

```

/*
 * 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 only if
 * update_alpha is TRUE
 * priorsvals = 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
 *
 * cParams - vector holding values employed in the cohesion
 *
 * OUTPUT
 * Si -
 * mu -
 * sig2 -
 * eta1 -
 * theta -
 * tau2 -
 * waic -
 * lpm1 -
 * phi0 -
 * phi1 -
 * gamma -
 * alpha.out -
 * like -
 * lpm1 -
 * 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 *charms, double *mh,
 *                 int *space_1, int *simpleModel, double *theta_tau2,
 *                 int *Si, double *mu, double *sig2, double *eta1, double *theta,
 *                 double *phi0, double *phi1, double *lam2, int *gamma, double *alpha_out,
 *                 double *fitted, double *like, 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", tout);
 * Rprintf("update_alpha = %d\n", *update_alpha);
 * Rprintf("update_eta1 = %d\n", *update_eta1);
 * Rprintf("update_phi1 = %d\n", *update_phi1);
 */


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// Memory vectors to hold MCMC iterates for non cluster specific parameters
// =====
// This variable is used to create a "buffer" zone of memory so that updating
// things on time boundary do not need special attention in the algorithm since
// I have to look at time period before and after when updating partitions
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 *theta1_iter = R_VectorInit(ntime1, rnorm(0,3));
double *tau2_iter = R_VectorInit(ntime1, runif(0, modelPriors[3]));
phi0_iter = rnorm(0,3);
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
// =====
// simpleModel1==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++){
        Si_iter[j*(ntime1) + t] = 1;
        gamma_iter[j*(ntime1) + t] = 0;
        nh[j*(ntime1) + t] = 0;
        if(t==0) Si_iter[j*ntime1 + t] = 1;
        if(t==*ntime) Si_iter[j*(ntime1) + t] = 0;
    }
}
// Note I am not initializing the "added time
memory"
1pp;

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// 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)*
(ntime1)+t] + 1;
    }
}
// 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
// =====
// stuff needed to update gamma vectors
int nclus_red=0, nh_red[*nsubject], nh_tmp[*nsubject];
int nh_redtmp[*nsubject], nh_tmp[*nsubject];
int nh_redtmp_no_zero[*nsubject],
nh_tmp_no_zero[*nsubject],nh_red_no_zero[*nsubject];
int nh_red_1[*nsubject],nh_red_1[*nsubject];
// int nclus_red_1;
int nh_redtmp_1[*nsubject], nh_tmp_1[*nsubject];
int nh_redtmp_no_zero_1[*nsubject],
nh_red_no_zero_1[*nsubject],nh_tmp_no_zero_1[*nsubject];
double *s1_red = R_VectorInit(*nsubject, 0.0);
double *s2_red = R_VectorInit(*nsubject, 0.0);
for(j=0; j<*nsubject; j++){
    nh_tmp[j] = 0; nh_red[j] = 0; nh_redtmp[j] = 0;
    nh_redtmp_no_zero[j] = 0; nh_tmp_no_zero[j] = 0;
    nh_red_no_zero[j] = 0; nh_red_no_zero_1[j] = 0;
}
nh_tmp_1[j] = 0; nh_red_1[j] = 0; nh_redtmp_1[j] = 0;
nh_redtmp_no_zero_1[j] = 0; nh_tmp_no_zero_1[j] = 0;
nh_red_no_zero_1[j] = 0; nh_red_no_zero_1[j] = 0;
}
// stuff that I need to update Si (the partition);
int comp1t[*nsubject],comptm1[*nsubject],comp2t[*nsubject];
int rho_tmpr[*nsubject], Si_tmpr[*nsubject], Si_tmpt2[*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 auxin, mudraw, sigdraw, maxph, denph, cprobh, uu, lco, lcn, lcn_1,
lpp;
double *ph = R_VectorInit(*nsubject, 0.0);
double *phtmp = R_VectorInit(*nsubject, 0.0);
double *probh = R_VectorInit(*nsubject, 0.0);
double *lweight = R_VectorInit(*nsubject, 0.0);
double *s10 = R_Vector(*nsubject);
double *s20 = R_Vector(*nsubject);

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double *s1n = R_Vector(*nsubject);
double *s2n = R_Vector(*nsubject);
for(j=0; j<(*nsubject); j++){
    compt1[j] = 0; compt1[j] = 0, compt2[j]=0, comptp1[j]=0;
}
// stuff I need to update eta1
double e10, e1n, logito, login, one_physq;
// stuff I need to update muh and sig2h
