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Bachelor's thesis

Probabilistic algorithms for computing the LTS estimate

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Acknowledgements THANKS to everybody

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In Prague on March 7, 2019

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Abstrakt

V několika větách shrňte obsah a přínos této práce v českém jazyce.

Klíčová slova LTS odhad, lineÃarnÃŋ regrese, optimalizace, nejmenÅaÃŋ usekanÃľ ÄDtvrece, metoda nejmenÅaÃŋch ÄDtvercÅŕ, outliers

Abstract

The least trimmed squares (LTS) method is a robust version of the classical method of least squares used to find an estimate of coefficients in the linear regression model. Computing the LTS estimate is known to be NP-hard, and hence suboptimal probabilistic algorithms are used in practice.

Keywords LTS, linear regressin, robust estimator, least trimmed squares, ordinary least squares, outliers, outliers detection

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Introduction

CHAPTER 1

Linear Regression

- 1.1 Description
- 1.2 Computation
- 1.3 Downfalls

The Least trimmed squares

2.0.1 Objective function

2.0.1.1 Problems

Algorithms

3.1 FAST-LTS

In this section we will introduce FAST-LTS algorithm[1]. It is, as well as in other cases, iterative algorithm. We will discuss all main components of the algorithm starting with its core idea called concentration step which authors simply call C-step.

3.1.1 C-step

We will show that from existing LTS estimate \hat{w}_{old} we can construct new LTS estimate \hat{w}_{new} which objective function is less or equal to the old one. Based on this property we will be able to create sequence of LTS estimates which will lead to better results.

Theorem 1. Consider dataset consisting of x_1, x_2, \ldots, x_n explanatory variables where $x_i \in \mathbb{R}^p$, $\forall x_i = (x_1^i, x_2^i, \ldots, x_p^i)$ where $x_1^i = 1$ and its corresponding y_1, y_2, \ldots, y_n response variables. Let's also have $\hat{\boldsymbol{w}}_0 \in \mathbb{R}^p$ any p-dimensional vector and $H_0 = \{h_i; h_i \in \mathbb{Z}, 1 \leq h_i \leq n\}, |H_0| = h$. Let's now mark $RSS(\hat{\boldsymbol{w}}_0) = \sum_{i \in H_0} (r_0(i))^2$ where $r_0(i) = y_i - (w_1^0 x_1^i + w_2^0 x_2^i + \ldots + w_p^0 x_p^i)$. Let's take $\hat{n} = \{1, 2, \ldots, n\}$ and mark $\pi : \hat{n} \to \hat{n}$ permutation of \hat{n} such that $|r_0(\pi(1))| \leq |r_0(\pi(2))| \leq \ldots \leq |r_0(\pi(n))|$ and mark $H_1 = \{\pi(1), \pi(2), \ldots \pi(h)\}$ set of h indexes corresponding to h smallest absolute residuals $r_0(i)$. Finally take $\hat{\boldsymbol{w}}_1^{OLS(H_1)}$ ordinary least squares fit on H_1 subset of observations and its corresponding $RSS(\hat{\boldsymbol{w}}_1) = \sum_{i \in H_1} (r_1(i))^2$ sum of least squares. Then

$$RSS(\hat{\boldsymbol{w}}_1) \le RSS(\hat{\boldsymbol{w}}_0) \tag{3.1}$$

Proof. Because we take h observations with smallest absolute residuals r_0 , then for sure $\sum_{i \in H_1} (r_0(i))^2 \leq \sum_{i \in H_0} (r_0(i))^2 = RSS(\hat{\boldsymbol{w}}_0)$. When we take into account that Ordinary least squares fit OLS_{H_1} minimize objective function of H_1 subset of observations, then for sure $RSS(\hat{\boldsymbol{w}}_1) = \sum_{i \in H_1} (r_1(i))^2 \leq C_{i}$

 $\sum_{i \in H_1} (r_0(i))^2$. Together we get

$$RSS(\hat{\boldsymbol{w}}_1) = \sum_{i \in H_1} (r_1(i))^2 \le \sum_{i \in H_1} (r_0(i))^2 \le \sum_{i \in H_0} (r_0(i))^2 = RSS(\hat{\boldsymbol{w}}_0)$$

Corollary 2. Based on previous theorem, using some $\hat{\boldsymbol{w}}^{OLS(H_{old})}$ on H_{old} subset of observations we can construct H_new subset with corresponding $\hat{\boldsymbol{w}}^{OLS(H_{new})}$ such that $RSS(\hat{\boldsymbol{w}}^{OLS(H_{new})}) \leq RSS(\hat{\boldsymbol{w}}^{OLS(H_{old})})$. With this we can apply above theorem again on $\hat{\boldsymbol{w}}^{OLS(H_{new})}$ with H_{new} . This will lead to the iterative sequence of $RSS(\hat{\boldsymbol{w}}_{old}) \leq RSS(\hat{\boldsymbol{w}}_{new}) \leq \ldots$ One step of this process is described by following pseudocode. Note that for C-step we actually need only $\hat{\boldsymbol{w}}$ without need of passing H.

