### Course 02402 Introduction to Statistics Lecture 9:

# Multiple linear regression

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- Warm up with some simple linear regression
- Multiple linear regression
- Model selection
- Residual analysis (model validation)
- Curvilinearity
- Confidence and prediction intervals
- Colinearity
- The overall regression method

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# Example: Ozon concentration

We have a set of observations of: logarithm to ozone concentration (log(ppm)), temperature, radiation and wind speed:

| ozone | radiation | wind | temperature | month | day |
|-------|-----------|------|-------------|-------|-----|
| 41    | 190       | 7.4  | 67          | 5     | 1   |
| 36    | 118       | 8.0  | 72          | 5     | 2   |
| :     | :         | :    | •           | :     | :   |
| 18    | 131       | 8.0  | 76          | 9     | 29  |
| 20    | 223       | 11.5 | 68          | 9     | 30  |

## Example: Ozone concentration

```
## Se info about data
?airquality
## Copy the data
Air <- airquality
## Remove rows with at least one NA value
Air <- na.omit(Air)
Air <- Air [-which(Air$0zone == 1), ]
## Check the empirical density
hist(Air$Ozone, probability=TRUE, xlab="Ozon", main="")
## Concentrations are positive and very skewed, let's
## log-transform right away:
## (although really one could wait and check residuals from models)
Air$logOzone <- log(Air$Ozone)
## Bedre epdf?
hist(Air$logOzone, probability=TRUE, xlab="log Ozone", main="")
## Make a time variable (R timeclass, se ?POSIXct)
Air$t <- ISOdate(1973, Air$Month, Air$Day)
## Keep only some of the columns
Air \leftarrow Air[,c(7,4,3,2,8)]
## New names of the columns
names(Air) <- c("logOzone", "temperature", "wind", "radiation", "t")</pre>
## What's in Air?
str(Air)
Air
head(Air)
tail(Air)
## Tupically one would begin with a pairs plot
pairs(Air, panel = panel.smooth, main = "airquality data")
```

# Example: Ozone concentration

- Let us first analyse the relation between ozone and temperature
- Apply a simple linear regressions model

$$Y_i = \beta_0 + \beta_1 x_i + \varepsilon_i$$
 ,  $\varepsilon_i \sim N(0, \sigma^2)$  og i.i.d.

#### where

- $Y_i$  is the (logarithm of) ozone concentration of observation i
- $x_i$  is the temperature at observation i

#### Fit the model in R

## Simple linear regression model for the other two

We can also make a simple linear regression model with each of the other two independent variables

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# Multiple linear regression

- Y is the dependent variable
- We are interested in modelling the Y's dependency of the *independent* or *explanatory* variables  $x_1, x_2, ..., x_p$
- We are modelling a *linear relation* between Y and  $x_1, x_2, ..., x_p$ , described with the regression model

$$Y_i = \beta_0 + \beta_1 x_{1,i} + \dots + \beta_p x_{p,i} + \varepsilon_i$$
 ,  $\varepsilon_i \sim N(0, \sigma^2)$  and i.i

•  $Y_i$  og  $\varepsilon_i$  are random variables and  $x_{i,i}$  are variables

# Least squares estimates

The coefficient estimates are found by minimizing:

$$RSS(\beta_0, \beta_1, \dots, \beta_p) = \sum_{i=1}^n \left[ y_i - (\beta_0 + \beta_1 x_{i,1} + \dots + \beta_p x_{i,p}) \right]^2$$

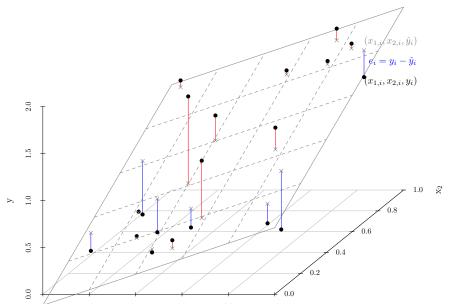
The "predicted" (= "fitted") are found as

$$\hat{y}_i = \hat{\beta}_0 + \hat{\beta}_1 x_{i,1} + \dots + \hat{\beta}_p x_{i,p}$$

And then the residuals are found as

$$e_i = y_i - \hat{y}_i$$
  
residual = observation – prediction

# Least squares estimates - The concept!



• Remark 6.6: Extract  $\hat{\beta}_i$  and  $\hat{\sigma}_{\beta_i}$  from R-output (summary(myfit))

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- Everything: THE SAME as for SIMPLE linear regression!

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- Everything: THE SAME as for SIMPLE linear regression!
- (In Section 6.6: Mathematical matrix based expressions including explicit formulas. Not syllabus in course 02402)

What dose  $\hat{\beta}_i$  express?

• The expected y-change with 1 unit  $x_i$ -change

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- The expected y-change with 1 unit  $x_i$ -change
- The effect of  $x_i$  given the other variables
- The effect of  $x_i$  corrected for the other variables
- The effect of  $x_i$  "other variables being equal"
- The unique effect of  $x_i$

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- Depends on what else is in the model!!

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- The effect of  $x_i$  "other variables being equal"
- The unique effect of  $x_i$
- Depends on what else is in the model!!
- Generally: NOT a causal/intervention effect!!

