Survival analysis

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1 Cox datafit

Let's $\mathbf{X} \in \mathbb{R}^{n \times p}$ be a matrix of p predictors and n samples $x_i \in \mathbb{R}^p$, $y \in \mathbb{R}^n$ a vector recording the time of events occurrences, and $s \in \{0, 1\}^n$ a binary vector where 1 means *event occurred*, and finally $\beta \in \mathbb{R}^p$ the vector of coefficient to be estimated.

1.1 Breslow

Proposition 1. [Lin, 2007, Section 2] The expression of the negative log-likelihood according to Breslow estimate reads

$$l(\beta) = \sum_{i=1}^{n} -s_i \langle x_i, \beta \rangle + s_i \log(\sum_{y_j \ge y_i} e^{\langle x_j, \beta \rangle}) \quad .$$
(1)

To get a more compact expression, we introduce the matrix $\mathbf{B} \in \mathbb{R}^{n \times n}$ defined as

$$\mathbf{B}_{i,j} = \mathbb{1}_{y_j \ge y_i} = \begin{cases} 1, & \text{if } y_j \ge y_i, \\ 0, & \text{otherwise} \end{cases},$$
(2)

and we let b_i be its *i*-th row.

Proposition 2. The expression in Equation (1) is equivalent to

$$l(\beta) = -\langle s, \mathbf{X}\beta \rangle + \langle s, \log(\mathbf{B}e^{\mathbf{X}\beta}) \rangle \quad . \tag{3}$$

Proof. We can observe that the sum can be split into two parts. The first one,

$$\sum_{i=1}^{n} -s_i \langle x_i, \beta \rangle = -\langle s, \mathbf{X}\beta \rangle \ .$$

And the second part,

$$\sum_{i=1}^{n} s_i \log(\sum_{y_j \ge y_i} e^{\langle x_j, \beta \rangle}) = \sum_{i=1}^{n} s_i \log(\sum_{j=1}^{n} \mathbb{1}_{y_j \ge y_i} e^{\langle x_j, \beta \rangle})$$
$$= \sum_{i=1}^{n} s_i \log(\sum_{j=1}^{n} b_{ij} e^{\langle x_j, \beta \rangle})$$
$$= \sum_{i=1}^{n} s_i \log \langle b_i, e^{\mathbf{X}\beta} \rangle$$
$$= \langle s, \log(\mathbf{B}e^{\mathbf{X}\beta}) \rangle .$$

Combining the two expressions we get the desired expression.

The latter defines the Cox datafit. We note that it only depends on $\mathbf{X}\beta$. Indeed, by considering $F : \mathbb{R}^n \to \mathbb{R}$ such that

$$F(u) = -\langle s, u \rangle + \langle s, \log(\mathbf{B}e^u) \rangle \quad , \tag{4}$$

it follows that $l(\beta) = F(\mathbf{X}\beta)$. Therefore, from now on, we focus on F to derive the gradient and Hessian of the datafit.

Proposition 3. For some u in \mathbb{R}^n , the gradient of F reads

$$\nabla F(u) = -s + [\operatorname{diag}(e^u)\mathbf{B}^{\top}] \frac{s}{\mathbf{B}e^u}$$
,

where the fraction is performed element-wise.

Proof. Deferred in the appendix.

Proposition 4. For some u in \mathbb{R}^n , the Hessian of F is

$$\nabla^2 F(u) = \operatorname{diag}(e^u \odot \mathbf{B}^\top \frac{s}{\mathbf{B}e^u}) - \operatorname{diag}(e^u) \mathbf{B}^\top \operatorname{diag}(\frac{s}{(\mathbf{B}e^u)^2}) \mathbf{B} \operatorname{diag}(e^u) \quad , \tag{5}$$

where the square and fraction operations are performed element-wise.

Proof. Deferred in the appendix.

The Hessian, as it is, is costly to evaluate because of the right hand-side term. Indeed, the latter involves, in particular, a $\mathcal{O}(n^3)$ operation. We overcome this limitation thanks to the proposition below.

Proposition 5. For some u in \mathbb{R}^n , the Hessian in Equation (5) can be overestimated as follows

$$\nabla^2 F(u) \preccurlyeq \operatorname{diag}(e^u \odot \mathbf{B}^\top \frac{s}{\mathbf{B}e^u})$$

Proof. We have to show that diag $(e^u \odot \mathbf{B}^\top \frac{s}{\mathbf{B}e^u}) - \nabla^2 F(u) := \Phi$ is positive semi-definite. Let u be in \mathbb{R}^n , we have

$$\begin{split} \langle \mathbf{\Phi} u, u \rangle &= \left\langle \operatorname{diag}(e^u) \mathbf{B}^\top \operatorname{diag}(\frac{s}{(\mathbf{B} e^u)^2}) \mathbf{B} \operatorname{diag}(e^u) u, u \right\rangle \\ &= \left\| \operatorname{diag}(\frac{\sqrt{s}}{\mathbf{B} e^u}) \mathbf{B} \operatorname{diag}(e^u) u \right\|^2 \geq 0 \end{split}$$

which enables us to conclude.

