

Untitled

April 11, 2024

0.1 basics statistik

```
[1]: # Geschwindigkeiten einer Liste Autos  
speed = [99,86,87,88,111,86,103,87,94,78,77,85,86]
```

```
[2]: import numpy
```

```
[3]: x = numpy.mean(speed)  
print(x)
```

89.76923076923077

```
[4]: x = numpy.median(speed)  
print(x)
```

87.0

```
[5]: x = numpy.std(speed)  
print(x)
```

9.258292301032677

0.2 erzeuge custom datasets

```
[14]: x = numpy.random.uniform(0,5,250) #uniformverteilung
```

```
[10]: print(x)
```

```
[0.06032085  3.41320662  4.50593933  2.94637761  0.1410724  0.03800528  
 0.61358343  3.04186485  4.51038572  0.98294072  4.58864088  3.90476907  
 4.41092348  0.38499656  1.78781045  2.80756224  4.84983127  2.92598717  
 4.65906916  2.11561923  2.01634911  0.85464482  3.13599283  3.02666343  
 1.81486456  3.863421    3.31720273  0.12463773  1.45120032  3.98421884  
 3.06859232  0.37501715  2.75653545  0.72639817  2.78597801  3.10733904  
 0.31204693  2.55718282  1.53773108  4.99270402  3.21495013  0.6232707  
 3.31362495  2.63479695  4.3938338   0.47324732  3.30203586  3.1500965  
 4.37854142  4.74025522  2.65357211  2.25992061  1.92873815  4.88152483]
```

```

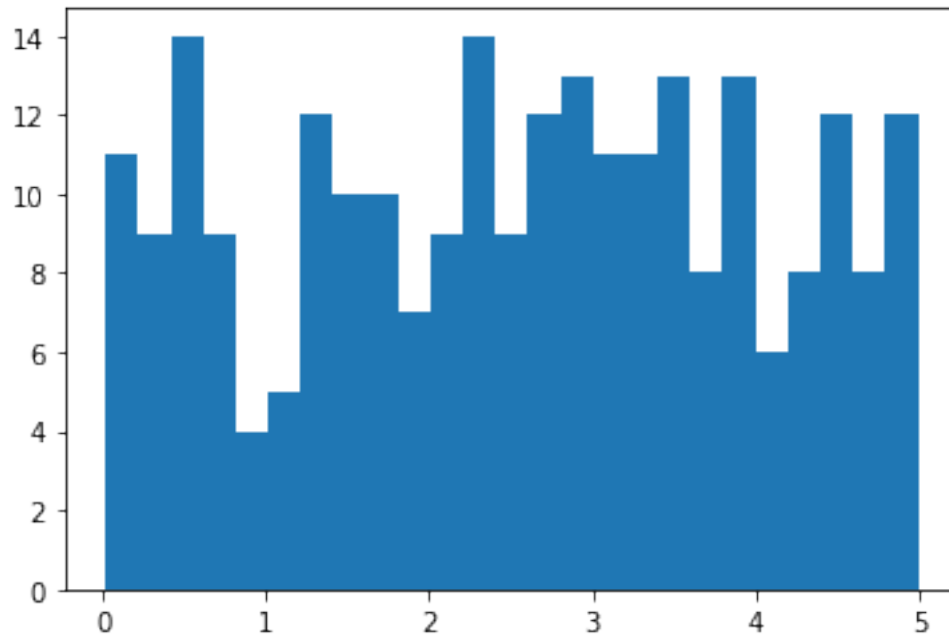
2.15434613 4.38998886 0.30736193 2.3632409 4.04011302 4.11126082
1.60374169 2.25418248 2.51992721 1.39224673 2.1363664 2.80107246
2.42204608 2.11361577 3.64254711 1.26549142 1.41635017 2.96530357
1.29425713 0.42862183 4.86972183 1.71812478 3.91118365 4.85783466
0.54021423 4.78779081 3.0101047 1.35116234 0.43672425 3.22778641
2.87084451 4.44010869 2.63860174 2.57718409 2.70430964 1.72733085
4.30738876 3.56836191 0.55241982 2.2541157 1.5061269 3.57981334
3.17810377 1.79708331 2.25928868 4.9927406 1.88717372 1.93011469
2.38665356 3.94196158 3.6756224 2.98491367 4.88482235 1.23242381
2.92971707 1.1780485 0.68346324 3.54768516 3.80014073 2.7203012
4.70417976 2.88833576 4.61005805 2.98429002 0.01971999 0.18833381
3.48840662 0.65998992 3.05333378 4.74968633 0.04010037 2.66009305
1.56150523 2.40952926 2.93374553 0.51797123 4.57704366 4.81856347
2.99439886 2.17979125 0.638195 3.89921225 3.08957522 0.51724147
1.9259026 1.67232498 4.56993758 0.50004429 3.53970306 4.28094638
3.83724574 0.90323611 0.81105364 4.15982666 0.27524222 1.32087187
1.6925015 2.33706245 3.80631326 3.65370665 0.01805888 3.5808555
0.58504058 1.66838627 1.40917914 4.57388805 3.81163106 2.51983384
2.20914894 4.58949015 3.60304497 2.88494161 3.53935739 0.56043038
3.9370061 1.63979289 2.32126025 2.83257741 0.02808154 2.23171813
0.03305873 0.31463212 0.61397136 1.80897995 1.59131484 1.76998698
4.31864891 3.9111814 1.77800511 3.58599208 0.63408624 4.5813537
2.62056688 4.49768174 1.26283547 4.62449891 0.32565243 4.90413336
2.39251465 0.76415258 3.30577558 4.14173351 3.60417479 3.31834342
2.61329669 1.09748908 0.52201105 3.65577271 3.39536917 0.95301561
3.63299523 4.33256744 1.5509261 3.87425944 3.22123705 1.18336251
4.16734426 3.57210594 4.76150162 0.25141063 3.71440155 3.30466563
0.2763223 1.33533215 3.59761502 3.46921096 1.24769155 2.45639717
4.84758404 2.61370374 1.21273137 3.44693113 0.75688368 0.18610356
1.08688544 1.45222248 3.39073385 4.12572989 2.36349516 2.57995574
2.18933009 1.41458555 2.30596671 2.38552105 1.01483024 0.45526327
1.21258965 4.26153392 1.99191466 3.09577282 4.81372709 2.06631544
2.46672297 4.83009026 4.52474431 2.03129534]

