

# Addendum: Modeling nonlinear effects of predictors (Bayesian GLMMs and GAMMs)

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## 1 Overview

This document builds on the GLMM homework and in-class material to use the same data set to model nonlinear effects of predictors. For convenience, I use the Bayesian GLMM developed at the end of the in-class GLMM materials as a starting point. As a reminder, this GLMM modeled the trial-level effects of the crowdedness condition, stimulus size and their interaction on the accuracy participants' responses. Here I am visualizing the models predictions both in proportion space (left) and log-odds (right):

```
my.priors <- c(
  prior(student_t(3, 0, 2.5), class = "b"),
  prior(cauchy(0,2.5), class = "sd"),
  prior(lkj(1), class = "cor")
)

Size.mu = mean(d$Size)
Size.sd = sd(d$Size)

# Standardize Size and make sure order of levels for Condition is as intended
d %<>%
  mutate(
    Condition = factor(Condition, levels = c("uncrowded", "crowded")),
    Size = (Size - mean(Size)) / (2 * sd(Size))
  )

# Sum-code condition
contrasts(d$Condition) = cbind("Crowded.vs.Uncrowded" = c(-.5,.5))

bm <- brm(
  formula = ResponseCorrect ~ 1 + Size * Condition +
    (1 + Size * Condition | Subject),
  data = d,
  family = bernoulli("logit"),
  iter = 2000,
  prior = my.priors,
  file = "../models/GLMM"
)
```

```
summary(bm)
```

```
Family: bernoulli
Links: mu = logit
Formula: ResponseCorrect ~ 1 + Size * Condition + (1 + Size * Condition | Subject)
Data: d (Number of observations: 3949)
Samples: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
         total post-warmup samples = 4000
```

```
Group-Level Effects:
~Subject (Number of levels: 10)
```

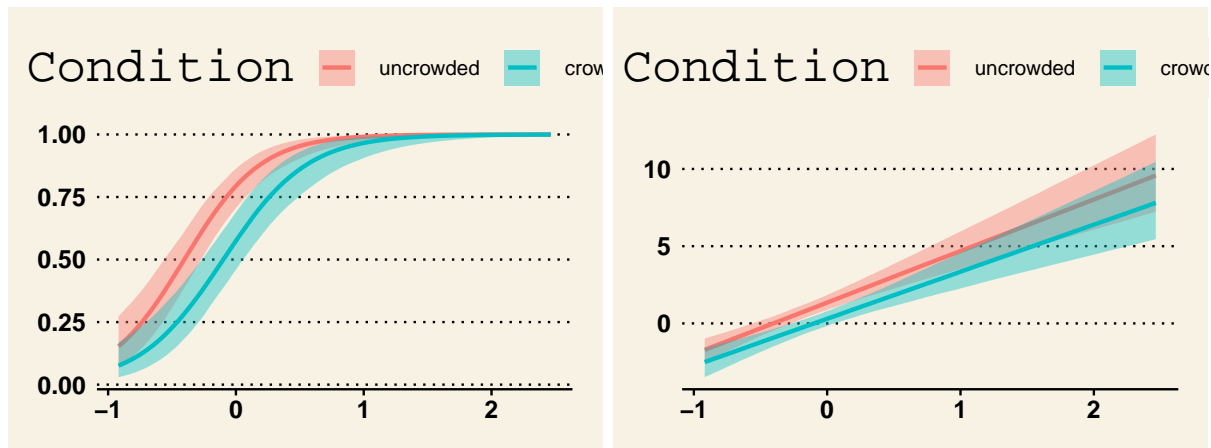
	Estimate	Est.Error	1-95% CI	u-95% CI	Rhat	Bulk_ESS	Tail_ESS
sd(Intercept)	0.67	0.19	0.40	1.15	1.00	1609	2342
sd(Size)	1.20	0.37	0.67	2.12	1.00	2237	2560
sd(ConditionCrowded.vs.Uncrowded)	0.41	0.22	0.04	0.89	1.00	864	755
sd(Size:ConditionCrowded.vs.Uncrowded)	0.42	0.35	0.02	1.29	1.00	2030	2511
cor(Intercept,Size)	0.36	0.29	-0.27	0.82	1.00	2358	2213
cor(Intercept,ConditionCrowded.vs.Uncrowded)	0.07	0.35	-0.62	0.69	1.00	4282	3089
cor(Size,ConditionCrowded.vs.Uncrowded)	-0.15	0.36	-0.77	0.58	1.00	3113	3045
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded)	-0.15	0.43	-0.85	0.72	1.00	5217	2741
cor(Size,Size:ConditionCrowded.vs.Uncrowded)	0.12	0.41	-0.71	0.83	1.00	4139	3018
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded)	0.04	0.44	-0.79	0.82	1.00	3992	3537

```
Population-Level Effects:
```

	Estimate	Est.Error	1-95% CI	u-95% CI	Rhat	Bulk_ESS	Tail_ESS
Intercept	0.83	0.22	0.40	1.29	1.00	1063	1522
Size	3.21	0.44	2.37	4.10	1.00	1722	2055
ConditionCrowded.vs.Uncrowded	-1.05	0.18	-1.40	-0.70	1.00	2816	2638
Size:ConditionCrowded.vs.Uncrowded	-0.29	0.35	-0.96	0.44	1.00	3746	2599

Samples were drawn using sampling(NUTS). For each parameter, Bulk\_ESS and Tail\_ESS are effective sample size measures, and Rhat is the potential scale reduction factor on split chains (at convergence, Rhat = 1).

```
plot(conditional_effects(bm, effects = "Size:Condition", method = "posterior_epred"), ask = F)
plot(conditional_effects(bm, effects = "Size:Condition", method = "posterior_linpred"), ask = F)
```



