

I python
 %time
 %timeit
 %ls
 %prun
 ?
 %rm
 %cd
 %mkdir

np.reshape (rows, columns)
 np.mean
 np.max, np.min
 np.std
 np.sum
 np.size, np.shape
 np [rows, columns] ~~shape~~
 np [first: last: step]
 np.indices
 np.transpose (array)
 np.random.shuffle
 np.invert
 np.zeros, np.ones
 np.count_nonzero (value)
 np.random.range

df.argmax
 df.index
 df.groupby()
 df.value_counts
 df.sort_values (ascending=, []
 df.plot (kind='hbar')
 df.drop
 df.dropna
 df.reset_index()

pd.crosstab (rows, columns)
 pd.pivot

100 man?
 10 zieh 100mal g, nicht
 90 nicht zieh

Series.values() = np.array

*df.index.weekday / month / day
 df.merge / df.join?

echt
 zieh nicht zieh
 Test zieh g
 nicht zieh 1 g

*str.contains
 *str.len
 *str.replace

1000
 echt
 zieh nicht zieh
 Test zieh g
 nicht zieh gg
 990-99 891

% ls magic

Shell met ! (pwd, ls, cd)

? access info

- * - wildcard options

dot $\sum_i (s_i \cdot S_i)$

• evaluation

Classification

→ accuracy

recall

$TP / (TP + FN)$

Precision

$TP / (TP + FP)$

RMSE = regression

NP → dir(np)

np.empty/zeros → array

np.shape/ndim/size

→ [i]
↓
[j]

np.info(np.ndarray.dtype)

math

aggregate

.exp → e^x

.sum

.min

.sqrt → \sqrt{x}

.std

.max

np.arange (start, stop, step)

[::-1] → backwards

[::2] → from 0 with step 2

a[2] → el. at 2nd index

a[1,2] → 2nd row, 3rd col

a[0:2] → items at 0 and 1

• boolean indexing

a[a < 2] → alles in a kleiner dan 1

• fancy

b[[1,0,1,2], [0,1,0,1]]

↳ [(1,0), (0,1), (1,0), (2,1)]

• transpose

np.T

np.transpose()[1]/[2] → add rows/cols

np.concatenate()

arr[i]=4 (maak 4 van index i)

np.add(arr1, arr2) → elementwise

(arr, 4) → 4 by elk element

pd → dir(pd) → help(^{pd.}Series, loc)
pd.read_csv('file.csv', header, rows)
df['a'] → get one element
df[:,2] → get subset
Position → df.iloc[0],[0]] + select
by label → df.loc[0],[country]]
by index → df.ix[2] → single row/sub
→ df.ix[:, ['Country']]

bool Select

S[S > 1] → S where S is not
S[(S < -1) | (S > 2)] → values under -1
& above 2

sort_index()

sort_values(by=...)

• Statistics

df.describe → Summary

df.mean → mean of all cols

df.corr → corr between cols

df.count → value count col

df.min/max/mean/median/std

• Cleaning

df.columns['a','b','c'] → rename cols

pd.isnull() check null, bool array

df.dropna() → drop row nan values

df.fillna(x) → fill nan with x

(s.mean()) → fill na with mean

s.replace([1,3], [one, three])

↳ replace all 1 → one 3 → three

S['col'].str.lower()

str.replace('vb', '7')

delete all met ...

S[S['col'].str.contains('...')]

