6,5 (10) No. 74 9) 1. The data points are centralized 70 their mean values. 2. Then using the covariance matrix, eigenvalues and eigenvectors, the correlation of values is calculated. Correlation of the 3. Choice of the main correlation by on = ? picking the largest eigenvalue. while using the word correlation 4. Filling the property matrix W with is not complexity wrong, the idea the eigenvectors Application of the property matrix that incorporate the highest amount to the individual data points of intermation 6. The closer the points are on the the highest correlation with (to some extend primary axis the more likely they the target variety are correlating. With what? b) x: [1,3,1,2,3,2] x2: [7,0,3,0,1,7] Megn values: X= - (7+0+3+0+1+1) x1 = 1 (1+3+1+2+3+2) New data points: [xi= x,-x,] caltulating Covariance Matrix: cor (xi, xi) = 1 2 (xix xyx) cov (x1,x1) = = -((-1)2+12+(-1)2+02+12+02)= +5 Cov (x2, x2) = = = ((0+ (-1)2+ (2)2+ (-1)2+02+04) = 6/5 BRUNNEN I COV (x1/x2) = (0V(x2,x1)= 7 · (-7·0 + 7·6·1)+1-1)·2+0·(-1)+1·0 + 0-0) = -35

$$S = \frac{1}{5} \begin{pmatrix} 4 & -3 \\ -3 & 6 \end{pmatrix}$$

$$Eigenvalues on M - vectors:$$

$$det(S - XI) = 0$$

$$\frac{1}{5} \cdot det(\frac{1}{3} + \frac{1}{3}) = 0 + \frac{1}{4} (X - \lambda 4)$$

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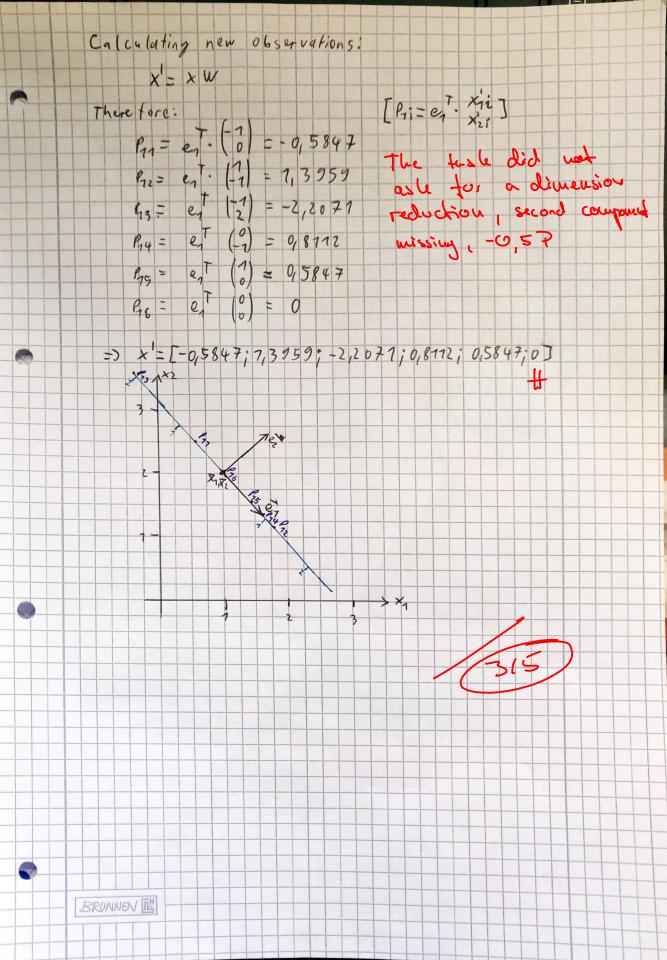
$$\frac{1}{5} \cdot det(\frac{1}{3} + \frac{1}{3}) = 0 + \frac{1}{4} (X - \lambda 4)$$