double mstan, s2star, sumy, sume2;
double nsig, osig, llo, lln, lrr;
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;
// double ssq;

// stuff that I need to update alpha
int sumg;
double astar, bstar, alpha_tmp;
// Stuff to compute lpm1, likelihood, and WAIC
int like0, nout_0=0;
double lpm1_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);
double *mnllike = R_VectorInit(*nsubject)*(ntime1), 0.0);
double *mmlike = R_VectorInit(*nsubject)*(ntime1), 0.0);
// stuff to predict
// int gpred[*nsubject], nh_pred[*nsubject];
// Prior values for sig2
double Asig_modelPriors[2];
double Atau=modelPriors[3];
double Alam=modelPriors[4];
// priors for phi0
double m0 = modelPriors[0], s20 = modelPriors[1];
// priors for alpha
double a = alphaPriors[0], b = alphaPriors[1];
// priors for eta1
double b_eta1 = modelPriors[5];
// DP weight parameter
double Map = *M;
Rprintf("Prior values: Asig = %.2f, Atau = %.2f, Alam = %.2f, \n m0 = %.2f, s20
= %.2f\n", Asig, Atau, Alam, m0, s20);
// Cohesion auxiliary model parameters for Cohesions 3 and 4
double k0=cParams[1], v0=cParams[2];
double *mu0 = R_VectorInit(2,cParams[0]);
double *L0 = R_VectorInit(2*2,0.0);
L0[0] = cParams[3]; L0[3] = cParams[3];
Rindx1 = Rindx1 + 1;
}

if(*sPPM==1){
    RprintVecAsMat("mu0", mu0, 1, 2);
    Rprintf("k0 = %f\n", k0);
    Rprintf("v0 = %f\n", v0);
    RprintVecAsMat("L0", L0, 2, 2);
}
// M-H step tuning parameter
double csigTG=mh[0], csigTAU=mh[1], csigLAM=mh[2], csigETA1=mh[3],
csigPHI1=mh[4];
GetRNGstate();
// start of the mcmc algorithm;
// begin by updating gamma (pegged) parameters
for(i = 0; i < *draws; i++){
    if((i+1) % 10000 == 0){
        time_t now;
        time(&now);
        Rprintf("mcmc iter = %d =====\n", i+1);
        Rprintf("%s",ctime(&now));
    }
    // Start updating gamma and partition for each time period
    for(t = 0; t < *ntime; t++){
        // at time period one, all gammas are zero (none are `pegged`)
        if(t == 0){
            gamma_iter[j*(ntime1) + t] = 0;
        } else {
            // find the reduced partition information
            // i.e., vector of cluster labels;
            Rindx1 = 0;
            for(jj = 0; jj < *nsubject; jj++){
                if(gamma_iter[jj*ntime1 + (t)] == 1){
                    if(jj != j){
                        Si_tmp[Rindx1] = Si_iter[jj*ntime1 + (t)];
                        Si_tmp2[Rindx1] = Si_iter[jj*ntime1 + (t)];
                        compt1[Rindx1] = Si_iter[jj*ntime1 + (t-1)];
                    }
                }
            }
            // Also get the reduced spatial coordinates if
            // space is included
            if(*sPPM==1){
                if((*space_1==1 & t == 0) | (*space_1==0)){
                    s1_red[Rindx1] = s1[jjj];
                    s2_red[Rindx1] = s2[jjj];
                }
            }
        }
    }
}
// Prior parameter values
// priors for alpha
double a = alphaPriors[0], b = alphaPriors[1];
// priors for eta1
double b_eta1 = modelPriors[5];
// DP weight parameter
double Map = *M;
Rprintf("Prior values: Asig = %.2f, Atau = %.2f, Alam = %.2f, \n m0 = %.2f, s20
= %.2f\n", Asig, Atau, Alam, m0, s20);
// Cohesion auxiliary model parameters for Cohesions 3 and 4
double k0=cParams[1], v0=cParams[2];
double *mu0 = R_VectorInit(2,cParams[0]);
double *L0 = R_VectorInit(2*2,0.0);
L0[0] = cParams[3]; L0[3] = cParams[3];
Rindx1 = Rindx1 + 1;
}

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        rho_comp = compatibility(Si_red_1, comptm1, Rindx1+1);
    }
}

Si_tmp2[Rindx1] = Si_iter[j*ntime1 + (t)];
comptm1[Rindx1] = Si_iter[j*ntime1 + (t-1)];
n_red = Rindx1;
n_red_1 = Rindx1 + 1;
relabel(Si_tmp, *nssubject, Si_red, reorder, oldLab);
relabel(Si_tmp2, *nssubject, Si_red_1, reorder, oldLab);
// I need to keep the relabeled cluster label for the pegged
// individual so that I know what lgweight to keep in the
// full conditional.
