

# MWM Analysis

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12 months vs. 18 months

```
## 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

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## 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

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```

## 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

```

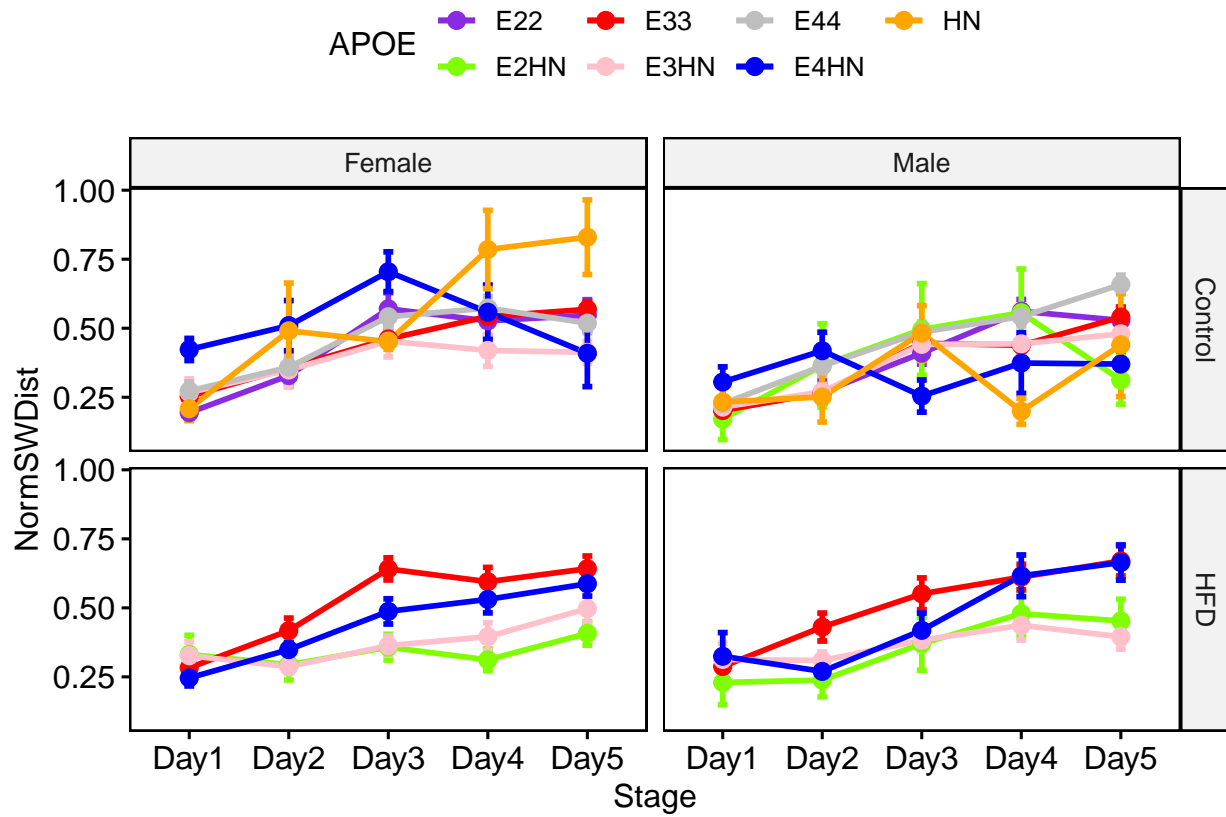
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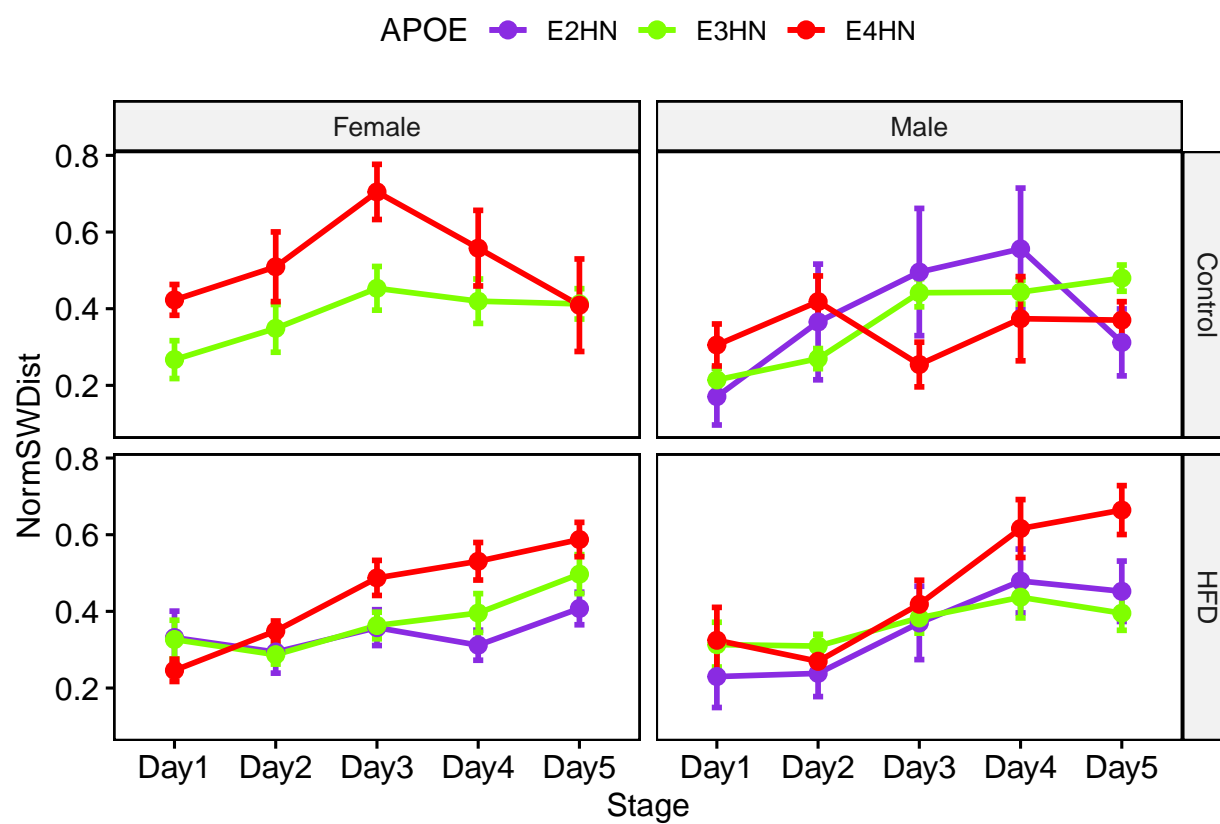
## contrast estimate SE df t.ratio p.value
## E22 - E2HN nonEst NA NA NA NA
## E22 - E33 nonEst NA NA NA NA
## E22 - E3HN nonEst NA NA NA NA
## E22 - E44 nonEst NA NA NA NA
## E22 - E4HN nonEst NA NA NA NA
## E22 - HN nonEst NA NA NA NA
## E2HN - E33 nonEst NA NA NA NA
## E2HN - E3HN nonEst NA NA NA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

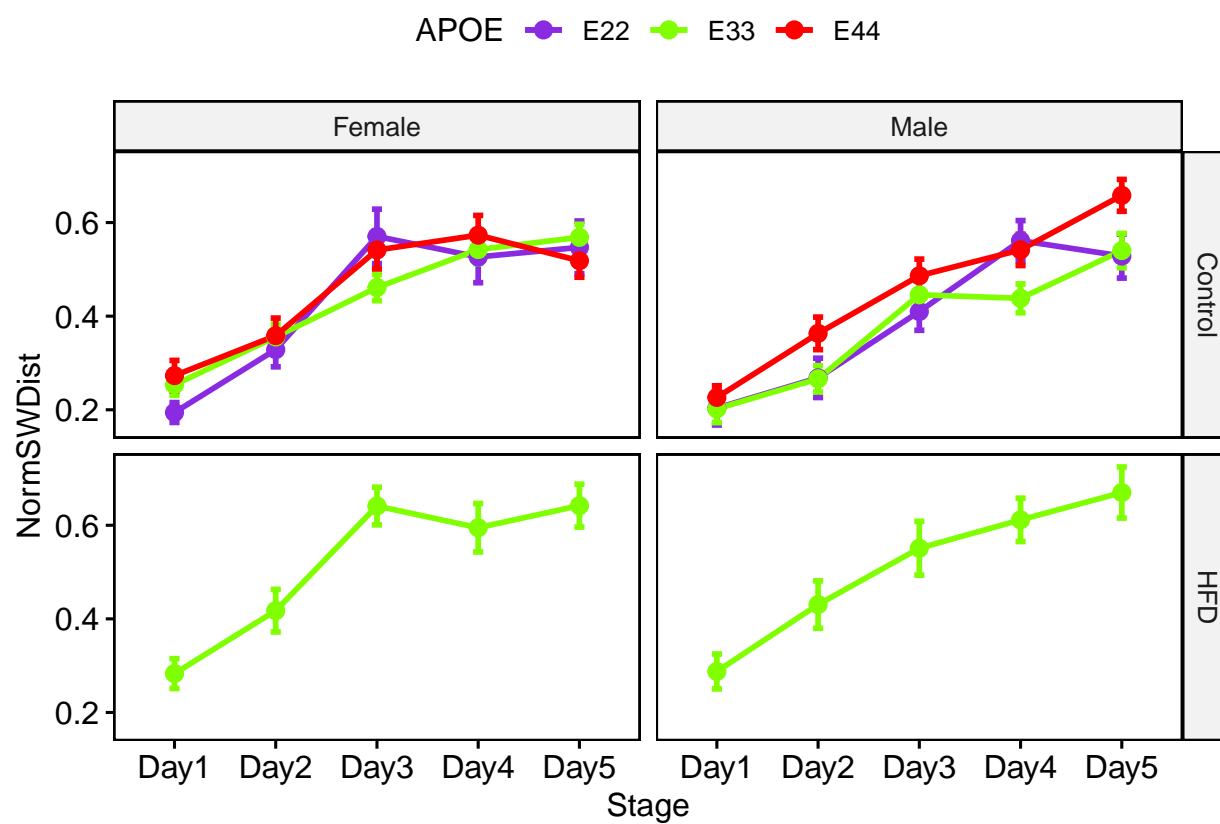
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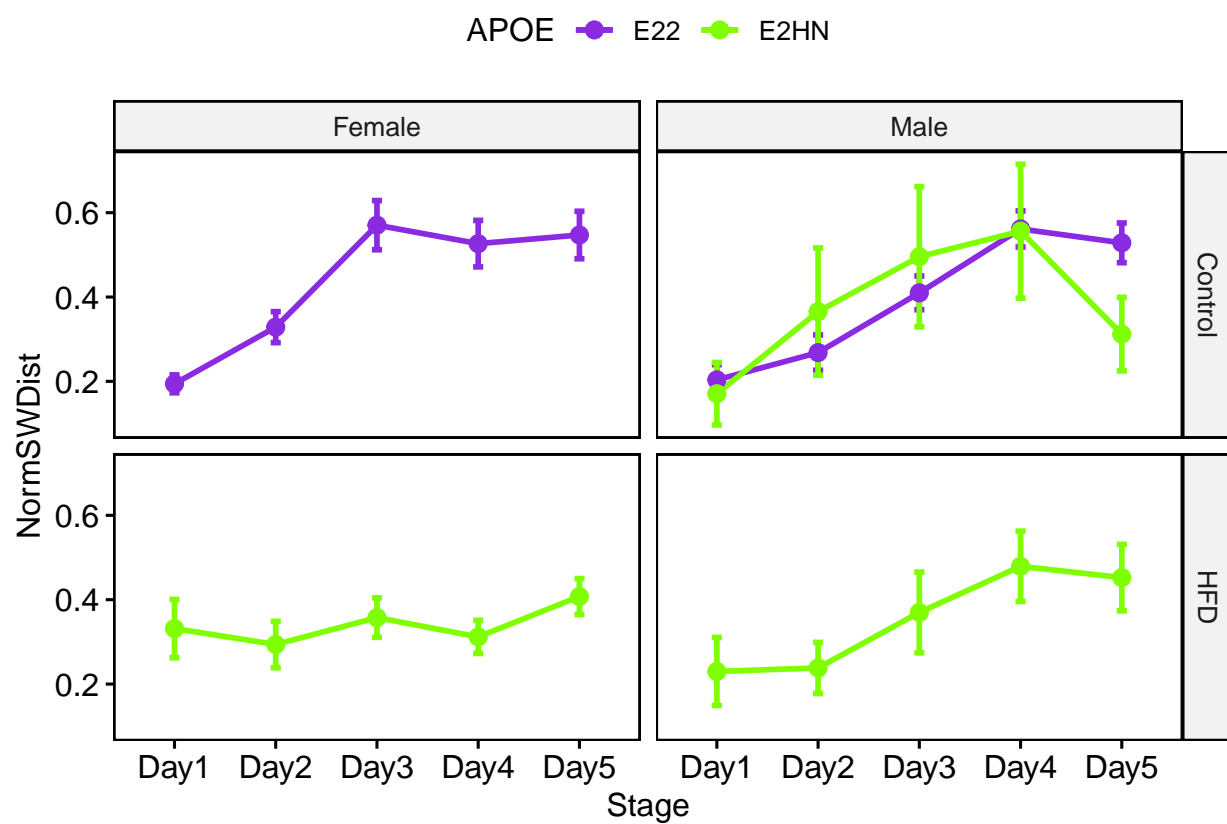
```
## E2HN - HN      nonEst      NA NA      NA      NA
## E33 - E3HN     0.1317 0.0315 129    4.176 0.0010
## E33 - E44      nonEst      NA NA      NA      NA
## E33 - E4HN     0.0747 0.0441 129    1.693 0.6222
## E33 - HN       nonEst      NA NA      NA      NA
## E3HN - E44     nonEst      NA NA      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
```

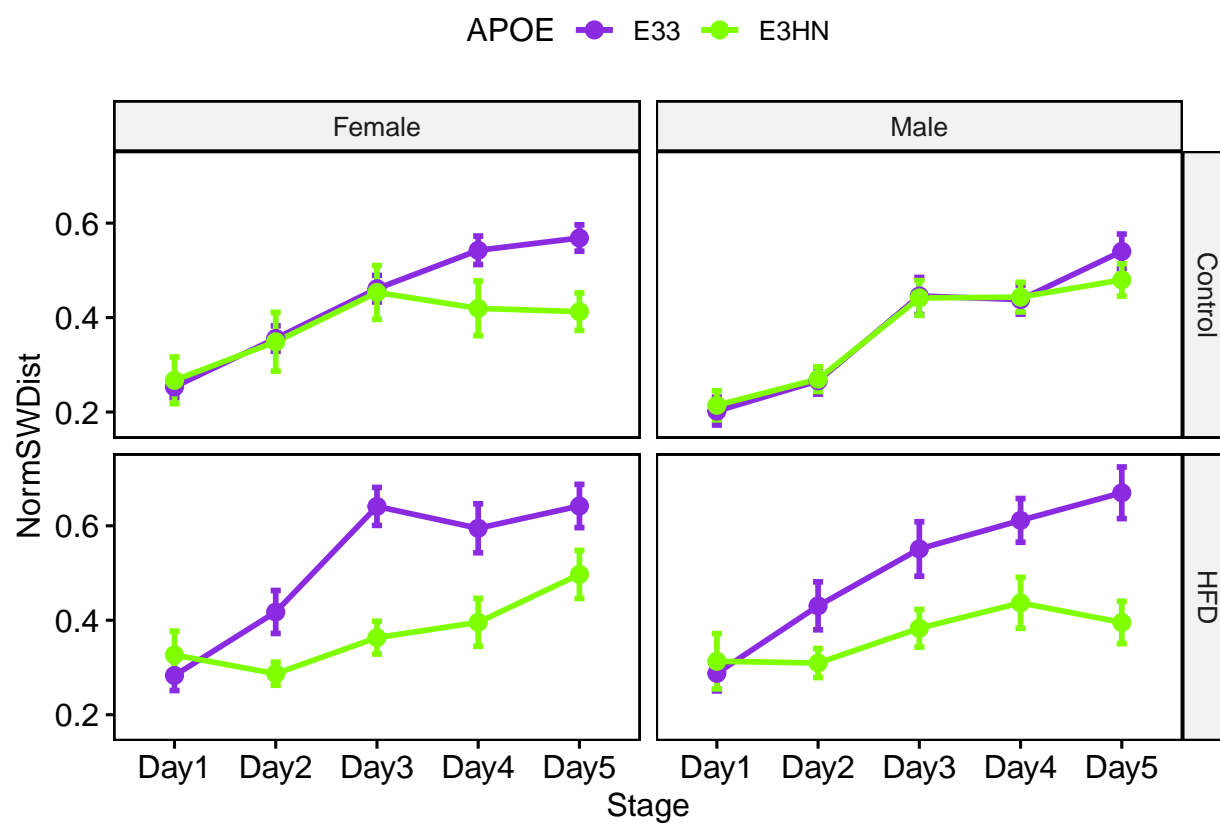
```
lm <- lm(Duration ~ SexAPOEDiet, reg_trials) summary(lm) anova(lm)
```



