Advanced Regression: 3b Penalised regression models (ridge, lasso, elastic net)

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Motivation for penalised regression

Penalised regression

- Ridge regression
- Lasso
- Elastic net
- How to tune the regularisation parameter?
- Prediction using penalised regression
- Penalised regression in R

Application

- Diabetes data
- Breast cancer data

The linear model

$$y = \alpha + x\beta + \epsilon$$

- y: Outcome, response, dependent variable
  Dimension: $n \times 1$
- x: Regressors, exposures, covariates, input, explanatory, or independent variables
  Dimension: $n \times p$
- ϵ: Residuals, error
- α: Intercept
- β: Regression coefficients, vector of length p

Classical regression

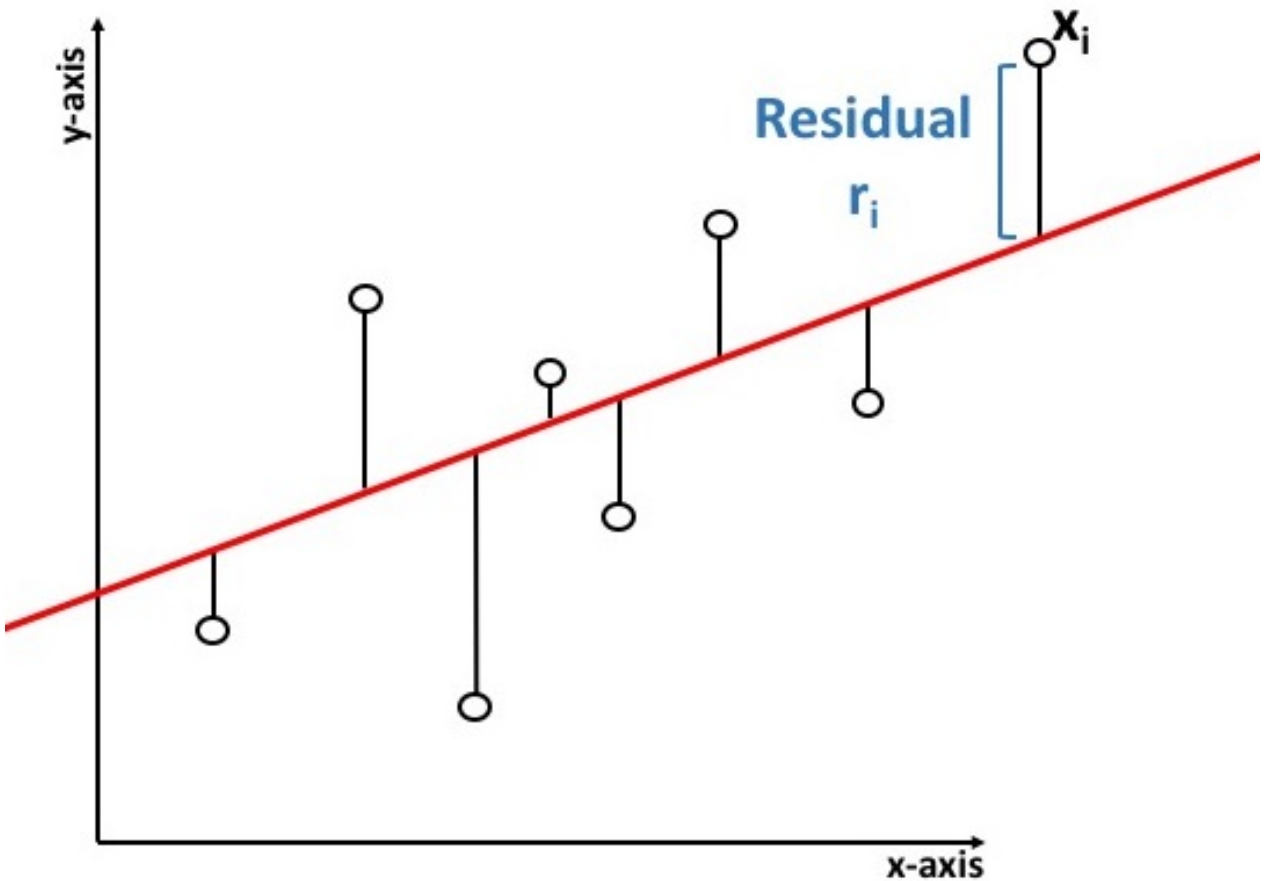
- The ordinary least squares $\hat{\beta}_{OLS}$ is defined as

$$\hat{\beta}_{OLS} = \underbrace{(x^t x)^{(-1)}}_{p \times p} \underbrace{x^t}_{p \times n} \underbrace{y}_{n \times 1}.$$

- The residual sum of squares (RSS) is minimised by the ordinary least squares estimate

$$\begin{aligned} RSS(\alpha, \beta) &= \epsilon_1^2 + ... + \epsilon_i^2 + ... + \epsilon_n^2 \\ &= \sum_{i=1}^{n} \epsilon_i^2 \\ &= \sum_{i=1}^{n} (y_i - \hat{y}_i)^2 \\ &= \sum_{i=1}^{n} (y_i - (\alpha + \beta x_i))^2 \end{aligned}$$

Residual sum of squares (RSS)



