### ABSTAT17

IGC, April 10-13, 2017

#### **EXERCISE:** EM Algoritm

The main locus for the blood type of mice is called Ag-B (B). Several alleles are associated to this locus but for some crossovers Mendel's laws do not seem to hold. A mating  $AaBb \times AaBb \equiv F_1 \times F_1$ , originated a  $F_2$  progeny, yielding

Genotype	Frequency	Probability
$\overline{AABB}$	11	$(1-\theta)^2/4$
AABb	14	$\theta(1-\theta)/2$
AAbb	1	$\theta^2/4$
AaBB	10	$\theta(1-\theta)/2$
AaBb	27	$(\theta^2/2) + [(1-\theta)^2]/2$
Aabb	12	$\theta(1-\theta)/2$
aaBB	3	$\theta^2/4$
aaBb	13	$\theta(1-\theta)/2$
aabb	11	$(1-\theta)^2/4$

Estimate the recombination fraction,  $\theta$ , from these data by the EM algorithm.

## Step 1

Read the data and state ao many recombinant gametes are there for each genotype.

```
nAABB<-11 # 0 recombinant gametes
nAABb<-14 # 1 recombinant gamete
nAAbb<-1 # 2 recombinant gametes
nAaBB<-10 # 1 recombinant gamete
nAaBb<-27 # 0 or 2 recombinant gametes
nAabb<-12 # 1 recombinant gamete
naaBB<-3 # 2 recombinant gametes
naaBb<-13 # 1 recombinant gamete
naabb<-11 # 0 recombinant gametes
```

Calculate  $n_1$ , the number of individuals from 1 recombinant gametes (n1).

Calculate  $n_2$ , the number of individuals from 2 recombinant gametes (n2). Note that  $n_{AaBb} = n_2^* + n_0^*$  (nAaBb = n2.star + n0.star).

Calculate n, the total number of individuals (n).

#### Step 2

Initialize  $\theta \in [0, 0.5]$  (r).

## Step 3 - E (Expectation)

Create function expected in order to calculate the expected value for  $N_2^*$ :

 $N_2^*$ : random variable representing the number of individuals from 2 recombinant gametes, among  $n_{AaBb}$  individuals.

$$N_2^* \frown Binomial(n_{AaBb}, p)$$
 with  $p = \frac{\theta^2}{\theta^2 + (1 - \theta)^2}$ 

then, 
$$n_2^* = E(N_2^*) = n_{AaBb} \times p$$

# Step 4 - M (Maximization)

Create function update.theta in order to update  $\theta$  according to:

$$\theta = \frac{n_1 + 2(n_{AAbb} + n_{aaBB} + n_2^*)}{2n}$$

meaning that the proportion of recombinant gametes is calculated as the total number of recombinant gametes (0, 1 or 2 for each individual) over the total number of gametes for n individuals.

# Step 5

EM algorithm: Iterative procedure.

# Step 6

Print the results.