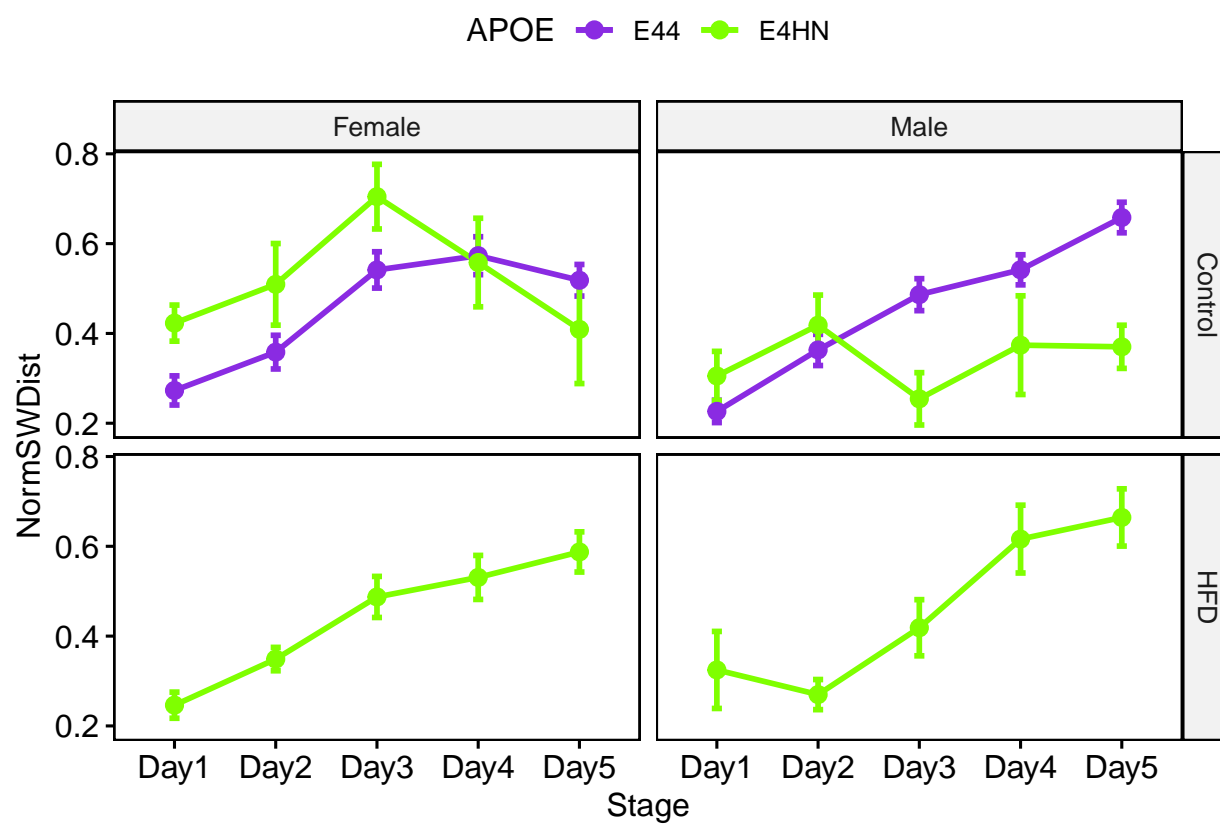


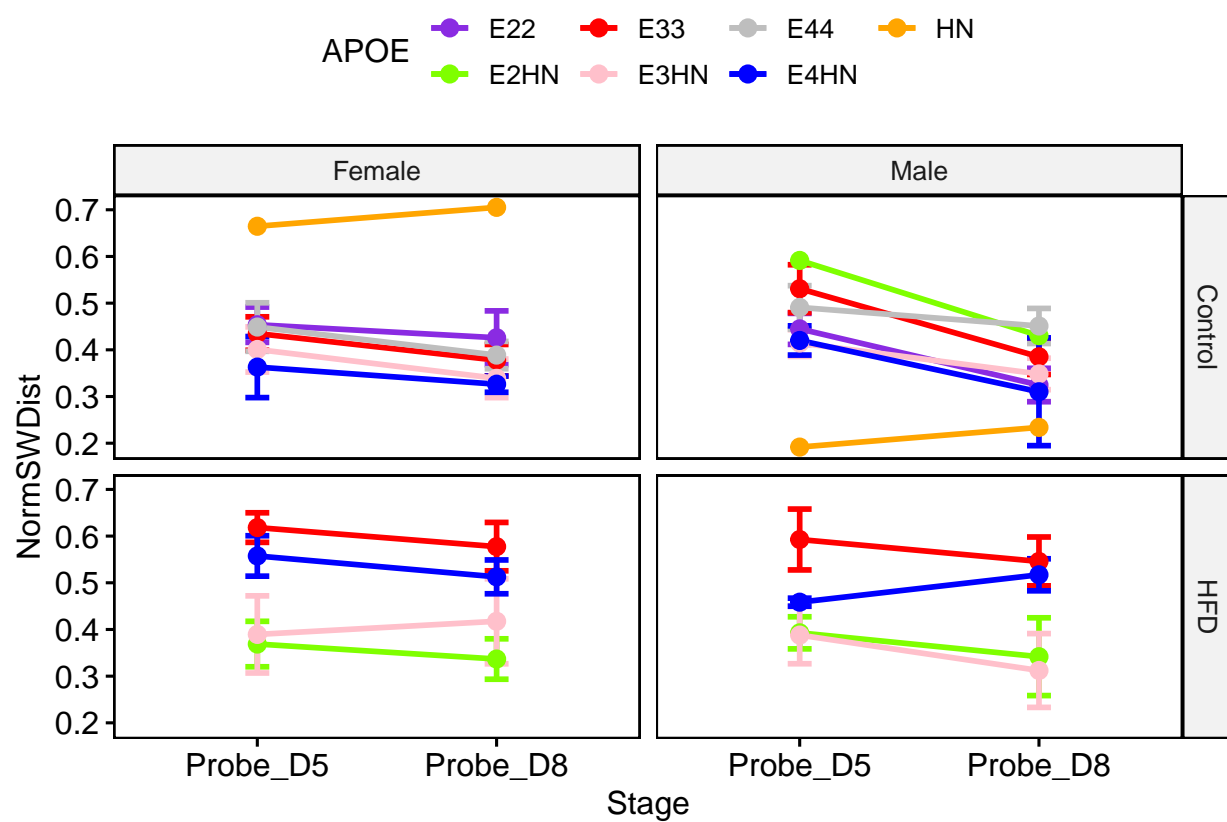


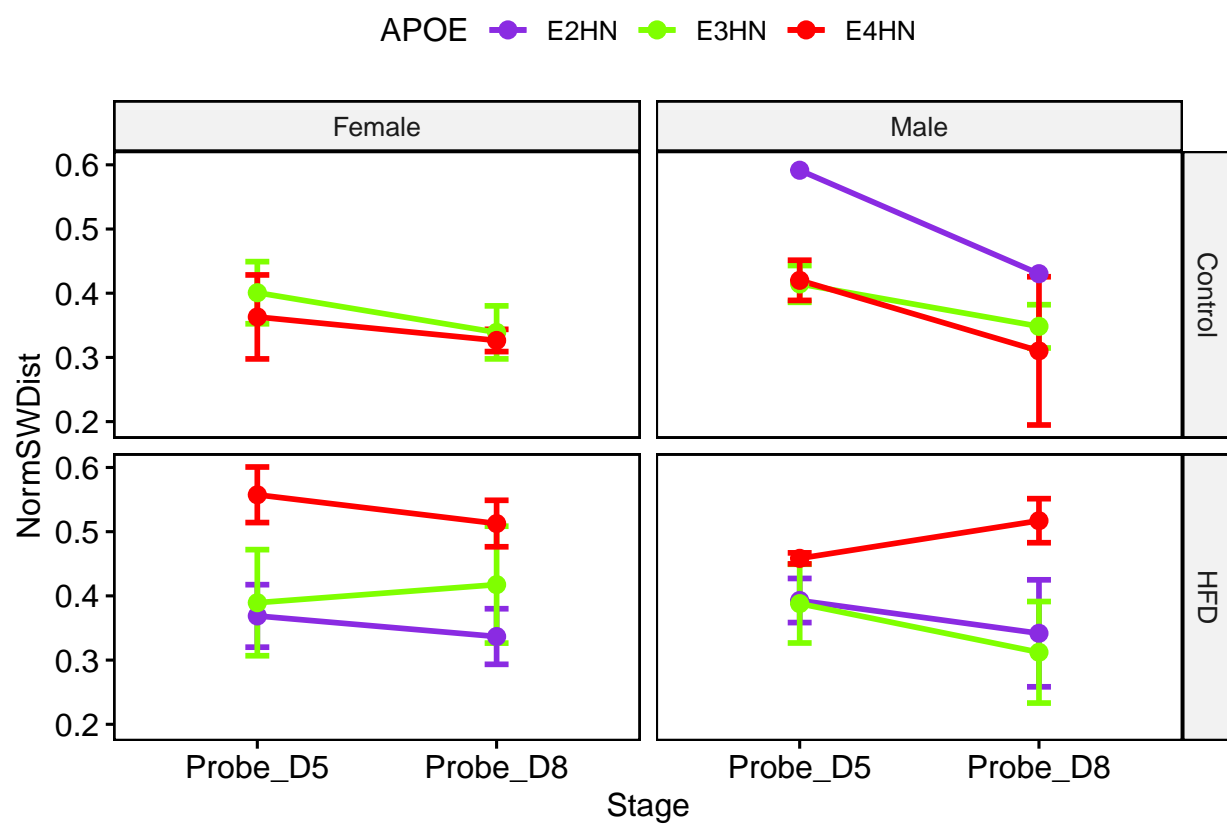


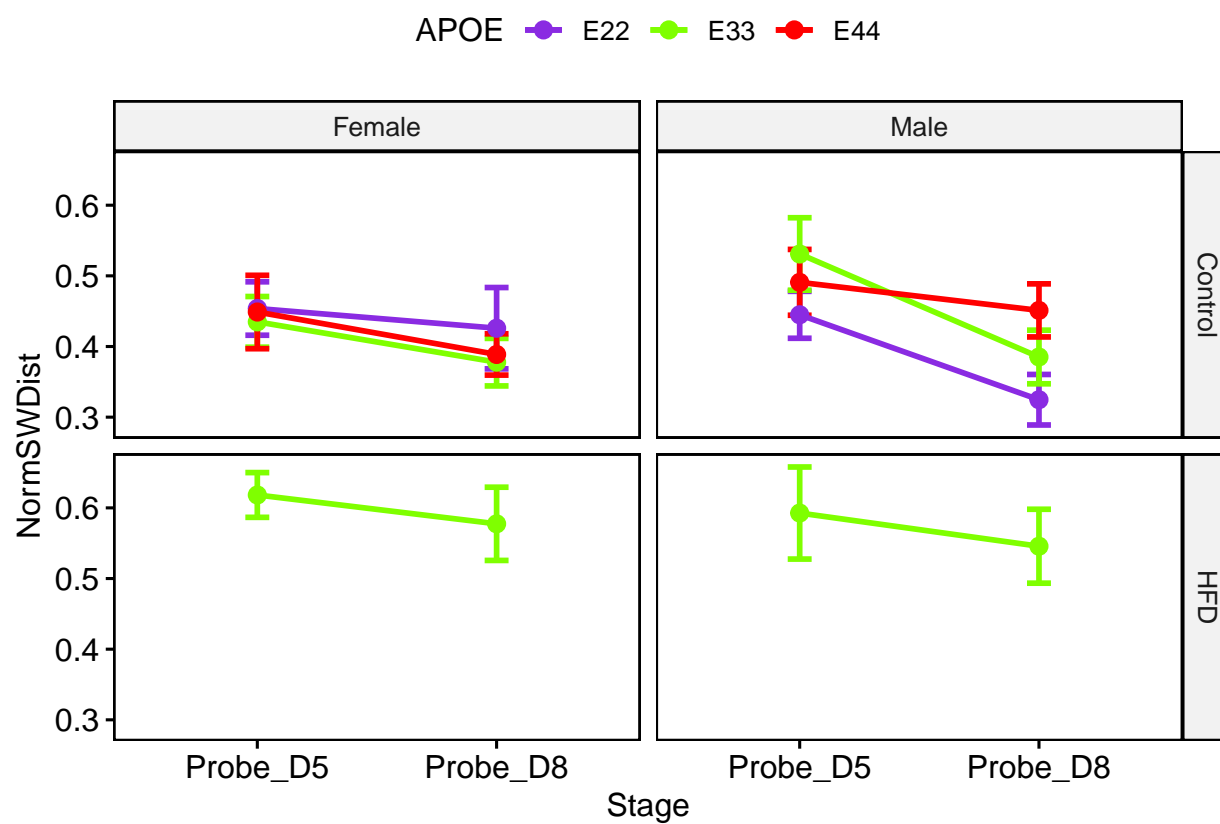


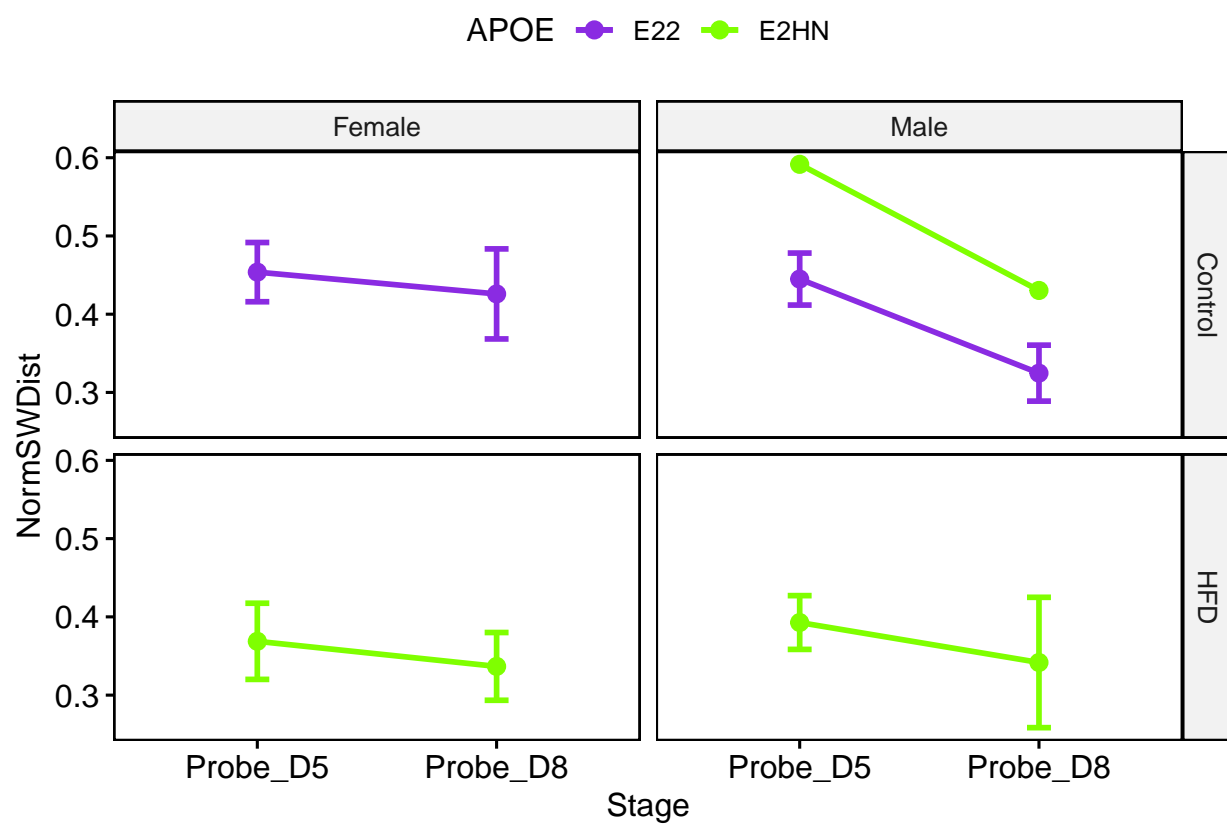


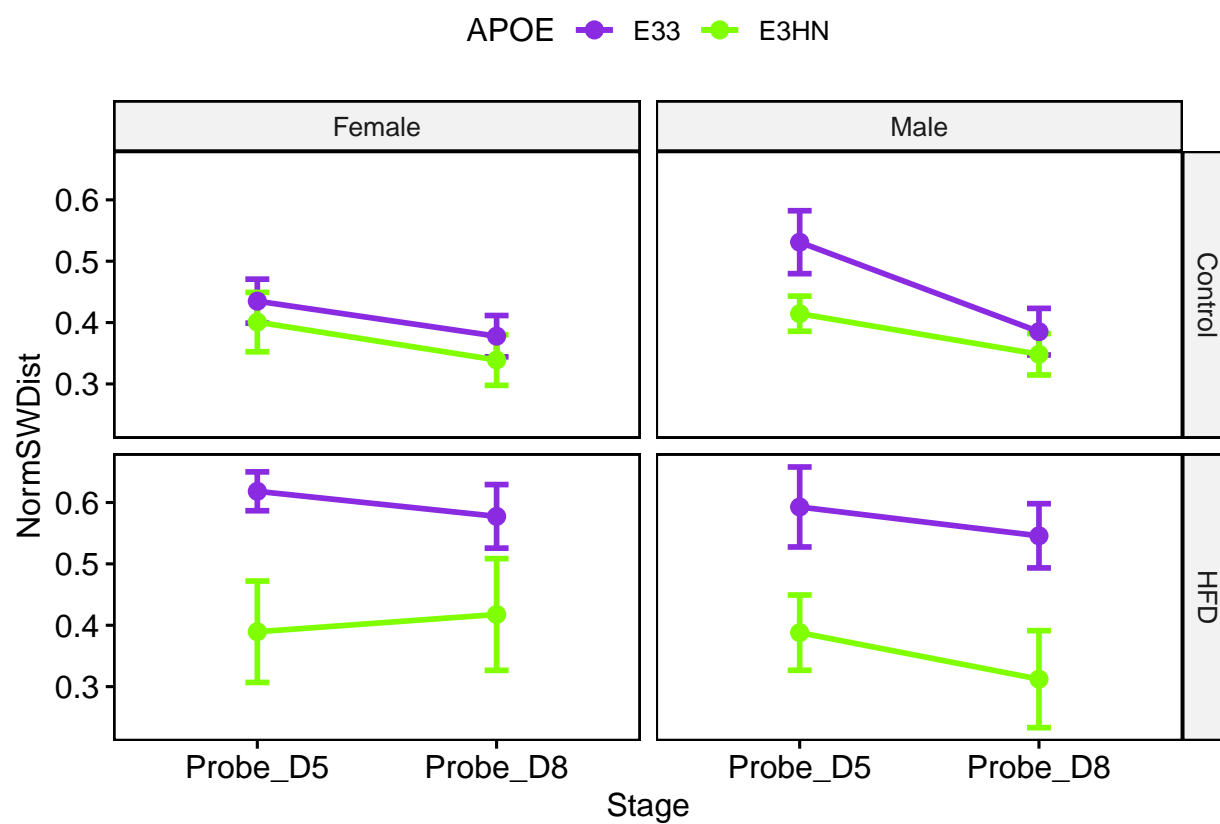