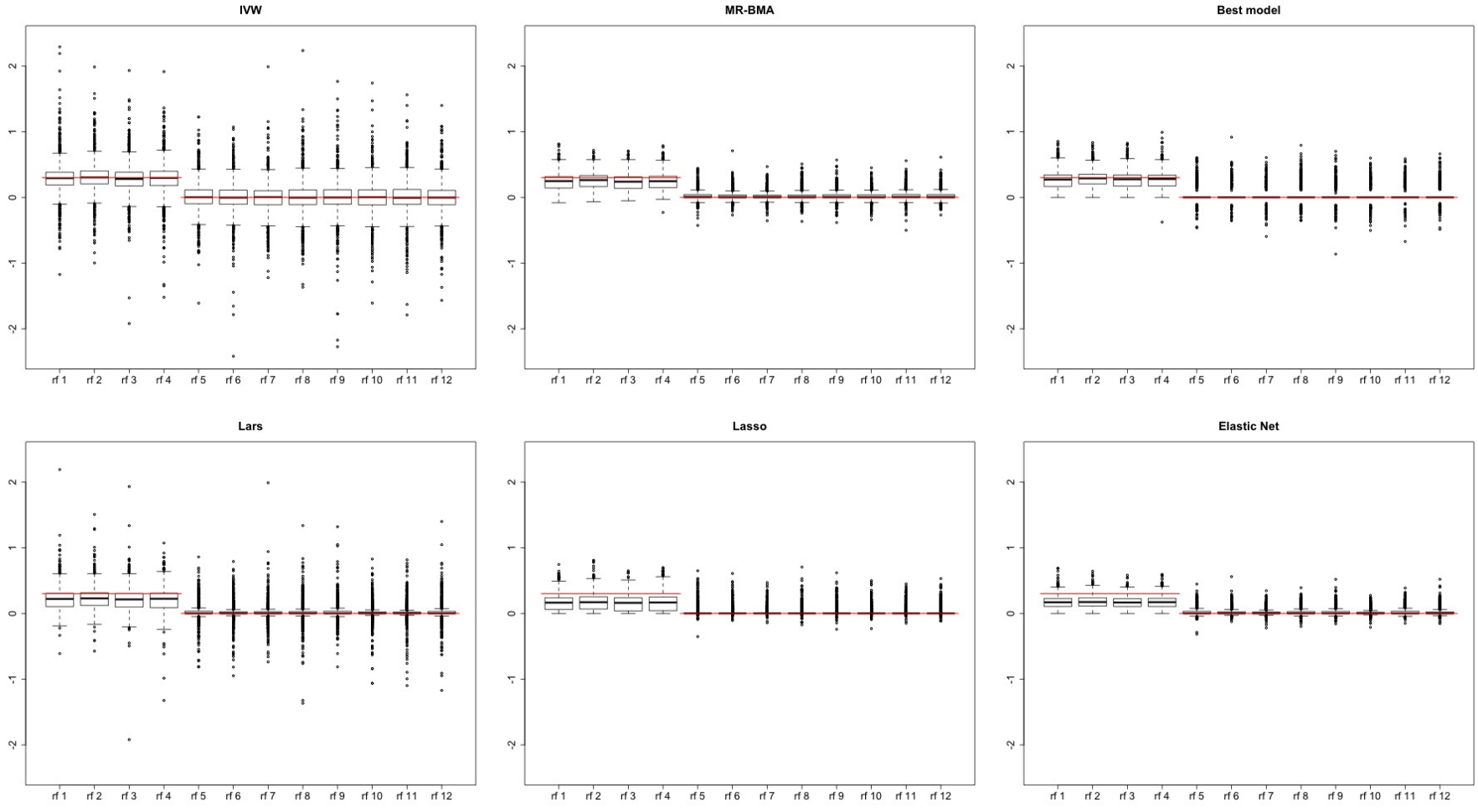
- Note $\sum_{i=1}^{n} \epsilon_i = 0$

Classical regression and high-dimensional data

- When $n << p$ the ordinary least squares cannot be computed because $\underbrace{(x^t x)}_{p \times p}$ is singular (rank n).
- Bias-variance trade-off:
  - The ordinary least squares estimate is best linear unbiased estimator (BLUE).
  - When considering high-dimensional data, the ordinary least squares estimate has a high variability.
  - We rather prefer an estimate that is biased (towards a sensible option, e.g. the Null), but is precise, (ie has low variance).

Bias-variance trade-off

Simulation study with 12 predictor variables



Motivation for penalised least squares

Minimise RSS but with penalty

$$\underset{\alpha,\beta}{argmin} = \underbrace{RSS(\alpha,\beta)}_{\text{Residual Sum of Squares}} + \underbrace{\lambda f(\beta)}_{\text{penalty}}$$

where

- Residual Sum of Squares:
  $RSS(\alpha,\beta) = \sum_{i=1}^{n} (y_i - (\alpha + \beta x_i))^2$
- Penalty term as a function of β: $f(\beta)$
- Regularistion parameter: λ
- The intercept is not penalised

Motivation for penalised least squares

The penalty introduces a bias, so why do it?

- Which variables do we include? Only those for which it is worth to take the penalty.
- Occam's razor: It induces sparsity and favours models with lower complexity (Lasso and elastic net).
- Regularises the inversion of $x^t x$ (Ridge regression).

Different penalty terms define different methods

$$\underset{\alpha,\beta}{\textit{argmin}} = RSS(\alpha, \beta) + \lambda f(\beta)$$

- Ridge regression: L2 penalty

$$\lambda f(\beta) = \lambda \sum_{j=1}^{p} \beta_j^2$$

- Lasso regression: L1 penalty

$$\lambda f(\beta) = \lambda \sum_{j=1}^{p} |\beta_j|$$

- Elastic net regression: L1 + L2 penalty

$$\lambda f(\beta) = \lambda_1 \sum_{j=1}^{p} |\beta_j| + \lambda_2 \sum_{j=1}^{p} \beta_j^2$$

Ridge regression

Ridge regression uses the L2 norm as penalty:

$$\underset{\alpha,\hat{\beta}_{Ridge}}{argmin} = \underbrace{RSS(\alpha,\beta)}_{\text{Residual Sum of Squares}} + \underbrace{\lambda \sum_{j=1}^{p} \beta_j^2}_{\text{penalty}}$$

Interpretation:

- The Ridge regression coefficient $\hat{\beta}_{Ridge}$ is a biased estimate, but has a reduced variance compared to $\hat{\beta}_{OLS}$.
- There is no intrinsic model selection in Ridge regression, all p variables will have $\hat{\beta}_{Ridge} \neq 0$.

Ridge regression and ordinary least squares

The ridge regression estimate is available in closed form

$$\hat{\beta}_{Ridge} = \underbrace{(x^t x + \lambda I)^{(-1)}}_{p \times p} \underbrace{x^t}_{p \times n} \underbrace{y}_{n \times 1},$$

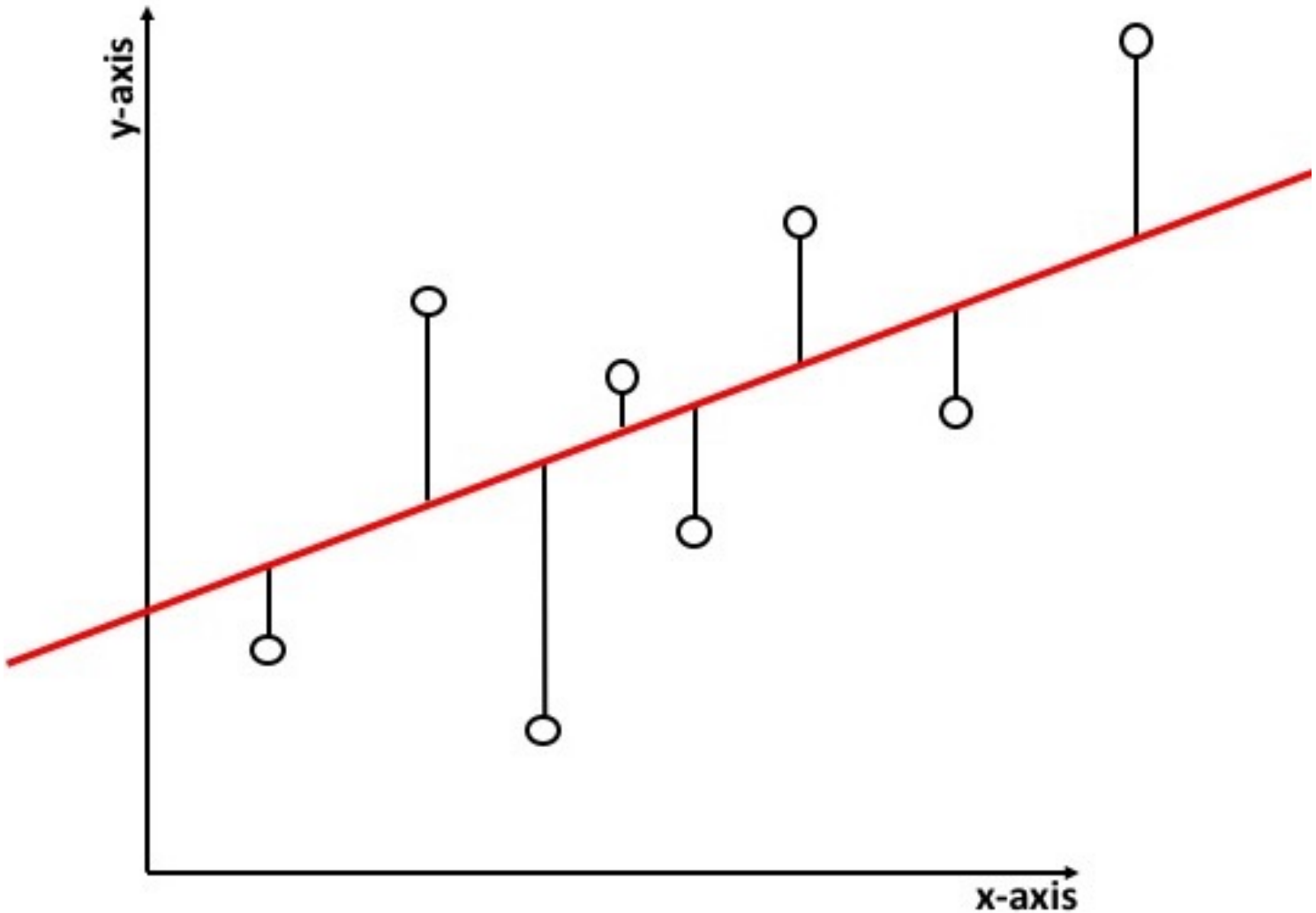
where I is a $p \times p$ diagonal matrix with ones on the diagonal and zero on the off-diagonal

$$x^t x + \lambda I = n \begin{bmatrix} cov(x_1) & cov(x_{12}) & cov(x_{13}) \\ cov(x_{21}) & cov(x_2) & cov(x_{23}) \\ cov(x_{31}) & cov(x_{23}) & cov(x_3) \end{bmatrix} + \begin{bmatrix} \lambda & 0 & 0 \\ 0 & \lambda & 0 \\ 0 & 0 & \lambda \end{bmatrix}.$$

