

# Multinomial Example

```
// Multinomial Example RCPP CODE
NumericVector multi_e(NumericVector x, NumericVector p){
  double n1;
  double n2;
  double n3;
  n1 = 38;
  n2 = 34;
  n3 = 125*(p[2]/(p[3]+p[2]));
  NumericVector n = {n1,n2,n3};
  return n;
}

NumericVector multi_m(NumericVector n){
  double theta = (34+n[2])/(72+n[2]);
  NumericVector p = {1.0/2.0-theta/2, theta/4, theta/4, 1.0/2.0};
  return p;
}

// [[Rcpp::export]]
NumericVector multi_out(NumericVector x, NumericVector p, int itr){

  for(int i = 0; i<itr; i++){
    NumericVector n = multi_e(x, p);
    p = multi_m(n);
  }

  return(p);
}
```

```
#Multinomial R Code
x = c(38,34,125)
n = rep(0,4)
theta = 1/2
p = c(1/2-theta/2, theta/4, theta/4, 1/2)
itr = 50

multi_out(x,p, itr)
```

결과값은

0.1865893 0.1567054 0.1567054 0.5000000

# PEPPERED MOTHS EXAMPLE

```
// RCPP
NumericVector pep_e1(NumericVector x, NumericVector p){
```

```

double n_cc = (x[0]*pow(p[0],2))/(pow(p[0],2)+2*p[0]*p[1]+2*p[0]*p[2]);
double n_ci = (2*x[0]*p[0]*p[1])/(pow(p[0],2)+2*p[0]*p[1]+2*p[0]*p[2]);
double n_ct = (2*x[0]*p[0]*p[2])/(pow(p[0],2)+2*p[0]*p[1]+2*p[0]*p[2]);
double n_ii = (x[1]*pow(p[1],2))/(pow(p[1],2)+2*p[1]*p[2]);
double n_it = (2*x[1]*p[1]*p[2])/(pow(p[1],2)+2*p[1]*p[2]);
NumericVector n = {n_cc,n_ci,n_ct,n_ii,n_it,x[2]};
return n;
}

NumericVector pep_m1(NumericVector x, NumericVector n){
double p_c = (2*n[0]+n[1]+n[2])/(2*sum(x));
double p_i = (2*n[3]+n[4]+n[1])/(2*sum(x));
double p_t = (2*n[5]+n[2]+n[4])/(2*sum(x));
NumericVector p = {p_c,p_i,p_t};
return p;
}

// [[Rcpp::export]]
NumericVector pep_out1(NumericVector x, NumericVector p, int itr){

for(int i = 0; i<itr; i++){
    NumericVector n = pep_e1(x, p);
    p = pep_m1(x, n);
}

return(p);
}

```

```

# R CODE
x = c(85, 196, 341) #observed data (carbonaria, insularia, typica)
n = rep(0,6) #expected number of each phenotype (CC, CI, CT, II, IT, TT)
p = rep(1/3,3) #probabilities of allele(carbonaria, insularia, typica)
itr = 50

pep_out1(x, p, itr)

```

결과값

0.07083691 0.18873652 0.74042657

## Problem 4.1

(a)

# Exercise 4.1

Complete Data:  $\sim$  Multinomial dist  $(p_{cc}, p_{cI}, p_{cT}, p_{II}, p_{IT}, p_{TT})$

$$y = (n_{cc}, n_{cI}, n_{cT}, n_{II}, n_{IT}, n_{TT})$$

Observed data

$$X = \begin{pmatrix} n_{cc} + n_{cI} + n_{cT} & n_{II} + n_{IT} & n_{TT} \\ \vdots & \vdots & \vdots \\ p_c & p_I & p_T \\ & p_{II} & p_{IT} \end{pmatrix} = (n_c, n_I, n_T, n_u)$$

$$(p_c^*, 2p_c p_I, 2p_c p_T, p_I^*, 2p_I p_T, p_T^*)$$

Complete-data log-likelihood

$$\mathcal{L}(p) = n_{cc} \log p_c^* + n_{cI} \log 2p_c p_I + n_{cT} \log 2p_c p_T + n_{II} \log p_I^* + n_{IT} \log 2p_I p_T + n_{TT} \log p_T^* + \text{Constant}$$

$$E\{n_{cc} | n_c, n_I, n_T, p^{(t)}\} = n_{cc}^{(t)} = n_c \times \frac{p_c^{(t)2}}{p_c^{(t)2} + 2p_c^{(t)} p_I^{(t)} + 2p_c^{(t)} p_T^{(t)}} \quad \downarrow \text{C와 관련된 것들}$$

$$E\{n_{cI} | n_c, n_I, n_T, p^{(t)}\} = n_{cI}^{(t)} = n_c \times \frac{2p_c^{(t)} p_I^{(t)}}{p_c^{(t)2} + 2p_c^{(t)} p_I^{(t)} + 2p_c^{(t)} p_T^{(t)}}$$