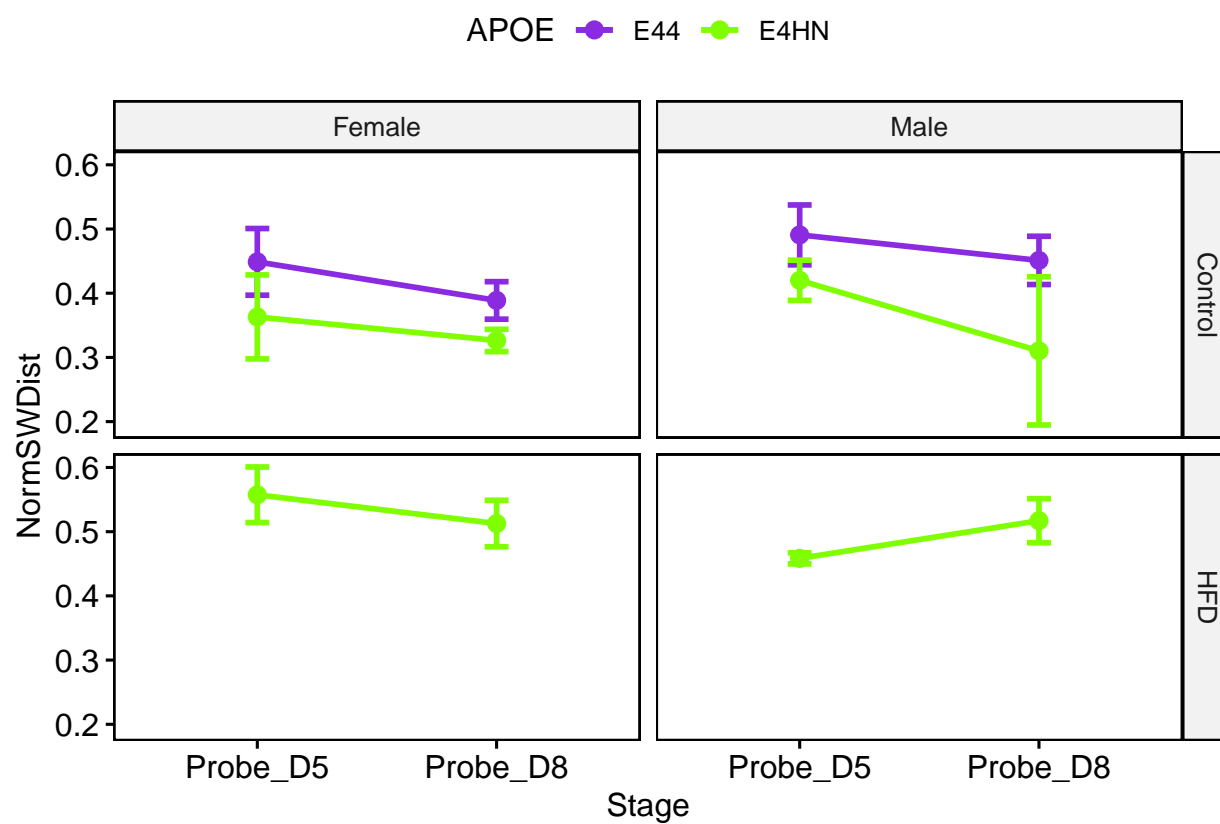




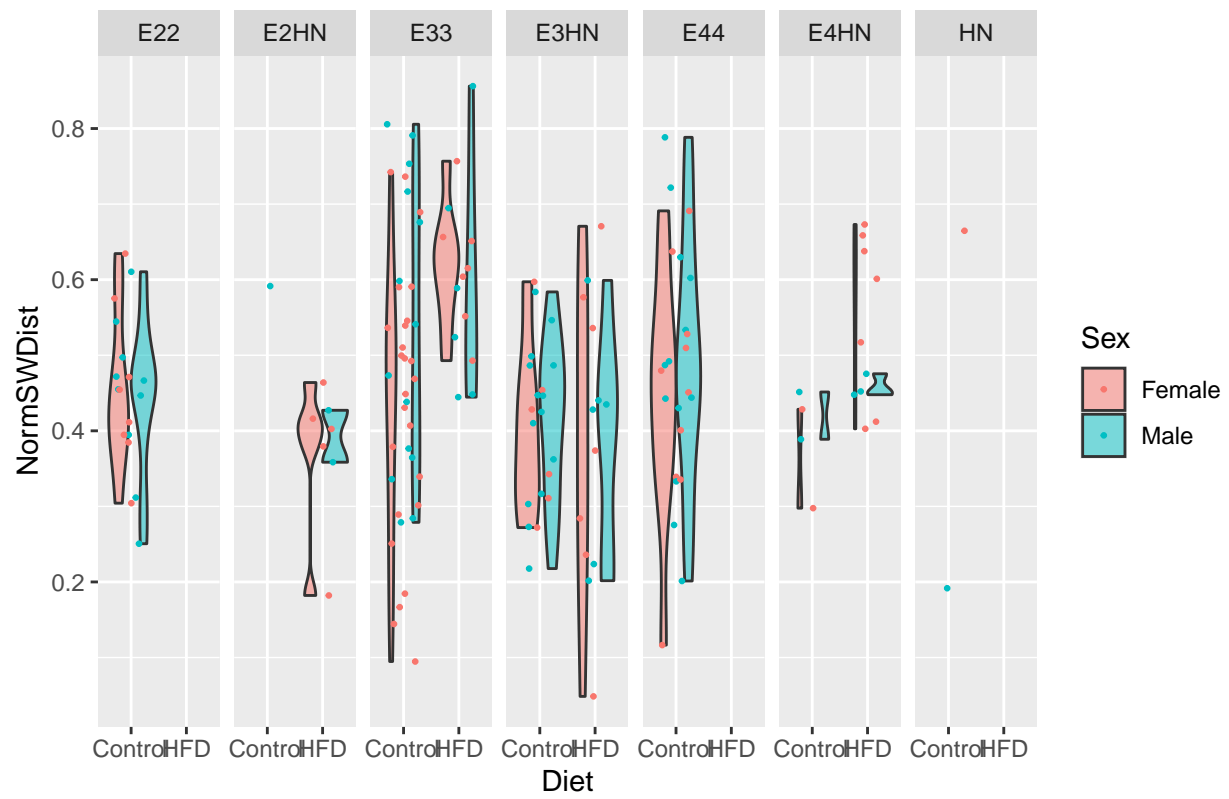






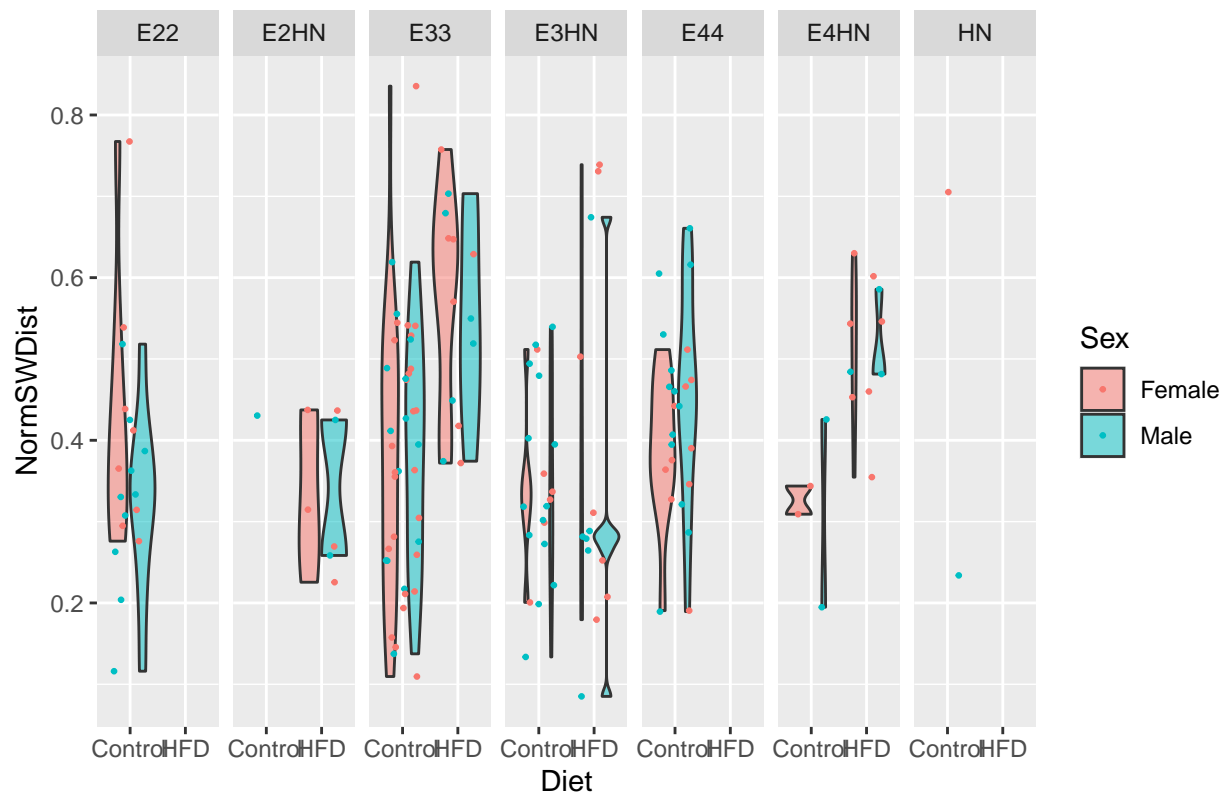


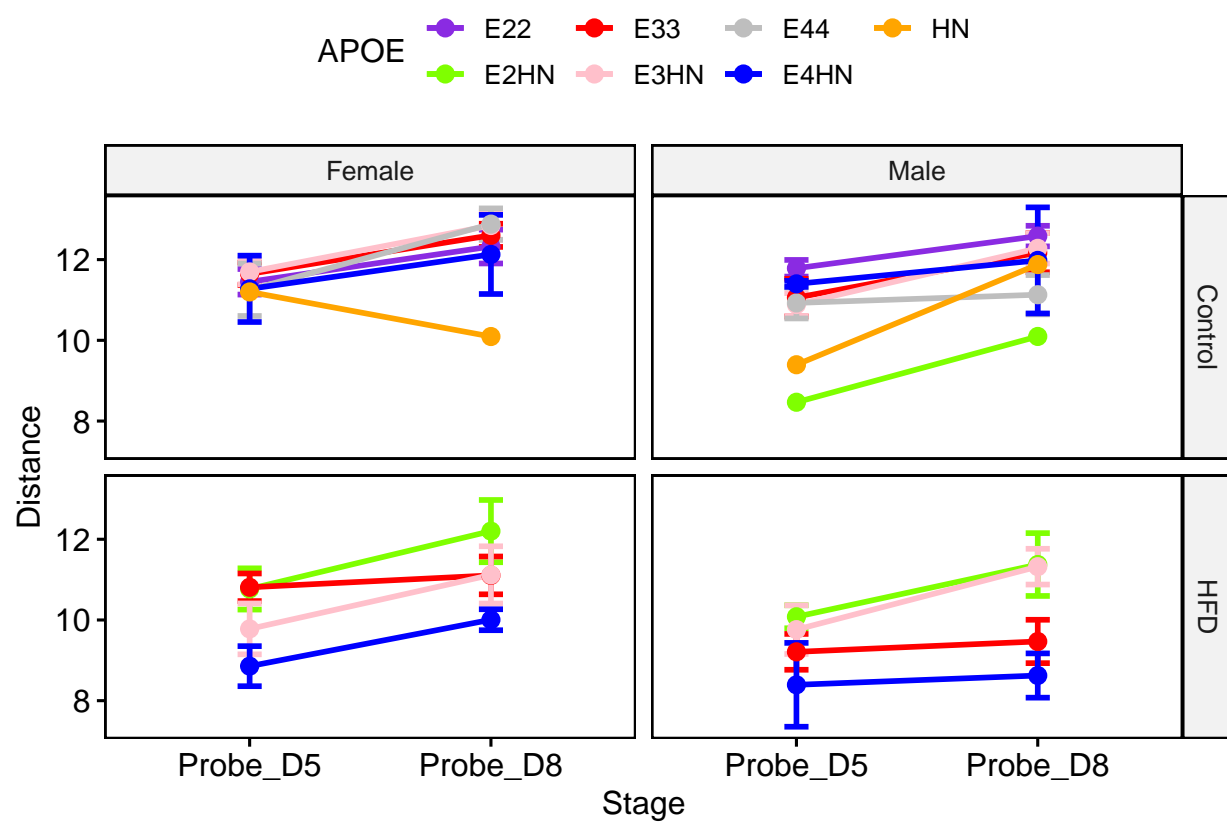
# Probe Trial: Day 5

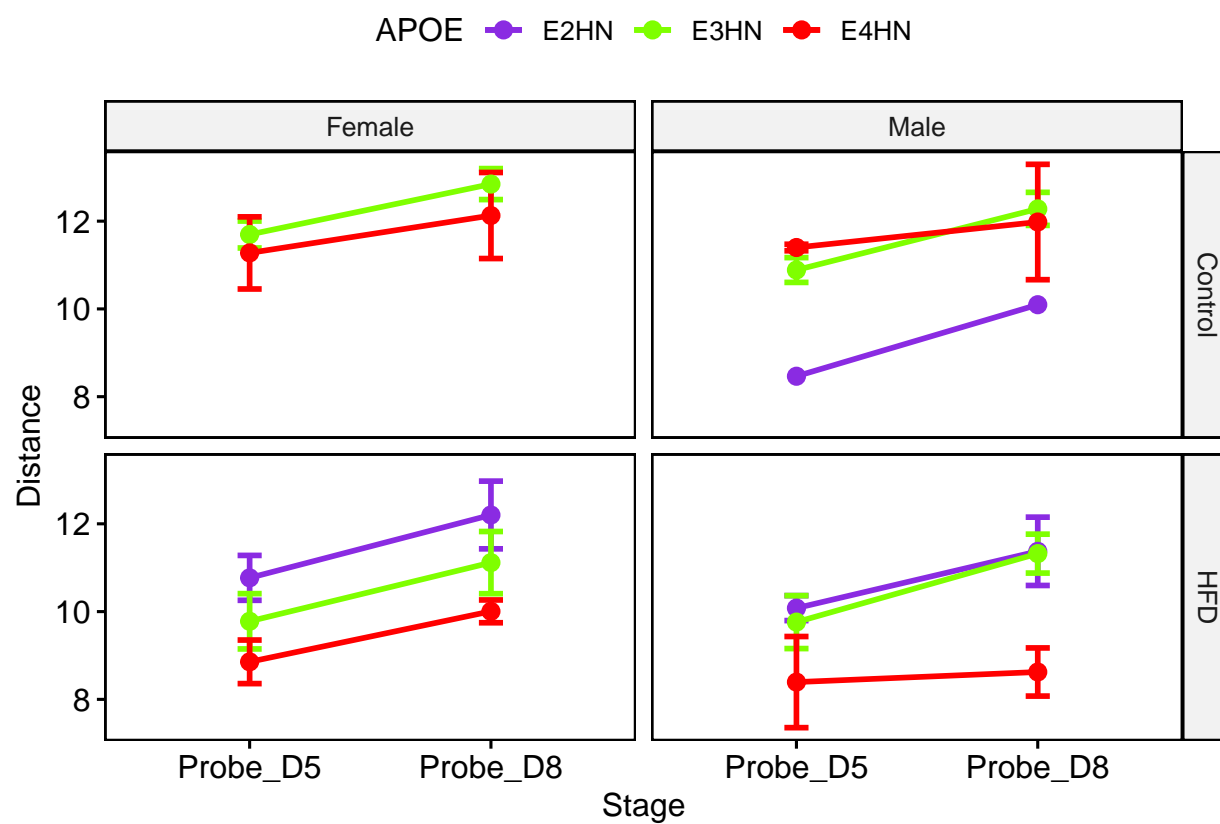


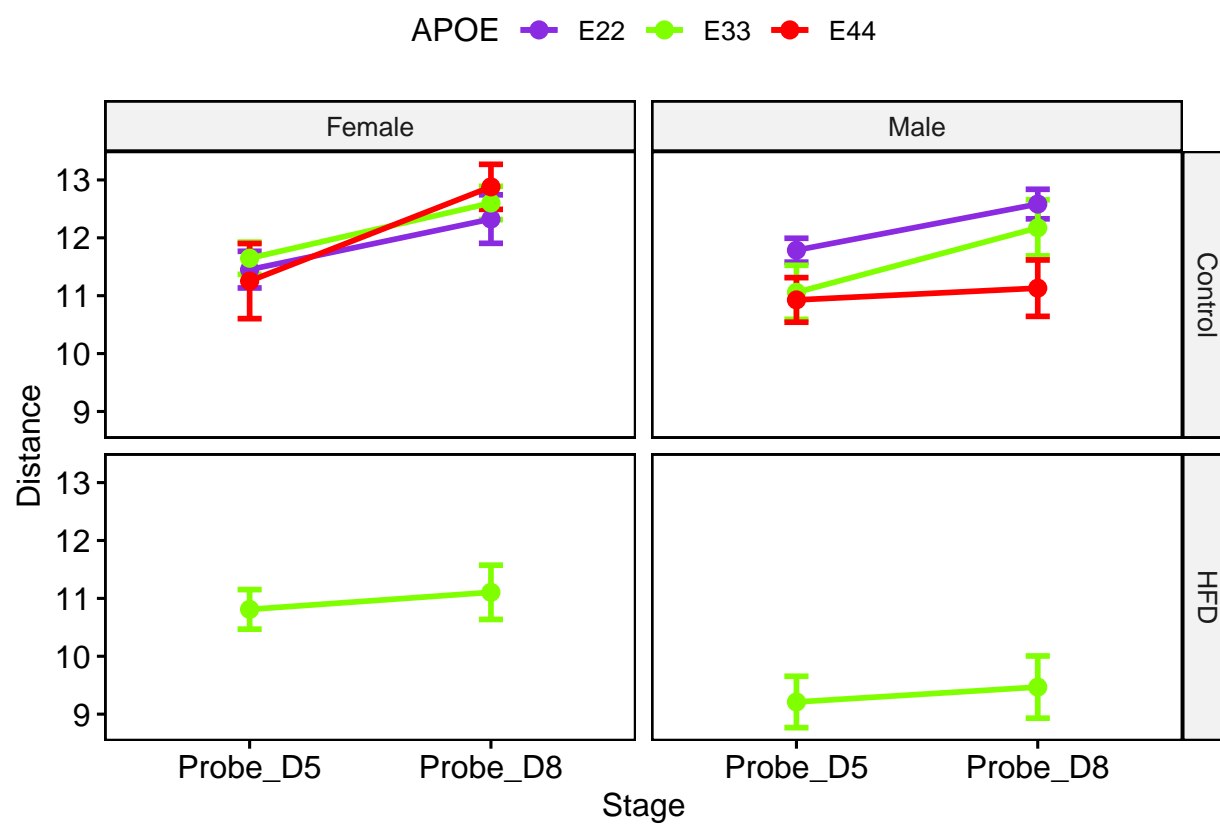


