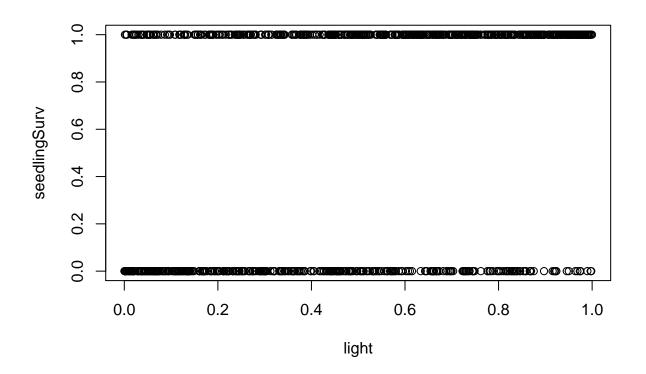
# Bayesian Logistic Regression

Maike Holthuijzen November 30, 2017

First, I used the optim function in R to estimate parameters for the logistic regression model. The function mle.logreg below estimates parameters using maximum likelihood.

```
#simulate data
set.seed(57)
beta0<- -1.0; beta1<-3.0
light<-runif(1000)
pVect<-exp(beta0+beta1*light)/(1+exp(beta0+beta1*light))
range(pVect)

## [1] 0.2690835 0.8806948
seedlingSurv<-rbinom(n=length(light),size=1,prob=pVect)
seedlingSurvData<-data.frame(light,seedlingSurv) # if you want a data frame
plot(light,seedlingSurv)</pre>
```



```
head(seedlingSurvData)
```

```
## light seedlingSurv
## 1 0.24391435 1
```

```
## 2 0.51294954
                            1
## 3 0.03862843
                            1
## 4 0.16617658
                            0
## 5 0.73320525
                            0
## 6 0.66280162
#solve using maximum liklihood
mle.logreg = function(fmla, data)
  # negative log likelihood function
  logl <- function(theta,x,y){</pre>
    у <- у
    x <- as.matrix(x)</pre>
    beta <- theta[1:ncol(x)]</pre>
    # Use the log-likelihood of the Bernoulli distribution, where p is
    # defined as the logistic transformation of a linear combination
    loglik <- sum(-y*log(1 + exp(-(x%*%beta))) - (1-y)*log(1 + exp(x%*%beta)))
    return(-loglik)
  }
  # Prepare the data
  outcome = rownames(attr(terms(fmla), "factors"))[1]
  dfrTmp = model.frame(data)
  x = as.matrix(model.matrix(fmla, data=dfrTmp))
  y = as.numeric(as.matrix(data[,match(outcome,colnames(data))]))
  # Define initial values for the parameters
  theta.start = rep(0, (dim(x)[2]))
  names(theta.start) = colnames(x)
  # Calculate the maximum likelihood
  mle = optim(theta.start,logl,x=x,y=y,hessian=T)
  out = list(beta=mle$par,vcov=solve(mle$hessian),ll=2*mle$value)
}
fmla = as.formula("seedlingSurv~light")
mylogit = mle.logreg(fmla, seedlingSurvData) #Estimate coefficients
```

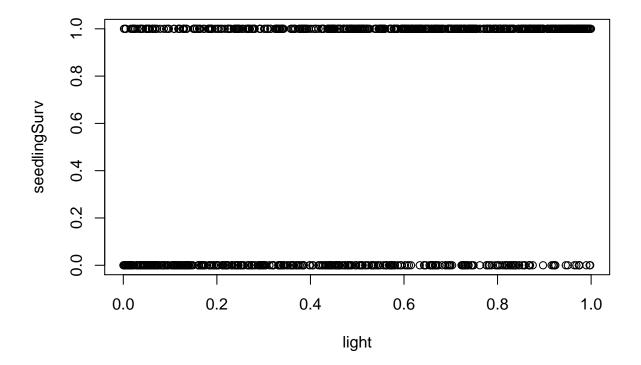
The parameter estimates for the intercept (beta0) and coefficient for  $\beta_1$  are mylogit

```
## $beta
## (Intercept)
                     light
   -0.9653514
                 2.8151933
##
##
## $vcov
##
               (Intercept)
                                  light
## (Intercept) 0.01841652 -0.02908678
               -0.02908678 0.06236447
## light
##
## $11
## [1] 1199.66
```

The estimates are fairly close to the actual values (-1 and 3 for the slope and intercept, respectively.). The

plot below shows a plot of the fitted data.