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# Extend the model (forward selection)

- Not included in the eNote
- Start with the linear regression model with the most significant independent variable
- Extend the model with the remaining independent variables (inputs) one at a time
- Stop when there is not any significant extensions possible

# Reduce the model (model reduction or backward selection)

- Described in the eNote, section 6.5
- Start with the full model
- Remove the most insignificant independent variable
- Stop when all prm. estimates are significant

### Model selection

- There is no fully certain method for finding the best model!
- It will require subjective decisions to select a model
- Different procedures: either forward or backward selection (or both), depends on the circumstances
- Statistical measures and tests to compare model fits
- In this course only backward selection is described

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# Residual analysis (model validation)

- Model validation: Analyze the residuals to check that the assumptions is met
- $e_i \sim N(0, \sigma^2)$  is independent and identically distributed (i.i.d.)
- Same as for the simple linear regression model

# Assumption of normal distributed residuals

 Make a qq-normalplot (normal score plot) to see if they seem normal distributed

### Assumption of identical distribution of residuals

• Plot the residuals  $(e_i)$  versus the predicted (fitted) values  $(\hat{y}_i)$ 

- Seems like the model kan be improved!
- Plot the residuals vs. the independent variables

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### Curvilinear model

If we want to estimate a model of the type

$$Y_i = \beta_0 + \beta_1 x_i + \beta_2 x_i^2 + \varepsilon_i$$

we can use a multiple linear regression model

$$Y_i = \beta_0 + \beta_1 x_{i,1} + \beta_2 x_{i,2} + \varepsilon_i$$

where

- $x_{i.1} = x_i$
- $x_{i,2} = x_i^2$

and apply the same methods as for multiple linear regression.

# Extend the ozone model with appropriate curvilinear regression

```
Air$windSq <- Air$wind^2
fitWindSq <- lm(logOzone ~ temperature + wind + windSq + radiation, data=Air)
summarv(fitWindSq)
## Equivalently for the temperature
Air$temperature2 <- Air$temperature^2
## Add it
fitTemperatureSq <- lm(logOzone ~ temperature + temperature2 + wind + radiation, data=Air)
summary(fitTemperatureSq)
Air$radiation2 <- Air$radiation^2
fitRadiationSq <- lm(logOzone ~ temperature + wind + radiation + radiation2, data=Air)
summary(fitRadiationSq)
## Which one was best?
fitWindSqTemperaturSq <- lm(logOzone ~ temperature + temperature2 + wind + windSq + radiation, data=Air)
summary(fitWindSqTemperaturSq)
## Model validation
qqnorm(fitWindSq$residuals)
ggline(fitWindSg$residuals)
plot(fitWindSq$residuals, fitWindSq$fitted.values, pch=19)
```

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# Confidence and prediction intervals for the plane, Method 6.9:

Extract Confidence and prediction intervals for the plane by R-function predict. Options for confidence og prediction exist.

```
## Confidence and prediction intervals for the curvilinear model
## Generate a new data. frame with constant temperature and radiation, but with varying wind speed
wind < -seq(1,20,3,bv=0.1)
AirForPred <- data.frame(temperature=mean(Air$temperature), wind=wind.
                         windSq=wind^2, radiation=mean(Air$radiation))
## Calculate confidence and prediction intervals (actually bands)
CI <- predict(fitWindSq, newdata=AirForPred, interval="confidence", level=0.95)
PI <- predict(fitWindSq, newdata=AirForPred, interval="prediction", level=0.95)
## Plot them
plot(wind, CI[, "fit"], vlim=range(CI,PI), type="l",
     main=paste("At temperature =".format(mean(Air$temperature).digits=3).
                "and radiation =", format(mean(Air$radiation),digits=3)))
lines(wind, CI[,"lwr"], lty=2, col=2)
lines(wind, CI[,"upr"], lty=2, col=2)
lines(wind, PI[,"lwr"], lty=2, col=3)
lines(wind, PI[, "upr"], lty=2, col=3)
## legend
legend("topright", c("Prediction","95% confidence band","95% prediction band"), lty=c(1,2,2), col=1:3)
```

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### Colinearity

- MLR breaks down if X-data has "exact linear redundancy"
  - Example: Both height in cm and height in m is in the data.

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  - Example: Both height in cm and height in m is in the data.
- Interpretation and model stability is challenged if X-data has "near redundancy" patterns
  - Example: Both weight and BMI are in the X-data (highly correlated)

# Colinearity

- MLR breaks down if X-data has "exact linear redundancy"
  - Example: Both height in cm and height in m is in the data.
- Interpretation and model stability is challenged if X-data has "near redundancy" patterns
  - Example: Both weight and BMI are in the X-data (highly correlated)
- With e.g. two highly correlated x-variables:
  - Together in the model for y none of them may have a unique effect
  - Separately they may have a strong effect each of them

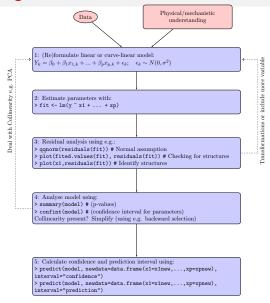
# Colinearity - an illustration in R

```
n <- 100
## First variable
x1 \leftarrow \sin(0:(n-1)/(n-1)*2*2*pi) + rnorm(n, 0, 0.1)
plot(x1, type="b")
## The second variable is the first plus a little noise
x2 <- x1 + rnorm(n, 0, 0, 1)
plot(x1,x2)
cor(x1,x2)
## Simulate an MLR
beta0=20; beta1=1; beta2=1; sigma=1
y <- beta0 + beta1 * x1 + beta2 * x2 + rnorm(n,0,sigma)
## See scatter plots for y vs. x1, and y vs. x2
par(mfrow=c(1,2))
plot(x1,y)
plot(x2,y)
## Fit an MLR
summary(lm(y ~ x1 + x2))
x1[1:(n/2)] \leftarrow 0
x2\lceil (n/2):n\rceil <- 0
plot(x1, type="b")
lines(x2, type="b", col="red")
cor(x1,x2)
y <- beta0 + beta1 * x1 + beta2 * x2 + rnorm(n,0,sigma)
## and fit MLR
summary(lm(y ~ x1 + x2))
```

It is important how experiments are designed!

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### The overall regression method box 6.16



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