

# **VR study script**

January 24, 2018

# 1 R setup

The first thing we do is set a random seed for R. This ensures we can reproduce any analysis that has a random component.

```
our_seed <- 123
set.seed(our_seed)
```

We'll use '123' as our seed. Next we need to load the libraries.

```
require(knitr)
require(lme4)

## Loading required package: lme4
## Loading required package: Matrix
## Loading required package: methods

require(afex)

## Loading required package: afex
## Loading required package: emmeans
## *****
## Welcome to afex. For support visit: http://afex.singmann.science/
## - Functions for ANOVAs: aov_car(), aov_ez(), and aov_4()
## - Methods for calculating p-values with mixed(): 'KR', 'S', 'LRT', and
## 'PB'
## - 'afex_aov' and 'mixed' objects can be passed to emmeans() for follow-up
## tests
## - Get and set global package options with: afex_options()
## - Set orthogonal sum-to-zero contrasts globally: set_sum_contrasts()
## - For example analyses see: browseVignettes("afex")
## *****
##
## Attaching package: 'afex'
## The following object is masked from 'package:lme4':
##
## lmer

require(cowplot)

## Loading required package: cowplot
## Loading required package: ggplot2
##
## Attaching package: 'cowplot'
## The following object is masked from 'package:ggplot2':
##
## ggsave
```

```

require(DHARMa)

## Loading required package: DHARMa

require(xtable)

## Loading required package: xtable

require(rockchalk)

## Loading required package: rockchalk

require(tikzDevice)

## Loading required package: tikzDevice

require(parallel)

## Loading required package: parallel

require(fifer)

## Loading required package: fifer
## Loading required package: MASS
##
## Attaching package: 'MASS'
## The following object is masked from 'package:rockchalk':
##
##     mvrnorm

require(multcomp)

## Loading required package: multcomp
## Loading required package: mvtnorm
## Loading required package: survival
## Loading required package: TH.data
##
## Attaching package: 'TH.data'
## The following object is masked from 'package:MASS':
##
##     geyser

require(vcdExtra)

## Loading required package: vcdExtra
## Loading required package: vcd
## Loading required package: grid
## Loading required package: gnm

```

## 2 Reading the files

We now read in the data files.

```
child.data <- read.csv("vr_data/childrenCSV.csv")
carsac.data <- read.csv("vr_data/selfSacCSV.csv")
sidewalk.data <- read.csv("vr_data/sidewalkCSV.csv")
```

Some of the variables need cleaning, so we rename them.

```
# rename gender variable
colnames(child.data)[2] <- "gender"

# rename driver variable
colnames(child.data)[3] <- "motorist"

# rename levels for driver
child.data$motorist <- factor(child.data$motorist,
                             levels = c("False", "True"),
                             labels = c("human", "self-driving"))

# set participant to factor
child.data$participant.ID <- factor(child.data$participant.ID)

# rename levels for gender
child.data$gender <- factor(child.data$gender,
                             levels = c("False", "True"),
                             labels = c("female", "male"))

# remove sanity check and perception check failures
child.sub <- subset(child.data, child.data$passedSanCheck == "True")
child.sub <- subset(child.sub, !(child.sub$perceivedCar == "Mensch" &
                                child.sub$motorist == "self-driving"))

# carsac

# rename gender variable
colnames(carsac.data)[2] <- "gender"

# rename driver variable
colnames(carsac.data)[3] <- "motorist"
```

```

# rename levels for driver
carsac.data$motorist <- factor(carsac.data$motorist,
                              levels = c("False", "True"),
                              labels = c("human", "self-driving"))

carsac.data$participant.ID <- factor(carsac.data$participant.ID)

# rename levels for gender
carsac.data$gender <- factor(carsac.data$gender,
                             levels = c("False", "True"),
                             labels = c("female", "male"))

carsac.sub <- subset(carsac.data, carsac.data$passedSanCheck == "True")
carsac.sub <- subset(carsac.sub, !(carsac.sub$perceivedCar == "Mensch" &
                                   carsac.sub$motorist == "self-driving"))

# sidewalk

# rename gender variable
colnames(sidewalk.data)[2] <- "gender"

# rename driver variable
colnames(sidewalk.data)[3] <- "motorist"

# set participant to factor
sidewalk.data$participant.ID <- factor(sidewalk.data$participant.ID)

# rename levels for driver
sidewalk.data$motorist <- factor(sidewalk.data$motorist,
                                 levels = c("False", "True"),
                                 labels = c("human", "self-driving"))

# rename levels for gender
sidewalk.data$gender <- factor(sidewalk.data$gender,
                               levels = c("False", "True"),
                               labels = c("female", "male"))

# sanity check
sidewalk.sub <- subset(sidewalk.data, sidewalk.data$passedSanCheck == "True")
sidewalk.sub <- subset(sidewalk.sub, !(sidewalk.sub$perceivedCar == "Mensch" &

```

```
sidewalk.sub$motorist == "self-driving"))
```

### 3 Manipulation check

We use a  $\chi^2$  goodness of fit test to see if perspective affects identification. First we create a subset of only one trial, and collapse the two pedestrian perspectives.

```
manip_check.sub <- subset(child.sub, trial == "largeGroups")

manip_check.sub$perspective <- combineLevels(manip_check.sub$perspective,
                                              levs = c("PedLarge", "PedSmall"),
                                              newLabel = c("Pedestrian"))

## The original levels Observer Passenger PedLarge PedSmall
## have been replaced by Observer Passenger Pedestrian
```

Next we create a contingency table.

```
manip_check.xtab <- xtabs(~perspective + perceivedIden, data = manip_check.sub)
print(xtable(manip_check.xtab))
```

	Beobachter	Fußgänger	Mitfahrer
Observer	30	5	10
Passenger	7	3	34
Pedestrian	33	57	3

Then we compute the test statistic.

```
manip_check.chisq <- chisq.test(manip_check.xtab, simulate.p.value = TRUE)
manip_check.chisq

##
## Pearson's Chi-squared test with simulated p-value (based on 2000
## replicates)
##
## data:  manip_check.xtab
## X-squared = 114.01, df = NA, p-value = 0.0004998
```

If the test statistic is significant, we perform follow-up comparisons.

```
alpha <- 0.05
if (manip_check.chisq$p.value <= alpha){
  manip_check.post <- chisq.post.hoc(manip_check.xtab)
}
```

