

```

only if update_alpha is TRUE
* priorvals = vector containing values for prior distributions as
follows
*
* time_specific_alpha = integer - logical indicating whether to make
alpha_time-specific or one global alpha.
* update_alpha = integer - logical indicating whether to update alpha
or not.
* update_eta1 = integer - logical indicating whether to update eta1
or set it to zero for all subjects.
* update_phi1 = integer - logical indicating whether to update phi1
or set it to zero.
* sppm = integer - logical indicating whether to use spatial
information or not
*
* SpatialCohesion = integer indication which cohesion to use
* 1 -Auxiliary
* 2- Double dipper
*
* cParms - vector holding values employed in the cohesion
*
* OUTPUT
* Si -
* mu -
* Sig2 -
* eta1
* theta -
* tau2 -
* phi0 -
* phi1
* gamma -
* alpha.out -
*
* Ipm1 -
* Waic -
*****
void drpm_ar1_sppm(int *draws, int *burn, int *thin, int *nsubject,
int *ntime,
double *y, double *s1, double *s2, double *M,
double *alpha, double *modelPriors, double *alphaPriors,
int *time_specific_alpha,
int *update_alpha, int *update_eta1, int *update_phi1,
int *sppm, int *SpatialCohesion, double *cParms, double *m,
int *space_1, int *simpleModel, double *theta_tau2,
int *Si, double *mu, double *sig2, double *eta1, double
*theta, double *tau2,
double *phi0, double *phi1, double *lam2, int *gamma, double
*alpha_out,

```

/\* Code is of the file DepdenetPartitionsAR1LikeAR1Theta\_SPPM2.C  
\* that one since the R function drpm\_fit calls the function which  
\* is defined there:  
initial\_partition <- 0  
C.out <- .C("drpm\_ar1\_sppm", // defined in that file  
as.integer(draws), as.integer(burn), as.integer(thin),  
as.integer(nsubject), as.i)  
\* so it should be the "main" file
\*/

\*\*\*\*\*  
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\*

\* This file contains C code for an MCMC algorithm constructed  
\* to fit a hierarchical model that incorporates the idea of  
\* temporally dependent partitions.
\*

\* I will include model details at a later date
\*

\*\*\*\*\*  
#include "matrix.h"  
#include "Rutil.h"  
#include <R\_ext/Lapack.h>  
#include <R.h>  
#include <Rmath.h>  
#include <stdio.h>  
#include <stdlib.h>  
#include <time.h>  
\*\*\*\*\*  
\* The following are the inputs of the function that are read from R
\*

\* draws = total number of MCMC draws
\* burn = number of MCMC draws discarded as burn-in
\* thin = indicates how much MCMC chain should be thinned
\*

\* nsubject = integer for the number of subjects in data set
\* ntime = integer for the number of time points
\* y = double nsubject x ntime matrix containing response for each
subject at time t
\* s1 = nsubject x 1 vector containing spatial coordinate one
\* s2 = nsubject x 1 vector containing spatial coordinate two
\*