• Filter, Sort, Group

df[df[col] > 1] → row where col val > 1

df[(df[col] > 1) & (df[col] < 2)]

df.sort_values(col, ascending)

df.groupby(col)

[col1], [col2]

df.pivot_table(index=col, values=col2,3, aggf)

→ group by col, aggfunc on 2,3

• Join / combine

df1.append(df2) → add row to end df1

.Concat(df1, df2, axis=1)

→ add cols to end df1

df.join(df2, on=col1, how=inner)

• View, inspect

df.head

df.tail

df.shape

df.info

df.value_counts

len(.unique) → unique len

vrg → len('Party') == 'pvdd'

- np slices and indices are views not copies so changing them changes original array, use `.copy()` for a copy.
- Broadcasting rules:
 1. If the two arrays differ in their number of dimension the shape of the one with fewer dimensions is padded with on its leading (left) side.
 2. If the shape of the two arrays does not match in a dimension, the array with shape equal to one in that dimension is stretched to match the other shape.
 3. If in any dimension the sizes disagree and neither is equal to 1, an error is raised.
- `x[::-1]` all elements, reversed
- Count amount of True entries in a Boolean array with `np.count_nonzero` (True = 1, False = 0)
- Quickly check any/all values are True with `np.any` / `np.all`
- Data science is: Based on patterns in data predicting a value or new unseen instances of data (classification/regression). Steps: interaction with outside world, preparation, transformation, modeling and computational presentation. Data Science is the study of the generalizable extraction of knowledge from data.

$$F_1 = 2 \times \frac{\text{precision} \cdot \text{recall}}{\text{precision} + \text{recall}}$$

$$\text{Precision} = \frac{TP}{TP+FP} \quad \text{Recall} = \frac{TP}{TP+FN}$$
- `np.nditer` to iterate through numpy arrays
 - `.value_counts()` `.index.values` `.str` (`['1\w|s'`)
 - `x[~x.y.str.contains(" " , na=False)]`
 - `.value_counts().sort_index()` `x.loc[x['x'] == '1']`
 - `normalize=True * 100` voor percentages

P N

P

N

row column

X[1, 2]

X[2, 3] # two rows, three columns

X[3, :2] # all rows, every other column

X[1, :-1] reverse

its divisible by 3
X[X % 3 == 0]

X[:, 0] first column

X[0] first row

incorporate
(X-X_{mean})/X_{std}(head)
= (X-X_{mean})/X_{std}(X_{mean})

masking
df[df > 0, 3] (cdf[k, g, 8])
X.std() std() X.std()

df.loc = explicit indexing

df.iloc = python style indexing

sort rows by height than width

res.sort_values(['height', 'width', ascending=False, True])
max length, sep spaces

res.groupby('species')['length'].max()

X.sort() of X.argsort
sort indices

K nearest 2:

k=2 z=np.argsort(X[:, k+1:axis-1])
for i in range(X.shape[0]):
for j in z[i:k+1]:

plt.plot(z[0:k+1] * X[:, k+1], color='black')

def shape (n=len(L), y):

return L.reshape(np.ma.count(L), y)
all others much less occupy 2d array
CS = pd.Series(C).sort_values(ascending=False)
C = Counter(text.split())
CS[CS*CS.index.str.len() == 20].index

df.columns [df.columns.str.contains('row')] / df.columns.str.contains('high')]

df.describe()
df['party'].value_counts()
df['party'].value_counts(normalize=True)
df['party'].value_counts(normalize=True).plot(kind='bar')

df['km2'] = np.sqrt(df['area'])
df['km2'].value_counts().plot(kind='bar')

df.index.str.lower().str.strip().str.replace(" ", "").sort_values()

df.groupby('area').count().sort_index()

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df.groupby('area').count().sort_index()

df.groupby('area').count().sort_index()

help(len) of len?

df.groupby('area').count().sort_index()

Data Science cheat sheet

NP

% !smagic = all magic commands

X [2, -3, 1] = last 3 columns first 2 rows | X + np.arange(len(X))

X.reshape(12, 1) = make one dimensional | X [0, 1, 2, 3] CO [1, 2, 3, 3] - pang index

(X - X.mean()) * X² sqr dif. each elem from mean of X

np.sqrt((X - X.mean())²).mean() = X.std() = ~~variance~~

X [X % 3 == 0] = apply bool. mask

X² [0, 1, 2, 3] CO [1, 2, 3] diagonal (pang index)

big-arr = np.random.randint(10^x, 10^{x+2}, size=10^{x+2})