## 2 Is there a *linear* effect of eye-movements on the probability of a correct answer?

We test whether the amount of eye-movements during a trial—operationalized as the diffusion constant—has an effect on the probability of a correct answer. For the purpose of this example, we intentionally do not assess whether there are outliers, despite the fact that the diffusion constant is known to be a very noisy measure at the trial level (previous analyses only assessed its effect at the subject-level, i.e., effects of the average diffusion constant across all trials of a subject in a given condition). We begin with a model that assumes a linear effect of diffusion constant (on the log-odds of a correct answer) in addition to the effects of crowdedness condition, stimulus size, and their interaction. To put the diffusion constant on the same scale as the other predictors—so that the priors on all

coefficients have the same weakly regularizing effect—we center and standardize the diffusion constant following the same procedure as applied to stimulus size in the in-class part of the GLMM tutorial. We visualize the effect of the diffusion constant on the proportion (left) and log-odds (right) of a correct answer:

```
DiffusionConstant.mu = mean(d$DiffusionConstant)
DiffusionConstant.sd = sd(d$DiffusionConstant)
```

```
d %<>%
  mutate(
    DiffusionConstant = (DiffusionConstant - mean(DiffusionConstant)) / (2 * sd(DiffusionConstant)))

bm.wDiffusionConstant <- brm(
  formula = ResponseCorrect ~ 1 + Size * Condition + DiffusionConstant +
    (1 + Size * Condition | Subject),
  data = d,
  family = bernoulli("logit"),
  iter = 2000,
  prior = my.priors,
  file = "../models/GLMM-with-DiffusionConstant"
)

summary(bm.wDiffusionConstant)
```

```
Family: bernoulli
Links: mu = logit
Formula: ResponseCorrect ~ 1 + Size * Condition + DiffusionConstant + (1 + Size * Condition | Subject)
Data: d (Number of observations: 6539)
Samples: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
         total post-warmup samples = 4000
```

Group-Level Effects:

~Subject (Number of levels: 10)

	Estimate
sd(Intercept)	0.64
sd(Size)	1.05
sd(ConditionCrowded.vs.Uncrowded)	0.38
sd(Size:ConditionCrowded.vs.Uncrowded)	0.36
cor(Intercept,Size)	0.44
cor(Intercept,ConditionCrowded.vs.Uncrowded)	0.07
cor(Size,ConditionCrowded.vs.Uncrowded)	-0.06
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded)	-0.11
cor(Size,Size:ConditionCrowded.vs.Uncrowded)	0.10
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded)	0.21
	Est.Error
sd(Intercept)	0.20
sd(Size)	0.33
sd(ConditionCrowded.vs.Uncrowded)	0.18
sd(Size:ConditionCrowded.vs.Uncrowded)	0.30
cor(Intercept,Size)	0.26
cor(Intercept,ConditionCrowded.vs.Uncrowded)	0.33
cor(Size,ConditionCrowded.vs.Uncrowded)	0.34
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded)	0.42
cor(Size,Size:ConditionCrowded.vs.Uncrowded)	0.40
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded)	0.43
	1-95% CI
sd(Intercept)	0.37
sd(Size)	0.60
sd(ConditionCrowded.vs.Uncrowded)	0.08
sd(Size:ConditionCrowded.vs.Uncrowded)	0.02
cor(Intercept,Size)	-0.16
cor(Intercept,ConditionCrowded.vs.Uncrowded)	-0.58
cor(Size,ConditionCrowded.vs.Uncrowded)	-0.66
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded)	-0.81
cor(Size,Size:ConditionCrowded.vs.Uncrowded)	-0.71
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded)	-0.68
	u-95% CI
sd(Intercept)	1.12
sd(Size)	1.88
sd(ConditionCrowded.vs.Uncrowded)	0.82
sd(Size:ConditionCrowded.vs.Uncrowded)	1.06
cor(Intercept,Size)	0.85

```

cor(Intercept,ConditionCrowded.vs.Uncrowded) 0.68
cor(Size,ConditionCrowded.vs.Uncrowded) 0.63
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded) 0.73
cor(Size,Size:ConditionCrowded.vs.Uncrowded) 0.80
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded) 0.89

Rhat
sd(Intercept) 1.00
sd(Size) 1.00
sd(ConditionCrowded.vs.Uncrowded) 1.00
sd(Size:ConditionCrowded.vs.Uncrowded) 1.00
cor(Intercept,Size) 1.00
cor(Intercept,ConditionCrowded.vs.Uncrowded) 1.00
cor(Size,ConditionCrowded.vs.Uncrowded) 1.00
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded) 1.00
cor(Size,Size:ConditionCrowded.vs.Uncrowded) 1.00
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded) 1.00

Bulk_ESS
sd(Intercept) 1282
sd(Size) 1811
sd(ConditionCrowded.vs.Uncrowded) 845
sd(Size:ConditionCrowded.vs.Uncrowded) 1635
cor(Intercept,Size) 2498
cor(Intercept,ConditionCrowded.vs.Uncrowded) 3260
cor(Size,ConditionCrowded.vs.Uncrowded) 2361
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded) 4815
cor(Size,Size:ConditionCrowded.vs.Uncrowded) 4465
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded) 3932

Tail_ESS
sd(Intercept) 2171
sd(Size) 2301
sd(ConditionCrowded.vs.Uncrowded) 640
sd(Size:ConditionCrowded.vs.Uncrowded) 1749
cor(Intercept,Size) 2544
cor(Intercept,ConditionCrowded.vs.Uncrowded) 2995
cor(Size,ConditionCrowded.vs.Uncrowded) 2262
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded) 3039
cor(Size,Size:ConditionCrowded.vs.Uncrowded) 2893
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded) 3288

```

#### Population-Level Effects:

	Estimate	Est.Error	1-95% CI	u-95% CI	Rhat
Intercept	0.81	0.21	0.36	1.23	1.00
Size	2.89	0.38	2.12	3.60	1.00
ConditionCrowded.vs.Uncrowded	-0.91	0.16	-1.22	-0.59	1.00
DiffusionConstant	-0.02	0.06	-0.14	0.09	1.00
Size:ConditionCrowded.vs.Uncrowded	-0.25	0.27	-0.73	0.32	1.00