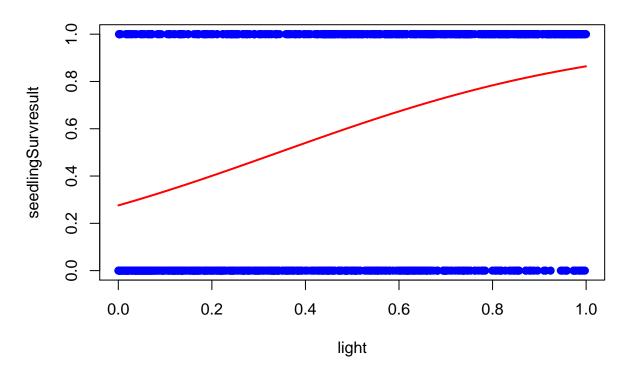
```
lightPreds<-light[order(light)]
plot(light,seedlingSurv)</pre>
```



```
outresult<-exp(mylogit[[1]][1]+mylogit[[1]][2]*light)/(1+exp(mylogit[[1]][1]+mylogit[[1]][2]*light))
seedlingSurvresult<-rbinom(n=length(light),size=1,prob=outresult)

plot(light,seedlingSurvresult, pch = 19, col = "blue", main = "Predicted values for logistic regression
f = function (x) 1 / -(1 + exp(mylogit[[1]][1] + mylogit[[1]][2] * x))
f2 = function (x) exp(mylogit[[1]][1]+mylogit[[1]][2]*x)/(1+exp(mylogit[[1]][1]+mylogit[[1]][2]*x))
lines(lightPreds, f2(lightPreds), col = "red", lwd= 2)</pre>
```

### Predicted values for logistic regression model



We can also compare these results to the output from R's glm function.

```
#compare to glm output
outGlmLogReg = glm(seedlingSurv~light, family=binomial,data=seedlingSurvData)
summary(outGlmLogReg)
##
## Call:
```

```
glm(formula = seedlingSurv ~ light, family = binomial, data = seedlingSurvData)
##
## Deviance Residuals:
##
      Min
                 1Q
                     Median
                                   3Q
                                          Max
## -1.9956 -1.0364
                     0.6186
                               0.9209
                                        1.6033
##
## Coefficients:
##
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) -0.9656
                            0.1357
                                   -7.115 1.12e-12 ***
## light
                 2.8156
                            0.2497
                                   11.274 < 2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
##
  (Dispersion parameter for binomial family taken to be 1)
##
      Null deviance: 1346.0 on 999 degrees of freedom
## Residual deviance: 1199.7 on 998 degrees of freedom
## AIC: 1203.7
##
```

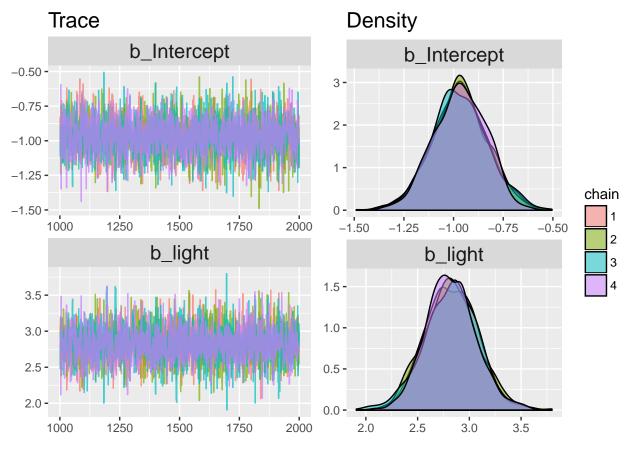
```
Next, I used the brms library for the logistic regression model. Again, the results are about the same as those
obtained from the maximum likelihood estimates.
## Loading required package: ggplot2
## Loading required package: StanHeaders
## rstan (Version 2.16.2, packaged: 2017-07-03 09:24:58 UTC, GitRev: 2e1f913d3ca3)
## For execution on a local, multicore CPU with excess RAM we recommend calling
## rstan_options(auto_write = TRUE)
## options(mc.cores = parallel::detectCores())
## Loading 'brms' package (version 1.1.0). Useful instructions
## can be found by typing help('brms'). A more detailed introduction
## to the package is available through vignette('brms').
## Attaching package: 'brms'
## The following object is masked from 'package:MicrosoftML':
##
##
       categorical
## Compiling the C++ model
## In file included from C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/config.hpp:39:0,
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/math/tools/config.
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/c
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/c
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/c
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/m
##
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math.hpp:4
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/src/stan/model/
##
                    from file3d1428eb7d67.cpp:8:
## C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/config/compiler/gcc.hpp:186:0: warning:
       define BOOST_NO_CXX11_RVALUE_REFERENCES
##
##
## <command-line>:0:0: note: this is the location of the previous definition
## In file included from C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/base.h
                    from C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array.hpp:21
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/numeric/odeint/uti
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/numeric/odeint.hpp
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/prim/
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/prim/
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/prim/s
##
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/m
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math.hpp:4
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/src/stan/model/s
##
##
                    from file3d1428eb7d67.cpp:8:
## C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/concept_checks.hpp: In stati
##
  C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/concept_checks.hpp:42:43: wa
##
          typedef typename Array::index_range index_range;
##
## C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/concept_checks.hpp:43:37: wa
##
          typedef typename Array::index index;
##
```