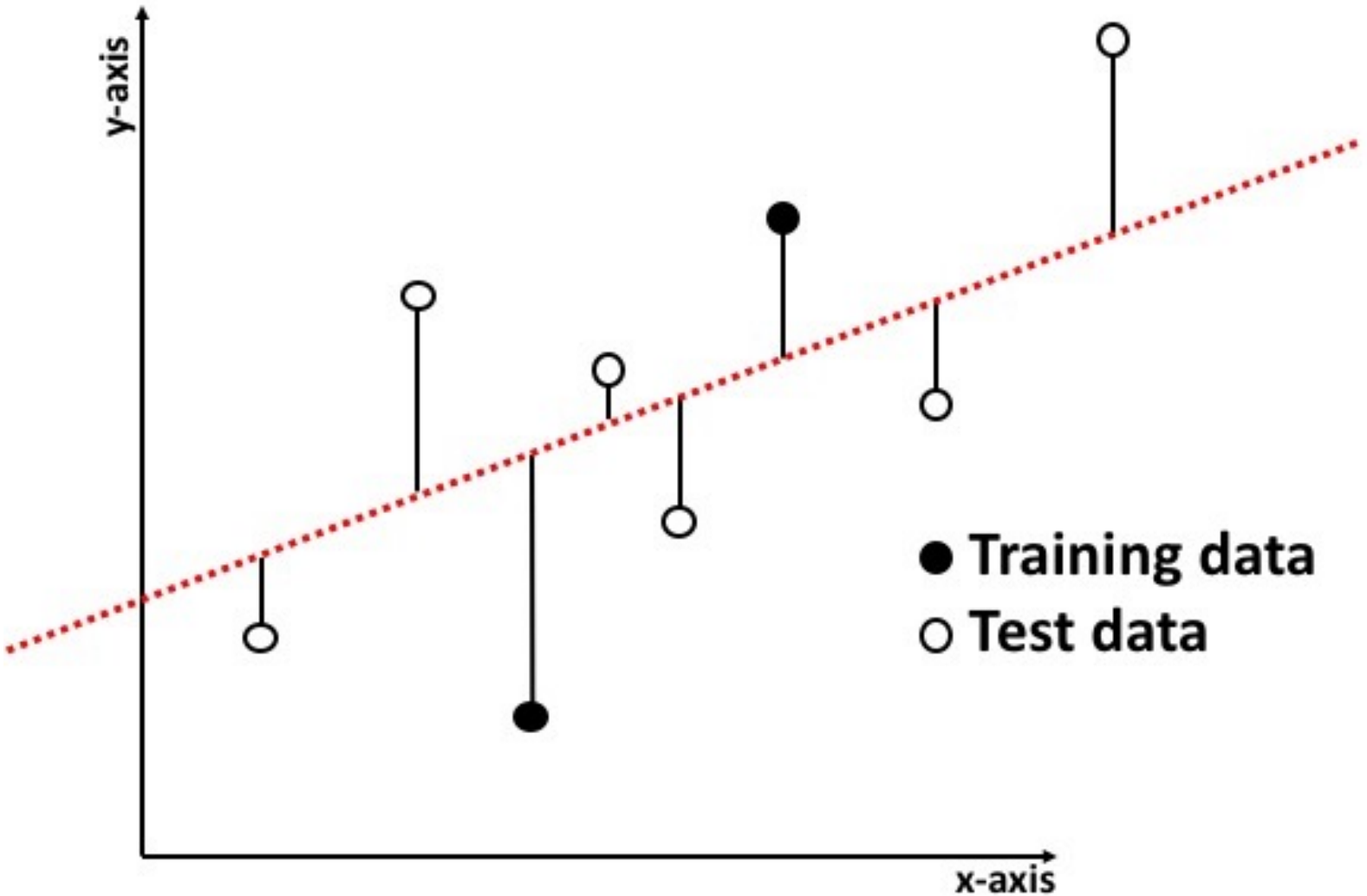
This resembles the OLS estimate apart from $+\lambda I$

$$\hat{\beta}_{OLS} = \underbrace{(x^t x)^{(-1)}}_{p \times p} \underbrace{x^t}_{p \times n} \underbrace{y}_{n \times 1}.$$

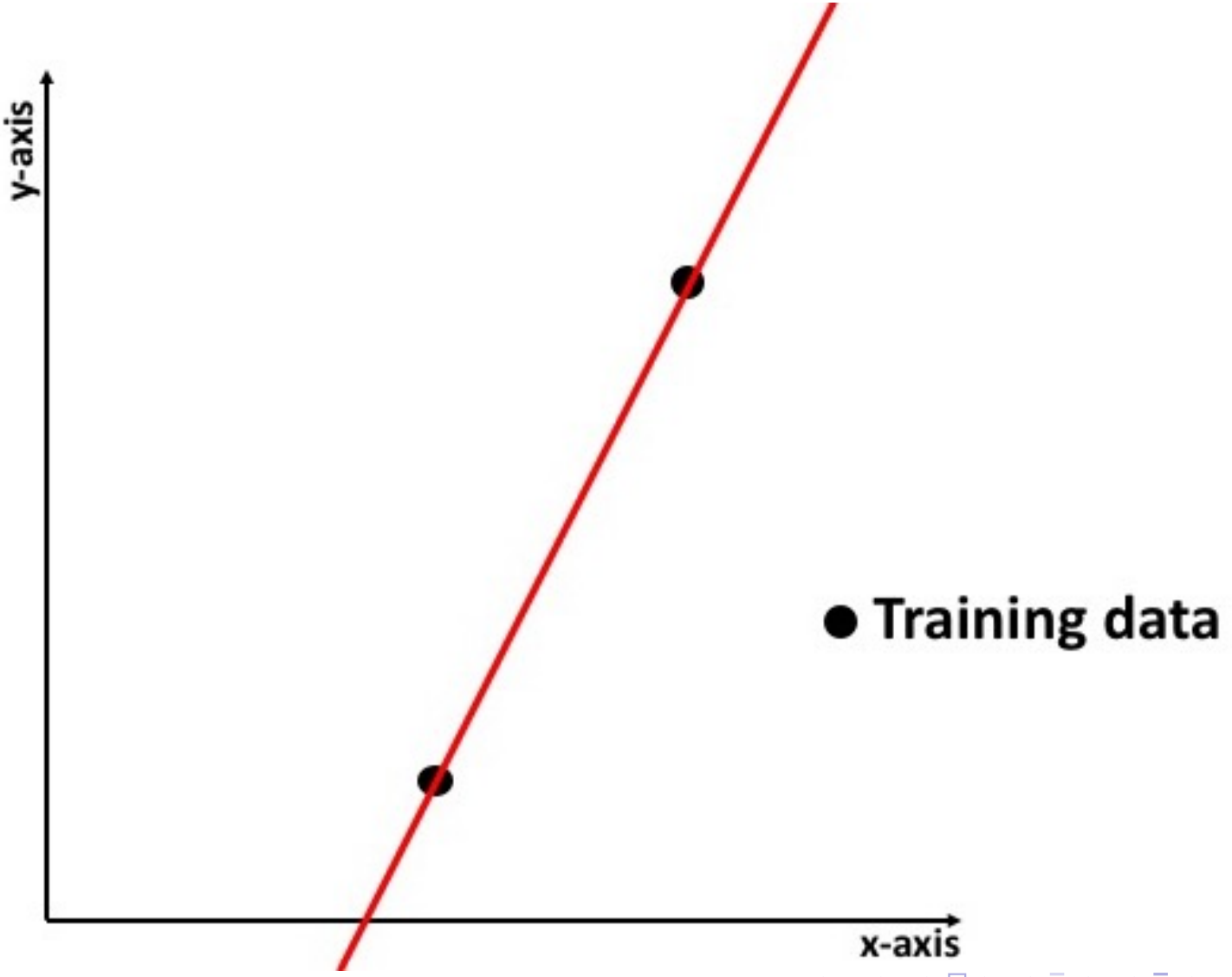
Ridge regression and ordinary least squares