Adjusted p-values used the fdr method.

```
print(xtable(manip_check.post))
```

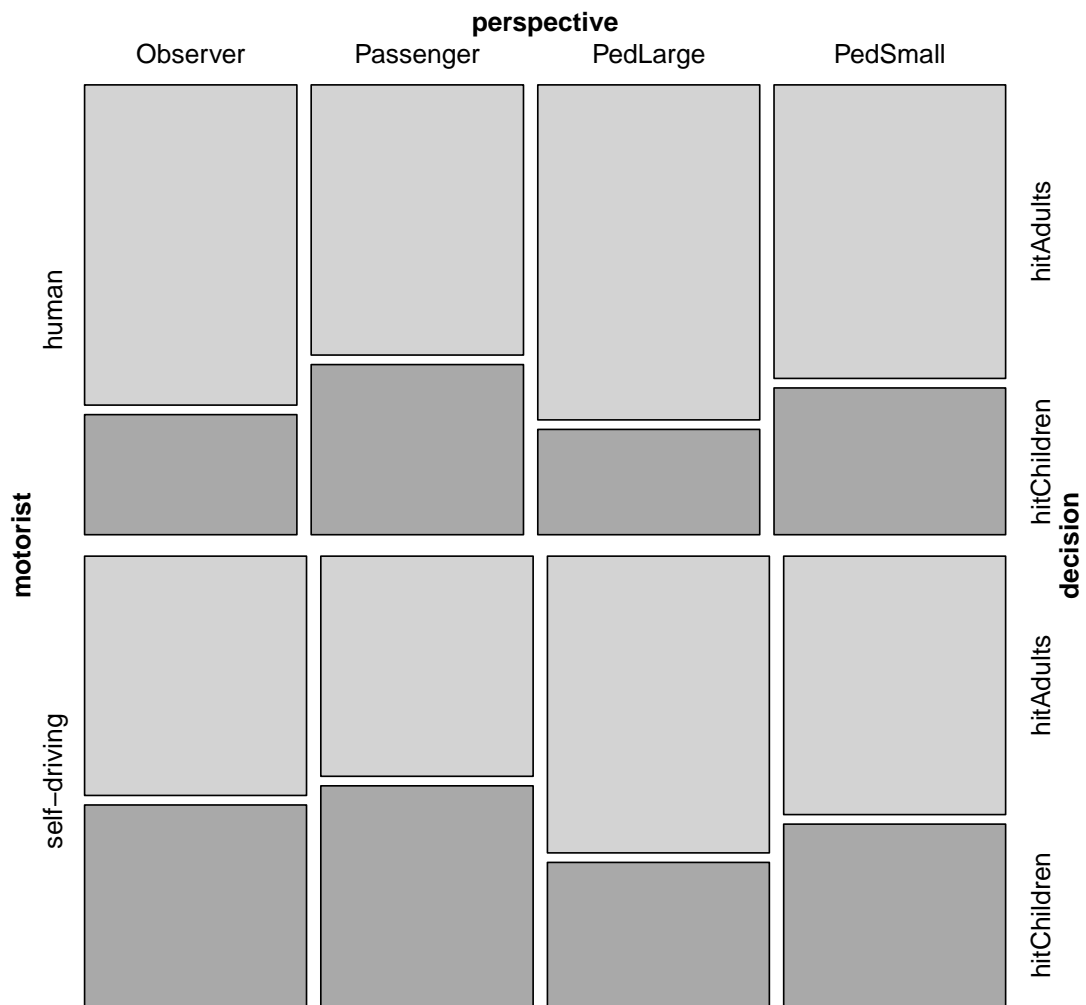
	comparison	raw.p	adj.p
1	Observer vs. Passenger	0.00	0.00
2	Observer vs. Pedestrian	0.00	0.00
3	Passenger vs. Pedestrian	0.00	0.00

## 4 Child vs adult dilemmas

### 4.1 Visualising the raw responses

First we look at the raw proportions of the responses.

```
child.xtabs <- xtabs(~ motorist + perspective + decision, data = child.sub)
child.mosaic <- mosaic(child.xtabs,
  gp = gpar(fill = c("light gray", "dark gray")))
```



## 4.2 The model

We run a logit mixed model to predict the decision of the participants.

```
(nc <- detectCores())
```

```
[1] 12
```

```
c1 <- makeCluster(rep("localhost", nc))
```

```
child_glmm <- mixed(decision ~ perspective + motorist +
  perspective:motorist +
```



```

trial + gender + #age +
(1 | participant.ID),
method = "PB", # change to PB for final
family = "binomial", data = child.sub,
args_test = list(nsim = 1000, cl = cl), cl = cl,
control = glmerControl(optimizer = "bobyqa",
                        optCtrl = list(maxfun = 2e5)))

```

*## Contrasts set to contr.sum for the following variables: decision, perspective, motorist, trial, gender, participant.ID*

Fitting 6 (g)lmer() models. Obtaining 5 p-values: [.....]

```

stopCluster(cl)
child_glmm_anova <- child_glmm$anova_table
print(xtable(child_glmm_anova))

```

	Chisq	Chi Df	Pr(>Chisq)	Pr(>PB)
perspective	3.90	3	0.2724	0.3986
motorist	4.23	1	0.0398	0.0675
trial	1.21	1	0.2713	0.2714
gender	0.30	1	0.5820	0.6128
perspective:motorist	0.45	3	0.9291	0.9328