\* M = double indicating value of M associated with cohesion (scale
parameter of DP).
\* alpha = double - prior probability of being pegged, starting value

```

double *fitted, double *llike, double *lpm1, double *waic){

// i - MCMC iterate
// ii - MCMC iterate that is saved
// j - subject iterate
// jj - second subject iterate
// t - time iterate
// k - cluster iterate
// kk - second cluster iterate
// p - prediction iterate

int i, ii, j, jj, t, k, kk;
ii = 0;

int tout = (*draws - *burn)/*thin*/;
Rprintf("nsubject = %d\n", *nsubject);
Rprintf("ntime = %d\n", *ntime);
Rprintf("nout = %d\n", nout);
Rprintf("update_alpha = %d\n", *update_alpha);
Rprintf("update_eta1 = %d\n", *update_eta1);
Rprintf("update_phi1 = %d\n", *update_phi1);

// Memory vectors to hold MCMC iterates for non cluster specific
parameters
// Memory vectors to hold MCMC iterates for cluster specific
parameters
// Memory vectors to hold MCMC iterates for cluster specific
parameters

double phi0_iter = rnorm(0,3);
double phi1_iter = runif(0,1);
double lam2_iter = runif(0, modelPriors[4]*modelPriors[4]);
double *alpha_iter = R_VectorInit(ntime1, *alpha);

// Memory vectors to hold MCMC iterates for cluster specific
parameters
// Memory vectors to hold MCMC iterates for cluster specific
parameters
// Memory vectors to hold MCMC iterates for cluster specific
parameters

double *muh = R_VectorInit(*nsubject)*(ntime1, 0.0);
double *sig2h = R_VectorInit(*nsubject)*(ntime1, 1.0);
if(*simpleModel==1){
    for(t = 0; t < ntime1; t++){
        theta_iter[t] = theta_tau2[0];
        tau2_iter[t] = theta_tau2[1];
    }
}
int nh[*nsubject]*(ntime1);

// Initialize a few parameter vectors
//
// Initialize Si according to covariates
// I am adding one time period here to have
// scratch memory (never filled in) so that
// I can avoid dealing with boundaries in algorithm
for(j = 0; j < *nsubject; j++){
    for(t = 0; t < ntime1; t++) {
        "added time memory"
        Si_iter[j*(ntime1) + t] = 1;
        gamma_iter[j*(ntime1) + t] = 0;
        nh[j*(ntime1) + t] = 0;
        if(t==1) Si_iter[j*ntime1 + t] = 1;
        if(t==*ntime) Si_iter[j*(ntime1) + t] = 0;
    }
}
// Initial enumeration of number of subjects per cluster;
for(j = 0; j < *nsubject; j++){
    for(t = 0; t < *ntime; t++){
        nh[(Si_iter[j*(ntime1)+t]-1)*(ntime1) + t] = nh[(Si_iter[j*(ntime1) + t] + 1];
    }
}

int ntime1 = *ntime + 1;
// I am adding one more year as an empty vector
// so that the C program does not crash.
int gamma_iter[*nsubject]*(ntime1);
int Si_iter[*nsubject)*(ntime1);
int nclus_iter[ntime1];
double *eta1_iter = R_VectorInit(*nsubject, runif(0,1));
double *theta_iter = R_VectorInit(ntime1, rnorm(0,3));
double *tau2_iter = R_VectorInit(ntime1, runif(0,
modelPriors[3]*modelPriors[3]));

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        double *ph = R_VectorInit(*nsubject, 0.0);
        double *phtmp = R_VectorInit(*nsubject, 0.0);
        double *probh = R_VectorInit(*nsubject, 0.0);
        double *lgweight = R_VectorInit(*nsubject, 0.0);
        double *s1o = R_Vector(*nsubject);
        double *s2o = R_Vector(*nsubject);
        double *s1n = R_Vector(*nsubject);
        double *s2n = R_Vector(*nsubject);
        for(j=0; j<(*nsubject); j++){
            comp1[j] = 0; comptp1[j] = 0, comp2t[j]=0;
        }
    } // stuff I need to update eta1
    double e1o, e1n, logito, logitn, one_phiisq;
    // stuff I need to update muh and sig2h
    double mstar, s2star, sumy, sume2;
    double nsig, osig, llo, lln, llr;
    double *mu_tmp = R_VectorInit(*nsubject, 0.0);
    double *sig2_tmp = R_VectorInit(*nsubject, 1.0);
    // stuff that I need for theta and lam2
    double summu, nt, ot, lam2tmp, phi1sq, sumt, op1, np1, ol, nl;
    // double ssq;
    // stuff that I need to update alpha
    int sumg;
    double astar, bstar, alpha_tmp;
    // Stuff to compute lpml, likelihood, and WAIC
    int like0, nout_0=0;
    double lpml_iter, elppdWAIC;
    double *CPO = R_VectorInit((*nsubject)*(ntime1), 0.0);
    double *like_iter = R_VectorInit((*nsubject)*(ntime1), 0.0);
    double *fitted_iter = R_VectorInit((*nsubject)*(ntime1), 0.0);
    double *mnlk = R_VectorInit((*nsubject)*(ntime1), 0.0);
    double *mllike = R_VectorInit((*nsubject)*(ntime1), 0.0);
    // stuff to predict
    // int gpred[*nsubject], nh_pred[*nsubject];
    // =====
    // Prior parameter values
    // =====
    // prior values for sig2
    double Asig=modelPriors[2];
    double Atau=modelPriors[3];
    double Alam=modelPriors[4];
}

// Initialize the number of clusters
for(t = 0; t < *ntime; t++){
    nclus_iter[t] = 0;
    for(j = 0; j < *nsubject; j++){
        if(nh[j*(ntime1) + t] > 0) nclus_iter[t] = nclus_iter[t] + 1;
    }
    nclus_iter[*ntime] = 0;
}

// scratch vectors of memory needed to update parameters
int nh_redtmp[*nsubject], nh_tmpr[*nsubject];
int nh_redtmp_no_zero[*nsubject],
nh_tmpr_no_zero[*nsubject], nh_red_no_zero[*nsubject];
int nh_red1[*nsubject];
// int nclus_red1;
int nh_redtmp_1[*nsubject], nh_tmpr_1[*nsubject];
int nh_redtmp_no_zero_1[*nsubject],
nh_red_no_zero_1[*nsubject], nh_tmpr_no_zero_1[*nsubject];
nh_red_no_zero_1[*nsubject], nh_tmpr_no_zero_1[*nsubject];
nh_red_no_zero[*nsubject], nh_tmpr_no_zero[*nsubject];
double *s1_red = R_VectorInit(*nsubject, 0.0);
double *s2_red = R_VectorInit(*nsubject, 0.0);
for(j=0; j<*nsubject; j++){
    nh_tmpr[j] = 0; nh_red[j] = 0; nh_redtmp[j] = 0;
    nh_redtmp_no_zero[j] = 0; nh_tmpr_no_zero[j] = 0;
    nh_red_no_zero[j] = 0; nh_tmpr_no_zero_1[j] = 0;
}
nh_tmpr_1[j] = 0; nh_red1[j] = 0; nh_redtmp_1[j] = 0;
nh_redtmp_no_zero_1[j] = 0; nh_tmpr_no_zero_1[j] = 0;
nh_tmpr_no_zero_1[j] = 0;

// stuff that I need to update Si (the partition);
int comp1t[*nsubject], comptm1[*nsubject], comp2t[*nsubject], comptp1[*nsubject];
int rho_tmpr[*nsubject], Si_tmpr[*nsubject], Si_tmpr2[*nsubject];
int Si_red[*nsubject], Si_red_1[*nsubject];
int oldLab[*nsubject], reorder[*nsubject];
int iaux, Rindx1, Rindx2, n_tmpr, nclus_tmpr, rho_comp, indx;
double auxm, auxs, mudraw, sigdraw, maxph, denph, cprobh, uu, lco,
lcn, lcn_1, lpp;
}

