

Learning about Kalman filter /

Resources

Kalman Filter from Ground Up ; author Alex Becker; <https://www.kalmanfilter.net>

Overview

This notebook covers the properties of the `multivariate normal distribution`. In large part the notebook is inspired by chapter 7.3 of the book `Kalman Filter from Ground Up`.

Covariance

The covariance provides a measure of the correlation of two or more random variables.

In its simplest form the properties of the covariance is studied using **two** random variables.

Data Sets / uncorrelated

Two example 2D data sets are created, displayed and it is shown numerically, that the x-y coordinates of the data set are uncorrelated.

Each data set shall have $N = 10000$ data points (x_n, y_n) . x_n and y_n are random variables.

For the first data set we assume that random variables x_n and y_n are drawn from uniform distributions. x_n is in the interval $-2, 3$ and y_n in the interval $-3, 1$.

For the second data set we assume that random variables x_n and y_n are drawn from normal distributions. The distribution for x_n has mean $\mu_x = 1$ and standard deviation $\sigma_x = 1$. The distribution for y_n has mean $\mu_y = 2$ and standard deviation $\sigma_y = 0.5$.

```
In [1]: import numpy as np
import scipy.stats as stats
import matplotlib.pyplot as plt
```

```
In [2]: # creating the data sets

N = 10000
x1 = stats.uniform.rvs(loc=-2, scale=5, size=N)
y1 = stats.uniform.rvs(loc=-3, scale=4, size=N)

x2 = stats.norm.rvs(loc=1, scale=1, size=N)
y2 = stats.norm.rvs(loc=2, scale=0.5, size=N)

var_x1 = np.var(x1)
```

```

var_y1 = np.var(y1)
cov_xy1 = np.cov(x1, y1)
trace_1 = np.trace(cov_xy1)

var_x2 = np.var(x2)
var_y2 = np.var(y2)
cov_xy2 = np.cov(x2, y2)
trace_2 = np.trace(cov_xy2)

print(f"var_x1: {var_x1:.3e}; var_y1: {var_y1:.3e}; var_x1+var_y1: {var_x1 + var_y1}
print(f"var_x2: {var_x2:.3e}; var_y2: {var_y2:.3e}; var_x2+var_y2: {var_x2 + var_y2}

```

```

var_x1: 2.089e+00; var_y1: 1.346e+00; var_x1+var_y1: 3.435e+00; cov_xy1: [[ 2.0893935
7 0.05035663]
[0.05035663 1.34621142]]; trace(cov_xy1):3.436e+00
var_x2: 9.999e-01; var_y2: 2.501e-01; var_x2+var_y2: 1.250e+00; cov_xy2: [[ 0.999994
4 -0.00361295]
[-0.00361295 0.25016953]]; trace(cov_xy2):1.250e+00

```

Notes

Apart from numerical artifacts / rounding errors the trace of the covariance matrix is identical to the sum of variances. This is the expected result because of the *uncorrelatedness* of the components of each data set.

Ideally the off-diagonal elements of the correlation matrix should be 0. Due to numerical effects these elements are quite small.

