

# K-sign depth

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# Definition

## $K$ -sign depth

Let  $K \geq 2$  be a natural number and  $u_1, \dots, u_N$  a sequence of residuals. Then their  $K$ -sign depth, or shortly  $K$ -depth, is defined as

$$d_K(u_1, \dots, u_N) = \frac{1}{\binom{N}{K}} \sum_{1 \leq n_1 < \dots < n_K \leq N} \left( \prod_{k=1}^K \mathbb{1} \{ (-1)^k u_{n_k} > 0 \} + \prod_{k=1}^K \mathbb{1} \{ (-1)^k u_{n_k} < 0 \} \right)$$

i.e. the relative number of  $K$ -element subsets with alternating signs.

# Assumptions

In this presentation, we will be working with a standard linear regression model

$$Y_i = \mathbf{X}_i^T \boldsymbol{\beta}_X + \varepsilon_i, \quad i = 1, \dots, N, \quad \boldsymbol{\beta}_X \in \mathbb{R}^p,$$

although  $K$ -depth can be extended to any models of the form

$$Y_i = g(\mathbf{X}_i, \boldsymbol{\beta}_X) + \varepsilon_i$$

where  $\varepsilon_i$  are mutually independent, absolutely continuous, and satisfy

$$\text{Med} [\varepsilon_i] = 0, \quad i = 1, \dots, N.$$

# Basic properties

For a chosen value of  $\beta$ , denote the residuals

$$U_i(\beta) = Y_i - \mathbf{X}_i^T \beta, \quad i = 1, \dots, N.$$

Then the expectation of the  $K$ -depth is given by

$$\mathbb{E} d_K(U_1(\beta_X), \dots, U_N(\beta_X)) = \left(\frac{1}{2}\right)^{K-1}$$

and it holds that

$$d_K(U_1(\beta_X), \dots, U_N(\beta_X)) \xrightarrow[N \rightarrow \infty]{a.s.} \left(\frac{1}{2}\right)^{K-1}.$$

# Computational complexity

There are multiple ways to calculate  $K$ -depth with various degrees of computational complexity. From slowest to fastest:

- Naive method using the definition. -  $\Theta(N^K)$
- Approximation method (Muller 2021). -  $\Theta(N)$
- Block implementation (Muller 2023). -  $\Theta(N)$
- Dynamic programming (Nagy 2024). -  $\Theta(N)$

# Hypothesis testing

$K$ -depth can be used as an alternative goodness-of-fit measure to test the regression model parameter, i.e. to test

$$H_0 : \beta_X \in \Theta^0.$$

This so-called  $K$ -depth test is based on the test statistic

$$T_K(\beta) = T_K(U_1(\beta), \dots, U_N(\beta)) = N \left( d_K(U_1(\beta), \dots, U_N(\beta)) - \left(\frac{1}{2}\right)^{K-1} \right).$$

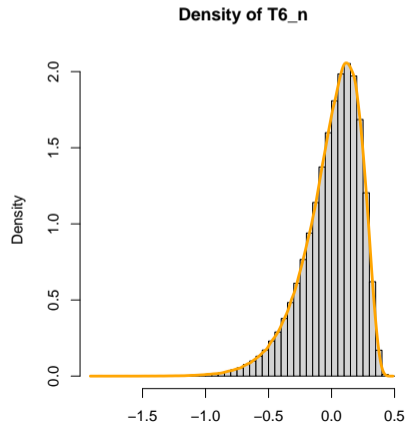
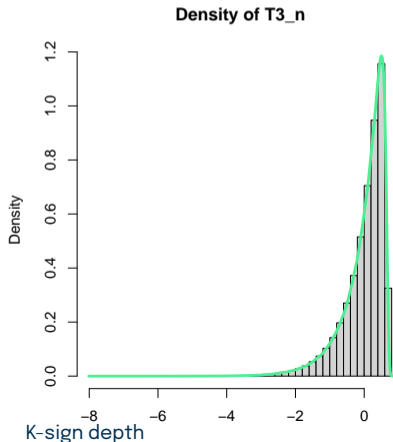
Low values of  $K$ -depth indicate that the assumption on the median has been violated, so consequently low values of the test statistic testify against the parameter  $\beta$ .

# Limiting distribution

It holds that

$$T_K(\beta_X) \xrightarrow[N \rightarrow \infty]{d} \Psi_K(W),$$

where  $\Psi_K(W)$  is a special function of a standard Brownian motion.



# Goodness of fit testing

Consider a simple linear regression model

$$Y_i = \beta_0 + \beta_1 x_i + \varepsilon_i, \quad i = 1, \dots, N$$

where  $\varepsilon_i \sim N(0, 1)$ ,  $\beta_0, \beta_1 \in \mathbb{R}$  and  $x_1, \dots, x_N$  represent realized regressors.



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Denote

$$\hat{\beta}^{LAD} = \operatorname{argmin}_{\beta_0, \beta_1} \sum_{i=1}^N |y_i - \beta_0 - \beta_1 x_i| \quad \hat{\beta}^{LSE} = \operatorname{argmin}_{\beta_0, \beta_1} \sum_{i=1}^N (y_i - \beta_0 - \beta_1 x_i)^2.$$

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Do the estimators  $\hat{\beta}^{LAD}$  and  $\hat{\beta}^{LSE}$  provide a “good” fit?