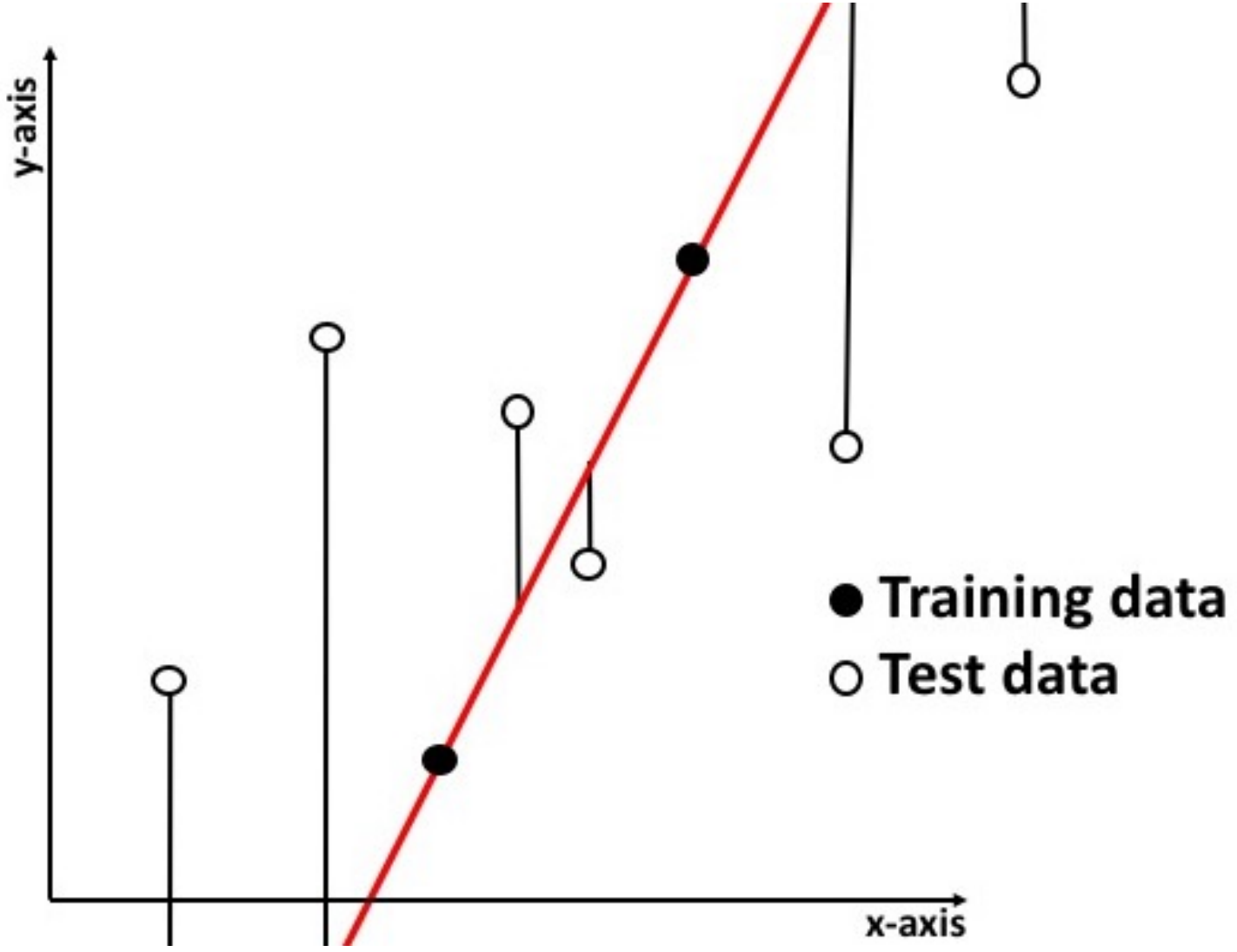
Ridge regression and ordinary least squares



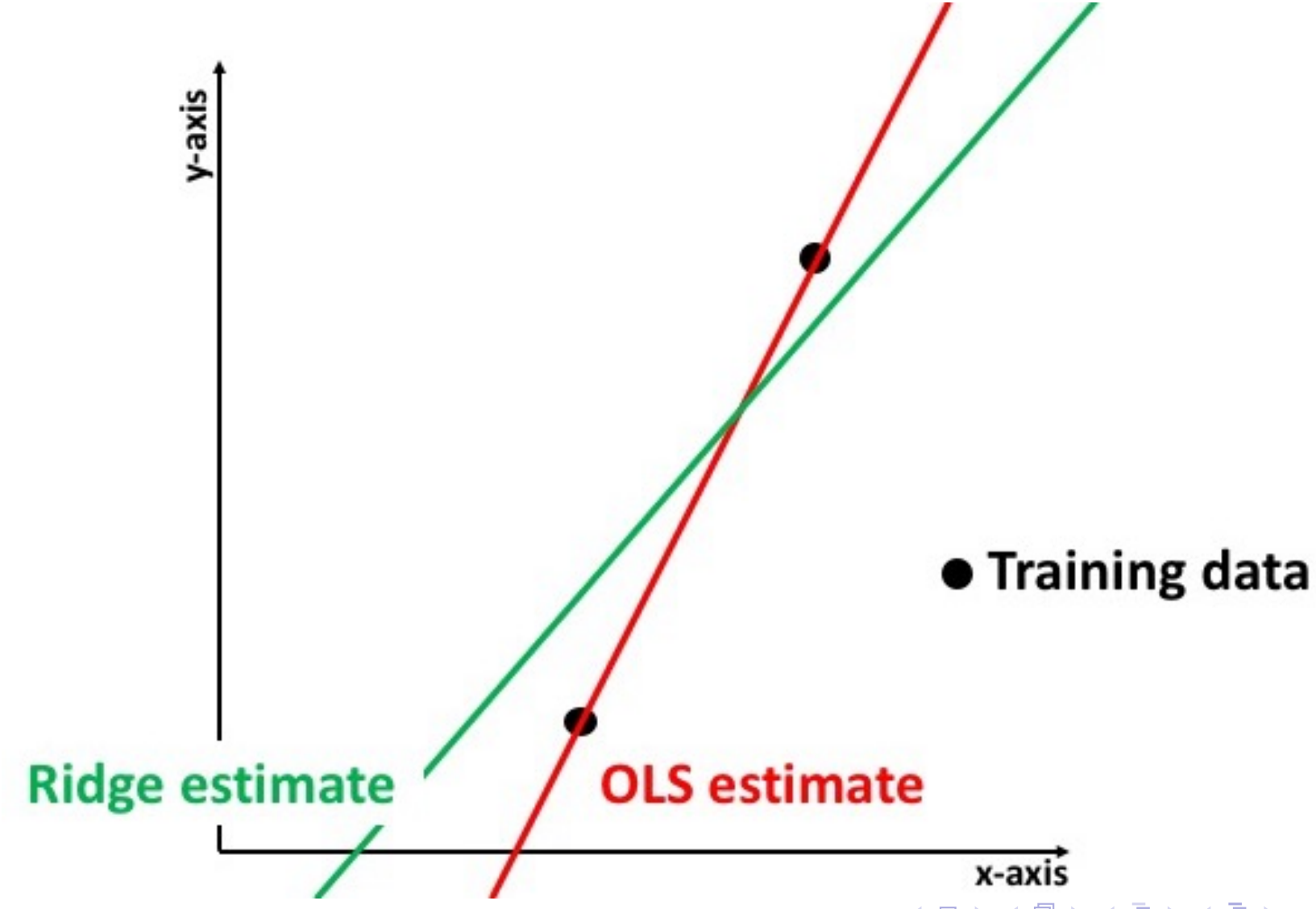
Ridge regression and ordinary least squares



