

**Supplementary Material of the paper:
A Data-Driven Allocation Sampler for Making Inference
on the Number of Clusters in the Dirichlet allocation
Model via Bayesian Mixture Modelling**

This supplementary material (SM) presents the acceptance probability for split-merge movements and some additional results obtained with application of the proposed SDAS algorithm to the simulated datasets.

Appendix 1: Acceptance Probability

From equation (14) of the paper, the acceptance probability for a split movement is $\Psi[\Phi^{sp}|\Phi] = \min(1, A^{sp})$, where

$$A^{sp} = \frac{P(\mathbf{y}|\mathbf{c}^{sp}, \boldsymbol{\theta}_{k_c^{sp}}, k_{\mathbf{c}^{sp}}) \pi(\mathbf{c}^{sp}|\gamma) \pi(\boldsymbol{\theta}^{sp}|\eta, k_{\mathbf{c}^{sp}})}{P(\mathbf{y}|\mathbf{c}, \boldsymbol{\theta}_{k_c}, k_c) \pi(\mathbf{c}|\gamma) \pi(\boldsymbol{\theta}^{sp}|\eta, k_c)} \frac{\prod_{j=k_{\mathbf{c}^{sp}}+1}^n \pi(\theta_j|\eta_j)}{\prod_{j=k_c+1}^n \pi(\theta_j|\eta_j)} \frac{q[\Phi|\Phi^{sp}]}{q[\Phi^{sp}|\Phi]},$$

The ratio of the joint probabilities of \mathbf{y} conditional on latent indicator variables and component parameters is given by

$$\frac{P(\mathbf{y}|\mathbf{c}^{sp}, \boldsymbol{\theta}_{k_c^{sp}}, k_{\mathbf{c}^{sp}})}{P(\mathbf{y}|\mathbf{c}, \boldsymbol{\theta}_{k_c}, k_c)} = \frac{L(\theta_{j_1}^{sp}|D_{j_1}) L(\theta_{j_2}^{sp}|D_{j_2})}{L(\theta_j|D_j)}, \quad (20)$$

where $L(\theta_m|D_m) = \prod_{D_m} f(y_i|\theta_m)$ is the likelihood function for component m , for $m = \{j, j_1, j_2\}$.

From Equation (7) of the paper, the ratio of the joint prior probability for latent indicator variables is

$$\frac{\pi(\mathbf{c}^{sp}|\gamma, k_{\mathbf{c}^{sp}})}{\pi(\mathbf{c}|\gamma, k_c)} = \frac{\Gamma(n_{j_1})\Gamma(n_{j_2})}{\Gamma(n_j)}. \quad (21)$$

The prior distributions ratio for component parameters is

$$\frac{\pi(\boldsymbol{\theta}_n|\boldsymbol{\eta}, k_{\mathbf{c}^{sp}})}{\pi(\boldsymbol{\theta}_n|\boldsymbol{\eta}, k_c)} = \frac{\prod_{j=1}^n \pi_G(\theta_j^{sp}|\eta_j)}{\prod_{j=1}^n \pi_G(\theta_j|\eta_j)} = \frac{\pi_G(\theta_{j_1}^{sp}|\eta_{j_1}) \pi_G(\theta_{j_2}^{sp}|\eta_{j_2})}{\pi_G(\theta_j|\eta_j) \pi_G(\theta_n|\eta_n)}. \quad (22)$$

From Equation (17) of the paper, the transition probability ratio for the split proposal is

$$\frac{q[\Phi|\Phi^{sp}]}{q[\Phi^{sp}|\Phi]} = \frac{P_{me|k_c^{sp}} P_{j_1, j_2|k_c^{sp}}}{P_{sp|k_c}} \frac{1}{P_{j|\kappa_2}} \frac{1}{P_{alloc}^{sp}} \frac{\pi(\theta_j|D_j, \eta_j)\pi(\theta_n|\eta_n)}{\pi(\theta_{j_1}^{sp}|D_{j_1}, \eta_{j_1})\pi(\theta_{j_2}^{sp}|D_{j_2}, \eta_{j_2})}, \quad (23)$$

where

$$\pi(\theta_m|D_m, \eta_m) = \frac{L(\theta_m|D_m)\pi_G(\theta_m|\eta_m)}{\int L(\theta_m|D_m)\pi_G(\theta_m|\eta_m)d\theta_m}$$

is the posterior distribution for component parameter θ_m , for $m = \{j, j_1, j_2\}$.