## Number of Fisher Scoring iterations: 4

```
## C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/concept_checks.hpp: In stati
## C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/concept_checks.hpp:53:43: wa
          typedef typename Array::index_range index_range;
##
##
## C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/concept_checks.hpp:54:37: wa
          typedef typename Array::index index;
##
##
## In file included from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/c
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/m
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math.hpp:4
##
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/src/stan/model/s
                    from file3d1428eb7d67.cpp:8:
##
## C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/core/set_zero_all_adjoi
## C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/core/set_zero_all_adjoi
        static void set_zero_all_adjoints() {
##
##
##
## SAMPLING FOR MODEL 'bernoulli(logit) brms-model' NOW (CHAIN 1).
##
## Gradient evaluation took 0 seconds
## 1000 transitions using 10 leapfrog steps per transition would take 0 seconds.
## Adjust your expectations accordingly!
##
##
## Iteration:
                 1 / 2000 [ 0%]
                                  (Warmup)
## Iteration: 200 / 2000 [ 10%]
                                  (Warmup)
## Iteration: 400 / 2000 [ 20%]
                                  (Warmup)
## Iteration: 600 / 2000 [ 30%]
                                  (Warmup)
## Iteration: 800 / 2000 [ 40%]
                                  (Warmup)
## Iteration: 1000 / 2000 [ 50%]
                                  (Warmup)
## Iteration: 1001 / 2000 [ 50%]
                                  (Sampling)
## Iteration: 1200 / 2000 [ 60%]
                                  (Sampling)
## Iteration: 1400 / 2000 [ 70%]
                                   (Sampling)
## Iteration: 1600 / 2000 [ 80%]
                                   (Sampling)
## Iteration: 1800 / 2000 [ 90%]
                                   (Sampling)
## Iteration: 2000 / 2000 [100%]
                                  (Sampling)
##
##
   Elapsed Time: 0.627 seconds (Warm-up)
##
                  0.568 seconds (Sampling)
##
                  1.195 seconds (Total)
##
##
## SAMPLING FOR MODEL 'bernoulli(logit) brms-model' NOW (CHAIN 2).
##
## Gradient evaluation took 0.001 seconds
## 1000 transitions using 10 leapfrog steps per transition would take 10 seconds.
## Adjust your expectations accordingly!
##
##
## Iteration:
                 1 / 2000 [ 0%]
                                  (Warmup)
## Iteration: 200 / 2000 [ 10%]
                                   (Warmup)
## Iteration: 400 / 2000 [ 20%]
                                   (Warmup)
## Iteration: 600 / 2000 [ 30%]
                                  (Warmup)
## Iteration: 800 / 2000 [ 40%]
```

```
## Iteration: 1000 / 2000 [ 50%]
                                   (Warmup)
## Iteration: 1001 / 2000 [ 50%]
                                   (Sampling)
## Iteration: 1200 / 2000 [ 60%]
                                   (Sampling)
## Iteration: 1400 / 2000 [ 70%]
                                   (Sampling)
## Iteration: 1600 / 2000 [ 80%]
                                   (Sampling)
## Iteration: 1800 / 2000 [ 90%]
                                   (Sampling)
## Iteration: 2000 / 2000 [100%]
                                   (Sampling)
##
##
    Elapsed Time: 0.645 seconds (Warm-up)
##
                  0.75 seconds (Sampling)
##
                  1.395 seconds (Total)
##
##
## SAMPLING FOR MODEL 'bernoulli(logit) brms-model' NOW (CHAIN 3).
##
## Gradient evaluation took 0.001 seconds
## 1000 transitions using 10 leapfrog steps per transition would take 10 seconds.
## Adjust your expectations accordingly!
##
##
## Iteration:
                 1 / 2000 [ 0%]
                                   (Warmup)
## Iteration: 200 / 2000 [ 10%]
                                   (Warmup)
## Iteration: 400 / 2000 [ 20%]
                                   (Warmup)
## Iteration: 600 / 2000 [ 30%]
                                   (Warmup)
## Iteration: 800 / 2000 [ 40%]
                                   (Warmup)
## Iteration: 1000 / 2000 [ 50%]
                                   (Warmup)
## Iteration: 1001 / 2000 [ 50%]
                                   (Sampling)
## Iteration: 1200 / 2000 [ 60%]
                                   (Sampling)
## Iteration: 1400 / 2000 [ 70%]
                                   (Sampling)
## Iteration: 1600 / 2000 [ 80%]
                                   (Sampling)
## Iteration: 1800 / 2000 [ 90%]
                                   (Sampling)
## Iteration: 2000 / 2000 [100%]
                                   (Sampling)
##
##
    Elapsed Time: 0.664 seconds (Warm-up)
##
                  0.6 seconds (Sampling)
##
                  1.264 seconds (Total)
##
##
## SAMPLING FOR MODEL 'bernoulli(logit) brms-model' NOW (CHAIN 4).
##
## Gradient evaluation took 0 seconds
## 1000 transitions using 10 leapfrog steps per transition would take 0 seconds.
## Adjust your expectations accordingly!
##
##
## Iteration:
                 1 / 2000 [ 0%]
                                   (Warmup)
               200 / 2000 [ 10%]
## Iteration:
                                   (Warmup)
## Iteration: 400 / 2000 [ 20%]
                                   (Warmup)
## Iteration:
               600 / 2000 [ 30%]
                                   (Warmup)
               800 / 2000 [ 40%]
## Iteration:
                                   (Warmup)
## Iteration: 1000 / 2000 [ 50%]
                                   (Warmup)
## Iteration: 1001 / 2000 [ 50%]
                                   (Sampling)
## Iteration: 1200 / 2000 [ 60%]
                                   (Sampling)
## Iteration: 1400 / 2000 [ 70%]
                                   (Sampling)
```

```
## Iteration: 1600 / 2000 [ 80%]
                                   (Sampling)
  Iteration: 1800 / 2000 [ 90%]
                                   (Sampling)
   Iteration: 2000 / 2000 [100%]
                                   (Sampling)
##
##
    Elapsed Time: 0.651 seconds (Warm-up)
##
                  0.816 seconds (Sampling)
##
                  1.467 seconds (Total)
##
    Family: bernoulli (logit)
##
  Formula: seedlingSurv ~ light
##
      Data: seedlingSurvData (Number of observations: 1000)
  Samples: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
##
##
            total post-warmup samples = 4000
##
      WAIC: Not computed
##
##
  Population-Level Effects:
             Estimate Est.Error 1-95% CI u-95% CI Eff.Sample Rhat
                                    -1.24
                                             -0.69
                                                          3964
## Intercept
                -0.97
                           0.14
                                                                  1
## light
                 2.82
                           0.25
                                     2.34
                                              3.32
                                                          3471
##
## Samples were drawn using sampling(NUTS). For each parameter, Eff.Sample
## is a crude measure of effective sample size, and Rhat is the potential
## scale reduction factor on split chains (at convergence, Rhat = 1).
```