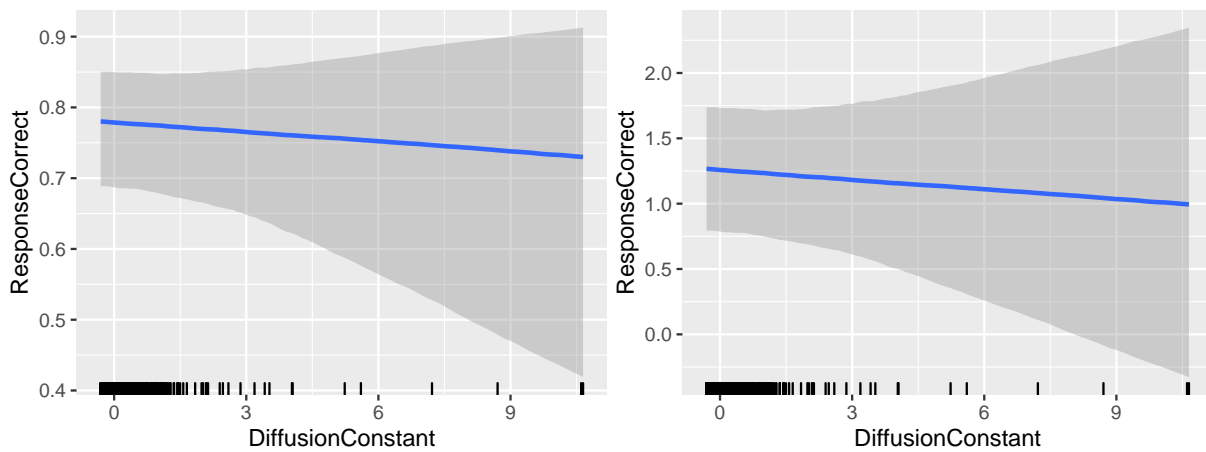
	Bulk_ESS	Tail_ESS
Intercept	951	1470
Size	1508	1786
ConditionCrowded.vs.Uncrowded	2150	2362
DiffusionConstant	6629	2817
Size:ConditionCrowded.vs.Uncrowded	3487	2346

Samples were drawn using sampling(NUTS). For each parameter, Bulk\_ESS and Tail\_ESS are effective sample size measures, and Rhat is the potential scale reduction factor on split chains (at convergence, Rhat = 1).

```

plot(
  conditional_effects(bm.wDiffusionConstant, effects = "DiffusionConstant", method = "posterior_epred"),
  ask = F, rug = T)
plot(
  conditional_effects(bm.wDiffusionConstant, effects = "DiffusionConstant", method = "posterior_linpred"),
  ask = F, rug = T)

```



One issue to expect from the plots shown above is the sparsity of data for high values of the diffusion constant, as evidenced in the data rug along the x-axis. Extreme values like this can be overly influential on the model fit, and this risk increases as we increase the functional flexibility of the effect of the diffusion constant in the next section (in order to entertain nonlinear effects).

### 3 Is there a *nonlinear* effect of eye-movements on the probability of a correct answer?

Next we entertain three ways of detecting non-linear effects of the diffusion constant. It should be noted though that a purely exploratory approach like this inflates the number of tests we conduct and thus the family-wise Type I error rate. In particular, since we are entertaining nonlinear effects a blind exploration of possible fits risks overfitting the model to the data. It is thus highly advisable to take a theory-driven approach, where the theory constraints the functional shape of the nonlinear effect. Whenever an exploratory approach is taken, it should be clearly indicated along with the consequences (risk of overfitting and inflated Type I error rate).

#### 3.1 Polynomials

One approach to modeling nonlinear effect are polynomials. Here we use R's `poly` function to obtain *orthogonal* polynomials of the third order. We again visualize the effect of the diffusion constant on the proportion (left) and log-odds (right) of a correct answer. Based on both the model output and the plots, there is little evidence of nonlinearities in the effect of the diffusion constant, and no evidence of any non-zero effect of diffusion constant. The model output, for example, shows that the 95% credible intervals of all three components of the polynomial—including the linear component—include zero:

```
bm.wDiffusionConstant.poly <- brm(
  formula = ResponseCorrect ~ 1 + Size * Condition + poly(DiffusionConstant,3) +
    (1 + Size * Condition | Subject),
  data = d,
  family = bernoulli("logit"),
  iter = 2000,
  prior = my.priors,
  file = "../models/GLMM-with-DiffusionConstant-poly",
  control = list(adapt_delta = .95)
)
```

```
summary(bm.wDiffusionConstant.poly)
```

```
Family: bernoulli
Links: mu = logit
Formula: ResponseCorrect ~ 1 + Size * Condition + poly(DiffusionConstant, 3) + (1 + Size * Condition | Subject)
Data: d (Number of observations: 6539)
Samples: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
         total post-warmup samples = 4000
```

Group-Level Effects:

~Subject (Number of levels: 10)