```



```

for(jj = 0; jj < n_red_1; jj++){
    nh_red[jj] = 0; nh_red_1[jj] = 0;
}

nclus_red = 0;
for(jj = 0; jj < n_red; jj++){
    nh_red[Si_red[jj]-1] = nh_red[Si_red[jj]-1] + 1;
    nh_red_1[Si_red_1[jj]-1] = nh_red_1[Si_red_1[jj]-1] + 1;
    if(Si_red[jj] > nclus_red) nclus_red = Si_red[jj];
}
}

nh_red_1[Si_red_1[n_red]-1] = nh_red_1[Si_red_1[n_red]-1] +
1; // this may need to be updated depending on if the value of
gamma changes
/nclus_red_1 = nclus_red;
/if(Si_red_1[n_red] > nclus_red) nclus_red_1 =
Si_red_1[n_red];
1Co=0.0, 1Cn=0.0;
for(k = 0; k < nclus_red; k++){
if(*sPPM==1){
// Note that if space is only included for first time
point
// then it does not inform gamma.
if((*space_1==1 & t == 0) | (*space_1==0)){
indx = 0;
for(jj = 0; jj < n_red; jj++){
if(Si_red[jj] == k+1){
s1o[indx] = s1_red[jj];
s2o[indx] = s2_red[jj];
}
s1n[indx] = s1_red[jj];
s2n[indx] = s2_red[jj];
indx = indx+1;
}
s1n[indx] = s1[j];
s2n[indx] = s2[j];
}
1Co = Cohesion3_4(s1o, s2o, mu0, k0, v0, L0,
1Cn = Cohesion3_4(s1n, s2n, mu0, k0, v0, L0,
nh_red[k], *SpatialCohesion, 1);
1Cn = Cohesion3_4(s1o, s2o, mu0, k0, v0, L0,
nh_red[k+1], *SpatialCohesion, 1);
}
}

lgweight[k] = Log(nh_red[k]) + 1Cn - 1Co;
Si_red_1[Rindx1] = k+1;
rho_comp = compatibility(Si_red_1, comptm1, Rindx1+1);
}
}

if((*space_1==1 & t == 0) | (*space_1==0)) {
    if(*sPPM==1){
        if((*space_1==1 & t == 0) | (*space_1==0)) {
            s1o[0] = s1[j];
            s2o[0] = s2[j];
            1Cn_1 = Cohesion3_4(s1o, s2o, mu0, k0, v0, L0,
1,*SpatialCohesion, 1);
        }
        lgweight[nclus_red] = Log(Map) + 1Cn_1;
        Si_red_1[Rindx1] = nclus_red+1;
        rho_comp = compatibility(Si_red_1, comptm1, Rindx1+1);
    }
    // Note that if space is only included for first time
    // then it does not inform gamma.
    if((*space_1==1 & t == 0) | (*space_1==0)){
        indx = 0;
        for(jj = 0; jj < nclus_red; jj++){
            if(Si_red[jj] == k+1){
                s1o[indx] = s1_red[jj];
                s2o[indx] = s2_red[jj];
            }
            s1n[indx] = s1_red[jj];
            s2n[indx] = s2_red[jj];
            indx = indx+1;
        }
        s1n[indx] = s1[j];
        s2n[indx] = s2[j];
    }
    1Co = Cohesion3_4(s1o, s2o, mu0, k0, v0, L0,
    1Cn = Cohesion3_4(s1n, s2n, mu0, k0, v0, L0,
    nh_red[k+1], *SpatialCohesion, 1);
}
}

probh[1] = alpha_iter[t]/(alpha_iter[t] + (1-
alpha_iter[t])*lgweight[cit1-1]);
// If gamma is 1 at current MCMC iterate, then there are no
// concerns about partitions being incompatible as gamma
}

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gt = rbinom(1,probh[1]);
gamma_iter[j*(ntime1) + t] = gt;
}

be taken when
// However, if gamma's current value is 0, then care must
partitions may
// trying to change from gamma=0 to gamma=1 as the
// no longer be compatible
if(gamma_iter[j*(ntime1) + t] == 0){

// To determine compatibility, I need to make sure that
// comparison of the reduced partitions is being made with
// correct cluster labeling. I try to do this by
identifying
// the sets of units and sequentially assigning "cluster
labels"
// starting with set that contains the first unit. I
wonder if
// there is a way to do this in C with out using loops?
who
// can I ask about this?
// Get rho_t | gamma_t = 1 and rho_{t-1} | gamma_t = 1
// when gamma_{it} = 1;
Rindx1 = 0;
for(jj = 0; jj < *nsubject; jj++){
if(gamma_iter[jj*ntime1 + (t)] == 1){
comptm1[Rindx1] = Si_iter[jj*ntime1 + (t-1)];
comp1[Rindx1] = Si_iter[jj*ntime1 + (t)];
Rindx1 = Rindx1 + 1;
}
}

// I need to include this because determine what
happens when
// gamma goes from 0 to 1;
if(jj == j){
comptm1[Rindx1] = Si_iter[jj*ntime1 + (t-1)];
comp1[Rindx1] = Si_iter[jj*ntime1 + (t)];
Rindx1 = Rindx1 + 1;
}

rho_comp = compatibility(comptm1, comp1, Rindx1);
// If rho_comp = 0 then not compatible and probability of
// pegging subject needs to be set to 0;
if(rho_comp==0){
probh[1] = 0;
}

// Observation belongs to a non-singleton :::
nh[(Si_iter[j*(ntime1) + t]-1)*(ntime1 + t) > 1] =

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nh[(Si_iter[j*(ntime1) + t]-1)*(ntime1) + t] - 1;
}

}else{
    // Observation is a member of a singleton cluster ...
    iaux = Si_iter[j*(ntime1) + t];
    if(iaux < nclus_iter[t]){
        // Need to relabel clusters. I will do this by swapping
        // cluster labels
        // Si_iter[j] and nclus_iter along with cluster
        // specific parameters;
        j's label
        for(jj = 0; jj < *nsubject; jj++){
            if(Si_iter[jj*(ntime1) + t] == nclus_iter[t]){
                Si_iter[jj*(ntime1) + t] = iaux;
            }
        }
        Si_iter[j*(ntime1) + t] = nclus_iter[t];
        // The following steps swaps order of cluster specific
        // parameters
        // so that the newly labeled subjects from previous
        // step retain
        // their correct cluster specific parameters
        auxs = sig2h[(iaux-1)*ntime1 + t];
        sig2h[(iaux-1)*ntime1 + t] = sig2h[(nclus_iter[t]-1)*
        (ntime1)+t];
        sig2h[(nclus_iter[t]-1)*(ntime1)+t] = auxs;
        auxm = muh[(iaux-1)*ntime1 + t];
        muh[(iaux-1)*ntime1 + t] = muh[(nclus_iter[t]-1)*
        (ntime1)+t];
        muh[(nclus_iter[t]-1)*(ntime1)+t] = auxm;
        // the number of members in cluster is also swapped
        // with the last
        nh[(iaux-1)*(ntime1)+t] = nh[(nclus_iter[t]-1)*
        (ntime1)+t];
        nh[(nclus_iter[t]-1)*(ntime1)+t] = 1;
    }
    // Now remove the ith obs and last cluster;
    nh[(nclus_iter[t]-1)*(ntime1)+t] = nh[(nclus_iter[t]-1)*
    (ntime1)+t] - 1;
    nclus_iter[t] = nclus_iter[t] - 1;

    }for(jj = 0; jj < *nsubject; jj++){
        rho_tmp[jj] = Si_iter[jj*(ntime1) + t];
    }
    for(k = 0; k < nclus_iter[t]; k++){
        rho_tmp[j] = k+1;
    }
    // First need to check compatibility
    Rindx2=0;
    for(jj = 0; jj < *nsubject; jj++){
        if(gamma_iter[jj*ntime1 + (t+1)] == 1){
            comp2t[Rindx2] = rho_tmp[jj];
            comp2t[Rindx2] = Si_iter[jj*ntime1 + (t+1)];
            Rindx2 = Rindx2 + 1;
        }
    }
    // check for compatibility
    rho_comp = compatibility(comp2t, comptp1, Rindx2);
    if(rho_comp != 1){
        ph[k] = Log(0); // Not compatible
    }else{
        // Need to compute Pr(rhot), Pr(rhot,R), Pr(rhot+1),
        Pr(rhot+1,R)
        for(jj = 0; jj < *nsubject; jj++){
            nh_tmp[jj] = 0;
        }
        n_tmp = 0;
        for(jj = 0; jj < *nsubject; jj++){
            nh_tmp[rho_tmp[jj]-1] = nh_tmp[rho_tmp[jj]-1]+1;
            n_tmp=n_tmp+1;
        }
        nclus_tmp=0;
        for(jj = 0; jj < *nsubject; jj++){
            if(nh_tmp[jj] > 0) nclus_tmp = nclus_tmp + 1;
        }
    }
}

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    }
    ph[k] = dnorm(y[j*(ntime) + t],
    muh[k*(ntime1) + t] +
    eta1_iter[j]*y[j*(ntime) + t]*  

    sqrt(sig2h[k*(ntime1) + t]*  

    (1-eta1_iter[j]*eta1_iter[j])), 1)+
    1pp;
}

// use this to test if MCMC draws from prior are
// Beginning of spatial part
if(*SPPM==1){
    if((*space_1==1 & t == 0) | (*space_1==0)){
        indx = 0;
        for(jj = 0; jj < *nsubject; jj++){
            if(rho_tmp[jj] == kk+1){
                s1n[indx] = s1[jjj];
                s2n[indx] = s2[jjj];
                indx = indx+1;
            }
        }
    }
}

// End of spatial part
nh_tmp[kk] + 1cn;
1pp = 1pp + nh_tmp[kk]*log(Mdp) + 1gamma((double)
nh_tmp[kk] + 1cn;
1pp = 1pp + (Log(Mdp) + Log((double) nh_tmp[kk]) +
1cn);

if(t==0){
    muh[k*(ntime1) + t],
    sqrt(sig2h[k*(ntime1) + t]), 1));
ph[k] = dnorm(y[j*(ntime) + t],
    muh[k*(ntime1) + t],
    sqrt(sig2h[k*(ntime1) + t]), 1) +
    sqrt(sig2h[j*(ntime1) + t]), 1);
}

if(t > 0){
    muh[k*(ntime1) + t] +
    eta1_iter[j]*y[j*(ntime) + t-1],
    sqrt(sig2h[k*(ntime1) + t]*  

    (1-eta1_iter[j]*eta1_iter[j])), 1)+
    1pp;
}

// use this to test if E.U. gets allocated to a new cluster
// Need to check compatibility first
rho_tmp[j] = nclus_iter[t]+1;

// First need to check compatibility
Rindx1 = 0, Rindx2=0;
for(jj = 0; jj < *nsubject; jj++){
    if(gamma_iter[jj*ntime1 + (t+1)] == 1){
        comp2[Rindx2] = rho_tmp[jj];
        comp1[Rindx2] = si_iter[jj*ntime1 + (t+1)];
        Rindx2 = Rindx2 + 1;
    }
}
// check for compatibility
rho_comp = compatibility(comp2t, comptp1, Rindx2);
if(rho_comp != 1){
    ph[nclus_iter[t]] = Log(0); // going to own cluster is
not compatible;
} else {
    mudraw = rnorm(theta_iter[t], sqrt(tau2_iter[t]));
    sigdraw = runif(0, Asig);
    for(jj = 0; jj < *nsubject; jj++){
        nh_tmp[jj] = 0;
    }
    nh_tmp = 0;
    for(jj = 0; jj < *nsubject; jj++){
        nh_tmp[rho_tmp[jj]-1] = nh_tmp[rho_tmp[jj]-1]+1;
    }
    n_tmp=n_tmp+1;
}
}

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nh[(Si_iter[j*(ntime1) + t]-1)*(ntime1)+t] = 1;
rho_tmp[j] = nclus_iter[t];
muh[(Si_iter[j*(ntime1) + t]-1)*(ntime1) + t] = mudraw;
sig2h[(Si_iter[j*(ntime1) + t]-1)*(ntime1) + t] =
sigdraw*sigdraw;
if(*simpleModel==1) sig2h[(Si_iter[j*(ntime1) +
t]-1)*(ntime1) + t] = 1.0;
}
for(jj = 0; jj < *nsubject; jj++){
Si_tmp[jj] = Si_iter[jj*(ntime1) + t];
Si_tmp2[jj] = 0;
reorder[jj] = 0;
}
// I believe that I have to make sure that groups are order
so that // EU one is always in the group one, and then the smallest
index not // with group 1 anchors group 2 etc.

relabel(Si_tmp, *nsubject, Si_tmp2, reorder, oldLab);

if(jj=0; jj<*nsubject; jj++){
Si_iter[jj*(ntime1) + t] = Si_tmp2[jj];
}
for(k = 0; k < nclus_iter[t]; k++){
muh[k*(ntime1)+t] = mu_tmp[(oldLab[k]-1)];
sig2_tmp[k] = sig2h[k*(ntime1)+t];
}
for(k = 0; k < nclus_iter[t]; k++){
nh[k*(ntime1)+t] = reorder[k];
muh[k*(ntime1)+t] = mu_tmp[(oldLab[k]-1)];
sig2h[k*(ntime1)+t] = sig2_tmp[(oldLab[k]-1)];
}

// I believe that I have to make sure that groups are order so
that // EU one is always in the group one, and then the smallest
index not // with group 1 anchors group 2 etc.

relabel(Si_tmp, *nsubject, Si_tmp2, reorder, oldLab);

if(t==0){
sumy = 0.0;
for(j = 0; j < *nsubject; j++){
if(Si_iter[j*(ntime1) + t] == k+1){
sumy = sumy + y[j*(ntime1)+t];
}
}
s2star = 1/((double) nh[k*(ntime1)+t]/sig2h[k*(ntime1) + t]
+ 1/tau2_iter[t]);
mstar = s2star*( 1/sig2h[k*(ntime1) + t]*sumy +
(1/tau2_iter[t])*theta_iter[t]);
}

if(t > 0){
sumy = 0.0;
sume2 = 0.0;
for(j = 0; j < *nsubject; j++){
if(Si_iter[j*(ntime1) + t] == k+1){
sume2 = sume2 + 1.0/(1-eta1_iter[j]*eta1_iter[j]);
sumy = sumy + (y[j*(ntime1)+t] - eta1_iter[j]*y[j*
(*ntime1)+t-1])/(
(1-eta1_iter[j])*eta1_iter[j]);
}
}
s2star = 1/(
(1.0/sig2h[k*(ntime1) + t])*sume2 +
)
}
}