Multiplying (20), (21), (22) and (23), the acceptance probability for a split proposal is $\Psi[\Phi^{sp}|\Phi] = \min(1, A^{sp})$ for

$$A^{sp} = \frac{\mathbf{I}(D_{j_1})\mathbf{I}(D_{j_2})}{\mathbf{I}(D_j)} \frac{\Gamma(n_{j_1})\Gamma(n_{j_2})}{\Gamma(n_j)} \frac{Q^{sp}}{P_{alloc}},$$

where

$$Q^{sp} = \frac{P_{me|k_c^{sp}} P_{j_1, j_2}}{P_{sp|k_c} P_{j|\mathbb{C}_2}} = \begin{cases} \frac{1}{2} & , \text{ if } k_c = 1; \\ \left(\frac{1}{2}\right)^{1-\mathbb{I}_{k_c^{sp}}(n)} \frac{3\mathbb{C}_2}{k_{c+1}} & , \text{ if } k_c \in \mathbb{K}_1; \\ 2^{\mathbb{I}_{k_c^{sp}}(n)} \frac{\mathbb{C}_2}{k_{c+1}} & , \text{ if } k_c \in \mathbb{K}_2; \end{cases}$$

with $\mathbb{I}_{k_c^{sp}}(n)$ being an indicator function and the sets $\mathbb{K}_1 = \{2 \leq k_c \leq k-1; \text{ and } j_1 = 1 \text{ or } j_2 = k_c\}$, $\mathbb{K}_2 = \{2 \leq k_c \leq k-1 \text{ and } j_1 \neq 1 \text{ or } j_2 \neq k_c\}$.

Similarly, the acceptance probability for a merge is $\Psi[\Phi^{me}|\Phi] = \min(1, A^{me}) = \frac{1}{A^{sp}}$, but with some obvious differences in the substitutions.

Appendix 2: Inference on \mathbf{c} and θ_{k_c}

In this Appendix we describe the procedure used to define a configuration for \mathbf{c} and estimate parameters $\theta_{\mathbf{c}}$ conditional on estimate $\tilde{k}_{\mathbf{c}}$ for the number of clusters.

Thus, consider $\tilde{k}_{\mathbf{c}}$ be the estimate for the number of clusters. Following [23], let

- (i) $L_{\tilde{k}_{\mathbf{c}}} = \sum_{l=B+1}^L \mathbb{I}_{k_{\mathbf{c}}^{(l)}}(\tilde{k}_{\mathbf{c}})$ be the number of iterations for which $k_{\mathbf{c}} = \tilde{k}_{\mathbf{c}}$ in $L - B$ iterations, where $\mathbb{I}_{k_{\mathbf{c}}^{(l)}} = 1$ if in l -iteration $k_{\mathbf{c}}^{(l)} = \tilde{k}_{\mathbf{c}}$ and $\mathbb{I}_{k_{\mathbf{c}}^{(l)}} = 0$ otherwise;
- (ii) $N_{ij} = \sum_{l=B+1}^L \mathbb{I}_{c_i^{(l)}}(j)\mathbb{I}_{k_{\mathbf{c}}^{(l)}}$ be the number of times that observation y_i is allocated in component j in $L_{\tilde{k}_{\mathbf{c}}}$ iterations, where $\mathbb{I}_{c_i^{(l)}}(j) = 1$ if in l -th iteration $c_i = j$ and $\mathbb{I}_{c_i^{(l)}}(j) = 0$ otherwise, for $i = 1, \dots, n$ and $j = 1, \dots, k$.

Let $\tilde{P}(c_i = j) = \frac{N_{ij}}{L_{\tilde{k}_{\mathbf{c}}}}$ be the posterior probability that the observation y_i belongs to cluster j , for $j = 1, \dots, \tilde{k}_{\mathbf{c}}$. If $\tilde{P}(c_i = j) = \max_{1 \leq j' \leq \tilde{k}_{\mathbf{c}}} (\tilde{P}(c_i = j'))$, then we consider that y_i belongs to cluster j , for $i = 1, \dots, n$ and $j = 1, \dots, k_{\mathbf{c}}$.

We estimate parameters θ_j , $j = 1, \dots, \tilde{k}_{\mathbf{c}}$, considering the average of the generated values, *i.e.*,

$$\tilde{\theta}_j | \tilde{k}_{\mathbf{c}} = \frac{1}{L_{\tilde{k}_{\mathbf{c}}}} \sum_{l=B+1}^L \theta_j^{(l)} \mathbb{I}_{\tilde{k}_{\mathbf{c}}^{(l)}}(\tilde{k}_{\mathbf{c}}).$$

Appendix 3: Generated values

Figure 8 shows the values generated by cluster for datasets A_1 to A_4 for $n = 500$.