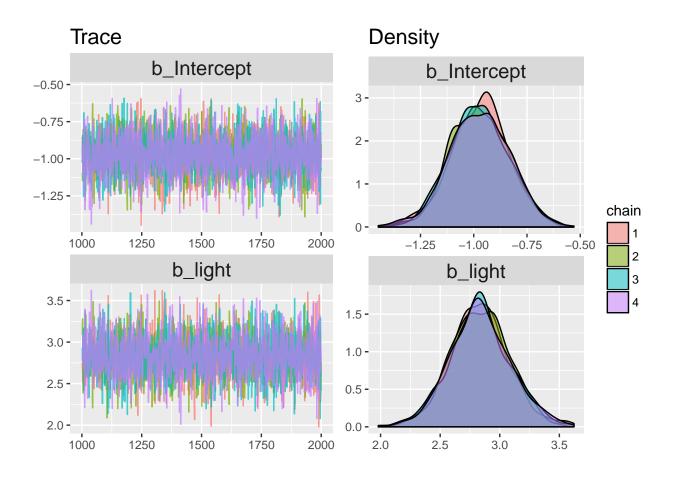
## Compiling the C++ model

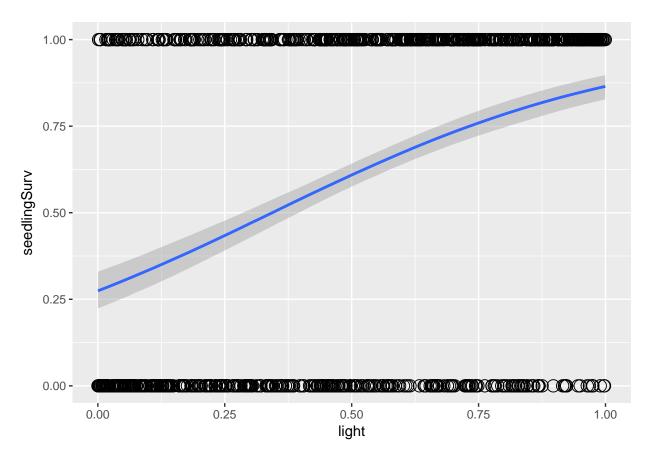
## In file included from C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/config.hpp:39:0,

```
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/math/tools/config.
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/c
##
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/c
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/c
##
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/m
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math.hpp:4
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/src/stan/model/s
##
##
                    from file3d143fd4394a.cpp:8:
  C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/config/compiler/gcc.hpp:186:0: warning:
##
##
       define BOOST_NO_CXX11_RVALUE_REFERENCES
##
  <command-line>:0:0: note: this is the location of the previous definition
  In file included from C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/base.h
                    from \ C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi\_array.hpp:21
##
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/numeric/odeint/uti
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/numeric/odeint.hpp
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/prim/
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/prim/
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/prim/s
##
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/m
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math.hpp:4
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/src/stan/model/
##
##
                    from file3d143fd4394a.cpp:8:
## C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/concept_checks.hpp: In stati
  C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/concept_checks.hpp:42:43: wa
##
          typedef typename Array::index_range index_range;
##
  C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/concept_checks.hpp:43:37: wa
##
##
          typedef typename Array::index index;
##
## C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/concept_checks.hpp: In stati
  C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/concept_checks.hpp:53:43: wa
##
          typedef typename Array::index_range index_range;
##
  C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/concept_checks.hpp:54:37: wa
##
##
          typedef typename Array::index index;
##
## In file included from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/c
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/m
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math.hpp:4
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/src/stan/model/s
##
                    from file3d143fd4394a.cpp:8:
##
## C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/core/set_zero_all_adjoi
  C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/core/set_zero_all_adjoints/
##
        static void set_zero_all_adjoints() {
##
##
  SAMPLING FOR MODEL 'bernoulli(logit) brms-model' NOW (CHAIN 1).
##
## Gradient evaluation took 0 seconds
## 1000 transitions using 10 leapfrog steps per transition would take 0 seconds.
## Adjust your expectations accordingly!
##
##
```

```
1 / 2000 [ 0%]
## Iteration:
                                   (Warmup)
## Iteration: 200 / 2000 [ 10%]
                                   (Warmup)
## Iteration: 400 / 2000 [ 20%]
                                   (Warmup)
## Iteration: 600 / 2000 [ 30%]
                                   (Warmup)
## Iteration: 800 / 2000 [ 40%]
                                   (Warmup)
## Iteration: 1000 / 2000 [ 50%]
                                   (Warmup)
## Iteration: 1001 / 2000 [ 50%]
                                   (Sampling)
## Iteration: 1200 / 2000 [ 60%]
                                   (Sampling)
## Iteration: 1400 / 2000 [ 70%]
                                   (Sampling)
## Iteration: 1600 / 2000 [ 80%]
                                   (Sampling)
## Iteration: 1800 / 2000 [ 90%]
                                   (Sampling)
## Iteration: 2000 / 2000 [100%]
                                   (Sampling)
##
   Elapsed Time: 0.684 seconds (Warm-up)
##
##
                  0.733 seconds (Sampling)
##
                  1.417 seconds (Total)
##
##
## SAMPLING FOR MODEL 'bernoulli(logit) brms-model' NOW (CHAIN 2).
## Gradient evaluation took 0 seconds
## 1000 transitions using 10 leapfrog steps per transition would take 0 seconds.
## Adjust your expectations accordingly!
##
##
## Iteration:
                 1 / 2000 [ 0%]
                                   (Warmup)
## Iteration:
               200 / 2000 [ 10%]