	Estimate
sd(Intercept)	0.63
sd(Size)	1.06
sd(ConditionCrowded.vs.Uncrowded)	0.39
sd(Size:ConditionCrowded.vs.Uncrowded)	0.36
cor(Intercept,Size)	0.44
cor(Intercept,ConditionCrowded.vs.Uncrowded)	0.07
cor(Size,ConditionCrowded.vs.Uncrowded)	-0.06
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded)	-0.11
cor(Size,Size:ConditionCrowded.vs.Uncrowded)	0.10
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded)	0.19
	Est.Error
sd(Intercept)	0.19
sd(Size)	0.32
sd(ConditionCrowded.vs.Uncrowded)	0.18
sd(Size:ConditionCrowded.vs.Uncrowded)	0.29
cor(Intercept,Size)	0.26
cor(Intercept,ConditionCrowded.vs.Uncrowded)	0.33
cor(Size,ConditionCrowded.vs.Uncrowded)	0.34
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded)	0.41
cor(Size,Size:ConditionCrowded.vs.Uncrowded)	0.41
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded)	0.43
	1-95% CI
sd(Intercept)	0.38
sd(Size)	0.60
sd(ConditionCrowded.vs.Uncrowded)	0.09
sd(Size:ConditionCrowded.vs.Uncrowded)	0.01
cor(Intercept,Size)	-0.15
cor(Intercept,ConditionCrowded.vs.Uncrowded)	-0.58
cor(Size,ConditionCrowded.vs.Uncrowded)	-0.65
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded)	-0.81
cor(Size,Size:ConditionCrowded.vs.Uncrowded)	-0.72
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded)	-0.70
	u-95% CI
sd(Intercept)	1.10
sd(Size)	1.84
sd(ConditionCrowded.vs.Uncrowded)	0.81
sd(Size:ConditionCrowded.vs.Uncrowded)	1.04
cor(Intercept,Size)	0.84
cor(Intercept,ConditionCrowded.vs.Uncrowded)	0.68
cor(Size,ConditionCrowded.vs.Uncrowded)	0.60
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded)	0.71
cor(Size,Size:ConditionCrowded.vs.Uncrowded)	0.82
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded)	0.88
	Rhat
sd(Intercept)	1.00
sd(Size)	1.00
sd(ConditionCrowded.vs.Uncrowded)	1.00
sd(Size:ConditionCrowded.vs.Uncrowded)	1.00
cor(Intercept,Size)	1.00
cor(Intercept,ConditionCrowded.vs.Uncrowded)	1.00
cor(Size,ConditionCrowded.vs.Uncrowded)	1.00
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded)	1.00
cor(Size,Size:ConditionCrowded.vs.Uncrowded)	1.00
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded)	1.00
	Bulk_ESS
sd(Intercept)	1289
sd(Size)	1718
sd(ConditionCrowded.vs.Uncrowded)	1219
sd(Size:ConditionCrowded.vs.Uncrowded)	1552
cor(Intercept,Size)	1851
cor(Intercept,ConditionCrowded.vs.Uncrowded)	2869
cor(Size,ConditionCrowded.vs.Uncrowded)	2339
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded)	4290
cor(Size,Size:ConditionCrowded.vs.Uncrowded)	3977
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded)	3237
	Tail_ESS
sd(Intercept)	1988
sd(Size)	2496

```

sd(ConditionCrowded.vs.Uncrowded)          1158
sd(Size:ConditionCrowded.vs.Uncrowded)      1347
cor(Intercept,Size)                        2288
cor(Intercept,ConditionCrowded.vs.Uncrowded) 2596
cor(Size,ConditionCrowded.vs.Uncrowded)      2832
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded) 2945
cor(Size,Size:ConditionCrowded.vs.Uncrowded) 3113
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded) 2933

```

Population-Level Effects:

	Estimate	Est.Error	l-95% CI	u-95% CI	Rhat
Intercept	0.82	0.21	0.40	1.24	1.01
Size	2.90	0.37	2.15	3.62	1.00
ConditionCrowded.vs.Uncrowded	-0.91	0.15	-1.23	-0.60	1.00
polyDiffusionConstant31	-0.56	1.82	-4.27	2.99	1.00
polyDiffusionConstant32	-0.31	1.78	-3.91	3.20	1.00
polyDiffusionConstant33	-1.25	1.83	-5.18	2.17	1.00
Size:ConditionCrowded.vs.Uncrowded	-0.24	0.26	-0.73	0.30	1.00

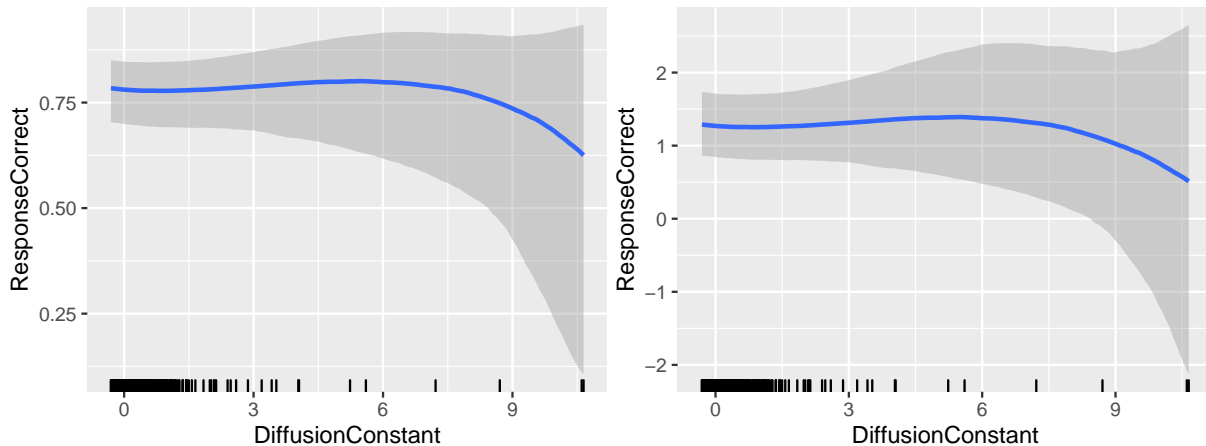
	Bulk_ESS	Tail_ESS
Intercept	742	1377
Size	1208	1525
ConditionCrowded.vs.Uncrowded	2091	2560
polyDiffusionConstant31	5001	2260
polyDiffusionConstant32	5568	2628
polyDiffusionConstant33	5457	2541
Size:ConditionCrowded.vs.Uncrowded	2723	2331

Samples were drawn using sampling(NUTS). For each parameter, Bulk\_ESS and Tail\_ESS are effective sample size measures, and Rhat is the potential scale reduction factor on split chains (at convergence, Rhat = 1).

```

plot(
  conditional_effects(bm.wDiffusionConstant.poly, effects = "DiffusionConstant", method = "posterior_epred"),
  ask = F, rug = T)
plot(
  conditional_effects(bm.wDiffusionConstant.poly, effects = "DiffusionConstant", method = "posterior_linpred"),
  ask = F, rug = T)