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$$\frac{1}{5} \cdot det(\frac$$



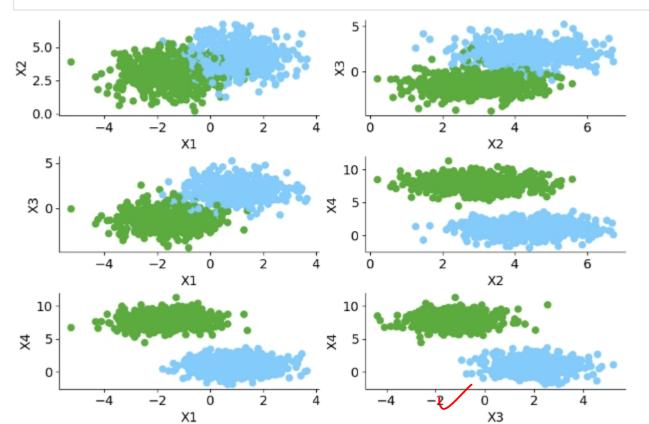
a)

This is headly readable

```
In [49]:
```

```
from ml import plots
import matplotlib
import matplotlib.pyplot as plt
import pandas as pd
import numpy as np
import itertools
plots.set_plot_style()
%matplotlib widget
colors = plots.colors
cmap = plots.cmap
from sklearn.datasets import make_blobs
X,y = make_blobs(n_samples=1000, centers=2, n_features=4, random_state=0)
fig, ax = plt.subplots(3,2)
feature_combinations = list(itertools.combinations(range(4), 2))
plot_pos = [(0,0), (1,0), (2,0), (0,1), (1,1), (2,1)]
for i in range(len(feature_combinations)):
    ax[plot\_pos[i]].scatter(X[:,feature\_combinations[i][0]], \ X[:,feature\_combinations[i][1]], \ c=y, \ s=50, \ cmap=cmap)
    ax[plot_pos[i]].set_xlabel('X'+str(feature_combinations[i][0]+1))
    ax[plot_pos[i]].set_ylabel('X'+str(feature_combinations[i][1]+1))
```





```
In [41]:
            from sklearn.decomposition import PCA
                                                      again not ask for
            pca = PCA(n_components = k)
            X_prime_sklearn = pca.fit_transform(X)
            cov_mat = pca.get_covariance()
            1, W = np.linalg.eigh(cov_mat)
            1 = 1[::-1]
            print('Eigenvalues', 1)
Out [41]:
          The eigenvalues describe the variance of the datapoints projected on the fisher discriminant (eigenvector) in that dimension
                                            ergemolies tell you about t
                                                       privcipal
                                                                     / component
In [57]:
            fig, ax = plt.subplots(3)
            for k in range(3):
               ax[k].hist(X_prime_sklearn[:,k])
               ax[k].set_xlabel("$x'$"+str(k+1))
               ax[k].set_ylabel('# of occurences')
                                                                 1 and 2
            None
          # of occurences
out [57]:
             200
                                                           -015 P
              100
                           -6
                                         -4
                                                      -2
                                                                    0
                                                                                 2
                                                                   x'1
          # of occurences
             200
              100
                0
                       -3
                                     -2
                                                    -1
                                                                    0
                                                                                   1
                                                                                                                 3
                                                                   x'2
          # of occurences
             200
             100
                0
                                             ×4 missing, -0,5 P
                            -3
                                                                                                          3
```

```
plt.xlabel("$x'_1$")
plt.ylabel("$x'_2$")
None
    3
    2
    1
    0
  ^{-1}
  -2
  -3
                                          -2
                                                                      2
               -6
                                                        0
                                                        x'_1
```

New Feature  $x_1'$  clearly delivers better information than the original  $x_1$ , a separation between the two populations is easier now

plt.scatter(X\_prime\_sklearn[:,0], X\_prime\_sklearn[:,1], c = y, cmap =cmap)

In [58]:

out [58]:

plt.figure()