cit_1 = Si_red_1[Rindx1];
}

for(jj = 0; jj < n_red_1; jj++){
    nh_red[jjj]=0;
    nh_red_1[jjj]=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];
lCo=0.0, lCn=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[jjj];
                    s2o_indx = s2_red[jjj];
                    s1n_indx = s1_red[jjj];
                    s2n_indx = s2_red[jjj];
                    indx = indx+1;
                }
            }
            lCo = Cohesion3_4(s1o, s2o, mu0, k0, v0, L0, nh_red[k], *SpatialCohesion, 1);
            lCn = Cohesion3_4(s1n, s2n, mu0, k0, v0, L0, nh_red[k+1], *SpatialCohesion, 1);
            nh_red_1[Rindx1] = Log(nh_red[k] + 1Cn - 1Co;
            Si_red_1[Rindx1] = k+1;
        }
    }
}
rho_comp = compatibility(Si_red_1, comptm1, Rindx1+1);
if(rho_comp==0) lgweight[k] = log(0);
lCn_1=0;
if(*sppm==1{
    if((*space_1==1 & t == 0) | (*space_1==0)) {
        s1o[0] = s1[j];
        s2o[0] = s2[j];
        lCn_1 = Cohesion3_4(s1o, s2o, mu0, k0, v0, L0, 1, *SpatialCohesion, 1);
    }
}
lgweight[nclus_red] = Log(Mdp) + lCn_1;
Si_red_1[Rindx1] = nclus_red+1;
rho_comp = compatibility(Si_red_1, comptm1, Rindx1+1);
if(rho_comp == 0) lgweight[nclus_red] = log(0);
// maxph = 0.0;
for(k = 0; k < nclus_red + 1; k++){
    denph = 0.0;
    phtmp[k] = lgweight[k];
}
R_rsort(phtmp, nclus_red + 1);
maxph = phtmp[nclus_red];
denph = 0.0;
for(k = 0; k < nclus_red + 1; k++){
    lgweight[k] = exp(lgweight[k] - maxph);
    denph = denph + lgweight[k];
}
for(k = 0; k < nclus_red + 1; k++){
    lgweight[k] = lgweight[k]/denph;
}
probh[1] = alpha_iter[t]/(alpha_iter[t] + (1-alpha_iter[t])*lgweight[cit_1-1]);
// If gamma is 1 at current MCMC iterate, then there are no
// concerns about partitions being incompatible as gamma changes
// from 1 to 0.
// However, if gamma's current value is 0, then care must be taken when
// trying to change from gamma=0 to gamma=1 as the partitions may
// 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"
}

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// 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} = 1
// when gamma_{it} = 1;
Rindx1 = 0;
for(jj = 0; jj < *nsubject; jj++){
    if(gamma_iter[jj*ntime1 + (t)] == 1){
        compt1[Rindx1] = Si_iter[jj*ntime1 + (t-1)];
        comp1t[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){
    compt1[Rindx1] = Si_iter[jj*ntime1 + (t-1)];
    comp1t[Rindx1] = Si_iter[jj*ntime1 + (t)];
    Rindx1 = Rindx1 + 1;
}

rho_comp = compatibility(comptmt, comp1t, 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;
}

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

/////////////////////////////////////////////////////////////////
// update partition
/////////////////////////////////////////////////////////////////
// The cluster probabilities depend on four partition probabilities
// leaning towards Pr(rho_t+1) and Pr(rho_t+1.R) can be absorbed. But I need
// to use rho_t.R and rho_t+1.R to check compatibility as I update rho_t.