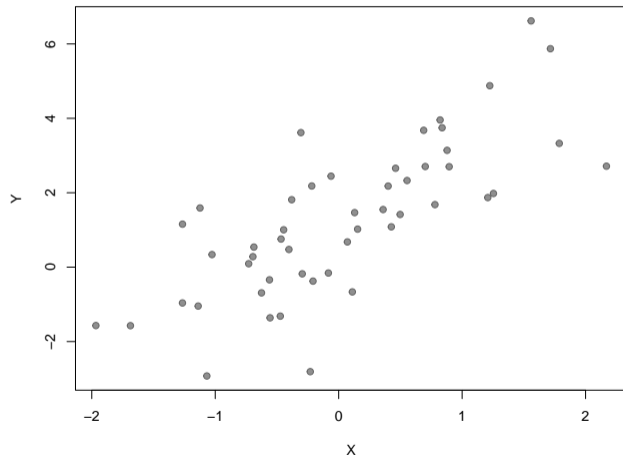
$$H_0^{LAD} : (\beta_0, \beta_1)^\top = \hat{\beta}^{LAD}$$

$$H_0^{LSE} : (\beta_0, \beta_1)^\top = \hat{\beta}^{LSE}$$

# Simulation

Generated dataset of size  $N = 50$ , with true parameters set to  $\beta_0 = 1, \beta_1 = 1.5$ .

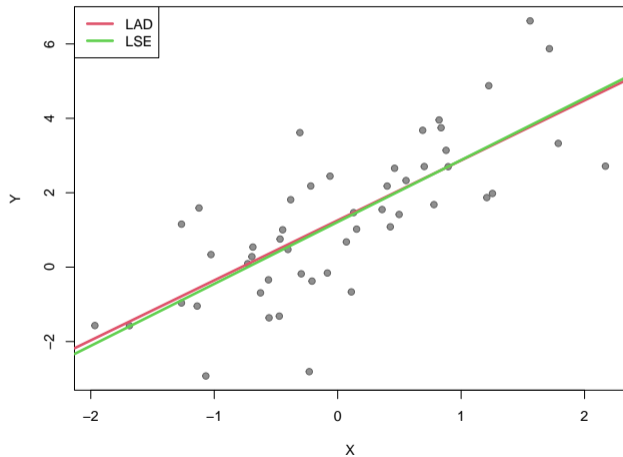
Original Data



# Simulation

Generated dataset of size  $N = 50$ , with true parameters set to  $\beta_0 = 1, \beta_1 = 1.5$ .

Original Data



Estimated coefficients and the respective 3-depths

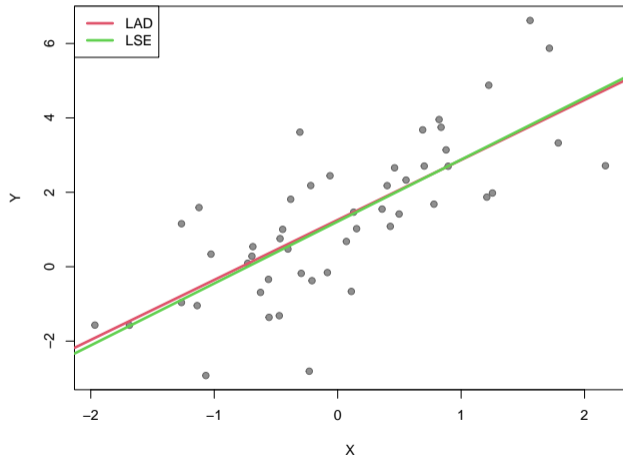
$$\hat{\beta}^{LAD} = (1.25, 1.61)^T; \quad d_3^{LAD} = 0.27$$

$$\hat{\beta}^{LSE} = (1.21, 1.66)^T; \quad d_3^{LSE} = 0.26$$

# Simulation

Null hypotheses not rejected for none of  $K \in \{3, 4, 5, 6, 7\}$ .

Original Data

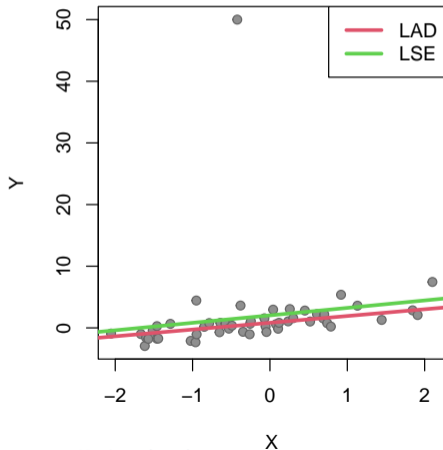


p-values	Estimates
$p_3^{LAD}$	0.8617
$p_4^{LAD}$	0.8296
$p_5^{LAD}$	0.7457
$p_6^{LAD}$	0.7425
$p_7^{LAD}$	0.7094
$p_3^{LSE}$	0.9450
$p_4^{LSE}$	0.8969
$p_5^{LSE}$	0.8452
$p_6^{LSE}$	0.9262
$p_7^{LSE}$	0.8489

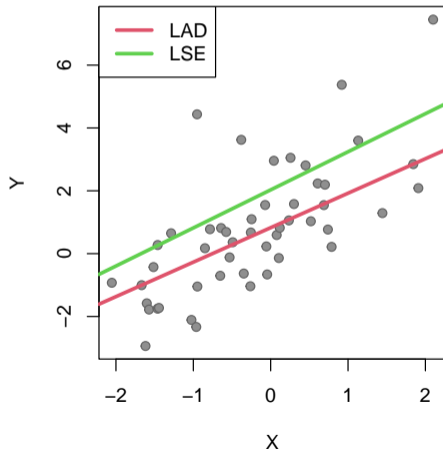
# Outliers in dataset

We added one outlying point in terms of  $Y$ .