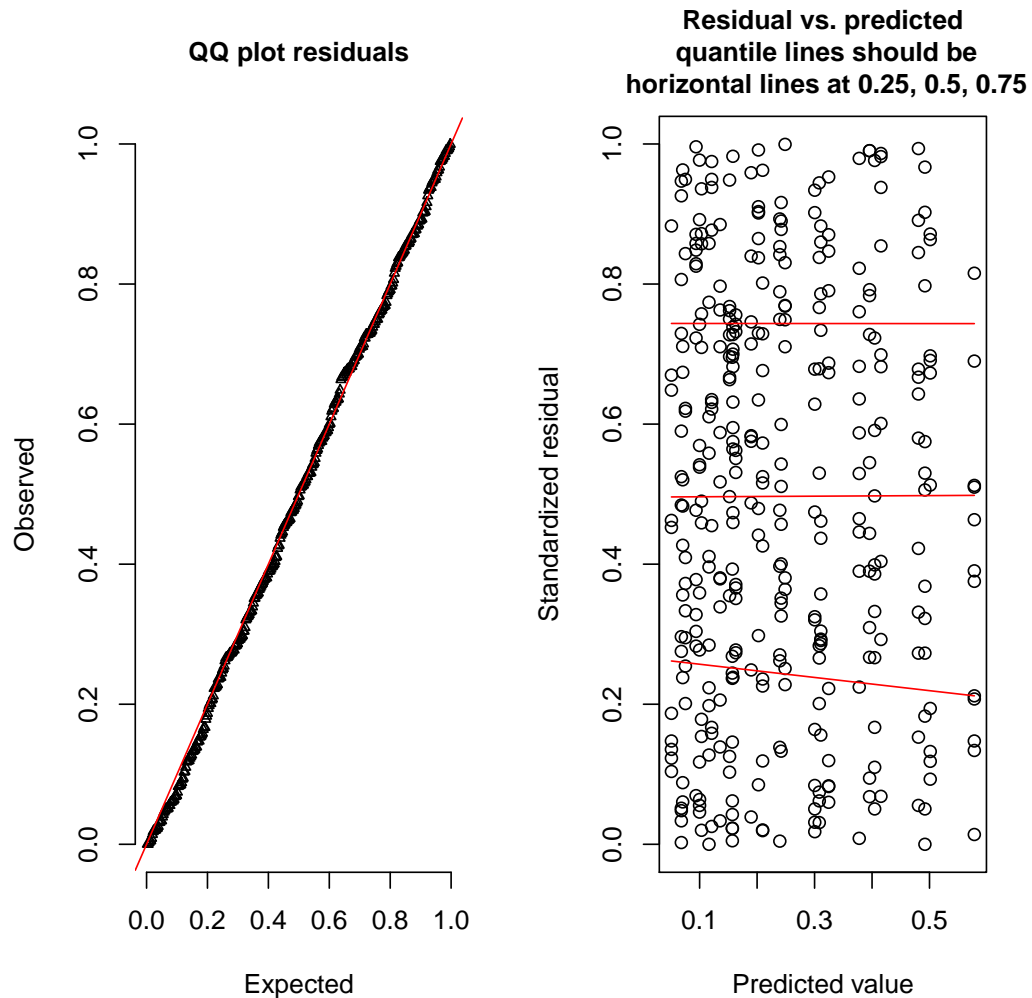
We then check the residuals of the model.

```

child_glmm.resid <- simulateResiduals(
  fittedModel = child_glmm$full_model,
  n = 2000)
child_glmm_resid.plot <- plotSimulatedResiduals(
  simulationOutput = child_glmm.resid)

```

## DHARMA scaled residual plots



If they look okay, we follow up any significant differences.

### 4.3 Plotting

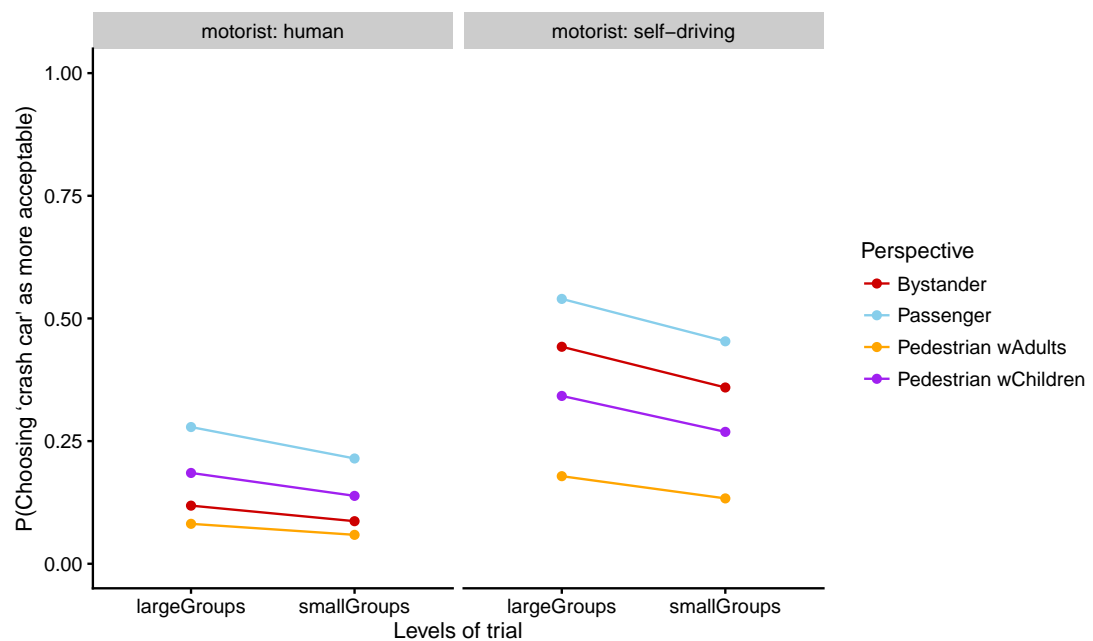
Depending on which effects or interactions we find, we probably want to change this plot. It currently just plots the Estimated Marginal Means for the 2-way interaction.

```
child_emm_plot <- emmip(child_glmm, perspective ~ trial | motorist, type = "response") +
  theme_cowplot(font_size = 10) +
  scale_y_continuous(name = "P(Choosing `crash car' as more acceptable)",
    limits = c(0, 1)) + coord_equal(ratio = 3) +
  scale_color_manual(name = "Perspective",
    labels = c("Bystander",
```

```

        "Passenger",
        "Pedestrian wAdults",
        "Pedestrian wChildren"),
    values = c("red3", "skyblue", "orange1", "purple"))
child_emm_plot

```



#### 4.4 Follow-up comparisons

For a good example, see <https://osf.io/q2a3b/>

If we find an interaction between perspective and motorist-type we should follow that up with comparisons of perspective within each level of motorist.

```
emm_child_i <- emmeans(child_glmm, ~ perspective | motorist, type = "response",
                        contr = "pairwise")
xtable(emm_child_i$contrasts)
```

contrast	odds.ratio	SE	df	z.ratio	p.value
motorist = human					
Observer / Passenger	0.3479	0.4008	Inf	-0.917	0.7960
Observer / PedLarge	1.5163	1.7108	Inf	0.369	0.9829
Observer / PedSmall	0.5917	0.6627	Inf	-0.469	0.9659
Passenger / PedLarge	4.3590	5.0289	Inf	1.276	0.5782
Passenger / PedSmall	1.7009	1.9212	Inf	0.470	0.9656
PedLarge / PedSmall	0.3902	0.4354	Inf	-0.843	0.8337
motorist = self-driving					
Observer / Passenger	0.6760	0.7553	Inf	-0.350	0.9852
Observer / PedLarge	3.6493	4.0984	Inf	1.153	0.6568
Observer / PedSmall	1.5251	1.6707	Inf	0.385	0.9806
Passenger / PedLarge	5.3985	6.3213	Inf	1.440	0.4743
Passenger / PedSmall	2.2562	2.5623	Inf	0.716	0.8906
PedLarge / PedSmall	0.4179	0.4686	Inf	-0.778	0.8644