/////////////////////////////////////////////////////////////////
// I have switched a number of times on which of these needs to be computed
// and which one can be absorbed in the normalizing constant. Right now I am
// leaning towards Pr(rho_t+1) and Pr(rho_t+1.R) can be absorbed. But I need
// to use rho_t.R and rho_t+1.R to check compatibility as I update rho_t.
/////////////////////////////////////////////////////////////////
// It seems to me that I can use some of the structure used to carry

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// out Algorithm 8 from previous code to keep track of empty clusters
// etc.
for(j = 0; j < *nsubject; j++){
    // Only need to update partition relative to units that are not pegged
    if(gamma_iter[j*(ntime1) + t] == 0){
        if(nh[(Si_iterjj*(ntime1) + t)-1]*(ntime1) + t] > 1){

            // Observation belongs to a non-singleton ...
            nh[(Si_iter[j*(ntime1) + t]-1)*(ntime1) + t] = 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;
                // All members of last cluster will be assigned subject j's label
                for(jj = 0; jj < *nsubject; jj++){
                    if(Si_iter[jj*(ntime1) + t] == nclus_iter[t]){

                        Si_iter[jj*(ntime1) + t] = iaux;

                    }
                }
            }

            rho_comp = compatibility(comptmt, comp1t, 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;
            }

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

/////////////////////////////////////////////////////////////////
// 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;
nclus_iter[t] = nclus_iter[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;

}

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    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];
        comptp1[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.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;
}

}

1Cn = Cohesion3_4(s1n, s2n, mu0, k0, v0, l0,
nh_tmp[kk], *SpatialCohesion, 1);
}
// End of spatial part

// Beginning of spatial part
1Cn;
// 1Cn;
1Cn;

lpp = lpp + nclus_tmp*log(Mdp) + lgamma((double) nh_tmp[kk]) +
1Cn;
lpp = lpp + nh_tmp[kk]*log(Mdp) + lgamma((double) nh_tmp[kk]) +
1Cn;

lpp = lpp + (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](*ntime1 + t],
    muh[k*(ntime1) + t],
    sqrt(sig2h[k*(ntime1) + t]), 1) +
lpp;

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

ph[k] = dnorm(y[j](*ntime1 + t],
    muh[k*(ntime1) + t] +
    eta1_iter[j]*y[j](*ntime1 + t-1],
    sqrt(sig2h[k*(ntime1) + t],
        (1-eta1_iter[j]*eta1_iter[j]), 1)) +
lpp;

}

// use this to test if MCMC draws from prior are correct
ph[k] = lpp;
}

}

// determine 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){
        comp2t[Rindx2] = rho_tmp[jj];
        comptp1[Rindx2] = Si_iter[jj*ntime1 + (t+1)];
        Rindx2 = Rindx2 + 1;
    }
}

// check for compatibility
rho_comp = compatibility(comp2t, comptp1, Rindx2);
if(rho_comp != 1){
    if[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;
        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;
        }
        lpp = 0.0;
        for(kk = 0; kk < nclus_tmp; kk++){
            // Beginning of spatial part
            lCn = 0.0;
            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[jj];
                            s2n[indx] = s2[jj];
                            indx = indx+1;
                        }
                    }