```

```

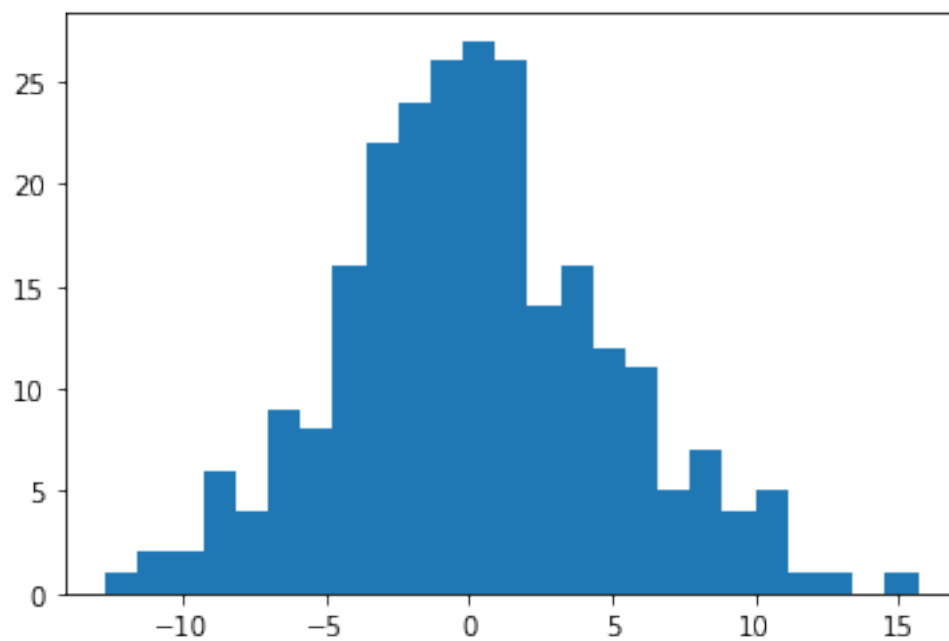
[13]: import matplotlib.pyplot as plt
plt.hist(x,25)
plt.show()

```



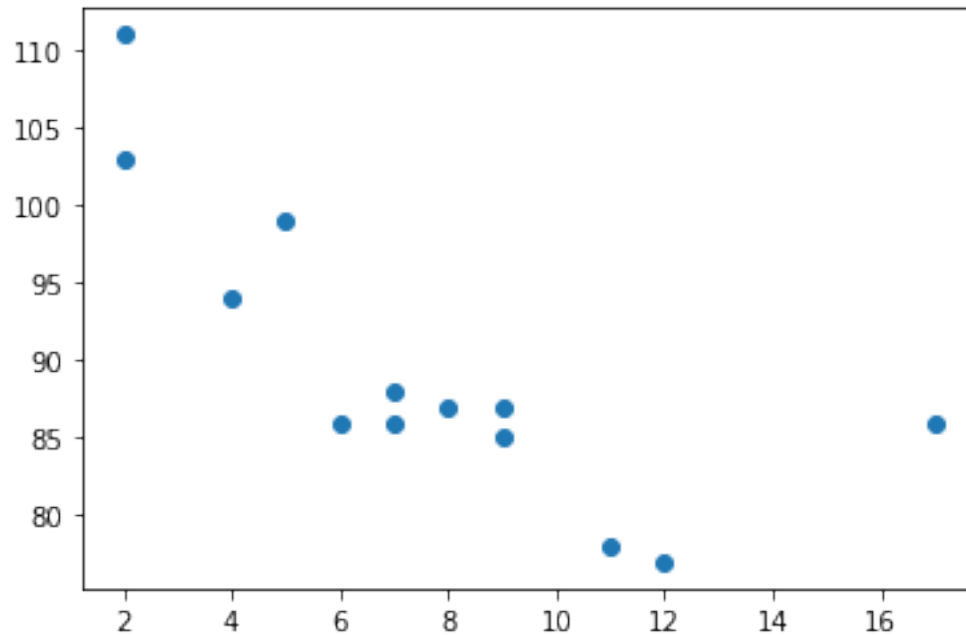
```
[15]: x = numpy.random.normal(0,5,250) #normalverteilung
```

```
[16]: import matplotlib.pyplot as plt
plt.hist(x,25)
plt.show()
```



```
[17]: x = [5,7,8,7,2,17,2,9,4,11,12,9,6]
      y = [99,86,87,88,111,86,103,87,94,78,77,85,86]

      plt.scatter(x,y)
      plt.show()
```



0.3 lineare regression

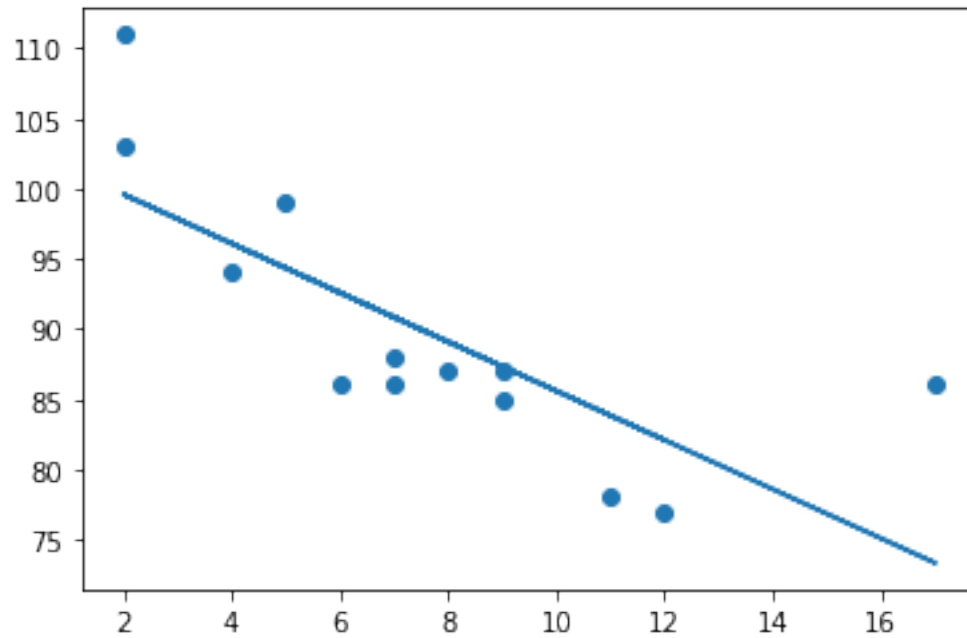
```
[18]: from scipy import stats
```

```
[21]: slope, intercept, r, p, std_err = stats.linregress(x,y)

      def myfunc(x):
          return slope *x + intercept

      mymodel = list(map(myfunc,x))

      plt.scatter(x,y)
      plt.plot(x,mymodel)
      plt.show()
```