```

```

1/tau2_iter[t];
mstar = s2star*( (1.0/sig2h[k*(ntime1) + t])*sumy +
(1./tau2_iter[t])*theta_iter[t];
}

muh[k*(ntime1) + t] = rnorm(mstar, sqrt(s2star));
// muh[k] = 0.0;
///////////////////////////////
// update sig2h
// ///////////////////////////////////////////////////
osig = sqrt(sig2h[k*(ntime1) + t]);
nsig = rnorm(osig,csigSIG);
if(nsig > 0.0 & nsig < Asig){

1ln = 0.0;
1lo = 0.0;
if(t == 0){
for(j = 0; j < *nsubject; j++){
if(Si_iter[j*(ntime1) + t] == k+1){
1lo = 1lo + dnorm(y[j*(ntime1)+t], muh[k*(ntime1) +
t], osig,1);
1ln = 1ln + dnorm(y[j*(ntime1)+t], muh[k*(ntime1) +
t], nsig,1);
}
}
if(t > 0){
for(j = 0; j < *nsubject; j++){
if(Si_iter[j*(ntime1) + t] == k+1){
1lo = 1lo + dnorm(y[j*(ntime1)+t], muh[k*(ntime1) +
t] +
eta1_iter[j]*eta1_iter[j],1);
1ln = 1ln + dnorm(y[j*(ntime1)+t], muh[k*(ntime1) +
t] +
eta1_iter[j]*y[j]*(ntime1) + t-1],
osig*sqrt(1-
eta1_iter[j]*eta1_iter[j],1));
}
}
eta1_iter[j]*eta1_iter[j],1);
1ln = 1ln + dnorm(y[j*(ntime1)+t], muh[k*(ntime1) +
t] +
eta1_iter[j]*y[j]*(ntime1) + t-1],
osig*sqrt(1-
eta1_iter[j]*eta1_iter[j],1));
}
}
else if(t==(*ntime-1)){
s2star = 1.0/((double) nclus_iter[t]/tau2_iter[t] +
1.0/1am2_iter + phi1sq/1am2tmp);
mstar = s2star*((1.0/tau2_iter[t])*summu +
(1.0/1am2_iter)*phi0_iter +
(1.0/1am2tmp)*phi1_iter*(theta_iter[t+1]-
phi0_iter*(1-phi1_iter)));
}
else if(t==0){
s2star = 1.0/((double) nclus_iter[t]/tau2_iter[t] +
1.0/1am2tmp);
mstar = s2star*((1.0/tau2_iter[t])*summu +
(1.0/1am2tmp)*(phi0_iter*(1-phi1_iter) +
phi1_iter*theta_iter[t-1]));
}
else {
s2star = 1.0/((double) nclus_iter[t]/tau2_iter[t] +
(1.0 +
phi1sq)/1am2tmp);
mstar = s2star*((1.0/tau2_iter[t])*summu +
(1.0/1am2tmp)*(phi0_iter*(1-phi1_iter) +
phi1_iter*theta_iter[t-1]));
}
}
}