```



### 3.2 Non-parametric smooths in a GAMM

Alternatively, we can use `brms`'s ability to fit generalized additive mixed models (GAMMs) and model the effect of the diffusion constant as a non-parametric smooth. If this method is applied in an actual research project, it is important to read up on the many different options you can select from when fitting such smooths, their consequences, and potential risks. We again visualize the effect of the diffusion constant on the proportion (left) and log-odds (right) of a correct answer. Based on this output, too, we see no evidence of linear or non-linear effects of the diffusion constant.

```

bm.wDiffusionConstant.smooth <- brm(
  formula = ResponseCorrect ~ 1 + Size * Condition + s(DiffusionConstant) +
    (1 + Size * Condition | Subject),

```

```

data = d,
family = bernoulli("logit"),
iter = 2000,
prior = my.priors,
file = "../models/GLMM-with-DiffusionConstant-smooth",
control = list(adapt_delta = .95)
)

summary(bm.wDiffusionConstant.smooth)

Family: bernoulli
Links: mu = logit
Formula: ResponseCorrect ~ 1 + Size * Condition + s(DiffusionConstant) + (1 + Size * Condition | Subject)
Data: d (Number of observations: 6539)
Samples: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
         total post-warmup samples = 4000

Smooth Terms:
              Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS
sds(sDiffusionConstant_1)    0.73    0.76    0.02    2.73 1.00    1864
              Tail_ESS
sds(sDiffusionConstant_1)    2047

Group-Level Effects:
~Subject (Number of levels: 10)

              Estimate
sd(Intercept)    0.64
sd(Size)         1.06
sd(ConditionCrowded.vs.Uncrowded) 0.38
sd(Size:ConditionCrowded.vs.Uncrowded) 0.36
cor(Intercept,Size) 0.44
cor(Intercept,ConditionCrowded.vs.Uncrowded) 0.06
cor(Size,ConditionCrowded.vs.Uncrowded) -0.07
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded) -0.12
cor(Size,Size:ConditionCrowded.vs.Uncrowded) 0.09
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded) 0.19

              Est.Error
sd(Intercept)    0.19
sd(Size)         0.33
sd(ConditionCrowded.vs.Uncrowded) 0.18
sd(Size:ConditionCrowded.vs.Uncrowded) 0.29
cor(Intercept,Size) 0.27
cor(Intercept,ConditionCrowded.vs.Uncrowded) 0.34
cor(Size,ConditionCrowded.vs.Uncrowded) 0.35
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded) 0.41
cor(Size,Size:ConditionCrowded.vs.Uncrowded) 0.41
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded) 0.44

              1-95% CI
sd(Intercept)    0.38
sd(Size)         0.59
sd(ConditionCrowded.vs.Uncrowded) 0.06
sd(Size:ConditionCrowded.vs.Uncrowded) 0.01
cor(Intercept,Size) -0.20
cor(Intercept,ConditionCrowded.vs.Uncrowded) -0.59
cor(Size,ConditionCrowded.vs.Uncrowded) -0.70
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded) -0.80
cor(Size,Size:ConditionCrowded.vs.Uncrowded) -0.70
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded) -0.72

              u-95% CI
sd(Intercept)    1.11
sd(Size)         1.88
sd(ConditionCrowded.vs.Uncrowded) 0.79
sd(Size:ConditionCrowded.vs.Uncrowded) 1.05
cor(Intercept,Size) 0.84
cor(Intercept,ConditionCrowded.vs.Uncrowded) 0.67
cor(Size,ConditionCrowded.vs.Uncrowded) 0.62
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded) 0.72
cor(Size,Size:ConditionCrowded.vs.Uncrowded) 0.80
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded) 0.89

              Rhat
sd(Intercept)    1.00

```



```

sd(Size) 1.00
sd(ConditionCrowded.vs.Uncrowded) 1.00
sd(Size:ConditionCrowded.vs.Uncrowded) 1.00
cor(Intercept,Size) 1.01
cor(Intercept,ConditionCrowded.vs.Uncrowded) 1.00
cor(Size,ConditionCrowded.vs.Uncrowded) 1.00
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded) 1.00
cor(Size,Size:ConditionCrowded.vs.Uncrowded) 1.00
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded) 1.00

Bulk_ESS
sd(Intercept) 1080
sd(Size) 1457
sd(ConditionCrowded.vs.Uncrowded) 1013
sd(Size:ConditionCrowded.vs.Uncrowded) 1757
cor(Intercept,Size) 1529
cor(Intercept,ConditionCrowded.vs.Uncrowded) 3037
cor(Size,ConditionCrowded.vs.Uncrowded) 1809
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded) 3921
cor(Size,Size:ConditionCrowded.vs.Uncrowded) 4372
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded) 3290

Tail_ESS
sd(Intercept) 1597
sd(Size) 1812
sd(ConditionCrowded.vs.Uncrowded) 959
sd(Size:ConditionCrowded.vs.Uncrowded) 1903
cor(Intercept,Size) 2216
cor(Intercept,ConditionCrowded.vs.Uncrowded) 2852
cor(Size,ConditionCrowded.vs.Uncrowded) 2328
cor(Intercept,Size:ConditionCrowded.vs.Uncrowded) 2815
cor(Size,Size:ConditionCrowded.vs.Uncrowded) 3142
cor(ConditionCrowded.vs.Uncrowded,Size:ConditionCrowded.vs.Uncrowded) 2955

```

#### Population-Level Effects:

	Estimate	Est.Error	l-95% CI	u-95% CI	Rhat
Intercept	0.81	0.22	0.34	1.24	1.01
Size	2.90	0.38	2.13	3.65	1.00
ConditionCrowded.vs.Uncrowded	-0.91	0.16	-1.23	-0.60	1.00
Size:ConditionCrowded.vs.Uncrowded	-0.25	0.26	-0.73	0.30	1.00
sDiffusionConstant_1	-0.41	1.35	-3.29	2.22	1.00

