Inverse Numerator Relationship Matrix

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Problem 1: Inverse Numerator Relationship Matrix

During the lecture the method of computing the inverse numerator relationship matrix A^{-1} directly was introduced. The computation is based on the LDL-decomposition. As a result, we can write

$$A^{-1} = (L^T)^{-1} \cdot D^{-1} \cdot L^{-1}$$

where $L^{-1} = I - P$, and D^{-1} is a diagonal matrix with $(D^{-1})_{ii} * \sigma_u^{-2} = var(m_i)^{-1}$.

Tasks

- Use the example pedigree given below and compute the matrices L^{-1} and D^{-1} to compute A^{-1}
- Verify your result using the function getAinv() from package pedigreemm.

Pedigree

```
nr_animal <- 6
tbl_pedigree <- tibble::tibble(Calf = c(1:nr_animal),</pre>
                                Sire = c(NA, NA, NA, 1, 3, 4),
                                Dam = c(NA, NA, NA, 2, 2, 5))
tbl_pedigree
## # A tibble: 6 x 3
      Calf Sire
     <int> <dbl> <dbl>
##
## 1
        1
              NA
                    NA
        2
## 2
              NA
                    NA
## 3
         3
              NA
                    NA
## 4
         4
              1
## 5
         5
               3
                     2
## 6
                     5
```

Solution

The matrix P comes from the simple decomposition and can be constructed using the pedigree.

```
P = matrix(0, nrow = nr_animal, ncol = nr_animal)
for (i in 1:nr_animal){
    s <- tbl_pedigree$Sire[i]
    d <- tbl_pedigree$Dam[i]
    if (!is.na(s)){
        P[i,s] <- 0.5
    }</pre>
```

```
if(!is.na(d)){
    P[i,d] \leftarrow 0.5
  }
}
Ρ
        [,1] [,2] [,3] [,4] [,5] [,6]
##
## [1,] 0.0 0.0 0.0 0.0 0.0
## [2,]
        0.0 0.0 0.0 0.0
                             0.0
                                    0
## [3,]
        0.0 0.0 0.0 0.0
                             0.0
## [4,]
        0.5 0.5 0.0 0.0
                             0.0
                                    0
        0.0 0.5 0.5 0.0
                                    0
## [5,]
                             0.0
## [6,]
        0.0 0.0 0.0 0.5 0.5
                                    0
With that the matrix L^{-1} is
I <- diag(1, nrow = nr_animal, ncol = nr_animal)</pre>
Linv <- I - P
Linv
        [,1] [,2] [,3] [,4] [,5] [,6]
## [1,] 1.0 0.0 0.0 0.0 0.0
## [2,] 0.0 1.0 0.0 0.0
                             0.0
                                    0
## [3,] 0.0 0.0 1.0 0.0
                             0.0
                                    0
## [4,] -0.5 -0.5 0.0 1.0 0.0
                                    0
## [5,] 0.0 -0.5 -0.5 0.0 1.0
                                    0
## [6,] 0.0 0.0 0.0 -0.5 -0.5
                                    1
The matrix D is obtained from package pedigreemm
ped <- pedigreemm::pedigree(sire = tbl_pedigree$Sire,</pre>
                            dam = tbl_pedigree$Dam,
                            label = as.character(1:nr_animal))
D <- pedigreemm::Dmat(ped = ped)</pre>
Dinv <- diag(1/D, nrow = nr_animal, ncol = nr_animal)</pre>
Dinv
        [,1] [,2] [,3] [,4] [,5] [,6]
## [1,]
                0
                          0
                               0
           1
                     0
## [2,]
           0
                     0
                          0
                               0
                                    0
                1
                                    0
## [3,]
           0
                0
                     1
                          0
                               0
## [4,]
                0
                     0
                          2
                               0
                                    0
           0
                          0
                               2
                                    0
## [5,]
           0
                0
                     0
## [6,]
                     0
                                    2
The inverse numerator relationship matrix is
Ainv <- t(Linv) %*% Dinv %*% Linv
Ainv
##
        [,1] [,2] [,3] [,4] [,5] [,6]
## [1,] 1.5 0.5 0.0 -1.0 0.0
## [2,]
        0.5 2.0 0.5 -1.0 -1.0
                                    0
## [3,] 0.0 0.5 1.5 0.0 -1.0
                                    0
## [4,] -1.0 -1.0 0.0 2.5 0.5
                                   -1
## [5,]
        0.0 -1.0 -1.0 0.5 2.5
                                   -1
## [6,] 0.0 0.0 0.0 -1.0 -1.0
```

Verification

```
pedigreemm::getAInv(ped = ped)

## 6 x 6 Matrix of class "dgeMatrix"

## 1 2 3 4 5 6

## 1 1.5 0.5 0.0 -1.0 0.0 0

## 2 0.5 2.0 0.5 -1.0 -1.0 0
```

3 0.0 0.5 1.5 0.0 -1.0 0 ## 4 -1.0 -1.0 0.0 2.5 0.5 -1

5 0.0 -1.0 -1.0 0.5 2.5 -1 ## 6 0.0 0.0 0.0 -1.0 -1.0 2

Problem 2: Rules

The following diagram helps to illustrate the rules for constructing ${\cal A}^{-1}$

1	D^{-1}	L^{-1}
	[,1] [,2] [,3] [,4] [,5] [,6] [1,] 1 0 0 0 0 0 [2,] 0 1 0 0 0 0 [3,] 0 0 1 0 0 0 [4,] 0 0 0 2 0 0 [5,] 0 0 0 0 2 0 [6,] 0 0 0 0 2	[,1] [,2] [,3] [,4] [,5] [,6] [1,] 1.0 0.0 0.0 0.0 0.0 0 [2,] 0.0 1.0 0.0 0.0 0.0 0 [3,] 0.0 0.0 1.0 0.0 0.0 0 [4,] -0.5 -0.5 0.0 1.0 0.0 0 [5,] 0.0 -0.5 -0.5 0.0 1.0 0 [6,] 0.0 0.0 0.0 -0.5 -0.5 1
[,1] [,2] [,3] [,4] [,5] [,6] [1,] 1 0 0 -0.5 0.0 0.0 [2,] 0 1 0 -0.5 -0.5 0.0 [3,] 0 0 1 0.0 -0.5 0.0 [4,] 0 0 0 1.0 0.0 -0.5 [5,] 0 0 0 0.0 1.0 -0.5 [6,] 0 0 0 0.0 0.0 1.0	[,1] [,2] [,3] [,4] [,5] [,6] [1,] 1 0 0 -1 0 0 [2,] 0 1 0 -1 -1 0 [3,] 0 0 1 0 -1 0 [4,] 0 0 0 2 0 -1 [5,] 0 0 0 0 2 -1 [6,] 0 0 0 0 2	[,1] [,2] [,3] [,4] [,5] [,6] [1,] 1.5 0.5 0.0 -1.0 0.0 0 [2,] 0.5 2.0 0.5 -1.0 -1.0 0 [3,] 0.0 0.5 1.5 0.0 -1.0 0 [4,] -1.0 -1.0 0.0 2.5 0.5 -1 [5,] 0.0 -1.0 -1.0 0.5 2.5 -1 [6,] 0.0 0.0 0.0 -1.0 -1.0 2
$(L^T)^{-1}$	$(L^T)^{-1}\cdot D^{-1}$	A^{-1}