3HN - E4HN     -0.0570 0.0456 129    -1.250 0.8729
## E3HN - HN       nonEst      NA  NA      NA      NA
## E44 - E4HN      nonEst      NA  NA      NA      NA
## E44 - HN        nonEst      NA  NA      NA      NA
## E4HN - HN       nonEst      NA  NA      NA      NA
##
## Results are averaged over the levels of: Stage, Sex, Diet
## Degrees-of-freedom method: kenward-roger
## P value adjustment: tukey method for comparing a family of 7 estimates
```

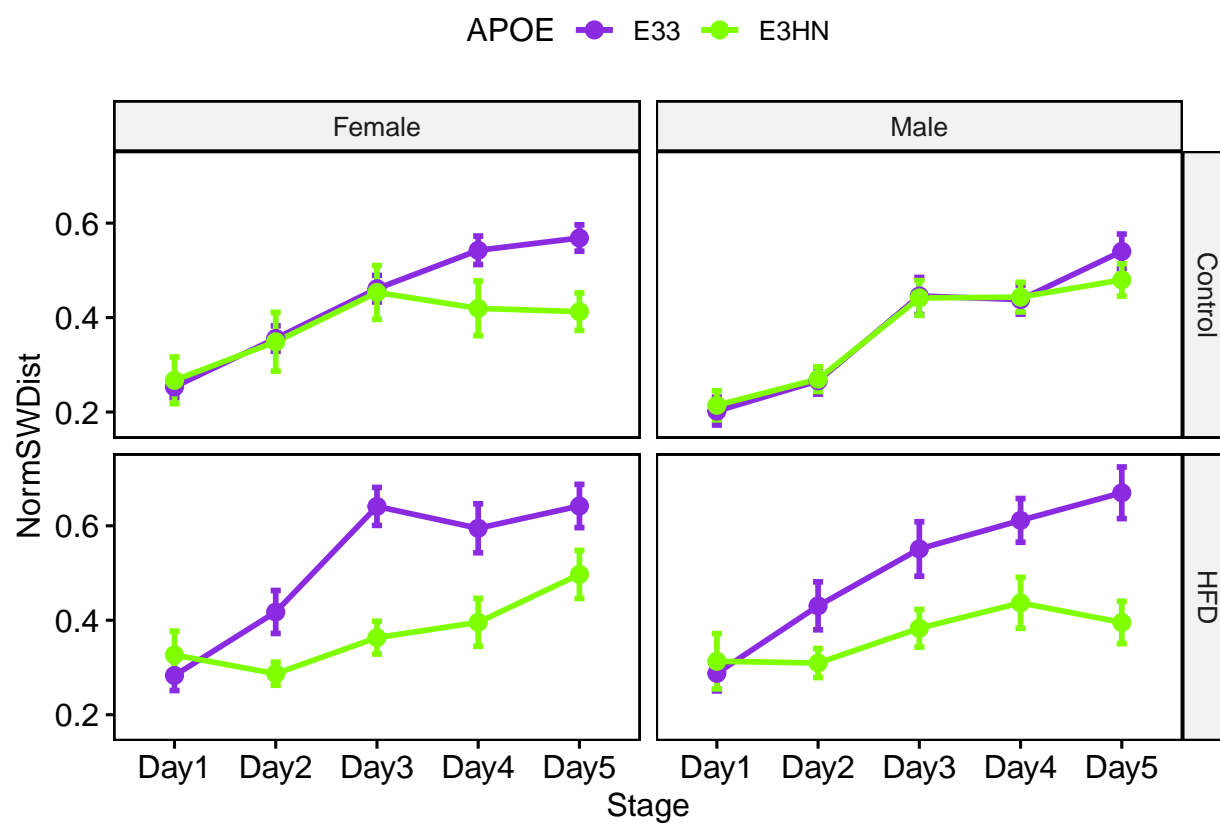
```
lm <- lm(Duration ~ SexAPOEDiet, reg_trials) summary(lm) anova(lm)
```

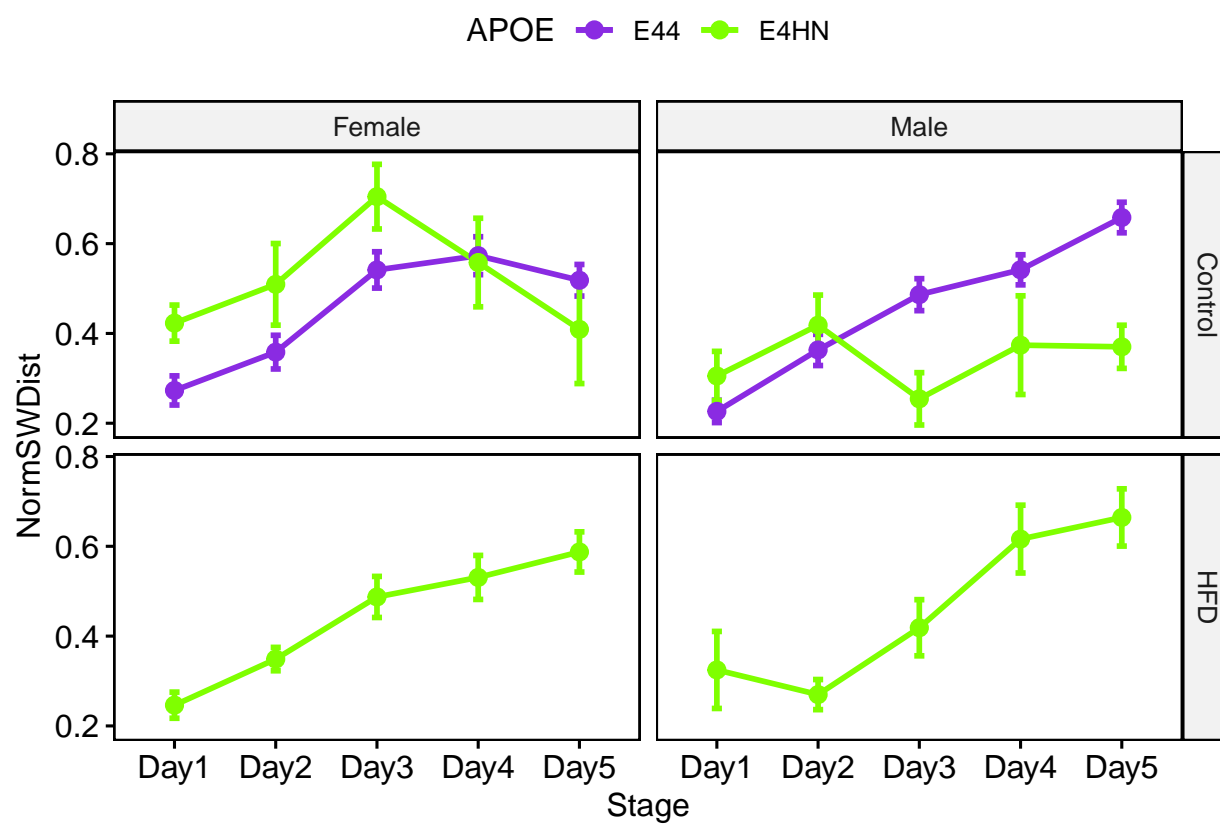


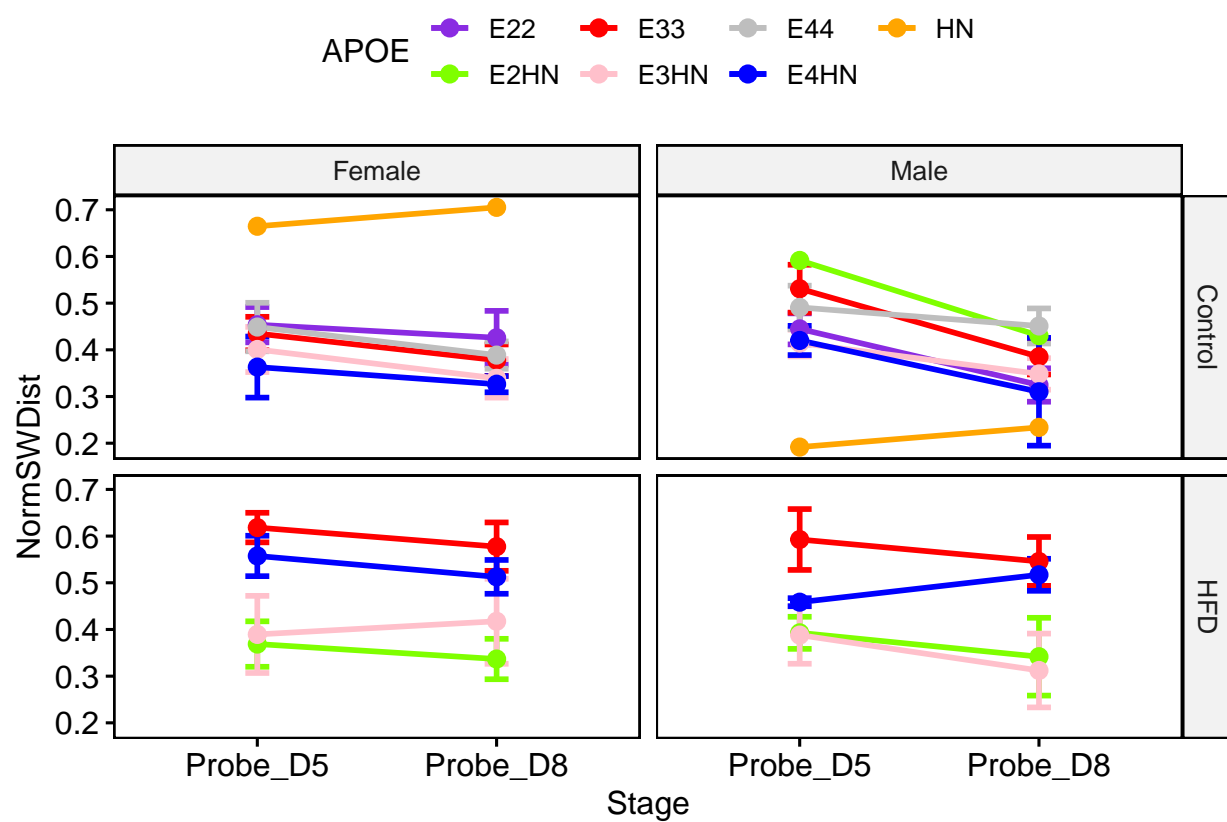