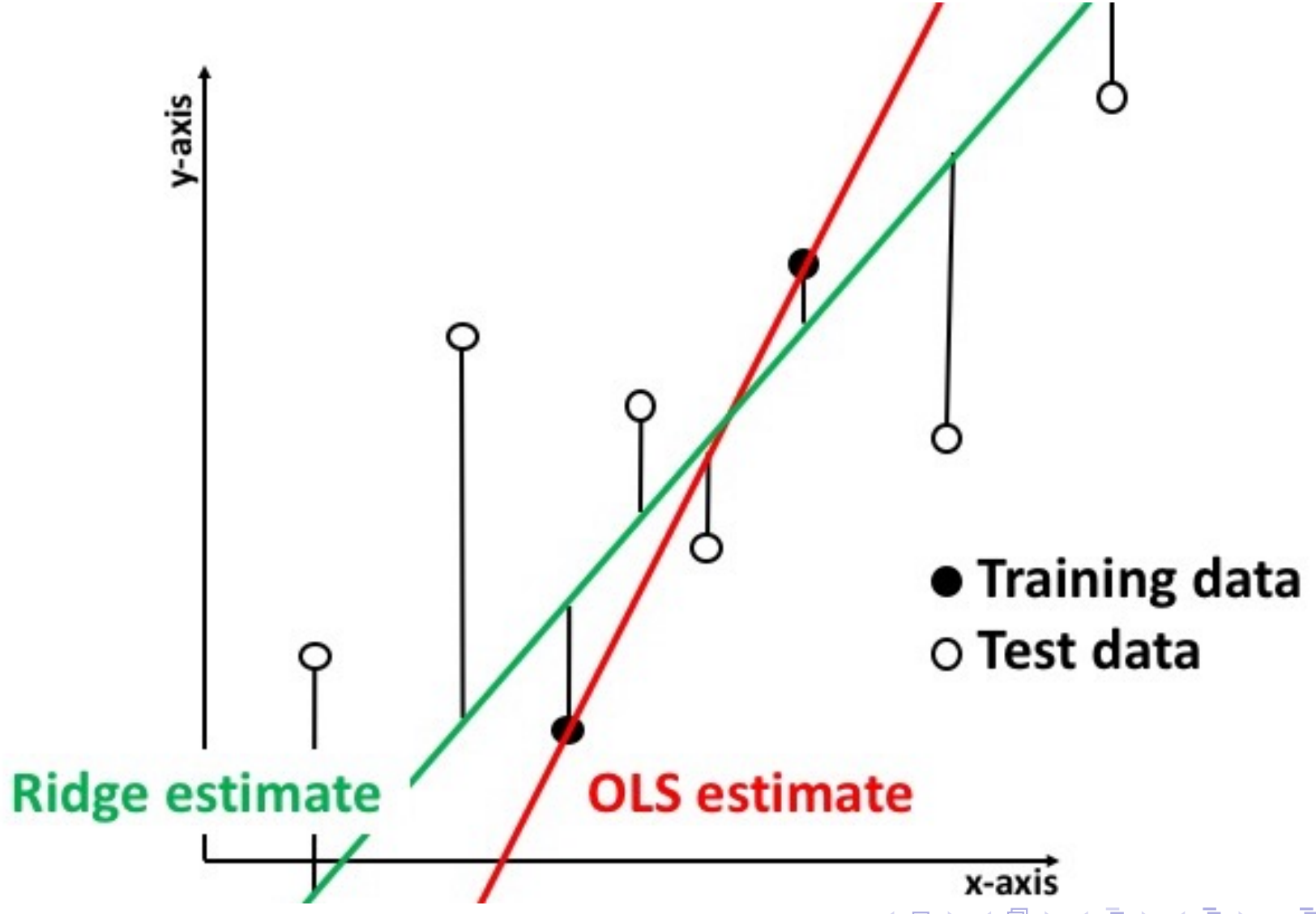
Ridge regression and ordinary least squares



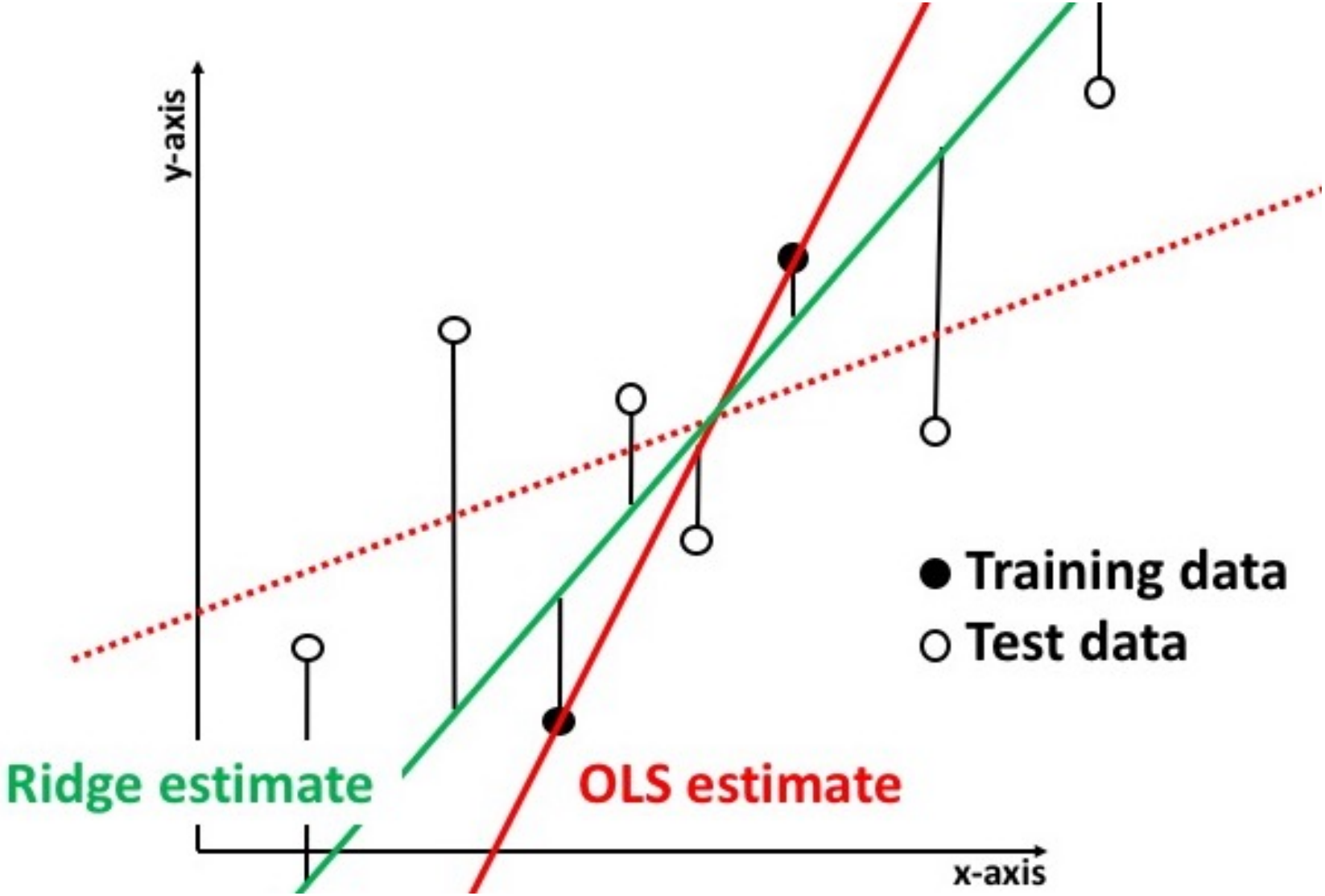
Ridge regression and ordinary least squares