Results are averaged over the levels of: trial, gender

P value adjustment: tukey method for comparing a family of 4 estimates

Tests are performed on the log odds ratio scale

If we find no interaction but a significant main effect of perspective, we need to do follow-up multiple comparisons to find out where the difference is.

```
emm_child_persp <- emmeans(child_glmm, pairwise ~ perspective,
                           type = 'response')

## NOTE: Results may be misleading due to involvement in interactions

xtable(emm_child_persp$contrasts)
```

If we find a main effect of motorist-type, we need to calculate the marginal means to see the effect.

```
emm_child_motorist <- emmeans(child_glmm, pairwise ~ motorist,
                              type = 'response')

## NOTE: Results may be misleading due to involvement in interactions

xtable(emm_child_motorist$contrasts)
```

contrast	odds.ratio	SE	df	z.ratio	p.value
Observer / Passenger	0.4849	0.3907	Inf	-0.898	0.8057
Observer / PedLarge	2.3523	1.8836	Inf	1.068	0.7089
Observer / PedSmall	0.9499	0.7427	Inf	-0.066	0.9999
Passenger / PedLarge	4.8510	4.0801	Inf	1.878	0.2377
Passenger / PedSmall	1.9590	1.5768	Inf	0.835	0.8376
PedLarge / PedSmall	0.4038	0.3218	Inf	-1.138	0.6661

Results are averaged over the levels of: motorist, trial, gender

P value adjustment: tukey method for comparing a family of 4 estimates

Tests are performed on the log odds ratio scale

contrast	odds.ratio	SE	df	z.ratio	p.value
human / self-driving	0.3158	0.1869	Inf	-1.948	0.0514

Results are averaged over the levels of: perspective, trial, gender

Tests are performed on the log odds ratio scale

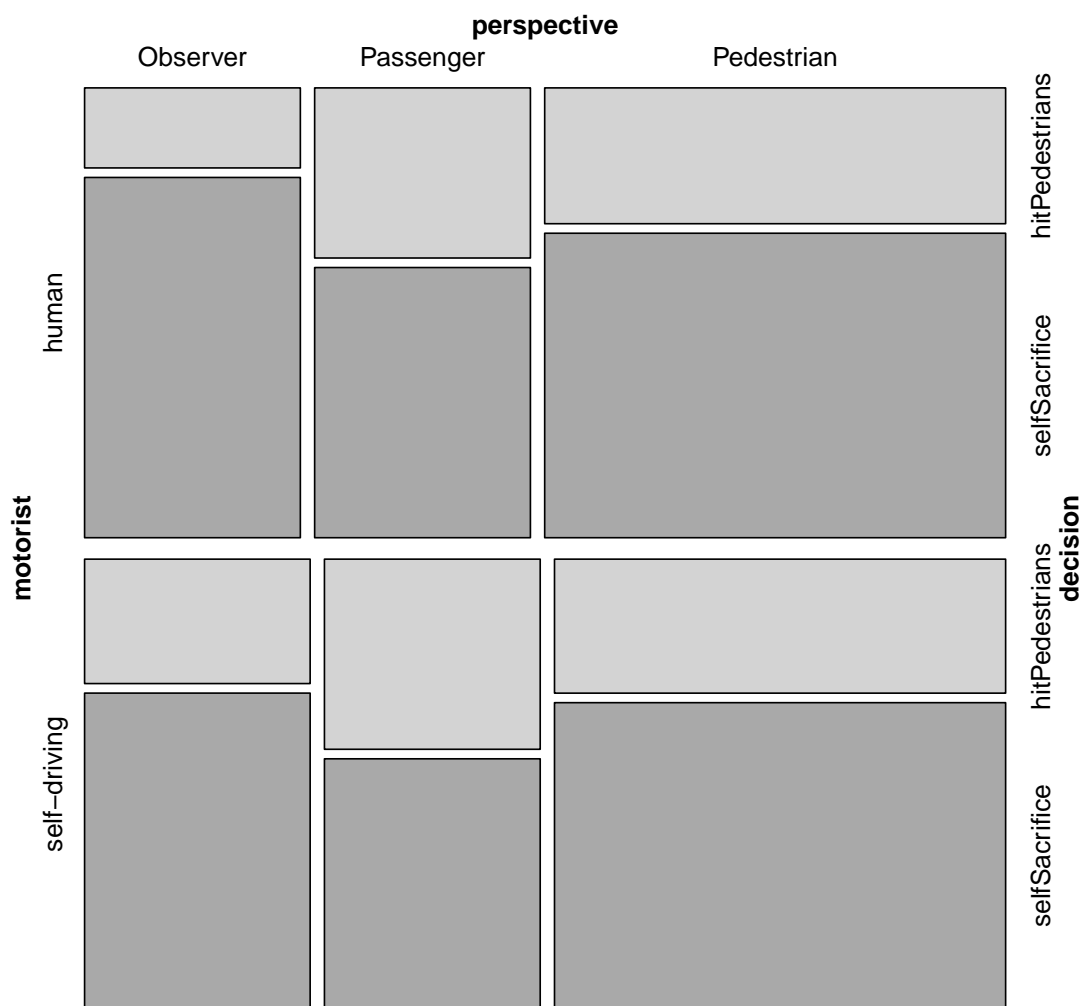
## 5 Car occupants vs pedestrians dilemma

We collapse both pedestrian perspectives for this dilemma.

```
carsac.sub$perspective <- combineLevels(carsac.sub$perspective,
                                       levs=c("PedLarge", "PedSmall"),
                                       newLabel = "Pedestrian")
```

```
## The original levels Observer Passenger PedLarge PedSmall
## have been replaced by Observer Passenger Pedestrian
```

```
carsac.xtabs <- xtabs(~ motorist + perspective + decision, data = carsac.sub)
carsac.mosaic <- mosaic(carsac.xtabs,
                       gp = gpar(fill = c("light gray", "dark gray")))
```