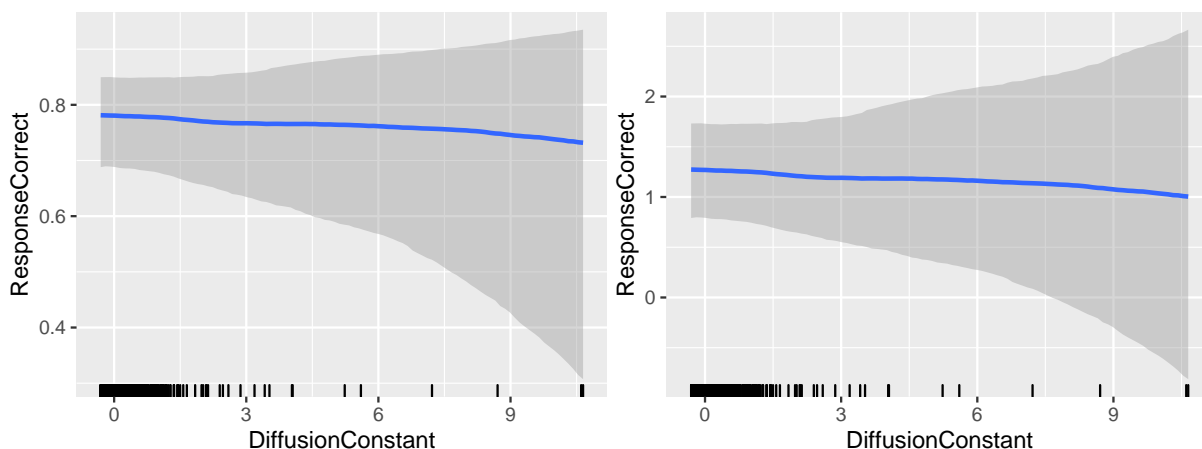
	Bulk_ESS	Tail_ESS
Intercept	801	1160
Size	1247	1453
ConditionCrowded.vs.Uncrowded	1939	2114
Size:ConditionCrowded.vs.Uncrowded	3083	1856
sDiffusionConstant_1	3334	2535

Samples were drawn using sampling(NUTS). For each parameter, Bulk\_ESS and Tail\_ESS are effective sample size measures, and Rhat is the potential scale reduction factor on split chains (at convergence, Rhat = 1).

```

plot(
  conditional_effects(bm.wDiffusionConstant.smooth, effects = "DiffusionConstant", method = "posterior_epred"),
  ask = F, rug = T)
plot(
  conditional_effects(bm.wDiffusionConstant.smooth, effects = "DiffusionConstant", method = "posterior_linpred"),
  ask = F, rug = T)

```



Notice further how this non-parametric smooth avoids the perhaps misleading impression that one might get from the polynomial in the previous section: the indication of a non-linear effect for large values of diffusion constant. Polynomial fits are well-known to be overly sensitive to data points with extreme values for the predictor, where the data is often sparse. This is one way in which polynomials are likely to overfit the data, yielding bad predictions in particular for novel data with predictor values that fall outside of the range observed in the sample the model was fit to.

## 4 Session info

```
devtools::session_info()
```

```

- Session info -----
setting  value
version  R version 4.0.2 (2020-06-22)
os       macOS High Sierra 10.13.6
system   x86_64, darwin17.0
ui       X11
language (EN)
collate  en_US.UTF-8
ctype    en_US.UTF-8
tz       America/New_York
date     2020-11-15

```

```

- Packages -----
package      * version      date      lib source
abind         1.4-5        2016-07-21 [1] CRAN (R 4.0.2)
arrayhelpers  1.1-0        2020-02-04 [1] CRAN (R 4.0.2)
assertthat    0.2.1        2019-03-21 [1] CRAN (R 4.0.2)
backports     1.1.10       2020-09-15 [1] CRAN (R 4.0.2)
base64enc     0.1-3        2015-07-28 [1] CRAN (R 4.0.2)
bayesplot     1.7.2        2020-05-28 [1] CRAN (R 4.0.2)
bayestestR    0.7.5        2020-10-22 [1] CRAN (R 4.0.2)
blob          1.2.1        2020-01-20 [1] CRAN (R 4.0.2)
boot          1.3-25       2020-04-26 [1] CRAN (R 4.0.2)
bridgesampling 1.0-0        2020-02-26 [1] CRAN (R 4.0.2)
brms          * 2.14.0       2020-10-08 [1] CRAN (R 4.0.2)
Brobdingnag   1.2-6        2018-08-13 [1] CRAN (R 4.0.2)
broom         0.7.2        2020-10-20 [1] CRAN (R 4.0.2)
broom.mixed   * 0.2.6        2020-05-17 [1] CRAN (R 4.0.2)
callr         3.5.1        2020-10-13 [1] CRAN (R 4.0.2)
cellranger    1.1.0        2016-07-27 [1] CRAN (R 4.0.2)
cli           2.1.0        2020-10-12 [1] CRAN (R 4.0.2)
coda          0.19-4       2020-09-30 [1] CRAN (R 4.0.2)
codetools     0.2-16       2018-12-24 [1] CRAN (R 4.0.2)
colorspace    1.4-1        2019-03-18 [1] CRAN (R 4.0.2)
colourpicker  1.1.0        2020-09-14 [1] CRAN (R 4.0.2)
crayon        1.3.4        2017-09-16 [1] CRAN (R 4.0.2)
crosstalk     1.1.0.1      2020-03-13 [1] CRAN (R 4.0.2)
curl          4.3          2019-12-02 [1] CRAN (R 4.0.1)