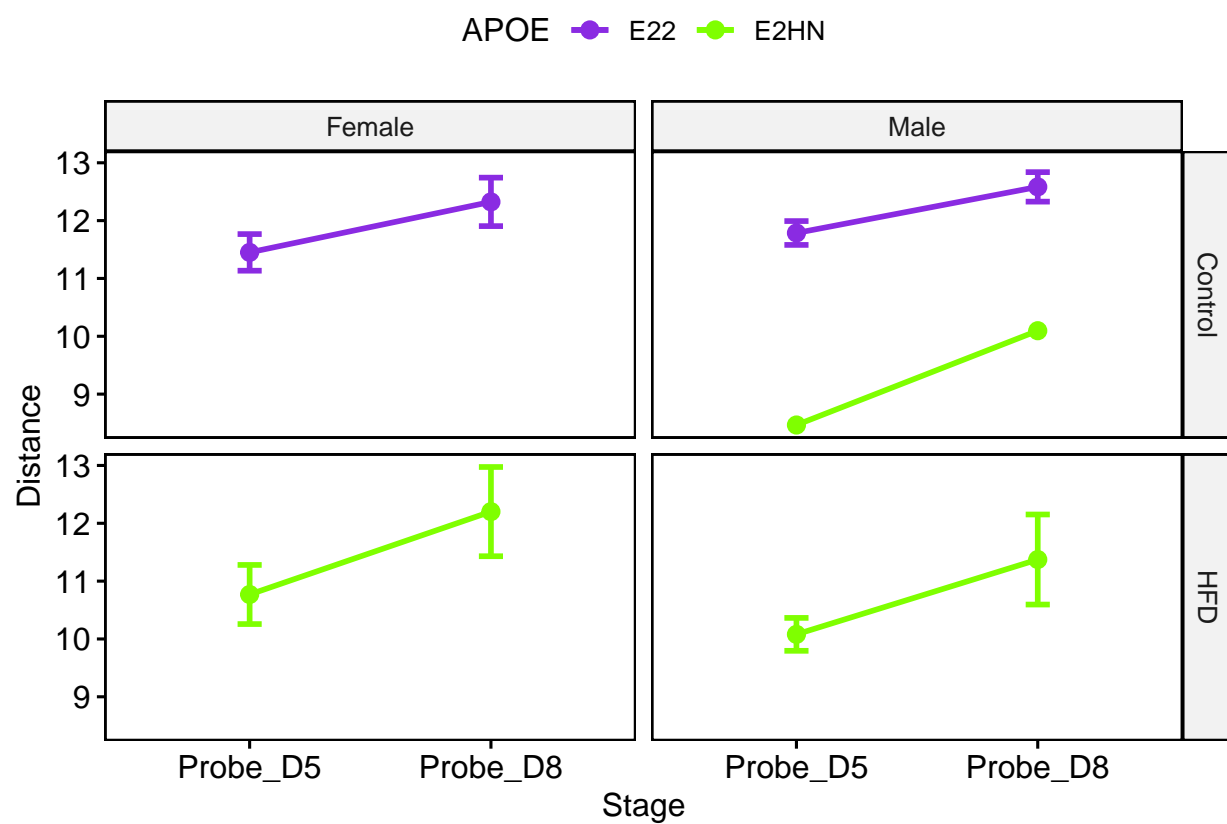
# Probe Trial: Day 8

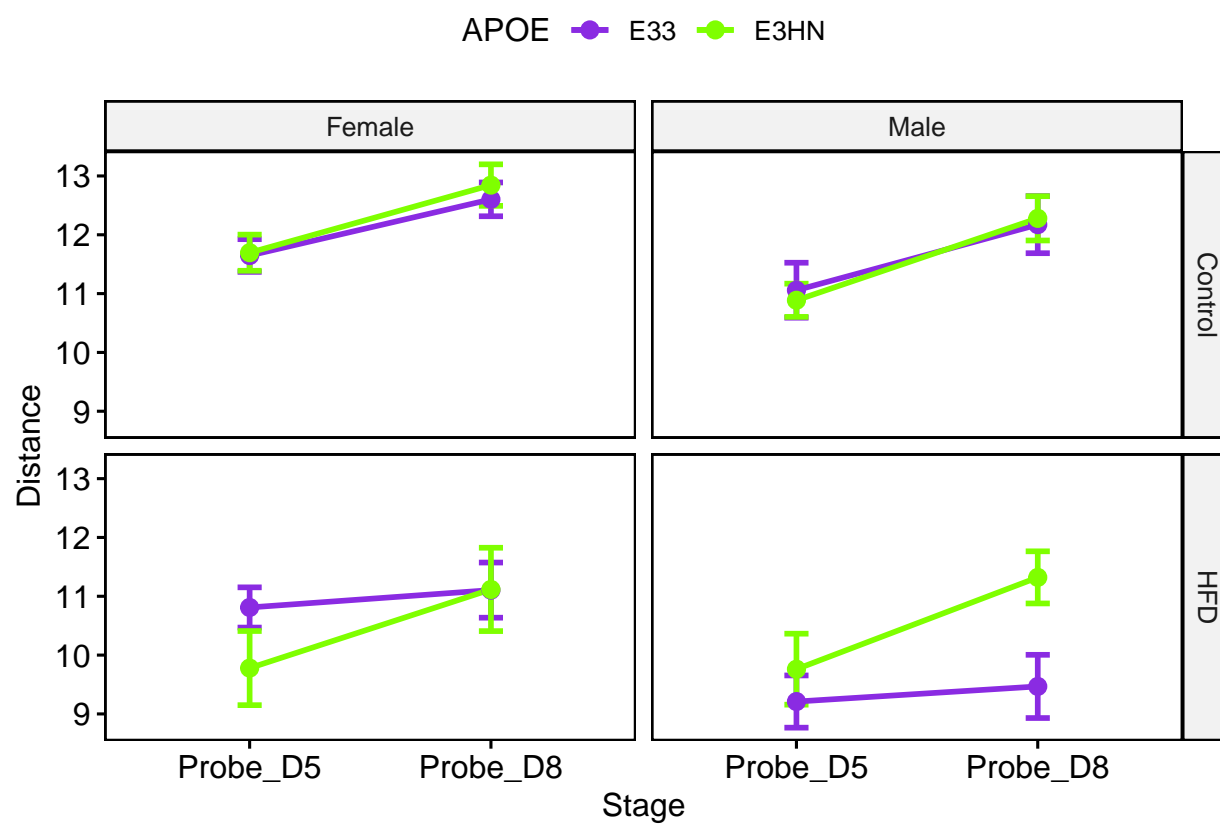


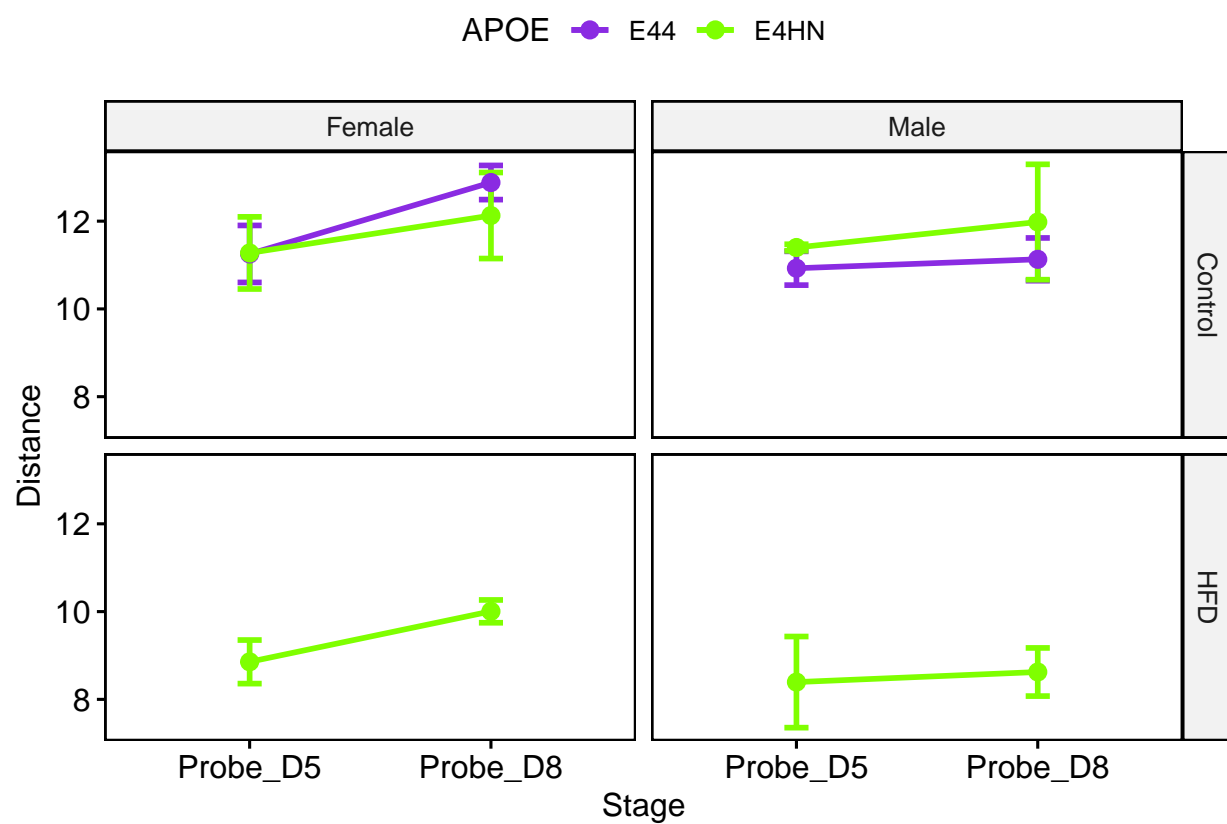


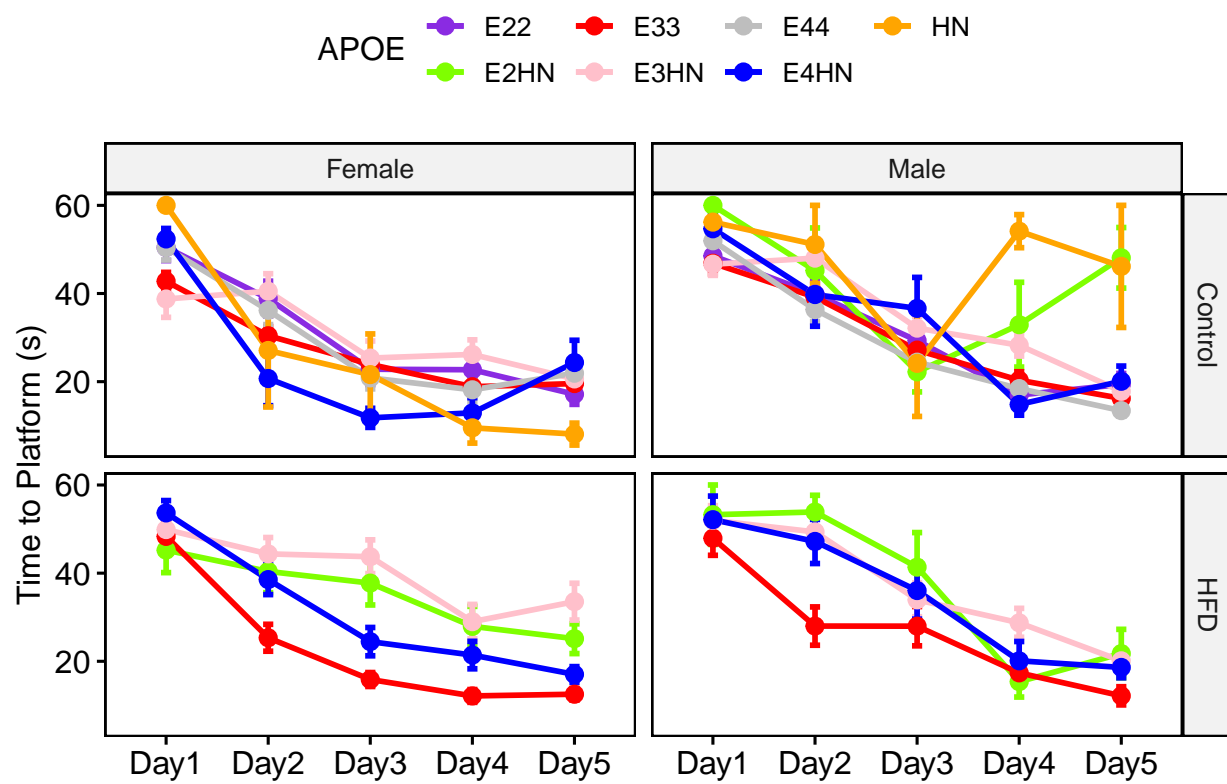




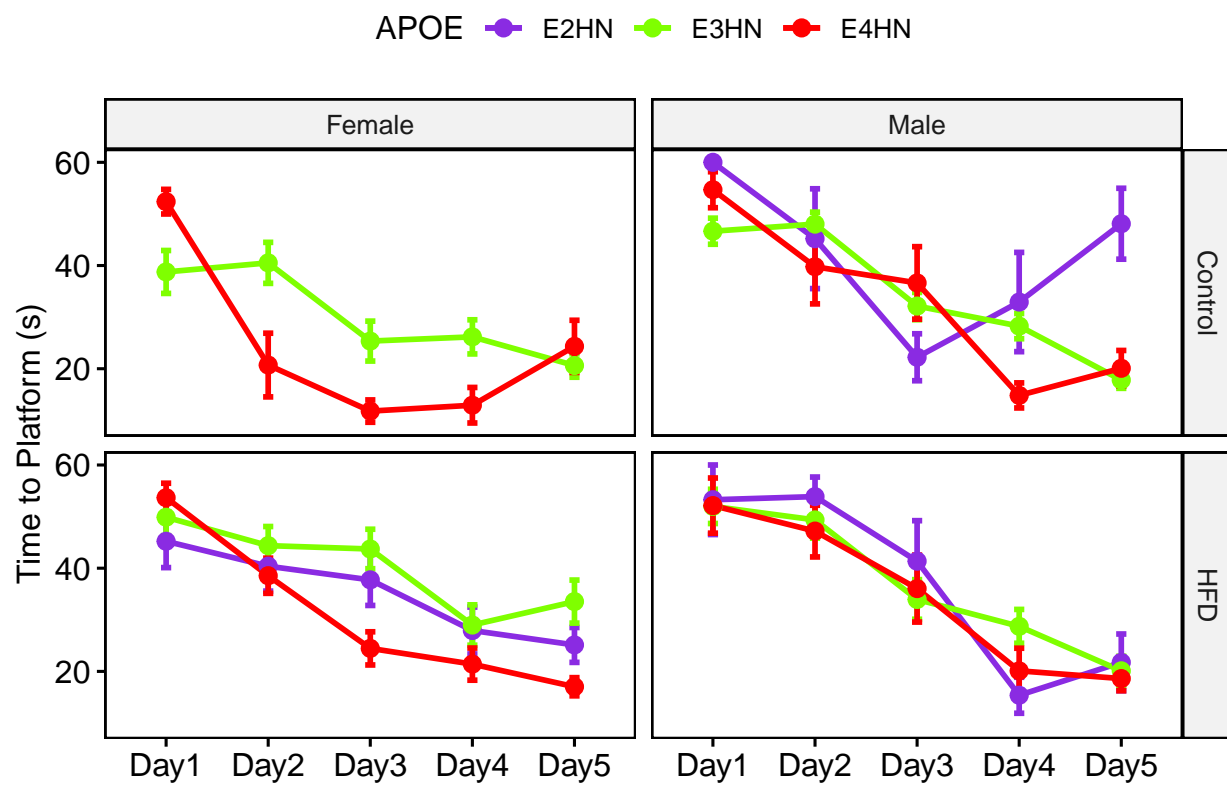


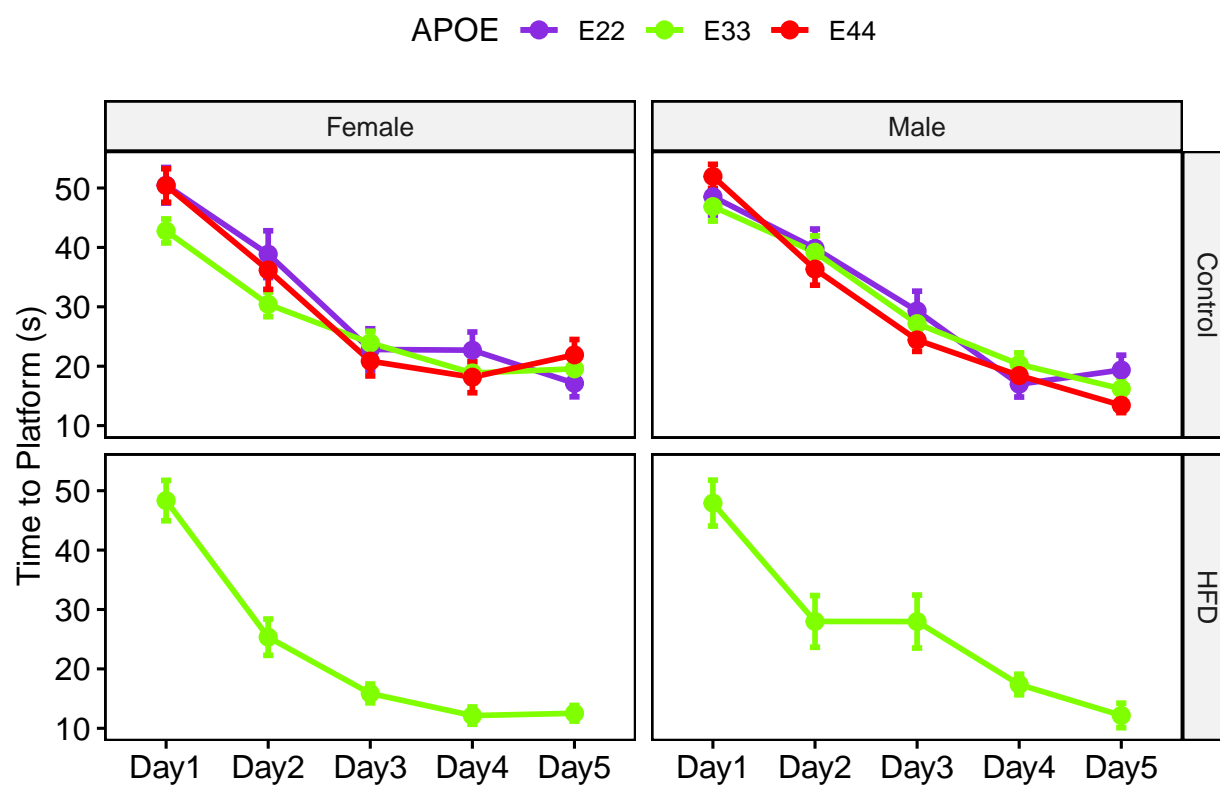