Ridge regression and ordinary least squares



Ridge regression and ordinary least squares



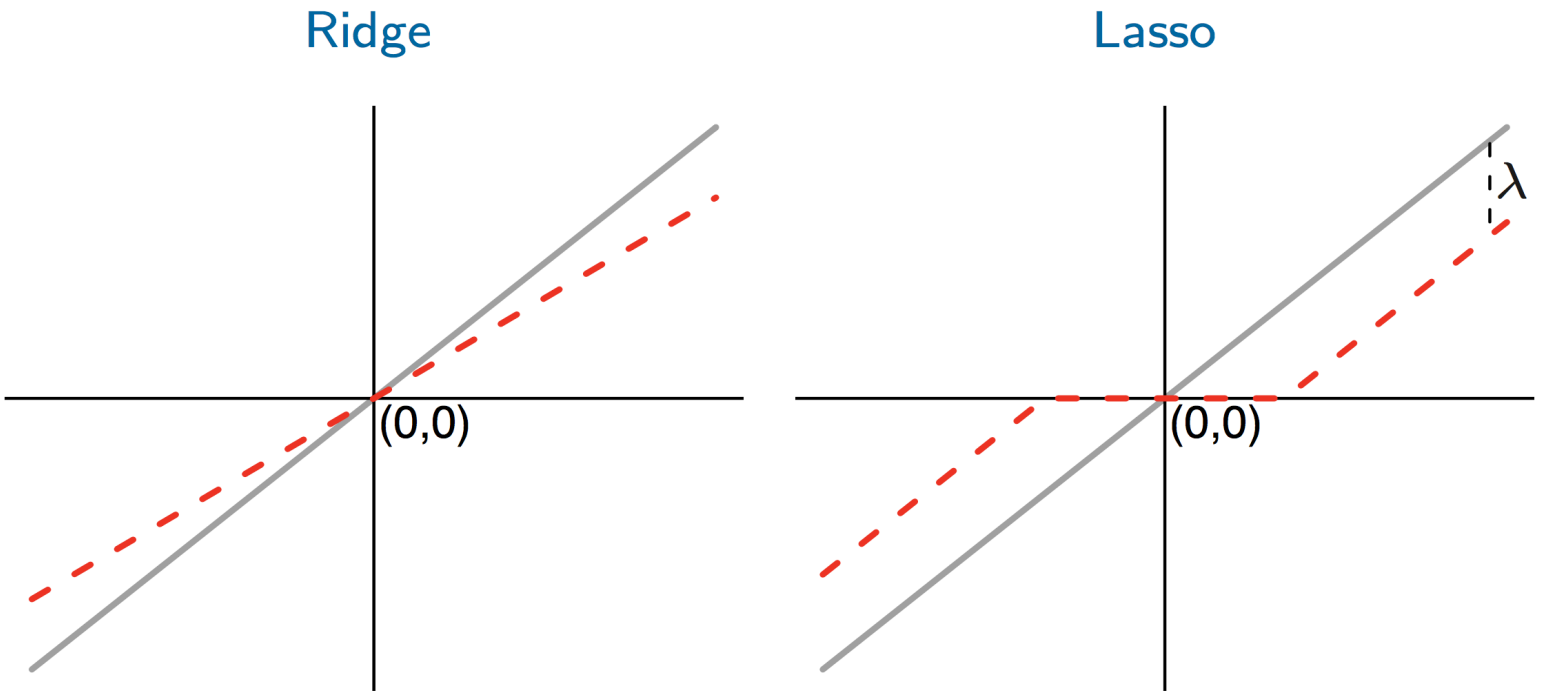
Lasso regression

$$\underset{\hat{\alpha},\hat{\beta}_{Lasso}}{argmin} = \underbrace{RSS(\alpha,\beta)}_{\text{Residual Sum of Squares}} \underbrace{+\lambda \sum_{j=1}^{p} |\beta_j|}_{\text{penalty}}$$

Interpretation:

- The Lasso regression coefficient $\hat{\beta}_{Lasso}$ is a biased estimate, but has a reduced variance compared to $\hat{\beta}_{OLS}$.
- There is an intrinsic model selection in Lasso regression, as it sets certain variables exactly to $\hat{\beta}_{Lasso} = 0$, and thus excludes them from the model.
- When two variables are highly correlated, Lasso includes only one (at random) and not both.

Ridge and lasso: Induced shrinkage



In case of orthogonal (independent) predictors there is a closed form for both estimators:

$$\hat{\beta}_{Ridge} = \hat{\beta}_{OLS}/(1+\lambda)$$
$$\hat{\beta}_{Lasso} = sign(\hat{\beta}_{OLS})(\|\hat{\beta}_{OLS}\| - \lambda)_+$$

Ridge and lasso: Geometric interpretation

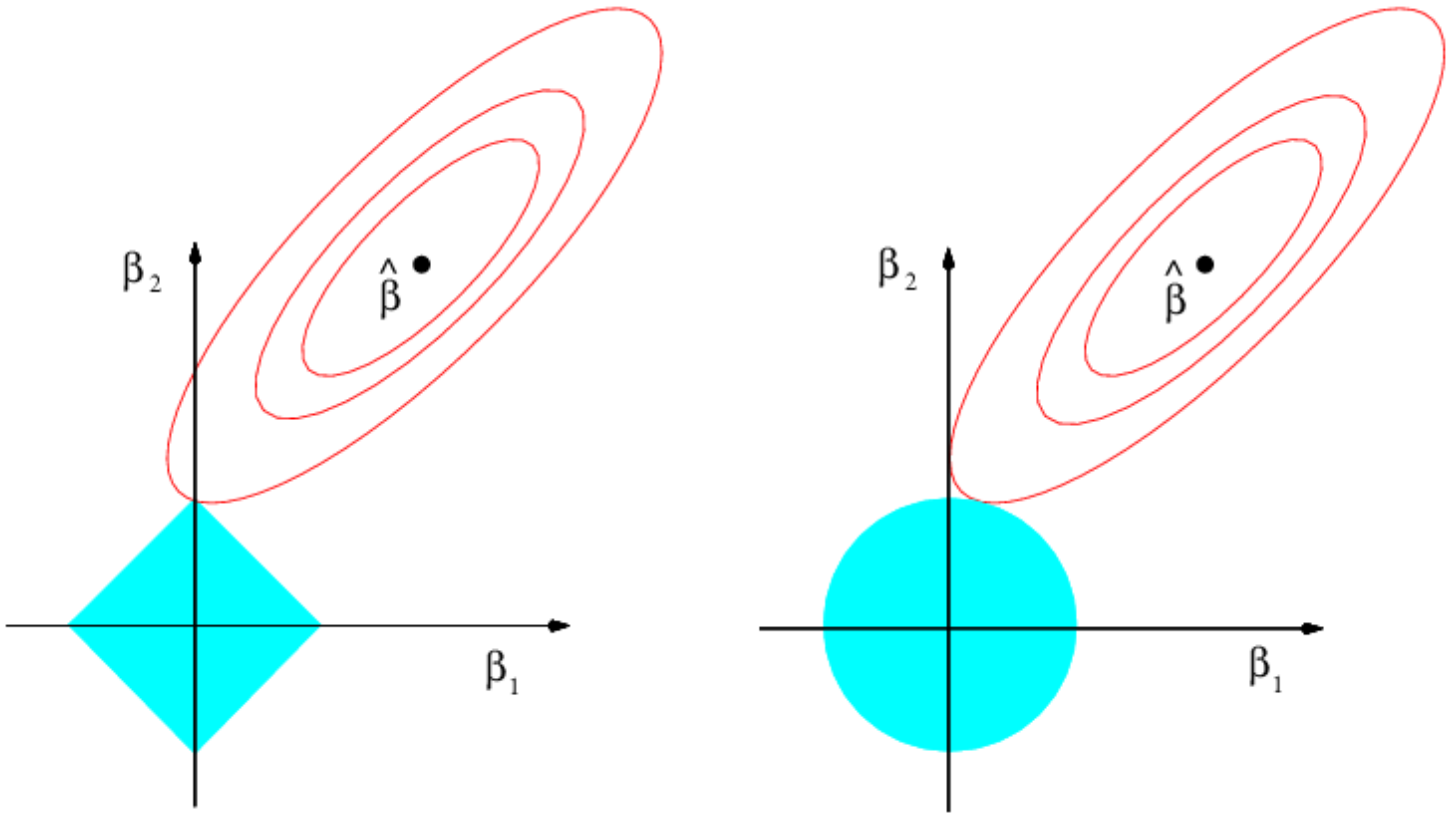
Both lasso and ridge regression minimise the same objective function