```

DBI	1.1.0	2019-12-15	[1]	CRAN	(R 4.0.2)
dbplyr	1.4.4	2020-05-27	[1]	CRAN	(R 4.0.2)
desc	1.2.0	2018-05-01	[1]	CRAN	(R 4.0.2)
devtools	2.3.2	2020-09-18	[1]	CRAN	(R 4.0.2)
digest	0.6.27	2020-10-24	[1]	CRAN	(R 4.0.2)
distributional	0.2.1	2020-10-06	[1]	CRAN	(R 4.0.2)
dplyr	* 1.0.2	2020-08-18	[1]	CRAN	(R 4.0.2)
DT	0.16	2020-10-13	[1]	CRAN	(R 4.0.2)
dygraphs	1.1.1.6	2018-07-11	[1]	CRAN	(R 4.0.2)
effectsize	0.4.0	2020-10-25	[1]	CRAN	(R 4.0.2)
ellipsis	0.3.1	2020-05-15	[1]	CRAN	(R 4.0.2)
emmeans	1.5.2-1	2020-10-25	[1]	CRAN	(R 4.0.2)
estimability	1.3	2018-02-11	[1]	CRAN	(R 4.0.2)
evaluate	0.14	2019-05-28	[1]	CRAN	(R 4.0.1)
fansi	0.4.1	2020-01-08	[1]	CRAN	(R 4.0.2)
farver	2.0.3	2020-01-16	[1]	CRAN	(R 4.0.2)
fastmap	1.0.1	2019-10-08	[1]	CRAN	(R 4.0.2)
forcats	* 0.5.0	2020-03-01	[1]	CRAN	(R 4.0.2)
fs	1.5.0	2020-07-31	[1]	CRAN	(R 4.0.2)
generics	0.1.0	2020-10-31	[1]	CRAN	(R 4.0.2)
ggdist	2.3.0	2020-10-31	[1]	CRAN	(R 4.0.2)
ggeffects	0.16.0	2020-09-13	[1]	CRAN	(R 4.0.2)
ggforce	* 0.3.2	2020-06-23	[1]	CRAN	(R 4.0.2)
ggplot2	* 3.3.2	2020-06-19	[1]	CRAN	(R 4.0.2)
ggridges	0.5.2	2020-01-12	[1]	CRAN	(R 4.0.2)
ggthemes	* 4.2.0	2019-05-13	[1]	CRAN	(R 4.0.2)
glue	1.4.2	2020-08-27	[1]	CRAN	(R 4.0.2)
gridExtra	2.3	2017-09-09	[1]	CRAN	(R 4.0.2)
gtable	0.3.0	2019-03-25	[1]	CRAN	(R 4.0.2)
gttools	3.8.2	2020-03-31	[1]	CRAN	(R 4.0.2)
haven	2.3.1	2020-06-01	[1]	CRAN	(R 4.0.2)
hms	0.5.3	2020-01-08	[1]	CRAN	(R 4.0.2)
htmltools	0.5.0	2020-06-16	[1]	CRAN	(R 4.0.2)
htmlwidgets	1.5.2	2020-10-03	[1]	CRAN	(R 4.0.2)
httpuv	1.5.4	2020-06-06	[1]	CRAN	(R 4.0.2)
httr	1.4.2	2020-07-20	[1]	CRAN	(R 4.0.2)
igraph	1.2.6	2020-10-06	[1]	CRAN	(R 4.0.2)
inline	0.3.16	2020-09-06	[1]	CRAN	(R 4.0.2)
insight	0.10.0	2020-10-22	[1]	CRAN	(R 4.0.2)
jsonlite	1.7.1	2020-09-07	[1]	CRAN	(R 4.0.2)
knitr	* 1.30	2020-09-22	[1]	CRAN	(R 4.0.2)
labeling	0.4.2	2020-10-20	[1]	CRAN	(R 4.0.2)
later	1.1.0.1	2020-06-05	[1]	CRAN	(R 4.0.2)
lattice	0.20-41	2020-04-02	[1]	CRAN	(R 4.0.2)
lifecycle	0.2.0	2020-03-06	[1]	CRAN	(R 4.0.2)
lme4	* 1.1-25	2020-10-23	[1]	CRAN	(R 4.0.2)
loo	2.3.1	2020-07-14	[1]	CRAN	(R 4.0.2)
lubridate	1.7.9	2020-06-08	[1]	CRAN	(R 4.0.2)
magrittr	* 1.5	2014-11-22	[1]	CRAN	(R 4.0.2)
markdown	1.1	2019-08-07	[1]	CRAN	(R 4.0.2)
MASS	7.3-53	2020-09-09	[1]	CRAN	(R 4.0.2)
Matrix	* 1.2-18	2019-11-27	[1]	CRAN	(R 4.0.2)
matrixStats	0.57.0	2020-09-25	[1]	CRAN	(R 4.0.2)
memoise	1.1.0	2017-04-21	[1]	CRAN	(R 4.0.2)
mgcv	1.8-33	2020-08-27	[1]	CRAN	(R 4.0.2)
mime	0.9	2020-02-04	[1]	CRAN	(R 4.0.2)
miniUI	0.1.1.1	2018-05-18	[1]	CRAN	(R 4.0.2)
minqa	1.2.4	2014-10-09	[1]	CRAN	(R 4.0.2)
modelr	0.1.8	2020-05-19	[1]	CRAN	(R 4.0.2)
multcomp	1.4-14	2020-09-23	[1]	CRAN	(R 4.0.2)
munSELL	0.5.0	2018-06-12	[1]	CRAN	(R 4.0.2)
mvtnorm	1.1-1	2020-06-09	[1]	CRAN	(R 4.0.2)
nlme	3.1-150	2020-10-24	[1]	CRAN	(R 4.0.2)
nloptr	1.2.2.2	2020-07-02	[1]	CRAN	(R 4.0.2)
parameters	0.9.0	2020-10-29	[1]	CRAN	(R 4.0.2)
performance	0.5.1	2020-10-29	[1]	CRAN	(R 4.0.2)
pillar	1.4.6	2020-07-10	[1]	CRAN	(R 4.0.2)
pkgbuild	1.1.0.9000	2020-08-06	[1]	Github	(r-lib/pkgbuild@3a87bd9)
pkgconfig	2.0.3	2019-09-22	[1]	CRAN	(R 4.0.2)
pkgload	1.1.0	2020-05-29	[1]	CRAN	(R 4.0.2)
plyr	1.8.6	2020-03-03	[1]	CRAN	(R 4.0.2)