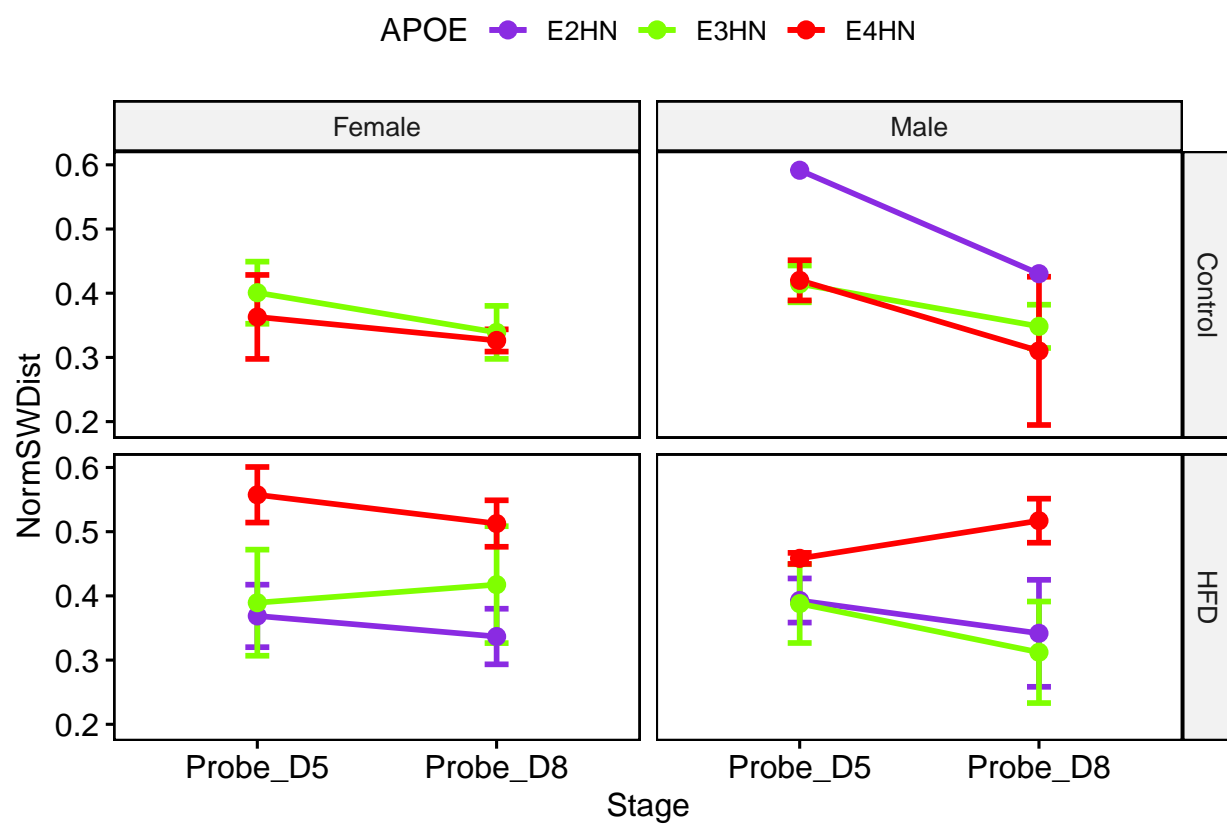


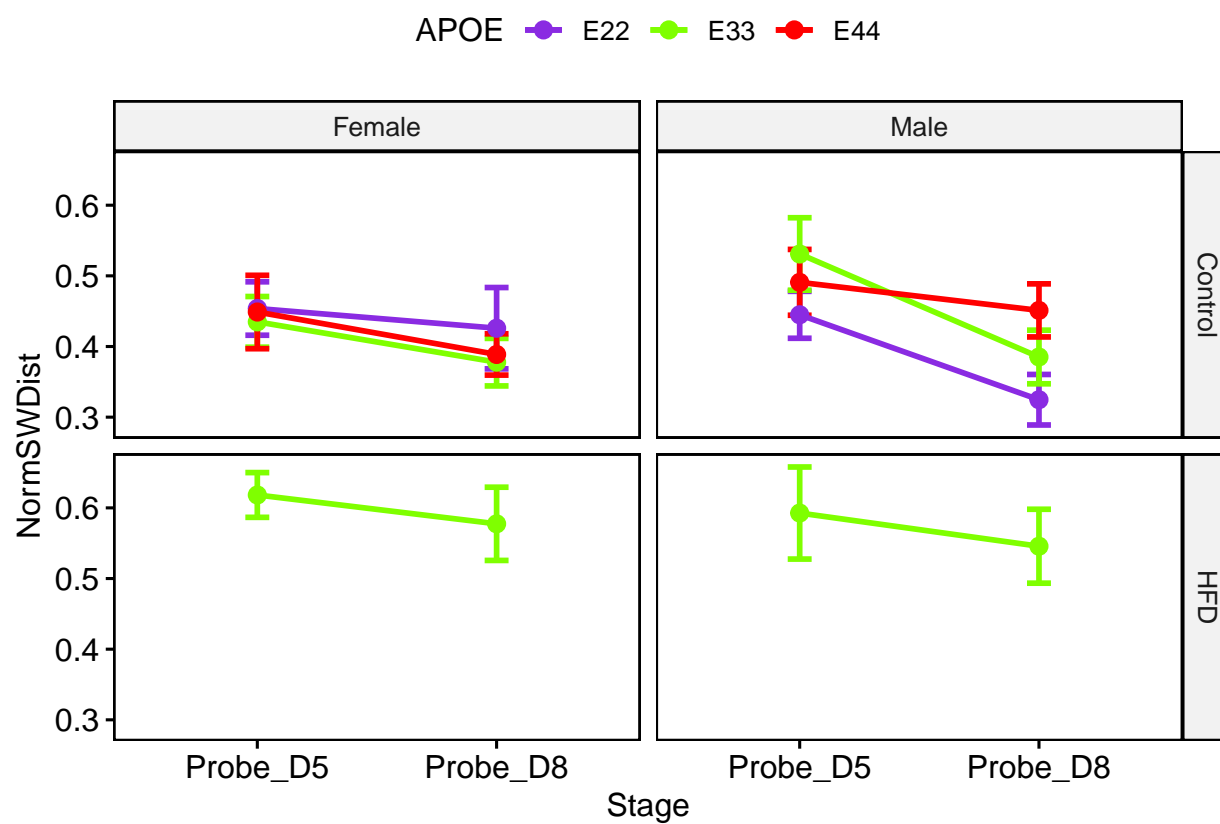


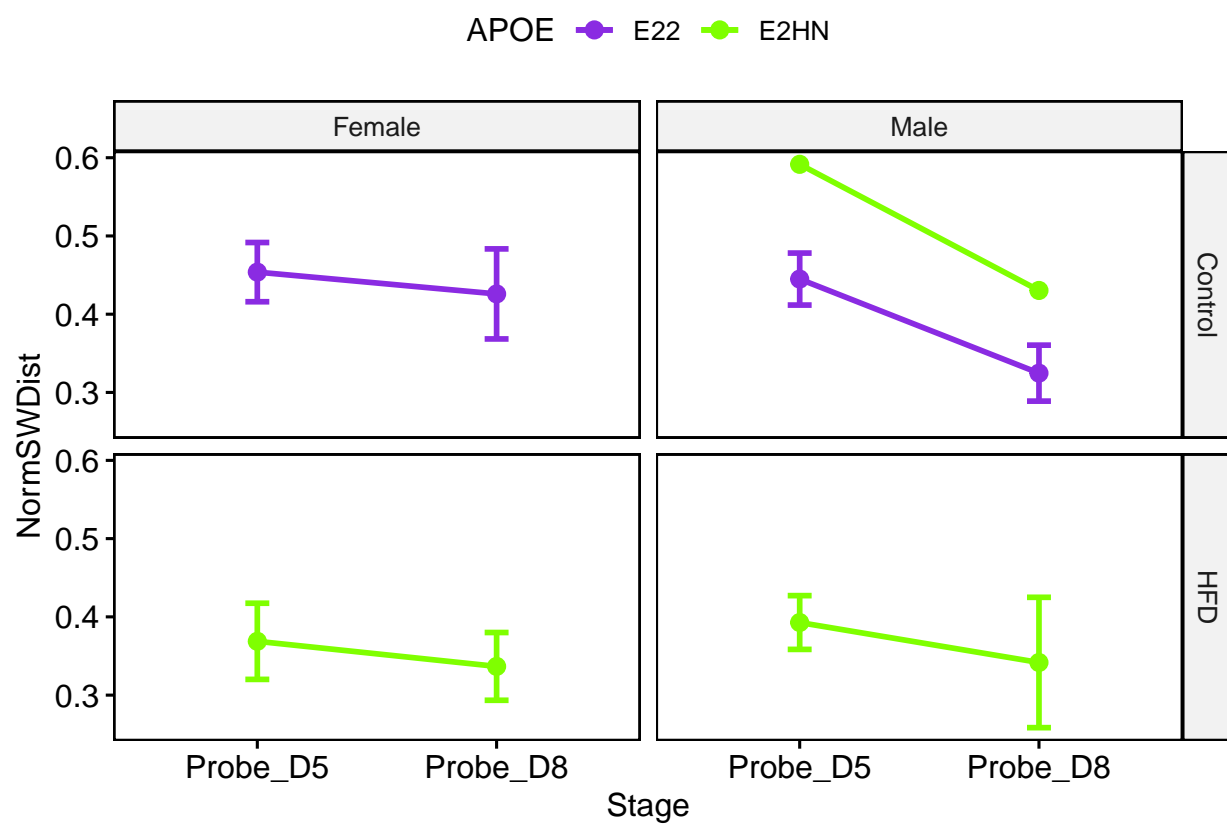


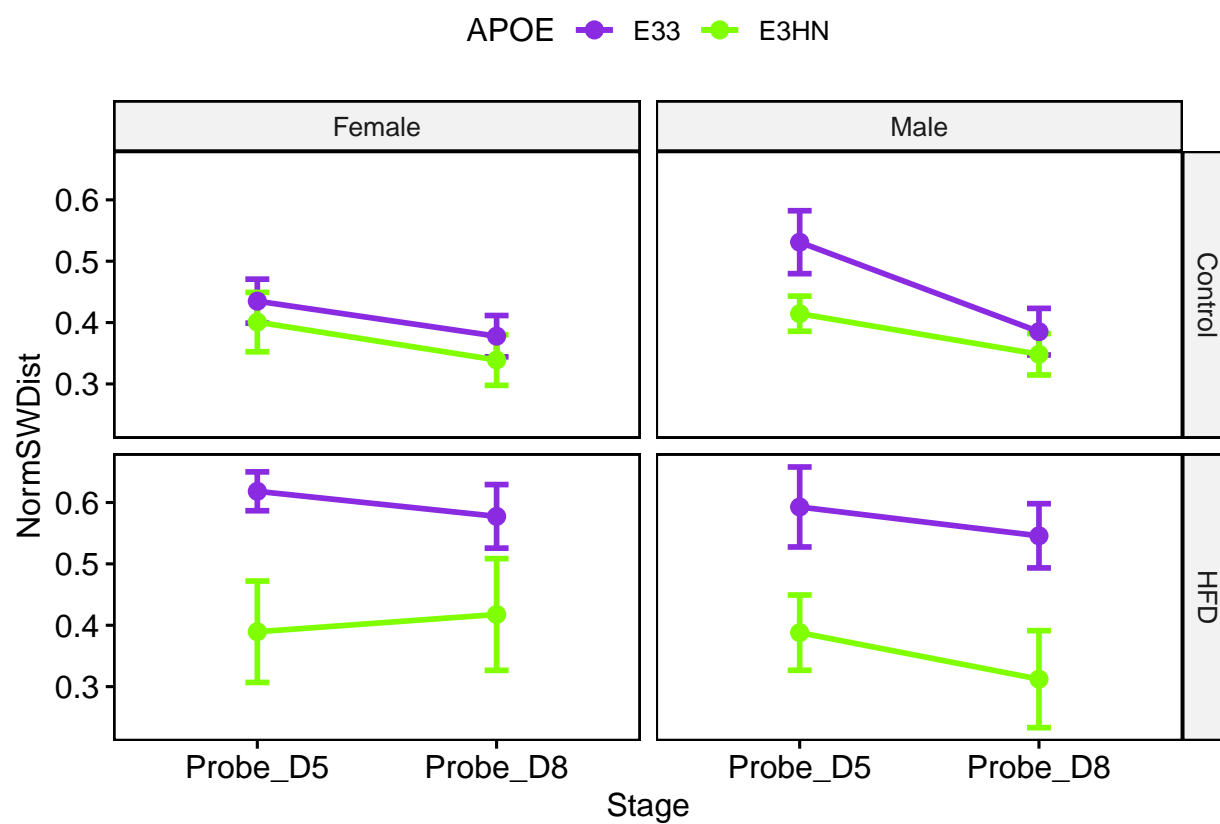


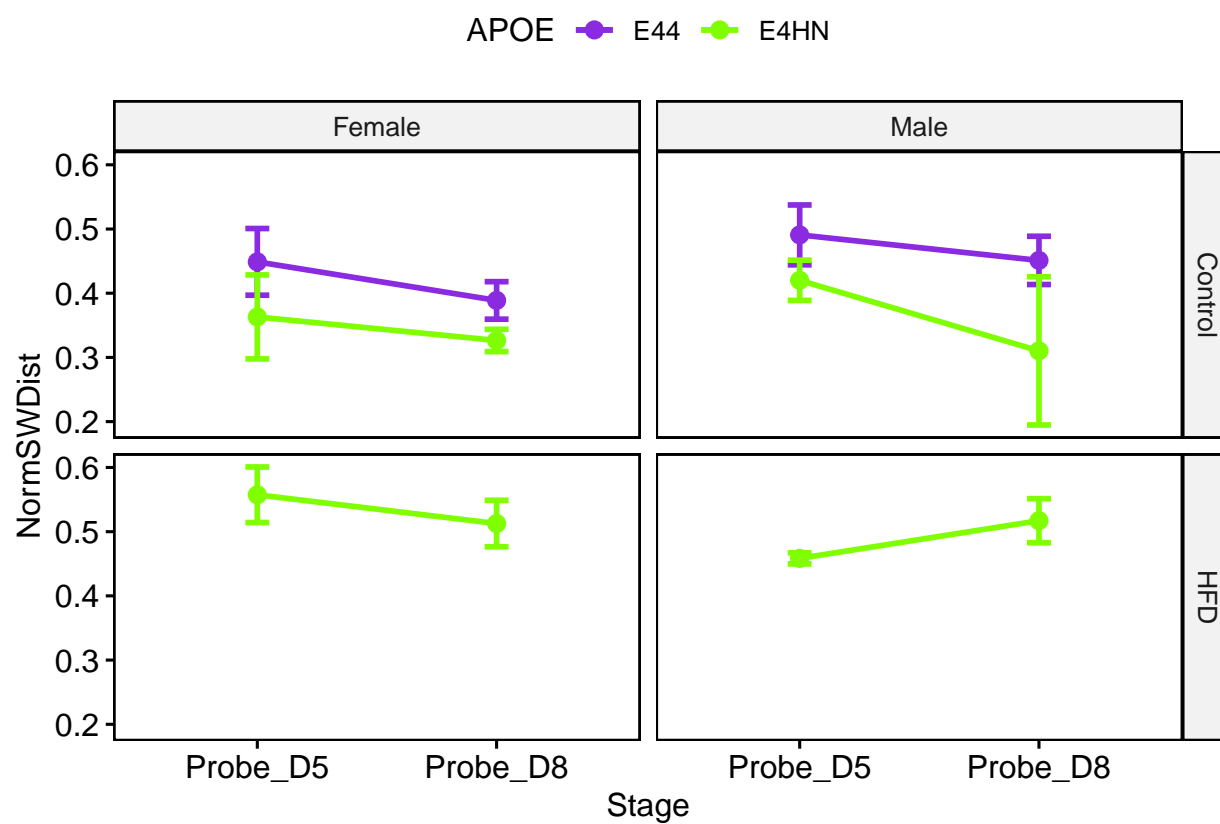




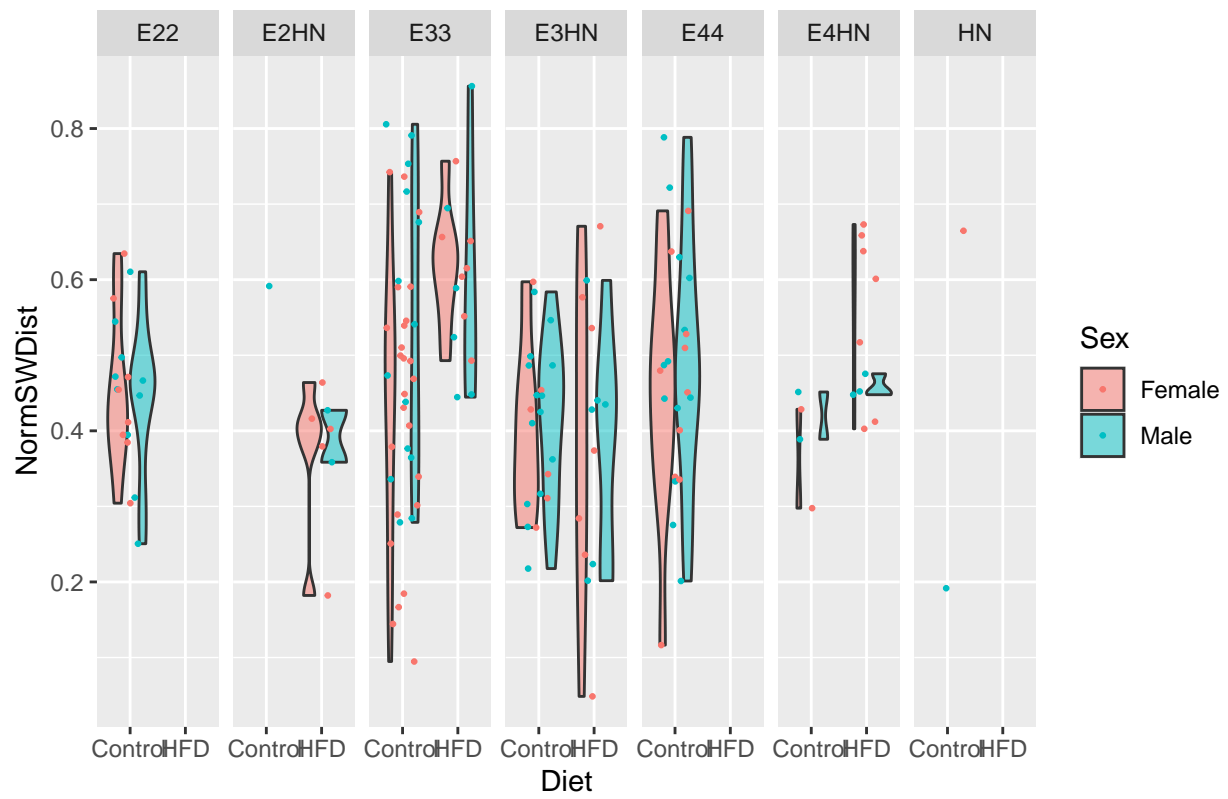




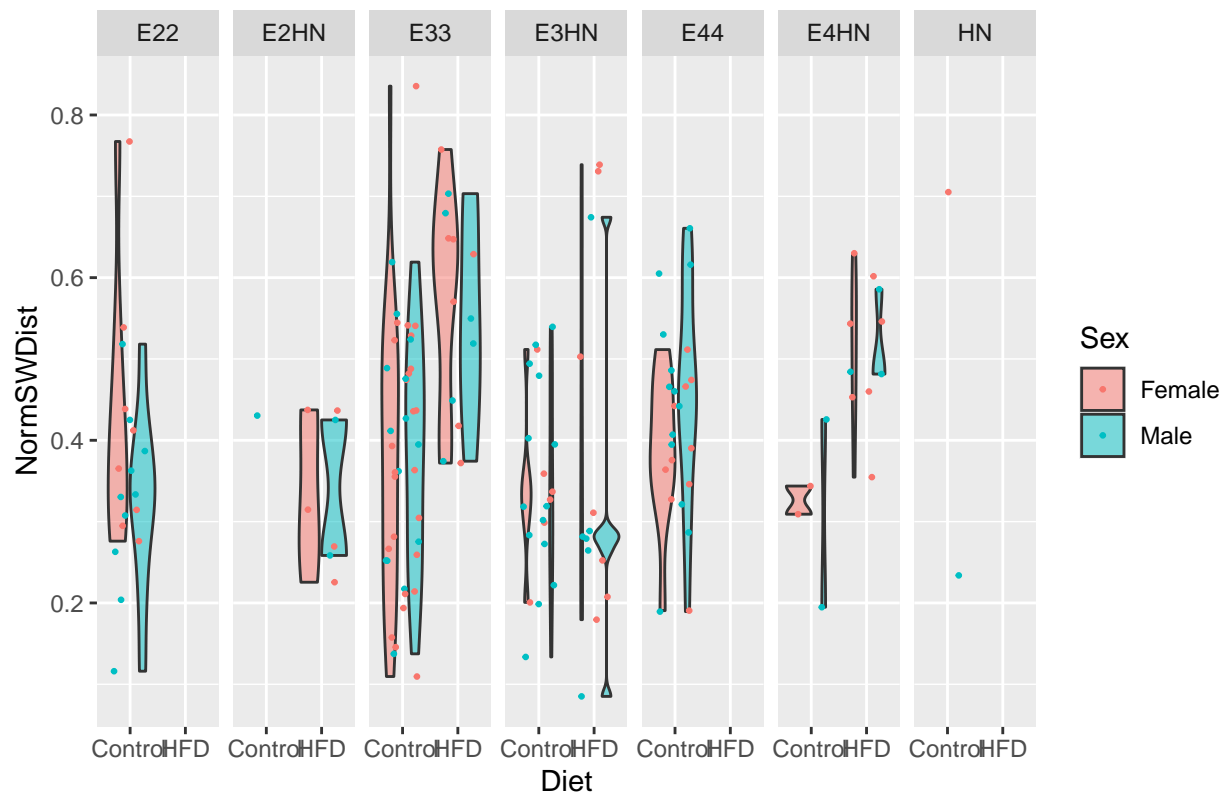


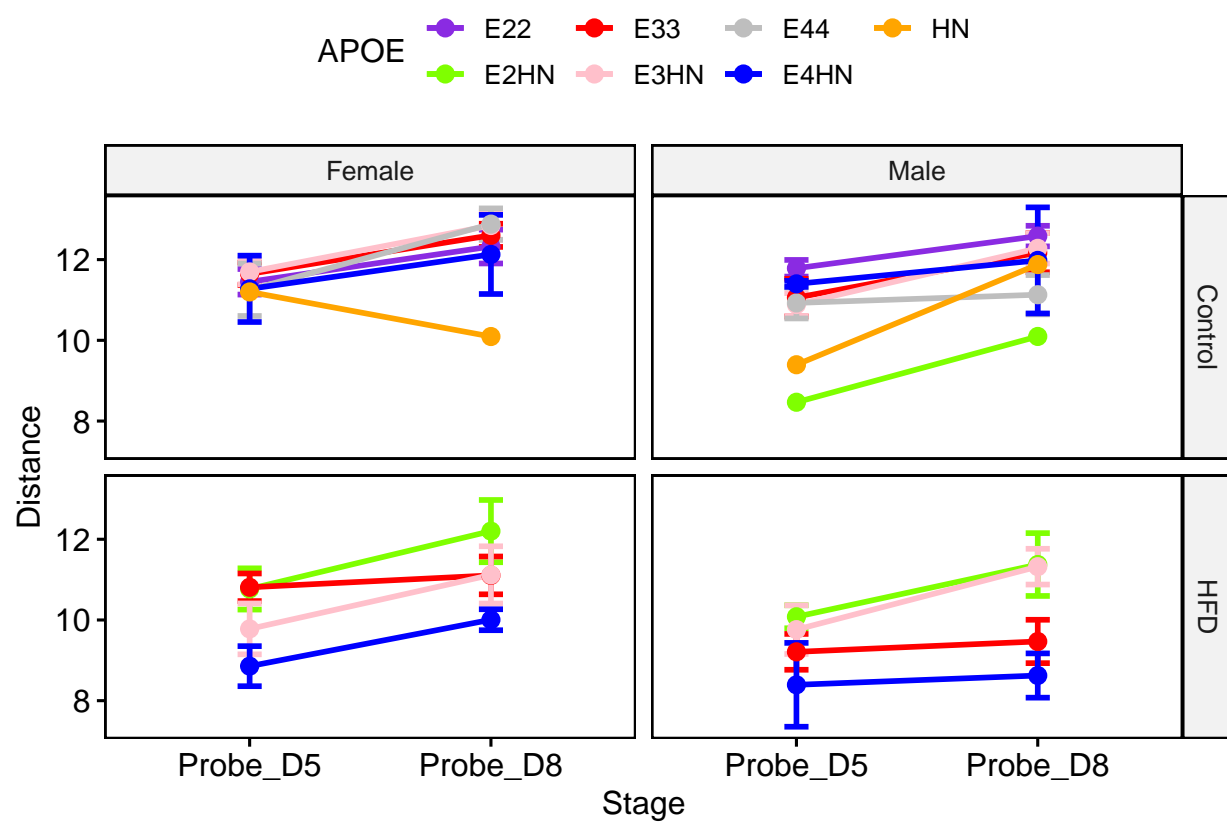