$$\underset{\alpha,\beta}{\textit{argmin}} = \underbrace{RSS(\alpha, \beta)}_{\text{Residual Sum of Squares}}$$

but with respect to different constraints:

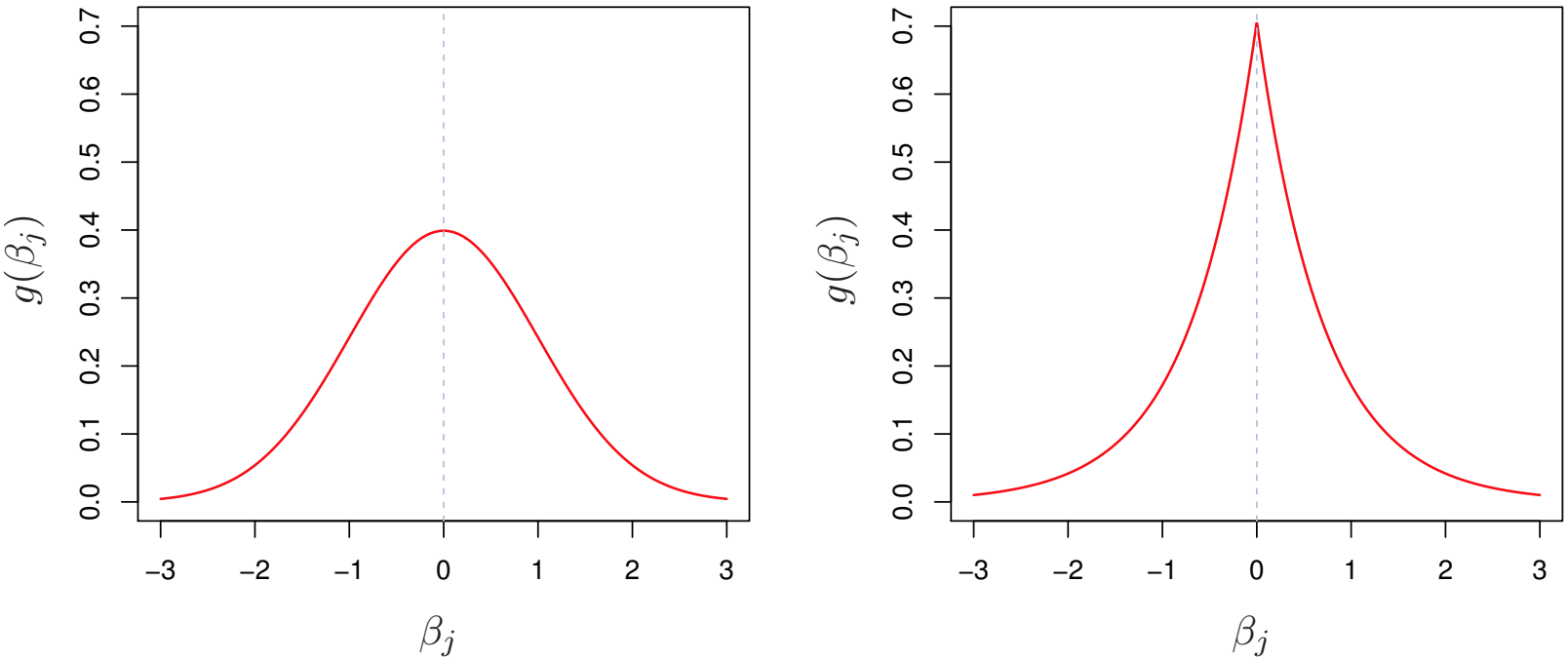
- Ridge regression: Subject to $\|\beta\|_2^2 \leq t$ (L2 norm)
- Lasso regression: Subject to $\|\beta\|_1 \leq t$ (L1 norm)

Ridge and lasso: Geometric interpretation



- Lasso regression: Subject to $\|\beta\|_1 \leq t$ (L1 norm)
  Constraint region defined by the L1 norm is a diamond.
- Ridge regression: Subject to $\|\beta\|_2^2 \leq t$ (L2 norm)
  Constraint region defined by the L2 norm is a circle.

Ridge and lasso: Bayesian interpretation



- Left: Ridge regression is the posterior mode for β under a Gaussian prior.
- Right: Lasso regression is the posterior mode for β under a double-exponential prior.

Elastic net regression

$$\underset{\hat{\alpha},\hat{\beta}_{\text{Elastic net}}}{\textit{argmin}} = \underbrace{RSS(\alpha,\beta)}_{\text{Residual Sum of Squares}} \underbrace{+\lambda_1 \sum_{j=1}^{p} |\beta_j| + \lambda_2 \sum_{j=1}^{p} \beta_j^2}_{\text{penalty}}$$

Interpretation:

- The Elastic net regression coefficient $\hat{\beta}_{\text{Elastic net}}$ is a biased estimate, but has a reduced variance compared to $\hat{\beta}_{OLS}$.
- There is an intrinsic model selection in Lasso regression, as it sets certain variables exactly to $\hat{\beta}_{\text{Elastic net}} = 0$, and thus excludes them from the model.
- When two variables are highly correlated, Elastic net includes both (Grouping property).

How to tune the regularisation parameter?

$$\underset{\alpha,\beta}{argmin} = RSS(\alpha, \beta) + \lambda f(\beta)$$

λ is the regularisation parameter

- $\lambda = 0$: No regularisation
- Small λ: Minimal regularisation
- Large λ: Strong regularisation
- How to choose the optimal λ?
  $\rightarrow$ Cross-validation (Lecture 3c)

Prediction using penalised regression

- Regularised regression is an ideal tool for prediction.
- We can define a prediction rule $\hat{f}(x)$ using the penalised regression coefficients

$$\hat{y} = \hat{f}(x) = \alpha + x\hat{\beta}_{\text{Penalised}}$$

  where $\hat{\beta}_{\text{Penalised}}$ is the regularised regression coefficient.
- Since Lasso and Elastic net force some $\hat{\beta}_{\text{Penalised}}$ to zero, variables with $\hat{\beta}_{\text{Penalised}} = 0$ are excluded from the model and do not contribute to the prediction rule.
- In contrast in Ridge regression all variables contribute to $\hat{f}(x)$.