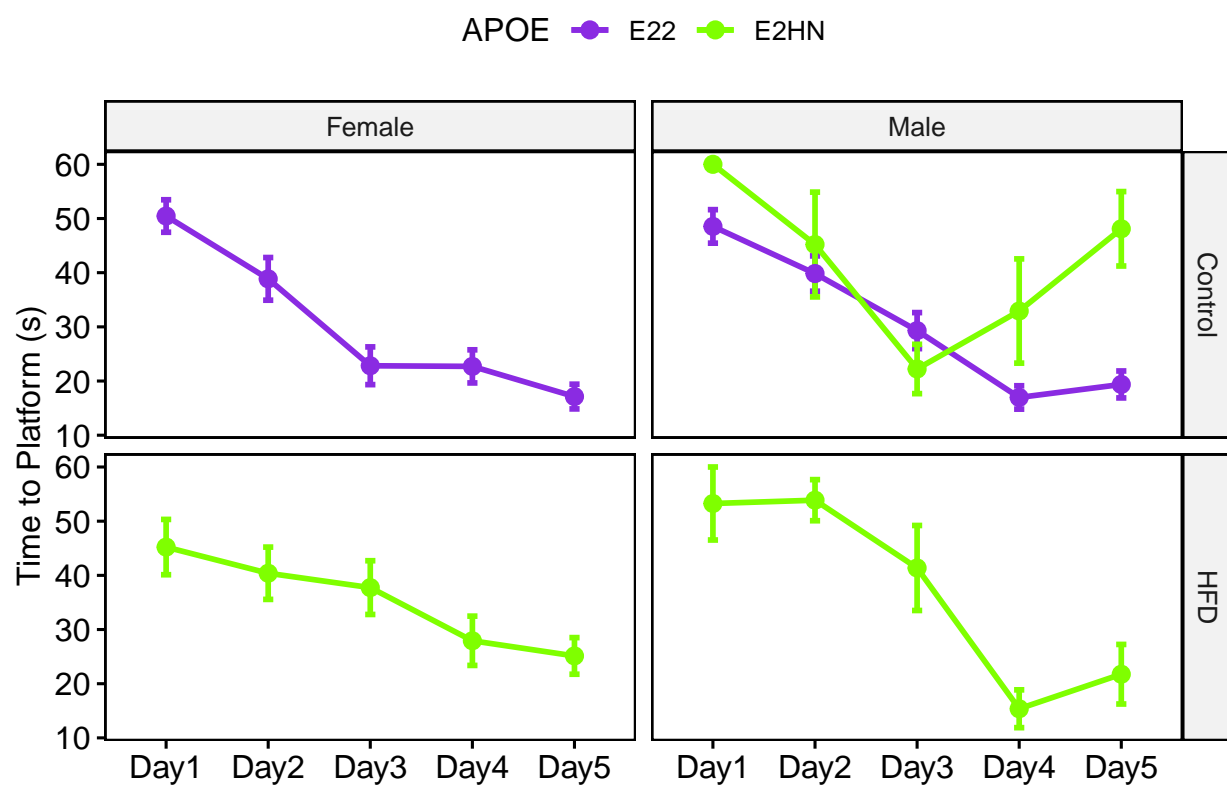


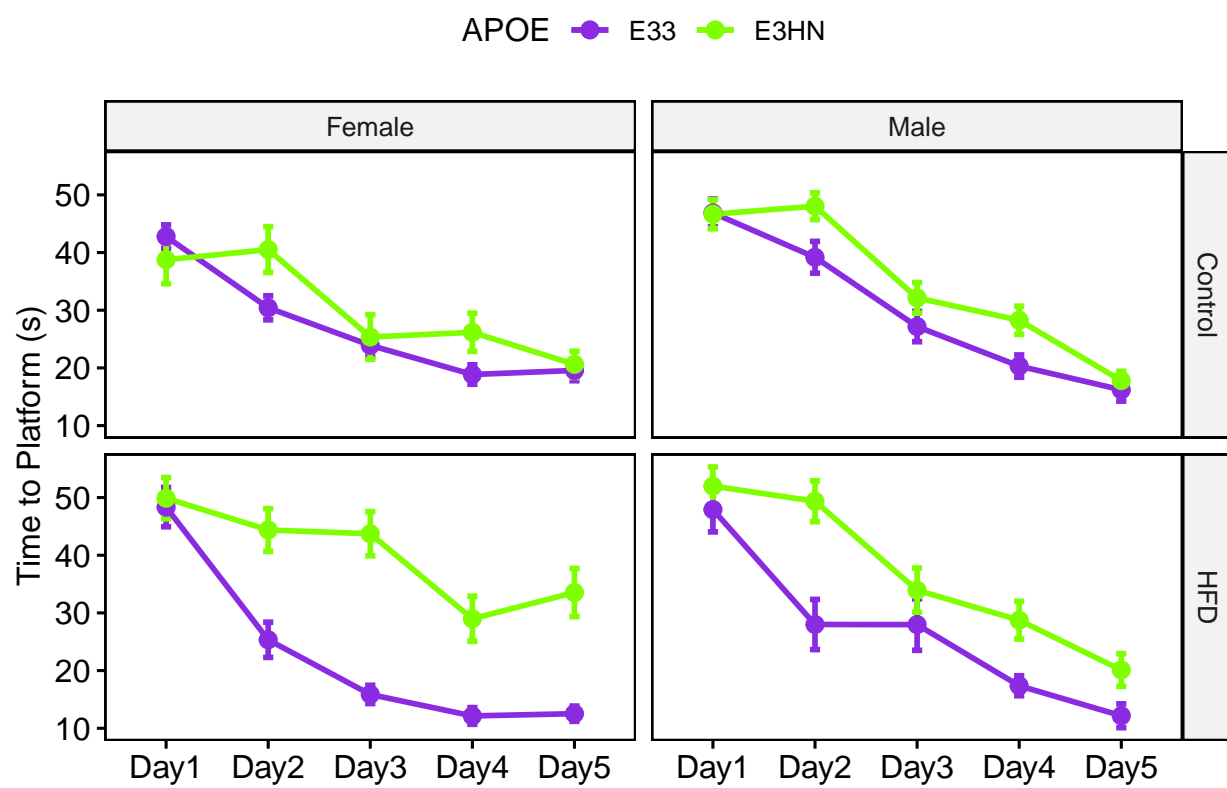


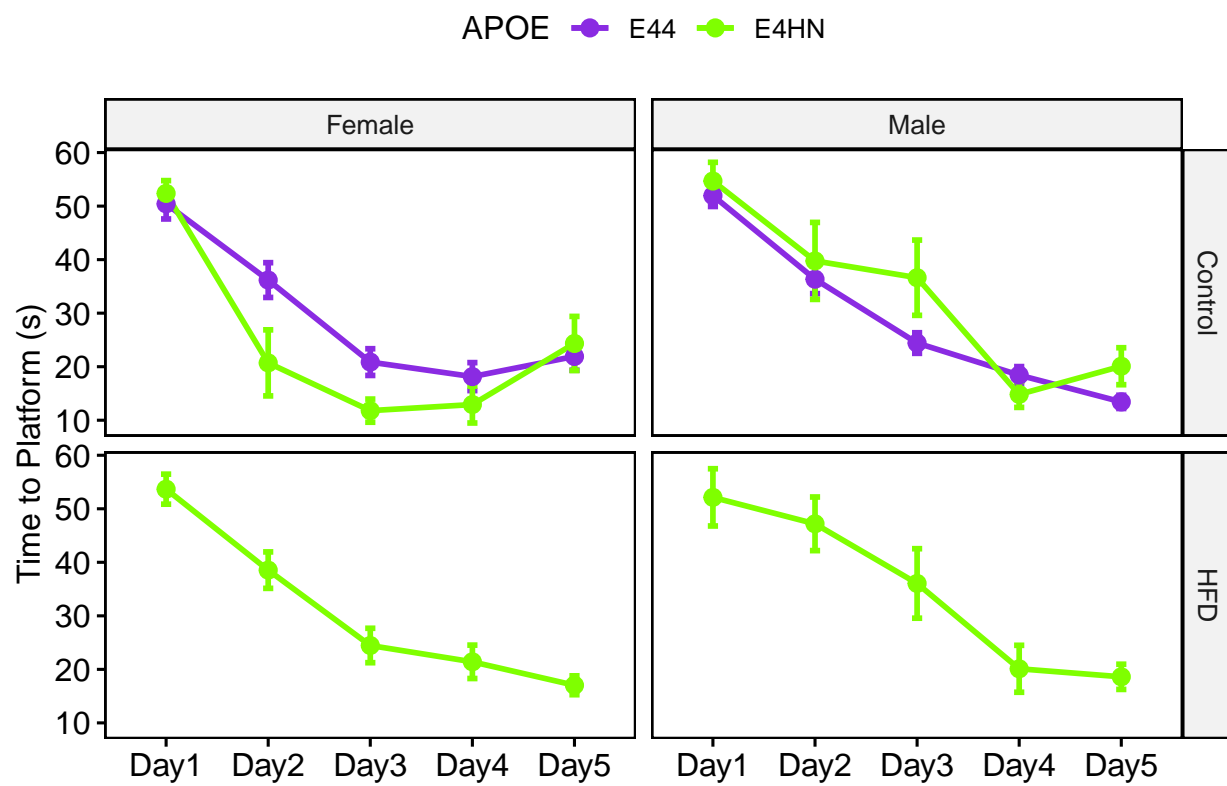






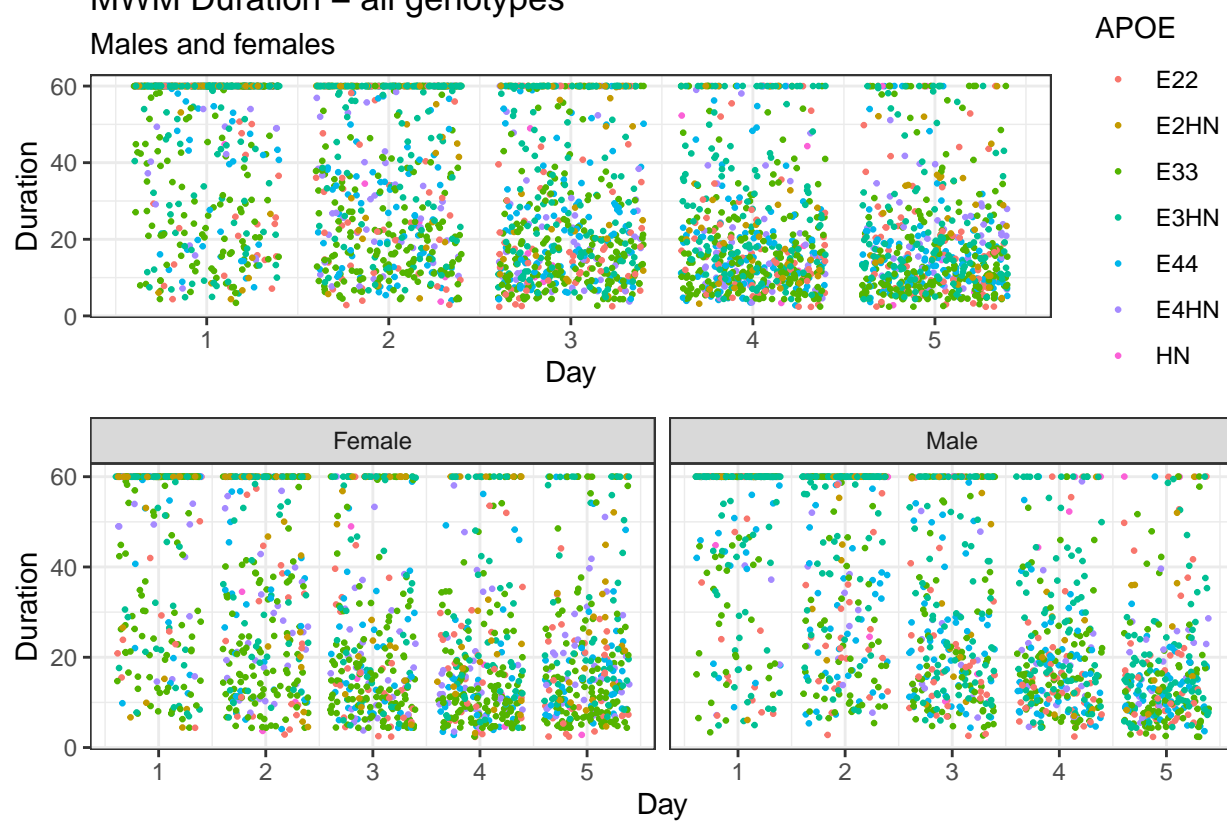






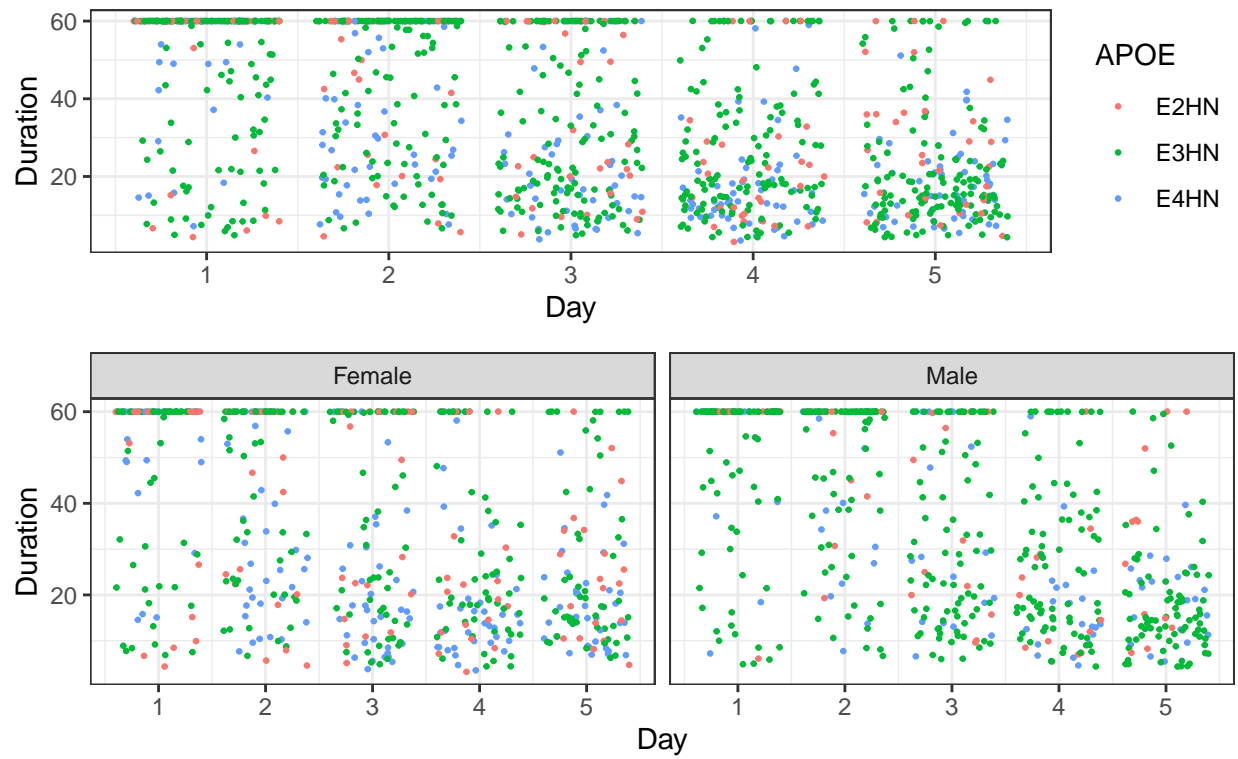
## MWM Duration – all genotypes

Males and females