## 5.1 The model

```
(nc <- detectCores())
```

```
[1] 12
```

```
cl <- makeCluster(rep("localhost", nc))
```

```
carsac_glmm <- mixed(decision ~ perspective + motorist +  
                      perspective:motorist +  
                      trial + gender + #age +
```

```

      (1 | participant.ID),
      method = "PB", # change to PB for final
      family = "binomial", data = carsac.sub,
      args_test = list(nsim = 1000, cl = cl), cl = cl,
      control = glmerControl(optimizer = "bobyqa",
                             optCtrl = list(maxfun = 2e5)))

```

*## Contrasts set to contr.sum for the following variables: decision, perspective, motorist, trial, gender, participant.ID*

Fitting 6 (g)lmer() models. Obtaining 5 p-values: [.....]

```

stopCluster(cl)
xtable(carsac_glmm$anova_table)

```

	Chisq	Chi Df	Pr(>Chisq)	Pr(>PB)
perspective	7.50	2	0.0235	0.0311
motorist	1.08	1	0.2994	0.3102
trial	61.24	1	0.0000	0.0010
gender	0.13	1	0.7228	0.7472
perspective:motorist	1.05	2	0.5903	0.6148

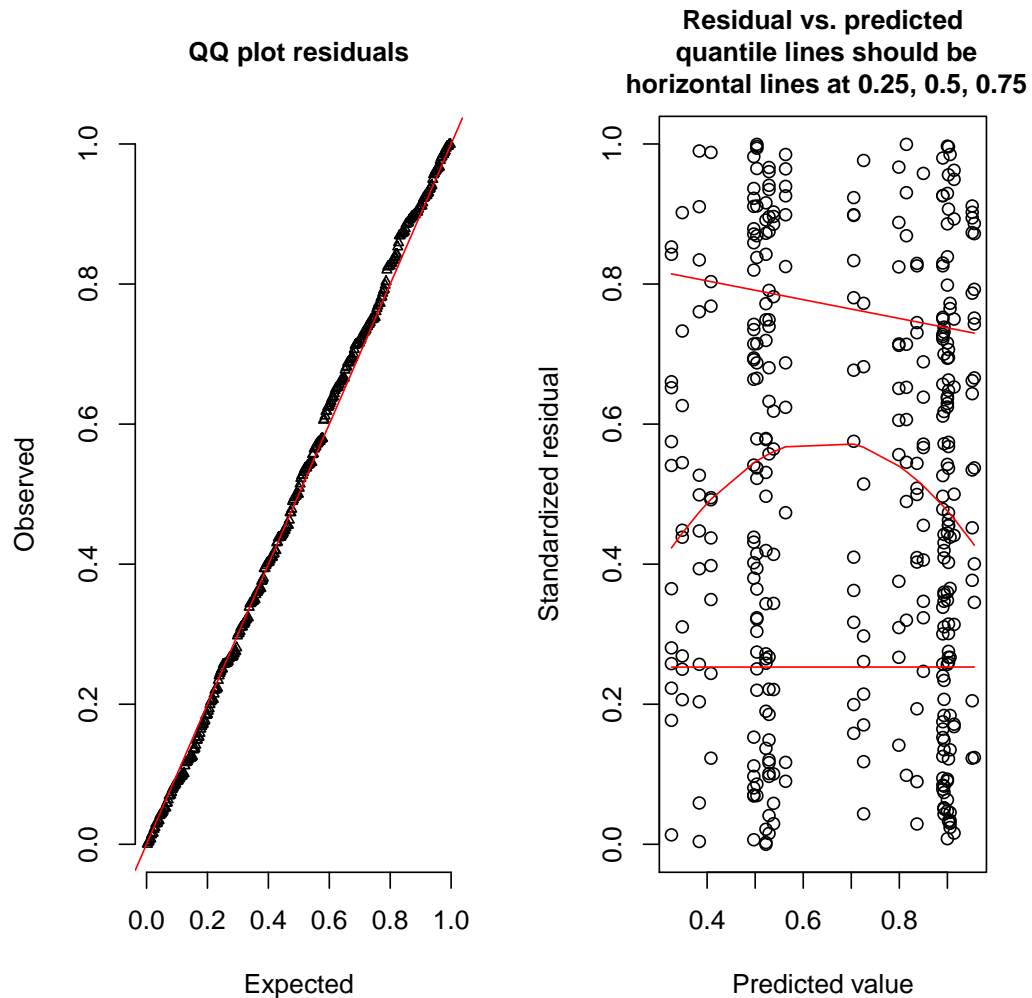
Then we check the residuals.

```

carsac_glmm.resid <- simulateResiduals(
  fittedModel = carsac_glmm$full_model, n = 2000)
carsac_glmm_resid.plot <- plotSimulatedResiduals(
  simulationOutput = carsac_glmm.resid)

```

## DHARMA scaled residual plots



## 5.2 Plotting

Depending on which effects or interactions we find, we probably want to change this plot. It currently just plots the Estimated Marginal Means for the 2-way interaction.

```
carsac.plot <- emmip(carsac_glmm, perspective ~ trial | motorist, type = "response") +
  theme_cowplot(font_size = 10) +
  scale_y_continuous(name = "P(Choosing `crash car' as more acceptable)",
    limits = c(0, 1)) + coord_equal(ratio = 3) +
  scale_color_manual(name = "Perspective",
    labels = c("Bystander",
      "Passenger",
```

