Livestock Breeding and Genomics - Solution 8

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Problem 1: Compute Inbreeding Coefficients

Given the following pedigree.

Animal	Sire	Dam
1	NA	NA
2	NA	NA
3	1	NA
4	3	2
5	4	2
6	4	5

Your Task

Compute the inbreeding coefficients F_i for all animals using the matrix R that comes from the cholesky decomposition of the numerator relationship matrix A

Solution

The cholesky decomposition of A corresponds to

$$A = R \cdot R^T$$

where R is a lower triangular matrix. The diagonal elements $(A)_{ii}$ of A can be computed as the sum of the squared elements of all elements of Matrix R on row i.

$$(A)_{ii} = \sum_{j=1}^{i} (R)_{ij}^{2}$$

Therefore to get $(A)_{ii}$, we have to compute all elements of R on row i. Due to the relation of the cholesky-decomposition to the LDL-decomposition, we can write the matrix R as a product of the two matrices L and S

$$R = L \cdot S \tag{1}$$

where L is the lower triangular matrix from the LDL-decomposition and S is a diagonal matrix with diagonal elements $(S)_{ii}$ corresponding to

$$(S)_{ii} = \sqrt{(D)_{ii}}$$

where $(D)_{ii}$ correspond to the diagonal elements of the matrix D from the LDL-decomposition. From the LDL, we know that

$$(D)_{ii} = \frac{1}{2} - \frac{1}{4}(F_s + F_d) = 1 - \frac{1}{4}(A_{ss} + A_{dd})$$

Based on the decomposition of R into the product of L and S given in (1) and based on the property of the matrix L which is to be shown in the additional problem 3 of this exercise, we can derive the following rules for computing the elements of the matrix R

• Diagnoal elements $(R)_{ii}$:

$$(R)_{ii} = (S)_{ii} = \sqrt{(D)_{ii}} = \sqrt{1 - \frac{1}{4}(A_{ss} + A_{dd})}$$

where s and d are parents of animal i and $(A)_{ss}$ and $(A)_{dd}$ are diagonal elements of the numerator relationship matrix A.

• Off-diagonal elements $(R)_{ij}$ $(i \neq j)$:

$$(R)_{ij} = \frac{1}{2}(R_{sj} + R_{dj})$$

where s and d are parents of animal i.

The solution of this exercise is to compute $(A)_{ii}$ for all animals in the pedigree using the above described rules.

• $(A)_{11}$

$$(A)_{11} = (R)_{11}^2 = 1$$

• $(A)_{22}$

$$(A)_{22} = (R)_{21}^2 + (R)_{22}^2 = 0 + 1 = 1$$

• $(A)_{33}$

$$(A)_{33} = (R)_{31}^2 + (R)_{32}^2 + (R)_{33}^2 = 0.25 + 0 + 0.75 = 1$$

• (A)₄₄

$$(A)_{44} = (R)_{41}^2 + (R)_{42}^2 + (R)_{43}^2 + (R)_{44}^2$$

= 0.0625 + 0.25 + 0.1875 + 0.5 = 1

• $(A)_{55}$

$$(A)_{55} = (R)_{51}^2 + (R)_{52}^2 + (R)_{53}^2 + (R)_{54}^2 + (R)_{55}^2$$

= 0.015625 + 0.5625 + 0.046875 + 0.125 + 0.5 = 1.25

• $(A)_{66}$

$$(A)_{66} = (R)_{61}^2 + (R)_{62}^2 + (R)_{63}^2 + (R)_{64}^2 + (R)_{65}^2 + (R)_{66}^2$$

= 0.03515625 + 0.390625 + 0.1054688 + 0.28125 + 0.125 + 0.4375 = 1.375

As a check, we can compute the inbreeding coefficients using the function pedigreemm::inbreeding()

```
pedigreemm::inbreeding(ped = ped_sol10p01)
```

[1] 0.000 0.000 0.000 0.000 0.250 0.375

Problem 2: Direct Construction of A^{-1}

Use the pedigree from problem 1 and the computed inbreeding coefficients from problem 1 to set up the inverse numerator relationship matrix A^{-1} using the general form of Henderson's rules for a pedigree with inbred animals. Compare your result using function pedigreemm::getAInv().

Solution

As a pre-requisite, we assume that the pedigree is sorted such that parents come before progeny. Henderson's rules contain the following steps

- Start with a matrix A^{-1} where all elements are set to 0.
- Let d^i be the *i*-th diagonal element of D^{-1} for animal *i*, assuming *i* has parents *s* and *d*.
- Then add the following contributions to A^{-1}
 - $-d^{i}$ to the element (i,i)
 - $-d^{i}/2$ to the elements (s,i), (i,s), (d,i), (i,d)
 - $-d^{i}/4$ to the elements (s,s), (s,d), (d,s), (d,d)

Applying these rules to the pedigree given in problem 1 leads to the following sequence of computations.