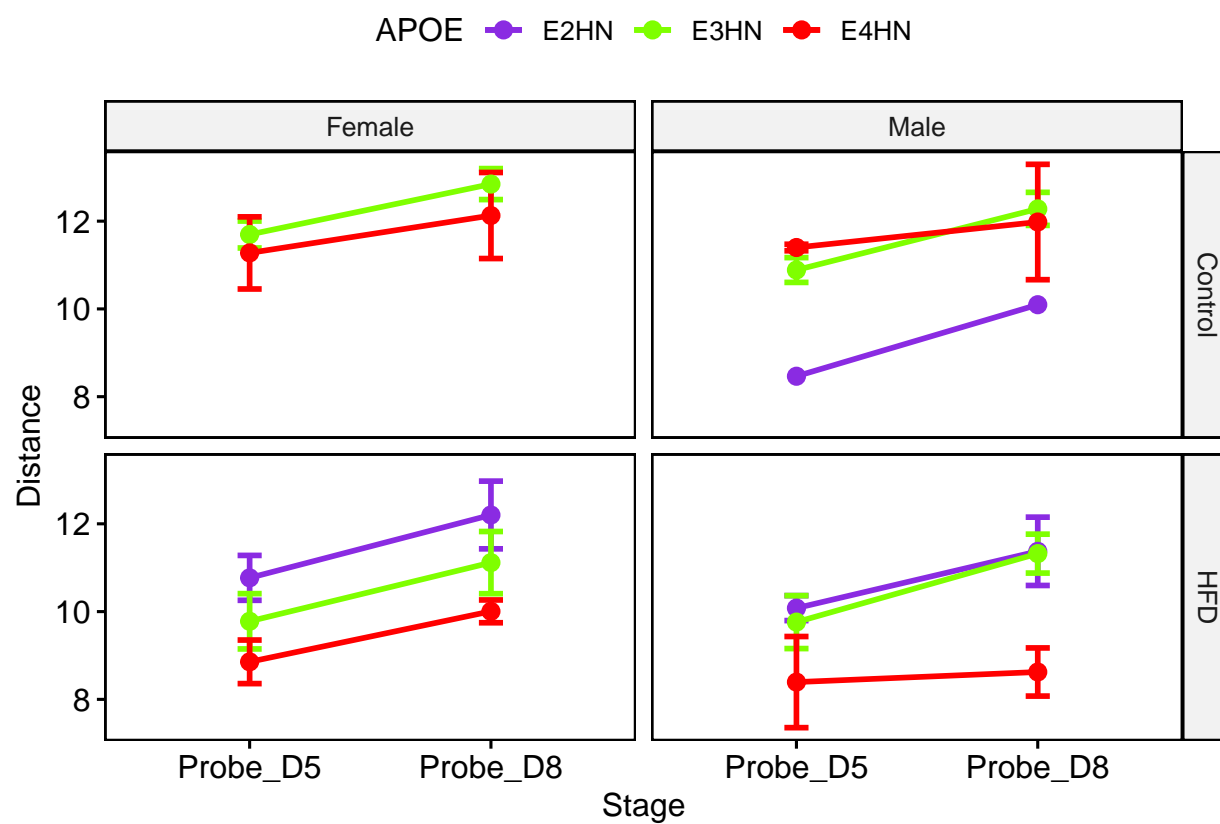
Probe Trial: Day 5

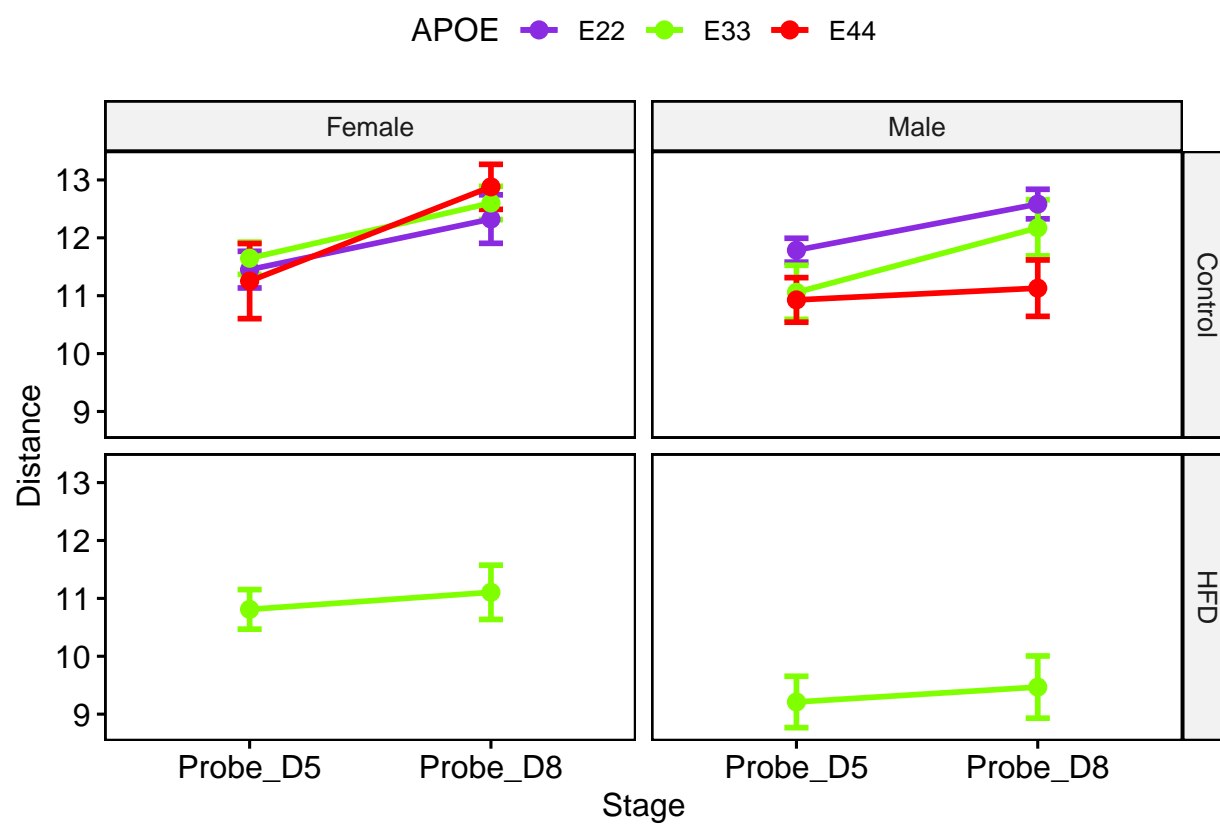


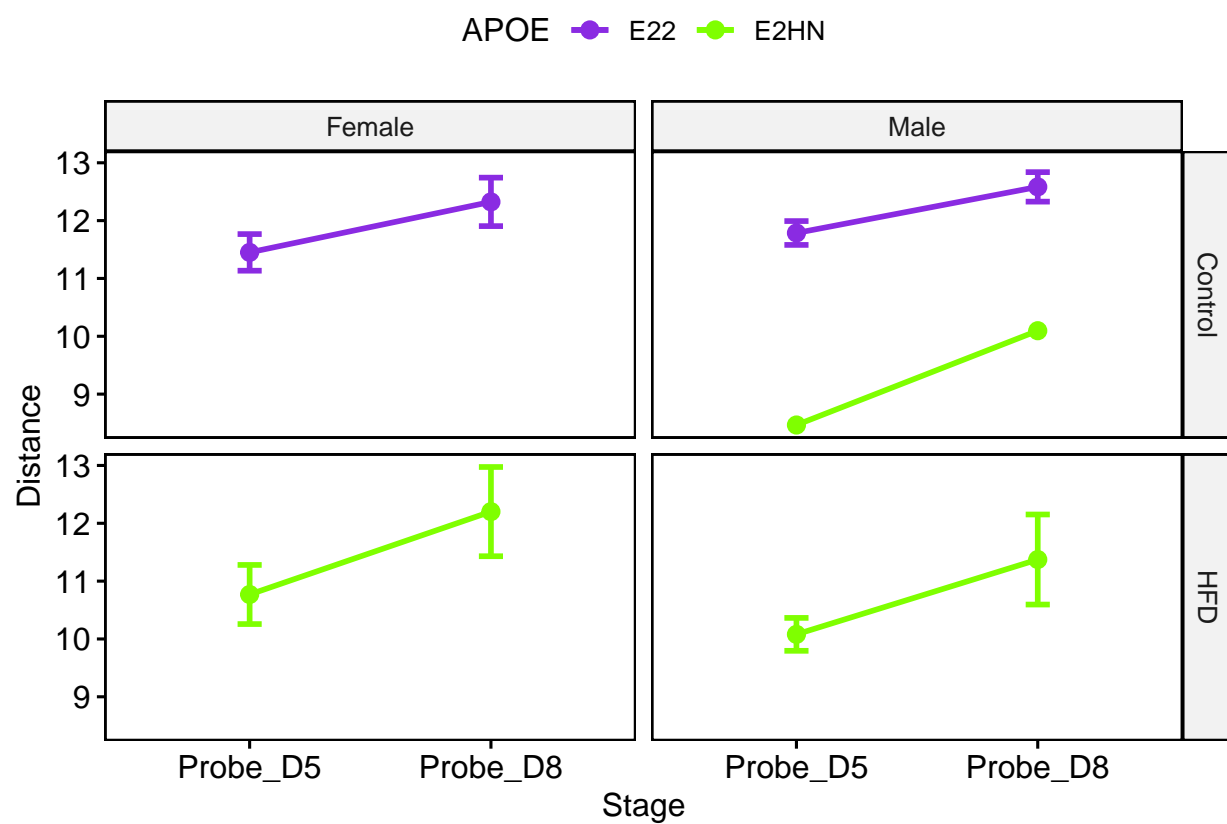
Probe Trial: Day 8

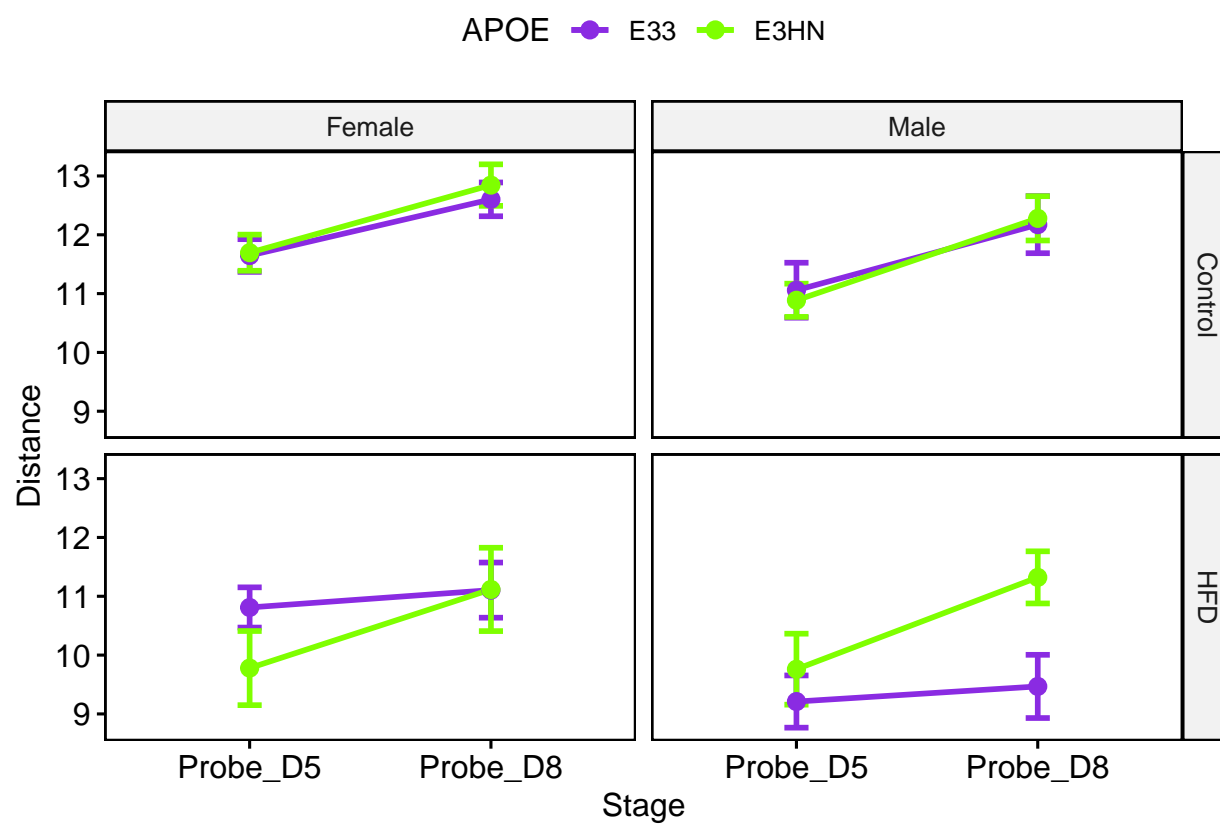


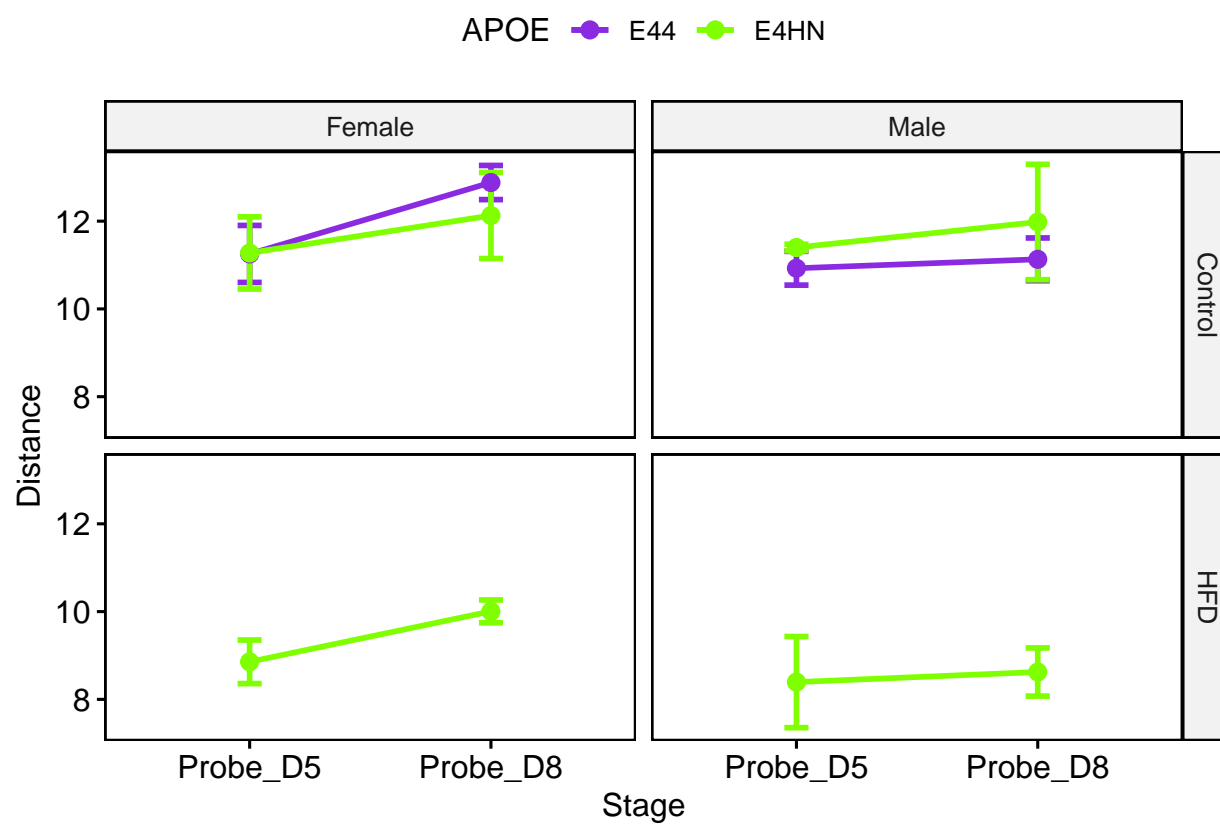


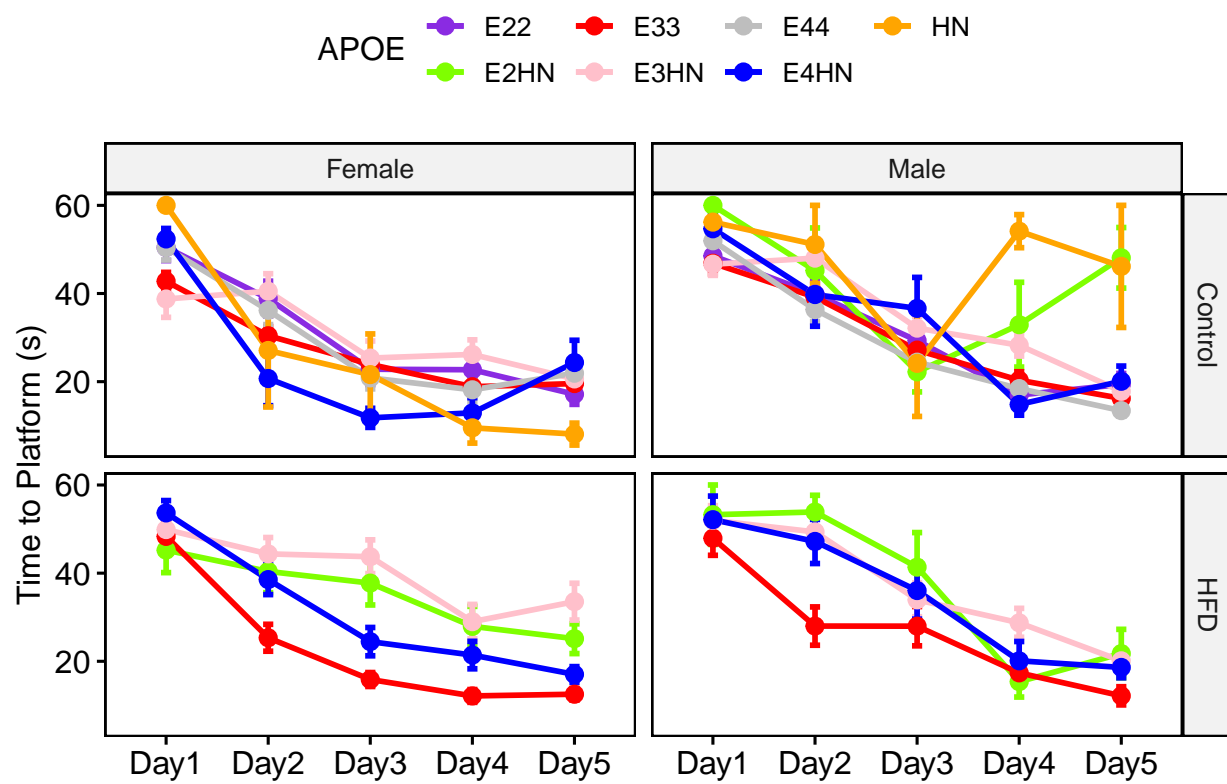


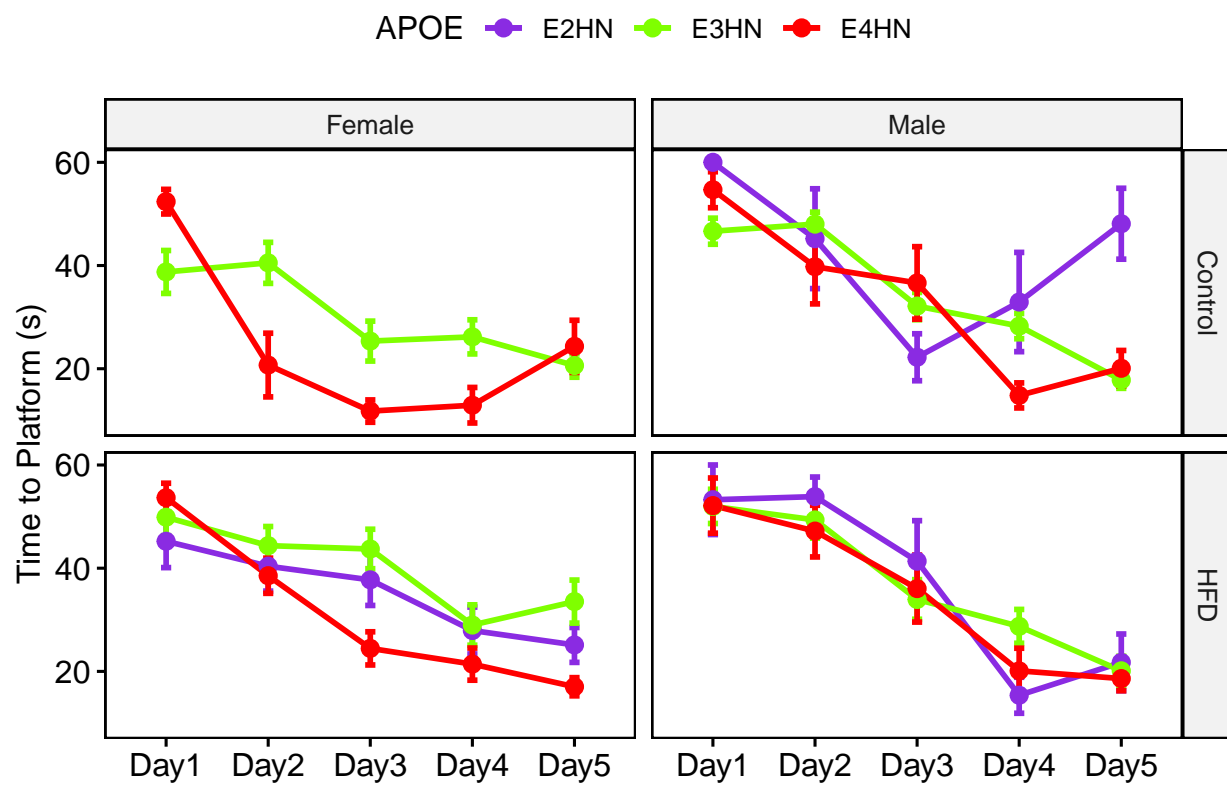