                                   (Warmup)
## Iteration: 400 / 2000 [ 20%]
                                   (Warmup)
## Iteration: 600 / 2000 [ 30%]
                                   (Warmup)
## Iteration: 800 / 2000 [ 40%]
                                   (Warmup)
## Iteration: 1000 / 2000 [ 50%]
                                   (Warmup)
## Iteration: 1001 / 2000 [ 50%]
                                   (Sampling)
## Iteration: 1200 / 2000 [ 60%]
                                   (Sampling)
## Iteration: 1400 / 2000 [ 70%]
                                   (Sampling)
## Iteration: 1600 / 2000 [ 80%]
                                   (Sampling)
## Iteration: 1800 / 2000 [ 90%]
                                   (Sampling)
## Iteration: 2000 / 2000 [100%]
                                   (Sampling)
##
   Elapsed Time: 0.633 seconds (Warm-up)
##
                  0.734 seconds (Sampling)
##
                  1.367 seconds (Total)
##
##
## SAMPLING FOR MODEL 'bernoulli(logit) brms-model' NOW (CHAIN 3).
## Gradient evaluation took 0 seconds
## 1000 transitions using 10 leapfrog steps per transition would take 0 seconds.
## Adjust your expectations accordingly!
##
##
## Iteration:
                 1 / 2000 [ 0%]
                                   (Warmup)
## Iteration: 200 / 2000 [ 10%]
                                   (Warmup)
## Iteration: 400 / 2000 [ 20%]
                                   (Warmup)
## Iteration: 600 / 2000 [ 30%]
                                   (Warmup)
```

```
## Iteration: 800 / 2000 [ 40%]
                                   (Warmup)
## Iteration: 1000 / 2000 [ 50%]
                                   (Warmup)
## Iteration: 1001 / 2000 [ 50%]
                                   (Sampling)
## Iteration: 1200 / 2000 [ 60%]
                                   (Sampling)
## Iteration: 1400 / 2000 [ 70%]
                                   (Sampling)
## Iteration: 1600 / 2000 [ 80%]
                                   (Sampling)
## Iteration: 1800 / 2000 [ 90%]
                                   (Sampling)
## Iteration: 2000 / 2000 [100%]
                                   (Sampling)
##
##
    Elapsed Time: 0.626 seconds (Warm-up)
                  0.517 seconds (Sampling)
                  1.143 seconds (Total)
##
##
##
## SAMPLING FOR MODEL 'bernoulli(logit) brms-model' NOW (CHAIN 4).
##
## Gradient evaluation took 0 seconds
## 1000 transitions using 10 leapfrog steps per transition would take 0 seconds.
## Adjust your expectations accordingly!
##
##
## Iteration:
                 1 / 2000 [ 0%]
                                   (Warmup)
## Iteration: 200 / 2000 [ 10%]
                                   (Warmup)
## Iteration: 400 / 2000 [ 20%]
                                   (Warmup)
## Iteration: 600 / 2000 [ 30%]
                                   (Warmup)
## Iteration: 800 / 2000 [ 40%]
                                   (Warmup)
## Iteration: 1000 / 2000 [ 50%]
                                   (Warmup)
## Iteration: 1001 / 2000 [ 50%]
                                   (Sampling)
## Iteration: 1200 / 2000 [ 60%]
                                   (Sampling)
## Iteration: 1400 / 2000 [ 70%]
                                   (Sampling)
## Iteration: 1600 / 2000 [ 80%]
                                   (Sampling)
## Iteration: 1800 / 2000 [ 90%]
                                   (Sampling)
## Iteration: 2000 / 2000 [100%]
                                   (Sampling)
##
##
    Elapsed Time: 0.659 seconds (Warm-up)
##
                  0.815 seconds (Sampling)
##
                  1.474 seconds (Total)
   Family: bernoulli (logit)
## Formula: seedlingSurv ~ light
      Data: seedlingSurvData (Number of observations: 1000)
## Samples: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
            total post-warmup samples = 4000
##
##
      WAIC: Not computed
##
## Population-Level Effects:
             Estimate Est.Error 1-95% CI u-95% CI Eff.Sample Rhat
                                             -0.71
## Intercept
                -0.97
                           0.14
                                    -1.24
                                                          3712
                                                                  1
                 2.83
                           0.25
                                     2.34
                                              3.34
                                                          3181
## light
##
## Samples were drawn using sampling(NUTS). For each parameter, Eff.Sample
## is a crude measure of effective sample size, and Rhat is the potential
## scale reduction factor on split chains (at convergence, Rhat = 1).
```