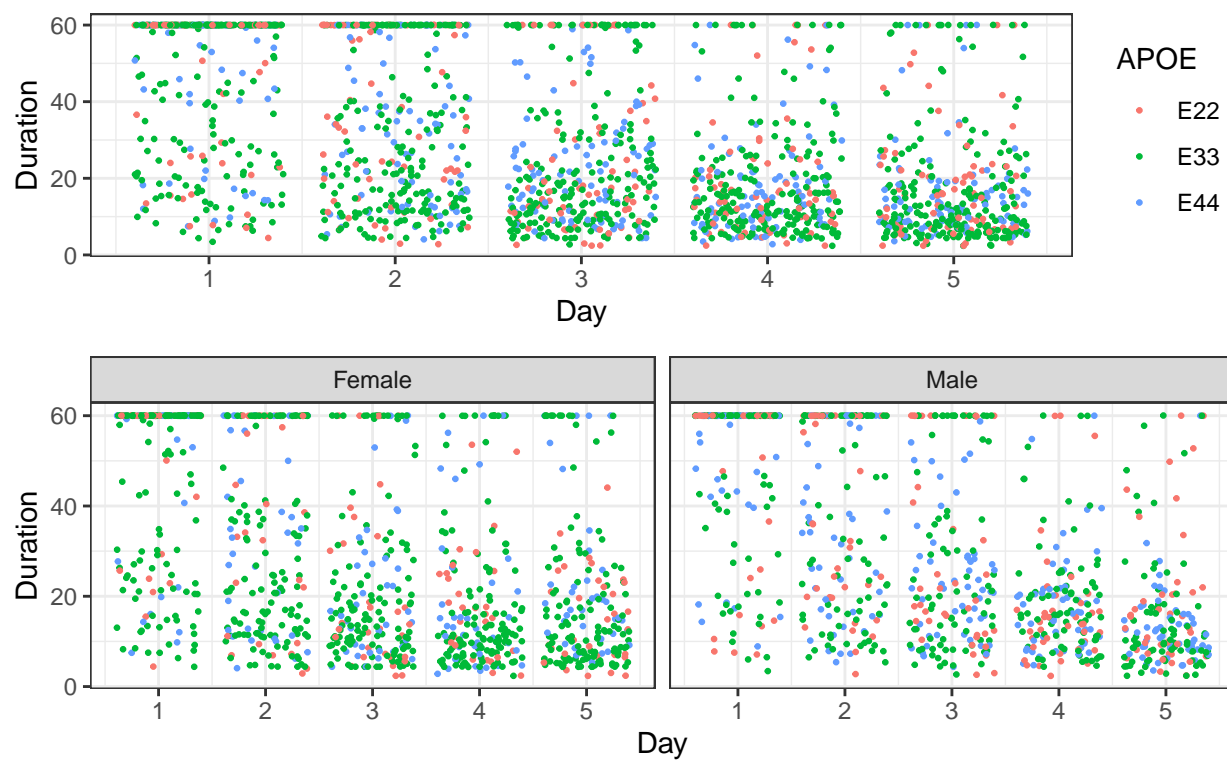
## MWM Duration – HN genotypes

Males and females

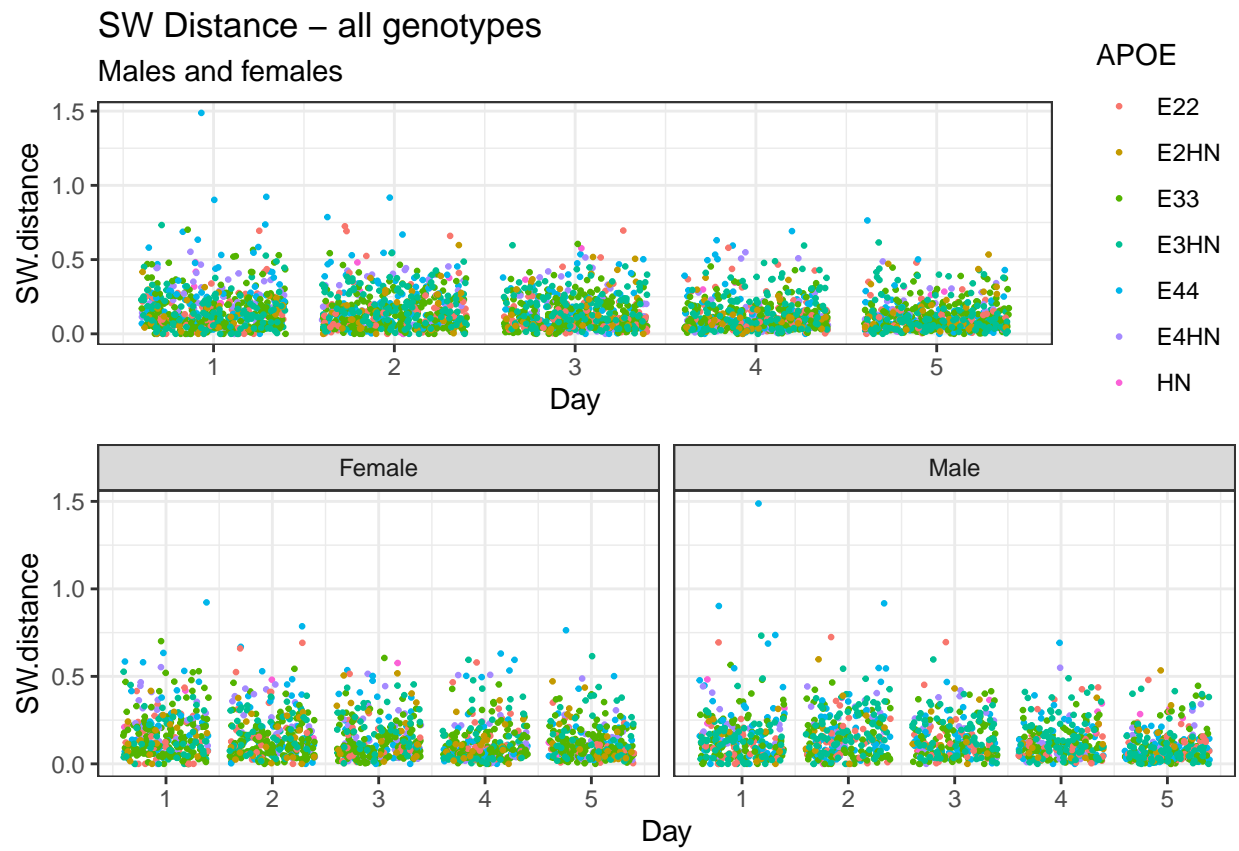


## MWM Duration – non-HN genotypes

Males and females

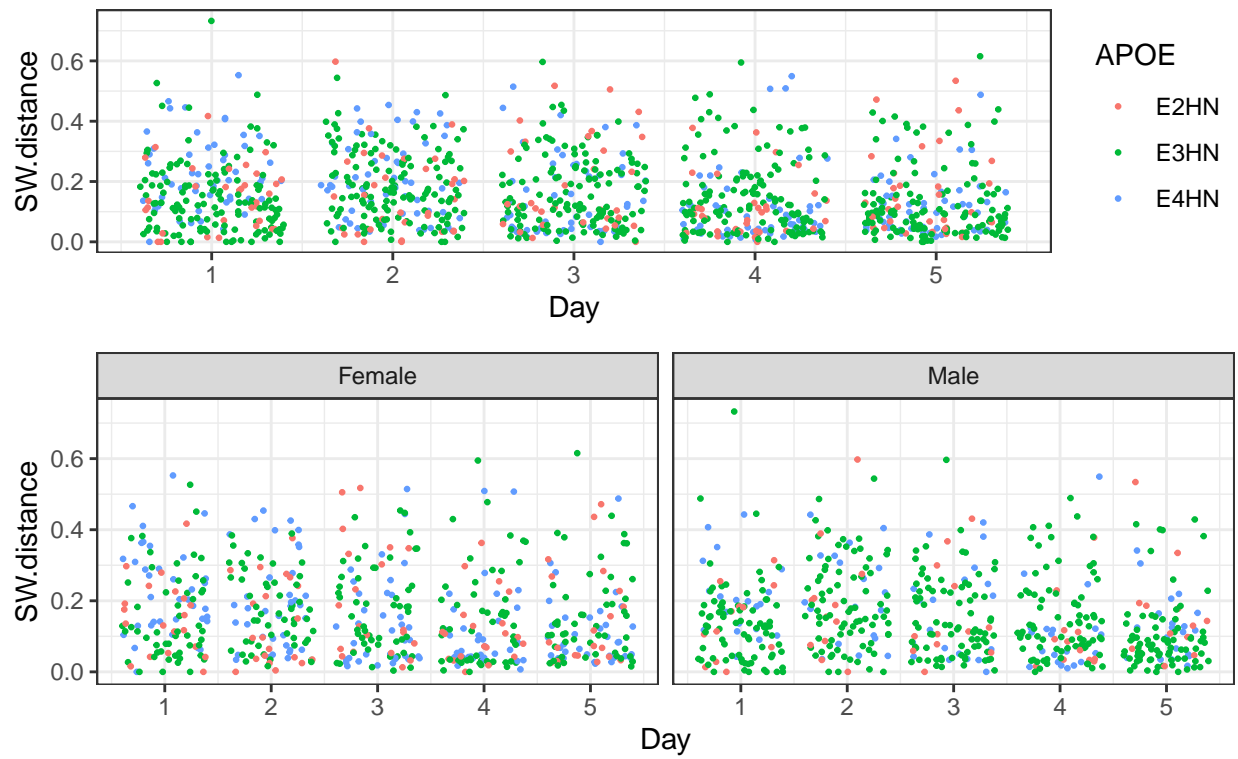






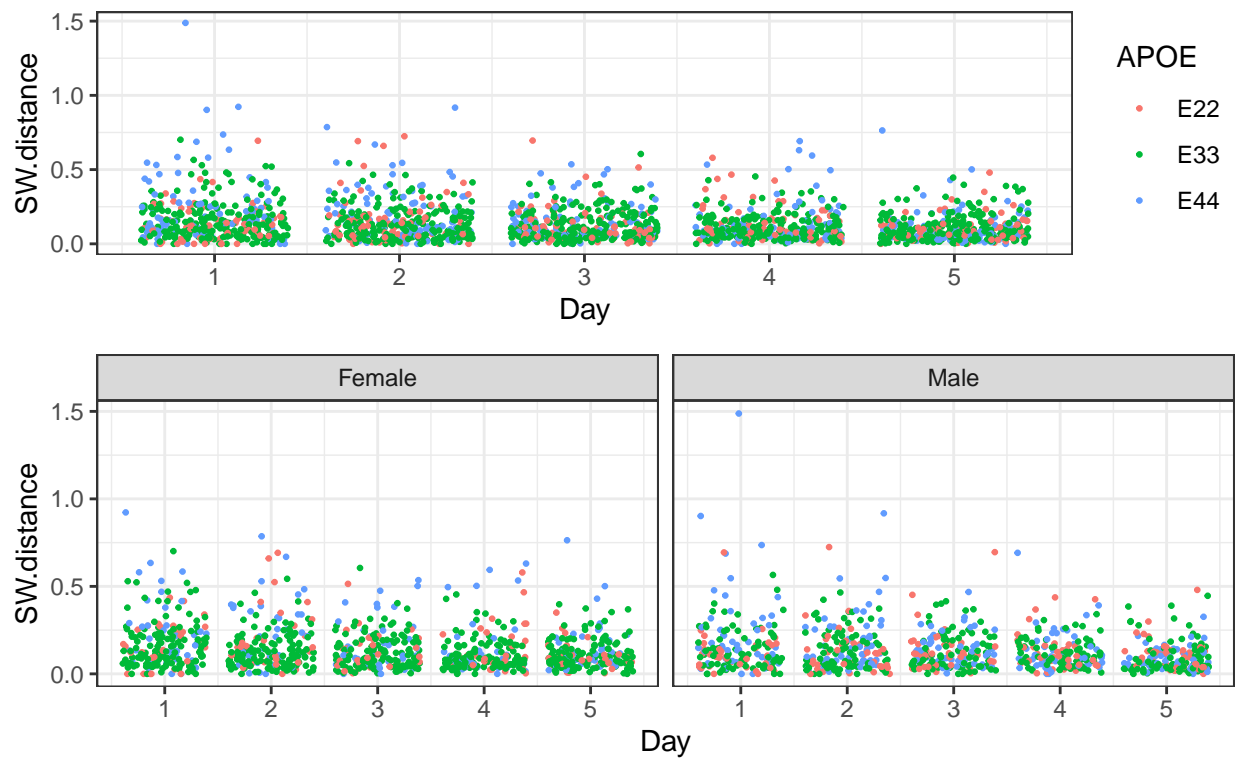