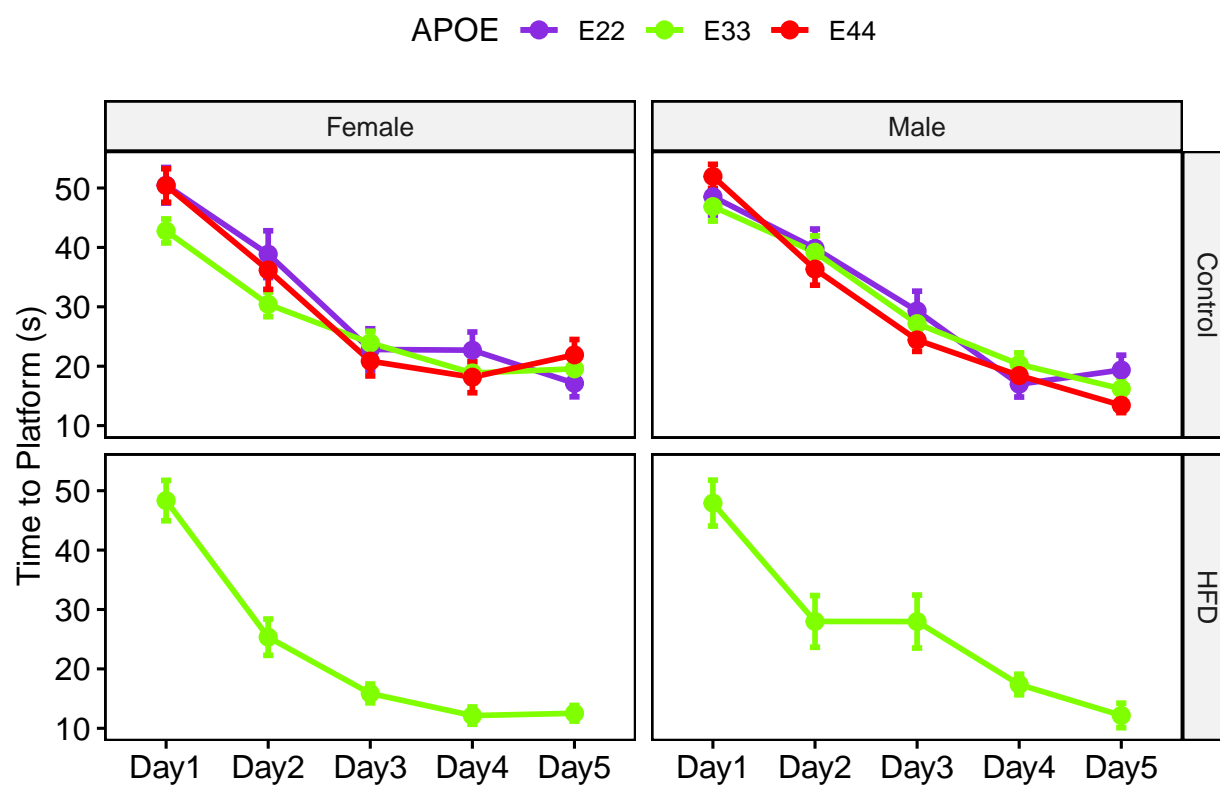


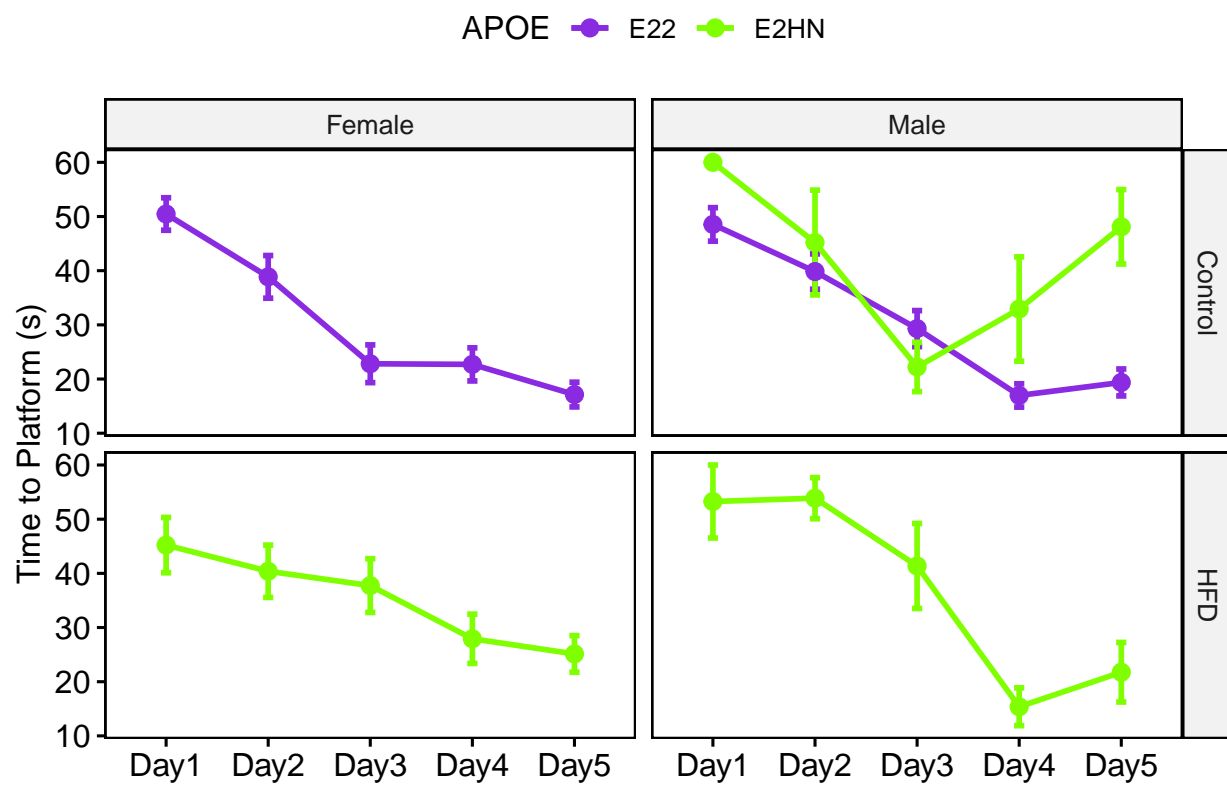


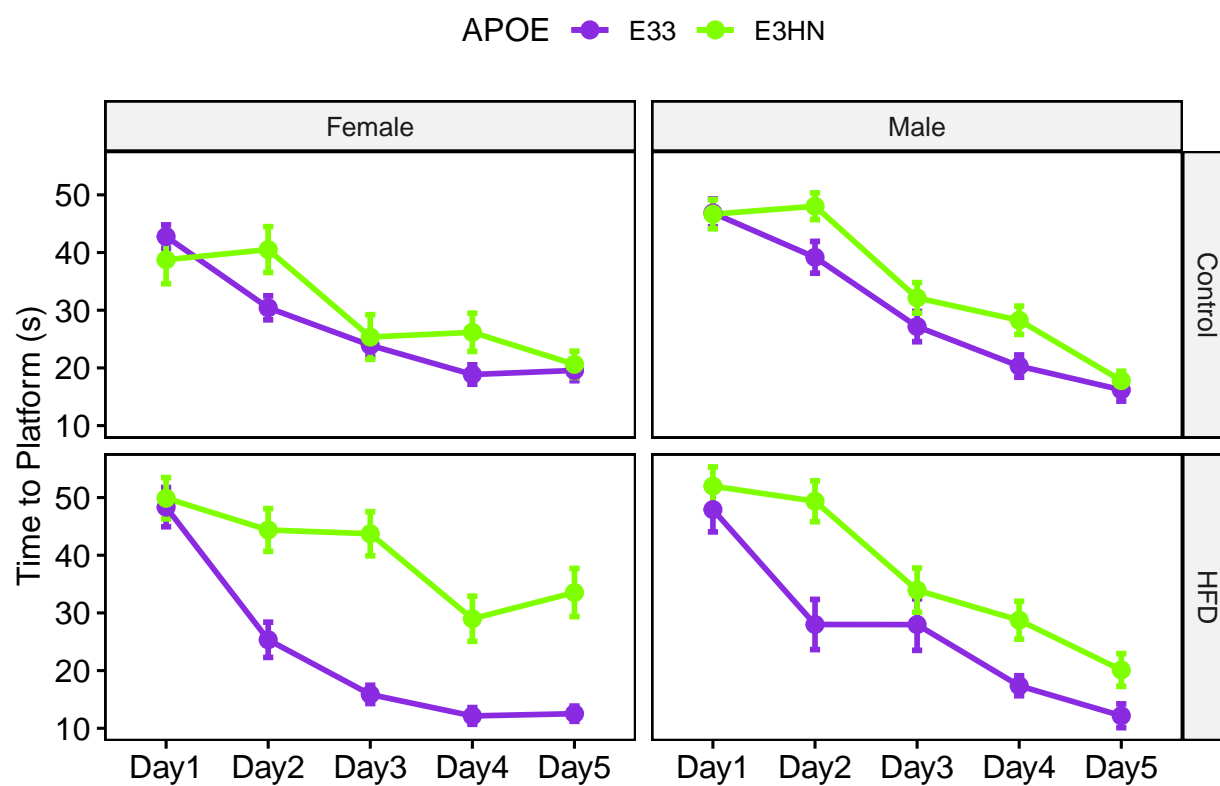


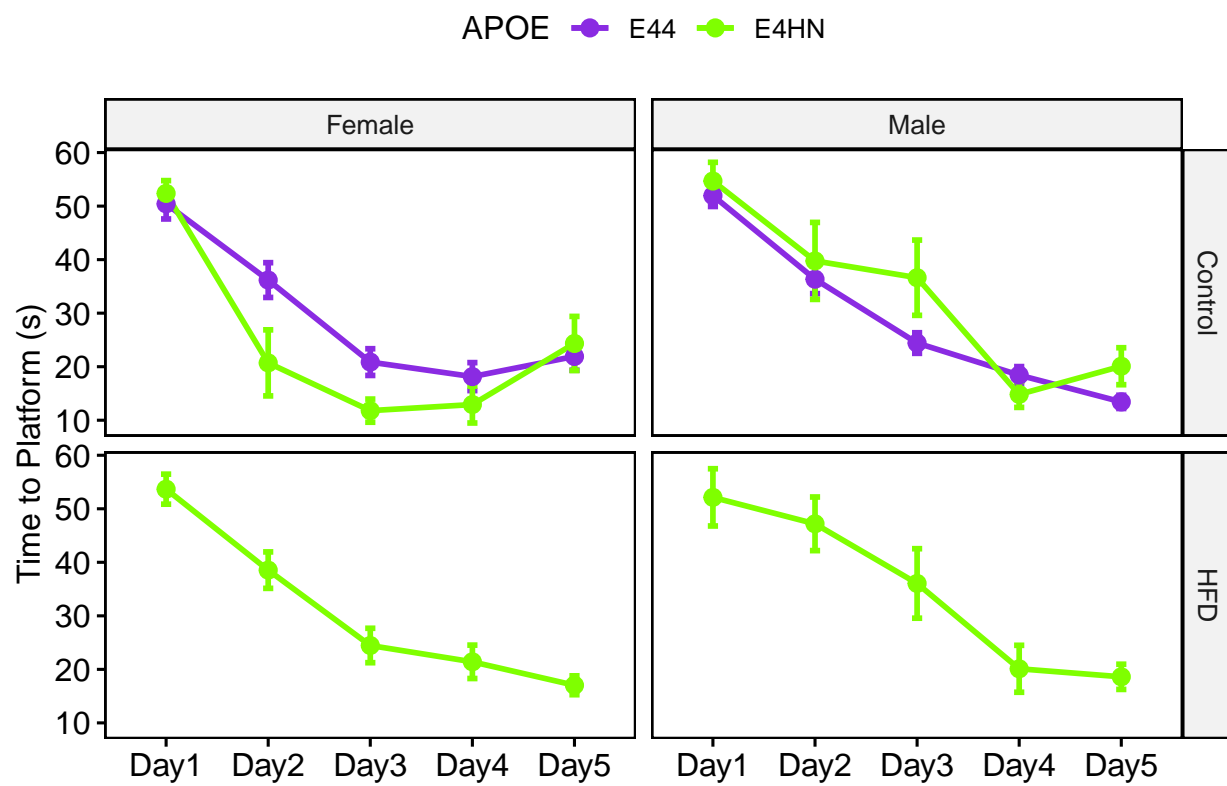


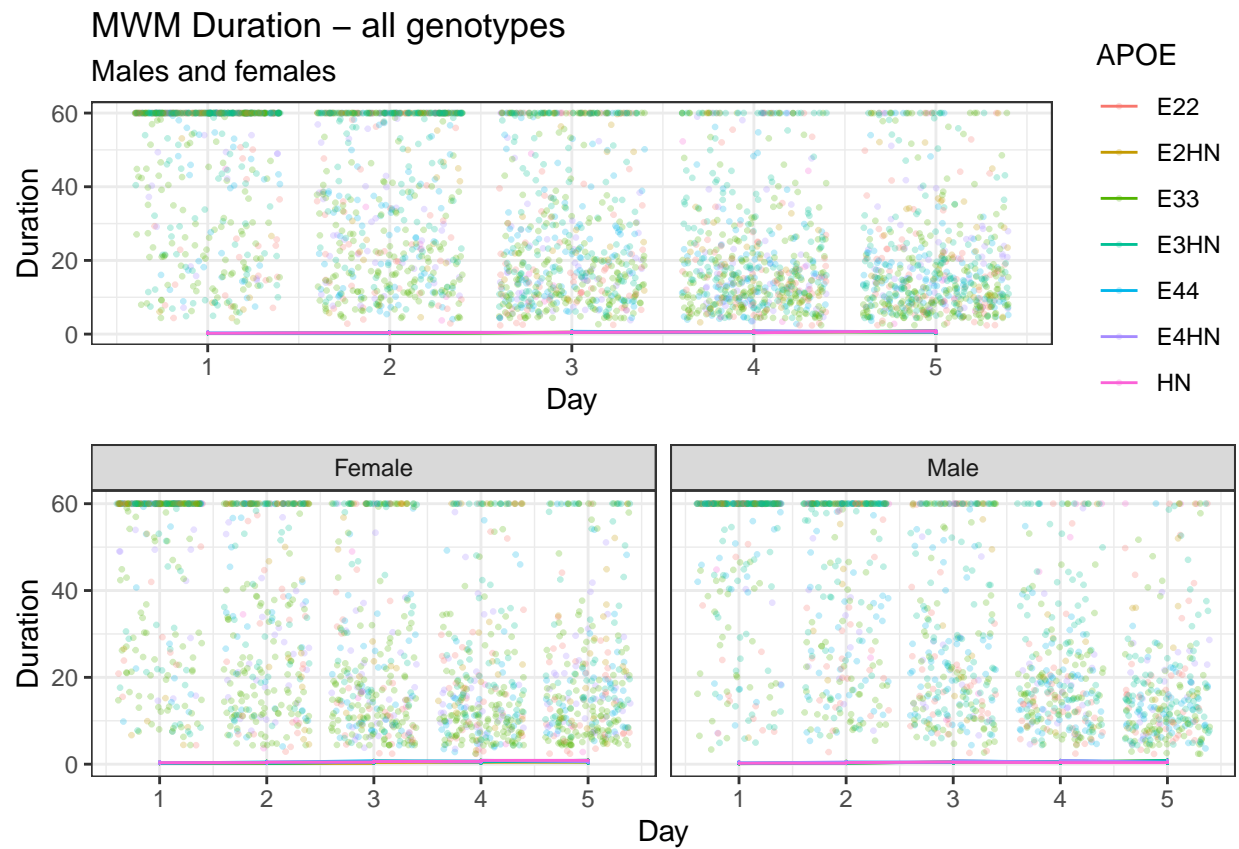






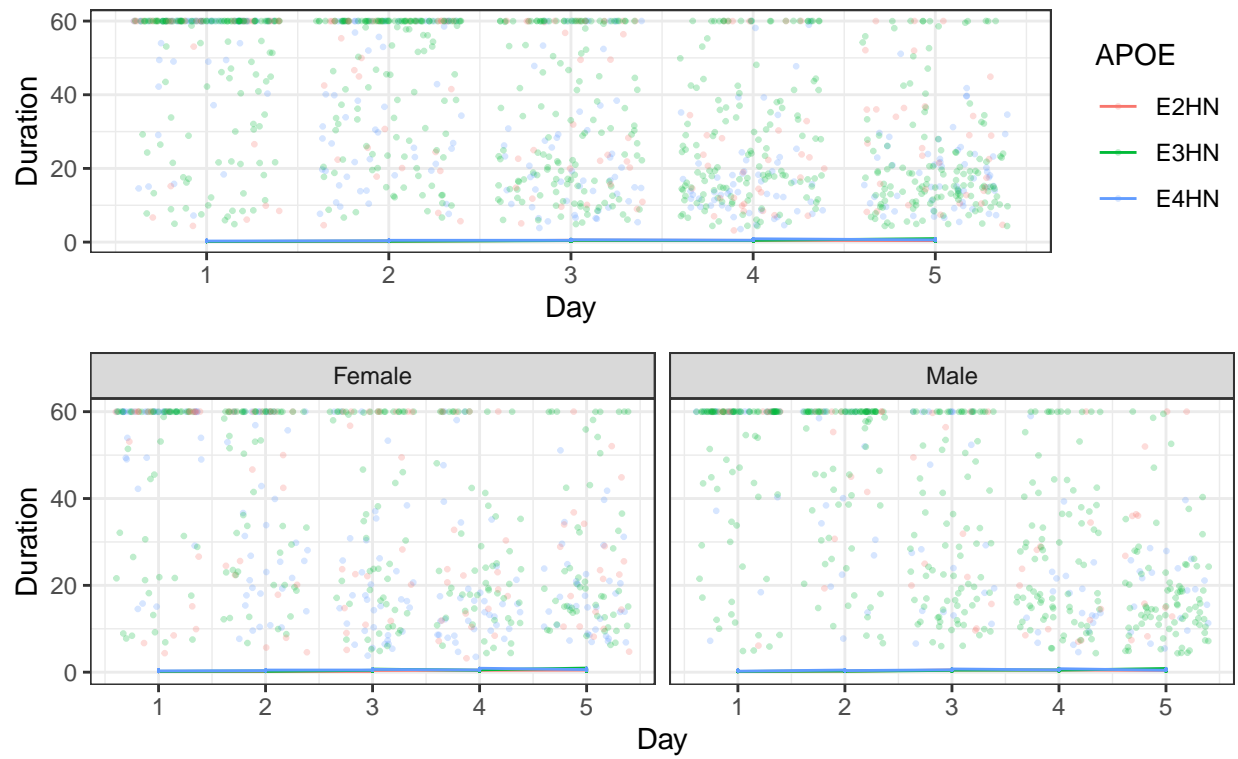






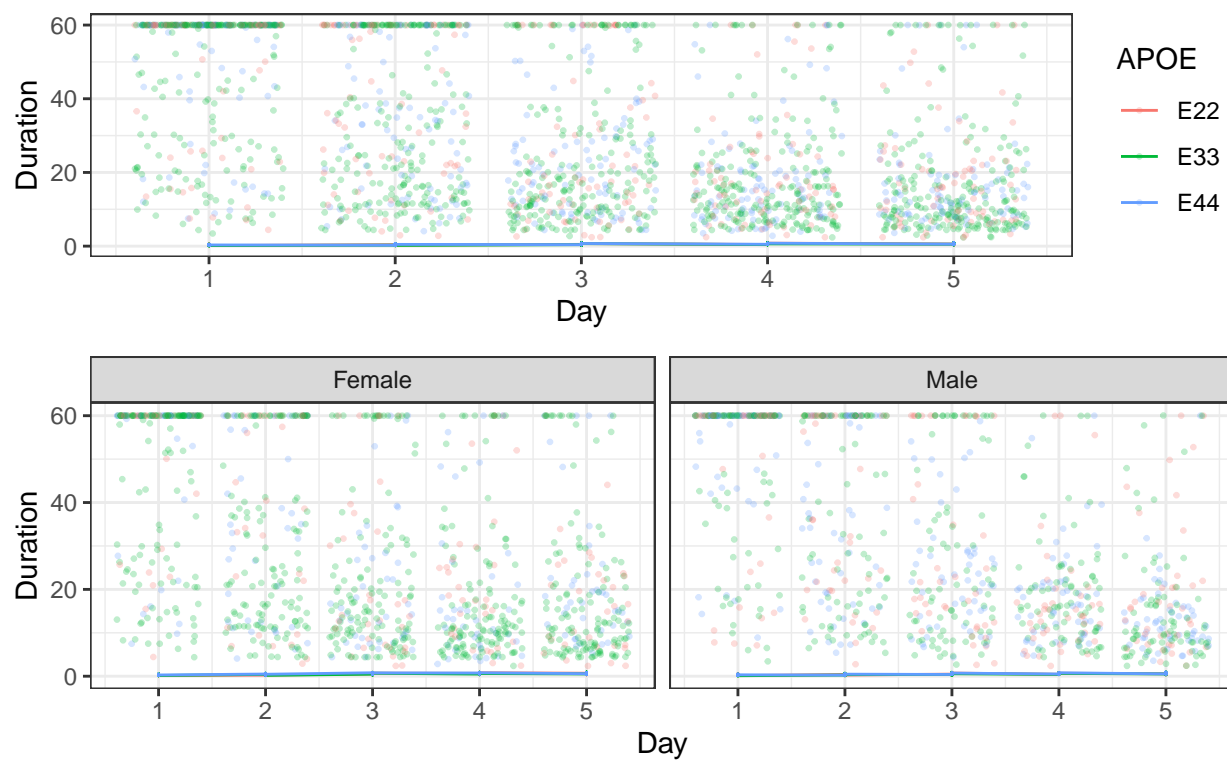