                    lCn = Cohesion3_4(sin, s2n, mutheta, ktheta, vtheta, Ltheta);
                }
                nh_tmp[kk],*SpatialCohesion, 1);
            }
            // End of spatial part
            lpp = lpp + Log(Mdp) + Gamma(double) nh_tmp[kk] + lCn;
            lpp = lpp + nh_tmp[kk]*log(Map) + log((double) nh_tmp[kk]) + lCn;
            if(t==0){
                ph[nclus_iter[t]] = dnorm(y[j*(ntime) + t], mudraw, sigdraw, 1) +
                lpp;
                if(t > 0){
                    ph[nclus_iter[t]] = dnorm(y[j*(ntime) + t], mudraw + eta1_iter[j]*y[j*(ntime) + t-1], mudraw + eta1_iter[j]*y[j*(ntime) + t-1], sigdraw*sqrt(1-eta1_iter[j]*eta1_iter[j]));
                }
            }
            // Now compute the probabilities
            for(k = 0; k < nclus_iter[t]+1; k++){
                phtmp[k] = ph[k];
            }
            R_rsort(phtmp, nclus_iter[t]+1);
            maxph = phtmp[nclus_iter[t]];
            denph = 0.0;
            for(k = 0; k < nclus_iter[t]+1; k++){
                ph[k] = exp(ph[k] - maxph);
                denph = denph + ph[k];
            }
            probh[k] = ph[k]/denph;
        }
        uu = runif(0.0,1.0);
        cprobh= 0.0;
        for(k = 0; k < nclus_iter[t]+1; k++){
            cprobh = cprobh + probh[k];
            if (uu < cprobh){
                iaux = k+1;
                break;
            }
        }
        if(iaux <= nclus_iter[t]){
            Si_iter[j*(ntime1) + t] = iaux;
            nh[(Si_iter[j*(ntime1) + t]-1)*(ntime1)+t] = nh[(Si_iter[j*(ntime1) + t]-1)*(ntime1)+t] + rho_tmp[j] + 1;
            rho_tmp[j] = iaux;
        }else{
            nclus_iter[t] = nclus_iter[t] + 1;
            Si_iter[j*(ntime1) + t] = nclus_iter[t];
            nh[(Si_iter[j*(ntime1) + t]-1)*(ntime1)+t] = 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;
        }
    }
    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, *nssubject, Si_tmp2, reorder, oldLab);

for(jj=0; jj<*nssubject; jj++){
    Si_iter[jj*(ntime1) + t] = Si_tmp2[jj];
}

for(k = 0; k < nclus_iter[t]; k++){
    mu_tmp[k] = muh[k*(ntime1)+t];
    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] = muh_tmp[(oldLab[k]-1)];
    sig2h[k*(ntime1)+t] = sig2_tmp[(oldLab[k]-1)];
}

for(j = 0; j < *nssubject; j++){
    Si_tmp[j] = Si_iter[j*(ntime1) + t];
    Si_tmp2[j] = 0;
    reorder[j] = 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, *nssubject, Si_tmp2, reorder, oldLab);

for(j=0; j<*nssubject; j++){
    Si_iter[j*(ntime1) + t] = Si_tmp2[j];
}

for(k = 0; k < nclus_iter[t]; k++){
    muh[k*(ntime1)+t] = reorder[k];
    sig2h[k*(ntime1)+t] = sig2h[k*(ntime1)+t];
}

for(k = 0; k < nclus_iter[t]; k++){
    nh[k*(ntime1)+t] = reorder[k];
    muh[k*(ntime1)+t] = muh_tmp[(oldLab[k]-1)];
    sig2h[k*(ntime1)+t] = sig2h_tmp[(oldLab[k]-1)];
}

for(k = 0; k < nclus_iter[t]; k++){
    sumy = 0.0;
    for(j = 0; j < *nssubject; j++){
        if(Si_iter[j*(ntime1) + t] == k+1){
            // update muh
            // sumy = sumy + y[j*(ntime1)+t];
        }
    }
}

if(t==0){
    sumy = 0.0;
    for(j = 0; j < *nssubject; j++){
        if(Si_iter[j*(ntime1) + t] == k+1){
            // update muh
            // sumy = sumy + y[j*(ntime1)+t];
        }
    }
}

```



```

        (1-e1n*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));

    // update phi0
    //

    phisq = phi1_iter*phi1_iter;
    one_phisq = (1-phi1_iter)*(1-phi1_iter);
    lam2tmp = lam2_iter*(1.0 - phi1sq);
    sumt = 0.0;
    for(t=1; t<*ntime; t++){