$$E\{n_{cT} | \quad \quad \quad \} = n_{cT}^{(t)} = n_c \times \frac{2p_c^{(t)} p_T^{(t)}}{p_c^{(t)2} + 2p_c^{(t)} p_I^{(t)} + 2p_c^{(t)} p_T^{(t)}}$$

$$E\{n_{II} | \quad \quad \quad \} = n_{II}^{(t)} = n_I \times \frac{p_I^{(t)2}}{p_I^{(t)2} + 2p_I^{(t)} p_T^{(t)}} + n_u \times \frac{p_I^{(t)2}}{p_I^{(t)2} + 2p_I^{(t)} p_T^{(t)} + p_T^{(t)2}}$$

$$E\{n_{IT} | \quad \quad \quad \} = n_{IT}^{(t)} = n_I \times \frac{2p_I^{(t)} p_T^{(t)}}{p_I^{(t)2} + 2p_I^{(t)} p_T^{(t)}} + n_u \times \frac{2p_I^{(t)} p_T^{(t)}}{p_I^{(t)2} + 2p_I^{(t)} p_T^{(t)} + p_T^{(t)2}}$$

$$E\{n_{TT} | \quad \quad \quad \} = n_{TT}^{(t)} = n_T + n_u \times \frac{p_T^{(t)2}}{p_I^{(t)2} + 2p_I^{(t)} p_T^{(t)} + p_T^{(t)2}}$$

$$Q(p|p^{(t)}) = n_{cc}^{(t)} \log p_c^2 + n_{ci}^{(t)} \log 2p_c p_i + n_{ct}^{(t)} \log 2p_c p_t + n_{ii}^{(t)} \log p_i^2 + n_{it}^{(t)} \log 2p_i p_t + n_{tt} \log p_t^2 + C.$$

M-Step.

$$\frac{dQ(p|p^{(t)})}{dp_c} = \frac{2n_{cc}^{(t)} + n_{ci}^{(t)} + n_{ct}^{(t)}}{p_c} + \frac{2n_{it} + n_{ct}^{(t)} + n_{it}^{(t)}}{1-p_c-p_i} \stackrel{\text{set}}{=} 0$$

$$\frac{dQ(p|p^{(t)})}{dp_i} = \frac{2n_{ii}^{(t)} + n_{ci}^{(t)} + n_{it}^{(t)}}{p_i} + \frac{2n_{it} + n_{ct}^{(t)} + n_{it}^{(t)}}{1-p_c-p_i} = 0$$

$$p_c^{(t+1)} = \frac{2n_{cc}^{(t)} + n_{ci}^{(t)} + n_{ct}^{(t)}}{2n} \quad p_t = \frac{2n_{tt}^{(t)} + n_{ct}^{(t)} + n_{it}^{(t)}}{2n}$$

$$p_i^{(t+1)} = \frac{2n_{ii}^{(t)} + n_{ci}^{(t)} + n_{it}^{(t)}}{2n}$$

(b)

```
// Exercise 4.1
// RCPP
// E-step
NumericVector pep_e2(NumericVector x, NumericVector p){
    double n_cc = (x[0]*pow(p[0],2))/(pow(p[0],2)+2*p[0]*p[1]+2*p[0]*p[2]);
    double n_ci = (2*x[0]*p[0]*p[1])/(pow(p[0],2)+2*p[0]*p[1]+2*p[0]*p[2]);
    double n_ct = (2*x[0]*p[0]*p[2])/(pow(p[0],2)+2*p[0]*p[1]+2*p[0]*p[2]);
    double n_ii = (x[1]*pow(p[1],2))/(pow(p[1],2)+2*p[1]*p[2]) +
    ((x[3]*pow(p[1],2))/(pow(p[1],2)+2*p[1]*p[2]+pow(p[2],2)));
    double n_it = (2*x[1]*p[1]*p[2])/(pow(p[1],2)+2*p[1]*p[2]) +
    (2*x[3]*p[1]*p[2])/(pow(p[1],2)+2*p[1]*p[2]+pow(p[2],2));
```

```

    double n_tt = x[2] +
((x[3])*pow(p[2],2))/(pow(p[1],2)+2*p[1]*p[2]+pow(p[2],2));
    NumericVector n = {n_cc,n_ci,n_ct,n_ii,n_it,n_tt};
    return n;
}

// M-step
NumericVector pep_m2(NumericVector x, NumericVector n){
    double p_c = (2*n[0]+n[1]+n[2])/(2*sum(x));
    double p_i = (2*n[3]+n[4]+n[1])/(2*sum(x));
    double p_t = (2*n[5]+n[2]+n[4])/(2*sum(x));
    NumericVector p = {p_c,p_i,p_t};
    return p;
}

// [[Rcpp::export]]
NumericVector pep_out2(NumericVector x, NumericVector p, int itr){

    for(int i = 0; i<itr; i++){
        NumericVector n = pep_e2(x, p);
        p = pep_m2(x, n);
    }

    return(p);
}