MWM Duration – HN genotypes

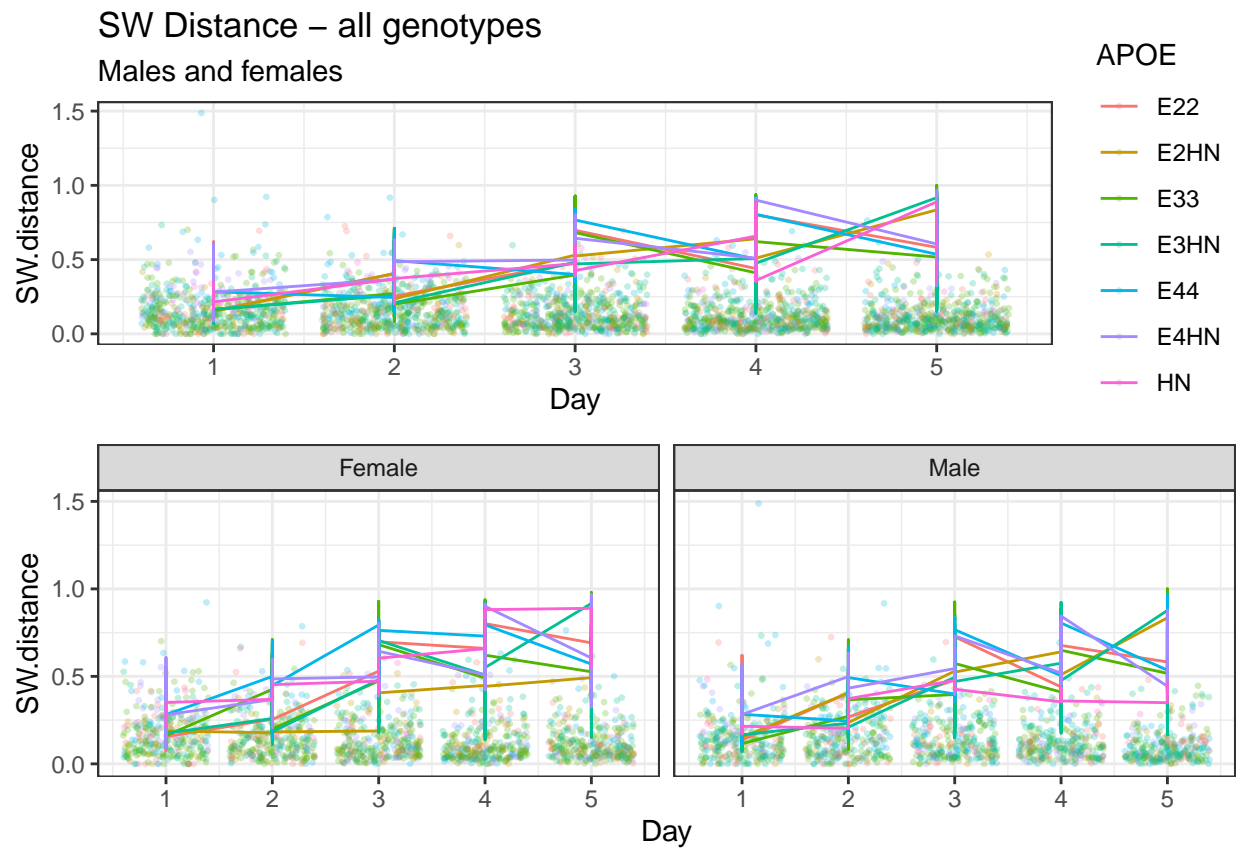
Males and females



MWM Duration – non-HN genotypes

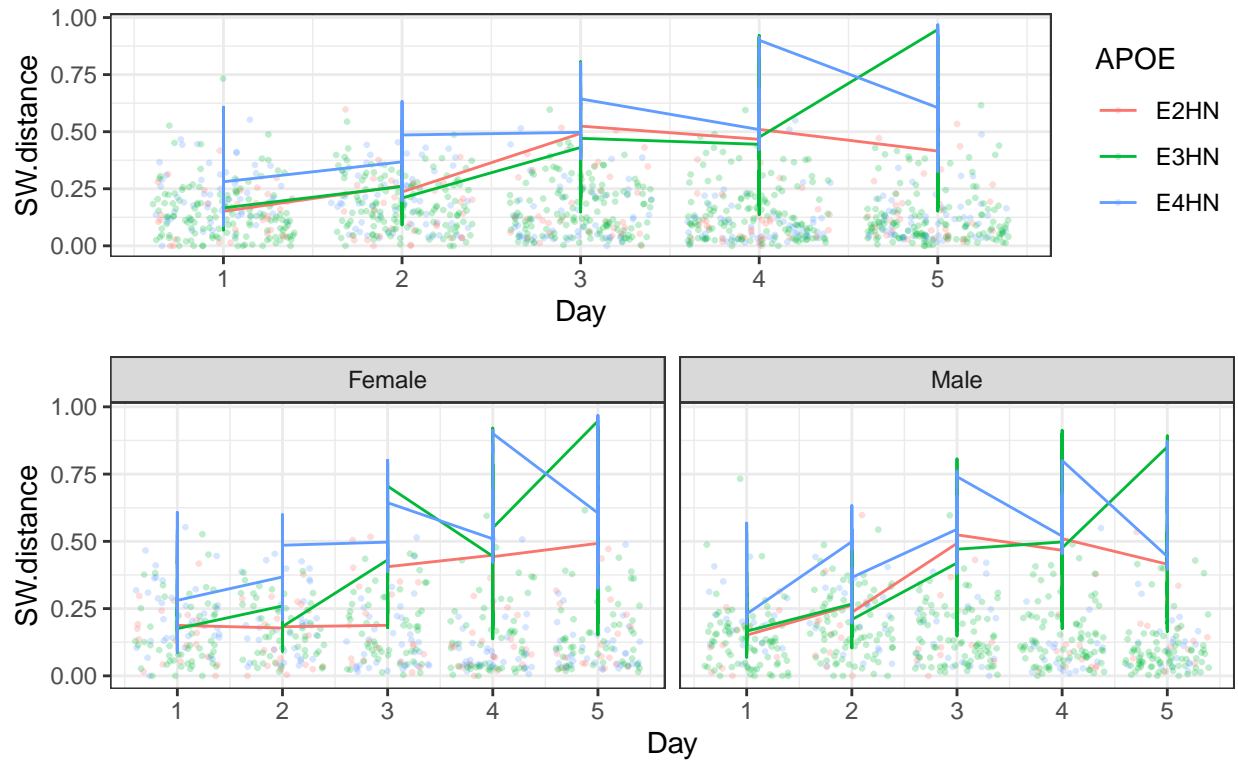
Males and females





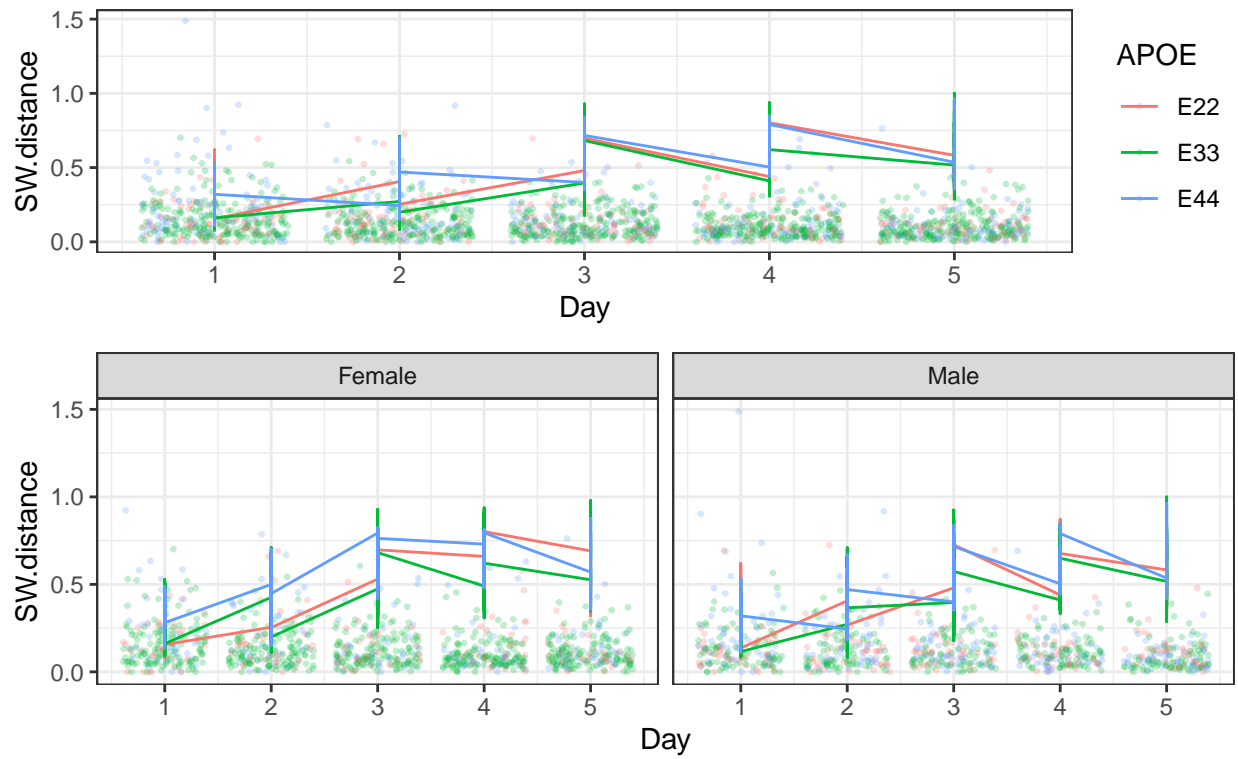
SW Distance – HN genotypes

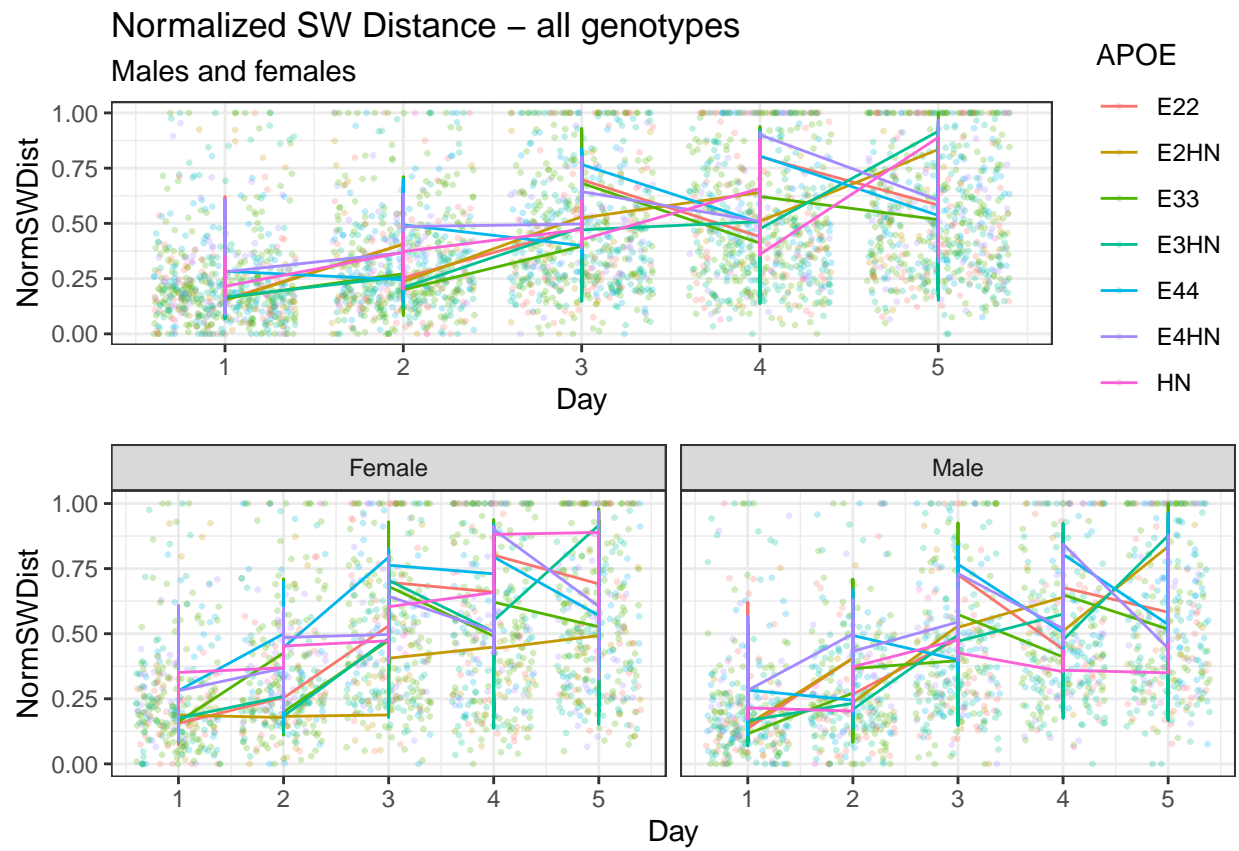
Males and females