Contamination in  $Y$

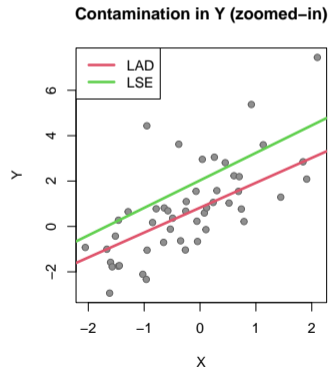
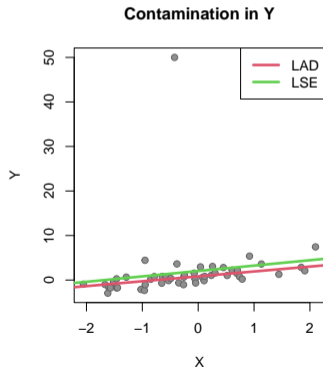


Contamination in  $Y$  (zoomed-in)



# Outliers in dataset

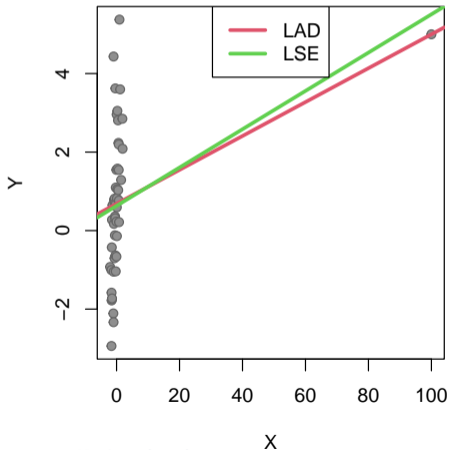
- The estimator  $\hat{\beta}^{LAD}$  is **robust** w.r.t.  $Y$  outliers, the estimator  $\hat{\beta}^{LSE}$  is **not robust**.
- $H_0$ 's for  $\hat{\beta}^{LAD}$  **not rejected** for all  $K$  with  $p$ -values  $\approx 0.9$ .
- $H_0$ 's for  $\hat{\beta}^{LSE}$  **rejected** for all  $K$  with  $p$ -values  $\approx 0.002$ .



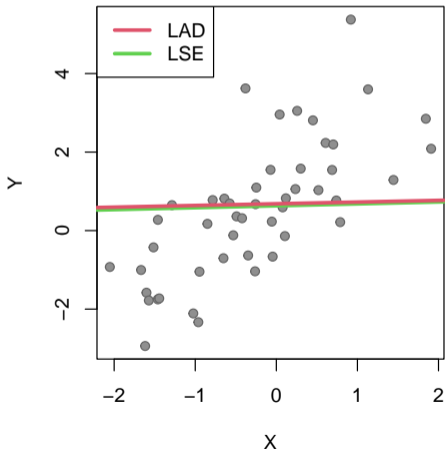
# Outliers in dataset

We added one outlying point in terms of X.

### Contamination in X



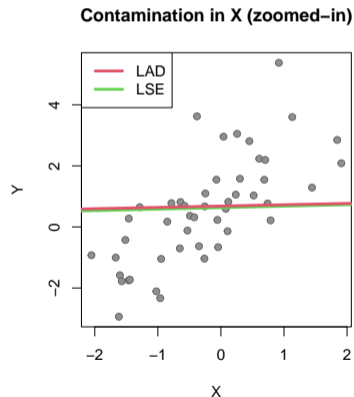
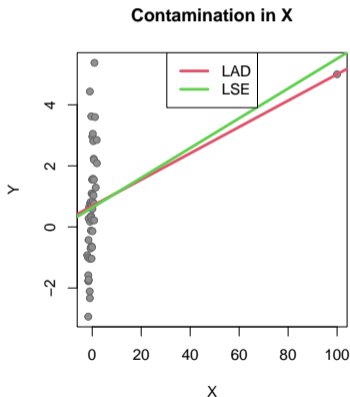
### Contamination in X (zoomed-in)





# Outliers in dataset

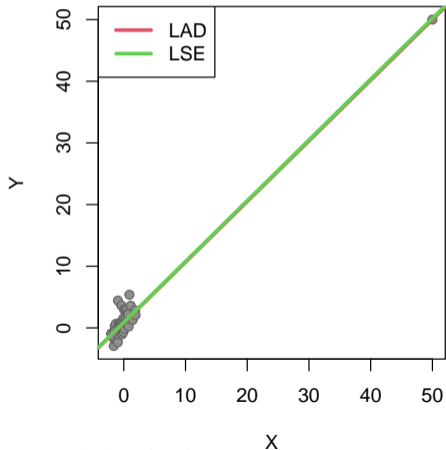
- The estimators  $\hat{\beta}^{LAD}$  and  $\hat{\beta}^{LSE}$  are **not robust** w.r.t.  $X$  outliers.
- $H_0$ 's for both  $\hat{\beta}^{LAD}$  and  $\hat{\beta}^{LSE}$  **rejected** for all  $K$  with  $p$ -values  $\approx 0.00001$ .



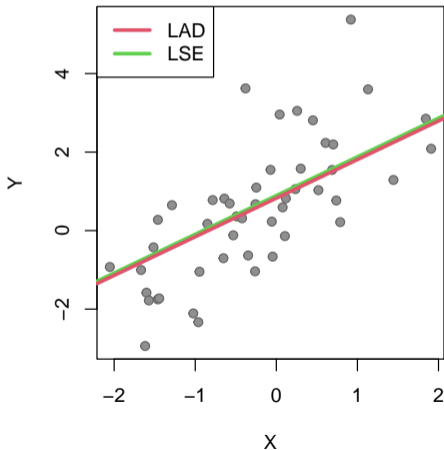
# Outliers in dataset

We added one outlying point in terms of both  $X$  and  $Y$ .