```

```

#R
#observed data (carbonaria, insularia, typica, in+ty)
x = c(85, 196, 341, 578)
#expected number of each phenotype (CC, CI, CT, II, IT, TT)
n = rep(0,6)
p = rep(1/3,3) #probabilities of allele(carbonaria, insularia, typica)
itr = 30

pep_out2(x, p, itr)

```

결과값

0.03606708 0.19579918 0.76813373

p\_c = 0.03606708

p\_i = 0.19579918

p\_t = 0.76813373

## Problem 4.2

(a)

## Exercise 4.2

$$N = 1500$$

$$\sum_{i=1}^{16} n_i = N$$

$$\text{Complete Data } Y = C(n_{z0}, n_{t0}, n_{t1}, n_{t2}, \dots, n_{t16}, n_{p0}, n_{p1}, \dots, n_{p16})$$

$$\text{Observed Data } X = M(Y) = C(n_0, n_1, n_2, \dots, n_{16})$$

$$\pi_i(\theta) = \alpha \cdot 1_{[i=0]} + \beta \mu \exp\{-\mu\} + (1-\alpha-\beta) \lambda \exp\{-\lambda\}$$

$$z_0(\theta) = \frac{\alpha}{\pi_0(\theta)}$$

$$t_i(\theta) = \frac{\beta \mu \exp\{-\mu\}}{\pi_i(\theta)}$$

$$p_i(\theta) = \frac{(1-\alpha-\beta) \lambda \exp\{-\lambda\}}{\pi_i(\theta)}$$

$$P = (z_0(\theta), t_0(\theta), t_1(\theta), \dots, t_{16}(\theta), p_0(\theta), p_1(\theta), \dots, p_{16}(\theta))$$

$$\text{Let } 1-\alpha-\beta = r$$

$$\text{Complete log likelihood} = n_{z,0} \log(\alpha) + \sum_{i=0}^{16} n_{t,i} \log \beta \mu \exp\{-\mu\} + \sum_{i=0}^{16} n_{p,i} \log r \lambda \exp\{-\lambda\}$$

$$= n_{z,0} \log(\alpha) + \sum_{i=0}^{16} n_{t,i} \{ \log(\beta) + i \log(\mu) - \mu \} + \sum_{i=0}^{16} n_{p,i} \{ \log(r) + i \log(\lambda) - \lambda \}$$

$$E[n_{z,0} | n_0, n_1, \dots, n_{16}, \theta^{(k)}] = n_0 \times \frac{\alpha^{(k)}}{\pi_0(\theta^{(k)})} = n_0 \times z_0(\theta^{(k)})$$

$$E[n_{t,i} | n_0, n_1, \dots, n_{16}, \theta^{(k)}] = n_i \times \frac{\beta^{(k)} (\mu^{(k)})^i \exp(-\mu^{(k)})}{\pi_i(\theta^{(k)})} = n_i \times t_i(\theta^{(k)})$$

$$E[n_{p,i} | n_0, n_1, \dots, n_{16}, \theta^{(k)}] = n_i \times \frac{r^{(k)} (\lambda^{(k)})^i \exp(-\lambda^{(k)})}{\pi_i(\theta^{(k)})} = n_i \times p_i(\theta^{(k)})$$

$$\therefore Q(\theta | \theta^{(k)}) = n_0 z_0(\theta^{(k)}) \log(\alpha) + \sum_{i=0}^{16} n_i t_i(\theta^{(k)}) \{ \log(\beta) + i \log(\mu) - \mu \} + \sum_{i=0}^{16} n_i p_i(\theta^{(k)}) \{ \log(r) + i \log(\lambda) - \lambda \}$$

Lagrange term을 이용해서 보자. ( $\alpha + \beta + r = 1$ 이라는 조건 존재).