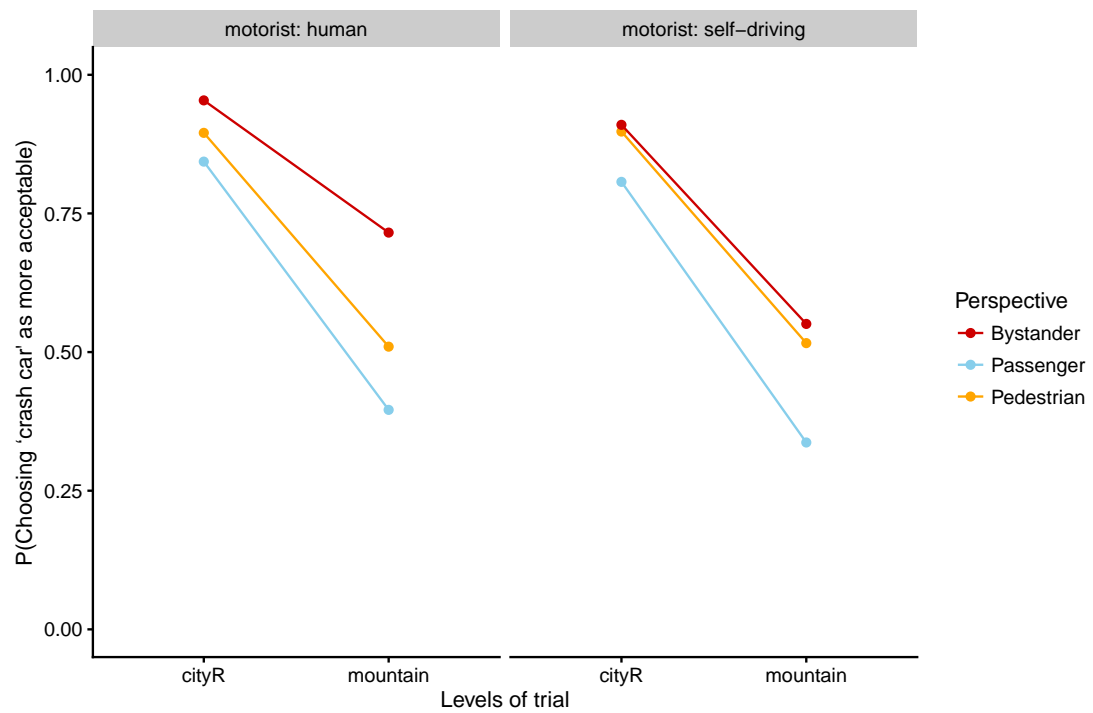


```

        "Pedestrian"),
    values = c("red3", "skyblue", "orange1" , "purple"))

carsac.plot

```



### 5.3 Follow-up analyses

If we find an interaction between perspective and motorist-type we should follow that up with comparisons of perspective within each level of motorist.

```
emm_carsac_i <- emmeans(carsac_glmm, pairwise ~ perspective | motorist,
                        type = 'response')
xtable(emm_carsac_i$contrasts)
```

contrast	odds.ratio	SE	df	z.ratio	p.value
motorist = human					
Observer / Passenger	3.8387	2.3773	Inf	2.172	0.0760
Observer / Pedestrian	2.4179	1.3164	Inf	1.622	0.2364
Passenger / Pedestrian	0.6299	0.3037	Inf	-0.959	0.6032
motorist = self-driving					
Observer / Passenger	2.4128	1.3701	Inf	1.551	0.2672
Observer / Pedestrian	1.1494	0.5673	Inf	0.282	0.9571
Passenger / Pedestrian	0.4764	0.2310	Inf	-1.529	0.2772
Results are averaged over the levels of: trial, gender					
P value adjustment: tukey method for comparing a family of 3 estimates					
Tests are performed on the log odds ratio scale					

If we find no interaction but a significant main effect of perspective, we need to do follow-up multiple comparisons to find out where the difference is.

```
emm_carsac_persp <- emmeans(carsac_glmm, pairwise ~ perspective,
                            type = 'response')

## NOTE: Results may be misleading due to involvement in interactions

xtable(emm_carsac_persp$contrasts)
```

contrast	odds.ratio	SE	df	z.ratio	p.value
Observer / Passenger	3.0433	1.3025	Inf	2.600	0.0252
Observer / Pedestrian	1.6671	0.6142	Inf	1.387	0.3475
Passenger / Pedestrian	0.5478	0.1889	Inf	-1.745	0.1885
Results are averaged over the levels of: motorist, trial, gender					
P value adjustment: tukey method for comparing a family of 3 estimates					
Tests are performed on the log odds ratio scale					

If we find a main effect of motorist-type, we need to calculate the marginal means to see the effect.

```
emm_carsac_motorist <- emmeans(carsac_glmm, pairwise ~ motorist,
                              type = 'response')

## NOTE: Results may be misleading due to involvement in interactions

xtable(emm_carsac_motorist$contrasts)
```

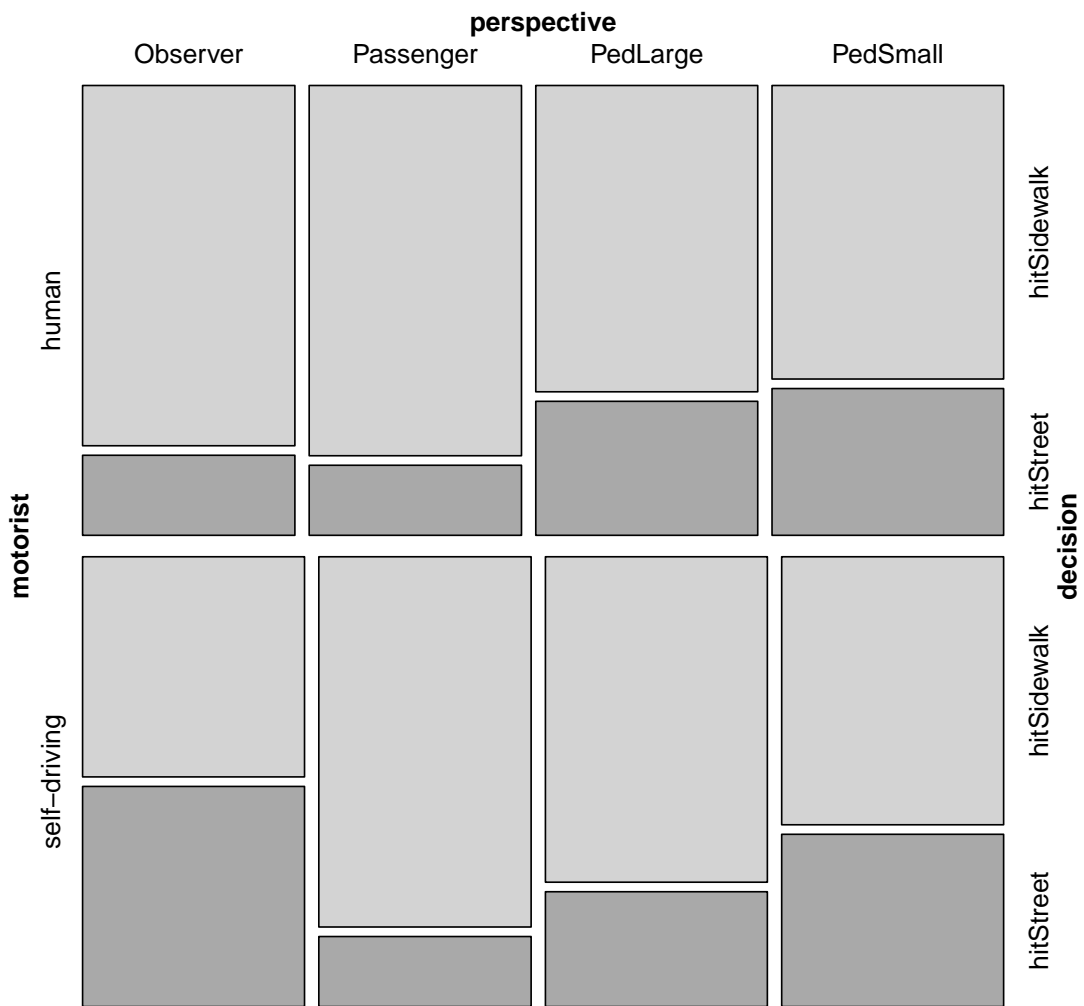
contrast	odds.ratio	SE	df	z.ratio	p.value
human / self-driving	1.3710	0.4191	Inf	1.032	0.3020