Contamination in X and Y

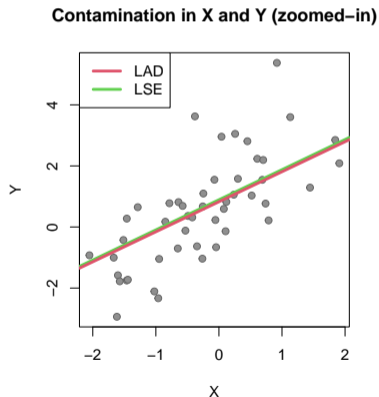
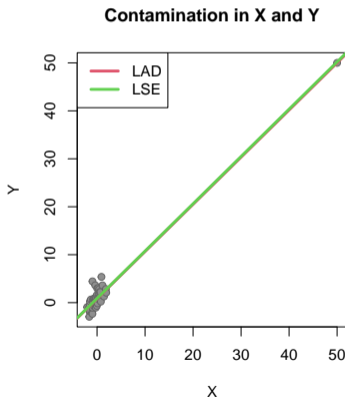


Contamination in X and Y (zoomed-in)



# Outliers in dataset

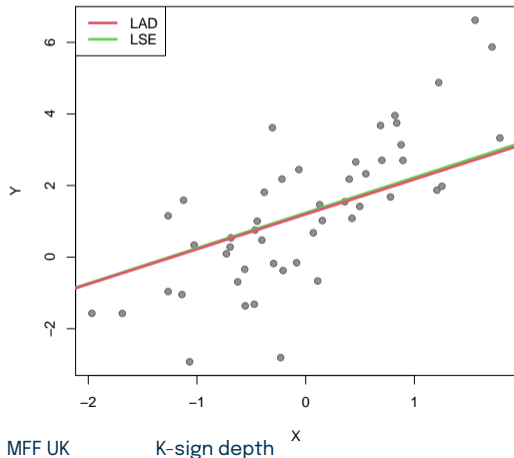
- Clearly a bad fit for both  $\hat{\beta}^{LAD} = (1.20, 0.98)^\top$  and  $\hat{\beta}^{LSE} = (1.24, 0.99)^\top$
- Interestingly enough, null hypotheses **not rejected** for all  $K$ , except for  $H_0 : (\beta_0, \beta_1)^\top = \hat{\beta}^{LSE}$  based on  $d_3$  (that is  $K = 3$ ).



# Further examination

The estimated  $p$ -values are very different between  $\hat{\beta}^{LAD}$  and  $\hat{\beta}^{LSE}$ , despite the two estimates being nearly identical.

Contamination in X and Y (zoomed-in)



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p-values	Estimates
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$p_3^{LAD}$  0.2472

$p_4^{LAD}$  0.5080

$p_5^{LAD}$  0.4508

$p_6^{LAD}$  0.5363

$p_7^{LAD}$  0.6201

$p_3^{LSE}$  **0.0333**

$p_4^{LSE}$  0.1477

$p_5^{LSE}$  0.0721

$p_6^{LSE}$  0.1680

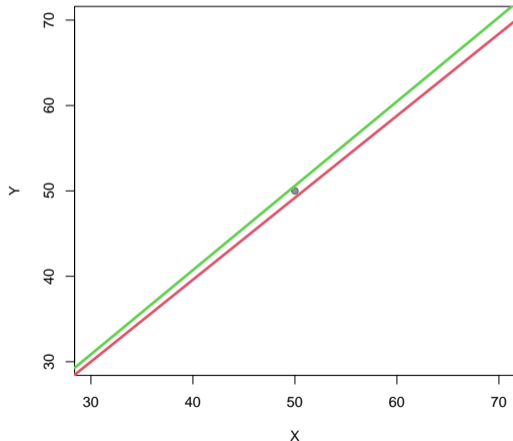
$p_7^{LSE}$  0.1301

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# Further examination

A change in the sign of any residual can make a big difference in  $K$ -depth.

Outlier (zoomed-in)



p-values	Estimates
$p_3^{LAD}$	0.2472
$p_4^{LAD}$	0.5080
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$p_7^{LAD}$	0.6201
$p_3^{LSE}$	<b>0.0333</b>
$p_4^{LSE}$	0.1477
$p_5^{LSE}$	0.0721
$p_6^{LSE}$	0.1680
$p_7^{LSE}$	0.1301

# K-depth estimator

What if instead  $\hat{\beta}^{LAD}$  or  $\hat{\beta}^{LSE}$  we define

$$\hat{\beta}^{d_k} = \operatorname{argmax}_{\beta_0, \beta_1} d_k((\beta_0, \beta_1)^\top)$$

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- Difficult to find already for a simple linear model.
- Not unique

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- What if we only consider regression lines passing through the point  $\bar{x}, \bar{y}$ ?



# K-depth estimator

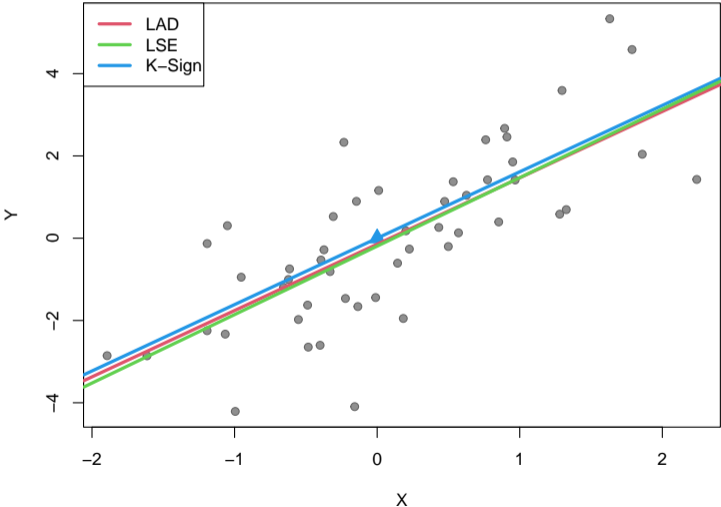
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- Difficult to find already for a simple linear model.
- Not unique
- What if we only consider regression lines passing through the point  $\bar{x}, \bar{y}$ ?
- How about  $\operatorname{median}(x_1, \dots, x_N), \operatorname{median}(y_1, \dots, y_N)$ ?