• Initialize the matrix A^{-1} with all 0

• Animal 1

• Animal 2

• Animal 3

$$A^{-1} = \begin{bmatrix} 1.3333 & 0.0000 & -0.6667 & 0.0000 & 0.0000 & 0.0000 \\ 0.0000 & 1.0000 & 0.0000 & 0.0000 & 0.0000 & 0.0000 \\ -0.6667 & 0.0000 & 1.3333 & 0.0000 & 0.0000 & 0.0000 \\ 0.0000 & 0.0000 & 0.0000 & 0.0000 & 0.0000 & 0.0000 \\ 0.0000 & 0.0000 & 0.0000 & 0.0000 & 0.0000 & 0.0000 \\ 0.0000 & 0.0000 & 0.0000 & 0.0000 & 0.0000 & 0.0000 \\ 0.0000 & 0.0000 & 0.0000 & 0.0000 & 0.0000 & 0.0000 \end{bmatrix}$$

• Animal 4

$$A^{-1} = \begin{bmatrix} 1.3333 & 0.0000 & -0.6667 & 0.0000 & 0.0000 & 0.0000 \\ 0.0000 & 1.5000 & 0.5000 & -1.0000 & 0.0000 & 0.0000 \\ -0.6667 & 0.5000 & 1.8333 & -1.0000 & 0.0000 & 0.0000 \\ 0.0000 & -1.0000 & -1.0000 & 2.0000 & 0.0000 & 0.0000 \\ 0.0000 & 0.0000 & 0.0000 & 0.0000 & 0.0000 & 0.0000 \\ 0.0000 & 0.0000 & 0.0000 & 0.0000 & 0.0000 & 0.0000 \end{bmatrix}$$

• Animal 5

$$A^{-1} = \begin{bmatrix} 1.3333 & 0.0000 & -0.6667 & 0.0000 & 0.0000 & 0.0000 \\ 0.0000 & 2.0000 & 0.5000 & -0.5000 & -1.0000 & 0.0000 \\ -0.6667 & 0.5000 & 1.8333 & -1.0000 & 0.0000 & 0.0000 \\ 0.0000 & -0.5000 & -1.0000 & 2.5000 & -1.0000 & 0.0000 \\ 0.0000 & -1.0000 & 0.0000 & -1.0000 & 2.0000 & 0.0000 \\ 0.0000 & 0.0000 & 0.0000 & 0.0000 & 0.0000 & 0.0000 \end{bmatrix}$$

• Animal 6

$$A^{-1} = \begin{bmatrix} 1.3333 & 0.0000 & -0.6667 & 0.0000 & 0.0000 & 0.0000 \\ 0.0000 & 2.0000 & 0.5000 & -0.5000 & -1.0000 & 0.0000 \\ -0.6667 & 0.5000 & 1.8333 & -1.0000 & 0.0000 & 0.0000 \\ 0.0000 & -0.5000 & -1.0000 & 3.0714 & -0.4286 & -1.1429 \\ 0.0000 & -1.0000 & 0.0000 & -0.4286 & 2.5714 & -1.1429 \\ 0.0000 & 0.0000 & 0.0000 & -1.1429 & -1.1429 & 2.2857 \end{bmatrix}$$

• Check with function pedigreemm::getAinv()

pedigreemm::getAInv(ped = ped_sol10p01)

```
## 6 x 6 Matrix of class "dgeMatrix"
##
                  2
                                                   5
                                                             6
                0.0 -0.6666667 0.0000000 0.0000000
     1.3333333
                                                      0.000000
     0.0000000
                2.0 0.5000000 -0.5000000 -1.0000000
                                                     0.000000
## 3 -0.6666667
                0.5 1.8333333 -1.0000000 0.0000000
                                                     0.000000
     0.0000000 -0.5 -1.0000000 3.0714286 -0.4285714 -1.142857
     0.0000000 -1.0 0.0000000 -0.4285714 2.5714286 -1.142857
     0.0000000 0.0 0.0000000 -1.1428571 -1.1428571 2.285714
```

The difference between the computed matrix and the matrix from pedigreemm::getAinv()

$$\begin{bmatrix} 0.00 & 0.00 & 0.00 & 0.00 & 0.00 & 0.00 \\ 0.00 & 0.00 & 0.00 & 0.00 & 0.00 & 0.00 \\ 0.00 & 0.00 & 0.00 & 0.00 & 0.00 & 0.00 \\ 0.00 & 0.00 & 0.00 & 0.00 & 0.00 & 0.00 \\ 0.00 & 0.00 & 0.00 & 0.00 & 0.00 & 0.00 \\ 0.00 & 0.00 & 0.00 & 0.00 & 0.00 & 0.00 \\ 0.00 & 0.00 & 0.00 & 0.00 & 0.00 & 0.00 \\ \end{bmatrix}$$