```

```

theta_iter[t+1] +          // update eta1 (temporal correlation parameter at likelihood)
(1.0/lam2tmp)*(phi1_iter*(theta_iter[t-1] +
phi0_iter*(1.0 - phi1_iter)));
}

theta_iter[t] = rnorm(mstar, sqrt(s2star));
if(*simpleModel==1) theta_iter[t] = 0.0;
// update tau2 (variance of m)
// update tau2 (variance of m)
// update tau2 (variance of m)

ot = sqrt(tau2_iter[t]);
nt = rnorm(ot,csigTAU);
if(nt > 0){
lln = 0.0;
for(k = 0; k < nclus_iter[t]; k++){
  ll0 = lln + dnorm(muh[k*(ntime1 + t], theta_iter[t],
  ot,1);
  lln = lln + dnorm(muh[k*(ntime1 + t], theta_iter[t],
  nt,1);
}
llr = lln + dunif(nt, 0.0, Atau, 1);
lln = lln + dunif(nt, 0.0, Atau, 1);
uu = runif(0,1);

if(Log(uu) < llr){
  tau2_iter[t] = nt*nt;
  tau2_iter[t] = 5*5;
}

if(*simpleModel==1) tau2_iter[t] = theta_tau[1];
// if(llr > Log(uu)) eta1_iter[j] = e1n;
}

if(*update_eta1==1{
  for(j = 0; j < *nsubject; j++){
    e1o = eta1_iter[j];
    e1n = rnorm(e1o, csigETA1);
    if(e1n < 1 & e1n > -1){

      lln=0.0;
      for(t=1; t<ntime; t++){
        ll0 = lln + dnorm(y[j*(ntime)+t],
        muh[(Si_iter[j*(ntime1 + t]-1)*(ntime1 + t]
        + e1o*y[j*(ntime)+t-1],
        sqrt(sig2h[(Si_iter[j*(ntime1 +
        t]-1)*(ntime1 + t]*e1o)*e1o)), 1];
        lln = lln + dnorm(y[j*(ntime)+t],
        muh[(Si_iter[j*(ntime1 + t]-1)*(ntime1 + t]
        + e1r*y[j*(ntime)+t-1],
        sqrt(sig2h[(Si_iter[j*(ntime1 +
        t]-1)*(ntime1 + t)*e1n)), 1];

      }
      logito = Log(0.5*(e1o + 1)) - Log(1 - 0.5*(e1o+1));
      logitn = Log(0.5*(e1n + 1)) - Log(1 - 0.5*(e1n+1));
    }
    llr = lln - ll0;
    uu = runif(0,1);
    if(llr > Log(uu)) eta1_iter[j] = e1n;
  }
}

if(*update_eta1==1{
  for(j = 0; j < *nsubject; j++){
    e1o = lln - ll0;
    llr = runif(0,1);
    if(llr > Log(uu)) eta1_iter[j] = e1n;
  }
}