Algorithm 1: C-step

```
Input: dataset consiting of X \in \mathbb{R}^{n \times p} and y \in \mathbb{R}^{n \times 1}, \hat{w}_{old} \in \mathbb{R}^{p \times 1}

Output: \hat{w}_{new}, H_{new}

1 R \leftarrow \emptyset;

2 for i \leftarrow 1 to n do

3 | R \leftarrow R \cup \{|y_i - \hat{w}_{old}x_i^T|\};

4 end

5 H_{new} \leftarrow select set of h smallest absolute residuals from R;

6 \hat{w}_{new} \leftarrow OLS(H_{new});

7 return \hat{w}_{new}, H_{new};
```

Observation 3. Time complexity of algorithm C-step 1 is same as time complexity as OLS. Thus $O(p^2n)$ **TODO**

Lemma 4. Time complexity of OLS on $X^{n \times p}$ and $Y^{n \times 1}$ is $O(p^2n)$.

Proof. Normal equation of OLS is $\hat{\boldsymbol{w}} = (\boldsymbol{X}^T\boldsymbol{X})^{-1}\boldsymbol{X}^T\boldsymbol{Y}$. Time complexity of matrix multiplication $\boldsymbol{A}^{m\times n}$ and $\boldsymbol{B}^{n\times p}$ is $\sim \mathcal{O}(mnp)$. Time complexity of matrix $\boldsymbol{C}^{m\times m}$ is $\sim \mathcal{O}(m^3)$ So we need to compute $\boldsymbol{A} = \boldsymbol{X}^T\boldsymbol{X} \sim \mathcal{O}(p^2n)$ and $\boldsymbol{B} = \boldsymbol{X}^T\boldsymbol{Y} \sim \mathcal{O}(pn)$ and $\boldsymbol{C} = \boldsymbol{A}^{-1} \sim \mathcal{O}(p^3)$ and finally $\boldsymbol{C}\boldsymbol{B} \sim \mathcal{O}(p^2)$. That give us $\mathcal{O}(p^2n + pn + p^3 + p^2)$. Because $\mathcal{O}(p^2n)$ and $\mathcal{O}(p^3)$ asymptotically dominates over $\mathcal{O}(p^2)$ and $\mathcal{O}(pn)$ we can write $\mathcal{O}(p^2n + p^3)$.

TODO CO zo toho je vic? Neni casove narocnejsi vynasobeni X^TX nez inverze, kdyz bereme v uvahu n >> p???

Proof. In C-step we must compute n absolute residuals. Computation of one absolute residual consist of matrix multiplication of shapes $\times p$ and $pt \times 1$ that give us $\mathcal{O}(p)$. Rest is in constant time. So time of computation n residuals is $\mathcal{O}(np)$. Next we must select set of h smallest residuals which can be done in $\mathcal{O}(n)$ using modification of algorithm QuickSelect. reference: TODO Finally we must compute \hat{w} OLS estimate on h subset of data. Because h is linearly

dependent on n, we can say that it is $\mathcal{O}(p^2n + p^3)$ which is asymptotically dominant against previous steps which are $\mathcal{O}(np+n)$.

As we stated above, repeating algorithm C-step will lead to sequence of $\hat{\boldsymbol{w}}_1, \hat{\boldsymbol{w}}_2 \dots$ on subsets $H_1, H_2 \dots$ with corresponding residual sum of squares $RSS(\hat{\boldsymbol{w}}_1) \geq RSS(\hat{\boldsymbol{w}}_2) \geq \dots$ One could ask if this sequence will converge, so that $RSS(\hat{\boldsymbol{w}}_i) == RSS(\hat{\boldsymbol{w}}_{i+1})$. Answer to this question will be answered by the following theorem.

Theorem 5. Sequence of C-step will converge to $\hat{\boldsymbol{w}}_{m}$ after maximum of $m = \binom{n}{h}$ so that $RSS(\hat{\boldsymbol{w}}_{m}) == RSS(\hat{\boldsymbol{w}}_{n}), \forall n \geq m$ where n is number of data samples and h is size of subset H_{i} .

Proof. Since the $RSS(\hat{\boldsymbol{w}}_i)$ is non-negative and $RSS(\hat{\boldsymbol{w}}_i) \leq RSS(\hat{\boldsymbol{w}}_{i+i})$ the sequence will converge. $\hat{\boldsymbol{w}}_i$ is computed out of subset $H_i \subset 1, 2, \ldots, n$. When there are finite number of subsets of size h out of n samples, namely $\binom{n}{h}$, the sequence will converge at the latest after this number of steps.

Above theorem gives us clue to create algorithm described by following pseudocode.