        sumt = sumt + (theta_iter[t] - phi1_iter*theta_iter[t-1]);
    }

    s2star = 1.0/(*ntime-1)*(one_phisq/lam2tmp) + (1/lam2_iter) + (1/s20);
    mstar = s2star*((1.0-phi1_iter)/lam2tmp)*sumt + (1/lam2_iter)*theta_iter[0] +
    (1/s20)*m0;

}

// update alpha
//

if(*update_alpha == 1){
    if(*time_specific_alpha != 1){
        sumg = 0;
        for(j = 0; j < *nsubject; j++){
            for(t = 1; t < *ntime; t++){
                sumg = sumg + gamma_iter[j*ntime1 + t];
            }
        }
        astar = (double) sumg + a;
        bstar = (double) (*nsubject)*(*ntime-1) - sumg + b;
    }

    alpha_tmp = rbeta(astar, bstar);
    for(t=0;t<*ntime;t++)alpha_iter[t] = alpha_tmp;
}
else {
    for(t = 0; t < *ntime; t++){
        sumg = 0;
        for(j = 0; j < *nsubject; j++){
            sumg = sumg + gamma_iter[j*ntime1 + t];
        }
        astar = (double) sumg + a;
        bstar = (double) (*nsubject) - sumg + b;
    }

    alpha_iter[t] = rbeta(astar, bstar);
}

astar = (double) sumg + a;
bstar = (double) (*nsubject) - sumg + b;
alpha_iter[t] = rbeta(astar, bstar);
}
else {
    for(t = 0; t < *ntime; t++){
        sumg = 0;
        for(j = 0; j < *nsubject; j++){
            sumg = sumg + gamma_iter[j*ntime1 + t];
        }
        astar = (double) sumg + a;
        bstar = (double) (*nsubject) - sumg + b;
    }

    alpha_iter[t] = rbeta(astar, bstar);
}

// update phi0_iter = rnorm(mstar, sqrt(s2star));
//
// update phi1
//
if(*update_phi1==1{
    op1 = phi1_iter;
    np1 = rnorm(op1, csigPHI1);

    if(np1 > -1 & np1 < 1){
        llo = 0.0, lln = 0.0;
        for(t=1; t < *ntime; t++){
            llo = llo + dnorm(theta_iter[t], phi0_iter*(1.0 - op1) + op1*theta_iter[t-1],
            sqrt(lam2_iter*(1.0 - op1)), 1);
            lln = lln + dnorm(theta_iter[t], phi0_iter*(1.0 - np1) + np1*theta_iter[t-1],
            sqrt(lam2_iter*(1.0 - np1)), 1);
        }

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

// update lambda with a MH step
phisq = phi1_iter*phi1_iter;
o1 = sqrt(lam2_iter);
n1 = rnorm(o1, csigLM);
}
alpha_iter[0] = 0.0;
}

```



```

        predSi_iter[j>(*npred) + p] = nclus_tmp;
        nh_pred[predSi_iter[j>(*npred) + p]-1](*npred)+p] = 1;
    }
    n_red = n_red + 1;
}
}

/////////////////////////////////////////////////////////////////
// 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_1ii>(*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];
    }
    for(j=0; j<*nsubject; j++) {
        eta1[ii(*nsubject) + j] = eta1_iter[j];
    }
    phi1[ii] = phi1_iter;
    phi0[ii] = phi0_iter;
    lam2[ii] = lam2_iter;
    ii = ii+1;
}
/* */
}

lpm1_iter=0.0;
for(t = 0; t < *ntime; t++){
    for(j = 0; j < *nsubject; j++){
        lpm1_iter = lpm1_iter - Log((1/(double) nout-nout_theta)*CP0[j(*ntime)+t]);
    }
    lpm1[0] = lpm1_iter;
    elppdWAIC = 0.0;
}

for(j = 0; j < *nsubject; j++){
    for(t = 0; t < *ntime; t++){
        elppdWAIC = elppdWAIC + (2*mnlle[j(*ntime)+t] - Log(mnlle[j(*ntime)+t]));
    }
}
waic[0] = -2*elppdWAIC;
PutRNGstate();
}

// These are needed for WAIC
mnlle[j(*ntime)+t] = mnlle[j(*ntime)+t] + exp(like_iter[j*
(*ntime)+t])(double) nout;
mnlle[j(*ntime)+t] = mnlle[j(*ntime)+t] + (like_iter[j*
(*ntime)+t])(double) nout;

if(exp(like_iter[j(*ntime)+t]) < 1e-320) like0=1;
}

}

if(like0==1) nout_theta = nout_theta + 1;
if(like0==0){
    for(j = 0; j < *nsubject; j++){
        for(t = 0; t < *ntime; t++){
            CP0[j(*ntime)+t] = CP0[j(*ntime)+t] + (1/exp(like_iter[j*
(*ntime)+t]));
        }
    }
}

```