```



```

llr = lln - ll0;
if(llr > Log(runif(0,1))) phi1_iter = np1;
}

/*
// Update lambda with a MH step
phi1sq = phi1_iter*phi1_iter;
ol = sqrt(lam2_iter);
nl = rnorm(ol, csigLAM);
ln = 0.0;
ll0 = 0.0;
for(t=1; t<*ntime; t++){
  lln = ll0 + dnorm(theta_iter[t],
    phi0_iter*(1-phi1_iter) +
    phi1_iter*theta_iter[t-1], ol*sqrt(1-phi1sq),1);
  ln = lln + dnorm(theta_iter[t],
    phi0_iter*(1-phi1_iter) +
    phi1_iter*theta_iter[t-1], nl*sqrt(1-phi1sq),1);
  ll0 = lln + dnorm(theta_iter[0], phi0_iter, ol, 1) + dunif(ol,
    0.0, Alam, 1);
  ln = lln + dnorm(theta_iter[0], phi0_iter, nl, 1) + dunif(nl,
    0.0, Alam, 1);
  llr = lln - ll0;
  uu = runif(0,1);
  if(Log(uu) < llr){
    lam2_iter = nl*nl;
  }
}
phi1sq = phi1_iter*phi1_iter;
ssq = 0.0;
for(t=1; t<*ntime; t++){
  ssq = ssq + (theta_iter[t] - (phi0_iter*(1-phi1_iter) +
    phi1_iter*theta_iter[t-1])* (theta_iter[t] - (phi0_iter*(1-phi1_iter) +
    phi1_iter*theta_iter[t-1])));
  ssq = 1.0/(1.0 - phi1sq)*ssq + (theta_iter[0]-phi0_iter)*
    (theta_iter[0]-phi0_iter);
  astar = 0.5*(*ntime) + 1;
  lam2_iter = 1.0/rgamma(astar, 1/bstar);
}

/*
// predict partition for new time period
*/
/*
/* predict partition for new time period
*/
/*
for(p = 0; p < *npred; p++){
  for(j=0; j<*nsubject; j++){
    nh_pred[j] = 0;
    predSi_iter[j*(*npred) + p] = 0;
  }
  if(*update_alpha == 0){
    n_red = 0;
    for(j=0;j<*nsubject;j++){
      gpred[j] = rbinom(1,*alpha);
      if(gpred[j] == 1){
        nh_pred[Si_iter[j*(ntime1)+(*ntime)-1] - 1] =
          nh_pred[Si_iter[j*(ntime1)+(*ntime)-1] - 1] + 1;
        n_red = n_red + 1;
        predSi_iter[j*(*npred) + p] = Si_iter[j*(ntime1) +
          (*ntime)-1];
      }
    }
    if(*update_alpha == 1){
      if(*time_specific_alpha == 1){
        n_red = 0;
        for(j=0;j<*nsubject;j++){
          gpred[j] = rbinom(1,alpha_iter[1]);
          if(gpred[j] == 1){
            nh_pred[Si_iter[j*(ntime1)+(*ntime)-1] - 1] =
              nh_pred[Si_iter[j*(ntime1)+(*ntime)-1] - 1] + 1;
            n_red = n_red + 1;
            predSi_iter[j*(*npred) + p] = Si_iter[j*(ntime1) +
              (*ntime)-1];
          }
        }
      }
    }
  }
}