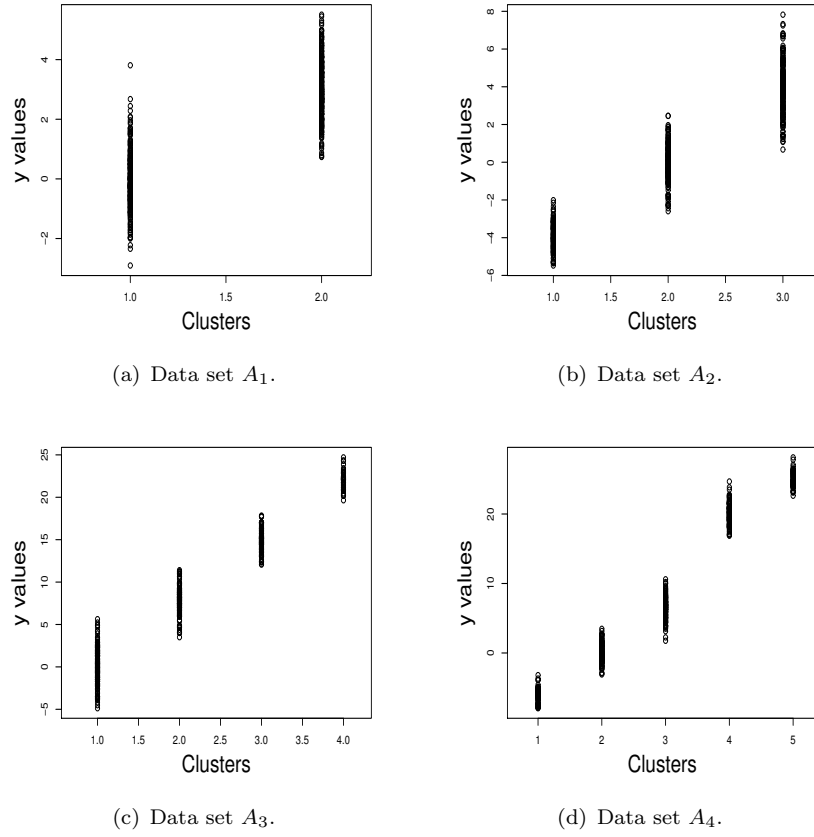


Figure 8. Values generated by cluster for datasets A_1 to A_4 with $n = 500$.

Appendix 4: Results from Artificial data sets

In this section, we present some additional results from simulation study. Figure 9 shows the generated values and the identified clusters by the SDAS algorithm for datasets A_1 to A_4 . As one can note, the cluster were satisfactorily identified.

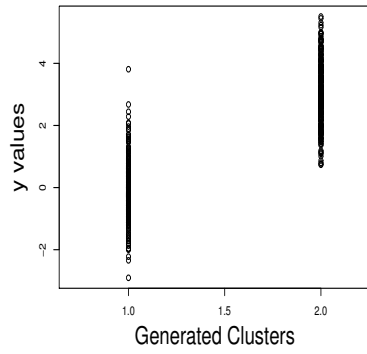
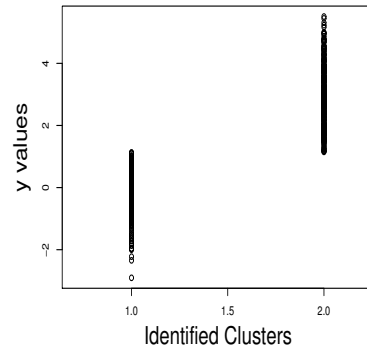
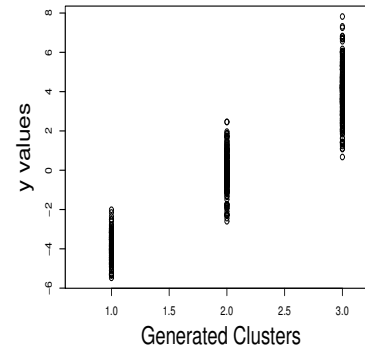
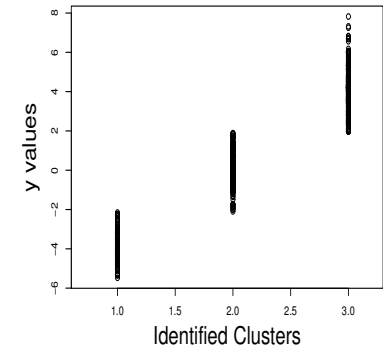
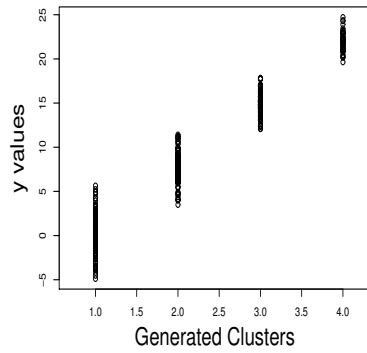
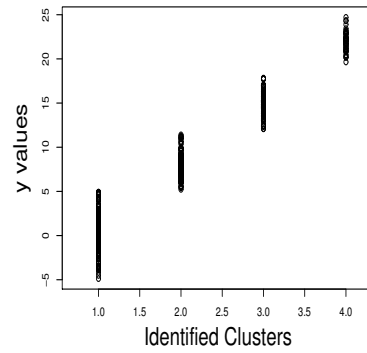
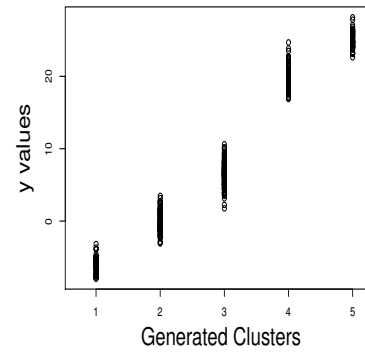
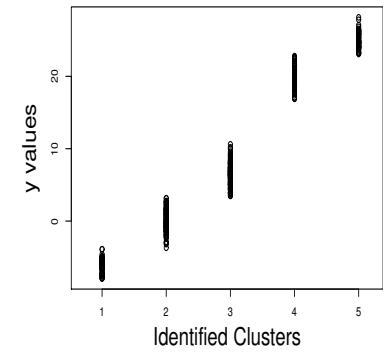
(a) Data set A_1 .(b) Data set A_1 .(c) Data set A_2 .(d) Data set A_2 .(e) Data set A_3 .(f) Data set A_3 .(g) Data set A_4 .(h) Data set A_4 .