## SW Distance – HN genotypes

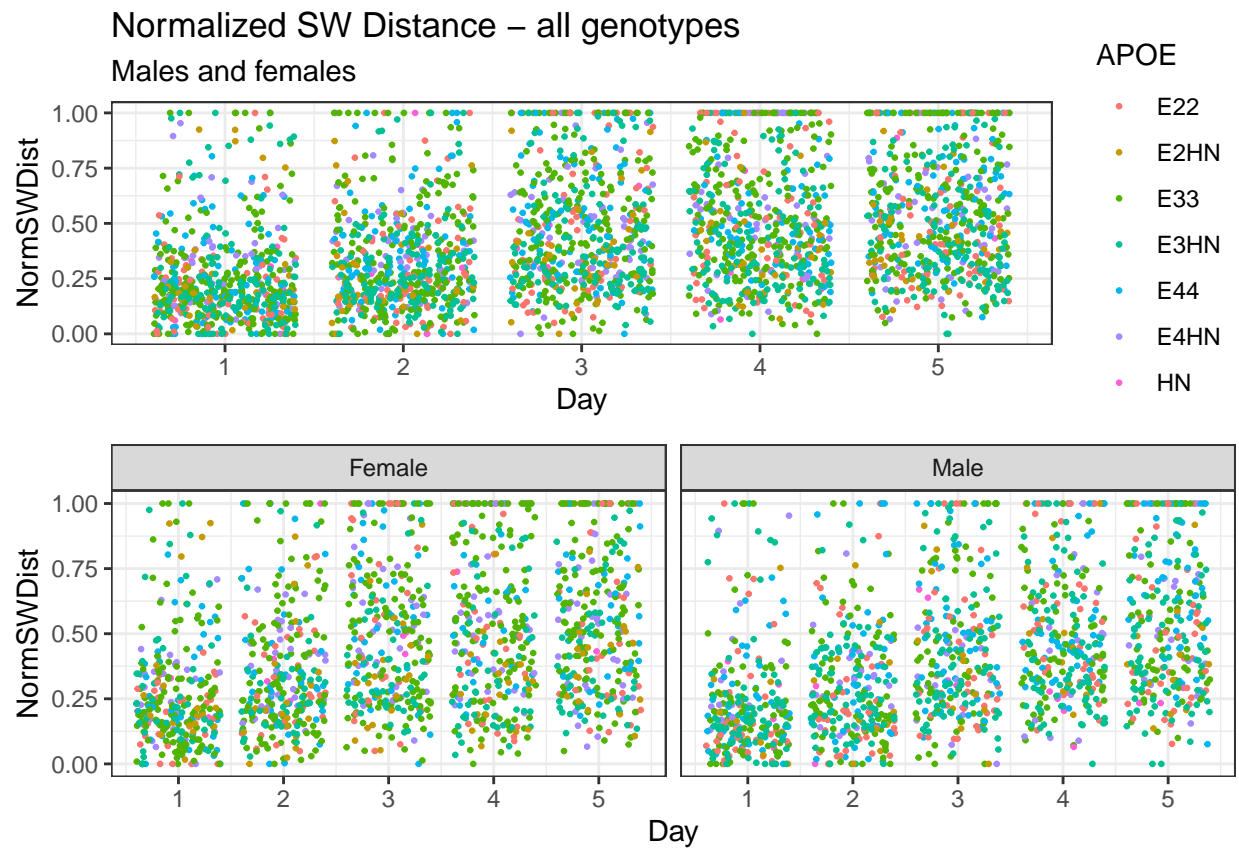
Males and females



## SW Distance – non-HN genotypes

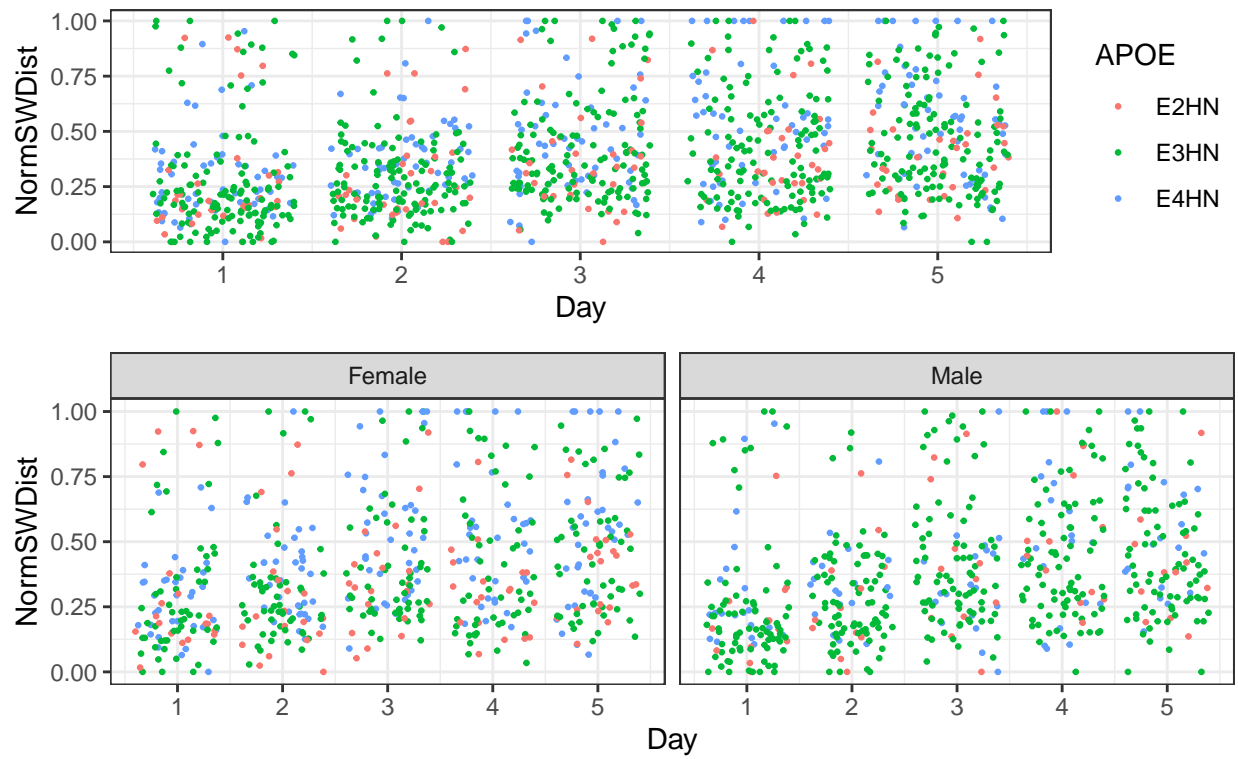
Males and females





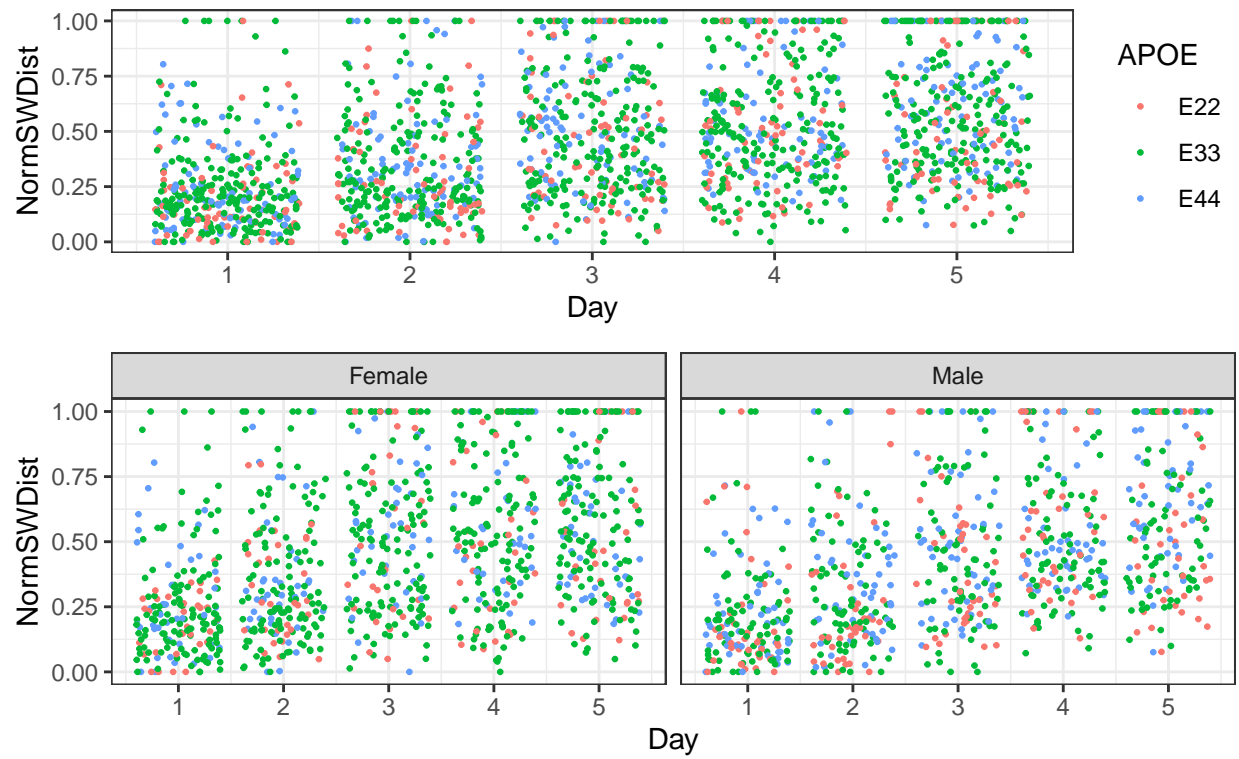
## Normalized SW Distance – HN genotypes