```

```

        }
    }

    }else {
        /*
        // evaluating likelihood that will be used to calculate LPML and
        WAIC?
        // (see page 81 Christensen Hansen and Johnson)
        //
        ///////////////////////////////////////////////////////////////////
        */

        remove_zero(nh_pred, *nsubject, nh_tmp_no_zero);
        nclus_tmp = 0;
        for(j=0; j<*nsubject;j++){
            if(nh_tmp_no_zero[j] > 0){
                nclus_tmp = nclus_tmp + 1;
            }else{
                break;
            }
        }
        if(gpred[j] == 0){
            for(k = 0; k < nclus_tmp; k++){
                probh[k] = nh_pred[k]/(n_red + Mdp);
            }
            probh[nclus_tmp] = Mdp/(n_red + Mdp);
            uu = runif(0.0,1.0);
            cprobh= 0.0;;
            for(k = 0; k < nclus_tmp+1; k++){
                cprobh = cprobh + probh[k];
            }
            if (uu < cprobh){
                iaux = k+1;
                break;
            }
        }
        if(iaux < nclus_tmp){
            predSi_iter[j*(npred) + p] = iaux;
            nh_pred[iaux-1] = nh_pred[iaux-1] + 1;
        }else{
            nclus_tmp = nclus_tmp + 1;
            predSi_iter[j*(npred) + p] = nclus_tmp;
            nh_pred[(predSi_iter[j*(npred) + p]-1)*(*npred)+p] = 1;
        }
        n_red = n_red + 1;
    }
}

if(likeθ==1) nout_θ = nout_θ + 1;
}