Figure 9. Generated values by cluster and identified clusters by SDAS algorithm, datasets A_1 to A_4 .

Table 6 shows the estimates for component parameters of the identified clusters. Estimates for parameters were obtained according to Equation described in page 14, lines 300-302.

Table 6. Estimates for component parameters of the \tilde{k}_c clusters.

Parameter	Data set			
	A_1	A_2	A_3	A_4
μ_1	-0.1884 (-0.4040, 0.0498)	-3.8162 (-4.0087, -3.6235)	0.3656 (-0.0066, 0.7621)	-6.1188 (-6.3457, -5.8726)
μ_2	2.9146 (2.6994, 3.1167)	0.0951 (-0.1558, 0.3615)	8.0019 (7.6710, 8.3148)	0.0335 (-0.3356, 0.4308)
μ_3	-	4.0610 (3.7350, 4.3675)	14.9097 (14.5892, 15.1972)	6.5935 (6.2640, 6.9024)
μ_4	-	-	22.0472 (21.7556, 22.3370)	20.0996 (19.7348, 20.5063)
μ_5	-	-	-	24.9200 (24.4953, 25.2850)
σ_1^2	0.8015 (0.5614, 1.1131)	0.6664 (0.4586, 0.9363)	4.9954 (3.7895, 6.4985)	0.8813 (0.6056, 1.2785)
σ_2^2	1.1406 (0.8754, 1.5045)	1.1427 (0.6555, 1.9545)	2.2255 (1.4298, 3.3258)	2.6316 (1.6968, 4.0146)
σ_3^2	-	1.6561 (1.1567, 2.3125)	1.9841 (1.4088, 2.8251)	2.4508 (1.7900, 3.3486)
σ_4^2	-	-	1.1314 (0.7625, 1/6702)	2.2023 (1.5074, 3.2496)
σ_5^2	-	-	-	1.4701 (0.9190, 2.3820)

Figures 10 and 11 show the histogram of the observed data and the estimated density function. As one can note the results are satisfactory.

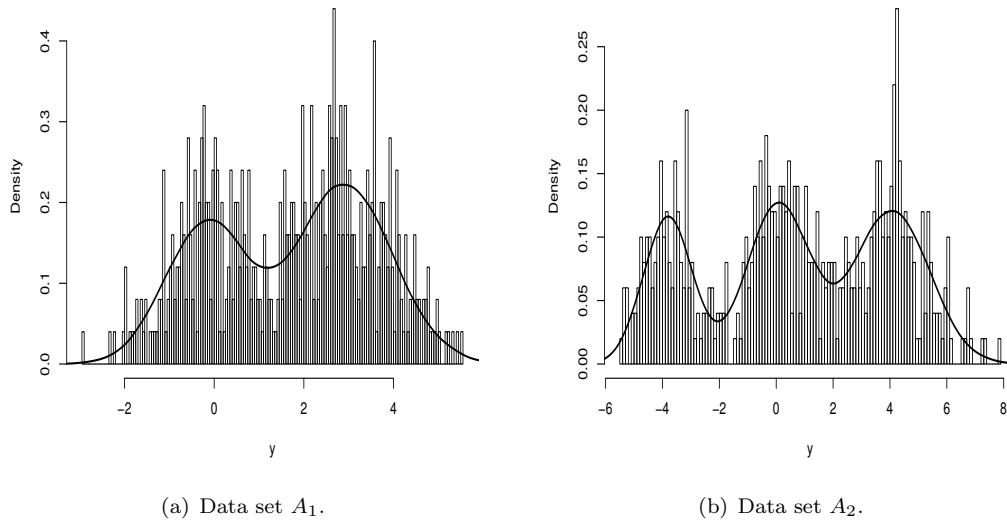


Figure 10. Histogram of observed data and Estimated density function for datasets A_1 and A_2 .

