Building General Random Effect Models with R and ADMB

Jeff Laake, Hans Skaug and Devin Johnson January 26, 2013

The ADMB and Otter Research web sites provide examples of specific random effect models. See http://admb-project.org/examples and http://www.admb-project.org/examples/miscellaneous/otter-reseach-collection. However, none of these examples discuss how to build more general TPL files to accommodate a class of random effect models. The R package glmmadmb (http://glmmadmb.r-forge.r-project.org/) is an example where R is used with a single TPL file that can fit generalized linear mixed models for a wide range of distributions. By examining the R code and TPL file for glmmadmb, you can work out how to develop a general TPL file and use R code to build the structures ADMB needs to fit mixed effect models. However, glmmadmb is fairly complex and sorting through the code can be challenging. Here we provide a simpler example with comments to demonstrate the process so others can incorporate this into their model building software.

We use the general modeling tools in R to create a DAT file that contains the structures needed for the TPL file for ADMB. We use the R2admb package as the R and ADMB interface. R and R2admb could be replaced with equivalent software that builds the DAT file but R is freely available and has a wealth of modeling and data manipulation tools. Using mixed logistic regression as an example, we'll begin by describing the TPL file structure to fit a random effects model for a user-specified dataframe and formula.

ADMB Code

We will describe the contents of the file mixed.tpl given in the Appendix. This simple skeleton file could be modified to compute a different likelihood or could be extended to incorporate mixed-effects modeling for several parameters contained in

a single likelihood. The TPL file is generalized to work with any data file or formula by specifying the model via design matrices that are passed as data. For mixed effect models, there is a design matrix for the fixed effect and one for the random effects. In addition, there is a vector of indices for the random effects that are passed as data as well.

We'll describe the DAT structure with a simple illustrative example that is too small to use in practice. The variable y is the response and id, sex and time are all factor variables.

У	id	sex	time
0	1	F	1
1	1	\mathbf{F}	2
0	2	Μ	1
1	2	Μ	2
1	3	\mathbf{F}	1
0	3	F	2

Consider building a logistic regression model for the response y in which sex is a fixed effect and id (individual heterogeneity) is a random intercept for the response. We begin by describing the DATA_SECTION of the TPL file and show the contents below of the DAT file for this example. The variables in parentheses are the names used in the TPL file (Appendix) and we have used underscore to show the range of the vector or matrix (e.g. y_i, i=1 to n):

```
# number of rows in data (n)
# vector of 6 responses (y_i, i=1 to n)
0 1 0 1 1 0
# number of columns in the fixed effect design matrix (kfixed)
2
# fixed effect design matrix (fixedDM_ij; i=1 to n; j=1 to kfixed) (first column is
intercept; second column is male effect)
10
10
1 1
1 1
10
10
# number of random effect values (nre) (for this model, one for each id)
3
# number of columns in the random effect design matrix (krand)
# random effect design matrix (randDM_ij; i=1 to n; j=1 to krand)
1
1
1
1
1
# random effect indices (randIndex_ij; i=1 to n; j=1 to krand)(values from 1 to nre)
1
1
2
2
3
3
```

There is a single random effect variable with nre=3 values (ie. one for each of the 3 individuals). The indices specify which random effect is used for the value in the random design matrix. For this model there are 3 parameters (kfixed(2) + krand(1)): Intercept and Male fixed effects and sigma (standard deviation) for id random effect. The likelihood is:

$$L(\beta_1, \beta_2, \tau_1, \boldsymbol{u} | \boldsymbol{y}) = \prod_{i=1}^{6} p_i^{y_i} (1 - p_i)^{(1-y_i)}$$

Below we specify the value of each p_i (as logit) in terms of the parameters and the 3 random effect values u_1, u_2, u_3 which are standard normal random variables. The variable mu (μ) is the logit of p_i and we compute the log of the likelihood which simplifies to $\sum_{i=1}^{6} [y_i \mu_i + \ln(1 + \exp(\mu_i))]$.