% timeit sum(X² per x in big-arr)

% timeit eig-arr

X [:, -1] = last column 2D array | X [X % n == 0] | X.mean(axis=0)

pd

dubbel = df.index.value_counts()

dubbel [dubbel > 1]

df.sort_index()

df.index.str.lower(), str.strip(), str.replace(' ', ''), str.replace('...', 'na')

df [~ df.index.str.lower().str.contains('string')].head(10)

df.describe

df.inkomen.sort_values(plots)

df [df.inkomen > 50000].gemeente.value_counts().plot(kind='bar')

kind = 'bar')

= gemeentes met scholen met inkomen > 50.000, gesorteerd op hoeveelheid in

for. bar plot

top-minis = kur [kurc 'party'] = 'pppp', Ministerie.value_counts()

top-minis [top-minis > 5].plot(kind='bar')

df.acepa(subset = ['party'])

deelvragen = kur.vraag.str.count('!')

no-outliers = deelvragen [deelvragen <= 50]

no-outliers.plot(kind='hist')

np.mean (no-outl)

median = statistics.median (no-outl)

modus = Counter (no-outliers).most-common(1) CO [50]

corr = no-outl.corr (kur.vraag.str.len())

corr = no-outl.corr (kur.vraag.str.len())

df.sort_values(['sepal-len', 'sepal-wid', 'petal-len', 'sepal-len', 'sepal-wid', 'petal-len'])

df.groupby('species')['sepal-len'].max()

df.query('sepal-len > 5 and sepal-wid < np.sqrt(5)')

C = Counter(text.split())

CS = pd.Series(C).sort_values(asc=False)

CSCS = CS.index.str.len() == 2117.index

```

import numpy
import matplotlib.pyplot as plt
import random
array
os
math
import matplotlib inline

```

```

np.random.seed(42)
randint(10, size=6) 1d
randint(10, size=(3,4)) 2d
randint(10, size=(4,4,6)) 3d
np.concatenate([X, Y])
np.vstack
np.hstack
np.split([3,5]) = tot 3/4 5/van 6
np.vsplit([X, [3]]) vanaf rij 3

```

indexing: $x[0]$ / $x[0:2]$ / $x[2:-1]$ / $x[-1]$ / $x[0, 2]$

reshaping: $X[np.newaxis, :]$ = row vector
 $X[:, np.newaxis]$ = column vector

computation: $2^x - x/2 - x//2 - np.log - np.log 2 - np.log 10(x)$

magic: %ls - the directory
 %row = remove
 %*% = double line

```

x = np.arange(1,5)
np.add.reduce(x) = returns it on all elements
np.multiply.outer(x)
np.linspace(start, end, spacing)
np.arange(start, end, step)
a = np.random.random(42)
a.vandint(10, size=(3,4)) -> a < 6 - a > 6 -
np.sort(x, axis=0/1)
np.argsort(x) = values of position
np.partition(x, 3) = 3 smallest values + remaining

```

Broadcasting: $val_i = padded\ to\ left$
 2 = 1 dimension array stretched
 3 = if dimension disagree = error

$m + A[:, np.newaxis].shape = right\ padding$

access simple = $x[0]$
 slicing = $x[0:5]$
 masking = $x[x > 5]$
 fancy indexing = $[x[3], x[5], x[6]]$
 fancy + simple = $[x[2, 0, 1]]$
 fancy + fancy = $[x[2, 1, 3, 2]]$
 fancy + masking = $mask = np.array([0, 1, 0, 1])$
 $x[rows[:, np.newaxis], mask]$

Theorie:
 Precision = # of predictions for class C covered
 Recall = # instances of C predicted to be C -> $TP / (TP + FN)$
 accuracy = # of all predictions being correct -> $TP / (TP + FP)$

accuracy = 95%	n = 10000	TP	FP
prec +	pos neg	590	940
neg +	5 995	910	10000

precision $95 / (95 + 95) = 0,16$

Pandas - good for data analysis and modeling without use of R

```

df = pd.read_csv('...'), names = ['kolom', 'human', 'index']
df = pd.read_html('...')[1]
%ls - ./Data/*.csv
%cat ./Data/foe.csv -> df = pd.read_csv('.. /Data/foe.csv')

```

- plot samples in dataset = iris.species, value = counts(). plot(kind='bar');
- create dataframe numerical data = x and y = X, y = iris.drop('species', axis=1), iris.species
- z-normalize by subtracting mean by std = $(X - X.mean()) / X.std()$. head(1), $(X - X.mean()) / X.std().std()$
- Sort iris by column and row inverse = iris.sort_values(['...'], ascending=[False, True])
- maximal lengths for each species = iris.groupby('species')['length'].max()
- larger than 5 and smaller than sqrt of 5 = a = np.sqrt(5) - iris.query('x > 5 & y < @a')
- tokenize the text by split, count taken, show head = C = Counter(text.split())
- Find all tokens which have >= 2 = CS = pd.Series(C).sort_values(ascending=False)
- what percentage of all unique words occur once = $CS[CS == 1].sum() / CS.count()$, $(CS == 1).mean$, $CS[CS == 1].sum() / CS.sum$
- make histogram which indicates for each number i - occur how many tokens in text occur i times
 pd.Series(Counter(CS.values)).plot(kind='bar'); CS.value_counts().sort_index().plot(kind='bar')

```

data = pd.read_html('...')[1], data.groupby('...').count().sort_values().tail(1)
data = pd.Series([1, 2, 3]), index=['a', 'b', 'c'], dict = {'ams': 100, 'utc': 50}
pop = pd.Series(dict)
pd.Series(5, index=[3, 6, 9])
pd.Series({'a': 1, 'b': 3}) -> index = [3, 1]
states = pd.DataFrame({'pop': pop, 'area': area})
states
indexing data['b'], 'a' in data, list(data.items())
slicing data['a': 'b'], data[0:3], data[(data > 0) & (data < 3)], data[['d', 'b']]
data.values = alle waarden
data.T = transpose
A.add(B, fill_value=0)

```

Series
 Pandas

Series: $x.size$, $x.dtype$ = similarity check
 $x.shape$, $x1 \times x2$ = add without same/union
 $x.ndim$, $x1 \wedge x2$ = symmetric difference

data.Loc [data.x > 100, ['pop', 'area']] = masking + fancy

slicing in 2d = $pop[('ams', 2010) : ('utc', 2000)]$

hard = select all values from 2010 -> $pop[[i for i in pop.index if i[1] == 2010]]$ -> pd.MultiIndex.from_tuples(index)

pd.concat([ser1, ser2], axis=0/1)

categories of join = 1-to-1
 many-to-one
 many-to-many
 need key column
 choose by pd.merge(x, y, on='...')
 (x, y, left_on='...', right_on='...')

pd.merge(x, y, how='outer', 'inner', 'left', 'right')

Lees de vragen goed! Kijk of je iets moet plotten!

iris = ... heeft niet verzameling set() a.tab → kkl

NUMPY

1D array: `x2.reshape((12,7))`

`x1.std()` = $\text{np.sqrt}(\frac{\sum (x1 - \text{mean}(x1))^2}{n})$
Squared difference
variance

diagonaal: `x2[[0,1,2], [0,1,2]]`

`%timeit sum([x**2 for x in bigarray])`

`%timeit sum(bigarray**2)`

- sum of squares in an array -

homogeen vs heterogeen. size = elementen ndim

PANDAS

`index_col = 'kolomnaam'` die je als index wilt

• `unique()` of `set()`

`str.strip()` haakt whitespace aan begin/ende weg

`str.contains('str')`

`df[naam]`

of `df.describe()` → statistiek data

→ in elke waarde het gem. uitre waarde in die kolom afkruken (normaaliseren)

`mean = cito.mean()`

`cito[mean.index] = mean`

`cito.gemeente[cito.inkomen > 50000].value_counts()`

• `sort_values()`, `plot(kind='bar')`

→ in welke gem. staan scholen waar inkomen meer dan 50000
`plot` aantal scholen per gemeente

• `drop(columns='naam')` of `(naam, axis=1)`

→ Z-normalize data by subtracting mean of column from each cell and divide by standard dev of each column

$(X - \text{mean}(X)) / \text{std}(X)$, `head()`

→ also show std of each col of the Z-normalized data • `std()` → moet gelijk zijn aan 1

→ Sort iris first by S-L & then by S-W, sort in reverse order (multindex)

`iris.sort_values(['S-L', 'S-W'], ascending=[False, True])`

→ max S-L for each species

`iris.groupby('species').sepal.length.max()`

→ Find all tokens in text that occupy 24 char in total

`CS[CS.index.str.len() * CS.values == 24].index`
(`CS = pd.Series(Counter(text.split()))`)

→ What % of all unique words in text occurs once

$\frac{CS[CS==1].sum()}{CS.count()}$
of $\text{len}(CS.index.unique()[CS.values==1]) / \text{len}(CS.index.unique())$

→ What % of all words in text occurrence

$\frac{CS[CS==1].sum()}{CS.sum()}$
of $\text{len}(CS.index.unique()[CS.values==1]) / \text{len}(text.split())$

→ Make a histogram (bar plot) which indicates for each number of occurrences how many tokens occur i times

`CS.value_counts().plot(kind='bar')`

→ land met meeste bendidagleden + aantal leden

`bdEO.value_counts().head(1)`

of `bd.groupby(0)[1].count().sort_values().tail(1)`

of `winnar` → `bd.groupby(0)[1].count()`

`winnar.argmax()`, `winnar.max()`

→ welke partij heeft grootste aantal leden 1 deelstaat

`bd.groupby(0)[2].value_counts().sort_values().tail(1)`

IPYTHON

`%ls files in directory` } %magic
`%cp file1 file2` } %magic
`%rm no-good.txt`

plod print work - array
cd naam / change dir
mkdir naam - make dir
mv .. / .. / move one dir up

`inspect resources: %timeit`
• `berschil` → %time ? ! %time?

`datatype of var. X` → `x?` inspect 1st line of output

`get documentation: help(...)`

`print(...)`

Supress output;

% debug interactief

% mode hoe exceptions zien

THEORY

precision: $\frac{TP}{TP+FP}$ accuracy: $\frac{TP+TN}{total}$ recall: $\frac{TP}{TP+FN}$

	T	F	
T	TP	FP	5%
F	FN	FN	
	5	945	1000

accuracy 95%

ziekte 1/100

precision = $(0.01 \times 0.95) / ((0.01 \times 0.95) + (0.99 \times 0.05)) = 0.16$

xml/json: cel + lijst (als er niet evenveel kolommen zijn → spreadsheet)

teken spreadsheet:

- insert / delete / update

- andere view (string / ding)

- combineren v. sheets

- aggregate / correlate

dot prod: `A.dot(B)`

$\sum (a[i] * b[i])$ for i in range(len(a))

no births p/year p/gender

`names.groupby(['year', 'sex'])['births'].sum().unstack()`

numpy hw wk2

• create array (n,0):
`return np.arange(0, n*0, 0)` eerste n getallen
deelbaar door 0

• add-per-column(L)
for i in range(L.shape[1]):
`L[:,i] = L[:,i] + i` tel i op voor item
in ide kolom

return L

• add-per-row(L)
`tp = np.transpose(L)`
return `add-per-column(tp).transpose()`

• mean-per-column(L)
return `np.mean(L, axis=0)`

alles reversed: `x[::-1, ::-1]`

1D → 2D `x.reshape((1,3))` of `x[0].newaxis`

`x1, x2 = np.split(x, [3,5])` → `[0:3]` `[3:5]` `[5:]`

`np.sum of L.sum()`

broadcasting - ele regels verbreden het om x1 naar de
shape van x2 te krijgen.

how many vals less than zero: `np.sum(x < 0)`

in each row: `np.sum(x < 6, axis=1)`

`np.any()` `np.all()`

combined indexing: `x[2, [2, 0, 1]] = x[2,2], x[2,0], x[2,1]`

get name from last row `data[-1]['name']`

`data.loc` expliciet → indices

`data.iloc` impliciet

`data.ix` hybride vorm

`data.T` transpose

→ zet dat ik heb
heb ik het ook

Numpy

- 1 Last 3 columns of first 2 rows: $x[2:-3:]$
- 2 Change each element in x_1 by its squared difference from the mean of x_1 .
 $(x_1 - x_1.mean())**2 = \text{variance}$
- 3 standard deviation = ~~np~~ sqrt of variance ($np.sqrt(\text{variance}) == x_1.std()$)
- 4 1D en 2D arrays vanwege broadcast regels niet samen te voegen, 1D kan niet naar 2D
- 5 boolean mask to x_1 and reduce x to ints divisible by 3: $x_1[x_1 \% 3 == 0]$
- 6 Fancy indexing for diagonal of x_2
 $x_2[[0,1,2], [0,1,2]]$
 TP / TP + FP
 TN / TN + FN

JPython

% ls -lh a = working directory ~~size~~ b = size
 % rm = remove
 % time, timeit, prun

Pandas

% cp = copy previous, new
 pd.read_excel(" ", index_col = " ")
 dimensies = .shape
 test voordubbelen = len(set(" ")) = len(" ")
 cito.index.value_counts() telt hoeraak index elk item in cito.index voorkomt
 >1 betekent dubbel (dubbel = cito.index.value_counts() | dubbel [dubbel > 1])
 sort_index()
 str.contains
 ~ = negation
 cito[M.index] - M.tail()
 M = cito.mean()
 cito[cito.inkomen > 50000].gemeente.value_counts() 7,2
 .sort_values() ascending = [6, 8
 .groupby .max() / .min()
 a = np.sqrt(5)
 iris.query('sepal_length > 7.5 and sepal_width < @a')
 C = Counter(text.split()) = text splitten en tellen 6,2 + 8,7c
 pd.Series(C)

10, 10: 10, 60, 10, 10 60 0,6 60 14,9 7,9 7,45
 10 70
 5,5 10 70/2 = 0,9

① import re
 import collections → (counter, most-common())
 import math → math.sqrt()

BASICS

random.random() → rand
 random.choice() → sel()
 [start: stop: step]
 L[-1:] → laatste rij
 L[:, -1] → laatste kolom

* $g^{\log(a)} = b \rightarrow g^b = a$
 $2^{\log(16)} = 4 \rightarrow 2^4 = 16$

Numpy

Pandas

.log2(), .info(),
 .shape(), .inplace(),
 .size(), .full(),
 .eye(),
 .subtract(),
 .multiply(),
 .sin(), .cos(), sqrt(),
 .corrcoef(),
 .transpose(),
 .mean(),
 .reshape(),
 .ravel(),
 .ones()

pd.read_csv()
 ↳ compression, skip initial space, sep, names= \mathbb{R} ,
 df.loc[" "].isin() of df[df[" "] == " "],
 .crosstab(L, R), .unique(),
 .dropna(subset=L) / .notnull(),
 df[df[" "], str.contains(" ")] , str.lower(),
 df.pivot_table(values, index, columns,
 aggfunc, margins=True),
 pivot.sort_values(L, ascending),
 df.column.sum(),
 df.column.index.values(),
 pivot[pivot[" "] >= 10.000] @ ((pivot["ratio"] >= 4,
 max_error = 1/15 * 100
 act = sum(error...) / sum(L"all") * 100%

df[L].grid, shape(),
 min(axis=) ↳ L[i-1% "], ...
 midden → L[int(len(L)/2)-2: "+2, "-2: "+2]

rel_error = hoogste error
 most_errors = meeste x post

accuracy = $\frac{TP + TN}{TP + TN + FP + FN}$

f_error = f.sum() + m_error * pivot.m.sum() / pivot.all.sum
 → = act_error

recall = $\frac{TP}{TP + FN}$ precision = $\frac{TP}{TP + FP}$

pd.head(), groupby(),
 .tail(),
 .plot()

plt.hist **PIT**
 plt.parch
 * plt.pcol([L], [R]), df.plot.scatter
 re.sub(r',', '', text) **Regex**
 re.findall()
 \w → word → [A-Z a-z]
 \d → digit → [0-9]
 \ → Escape spec. char.
 *? → 0 of meer, 1 of meer
 ↳ any char
 () → capture {} → wantat

```
np.linspace(0, 1, 5) - 0.2, 0.4
```

```
Linear(axis=0)
```

```
np.concatenate([a, b], axis=1)
```

```
np.full((3, 5), 3.14)
```

```
np.all(x==6)
```

```
x[[2, 1, 0]] == 10
```

```
np.argmax / np.argmin index of min value
```

Rule 3: if any dimension the sizes disagree and neither is equal to 1, an error is raised

```
data.loc[data.density > 100, ['pop', 'density']]
```

```
np.nansum(), pd.merge(df1, df2, on='employee')
```

```
df.dropna(), pd.merge(df1, df2, left_on='employee', right_on='name')
```

```
.drop('name', axis=1)
```

pd.join is selfe als merge, alleen autom-op index

```
df.pivot_table(values='births', columns='name', index='year',
```

```
aggfunc=sum)
```

Madekn excel: schouft niet, beperkte functionaliteit, integratie met andere tools

```
cross=crossstab(df.jaar, df.party)[party in [0, 10].index.values]
```

1. Panda helps filling in the gap with Python concerning data analysis and modeling, without having to switch to a more domain specific language like R

2. Python data analyze excels in performance, productivity and the ability to collaborate

```
np.arange(0, 14, 2), create_array(14, 7), np.reshape(L, (y, x))
```

```
laatste kolom = L[:, -1], L[begin: eind]
```

```
df.columns, len(df.index) geeft hoeveelheid rijen
```

index, value - counts

```
cito.index.str.lower(), str.strip(), str.replace(" ", "").sort_values()
```

```
cito[n.index.str.contains('school')], cito.describe()
```

```
cito['RMSE'] = np.sqrt((cito - cito.verwacht)**2), arr[:, -1]
```

```
np.random(0, 10, (3, 3))
```

```
x[1::2] every other elem, x[5:: -2]
```

```
np.split(x, [3, 5]), spliten op 3 en 5
```

```
x < 3 geeft array, delta(i) loc [ ]
```

```
data.nansum, data.fillna(), data.reset()
```

```
df.groupby('key').sum(), pd.read_csv/html
```

```
data.plot(x='', y='', kind='bar')
```

Kom met hypothese
verzamel data
schoon op
structureer
analyseer
rapporteer

2de ingesloten
gise pas en klopt niet zelf en klopt
ziet in font niet zelf en font

$$\frac{g}{g_0} \Big/ 1 \quad acc = \frac{TP + TN}{TP + FP + TN + FN}$$

g ₀	g ₁	
10	10	g ₂
g ₀	g ₁	10

$$gg \frac{g}{g_0} \Big/ 1$$