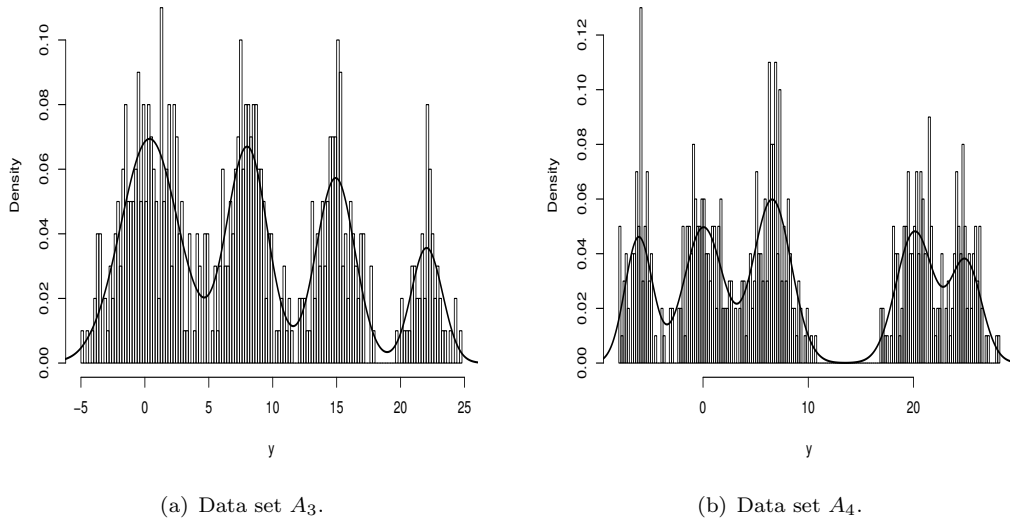


Figure 11. Histogram of observed data and Estimated density function for datasets A_3 and A_4 .

Appendix 4: SDAS codes for Galaxy Dataset

In this Appendix, we present the R codes used in the application of the SDAS algorithm to the Galaxy dataset. The codes in format .txt can be obtained by emailing the first author.

```

rm(list = ls(all = TRUE))
set.seed(19)
library(compiler)
enableJIT(3)
library(MCMCpack)
L = 5000
B = 5000
salto = 6
tx.acs = 0
tx.acm = 0
km = 10
y = c(9172, 9350, 9483, 9558, 9775, 10227, 10406, 16084, 16170, 18419, 18552, 18600, 18927, 19052, 19070, 19100)
n = length(y)
y = y/1000
gama = 0.1
mu0 = (min(y) + max(y))/2
lamb = 0.01
halfa = 1
R = max(y) - min(y)
hbeta = halfa/(R2)
ID = function(dx, nx){
  DA1 = -(nx/2) * log(2 * hbeta * pi)
  DA2 = 0.5 * log(lamb) - 0.5 * log(nx + lamb)
}

```

```

DA3 = lgamma(halfa + ((nx + 1)/2)) - lgamma(halfa)
DA4 = -(halfa + (nx/2)) * log(1 + ((sum(dx^2) + (lamb * (mu0^2)))/(2 * hbeta)) -
((sum(dx) + (lamb * mu0))^2)/(2 * hbeta * (nx + lamb)))
(DA1 + DA2 + DA3 + DA4) }
KL = function(dx1, dx){
  nx1 = length(dx1)
  nx = length(dx)
  alfa1e = halfa + ((nx1 + 1)/2)
  alfa0e = halfa + ((nx + 1)/2)
  beta1e = hbeta + ((sum(dx1^2) + (lamb * (mu0^2)))/2) - (((sum(dx1) + (lamb *
mu0))^2)/(2 * (nx1 + lamb)))
  beta0e = hbeta + ((sum(dx^2) + (lamb * (mu0^2)))/2) - (((sum(dx) + (lamb *
mu0))^2)/(2 * (nx + lamb)))
  mi1e = (sum(dx1) + (lamb * mu0))/(nx1 + lamb)
  mi0e = (sum(dx) + (lamb * mu0))/(nx + lamb)
  KL1 = 0.5 * log(nx1 + lamb) - 0.5 * log(nx + lamb) - 0.5 + ((nx + lamb)/(2 * (nx1 +
lamb)))
  KL2 = (((nx + lamb) * ((mi1e - mi0e)^2))/2) * (alfa1e/beta1e)
  KL3 = alfa0e * log(beta1e) - alfa0e * log(beta0e)
  KL4 = lgamma(alfa0e) - lgamma(alfa1e)
  KL5 = (alfa1e - alfa0e) * digamma(alfa1e)
  KL6 = -(beta1e - beta0e) * (alfa1e/beta1e)
  (KL1 + KL2 + KL3 + KL4 + KL5 + KL6) }
mu = rep(1000, km) sig = rep(1000, km)
kc = numeric()
z = matrix(0, nrow = n, ncol = km + 1)
ind.y = matrix(0, nrow = n, ncol = km)
nj = rep(0, km)
syj = rep(0, km)
syj2 = rep(0, km)
prop = matrix(0, nrow = n, ncol = km)
kc[1] = 1
z[, 1] = 1
ind.y[, 1] = y
nj[1] = n
syj[1] = sum(y)
syj2[1] = sum(y^2)
mu[1] = mean(y)
sig[1] = var(y)
kc.est = numeric()
kc.est[1] = 1
l0 = B + 1
l1 = 1
for(lin2 : L){
  zt = z
  for(iin1 : n){
    p = rep(0, km)
    for(jin1 : kc){
      if(z[i, j] == 1 & nj[j] > 1)
        p[j] = ((nj[j] - 1)/(n + gama - 1)) * dnorm(y[i], mu[j], sqrt(sig[j]))