Penalised regression in R: glmnet()

```
glmnet(x, y, family, alpha,nlambda = 100,
lambda.min.ratio = ifelse(nobs<nvars,0.01,0.0001),
lambda=NULL, standardize = TRUE, intercept=TRUE)
```

Input

- y: Outcome or response
- x: Predictors, formated `as.matrix(x)`

Generalised linear models included

- Linear regression: family = 'gaussian'
- Logistic regression: family = 'binomial'
- Count regression: family = 'poisson'
- Categorical outcome: family = 'multinomial'
- Survival model: family = 'cox'
- Multivariate linear model: family = 'mgaussian'

Penalised regression in R: glmnet()

```
glmnet(x, y, family, alpha, nlambda = 100,
lambda.min.ratio = ifelse(nobs<nvars,0.01,0.0001),
lambda=NULL, standardize = TRUE, intercept=TRUE)
```

Penalised regression models:

- Ridge regression: alpha = 0
- Lasso regression: alpha = 1
- Elastic net: 0<alpha<1

Regularisation parameter:

- Specify lambda for a pre-defined regularisation parameter
- Recommended: Perform cross-validation
  - nlambda: Grid length
  - lambda.min.ratio = ifelse(nobs<nvars,0.01,0.0001)

Penalised regression in R: glmnet

```
glmnet.out = glmnet(x, y, family, alpha)
```

Values:

- Intercept: glmnet.out$a0
- Regression coefficient estimates: glmnet.out$beta
- Regularisation parameters used: glmnet.out$lambda

Functions:

- Cross-validation: `cv.glmnet()`
- Regression coefficients: `coef(glmnet.out)`
- Prediction: `predict(glmnet.out, newx)`

Penalised regression in R: glmnet

For more details on glmnet, see the useful vignette: `http://web.stanford.edu/~hastie/glmnet/glmnet_alpha.html`

Other packages in R

- lm.ridge in the MASS package
- lars in the lars package
- penalized in the penalized package

Example: Diabetes data

- y: quantitative measure of disease progression one year after baseline (vector)
- x: predictor matrix
  - clinical parameters: age, sex, bmi
  - map: blood pressure
  - tc: total cholesterol
  - ldl: low-density lipoprotein
  - hdl: high-density lipoprotein
  - tch: total cholesterol over hdl
  - ltg: triglycerides
  - glu: glucose
- $n = 442$: sample size

Ridge regression and diabetes data

1. `lm(y~x)`
2. `glmnet(x,y,family="gaussian",alpha=0,lambda=0.1)`
3. `glmnet(x,y,family="gaussian",alpha=0,lambda=1)`

	lm	ridge01	ridge1
(Intercept)	152.13348	152.133484	152.133484
age	-10.01220	-9.358965	-6.698824
sex	-239.81909	-238.769510	-233.325125
bmi	519.83979	520.713588	520.019917
map	324.39043	323.548886	319.639206
tc	-792.18416	-666.737488	-320.594626
ldl	476.74584	377.500940	103.343333
hdl	101.04457	45.579036	-104.230542
tch	177.06418	161.855769	124.122091
ltg	751.27932	703.885551	568.507179
glu	67.62539	68.277472	71.865726

Lasso regression and diabetes data

1. `lm(y~x)`
2. `glmnet(x,y,family="gaussian",alpha=1,lambda=0.1)`
3. `glmnet(x,y,family="gaussian",alpha=1,lambda=40)`

	lm	lasso01	lasso40
(Intercept)	152.13348	152.133484	152.13348
age	-10.01220	-5.789635	.
sex	-239.81909	-234.457334	.
bmi	519.83979	522.819506	93.58588
map	324.39043	320.347881	.
tc	-792.18416	-534.397332	.
ldl	476.74584	271.305848	.
hdl	101.04457	-9.067565	.
tch	177.06418	146.255119	.
ltg	751.27932	655.715819	33.43273
glu	67.62539	66.410644	.

Elastic regression and diabetes data

1. `lm(y~x)`
2. `glmnet(x,y,family="gaussian",alpha=0.5,lambda=0.1)`
3. `glmnet(x,y,family="gaussian",alpha=0.5,lambda=40)`

	lm	enet01	enet40
(Intercept)	152.13348	152.133484	152.13348
age	-10.01220	-7.373073	.
sex	-239.81909	-236.908421	.
bmi	519.83979	521.524719	308.42812
map	324.39043	321.784878	53.18902
tc	-792.18416	-570.166854	.
ldl	476.74584	302.943444	.
hdl	101.04457	.	.
tch	177.06418	144.752485	.
ltg	751.27932	669.554762	267.61977
glu	67.62539	67.483126	.

Example: Breast cancer data

- y: benign or aggressive tumour (binary)

Benign	Aggressive	Total
185	121	306

- x: gene expression of $p = 22,283$ genes
- $n = 306$: sample size
- Truly big data $n << p$
- Data taken from Hatzis et al 2011 `https://jamanetwork.com/journals/jama/fullarticle/899864`

Breast cancer data and glm

1. `glm(severity~as.matrix(x), family='binomial')`

```
> glm.out = glm(severity~as.matrix(x), family="binomial")
glm.out$converged
```

Error: vector memory exhausted (limit reached?)
In addition: Warning message:
glm.fit: algorithm did not converge

```
> glm.out$converged
[1] FALSE
```

Breast cancer data and lasso

```
> lasso.out00015 = glmnet(as.matrix(x),y=severity,family="binomial"
,alpha=1,lambda=0.0015)
> sum(abs(coef(lasso.out00015))>0)
[1] 242
>
> lasso.out012 = glmnet(as.matrix(x),y=severity,family="binomial",a
lpha=1,lambda=0.12)
> sum(abs(coef(lasso.out012))>0)
[1] 4
>
> lasso.out012$a0
        s0
-0.229678
> summary(lasso.out012$beta)
22283 x 1 sparse Matrix of class "dgCMatrix", with 3 entries
      i j             x
1   411 1 -0.024275637
2  5307 1 -0.001092753
3 18933 1  0.008228413
```

Breast cancer data and elastic net

```
> enet.out003 = glmnet(as.matrix(x),y=severity,family="binomial"
,alpha=0.5,lambda=0.003)
> sum(abs(coef(enet.out003))>0)
[1] 442
>
> enet.out024 = glmnet(as.matrix(x),y=severity,family="binomial"
,alpha=0.5,lambda=0.24)
> sum(abs(coef(enet.out024))>0)
[1] 5
>
> enet.out024$a0
         s0
-0.2758569
> summary(enet.out024$beta)
22283 x 1 sparse Matrix of class "dgCMatrix", with 4 entries
      i j             x
1   411 1 -0.015083256
2   904 1 -0.002995031
3  5307 1 -0.001209533
4 18933 1  0.006024934
```

Take away: Penalised regression models

- Regularised regression approaches minimise the residual sum of squares and an additional penalty function.
- Different penalties imply different approaches
  - Ridge regression: $L2$
  - Lasso regression: $L1$
  - Elastic net regression: $L1 + L2$
- Penalised regression approaches are biased, but reduce the variance of the estimate and the prediction rule.
- Lasso and Elastic net perform an intrinsic model selection.
- The regularisation parameter λ can be choosen using cross-validation.

Further reading:

- An Introduction to Statistical Learning: Chapter 6 Linear Model Selection and Regularization `http://www-bcf.usc.edu/~gareth/ISL/index.html`
- The epigenetic clock: 'A multi-tissue full lifespan epigenetic clock for mice' `https://www.ncbi.nlm.nih.gov/pmc/articles/PMC6224226/` *Using DNA methylation data from previous publications with data collected in house for a total 1189 samples spanning 193,651 CpG sites, we developed 4 novel epigenetic clocks by choosing different regression models (elastic net- versus ridge-regression) and by considering different sets of CpGs (all CpGs vs highly conserved CpGs).*