Then, I used the stanarm library. Again, results were similar to the previous approaches.

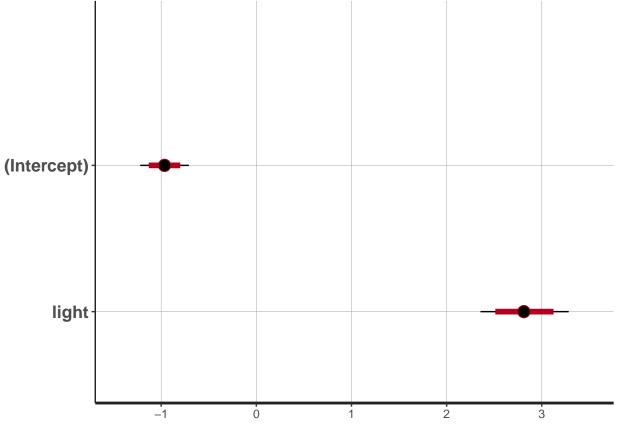
library(rstanarm)

summary(outrstanarmLogReg)

```
## stan_glm(formula = seedlingSurv ~ light, family = binomial, data = seedlingSurvData,
## iter = 2000, warmup = 1000, cores = 4)
```

## trying deprecated constructor; please alert package maintainer

```
##
## Family: binomial (logit)
## Algorithm: sampling
## Posterior sample size: 4000
## Observations: 1000
##
## Estimates:
##
                                 2.5%
                                        25%
                                                50%
                                                       75%
                                                              97.5%
                   mean
                          sd
## (Intercept)
                   -1.0
                           0.1
                                 -1.2
                                         -1.1
                                                -1.0
                                                       -0.9
                                                              -0.7
## light
                    2.8
                           0.2
                                  2.4
                                         2.7
                                                 2.8
                                                        3.0
                                                               3.3
## mean_PPD
                    0.6
                           0.0
                                  0.6
                                         0.6
                                                 0.6
                                                        0.6
                                                               0.6
                           0.9 -607.7 -605.4 -604.8 -604.5 -604.2
## log-posterior -605.1
## Diagnostics:
##
                 mcse Rhat n_eff
## (Intercept)
                 0.0 1.0 3852
## light
                 0.0 1.0 3369
## mean_PPD
                 0.0 1.0
                           3554
## log-posterior 0.0 1.0 2197
## For each parameter, mcse is Monte Carlo standard error, n_eff is a crude measure of effective sample
plot(outrstanarmLogReg)
## ci_level: 0.8 (80% intervals)
## outer_level: 0.95 (95% intervals)
```