polyclip	1.10-0	2019-03-14	[1]	CRAN	(R 4.0.2)
prettyunits	1.1.1	2020-01-24	[1]	CRAN	(R 4.0.2)
processx	3.4.4	2020-09-03	[1]	CRAN	(R 4.0.2)
promises	1.1.1	2020-06-09	[1]	CRAN	(R 4.0.2)
ps	1.4.0	2020-10-07	[1]	CRAN	(R 4.0.2)
purrr	* 0.3.4	2020-04-17	[1]	CRAN	(R 4.0.2)
R6	2.5.0	2020-10-28	[1]	CRAN	(R 4.0.2)
Rcpp	* 1.0.5	2020-07-06	[1]	CRAN	(R 4.0.2)
RcppParallel	5.0.2	2020-06-24	[1]	CRAN	(R 4.0.2)
readr	* 1.4.0	2020-10-05	[1]	CRAN	(R 4.0.2)
readxl	1.3.1	2019-03-13	[1]	CRAN	(R 4.0.2)
remotes	2.2.0	2020-07-21	[1]	CRAN	(R 4.0.2)
reprer	0.3.0	2019-05-16	[1]	CRAN	(R 4.0.2)
reshape2	1.4.4	2020-04-09	[1]	CRAN	(R 4.0.2)
rlang	0.4.8	2020-10-08	[1]	CRAN	(R 4.0.2)
rmarkdown	2.5	2020-10-21	[1]	CRAN	(R 4.0.2)
rprojroot	1.3-2	2018-01-03	[1]	CRAN	(R 4.0.2)
rsconnect	0.8.16	2019-12-13	[1]	CRAN	(R 4.0.2)
rstan	2.21.2	2020-08-04	[1]	Github (stan-dev/rstan@a9a44bb)	
rstantools	2.1.1	2020-07-06	[1]	CRAN	(R 4.0.2)
rstudioapi	0.11	2020-02-07	[1]	CRAN	(R 4.0.2)
rvest	0.3.6	2020-07-25	[1]	CRAN	(R 4.0.2)
sandwich	3.0-0	2020-10-02	[1]	CRAN	(R 4.0.2)
scales	1.1.1	2020-05-11	[1]	CRAN	(R 4.0.2)
sessioninfo	1.1.1	2018-11-05	[1]	CRAN	(R 4.0.2)
shiny	1.5.0	2020-06-23	[1]	CRAN	(R 4.0.2)
shinyjs	2.0.0	2020-09-09	[1]	CRAN	(R 4.0.2)
shinytan	2.5.0	2018-05-01	[1]	CRAN	(R 4.0.2)
shinythemes	1.1.2	2018-11-06	[1]	CRAN	(R 4.0.2)
sjlabelled	1.1.7	2020-09-24	[1]	CRAN	(R 4.0.2)
sjmisc	2.8.5	2020-05-28	[1]	CRAN	(R 4.0.2)
sjPlot	* 2.8.6	2020-10-28	[1]	CRAN	(R 4.0.2)
sjstats	0.18.0	2020-05-06	[1]	CRAN	(R 4.0.2)
SparseM	1.78	2019-12-13	[1]	CRAN	(R 4.0.2)
StanHeaders	2.21.0-6	2020-08-16	[1]	CRAN	(R 4.0.2)
statmod	1.4.35	2020-10-19	[1]	CRAN	(R 4.0.2)
stringi	1.5.3	2020-09-09	[1]	CRAN	(R 4.0.2)
stringr	* 1.4.0	2019-02-10	[1]	CRAN	(R 4.0.2)
survival	3.2-7	2020-09-28	[1]	CRAN	(R 4.0.2)
svUnit	1.0.3	2020-04-20	[1]	CRAN	(R 4.0.2)
testthat	3.0.0	2020-10-31	[1]	CRAN	(R 4.0.2)
TH.data	1.0-10	2019-01-21	[1]	CRAN	(R 4.0.2)
threejs	0.3.3	2020-01-21	[1]	CRAN	(R 4.0.2)
tibble	* 3.0.4	2020-10-12	[1]	CRAN	(R 4.0.2)
tidybayes	* 2.3.1	2020-11-02	[1]	CRAN	(R 4.0.2)
tidyr	* 1.1.2	2020-08-27	[1]	CRAN	(R 4.0.2)
tidyselect	1.1.0	2020-05-11	[1]	CRAN	(R 4.0.2)
tidyverse	* 1.3.0	2019-11-21	[1]	CRAN	(R 4.0.2)
TMB	1.7.18	2020-07-27	[1]	CRAN	(R 4.0.2)
tweenr	1.0.1	2018-12-14	[1]	CRAN	(R 4.0.2)
usethis	1.6.3	2020-09-17	[1]	CRAN	(R 4.0.2)
V8	3.3.1	2020-10-26	[1]	CRAN	(R 4.0.2)
vctrs	0.3.4	2020-08-29	[1]	CRAN	(R 4.0.2)
withr	2.3.0	2020-09-22	[1]	CRAN	(R 4.0.2)
xfun	0.19	2020-10-30	[1]	CRAN	(R 4.0.2)
xml2	1.3.2	2020-04-23	[1]	CRAN	(R 4.0.2)
xtable	1.8-4	2019-04-21	[1]	CRAN	(R 4.0.2)
xts	0.12.1	2020-09-09	[1]	CRAN	(R 4.0.2)
yaml	2.2.1	2020-02-01	[1]	CRAN	(R 4.0.2)
zoo	1.8-8	2020-05-02	[1]	CRAN	(R 4.0.2)

[1] /Library/Frameworks/R.framework/Versions/4.0/Resources/library