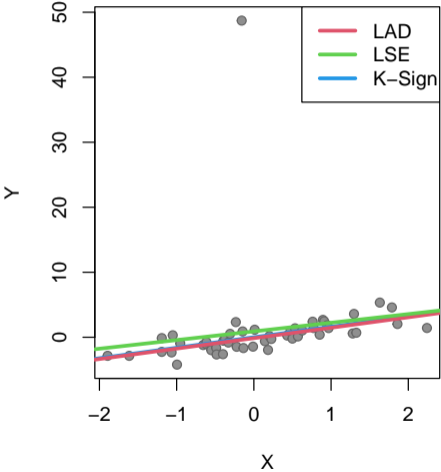
# Outliers from before

Original Data

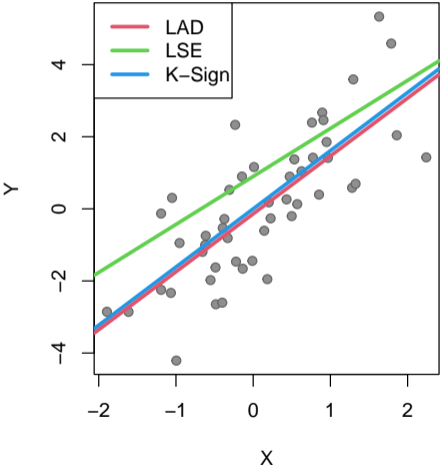


# Outliers from before

Contamination in Y

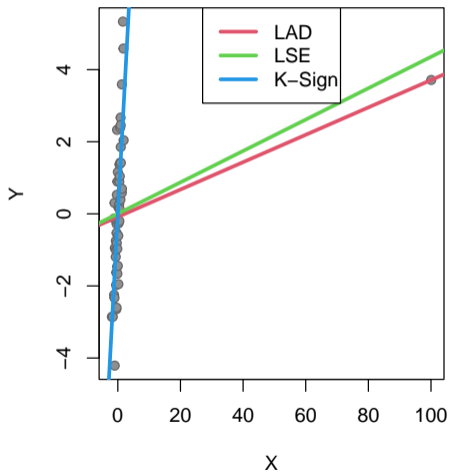


Contamination in Y (zoomed-in)

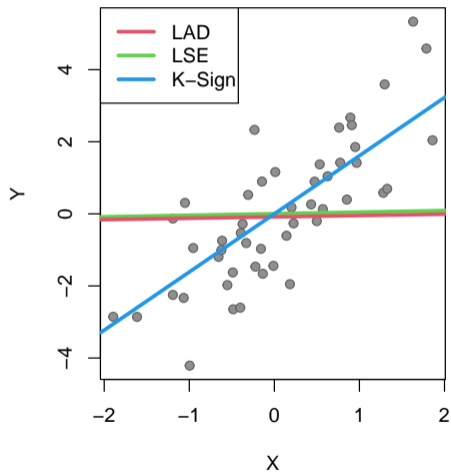


# Outliers from before

Contamination in X

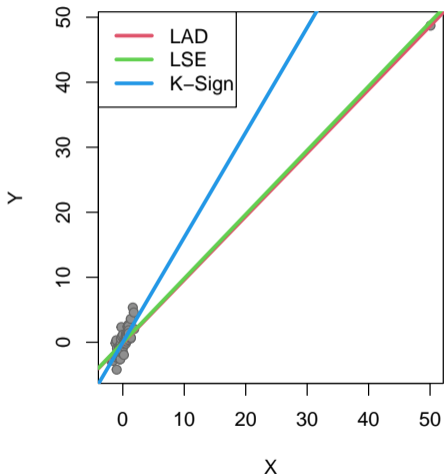


Contamination in X (zoomed-in)

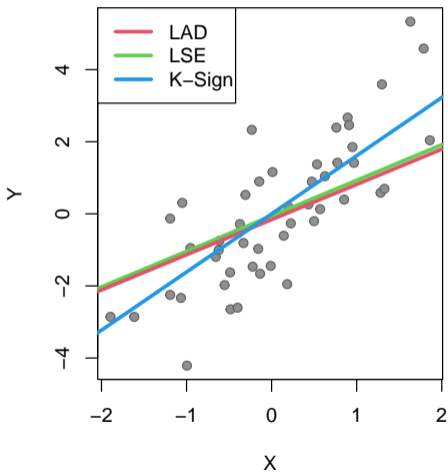


# Outliers from before

Contamination in X and Y



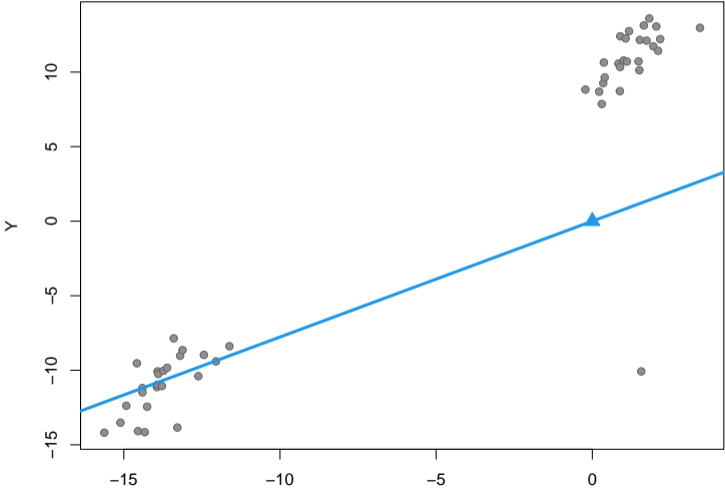
Contamination in X and Y (zoomed-in)



# Is it unbreakable?

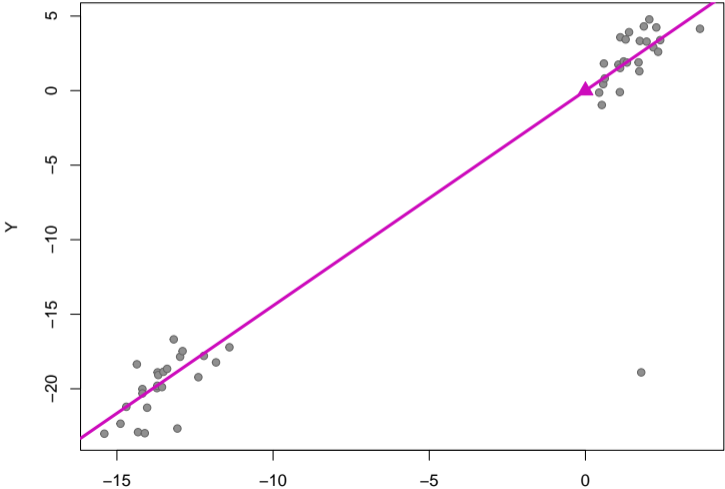
# Is it unbreakable? No.