```

```

if(z[i,j] == 0 & nj[j] > 0)
p[j] = (nj[j]/(n + gama - 1)) * dnorm(y[i], mu[j], sqrt(sig[j])) }
p0 = (gama/(n + gama - 1)) * exp(ID(y[i], 1))
b = 1/(sum(p) + p0)
Pr = c(b * p, b * p0)
z[i,] = rmultinom(1, 1, Pr)
if(z[i, (km + 1)] == 0)
ind.y[i,] = z[i, 1 : km] * y[i] }
if(z[i, (km + 1)] == 1) {
kc = kc + 1
z[i,] = 0
z[i, kc] = 1
ind.y[i,] = 0
ind.y[i, kc] = y[i]
alfaj = halfa + 1
betaj = hbeta + (((y[i]^2) + (lamb*(mu0^2)))/2) - ((y[i] + lamb*mu0)^2)/(2*(1+lamb))
sig[kc] = 1/rgamma(1, alfaj, betaj)
mu[j] = (y[i] + lamb * mu0)/(1 + lamb)
sig[j] = sig[kc]/(1 + lamb)
mu[kc] = rnorm(1, mu[j], sqrt(sig[j])) }
for(jin1 : kc){
nj[j] = sum(z[, j]) } }
kc = 0
for(jin1 : km){
nj[j] = sum(z[, j])
if(nj[j] > 0)
kc = kc + 1
if(nj[j] == 0) {
mu[j] = 1000
sig[j] = 1000 } }
mu.ord = sort(mu)
z.ord = z
sig.ord = sig
indy.ord = ind.y
for(gin1 : kc){
for(win1 : km){
if(mu.ord[g] == mu[w]) {
sig.ord[g] = sig[w]
z.ord[, g] = z[, w]
indy.ord[, g] = ind.y[, w] } } }
mu = mu.ord
sig = sig.ord
z = z.ord
ind.y = indy.ord
mu[(kc + 1) : km] = 1000
sig[(kc + 1) : km] = 1000
for(gin1 : kc){
nj[g] = sum(z[, g])
syj[g] = sum(ind.y[, g])
syj2[g] = sum(ind.y[, g]^2) }

```



```

sp < -0.5
me < -0.5
if(kc == 1) {
  sp = 1
  mg = 0 }
ind.mov = rbinom(1, 1, sp)
if(ind.mov == 1) {
  if(kc == 1) {
    k1 = 1
    k2 = 2 }
  if(kc > 1) {
    k1 = sample(seq(1, kc, 1), 1) k2 = k1 + 1 }
  if(nj[k1] > 1) {
    kc = kc + 1
    ind.ysp = ind.y
    i1 = k2 + 1
    i2 = km
    ind.ysp[, i1 : i2] = ind.y[, k2 : (km - 1)]
    ind.ysp[, k1 : k2] = 0
    z.sp = z
    z.sp[, i1 : i2] = z[, k2 : (km - 1)]
    z.sp[, k1 : k2] = 0
    y.sp = ind.y[, k1]
    y.sp = y.sp[y.sp! = 0]
    isp = match(y.sp, ind.y[, k1])
    ns = length(y.sp)
    Palloc < -1
    if(ns > 2) {
      cl = rep(1, ns)
      ymin = min(y.sp)
      ymax = max(y.sp)
      Dj1 = ymin
      Dj2 = ymax
      nj1 = 1
      nj2 = 1
      Pm = rep(1, ns)
      for(iin1 : ns){
        if(y.sp[i] == ymax)
          cl[i] < -2
        if(y.sp[i]! = ymin & y.sp[i]! = ymax) {
          xj1 = c(Dj1, y.sp[i]) xj2 = c(Dj2, y.sp[i]) pj1 = KL(xj1, Dj1) pj2 =
          KL(xj2, Dj2) P1 = pj2/(pj1 + pj2)
          P2 = 1 - P1
          Pm[i] = P1
          Im = rbinom(1, 1, P1)
          if(Im == 1) {
            Dj1 = c(Dj1, y.sp[i])
            nj1 < -nj1 + 1 }
          if(Im == 0) {
            Dj2 = c(Dj2, y.sp[i])