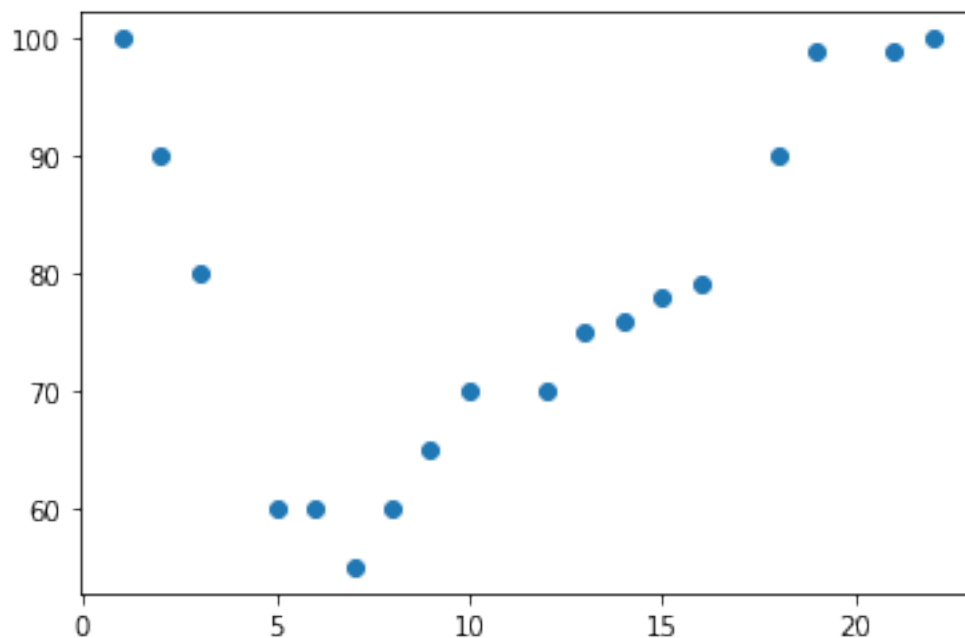


```
[22]: speed = myfunc(10)
      print(speed)
```

85.59308314937454

0.4 polynomische regression

```
[23]: import matplotlib.pyplot as plt
      x = [1,2,3,5,6,7,8,9,10,12,13,14,15,16,18,19,21,22]
      y = [100,90,80,60,60,55,60,65,70,70,75,76,78,79,90,99,99,100]
      plt.scatter(x,y)
      plt.show()
```



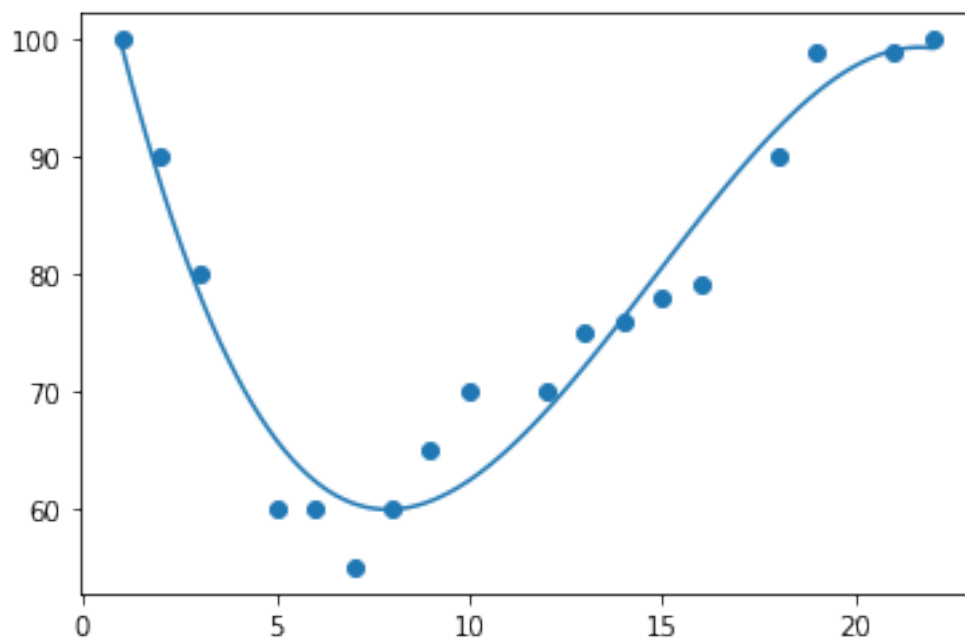
```
[29]: mymodel = numpy.poly1d(numpy.polyfit(x,y,3))
```

```
myline = numpy.linspace(1,22,100)
```

```
plt.scatter(x,y)
```

```
plt.plot(myline,mymodel(myline))
```

```
plt.show()
```



0.5 echten Datensatz

```
[30]: import pandas as pd
      df = pd.read_csv('/Users/h4/desktop/cars.csv')
```

```
[31]: df.head()
```

```
[31]:
```