Counterexample



# Some other possibilities

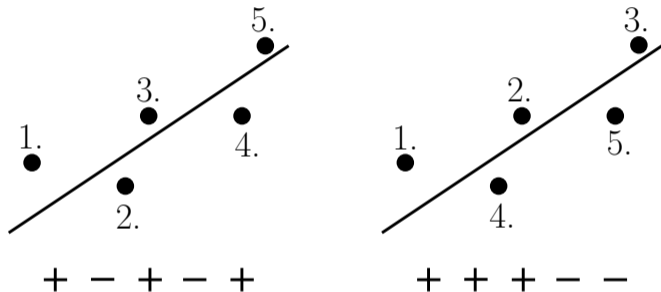
Simplicial median instead of element-wise median





# Multiple regression

The concept of  $K$ -sign depth relies heavily on the **order** of the residuals.



- Straightforward to define in a one-dimensional case.
- Can it be generalized to higher dimensions?

# Ordering in the bivariate case

Consider a multiple linear regression model

$$Y_i = \beta_0 x_i^{(1)} + \beta_1 x_i^{(2)} + \varepsilon_i, \quad i = 1, \dots, N.$$

What if we order residuals based on one of the regressors?

# Ordering in the bivariate case

What if we order the residuals based on one of the regressors?

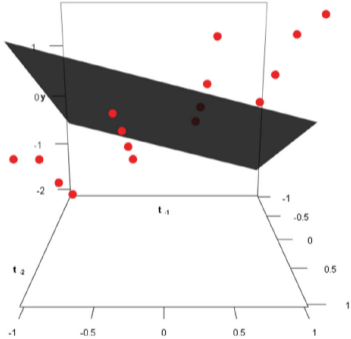
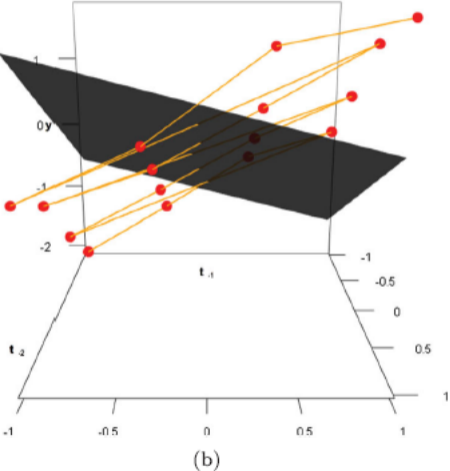
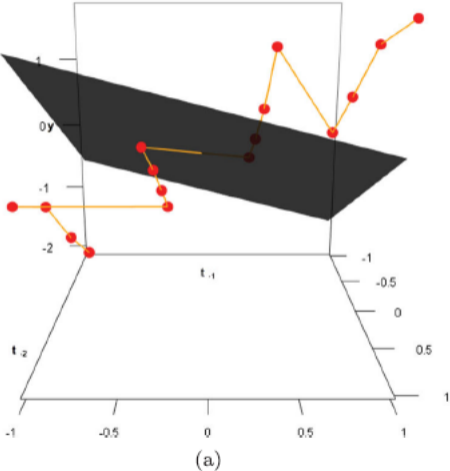


Figure 2. Visualization of the 16 data points and the fitted model (dark grey area).