```

```

cl[i] = 2
Pm[i] = P2
nj2 = nj2 + 1 } } }
Palloc < -prod(Pm) }
if(ns == 2)
cl = c(1, 2)
for(j1in1 : ns){
if(cl[j1] == 1) {
ind.yisp[isp[j1], k1] = y[isp[j1]]
z.sp[isp[j1], k1] = 1 }
if(cl[j1] == 2) {
ind.yisp[isp[j1], k2] = y[isp[j1]]
z.sp[isp[j1], k2] = 1 } }
nj.sp = nj
syj.sp = syj
syj2.sp = syj2
for(gin1 : kc){
nj.sp[g] = sum(z.sp[, g])
syj.sp[g] = sum(ind.yisp[, g])
syj2.sp[g] = sum(ind.yisp[, g]2) }
mu.sp = mu
sig.sp = sig
mu.sp[i1 : i2] = mu[k2 : (km - 1)]
sig.sp[i1 : i2] = sig[k2 : (km - 1)]
mu.sp[k1 : k2] = 0
sig.sp[k1 : k2] = 0
alfaj = halfa + ((nj.sp[k1] + 1)/2)
betaj = hbeta + ((syj2.sp[k1] + (lamb * (mu02))))/2 - (((syj.sp[k1] + (lamb *
mu0)2))/(2 * (nj.sp[k1] + lamb)))
sig.sp[k1] = 1/rgamma(1, alfaj, betaj)
alfaj = halfa + ((nj.sp[k2] + 1)/2)
betaj = hbeta + ((syj2.sp[k2] + (lamb * (mu02))))/2 - (((syj.sp[k2] + (lamb *
mu0)2))/(2 * (nj.sp[k2] + lamb)))
sig.sp[k2] = 1/rgamma(1, alfaj, betaj)
muj = (syj.sp[k1] + lamb * mu0)/(nj.sp[k1] + lamb)
sigj = sig.sp[k1]/(nj.sp[k1] + lamb)
mu.sp[k1] = rnorm(1, muj, sqrt(sigj))
muj = (syj.sp[k2] + lamb * mu0)/(nj.sp[k2] + lamb)
sigj = sig.sp[k2]/(nj.sp[k2] + lamb)
mu.sp[k2] = rnorm(1, muj, sqrt(sigj))
if(mu.sp[k1] > mu.sp[k2]) {
zz1 = z.sp[, k2]
zz2 = z.sp[, k1]
iy1 = ind.yisp[, k2]
iy2 = ind.yisp[, k1]
m1 = mu.sp[k2]
m2 = mu.sp[k1]
si1 = sig.sp[k2]
si2 = sig.sp[k1]
nj1 = nj.sp[k2]

```

```

nj2 = nj.sp[k1]
sy1 = syj.sp[k2]
sy2 = syj.sp[k1]
sy21 = syj2.sp[k2]
sy22 = syj2.sp[k1]
z.sp[, k1] = zz1
z.sp[, k2] = zz2
ind.ysp[, k1] = iy1
ind.ysp[, k2] = iy2
mu.sp[k1] = m1
mu.sp[k2] = m2
sig.sp[k1] = si1
sig.sp[k2] = si2
nj.sp[k1] = nj1
nj.sp[k2] = nj2
syj.sp[k1] = sy1
syj.sp[k2] = sy2
syj2.sp[k1] = sy21
syj2.sp[k2] = sy22 }
ind.nad = 0
for(jin1 : kc){
if(mu.sp[j] > mu.sp[(j + 1)])
ind.nad = 1 }
As = 0
if(ind.nad == 0) {
Q.sp = 0.5
if((kc - 1) > 1) {
if(k1 == 1)
Q.sp = 3 * (kc - 1)/(2 * kc)
if(k1! = 1)
Q.sp = (kc - 1)/kc }
Ga = lgamma(nj.sp[k1]) + lgamma(nj.sp[k2]) - lgamma(nj[k1]) As =
Q.sp * (1/Palloc) * exp(ID(ind.ysp[, k1], nj.sp[k1]) + ID(ind.ysp[, k2], nj.sp[k2]) -
ID(ind.y[, k1], nj[k1]) + Ga) }
if(As == "NaN")
As = 0
alpha.s = min(1, As)
u.ts = runif(1)
if(u.ts < alpha.s) {
z = z.sp
ind.y = ind.ysp
mumu.sp
sig = sig.sp
tx.acs = tx.acs + 1
nj = nj.sp
syj = syj.sp
syj2 = syj2.sp }
if(u.ts >= alpha.s)
kc = kc - 1 } }
if(ind.mov == 0) {