Next, I used actual Stan code within the rstan library.

```
## In file included from C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/config.hpp:39:0,
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/math/tools/config.
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/c
##
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/c
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/c
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/m
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math.hpp:4
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/src/stan/model/s
##
##
                    from file3d1412be627d.cpp:8:
## C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/config/compiler/gcc.hpp:186:0: warning:
##
       define BOOST_NO_CXX11_RVALUE_REFERENCES
##
## <command-line>:0:0: note: this is the location of the previous definition
  In file included from C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/base.h
##
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array.hpp:21
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/numeric/odeint/uti
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/numeric/odeint.hpp
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/prim/
##
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/prim/
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/prim/
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/m
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math.hpp:4
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/src/stan/model/s
##
##
                    from file3d1412be627d.cpp:8:
## C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/concept_checks.hpp: In stati
  C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/concept_checks.hpp:42:43: wa
##
          typedef typename Array::index_range index_range;
##
##
  C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/concept_checks.hpp:43:37: wa
##
          typedef typename Array::index index;
##
## C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/concept_checks.hpp: In stati
  C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/concept_checks.hpp:53:43: wa
##
          typedef typename Array::index_range index_range;
##
##
  C:/Users/Maike/Documents/R/win-library/3.3/BH/include/boost/multi_array/concept_checks.hpp:54:37: wa
##
          typedef typename Array::index index;
##
## In file included from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/c
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/m
##
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math.hpp:4
##
                    from C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/src/stan/model/
##
                    from file3d1412be627d.cpp:8:
  C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/core/set_zero_all_adjoi
  C:/Users/Maike/Documents/R/win-library/3.3/StanHeaders/include/stan/math/rev/core/set_zero_all_adjoints/
##
        static void set_zero_all_adjoints() {
##
## SAMPLING FOR MODEL 'fe004f478f8ad74110963f3663c16e41' NOW (CHAIN 1).
##
## Gradient evaluation took 0 seconds
## 1000 transitions using 10 leapfrog steps per transition would take 0 seconds.
```

```
## Adjust your expectations accordingly!
##
##
## Iteration:
                 1 / 3000 [ 0%]
                                  (Warmup)
## Iteration: 300 / 3000 [ 10%]
                                   (Warmup)
## Iteration: 501 / 3000 [ 16%]
                                   (Sampling)
## Iteration: 800 / 3000 [ 26%]
                                   (Sampling)
## Iteration: 1100 / 3000 [ 36%]
                                   (Sampling)
## Iteration: 1400 / 3000 [ 46%]
                                   (Sampling)
## Iteration: 1700 / 3000 [ 56%]
                                   (Sampling)
## Iteration: 2000 / 3000 [ 66%]
                                   (Sampling)
## Iteration: 2300 / 3000 [ 76%]
                                   (Sampling)
## Iteration: 2600 / 3000 [ 86%]
                                   (Sampling)
## Iteration: 2900 / 3000 [ 96%]
                                   (Sampling)
## Iteration: 3000 / 3000 [100%]
                                   (Sampling)
##
##
   Elapsed Time: 0.668 seconds (Warm-up)
##
                  2.751 seconds (Sampling)
##
                  3.419 seconds (Total)
##
##
## SAMPLING FOR MODEL 'fe004f478f8ad74110963f3663c16e41' NOW (CHAIN 2).
##
## Gradient evaluation took 0 seconds
## 1000 transitions using 10 leapfrog steps per transition would take 0 seconds.
## Adjust your expectations accordingly!
##
## Iteration:
                 1 / 3000 [ 0%]
                                   (Warmup)
## Iteration: 300 / 3000 [ 10%]
                                   (Warmup)
## Iteration: 501 / 3000 [ 16%]
                                   (Sampling)
## Iteration: 800 / 3000 [ 26%]
                                   (Sampling)
## Iteration: 1100 / 3000 [ 36%]
                                   (Sampling)
## Iteration: 1400 / 3000 [ 46%]
                                   (Sampling)
## Iteration: 1700 / 3000 [ 56%]
                                   (Sampling)
## Iteration: 2000 / 3000 [ 66%]
                                   (Sampling)
## Iteration: 2300 / 3000 [ 76%]
                                   (Sampling)
## Iteration: 2600 / 3000 [ 86%]
                                   (Sampling)
## Iteration: 2900 / 3000 [ 96%]
                                   (Sampling)
## Iteration: 3000 / 3000 [100%]
                                   (Sampling)
##
##
   Elapsed Time: 0.597 seconds (Warm-up)
                  2.485 seconds (Sampling)
##
##
                  3.082 seconds (Total)
##
##
## SAMPLING FOR MODEL 'fe004f478f8ad74110963f3663c16e41' NOW (CHAIN 3).
##
## Gradient evaluation took 0 seconds
## 1000 transitions using 10 leapfrog steps per transition would take 0 seconds.
## Adjust your expectations accordingly!
##
##
                 1 / 3000 [ 0%] (Warmup)
## Iteration:
```

```
## Iteration: 300 / 3000 [ 10%]
                                  (Warmup)
                                  (Sampling)
## Iteration: 501 / 3000 [ 16%]
## Iteration: 800 / 3000 [ 26%]
                                  (Sampling)
## Iteration: 1100 / 3000 [ 36%]
                                  (Sampling)
## Iteration: 1400 / 3000 [ 46%]
                                   (Sampling)
## Iteration: 1700 / 3000 [ 56%]
                                  (Sampling)
## Iteration: 2000 / 3000 [ 66%]
                                   (Sampling)
## Iteration: 2300 / 3000 [ 76%]
                                   (Sampling)
## Iteration: 2600 / 3000 [ 86%]
                                   (Sampling)
## Iteration: 2900 / 3000 [ 96%]
                                   (Sampling)
## Iteration: 3000 / 3000 [100%]
                                  (Sampling)
##
   Elapsed Time: 0.583 seconds (Warm-up)
##
                  2.767 seconds (Sampling)
##
                  3.35 seconds (Total)
## $summary
##
               mean
                        se_mean
                                       sd
                                               2.5%
                                                           25%
                                                                      50%
## beta0 -0.9722504 0.004824192 0.1321159 -1.224641 -1.064865 -0.9703983
## beta1 2.8291668 0.008657310 0.2370902 2.384171 2.662065 2.8343223
                        97.5% n_eff
                75%
                                         Rhat
## beta0 -0.8821296 -0.722219
                                750 0.9980000
## beta1 2.9876787 3.267216
                               750 0.9986143
##
## $c_summary
## , , chains = chain:1
##
##
            stats
                                       2.5%
                                                  25%
                                                              50%
                                                                         75%
## parameter
                               sd
                   mean
##
       beta0 -0.9691986 0.1327635 -1.215911 -1.067099 -0.9699606 -0.8671854
       beta1 2.8275144 0.2411156 2.367019 2.656022 2.8317811 2.9974880
##
##
            stats
## parameter
                 97.5%
       beta0 -0.750219
##
       beta1 3.283041
##
##
  , , chains = chain:2
##
            stats
## parameter
                               sd
                                       2.5%
                                                  25%
                                                              50%
                   mean
       beta0 -0.9753559 0.1332142 -1.213840 -1.070007 -0.9757403 -0.8862542
##
       beta1 2.8367076 0.2375048 2.379688 2.669312 2.8282827 2.9955967
##
##
            stats
##
  parameter
                  97.5%
##
       beta0 -0.6999382
##
       beta1 3.2781272
##
##
   , , chains = chain:3
##
##
            stats
##
                               sd
                                       2.5%
                                                  25%
                                                              50%
                                                                        75%
  parameter
                   mean
       beta0 -0.9721967 0.1308162 -1.256152 -1.050601 -0.9640484 -0.886754
##
##
       beta1 2.8232784 0.2333386 2.411116 2.659894 2.8413074 2.968733
##
            stats
```

```
## parameter 97.5%
## beta0 -0.7258352
## beta1 3.2409186
## Warning: package 'coda' was built under R version 3.3.3
##
## Attaching package: 'coda'
## The following object is masked from 'package:rstan':
## traceplot
```