SW Distance – non-HN genotypes

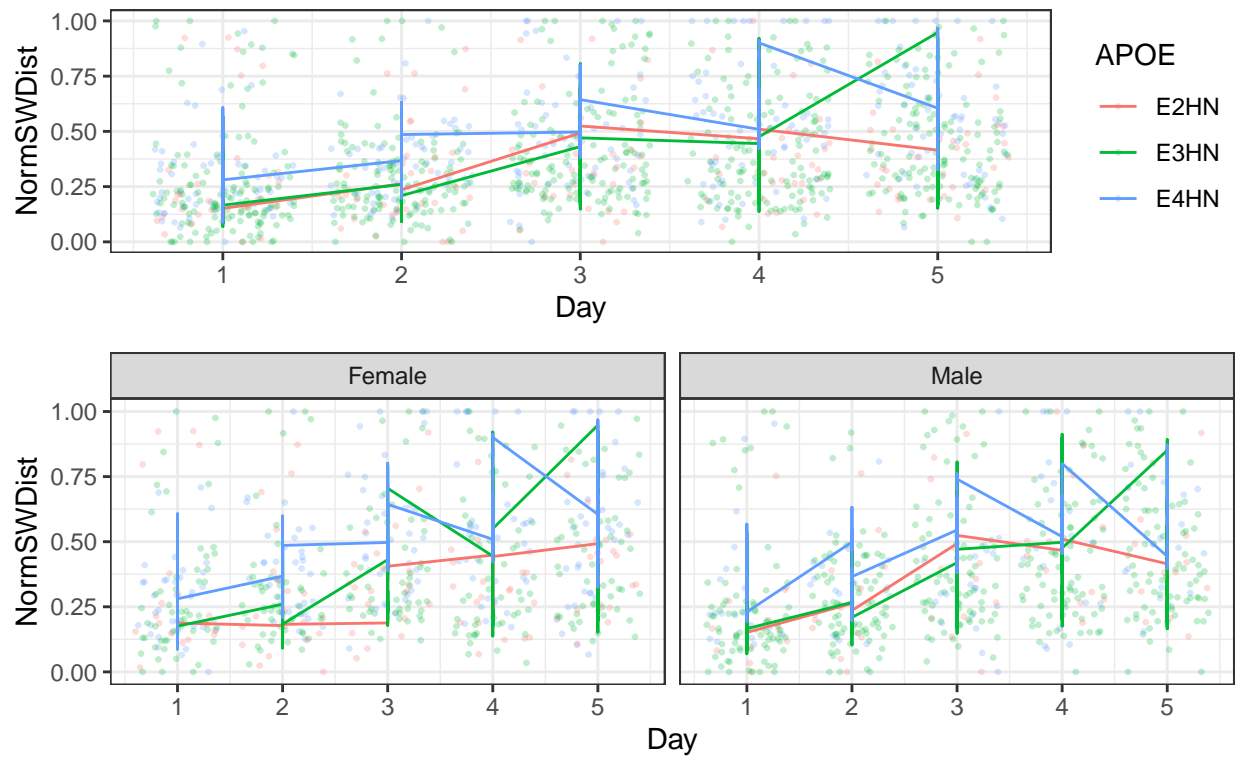
Males and females



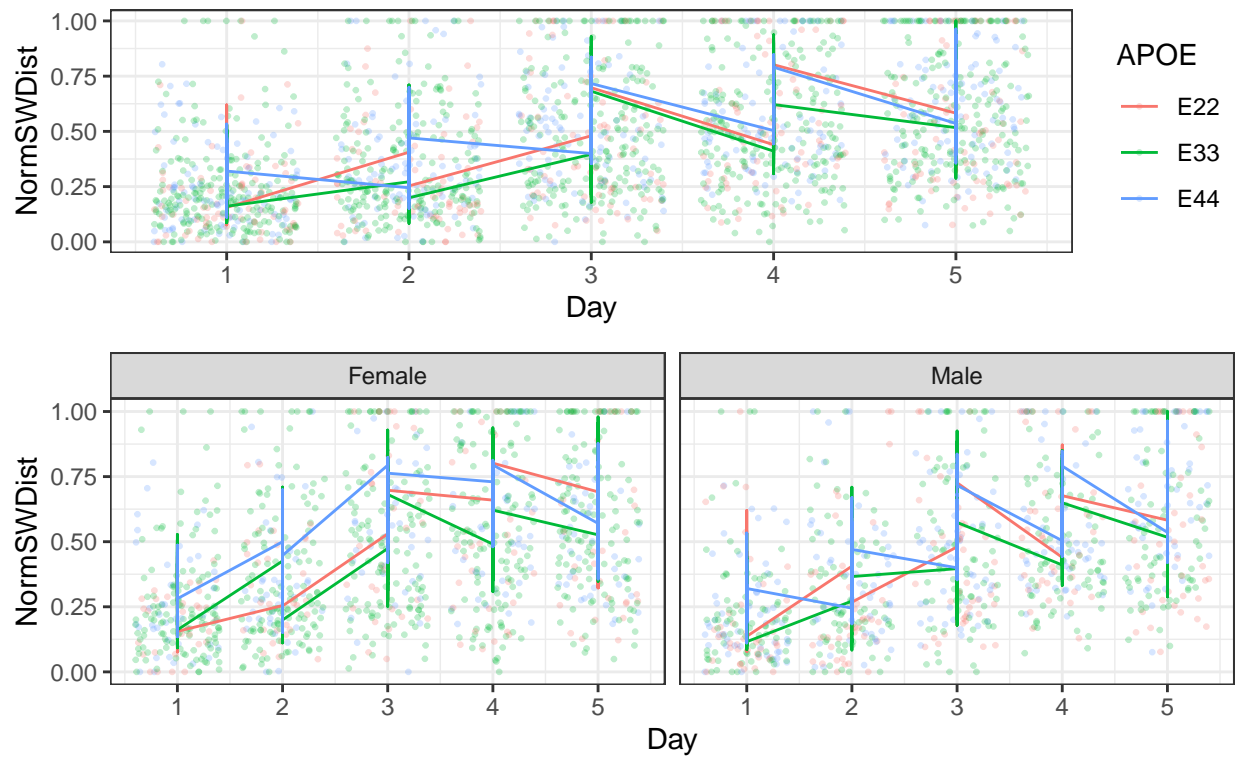


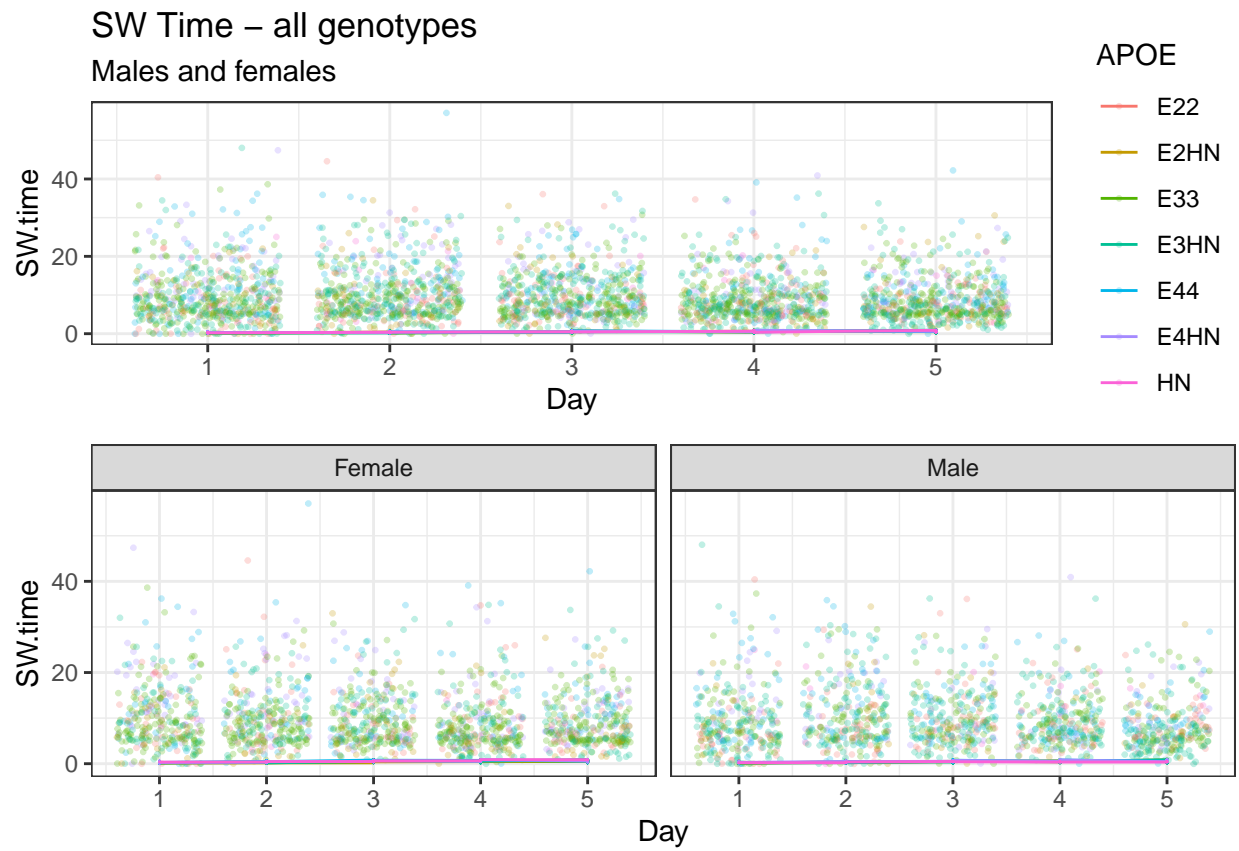
Normalized SW Distance – HN genotypes

Males and females



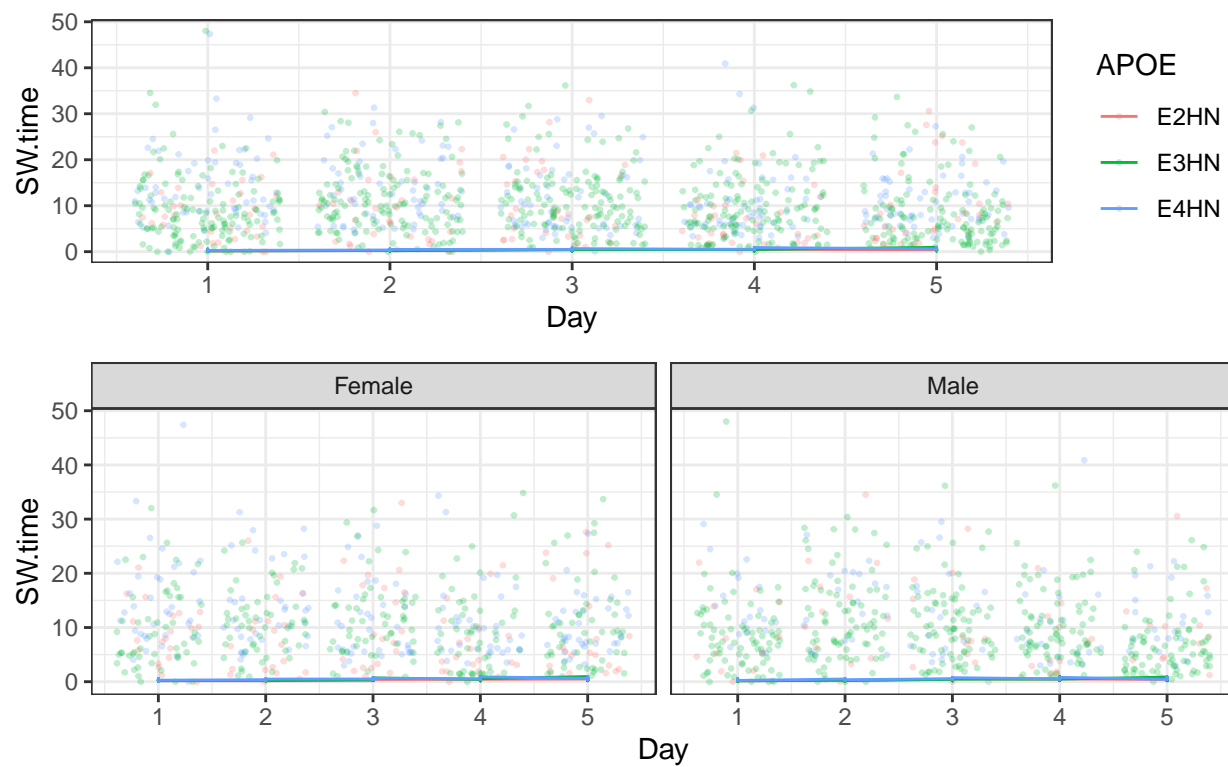
Normalized SW Distance – non-HN genotypes Males and females