```

```

kc = kc - 1
k1 <- sample(seq(1, kc, 1), 1)
if(k1 == 1)
k2 <- -2
if(k1 != 1) {
co <- rbinom(1, 1, 0.5)
if(co == 1) {
k2 = k1
k1 = k1 - 1 }
if(co == 0)
k2 = k1 + 1 }
z.me = z
z.me[, k1] = z[, k1] + z[, k2]
me2km - 1
z.me[, (k2 : me2)] = z[, (k2 + 1) : km]
ind.y.me = ind.y
ind.y.me[, k1] = ind.y[, k1] + ind.y[, k2]
ind.y.me[, k2 : me2] = ind.y[, (k2 + 1) : km]
nj.me = nj
syj.me = syj
syj2.me = syj2
nj.me[k1] = nj[k1] + nj[k2]
syj.me[k1] = syj[k1] + syj[k2]
syj2.me[k1] = syj2[k1] + syj2[k2]
nj.me[k2 : me2] = nj[(k2 + 1) : km]
syj.me[k2 : me2] = syj[(k2 + 1) : km]
syj2.me[k2 : me2] = syj2[(k2 + 1) : km]
mu.me = mu
sig.me = sig
mu.me[k2 : me2] = mu[(k2 + 1) : km]
sig.me[k2 : me2] = sig[(k2 + 1) : km]
alfaj = halfa + ((nj.me[k1] + 1)/2)
betaj = hbeta + ((syj2.me[k1] + (lamb * (mu02)))/2) - (((syj.me[k1] + (lamb *
mu0)2)/2 * (nj.me[k1] + lamb)))
sig.me[k1] = 1/rgamma(1, alfaj, betaj)
mu.j = (syj.me[k1] + lamb * mu0)/(nj.me[k1] + lamb)
sig.j = sig.me[k1]/(nj.me[k1] + lamb)
mu.me[k1] = rnorm(1, mu.j, sqrt(sig.j))
ind.nad <- -0
for(jin1 : kc){
if(mu.me[j] > mu.me[(j + 1)])
ind.nad <- -1 }
y.me1 = ind.y[, k1]
y.me1 = y.me1[y.me1! = 0]
y.min = min(y.me1)
y.me2 = ind.y[, k2]
y.me2 = y.me2[y.me2! = 0]
y.max = max(y.me2)
y.me = ind.y.me[, k1]
y.me = y.me[y.me! = 0]

```

```

isp = match(y.me, ind.y.me[, k1])
ns = length(y.me)
Palloc = 1
cl = rep(1, ns)
Dj1 = ymin
Dj2 = ymax
nj1 = 1
nj2 = 1
Pm = rep(1, ns)
for(iin1 : ns){
  if(y.me[i]! = ymin & y.me[i]! = ymax) {
    xj1 = c(Dj1, y.me[i])
    xj2 = c(Dj2, y.me[i])
    pj1 = KL(xj1, Dj1)
    pj2 = KL(xj2, Dj2)
    P1 = pj2/(pj1 + pj2)
    P2 = 1 - P1
    if(z[isp[i], k1] == 1) {
      Pm[i] < -P1
      Dj1 < -c(Dj1, y.me[i])
      nj1 < -nj1 + 1 }
    if(z[isp[i], k2] == 1) {
      Pm[i] = P2
      Dj2 = c(Dj2, y.me[i])
      cl[i] = 2
      nj2 = nj2 + 1 } }
  Palloc = prod(Pm)
  Am = 0
  if(ind.nad == 0) {
    if(kc == 1)
      Q.me = 2
    if(kc > 1) {
      if(k1 == 1)
        Q.me = 2 * (kc + 1) / (3 * kc)
      if(k1! = 1)
        Q.me = (kc + 1) / kc }
    Ga = lgamma(nj.me[k1]) - lgamma(nj[k1]) - lgamma(nj[k2])
    Am = Q.me * Palloc * exp(ID(ind.y.me[, k1], nj.me[k1]) - ID(ind.y[, k1], nj[k1]) -
ID(ind.y[, k2], nj[k2]) + Ga) }
    if(Am == "NaN")
      Am = 0
    alpha.m = min(1, Am)
    u.tm = runif(1)
    if(u.tm < alpha.m) {
      z = z.me
      ind.y = ind.y.me
      mu = mu.me
      sig = sig.me
      tx.acm = tx.acm + 1
      nj = nj.me

```

```

syj = syj.me
syj2 = syj2.me }
if(u.tm > alpha.m)
kc = kc + 1 }
if(l == l0){
kc.est[l1] = kc
l0 = l0 + salto
l1 = l1 + 1 }
if(kc == 3){
prop3 = prop3 + z[, 1 : 3]
mu.est = mu[1 : 3]
sig.est < -sig[1 : 3] }
mu.prop = mu
sig.prop = sig
ind.ord = 0
for(jin1 : kc){
alfaj = halfa + ((nj[j] + 1)/2)
betaj = hbeta + ((syj2[j] + (lamb * (mu02)))/2) - (((syj[j] + lamb * mu0)2)/(2 *
(nj[j] + lamb)))
sig.prop[j] = 1/rgamma(1, alfaj, betaj)
mu[j] = (syj[j] + lamb * mu0)/(nj[j] + lamb)
sig[j] = sig[j]/(nj[j] + lamb)
mu.prop[j] = rnorm(1, mu[j], sqrt(sig[j]))
if(j >= 2){
if(mu.prop[j] < mu.prop[j - 1])
ind.ord = 1 } }
if(ind.ord == 0){
mu = mu.prop
sig = sig.prop } }

```