	Car	Model	Volume	Weight	CO2
0	Toyoty	Aygo	1000	790	99
1	Mitsubishi	Space Star	1200	1160	95
2	Skoda	Citigo	1000	929	95
3	Fiat	500	900	865	90
4	Mini	Cooper	1500	1140	105

```
[32]: X = df[['Weight', 'Volume']]
      y = df['CO2']
```

```
[33]: from sklearn import linear_model
```

```
[35]: # !pip install sklearn (falls library nicht vorhanden)
```

```
[36]: regression = linear_model.LinearRegression()
      regression.fit(X,y)
```

```
[36]: LinearRegression()
```

```
[37]: predictedCO2 = regression.predict([[2300, 1300]])
```

```
/Users/h4/anaconda3/lib/python3.9/site-packages/sklearn/base.py:450:
UserWarning: X does not have valid feature names, but LinearRegression was
fitted with feature names
  warnings.warn(
```

```
[38]: predictedCO2
```

```
[38]: array([107.2087328])
```

0.6 clustering

```
[40]: wine = pd.read_csv('/Users/h4/desktop/wine.csv')
```

```
[41]: wine.head()
```

```
[41]:
```

	Cultivar	Alcohol	Malic acid	Ash	Alcalinity of ash	Magnesium	\
0	1	14.23	1.71	2.43		15.6	127
1	1	13.20	1.78	2.14		11.2	100
2	1	13.16	2.36	2.67		18.6	101
3	1	14.37	1.95	2.50		16.8	113
4	1	13.24	2.59	2.87		21.0	118

	Total phenols	Flavanoids	Nonflavanoid phenols	Proanthocyanins	\
0	2.80	3.06		0.28	2.29
1	2.65	2.76		0.26	1.28
2	2.80	3.24		0.30	2.81
3	3.85	3.49		0.24	2.18
4	2.80	2.69		0.39	1.82

	Color intensity	Hue	OD280/OD315 of diluted wines	Proline	
0	5.64	1.04		3.92	1065
1	4.38	1.05		3.40	1050
2	5.68	1.03		3.17	1185
3	7.80	0.86		3.45	1480
4	4.32	1.04		2.93	735

```
[42]: # drop cultivar, da nicht notwendig für die clusterung
```

```
wine = wine.drop('Cultivar', axis=1)
```

```
[43]: wine.head()
```

```
[43]:
```

	Alcohol	Malic acid	Ash	Alcalinity of ash	Magnesium	Total phenols	\
0	14.23	1.71	2.43		15.6	127	2.80
1	13.20	1.78	2.14		11.2	100	2.65
2	13.16	2.36	2.67		18.6	101	2.80
3	14.37	1.95	2.50		16.8	113	3.85
4	13.24	2.59	2.87		21.0	118	2.80

	Flavanoids	Nonflavanoid phenols	Proanthocyanins	Color intensity	Hue	\
0	3.06		0.28		2.29	5.64 1.04
1	2.76		0.26		1.28	4.38 1.05
2	3.24		0.30		2.81	5.68 1.03
3	3.49		0.24		2.18	7.80 0.86
4	2.69		0.39		1.82	4.32 1.04

	OD280/OD315 of diluted wines	Proline
0		3.92 1065
1		3.40 1050
2		3.17 1185
3		3.45 1480
4		2.93 735


```
[44]: from sklearn.cluster import KMeans
```

```
[46]: kmeans = KMeans(n_clusters = 3, random_state=42).fit(wine.values)
```

```
[47]: kmeans_3 = pd.DataFrame(kmeans.labels_, columns = ['cluster'])  
print(kmeans_3)
```

	cluster
0	1
1	1
2	1
3	1
4	2
...	...
173	2
174	2
175	2
176	2
177	0

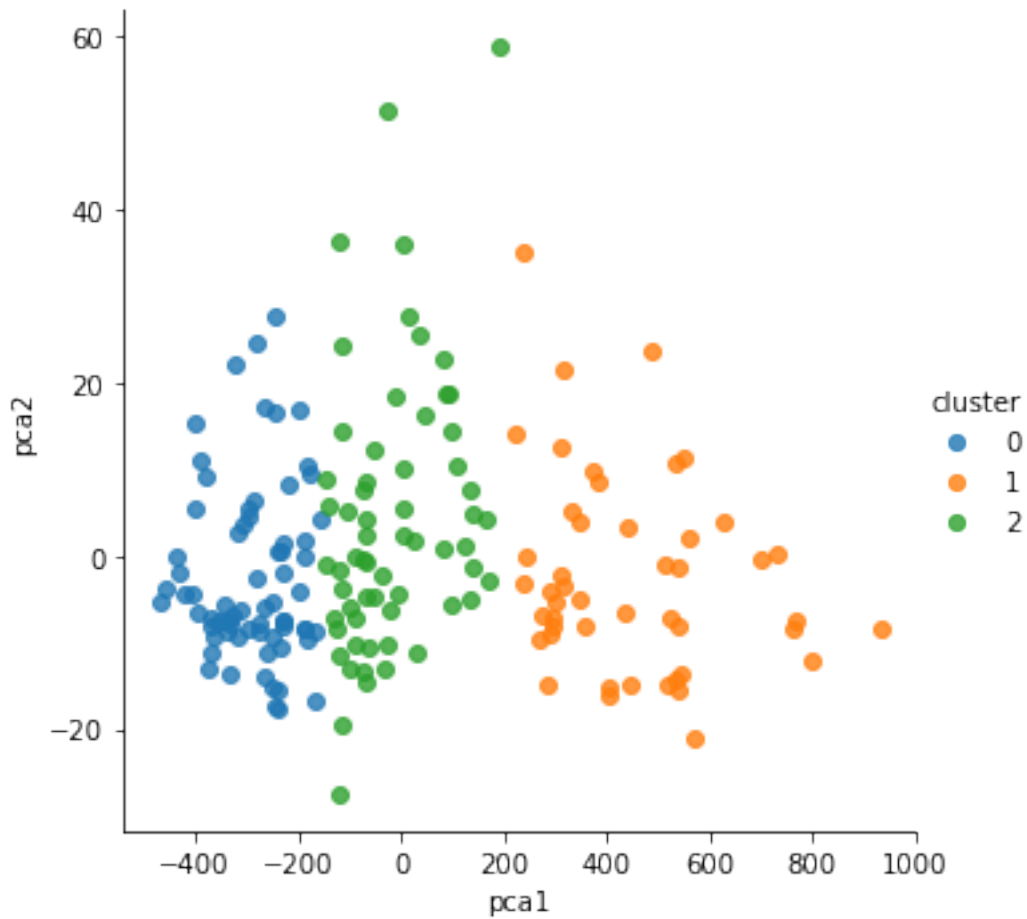
[178 rows x 1 columns]