Results are averaged over the levels of: perspective, trial, gender

Tests are performed on the log odds ratio scale

## 6 Sidewalk vs road dilemma

```
sidewalk.xtabs <- xtabs(~ motorist + perspective + decision, data = sidewalk.sub)
sidewalk.mosaic <- mosaic(sidewalk.xtabs,
  gp = gpar(fill = c("light gray", "dark gray")))
```



## 6.1 The model

```
(nc <- detectCores())
```

```
[1] 12
```

```
cl <- makeCluster(rep("localhost", nc))
```

```
sidewalk_glmm <- mixed(decision == 'hitSidewalk' ~ perspective +  
  motorist +  
  perspective:motorist +  
  trial + gender + #age +  
  (1 | participant.ID),  
  method = "PB", # change to PB for final  
  family = "binomial", data = sidewalk.sub,  
  args_test = list(nsim = 1000, cl = cl), cl = cl,  
  control = glmerControl(optimizer = "bobyqa",  
    optCtrl = list(maxfun = 2e5)))
```

*## Contrasts set to contr.sum for the following variables: decision, perspective, motorist, trial, gender, participant.ID*

Fitting 6 (g)lmer() models. Obtaining 5 p-values: [.....]

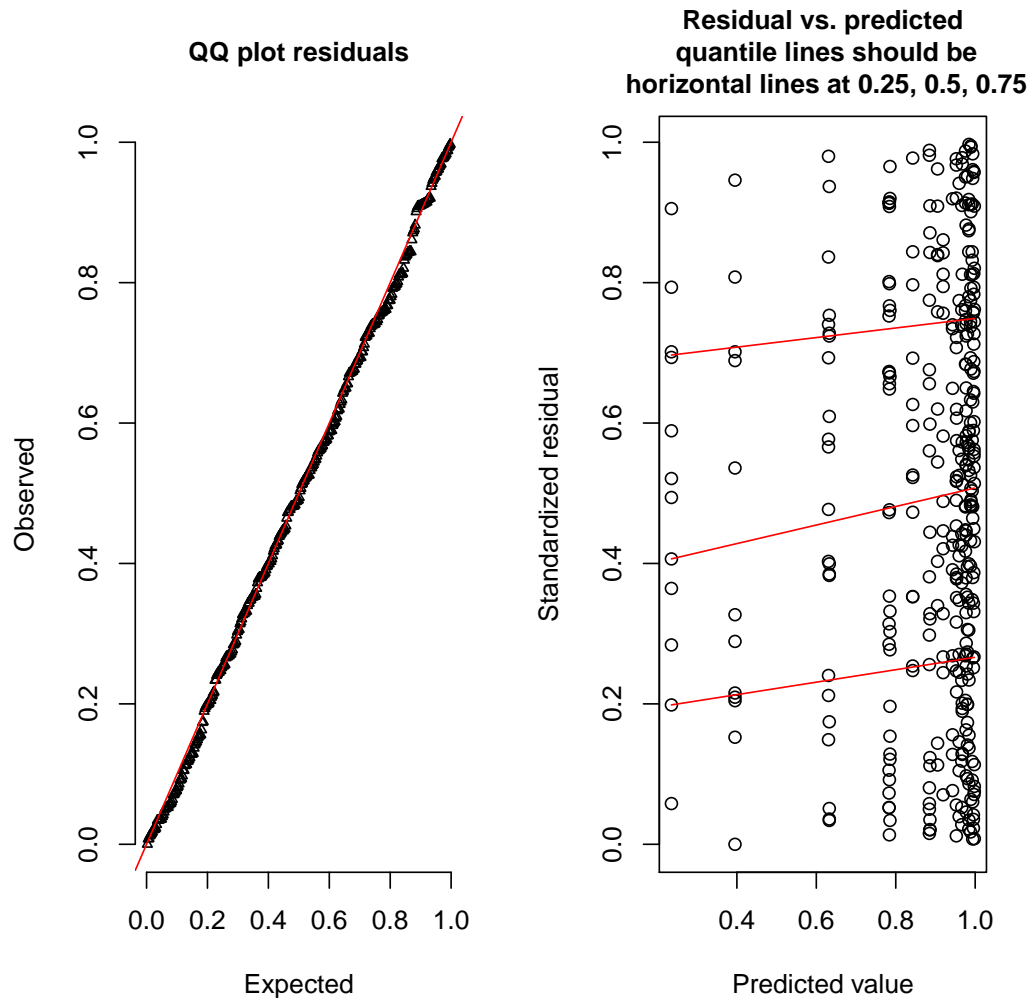
```
stopCluster(cl)  
xtable(sidewalk_glmm$anova_table)
```

	Chisq	Chi Df	Pr(>Chisq)	Pr(>PB)
perspective	7.93	3	0.0475	0.0526
motorist	2.12	1	0.1455	0.2086
trial	4.00	1	0.0455	0.0909
gender	4.90	1	0.0268	0.0058
perspective:motorist	6.64	3	0.0842	0.0536

Then we check the residuals.

```
sidewalk_glmm.resid <- simulateResiduals(  
  fittedModel = sidewalk_glmm$full_model, n = 2000)  
sidewalk_glmm_resid.plot <- plotSimulatedResiduals(  
  simulationOutput = sidewalk_glmm.resid)
```

## DHARMA scaled residual plots



## 6.2 Plotting

Depending on which effects or interactions we find, we probably want to change this plot. It currently just plots the Estimated Marginal Means for the 2-way interaction.