Lastly, I used the rjags library to fit the logistic regression model. The code is similar to the rstan model, but it is a little simpler. Again, the posterior distributions for the slope and intercept parameters show a high density around the actual parameters, and the trace plot (obtained via the coda library) shows that mixing within the MCMC chain was also sufficient

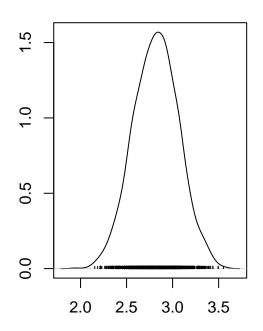
```
## Warning: package 'rjags' was built under R version 3.3.3
## Linked to JAGS 4.3.0
## Loaded modules: basemod, bugs
## Compiling model graph
##
      Resolving undeclared variables
##
      Allocating nodes
## Graph information:
      Observed stochastic nodes: 1000
##
##
      Unobserved stochastic nodes: 2
##
      Total graph size: 5005
##
## Initializing model
## $beta0
## mcarray:
## [1] -0.9746154
## Marginalizing over: iteration(1000), chain(4)
##
## $beta1
## mcarray:
## [1] 2.835911
## Marginalizing over: iteration(1000), chain(4)
```

## Density of beta0

# -1.4 -1.0 -0.6

N = 1000 Bandwidth = 0.02647

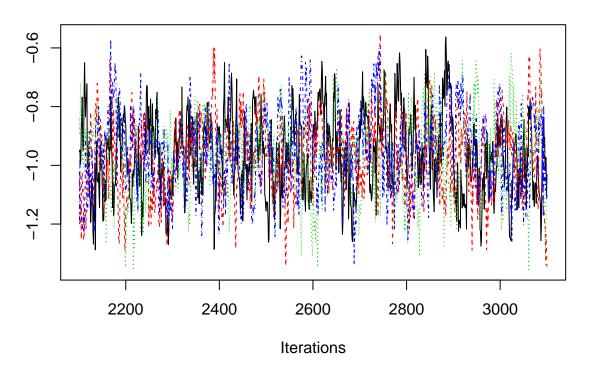
## **Density of beta1**



N = 1000 Bandwidth = 0.04933

```
##
## Iterations = 2101:3100
## Thinning interval = 1
## Number of chains = 4
## Sample size per chain = 1000
##
## 1. Empirical mean and standard deviation for each variable,
      plus standard error of the mean:
##
##
                     SD Naive SE Time-series SE
##
            Mean
## beta0 -0.9651 0.1312 0.002074
                                       0.006165
## beta1 2.8195 0.2445 0.003866
                                       0.011588
##
## 2. Quantiles for each variable:
##
                           50%
##
           2.5%
                   25%
                                   75%
                                         97.5%
## beta0 -1.216 -1.057 -0.9657 -0.8732 -0.7055
## beta1 2.338 2.651 2.8214 2.9864 3.3054
```

# Trace of beta0



# Trace of beta1