# Ordering in the bivariate case

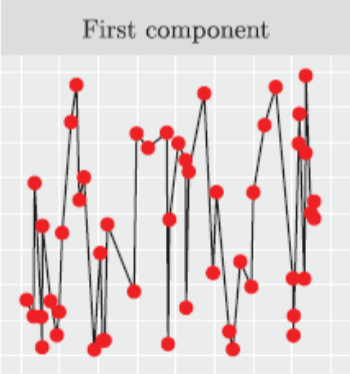


# Ordering in the bivariate case

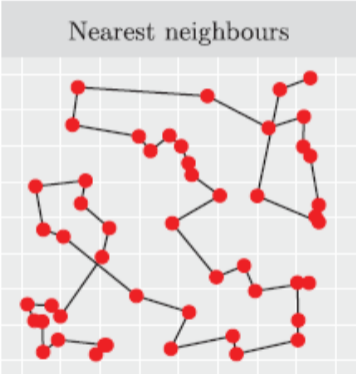
Random order



First component



Nearest neighbours

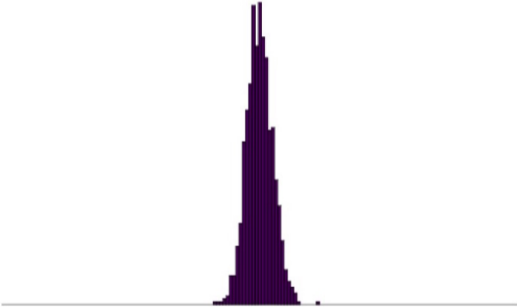


# Cauchy versus Normal errors

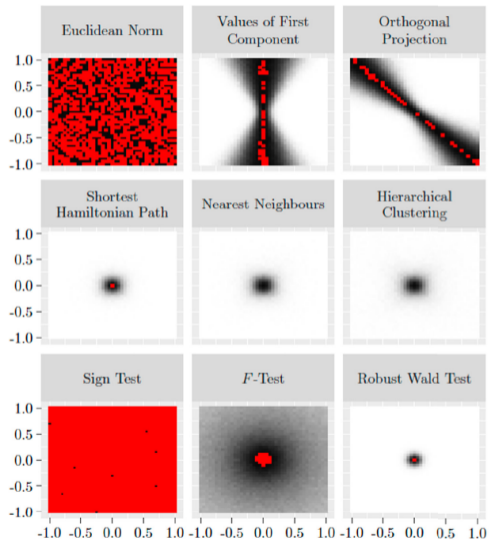
Cauchy



Gaussian



# Power of the tests - simulation



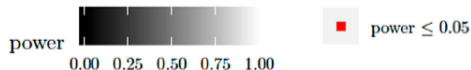
## Model

$$Y_i = \beta_0 x_i^{(1)} + \beta_1 x_i^{(2)} + \varepsilon_i, \quad i = 1, \dots, N,$$

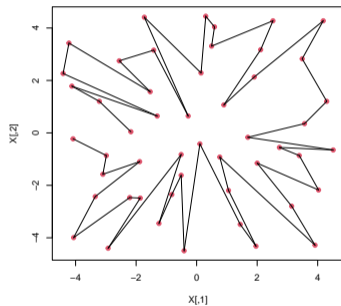
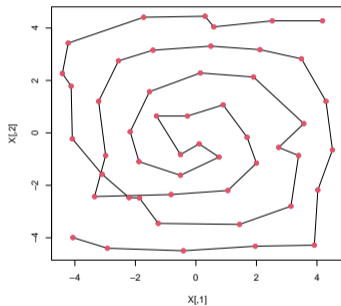
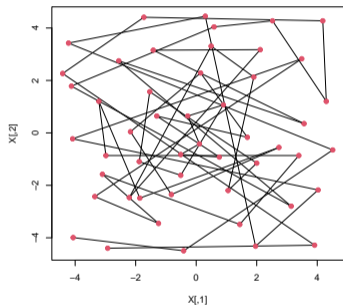
$$\varepsilon_i \sim \text{Cauchy}(0, \sigma^2); \quad x_i^{(1)}, x_i^{(2)} \sim \text{Unif}(-1, 1).$$

## Null Hypothesis

$$H_0 : \beta_0 = \beta_1 = 0.$$

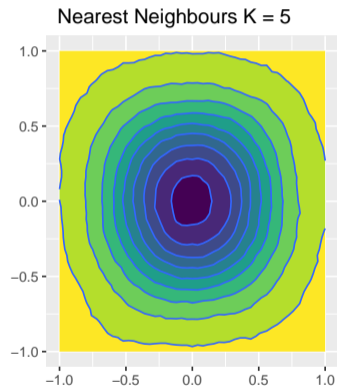
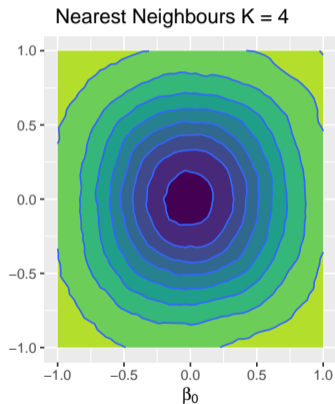
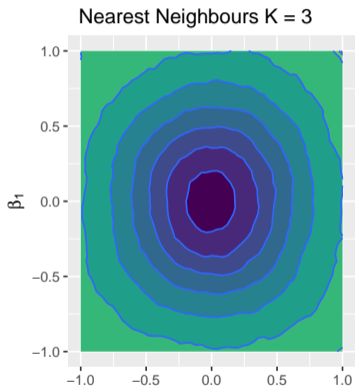


# Experimenting with bivariate ordering

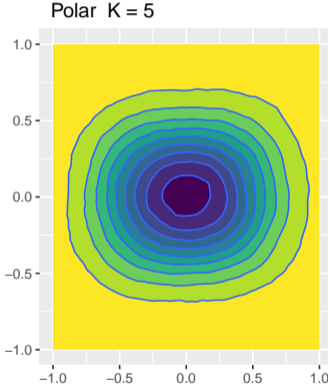
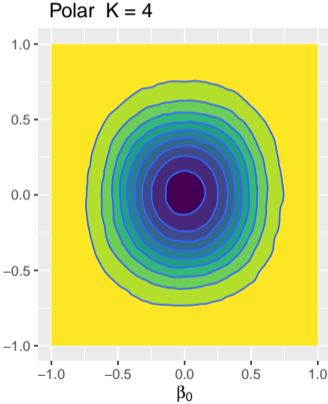
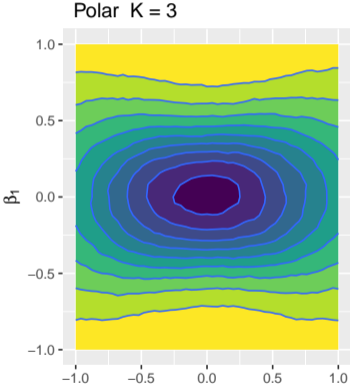




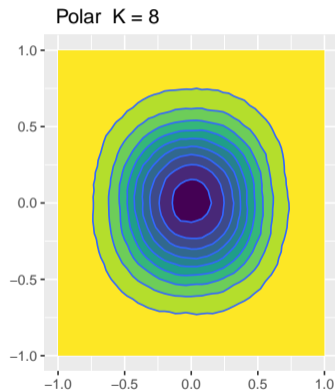
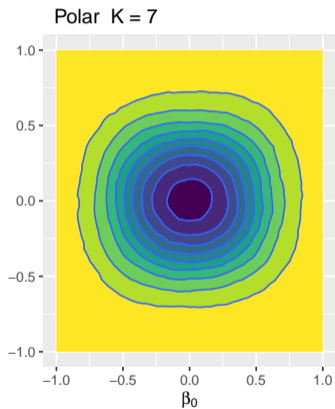
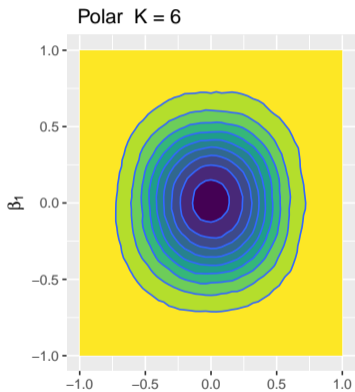
# Power for different $K$ and order