```

```

if(likeθ==0){
    for(j = 0; j < *nsubject; j++){
        for(t = 0; t < *ntime; t++){
            CPO[j*(*ntime)+t] = CPO[j*(*ntime)+t] +
                (1/exp(like_iter[j*(*ntime)+t]));
    }
}

///////////////////////////////
// Save MCMC iterates
//



if((i > (*burn-1) & ((i+1) % *thin == 0)){
    for(t = 0; t < *ntime; t++){
        alpha_out[ii>(*ntime) + t] = alpha_iter[t];
        theta[ii>(*ntime) + t] = theta_iter[t];
        tau2[ii>(*ntime) + t] = tau2_iter[t];
        for(j = 0; j < *nsubject; j++){
            sig2[(ii>(*nsubject) + j)*(*ntime) + t] = sig2h[(Si_iter[j]
                (*ntime1) + t]-1)*(ntime1) + t];
            mu[(ii>(*nsubject) + j)*(*ntime) + t] = muh[(Si_iter[j]
                (*ntime1) + t]-1)*(ntime1) + t];
            Si[(ii)(*nsubject) + j)*(*ntime) + t] = si_iter[j*ntime1 +
                t];
            gamma[(ii>(*nsubject) + j)*(*ntime) + t] =
                gamma_iter[j*ntime1 + t];
            like[(ii)(*nsubject) + j)*(*ntime) + t] = like_iter[j*
                (*ntime)+t];
            fitted[(ii)(*nsubject) + j)*(*ntime) + t] = fitted_iter[j*
                (*ntime)+t];
        }
    }
}

eta1[ii>(*nsubject) + j] = eta1_iter[j];
}
phi1[ii] = phi1_iter;
phi0[ii] = phi0_iter;
lam2[ii] = lam2_iter;
ii = ii+1;
}
*/
}

```