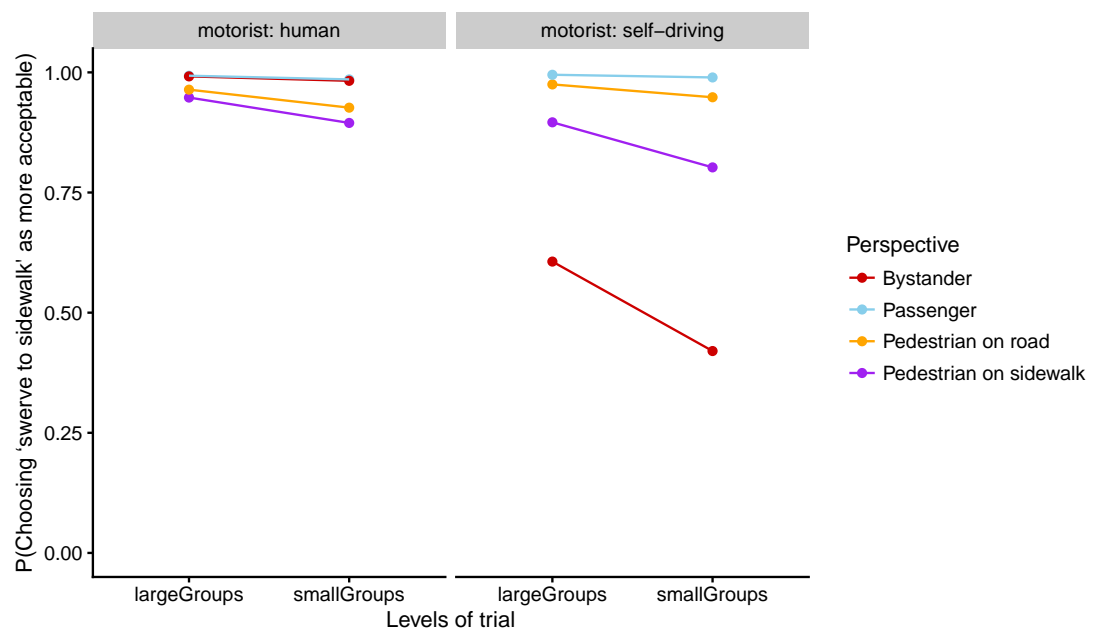
```
sidewalk.plot <- emmip(sidewalk_glmm, perspective ~ trial | motorist,
  type = "response") +
  theme_cowplot(font_size = 10) +
  scale_y_continuous(
    name = "P(Choosing `swerve to sidewalk' as more acceptable)",
    limits = c(0, 1)) + coord_equal(ratio = 3) +
  scale_color_manual(name = "Perspective",
```

```

labels = c("Bystander",
           "Passenger",
           "Pedestrian on road",
           "Pedestrian on sidewalk"),
values = c("red3", "skyblue",
           "orange1", "purple")

```

sidewalk.plot



### 6.3 Follow-up analyses

If we find an interaction between perspective and motorist-type we should follow that up with comparisons of perspective within each level of motorist.

```
emm_sidewalk_i <- emmeans(sidewalk_glmm, pairwise ~ perspective | motorist,  
                           type = 'response')  
xtable(emm_sidewalk_i$contrasts)
```

contrast	odds.ratio	SE	df	z.ratio	p.value
motorist = human					
Observer / Passenger	0.8300	1.1971	Inf	0.693	0.8997
Observer / PedLarge	4.4380	6.5893	Inf	0.674	0.9071
Observer / PedSmall	6.5827	9.7895	Inf	0.672	0.9075
Passenger / PedLarge	5.3471	7.9321	Inf	0.674	0.9069
Passenger / PedSmall	7.9310	11.7982	Inf	0.672	0.9076
PedLarge / PedSmall	1.4832	2.2475	Inf	0.660	0.9121
motorist = self-driving					
Observer / Passenger	0.0077	0.0151	Inf	0.510	0.9568
Observer / PedLarge	0.0395	0.0721	Inf	0.549	0.9469
Observer / PedSmall	0.1785	0.3171	Inf	0.563	0.9430
Passenger / PedLarge	5.1422	7.5438	Inf	0.682	0.9041
Passenger / PedSmall	23.2135	37.5966	Inf	0.617	0.9266
PedLarge / PedSmall	4.5143	7.0392	Inf	0.641	0.9186
Results are averaged over the levels of: trial, gender					
Unknown transformation "==" : no transformation done					
P value adjustment: tukey method for comparing a family of 4 estimates					

If we find no interaction but a significant main effect of perspective, we need to do follow-up multiple comparisons to find out where the difference is.

```
emm_sidewalk_persp <- emmeans(sidewalk_glmm, pairwise ~ perspective,  
                              type = 'response')  
  
## NOTE: Results may be misleading due to involvement in interactions  
  
xtable(emm_sidewalk_persp$contrasts)
```

If we find a main effect of motorist-type, we need to calculate the marginal means to see the effect.

```
emm_sidewalk_motorist <- emmeans(sidewalk_glmm, pairwise ~ motorist,  
                                 type = 'response')
```

contrast	odds.ratio	SE	df	z.ratio	p.value
Observer / Passenger	0.0799	0.0977	Inf	0.818	0.8462
Observer / PedLarge	0.4189	0.4855	Inf	0.863	0.8240
Observer / PedSmall	1.0841	1.2186	Inf	0.890	0.8103
Passenger / PedLarge	5.2436	5.5014	Inf	0.953	0.7760
Passenger / PedSmall	13.5686	15.2436	Inf	0.890	0.8100
PedLarge / PedSmall	2.5876	2.8321	Inf	0.914	0.7976

Results are averaged over the levels of: motorist, trial, gender

Unknown transformation "==" : no transformation done

P value adjustment: tukey method for comparing a family of 4 estimates

**## NOTE: Results may be misleading due to involvement in interactions**

```
xtable(emm_sidewalk_motorist$contrasts)
```

contrast	odds.ratio	SE	df	z.ratio	p.value
human / self-driving	2.9910	2.4391	Inf	1.226	0.2201

Results are averaged over the levels of: perspective, trial, gender

Unknown transformation "==" : no transformation done