0.7 hauptkomponentenanalyse (principal component analysis)

```
[48]: from sklearn.decomposition import PCA  
pca = PCA(n_components=2).fit(wine)
```

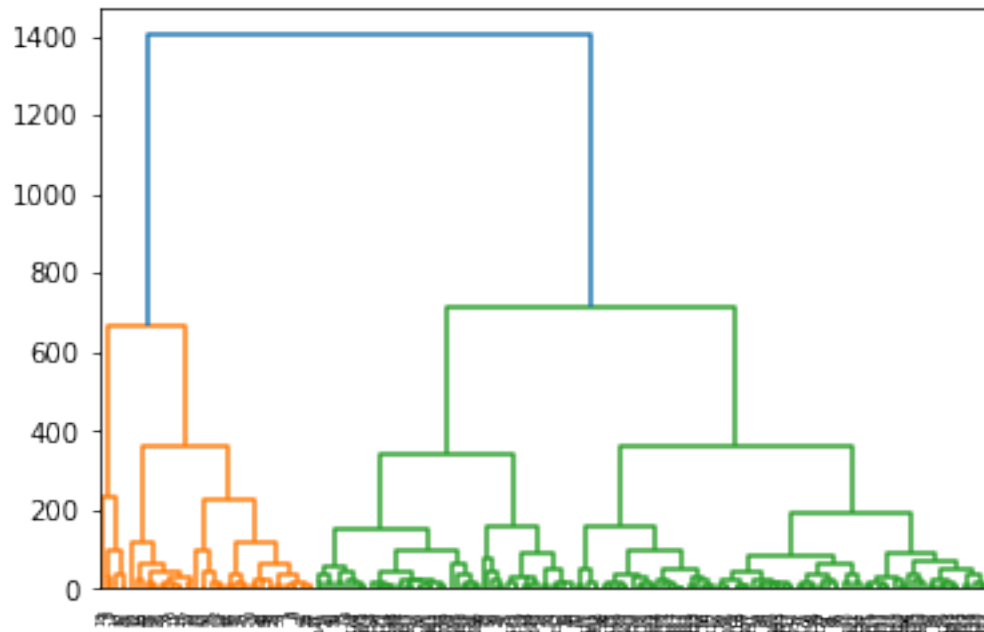
```
[50]: pca_trans = pca.transform(wine)  
pca_trans_df = pd.DataFrame(pca_trans, columns = ['pca1', 'pca2'])  
kmeans_3 = pd.concat([kmeans_3, pca_trans_df], axis= 1)
```

```
[51]: import seaborn as sns  
import matplotlib.pyplot as plt  
  
fig = sns.lmplot(x='pca1', y='pca2', data=kmeans_3, hue='cluster',  
                ↪fit_reg=False)  
plt.show()
```



0.8 hierarchical clustering

```
[53]: from scipy.cluster import hierarchy
      wine_complete = hierarchy.complete(wine)
      fig = plt.figure()
      dn = hierarchy.dendrogram(wine_complete)
      plt.show()
```



0.9 face recognition mit support vector machines

```
[54]: from sklearn.datasets import fetch_lfw_people
faces = fetch_lfw_people(min_faces_per_person=60)
print(faces.target_names)
print(faces.images.shape)
```

```
['Ariel Sharon' 'Colin Powell' 'Donald Rumsfeld' 'George W Bush'
 'Gerhard Schroeder' 'Hugo Chavez' 'Junichiro Koizumi' 'Tony Blair']
(1348, 62, 47)
```

```
[56]: import matplotlib.pyplot as plt
fig, ax = plt.subplots(3,5)
for i, axi in enumerate(ax.flat):
    axi.imshow(faces.images[i], cmap='bone')
    axi.set(xticks=[], yticks=[],
            xlabel=faces.target_names[faces.target[i]])
```



```
[58]: from sklearn.svm import SVC
      from sklearn.decomposition import PCA as RandomizedPCA
      from sklearn.pipeline import make_pipeline

      pca = RandomizedPCA(n_components = 90, whiten = True, random_state = 42)
      svc = SVC(kernel='rbf', class_weight='balanced')
      model = make_pipeline(pca,svc)

[59]: from sklearn.model_selection import train_test_split
      X_train, X_test, y_train, y_test = train_test_split(faces.data, faces.target,
      ↪random_state=42)

[61]: # Zeile nicht prüfungsrelevant

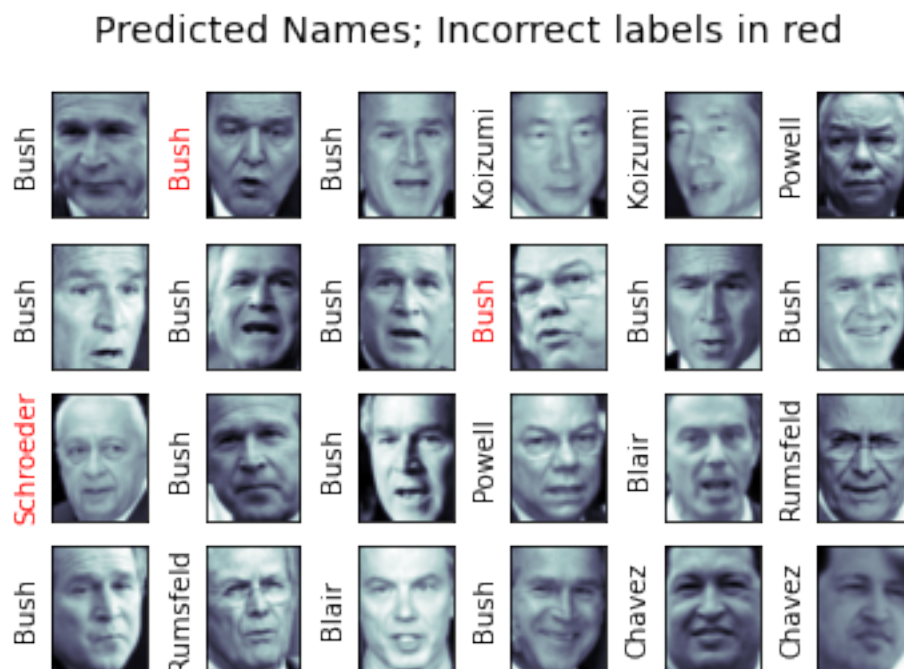
      from sklearn.model_selection import GridSearchCV
      param_grid = {'svc__C': [1, 5, 10, 50], 'svc__gamma': [0.0001, 0.0005, 0.001, 0.
      ↪0.005]}
      grid = GridSearchCV(model, param_grid)
      %time grid.fit(X_train, y_train)
      print(grid.best_params_)

CPU times: user 31min 45s, sys: 25min, total: 56min 46s
Wall time: 6min 35s
{'svc__C': 5, 'svc__gamma': 0.005}
```

```
[62]: model = grid.best_estimator_
yfit = model.predict(X_test)
```

```
[66]: fig, ax = plt.subplots(4,6)
for i, axi in enumerate(ax.flat):
    axi.imshow(X_test[i].reshape(62,47), cmap='bone')
    axi.set(xticks=[], yticks=[])
    axi.set_ylabel(faces.target_names[yfit[i]].split()[-1],
                  color='black' if yfit[i]==y_test[i] else 'red')
fig.suptitle('Predicted Names; Incorrect labels in red', size=14)
```

```
[66]: Text(0.5, 0.98, 'Predicted Names; Incorrect labels in red')
```