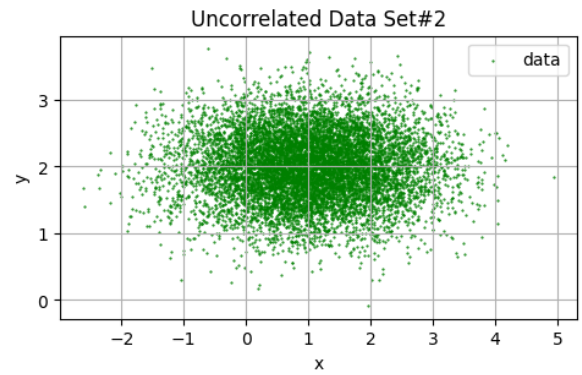
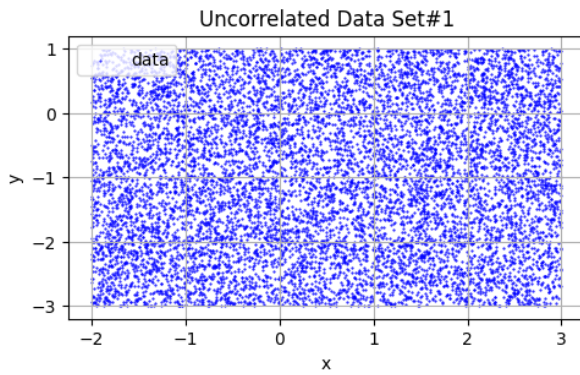
```

In [3]: fig1, ax1 = plt.subplots(nrows=1, ncols=2, figsize=(12,3))

ax1[0].scatter(x1, y1, color='b', s=1, marker='.', label='data')
ax1[0].set_xlabel('x')
ax1[0].set_ylabel('y')
ax1[0].set_title('Uncorrelated Data Set#1')
ax1[0].legend()
ax1[0].grid(True)

ax1[1].scatter(x2, y2, s=1, color='g', marker='.', label='data')
ax1[1].set_xlabel('x')
ax1[1].set_ylabel('y')
ax1[1].set_title('Uncorrelated Data Set#2')
ax1[1].legend()
ax1[1].grid(True)

```



Data Sets / correlated

Here we create a data set with uncorrelated components and another one which is correlated. The correlation is accomplished via a mixing process.

recipe

Generate a random vector $\mathbf{x}[k]$ with two elements $x_1[k]$, $x_2[k]$:

$$\mathbf{x}[k] = \begin{bmatrix} x_1[k] \\ x_2[k] \end{bmatrix} : k := 1, \dots, K$$

There shall be K samples from this vector. These sample are stored in a matrix $\mathbf{X} : \in \mathbb{R}^{2 \times K}$

$$\mathbf{X} = \begin{bmatrix} | & | & \dots & | \\ \mathbf{x}[1] & \mathbf{x}[2] & \dots & \mathbf{x}[K] \\ | & | & \dots & | \end{bmatrix} = \begin{bmatrix} x_1[1] & x_1[2] & \dots & x_1[K] \\ x_2[1] & x_2[2] & \dots & x_2[K] \end{bmatrix}$$

The matrix \mathbf{X} is right multiplied to a weighting matrix $\mathbf{W} : \in \mathbb{R}^{2 \times 2}$ to obtain a new random matrix \mathbf{Y} :

$$\mathbf{W} = \begin{bmatrix} w_{1,1} & w_{1,2} \\ w_{2,1} & w_{2,2} \end{bmatrix}$$

$$\mathbf{y} = \mathbf{W} \cdot \mathbf{X}$$

The covariance of matrices \mathbf{X} and \mathbf{Y} are computed. While the covariance matrix of \mathbf{X} has only significant entries for the main diagonal, the covariance of \mathbf{Y} has significant entries everywhere due to the processing of the weighting matrix.

A scatter plot of the data in the rows of matrix \mathbf{X} show the de-correlation of samples. However a scatter plot of rows of matrix \mathbf{Y} shows a diagonal orientation due to the mutual correlation.

```
In [4]: # two independent random variables
K = 100000
x1 = np.random.randn(K)
x2 = np.random.randn(K)
```

```

# each row represents a series of randomly distributed numbers
Xmat = np.vstack((x1, x2))

# weighting matrix: defines the mixing process
Wmat = np.array([[1, 2],[3, 1]])

# build a new data set by mixing
Ymat = Wmat @ Xmat

# compute covariance matrices
Cmat_x = np.cov(Xmat)
Cmat_y = np.cov(Ymat)

print("covariance matrices\n")
print(f"Cmat_x   :\n{Cmat_x}\n")
print(f"Cmat_y   :\n{Cmat_y}")

```

covariance matrices

```

Cmat_x  :
[[0.99472722 0.00317491]
 [0.00317491 0.99650014]]