$logit(p_1) = \mu_1 =$	$\beta_1 + u_1 e^{\tau_1}$
$logit(p_2) = \mu_2 =$	$\beta_1 + u_1 e^{\tau_1}$
$logit(p_3) = \mu_3 =$	$\beta_1 + \beta_2 + u_2 e^{\tau_1}$
$logit(p_4) = \mu_4 =$	$\beta_1 + \beta_2 + u_2 e^{\tau_1}$
$logit(p_5) = \mu_5 =$	$\beta_1 + u_3 e^{\tau_1}$
$logit(p_6) = \mu_6 =$	$\beta_1 + u_3 e^{\tau_1}$

The PARAMETER_SECTION contains a vector Beta which are the values of the fixed parameters from 1 to kfixed that are estimated in phase 1. It also contains a vector Tau which are the random parameters from 1 to krand and a random_effects_vector from 1 to nre. The random effect parameters and values are estimated in phase 2. If there is no random component in the model, then the phase is set to -1 so it will be ignored and a fixed effects model is fitted. The negative log-likelihood value (objective_function_value) is held in g.

The PROCEDURE_SECTION of the TPL file for computation of the objective function is fairly simple. It contains a loop over each of the nre random effects that calls the SEPARABLE_FUNCTION no1_prior which sums the negative log-likelihood of a standard normal distribution. That loop is only called if there are any random effects. The next loop is over each row in the data and it calls the SEPARABLE_FUNCTION 11_i to compute and sum the negative log-likelihood contribution for each response. In the call to 11_i you must pass the parameters and the random effects used in the calculation. Notice that the code specifies u(randIndex(i)) which passes only the random effect values needed for the each row in the data. When used with the -shess run argument this improves execution speed (Skaug and Fournier 2011).

The SEPARABLE_FUNCTION 11_i computes the value of mu which is fixedDM(i)*Beta plus randDM(i,j)*u(j)*mfexp(Tau(j)) for each column in the random design matrix (krand). For our simple example, krand=1 and randDM(i,j)=1, so the code simplifies to the calculations shown above.

Now let's expand on the example with a model that has both an individual and temporal random intercept. The TPL file is the same and only the DAT file changes. Now we have nre=5 random values (3 for id and 2 for time) and we have krand=2 columns in the random design matrix, one for the intercept of each random effect.

```
# number of rows in data (n)
# vector of 6 responses (y_i, i=1 to n)
010110
# number of columns in the fixed effect design matrix (kfixed)
# fixed effect design matrix (fixedDM_ij; i=1 to n; j=1 to kfixed) (first column is
intercept; second column is male effect)
10
10
1 1
1 1
10
10
# number of random effect values (nre) (for this model, one for each id)
# number of columns in the random effect design matrix (krand)
# random effect design matrix (randDM_ij; i=1 to n; j=1 to krand)
1 1
1 1
1 1
1 1
1 1
1 1
# random effect indices (randIndex_ij; i=1 to n; j=1 to krand)(values from 1 to nre)
14
15
2 4
2 5
3 4
3 5
```

The important design aspect is the use of a single random effect vector to contain the values for each type of random effect (e.g. id and time). This is important because random effects must be passed as arguments to the SEPARABLE_FUNCTION. Had we decided to use a different vector for each type of random effect, we would have to change the definition of the SEPARABLE_FUNCTION and its call in the PROCEDURE_SECTION. Modification of the code is avoided by using a single vector and using indices to specify which to use with the random design matrix values. For this model there are 4 parameters $(\beta_1, \beta_2, \tau_1, \tau_2)$ and the values of mu are:

$\log it(p_1) = \mu_1 =$	$\beta_1 + u_1 e^{\tau_1} + u_4 e^{\tau_2}$
$logit(p_2) = \mu_2 =$	$\beta_1 + u_1 e^{\tau_1} + u_5 e^{\tau_2}$
$logit(p_3) = \mu_3 =$	$\beta_1 + \beta_2 + u_2 e^{\tau_1} + u_4 e^{\tau_2}$
$logit(p_4) = \mu_4 =$	$\beta_1 + \beta_2 + u_2 e^{\tau_1} + u_5 e^{\tau_2}$
$logit(p_5) = \mu_5 =$	$\beta_1 + u_3 e^{\tau_1} + u_4 e^{\tau_2}$
$logit(p_6) = \mu_6 =$	$\beta_1 + u_3 e^{\tau_1} + u_5 e^{\tau_2}$

R Code

Now you certainly would not want to construct the DAT file by hand for anything other than a simple illustrative example like this one. It would be tedious and error prone. This is where R and the R package R2admb become an invaluable tool. We have used functions in R listed in the Appendix: 1) proc.form - parses a formula using the structure specified in the R package lme4, 2) mixed.model.admb - calls proc.form with a formula and then with a data frame constructs the design matrices and indices needed for the DAT file, and 3) mixed.model.dat - writes the model components structures to the DAT file. The function mixed.model.dat only writes out a mixed effect model structure and not the complete DAT file. We chose to do this because it may be included as part of a DAT file that may have other components.

Below are code that construct a simulated data set, fit a mixed-effects logistic model with ADMB using the code described and then fit the same model with the function lmer in the package lme4 to demonstrate that the same results are obtained.

- > # Load R2admb package from the library
- > library(R2admb)
- > # Set environment variables PATH and ADMB_HOME to enable ADMB connection

```
> Sys.setenv(PATH = paste("c:/admb/bin;c:admb/utilities;c:/MinGW/bin;",
                 Sys.getenv("PATH"), sep = ";"))
> Sys.setenv(ADMB_HOME = "c:/admb")
> # create simulated data with 2 factor variables f and g with
> # N(0,.3) and N(0,1) random effects
> fn=rnorm(10,0,.3)
> gn=rnorm(20,0,1)
> data=cbind(data.frame(y=rbinom(400,1,
        p=plogis(log(.3/.7)+rowSums(expand.grid(fn,gn))))),
        rbind(expand.grid(f=factor(1:10),g=factor(1:20)),
        expand.grid(f=factor(1:10),g=factor(1:20))))
> # Use R2admb functions to compile the ADMB TPL file
> compile_admb("mixed",re=TRUE)
> # Remove any leftover files
> clean_admb("mixed")
> # Create a connection to a file mixed.dat (DAT file)
> con=file("mixed.dat",open="wt")
> # Write out n and responses y to DAT file
> write(nrow(data),con,append=FALSE)
> write(data$y,con,append=TRUE)
> clean_admb("mixed")
> con=file("mixed.dat",open="wt")
> write(nrow(data),con,append=FALSE)
> write(data$y,con,append=TRUE)
> # Write mixed model structures to DAT file
> mixed.model.dat(mixed.model.admb(~1 +(1|f) +(1|g),data),con)
> # Close connection
> close(con)
> # Run program
> run_admb("mixed",extra.args="-shess")
> # Read results into R object
> results=read_admb("mixed")
> results
Model file: mixed
Negative log-likelihood: 230.449
Coefficients:
     Beta
             Tau.1
                       Tau.2
-0.904190 -0.743352 0.038662
```

```
> # The value of exp(Tau)^2 should match the variances from lmer
> exp(results$coeflist$Tau*2)
[1] 0.2261167 1.0803921
> # Compare to lmer from lme4
> library(lme4)
> lme4res=lmer(y~1 +(1|f) +(1|g),data,family="binomial",REML=FALSE)
> lme4res
Generalized linear mixed model fit by the Laplace approximation
Formula: y ~ 1 + (1 | f) + (1 | g)
  Data: data
        BIC logLik deviance
  AIC
 466.9 478.9 -230.4
                      460.9
Random effects:
 Groups Name
                   Variance Std.Dev.
        (Intercept) 1.08039 1.03942
        (Intercept) 0.22613 0.47553
Number of obs: 400, groups: g, 20; f, 10
Fixed effects:
            Estimate Std. Error z value Pr(>|z|)
(Intercept) -0.9042 0.3020 -2.994 0.00275 **
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
>
```

References

[1] Skaug, H. and D. Fournier. 2011. Random effects in AD Model Builder: ADMB-RE User Guide. Version 10.0. http://admb-project.googlecode.com/files/admbre-10.0-rev1.pdf.

9

```
// Skeleton template for a mixed effects model structure; with logistic regression likelihood
// Jeff Laake; 22 Jan 2013
DATA_SECTION
    init_int n;
                                             // number of rows in data
    init_vector y(1,n);
                                             // vector of responses
    init_int kfixed:
                                            // number of columns in design matrix(DM) for fixed effects // fixed effect DM
    init_matrix fixedDM(1,n,1,kfixed);
    init_int nre;
                                            // number of random effects
    int phase;
                                            // if no random effects set phase to -1 otherwise 2
         !! phase=2;
        !! if (nre==0)phase=-1;
    init_int krand;
                                            // number of columns in random effect DM
    init_matrix randDM(1.n.1.krand):
                                            // random effect DM
// random effect indices for DM
    init_imatrix randIndex(1,n,1,krand);
PARAMETER_SECTION
    init_vector Beta(1, kfixed, 1);
                                            // parameter vector for fixed effects
                                            // parameter vector for log(sigma)
    init_vector Tau(1, krand, phase);
    objective_function_value g;
                                            // objective function - negative log-likelihood
    random_effects_vector u(1, nre, phase); // random effects vector
PROCEDURE_SECTION
                                                             // index variable
// initialize negative log-likelihood
    int i;
    g=0;
        cout <<"nre = "<< nre <<endl;
        cout << "kfixed = "<< kfixed << endl;
        cout << "krand = "<< krand << endl;
    for (i=1;i<=nre;i++)
                                                              // if any random effects compute likelihood contribution for each
        n01_prior(u(i));
                                                              // u's are N(0,1) distributed
        cout << "g = "<< g << endl;
                                                              // loop over rows in data
    for (i = 1; i \le n; i ++)
          ll_i (i, Beta, Tau, u(randIndex(i)));
                                                              //compute negative log-likelihood for each row with separable function
        cout << "g = "<< g << endl;
SEPARABLE-FUNCTION void n01_prior(const prevariable& u) // taken from glmmadmb.tpl; uses PI
 g = -0.5*\log(2.0*PI) - 0.5*square(u);
SEPARABLE_FUNCTION void 11_i (const int i, const dvar_vector& Beta, const dvar_vector& Tau, const dvar_vector& u)
    int j;
    dvariable mu;
                                                                 link value
    mu=fixedDM(i)*Beta;
                                                              // fixed portion
    for (j=1; j \le krand; j++)
                                                              // random portion (if any)
        mu+=randDM(i,j)*u(j)*mfexp(Tau(j));
    g=y(i)*mu - log(1+exp(mu));
                                                             // Bernoulli likelihood with logit link
```

Appendix - proc.form, mixed.model.admb, mixed.model.dat

```
Mixed effect model formula parser
   Parses a mixed effect model in the lme4 structure of fixed +(re1|g1) +...+(ren|gn)
   @param f formula for mixed effect mode in the form used in lme4; ~fixed +(re1|g1) +...+(ren|gn)
   @return A list with elements fix.model and re.model. fix.model contains the formula for the fixed effects;
  re.model contains elements sub, the grouping formula and model the design formula for the
# random effect. Each formula is of type character and must be wrapped with as formula in use with model matrix
#' @author Devin Johnson <devin.johnson@@noaa.gov>
proc.form <- function(f){
         tms <- terms(f)
         tms.lab <- attr(tms, "term.labels")
tms.lst <- strsplit(tms.lab, rep(" | ",length(tms.lab)), fixed=TRUE)
         fix.var <- attr(tms, "term.labels")[sapply(tms.lst, "length")==1]
         if (length (fix.var)==0)
fix.model="~1"
                   \label{eq:fix.model} \textit{fix.model} <- \ \textit{paste}("`", \textit{paste}(\textit{fix.var}, \ \textit{collapse}=""+"))
         {\tt re.lst} \; \leftarrow \; {\tt tms.lst} \; [\; {\tt sapply} \; (\; {\tt tms.lst} \; , \; \; "\; {\tt length}") {\tt ==} 2]
         if(length(re.lst)==0) re.model <- NULL
         else {
                   re.model \leftarrow lapply (re.lst, function(x) \{list(model=paste(""",x[1]), sub=paste(""",x[2],"-1", collapse=""))\})
         return(list(fix.model=fix.model, re.model=re.model))
```

```
Mixed effect model construction
   Functions that develop structures needed for a mixed effect model
#'
   mixed.model.admb - creates design matrices and supporting index matrices
   for use of mixed model in ADMB
#'
   mixed.model.dat - writes to data file (con) for fixed and random effect stuctures
   @aliases mixed.model.admb mixed.model.dat
   Qusage mixed.model.admb(formula,data)
          mixed.model.dat(x,con)
   @param formula formula for mixed effect mode in the form used in lme4; fixed +(re1|g1) +...+(ren|gn)
   @param data dataframe used to construct the design matrices from the formula
   @param x list structure created by mixed.model.admb
   @param con connection to data file which contents will be appended
   @return mixed.model.admb returns a list with elements fixed.dm, the design matrix for
   the fixed effects; re.dm, a combined design matrix for all of the random effects; and
  re.indices, matrix of indices into a single vector of random effects to be applied to the
  design matrix location.
#' @author Jeff Laake < jeff.laake@@noaa.gov>
mixed.model.admb=function(formula, data)
# parse formula for fixed and random effects
  mlist=proc.form(formula)
# construct design matrix for fixed effects
  fixed.dm=model.matrix(as.formula(mlist$fix.model),data)
# remainder of code is for random effects unless NULL
  re.dm=NULL
  re.indices=NULL
  if (!is.null(mlist$re.model))
#
     Loop over each random effect component
     for (i in 1:length (mlist$re.model))
       Make sure each variable used to define random effect group is a factor variable
#
           if (! all(sapply(model.frame(as.formula(mlist$re.model[||i|]| sub),data),is.factor)))
               stop(paste("\n one or more variables in", mlist$re.model[[i]]$sub," is not a factor variable\n"))
          Compute design matrix for grouping variables
          zz=model.matrix(as.formula(mlist$re.model[[i]]$sub),data)
          Not all combinations of factor variable(s) may be used so only use those observed in the data
#
                   used.columns=which(colSums(zz)>0)
                   nre=length (used.columns)
#
          Compute the indices for this particular grouping structure and reindex if any missing
          indices=rowSums(zz*col(zz))
                   if (nre!=ncol(zz)) indices=match (indices, used.columns)
          Compute the design matrix for the random effect formula
          zz=model.matrix(as.formula(mlist$re.model[[i]]$model),data)
          Now shift indices to refer to a single vector of random effects across all re groupings
                   ng=max(indices)
          \verb|indices| = \verb|matrix| (\verb|indices|), \verb|nrow| = \verb|length| (\verb|indices|), \verb|ncol| = \verb|ncol| (\verb|zz|) + \verb|reindex|
          indices=t(t(indices)+cumsum(c(0,rep(ng,ncol(zz)-1))))
          reindex=max(indices)
          Bind random effect design matrices (re.dm), indices into random effects vector (re.indices) and
          index for the random effect sigma parameter (re.sigma)
                   re.dm=cbind(re.dm,zz)
```

```
re.indices=cbind(re.indices, indices)
   return(list(fixed.dm=fixed.dm,re.dm=re.dm,re.indices=re.indices))
mixed.model.dat=function(x,con)
         # number of columns of fixed dm
         write (ncol(x$fixed.dm),con,append=TRUE)
         # fixed dm
         \label{eq:write} \begin{array}{l} \text{write} \left( \text{t} \left( \text{x\$fixed.dm} \right), \text{con}, \text{ncolumns=ncol} \left( \text{x\$fixed.dm} \right), \text{append=TRUE} \right) \end{array}
    if (!is.null(x$re.dm))
                  # number of random effects
                  # number of columns of re dm
                  write(ncol(x$re.dm),con,append=TRUE)
                  # re dm
                  write(t(x$re.dm),con,ncolumns=ncol(x$re.dm),append=TRUE)
                  # re indices
                  write(t(x$re.indices),con,ncolumns=ncol(x$re.indices),append=TRUE)
         }
else
                  \# number of re =0
                   write (0, con, append=TRUE)
                  # number of columns of re dm=0
write(0,con,append=TRUE)
    invisible ()
```

12