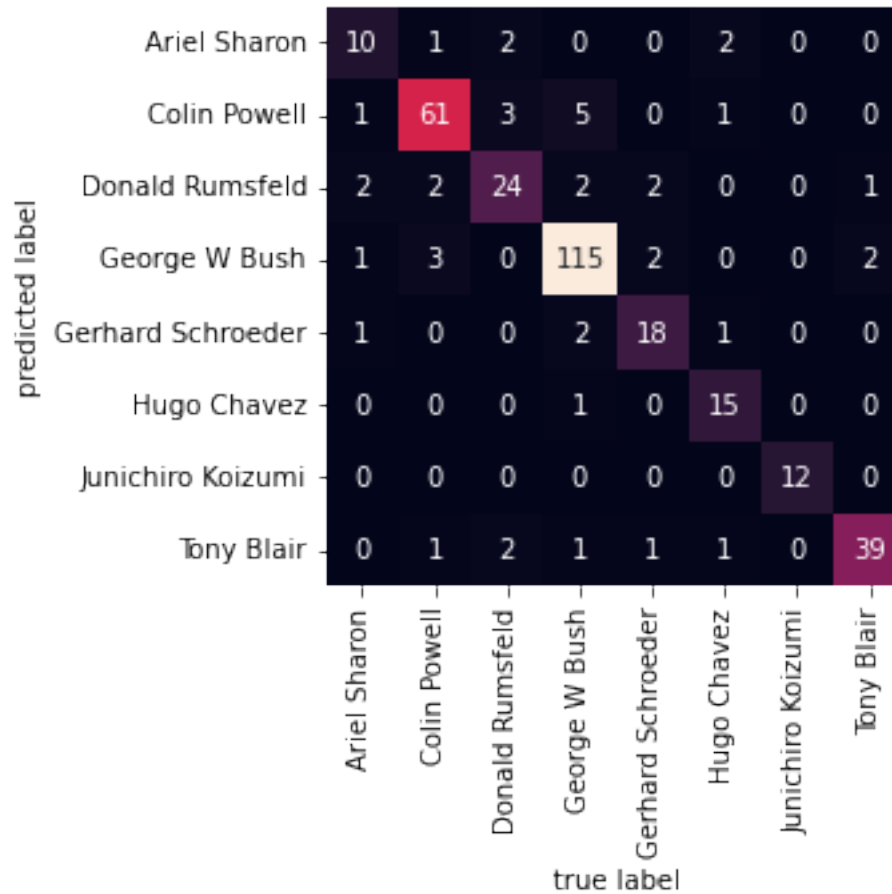
```
[67]: # confusion matrix

from sklearn.metrics import confusion_matrix

mat = confusion_matrix(y_test, yfit)

sns.heatmap(mat.T, square=True, annot=True, fmt='d', cbar=False,
            xticklabels=faces.target_names,
            yticklabels=faces.target_names)
plt.xlabel('true label')
plt.ylabel('predicted label')
```

```
[67]: Text(91.68, 0.5, 'predicted label')
```