1.2 Efron

Efron estimate refines Breslow by handling tied observations, observations with identical occurrences' time. Let's define $H_k = \{i \mid s_i = 1 ; y_i = y_k\}$, the set of uncensored observations with the same occurrence time y_k , and denote y_{i_1}, \ldots, y_{i_m} the unique times, assumed to be in total equal to m.

Proposition 6. [Efron, 1977, Section 6, equation (6.7)] The minus log-likelihood according to Efron estimate is

$$l(\beta) = \sum_{l=1}^{m} \left(\sum_{i \in H_{i_l}} -\langle x_i, \beta \rangle + \log \left(\sum_{y_j \ge y_{i_l}} e^{\langle x_j, \beta \rangle} - \frac{\#(i) - 1}{|H_{i_l}|} \sum_{j \in H_{i_l}} e^{\langle x_j, \beta \rangle} \right) \right) , \qquad (6)$$

where $|H_{i_l}|$ stands for the cardinal of H_{i_l} , and #(i) the index of observation i in H_{i_l} .

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Ideally, we would like to rewrite this expression like Equation (3) to leverage the established results about the gradient and Hessian. What distinguishes both expressions is the presence of a double sum and second term within the log.

Proposition 7. The expression in Equation (6) is equivalent to

$$l(\beta) = -\langle s, \mathbf{X}\beta \rangle + \langle s, \log(\mathbf{B}e^{\mathbf{X}\beta} - \mathbf{A}e^{\mathbf{X}\beta}) \rangle \quad , \tag{7}$$

where \mathbf{A} is linear operator defined by Algorithm 1.

Proof. First, we can observe that $\bigcup_{l=1}^{m} H_{i_l} = \{i \mid s_i = 1\}$, which enables us to write the double sum as a single one. Therefore,

$$\sum_{l=1}^{m} \sum_{i \in H_{i_l}} -\langle x_i, \beta \rangle = \sum_{i:s_i=1}^{n} -\langle x_i, \beta \rangle = \sum_{i=1}^{n} -s_i \langle x_i, \beta \rangle$$
$$= -\langle s, \mathbf{X}\beta \rangle \quad .$$

On the other hand, we have

$$\begin{aligned} -\frac{\#(i)-1}{|H_{i_l}|} \sum_{j \in H_{i_l}} e^{\langle x_j, \beta \rangle} &= \sum_{j=1}^n -\frac{\#(i)-1}{|H_{i_l}|} \ \mathbb{1}_{j \in H_{i_l}} \ e^{\langle x_j, \beta \rangle} \\ &= \sum_{j=1}^n a_{i,j} e^{\langle x_j, \beta \rangle} \\ &= \langle a_i, e^{\mathbf{X}\beta} \rangle \ , \end{aligned}$$

where a_i is a vector in \mathbb{R}^n chosen accordingly to preform the linear operation.

Defining the matrix **A** with rows $(a_i)_{i \in [n]}$, and combining that with the first result enable us to conclude.

Since the expression of the gradient and Hessian involve the adjoint of \mathbf{A} , we present also Algorithm 2 to evaluate $\mathbf{A}^{\top} v$, for some v in \mathbb{R}^{n} .

We notice that the complexity of both algorithms is $\mathcal{O}(n)$ despite intervening a matrix multiplication. This is due to the special structure of **A** which in the case of sorted observations has a block diagonal structure with each block having equal columns.

$$\begin{bmatrix}
H_{i_1} & H_{i_2} & H_{i_3} & H_{i_4} \\
0 & 0 & 0 \\
1/3 & 1/3 & 1/3 \\
2/3 & 2/3 & 2/3
\end{bmatrix}$$

$$\begin{bmatrix}
0 & 0 \\
1/2 & 1/2
\end{bmatrix}$$

$$\begin{bmatrix}
0 & 0 \\
1/2 & 1/2
\end{bmatrix}$$

$$\begin{bmatrix}
0 & 0 \\
1/3 & 1/3 & 1/3 \\
2/3 & 2/3 & 2/3
\end{bmatrix}$$

Figure 1: Strucutre of \mathbf{A} in the case of sorted observations with group sizes of identical occurrences times being 3, 2, 1, 3 respectively.

 Algorithm 1 Evaluate Av

 input: $v \in \mathbb{R}^n$

 init : $o \in \mathbb{R}^n$

 1 for $l = 1, \dots, m$ do

 2 $\left| \begin{array}{c} \bar{v}_{H_{i_l}} \leftarrow \operatorname{sum}(v_{H_{i_l}}) \\ o_{H_{i_l}} \leftarrow \bar{v}_{H_{i_l}} \times [0, \frac{1}{|H_{i_l}|}, \dots, \frac{|H_{i_l}|-1}{|H_{i_l}|}] \\ a ext{ return } o \end{array} \right|$

 Algorithm 2 Evaluate $\mathbf{A}^{\top} v$

 input: $v \in \mathbb{R}^n$

 init : $o \in \mathbb{R}^n$

 1 for $l = 1, \cdots, m$ do

 2
 $w_{H_{i_l}} \leftarrow \langle v_{H_{i_l}}, [0, \frac{1}{|H_{i_l}|}, \dots, \frac{|H_{i_l}|-1}{|H_{i_l}|}] \rangle$

 3
 $o_{H_{i_l}} \leftarrow w_{H_{i_l}} \times [1, \dots, 1]$

 4 return o

References

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