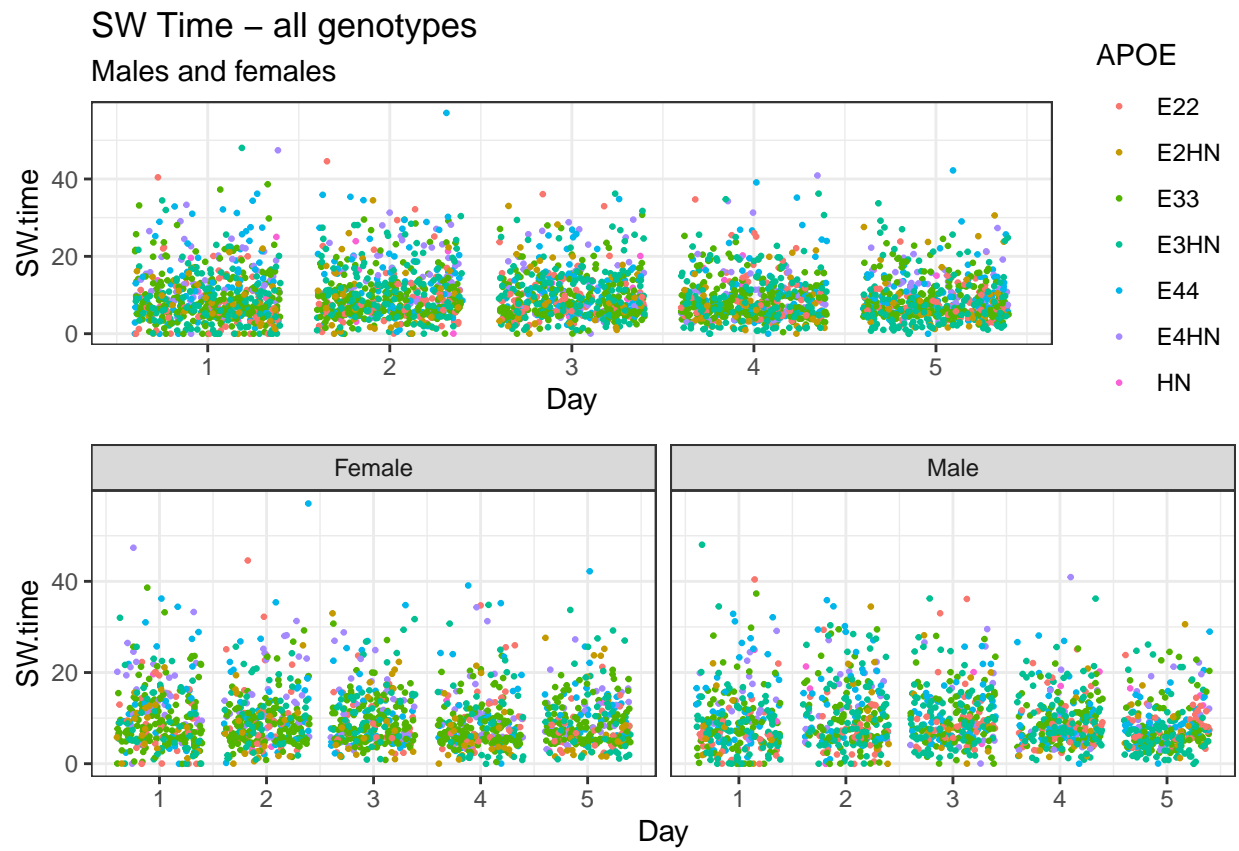
Males and females



## Normalized SW Distance – non-HN genotypes

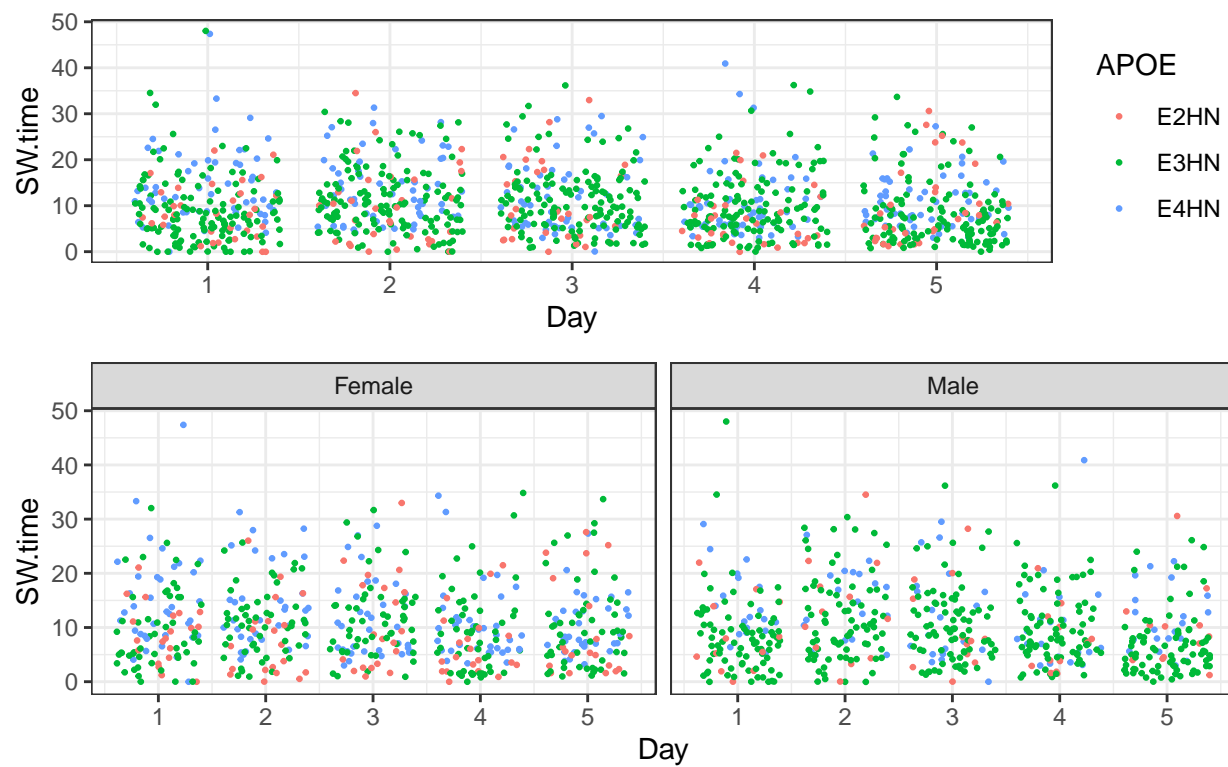
Males and females





## SW Time – HN genotypes

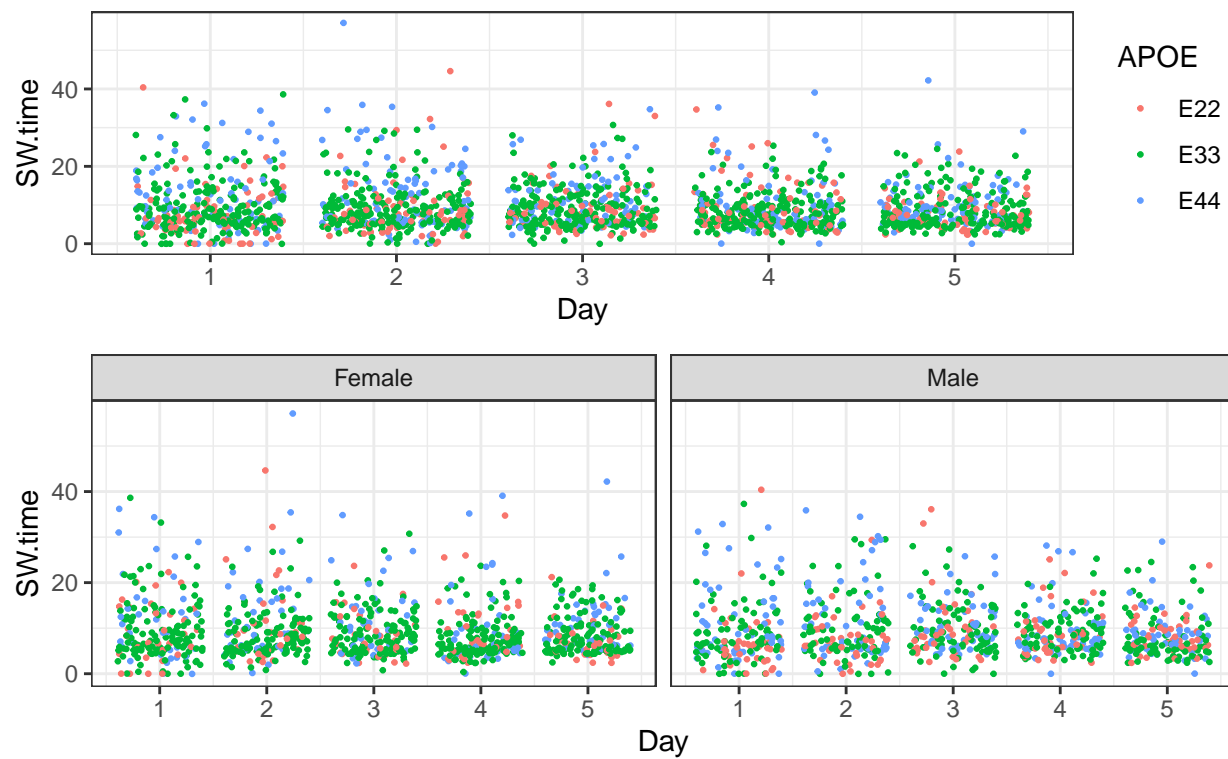
Males and females

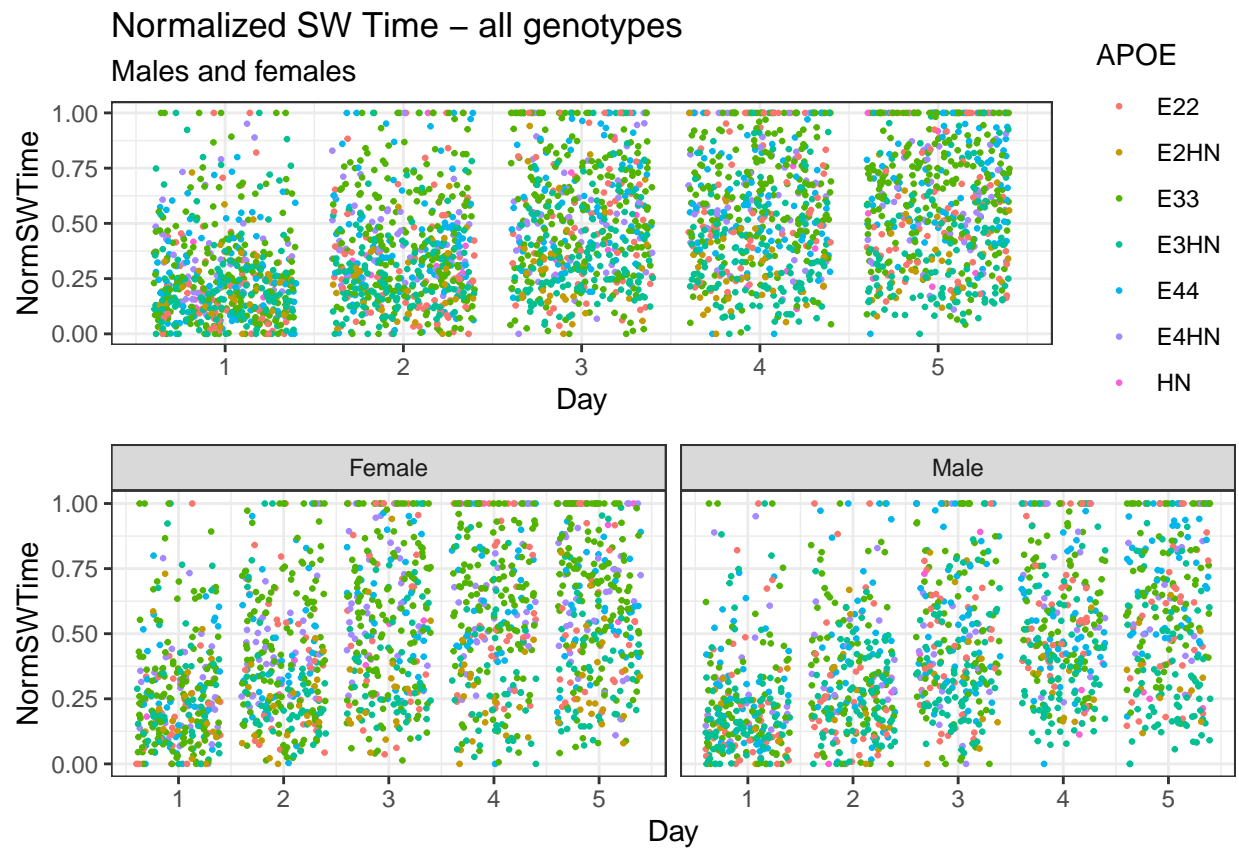




## SW Time – non-HN genotypes

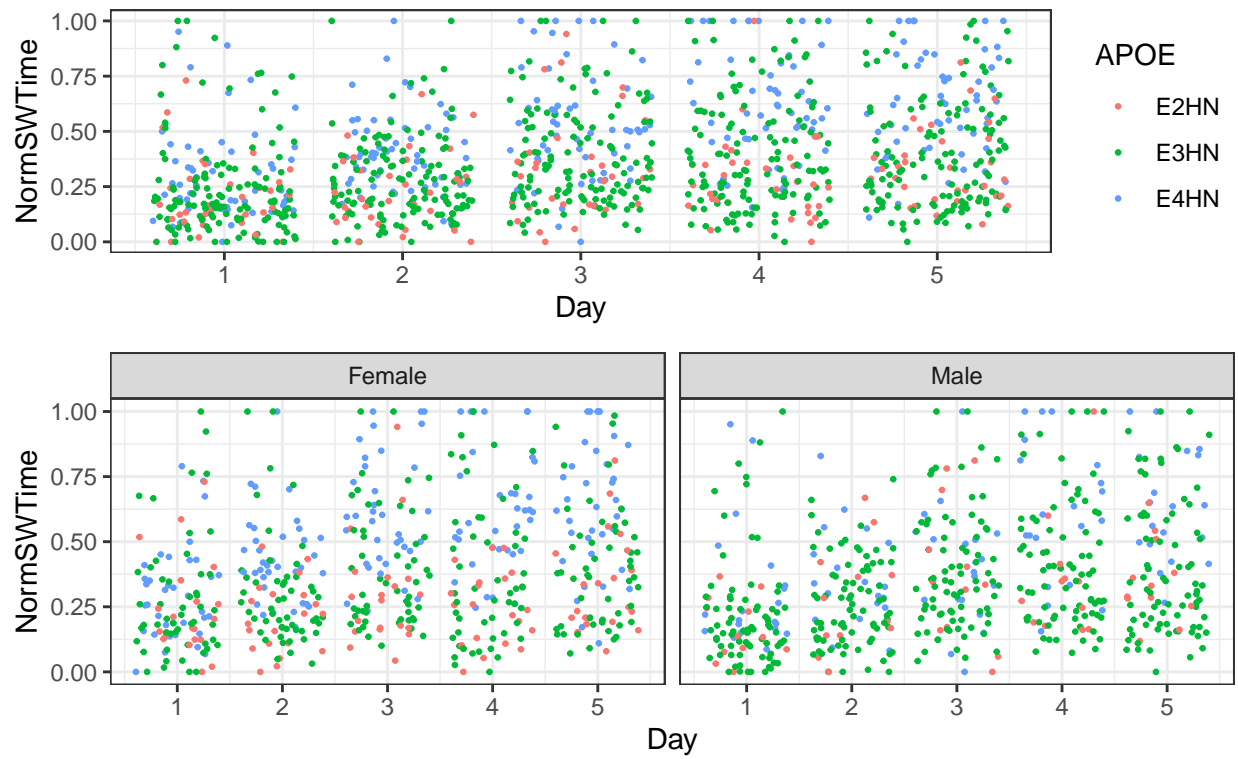
Males and females





## Normalized SW Time – HN genotypes

Males and females



## Normalized SW Time – non-HN genotypes

Males and females