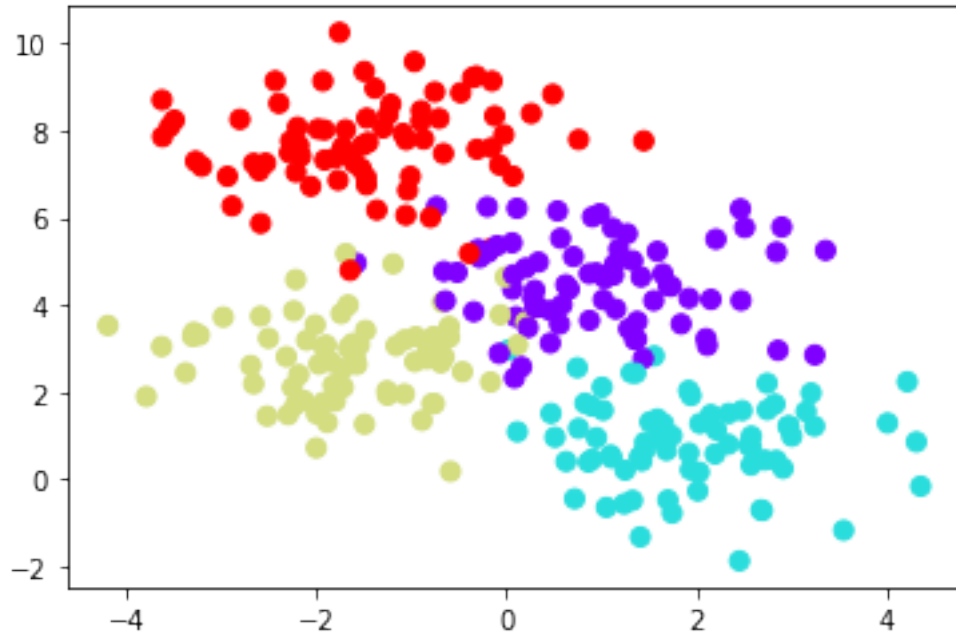


0.10 decision tree

```
[70]: from sklearn.datasets import make_blobs

X, y = make_blobs(n_samples=300, centers=4, random_state=0, cluster_std=1)
plt.scatter(X[:,0], X[:,1], c=y, s=50, cmap='rainbow')
```

```
[70]: <matplotlib.collections.PathCollection at 0x32fcafa90>
```



```
[71]: from sklearn.tree import DecisionTreeClassifier
tree = DecisionTreeClassifier().fit(X,y)
```

```
[76]: #Zeile ist nicht prüfungsrelevant
```

```
import numpy as np

def visualize_classifier(model, X, y, ax=None, cmap='rainbow'):
    ax = ax or plt.gca()
    # Plot the training points
    ax.scatter(X[:, 0], X[:, 1], c=y, s=30, cmap=cmap,
               clim=(y.min(), y.max()), zorder=3)
    ax.axis('tight')
    ax.axis('off')
    xlim = ax.get_xlim()
    ylim = ax.get_ylim()
    # fit the estimator
    model.fit(X, y)
    xx, yy = np.meshgrid(np.linspace(*xlim, num=200),
                         np.linspace(*ylim, num=200))
    Z = model.predict(np.c_[xx.ravel(), yy.ravel()]).reshape(xx.shape)
    # Create a color plot with the results
    n_classes = len(np.unique(y))
    contours = ax.contourf(xx, yy, Z, alpha=0.3,
                          levels=np.arange(n_classes + 1) - 0.5,
                          cmap=cmap, clim=(y.min(), y.max()),
```

```

        zorder=1)
    ax.set(xlim=xlim, ylim=ylim)

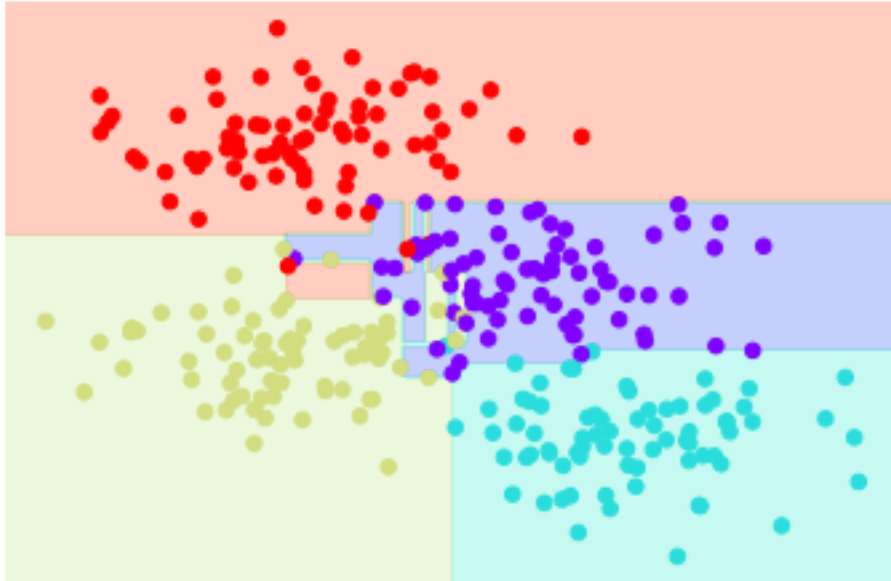
```

```
[77]: visualize_classifier(DecisionTreeClassifier(), X,y)
```

```

/var/folders/nw/k_k0_cbj7vl_npdmyvhl53c0000gn/T/ipykernel_46393/2167791633.py:2
1: UserWarning: The following kwargs were not used by contour: 'clim'
    contours = ax.contourf(xx, yy, Z, alpha=0.3,

```



0.11 random forest für Bilder Klassifizierung

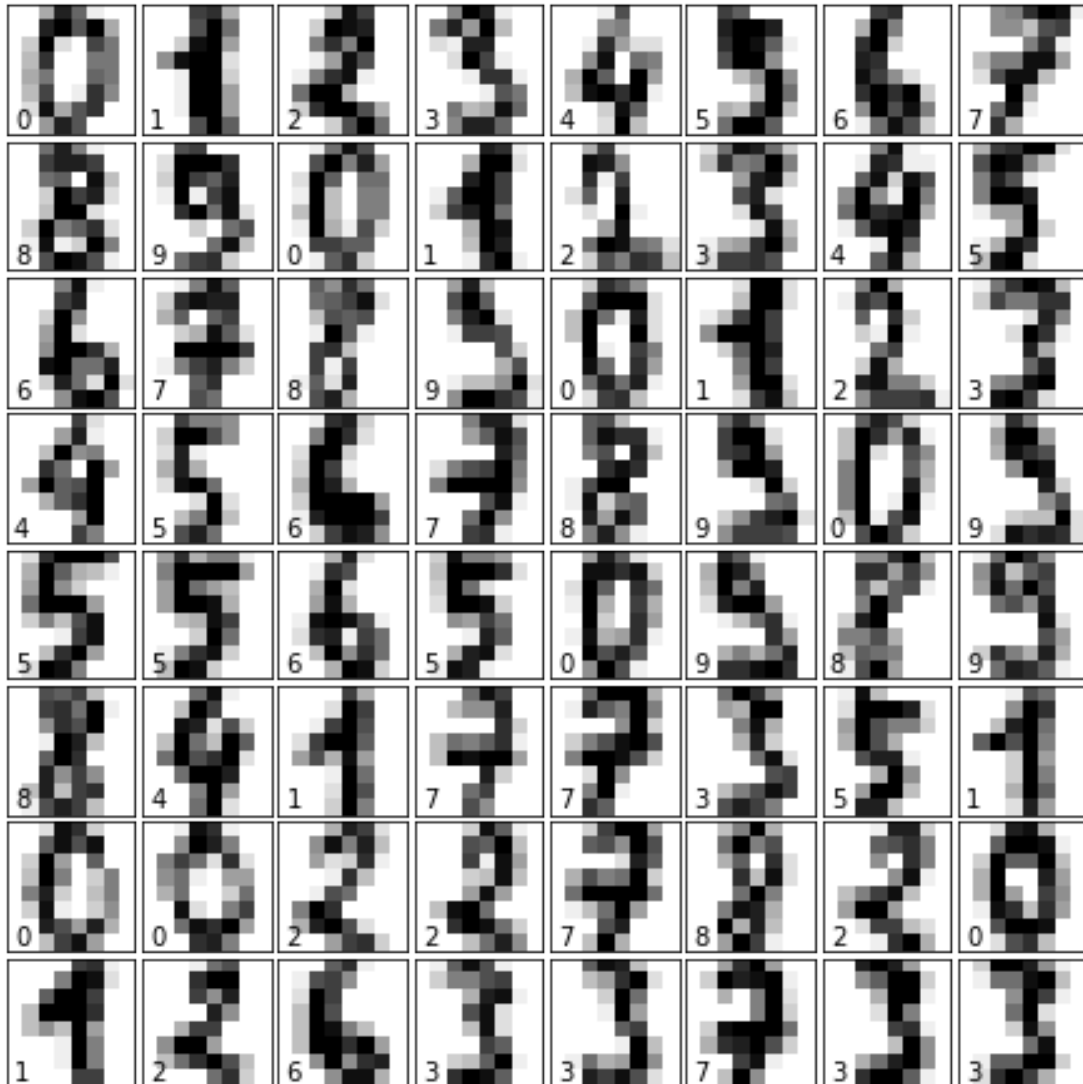
```
[78]: from sklearn.datasets import load_digits
      digits = load_digits()
      digits.keys()
```

```
[78]: dict_keys(['data', 'target', 'frame', 'feature_names', 'target_names', 'images',
               'DESCR'])
```

```
[80]: # Ziele nicht Prüfungsrelevant

fig = plt.figure(figsize=(6, 6))
fig.subplots_adjust(left=0, right=1, bottom=0, top=1, hspace=0.05, wspace=0.05)
for i in range(64):
    ax = fig.add_subplot(8, 8, i + 1, xticks=[], yticks=[])
    ax.imshow(digits.images[i], cmap=plt.cm.binary, interpolation='nearest')
    ax.text(0, 7, str(digits.target[i]))

```

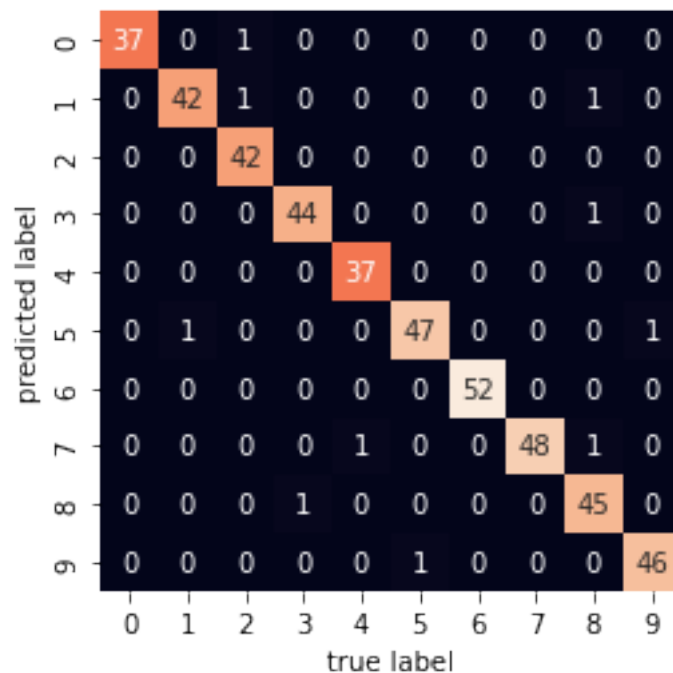
```
[82]: from sklearn.model_selection import train_test_split
```

```
[83]: Xtrain, Xtest, ytrain, ytest = train_test_split(digits.data, digits.target,
↳ random_state=0)
```

```
[86]: from sklearn.ensemble import RandomForestClassifier
model = RandomForestClassifier(n_estimators=1000)
model.fit(Xtrain, ytrain)
ypred = model.predict(Xtest)
```

```
[87]: from sklearn.metrics import confusion_matrix
mat = confusion_matrix(ytest, ypred)
sns.heatmap(mat.T, square=True, annot=True, fmt='d', cbar=False)
```

```
plt.xlabel('true label')
plt.ylabel('predicted label');
```



```
[88]: #####
```

```
[ ]: # Maschine 1.
```

```
# Erzeugen Sie bitte einen Dataset mit 10 Dimensionen (10 KPIs).
# Die Dimensionen sollten 10 Variablen von einer Maschine darstellen.
# Die Variablen sind mit einer Weibul Verteilung ausgelegt mit Formparameter = 4,8 bei 3 Varibeln, 3,6 bei den Rest.
# Skalenparameter für Alle Variablen 1.5.
# Pro KPI gibt es 2400 Datensätze.

# Bitte Erstellen einen Netzwerk mit den 10 dazu gehörigen Sensoren mit einem CC=2, und ein APL=2*ln(10).

# Bitte berechnen Sie den Laplacian vom Netzwerk. Dieser Lplacian Matrix sind die X und Y Dimensionen vom Zieldatset.
# Die Z-Dimension wird durch die KPIs dargestellt, jeweils auf dem i=i Diagonale vom Laplacian.

# Somit ist unser Zieldatset (10,10, 2400).
```

```
# Maschine 2.  
# wie Maschine 1 aber:  
# Formparameter = 1.2 für 8 Sensoren und 2.3 für den Rest.  
# CC=4, APL=ln(10)  
  
## Sobald die Datensätze definiert wurden, bitte nutzen Sie PCA und K-Means  
↳ Cluster um beide Maschinen in einem 2 dimensionalen Raum darzustellen.  
## Die Clusterung sollte erfolgen nach dem Formparameter der Weibul Verteilung.
```