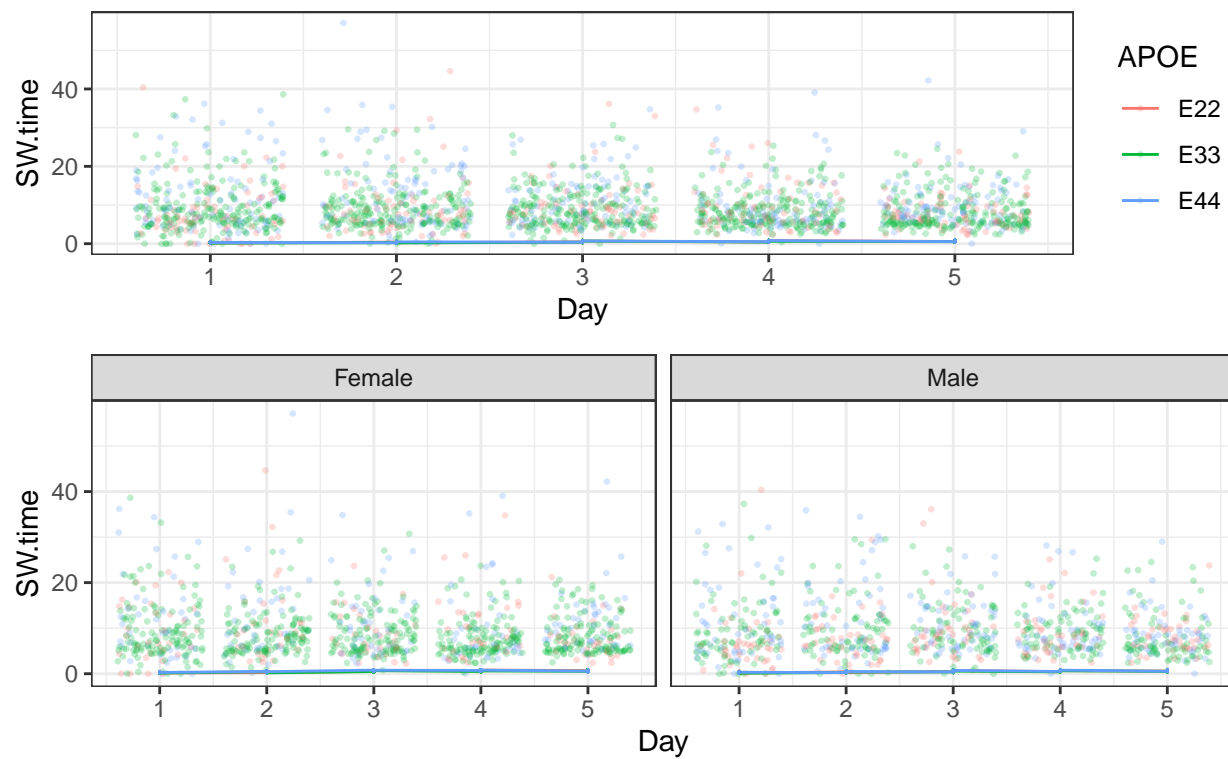
SW Time – HN genotypes

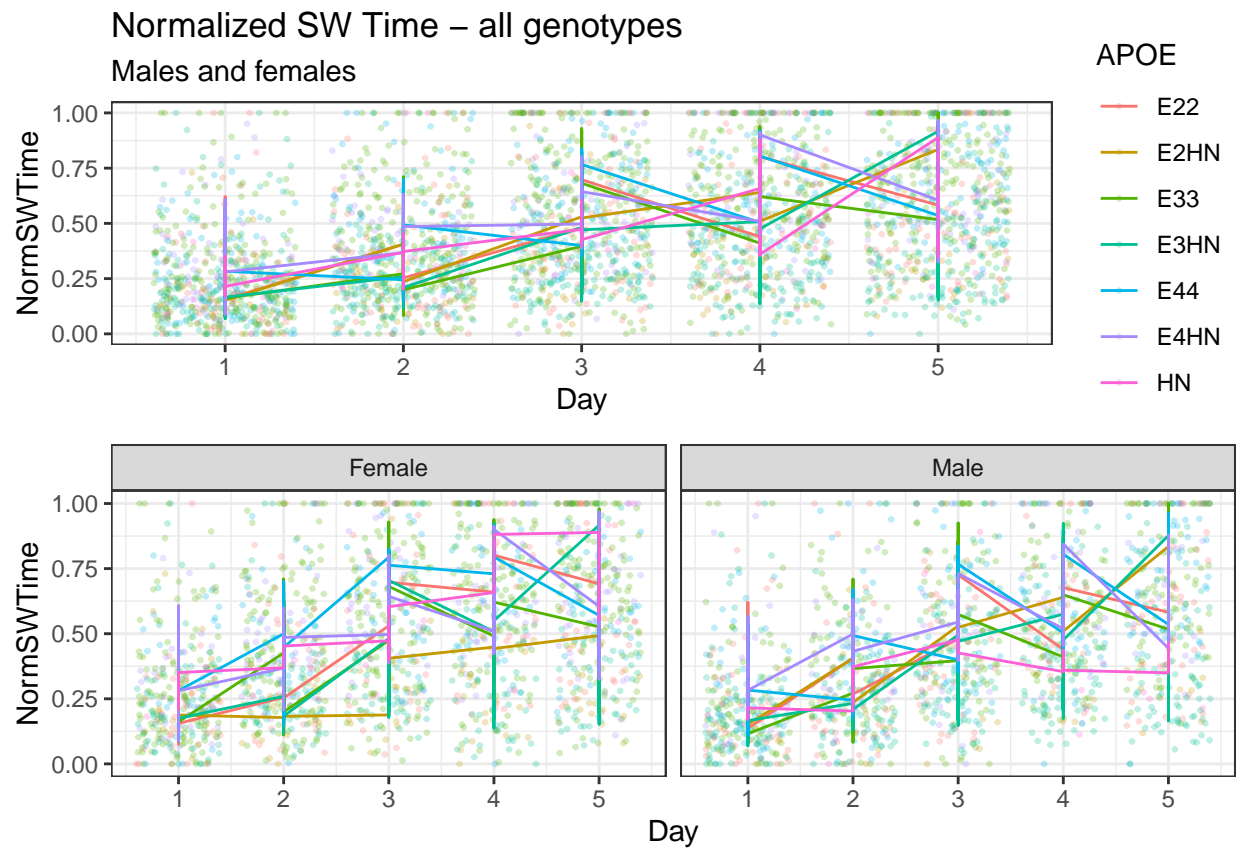
Males and females



SW Time – non-HN genotypes

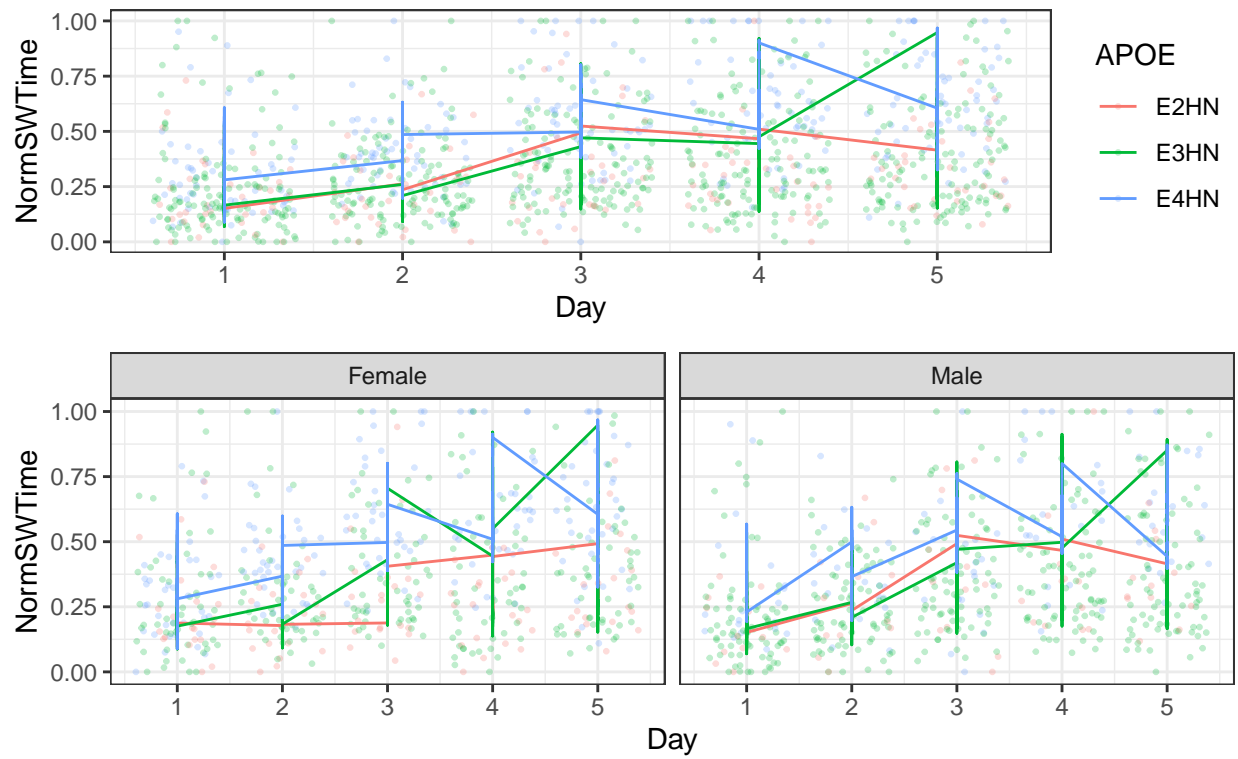
Males and females





Normalized SW Time – HN genotypes

Males and females



Normalized SW Time – non-HN genotypes

Males and females