$$Q_{\text{logr}}(\theta | \theta^{(k)}) = Q(\theta | \theta^{(k)}) + \phi(1-\alpha-\beta-r)$$

$$\frac{\partial Q_{\text{logr}}(\theta | \theta^{(k)})}{\partial \alpha} = n_0 z_0(\theta^{(k)}) \times \frac{1}{\alpha} - \phi$$

$$\therefore \alpha^{(t+1)} = \frac{1}{\phi} n_0 z_0(\theta^{(t)})$$

$$\frac{\partial Q_{\log r}(\theta|\theta^{(t)})}{\partial \beta} = \sum_{i=0}^{16} n_i t_i(\theta^{(t)}) \frac{1}{\beta} - \phi$$

$$\therefore \beta^{(t+1)} = \frac{1}{\phi} \sum_{i=0}^{16} n_i t_i(\theta^{(t)})$$

$$\frac{\partial Q_{\log r}(\theta|\theta^{(t)})}{\partial r} = \sum_{i=0}^{16} n_i p_i(\theta^{(t)}) \frac{1}{r} - \phi$$

$$\therefore r^{(t+1)} = \frac{1}{\phi} \sum_{i=0}^{16} n_i p_i(\theta^{(t)})$$

$$\alpha^{(t+1)} + \beta^{(t+1)} + r^{(t+1)} = 1 = \frac{1}{\phi} \left\{ n_0 z_0(\theta^{(t)}) + \sum_{i=0}^{16} n_i t_i(\theta^{(t)}) + \sum_{i=0}^{16} n_i p_i(\theta^{(t)}) \right\} = \frac{N}{\phi}$$

$$\phi = N$$

$$\therefore \alpha^{(t+1)} = \frac{1}{N} n_0 z_0(\theta^{(t)})$$

$$\beta^{(t+1)} = \frac{1}{N} \sum_{i=0}^{16} n_i t_i(\theta^{(t)}) = \frac{1}{N} \frac{\sum_{i=0}^{16} n_i t_i(\theta^{(t)})}{N}$$

$$\frac{\partial Q_{\log r}(\theta|\theta^{(t)})}{\partial \mu} = \sum_{i=0}^{16} n_i t_i(\theta^{(t)}) \left( \frac{i}{\mu} - 1 \right)$$

$$\therefore \mu^{(t+1)} = \frac{\sum_{i=0}^{16} i n_i t_i(\theta^{(t)})}{\sum_{i=0}^{16} n_i t_i(\theta^{(t)})}$$

$$\frac{\partial Q_{\log r}(\theta|\theta^{(t)})}{\partial \lambda} = \sum_{i=0}^{16} n_i p_i(\theta^{(t)}) \left( \frac{i}{\lambda} - 1 \right)$$

$$\therefore \lambda^{(t+1)} = \frac{\sum_{i=0}^{16} i n_i p_i(\theta^{(t)})}{\sum_{i=0}^{16} n_i p_i(\theta^{(t)})}$$

(b)

```
// RCPP
// [[Rcpp::export]]
vec hiv_out(vec param, vec obs, int itr){
  vec new_param = zeros(4);
  double len = obs.size();
  double N = sum(obs);
  double z_0, t_sum, p_sum, z_sum;
  vec t(len);
  vec p(len);
  vec pi_f(len);
```

```

for(int i = 0; i<itr; i++){
    //E-step
    //statistics update
    //0 group setting
    pi_f[0] = param[0] + param[1]*exp(-param[2]) + (1-param[0]-param[1])*exp(-
param[3]);
    z_0 = param[0] / pi_f[0];
    t[0] = (param[1]*exp(-param[2])) / pi_f[0];
    p[0] = ( (1-param[0]-param[1])*exp(-param[3]) ) / pi_f[0];

    //1~17 group update
    for(int j = 1; j < len; j++){
        pi_f[j] = param[1]*pow(param[2], j)*exp(-param[2]) + (1-param[0]-
param[1])*pow(param[3], j)*exp(-param[3]);
        t[j] = (param[1]*pow(param[2], j)*exp(-param[2])) / pi_f[j];
        p[j] = ((1-param[0]-param[1])*pow(param[3], j)*exp(-param[3])) / pi_f[j];
    }

    t_sum = 0;
    p_sum = 0;
    z_sum = 0;

    z_sum = obs[0] * z_0;

    for(int k=0; k < len; k++){
        t_sum = t_sum + obs[k]*t[k];
        p_sum = p_sum + obs[k]*p[k];
    }

    // M-step
    new_param = {0, 0, 0, 0};
    new_param[0] = z_sum / N ;
    for(int l=0; l<len; l++){
        new_param[1] = new_param[1] + (obs[l]*t[l]) / N ;
        new_param[2] = new_param[2] + (l*obs[l]*t[l]) / t_sum ;
        new_param[3] = new_param[3] + (l*obs[l]*p[l]) / p_sum ;
    }

    param = new_param;
}

return(param);
}

```



```
#R
#data
obs =
c(379.0,299.0,222.0,145.0,109.0,95.0,73.0,59.0,45.0,30.0,24.0,12.0,4.0,2.0,0.0,1
.0,1.0)
itr = 100
alpha = 0.5
beta = 0.2
mu = 1.0
lambda = 5.0
param = c(alpha, beta, mu, lambda)

#RCPP
hiv_out(param, obs, itr)
```

결과값

alpha: 0.1221626  
beta: 0.5625419  
mu: 1.4674525  
lambda: 5.9388614