Algorithm 2: Repeat-C-step

```
Input: dataset consiting of X \in \mathbb{R}^{n \times p} and y \in \mathbb{R}^{n \times 1}, \hat{w}_{old} \in \mathbb{R}^{p \times 1}, H_0
      Output: \hat{\boldsymbol{w}}_{final}, H_{final}
  1 \hat{w}_{new} \leftarrow \emptyset;
  2 H_{new} \leftarrow \emptyset;
  3 RSS_{new} \leftarrow \infty;
  4 while True do
              RSS_{old} \leftarrow RSS(\hat{\boldsymbol{w}}_{old});
  5
             \hat{\boldsymbol{w}}_{new}, H_{new} \leftarrow \boldsymbol{X}, \boldsymbol{y}, \hat{\boldsymbol{w}}_{old};
  6
             RSS_{new} \leftarrow RSS(\boldsymbol{\hat{w}_{new}});
  7
             if RSS_{old} == RSS_{new} then
  8
                    break
  9
10
             end
             \hat{w}_{old} \leftarrow \hat{w}_{new}
11
12 end
13 return \hat{\boldsymbol{w}}_{new}, H_{new};
```

It is important to note, that although maximum number of steps of this algorithm is $\binom{n}{h}$ in practice it is very low, most often under 20 steps. TODO nejaky hezky grafik ktery to ukazuje.... That is not enough for algorithm Repeat-C-step to converge to global minimum, but it is necessary condition. That gives us, as stated in [1]

 $Choose lot of initial subsets H_1 and one a chof the mapply algorithm Repeat-C-step. From all convergeds (3.2)$

Before we can construct final algorithm we must decide how to choose initial subset H_1 and how many of them means "lot of". Let's focus first on how to choose initial subset H_1 .

3.1.2 Choosing initial H_1 subset

We have couple of different options.

Random

Not random

```
Input: A finite set A = \{a_1, a_2, \dots, a_n\} of integers

Output: The largest element in the set

1 max \leftarrow a_1

2 for i \leftarrow 2 to n do

3 | if a_i > max then

4 | max \leftarrow a_i

5 | end

6 end

7 return max
```

Algorithm 8 is a greedy change-making algorithm (Slide 19 in Class Slides).

```
Input: A set C = \{c_1, c_2, \dots, c_r\} of denominations of coins, where c_i > c_2 > \dots > c_r and a positive number n
Output: A list of coins d_1, d_2, \dots, d_k, such that \sum_{i=1}^k d_i = n and k is minimized

1 C \leftarrow \emptyset
2 for i \leftarrow 1 to r do

3 | while n \geq c_i do

4 | C \leftarrow C \cup \{c_i\}
5 | n \leftarrow n - c_i
6 | end
7 end
8 return C
```

Algorithm 14 and Algorithm 13 will find the first duplicate element in a sequence of integers.

In this section we will describe FAST-LTS algorithm and its main properties. The main idea of this algorithm is based on the fact that from one approximation of the algorithm we can compute another which can have lower

```
Input: A sequence of integers \langle a_1, a_2, \dots, a_n \rangle
   Output: The index of first location with the same value as in a
                previous location in the sequence
 \mathbf{1} \ location \leftarrow 0
 i \leftarrow 2
 3 while i \leq n and location = 0 do
        j \leftarrow 1
 4
        while j < i and location = 0 do
 \mathbf{5}
            if a_i = a_j then
 6
                location \leftarrow i
 7
            else
 8
             j \leftarrow j + 1
 9
            end
10
        end
11
        i \leftarrow i+1
13 end
14 return location
```

```
Input: A sequence of integers \langle a_1, a_2, \dots, a_n \rangle
    Output: The index of first location with the same value as in a
                previous location in the sequence
 1 location \leftarrow 0
 i \leftarrow 2
 3 while i \leq n \wedge location = 0 do
        j \leftarrow 1
 4
        while j < i \land location = 0 do
 5
            if a_i = a_j then location \leftarrow i
 6
 7
            else j \leftarrow j+1
 9
        end
10
        i \leftarrow i+1
12 end
13 return location
```

3. Algorithms

objective function. TA DAAAAAAAAAAAAAAAAA Theorem 1: [2] Let w0 ... wp be the LTS estimate. for each data sample we can compute —y-wx—

- 3.2 Exact algorithm
- 3.3 Feasible solution
- **3.4** MMEA
- 3.5 Branch and bound
- 3.6 Adding row

CHAPTER 4

Experiments

- 4.1 Data
- 4.2 Results
- 4.3 Outlier detection

Conclusion

Bibliography

- [1] Rousseeuw, P. J.; Driessen, K. V. An Algorithm for Positive-Breakdown Regression Based on Concentration Steps. In *Data Analysis: Scientific Modeling and Practical Application*, edited by M. S. W. Gaul, O. Opitz, Springer-Verlag Berlin Heidelberg, 2000, pp. 335–346.
- [2] Rybicka, J. LaTeX pro začátečníky. Brno: Konvoj, third edition, ISBN 80-7302-049-1.

Appendix A

Datasets

 ${\bf GUI}$ Graphical user interface

XML Extensible markup language

 $_{\text{APPENDIX}}\,B$

Contents of enclosed CD

:	readme.txt	. the file with CD contents description
	exe	the directory with executables
	src	the directory of source codes
	wbdcm	implementation sources
	thesisthe direct	ory of LATEX source codes of the thesis
	text	the thesis text directory
	thesis.pdf	the thesis text in PDF format
	thesis ns	the thesis text in PS format