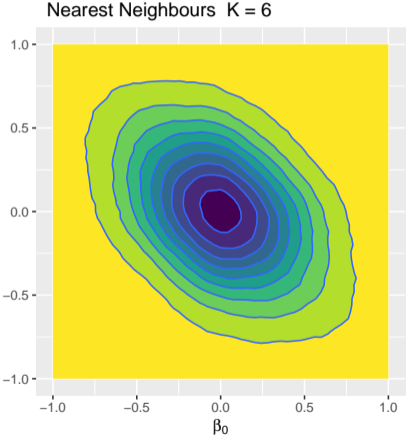
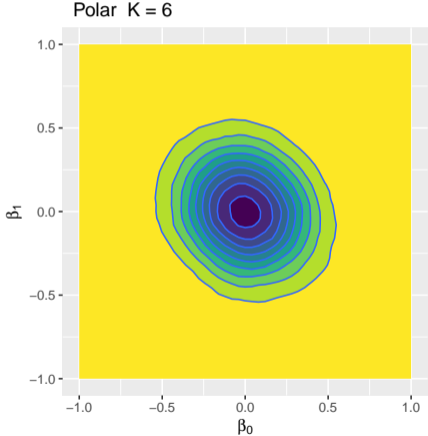
# Power for different $K$ and order



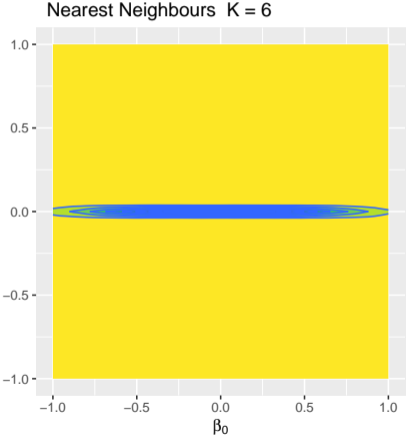
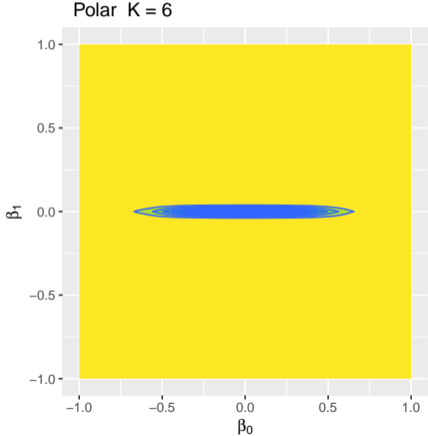
# Power for different $K$ and order



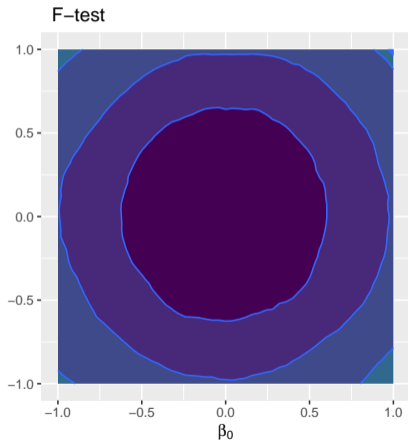
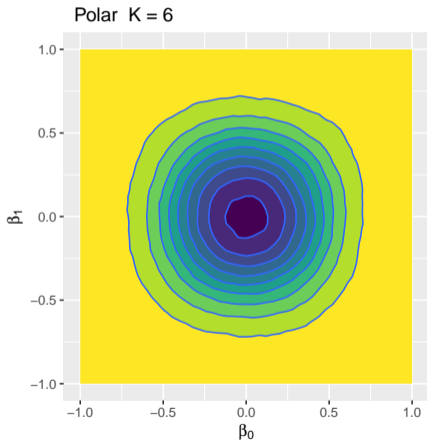
# Exponentially distributed regressors



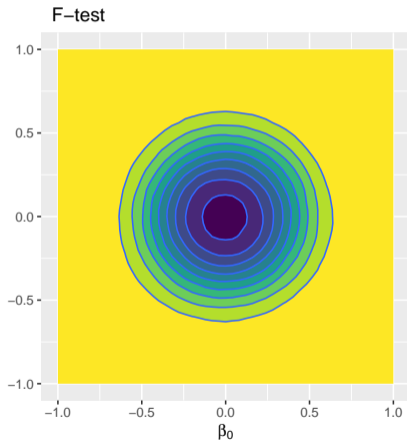
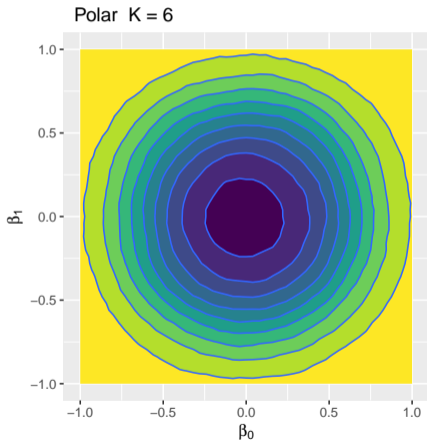
# Bimodal regressors



# K-sign depth versus F-test



# K-sign depth versus F-test



# Power of the tests - high dimensional data