```

```

Cmat_y  :
[[4.99342744 4.99940632]
 [4.99940632 9.96809459]]

```

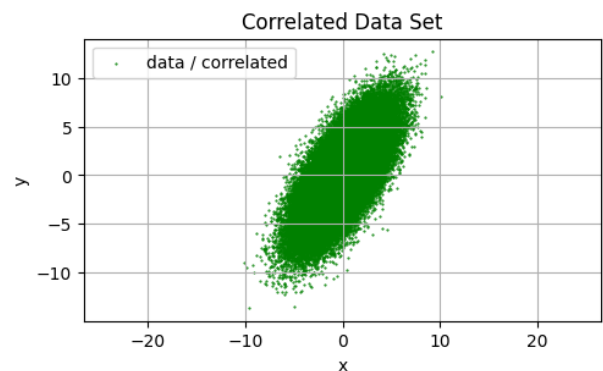
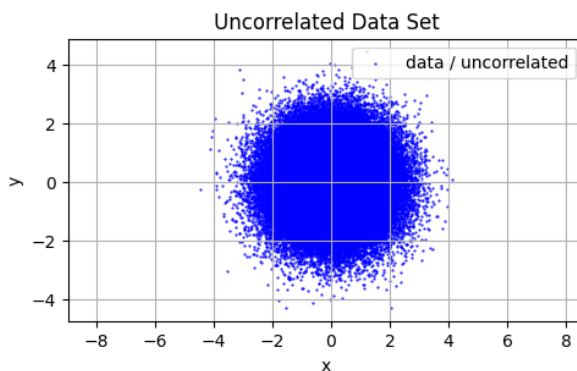
```

In [5]: fig2, ax2 = plt.subplots(nrows=1, ncols=2, figsize=(12,3))

ax2[0].scatter(Xmat[0,:], Xmat[1,:], color='b', s=1, marker='.', label='data / unc
ax2[0].set_xlabel('x')
ax2[0].set_ylabel('y')
ax2[0].set_title('Uncorrelated Data Set')
ax2[0].legend()
ax2[0].axis('equal')
ax2[0].grid(True)

ax2[1].scatter(Ymat[0,:], Ymat[1,:], s=1, color='g', marker='.', label='data / corr
ax2[1].set_xlabel('x')
ax2[1].set_ylabel('y')
ax2[1].set_title('Correlated Data Set')
ax2[1].legend()
ax2[1].axis('equal')
ax2[1].grid(True)

```



The multivariate normal distribution

In one dimension the normal distribution has a probability density function $p(x)$ defined by this formula:

$$p(x) = \frac{1}{\sqrt{2\pi\sigma^2}} \cdot \exp\left[-\frac{(x - \mu)^2}{2\sigma^2}\right]$$

In more than one dimension the probability density of the multivariate normal distribution becomes:

$$p(\mathbf{x}) = \frac{1}{\sqrt{(2\pi)^n |\boldsymbol{\Sigma}|}} \cdot \exp\left[-\frac{1}{2} \cdot (\mathbf{x} - \boldsymbol{\mu})^T \cdot \boldsymbol{\Sigma}^{-1} \cdot (\mathbf{x} - \boldsymbol{\mu})\right]$$

| property | description | |-----|-----| | $\boldsymbol{\Sigma}$ | covariance matrix $\in \mathbb{R}^{n \times n}$ | | $|\boldsymbol{\Sigma}|$ | determinant of covariance matrix | | $\boldsymbol{\Sigma}^{-1}$ | inverse of covariance matrix | | \mathbf{x} | vector $\in \mathbb{R}^{n \times 1}$ | | $\boldsymbol{\mu}$ | vector $\in \mathbb{R}^{n \times 1}$ of mean values of each component of vector \mathbf{x} |

But before we look at some applications which involve the multivariate normal distribution we need to know some more details about the concept of covariance and its generalisation to its matrix equivalent.

The material is covered in a separate notebook.

In []: