## Multivariate Gaussian Draw function

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## 1 Theorm of linear translation of Gaussian Distribution

It is possible to transform a multivariate normal distribution into a new normal distribution with an affine transformation. More specifically if  $X \sim \mathcal{N}(\mu_x, \Sigma_x)$  and Y = LX + C with L a linear transformation and C a vector then  $Y \sim \mathcal{N}(\mu_y, \Sigma_y)$  with  $\mu_y = L\mu_x + C$  and  $\Sigma_y = L\Sigma_x L^T$ 

## 1.1 Drawing from Multivariate Gaussian Distribution

An alternative way of presenting Multivariate Gaussian variable with  $p(\underline{x}) = \mathcal{N}(\underline{x}; \mu, \underline{\Sigma})$  is as below:

$$\underline{x} = \underline{\mu} + \underline{\underline{\Sigma}}^{1/2} \underline{\epsilon}$$

with

$$\underline{\epsilon} \sim \mathcal{N}(\underline{0}, \underline{\underline{I}})$$

This is based on the linear transformation theorm presented above. This makes sampling easier as we sample first from  $\underline{\epsilon}$  and then apply the linear transformation. Sampling from  $\underline{\epsilon}$  is easier because each variable in  $\underline{\epsilon}$  is independent from all other variables. This means we can just draw each variable separately.

## 1.2 Example for bivariate Gaussian:

Let's start with an example of drawing 100 samples from variable X with the following bivariate Gaussian distribution:

$$X \sim \mathcal{N}\left(\left[\begin{array}{c} 0\\1 \end{array}\right], \left[\begin{array}{cc} 1 & 0.6\\0.6 & 1 \end{array}\right]\right)$$

The first step is to define mean and covariance matrices.

We can now use the linear transformation that we discussed above with  $\underline{\mu} = \begin{bmatrix} 0 \\ 2 \end{bmatrix}$  and  $\underline{\underline{\Sigma}}^{1/2}$  as the choloskey decomposition of the covariance matrix:  $\begin{bmatrix} 0.3 & -1 \\ -1 & 5 \end{bmatrix}$ 

```
[48]: L = np.linalg.cholesky(cov) # get the choloskey decomposition of cov matrix
d = cov.shape[0] # get the dimension
n = 100 # total samples
SN = np.random.normal(size=(d,n)) # draw n samples from the standard normal

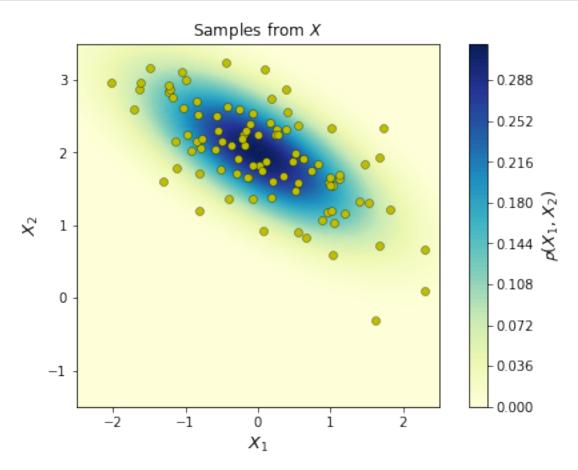
distribution, each for d minsions (i.e. d independent sample due to 0

correlation in underlined epsilon)
X = L.dot(SN)+mean
```

Let's now plot the sample and distribution

```
[50]: def multivariate_normal(x, d, mean, covariance):
          """pdf of the multivariate normal distribution."""
          import numpy as np
          xm = x - mean
          return (1. / (np.sqrt((2 * np.pi)**d * np.linalg.det(covariance))) *
                  np.exp(-(np.linalg.solve(covariance, xm).T.dot(xm)) / 2))
      def surface(mean, covariance, d):
          """A function to generate surface for density."""
          import numpy as np
          nb of x = 100 \# qrid size
          x1s = np.linspace(-5, 5, num=nb_of_x)
          x2s = np.linspace(-5, 5, num=nb_of_x)
          x1, x2 = np.meshgrid(x1s, x2s) # Generate grid
          pdf = np.zeros((nb_of_x, nb_of_x))
          # Fill the cost matrix for each combination of weights
          for i in range(nb_of_x):
              for j in range(nb_of_x):
                  pdf[i,j] = multivariate_normal(
                      np.matrix([[x1[i,j]], [x2[i,j]]]),
                      d, mean, covariance)
          return x1, x2, pdf # x1, x2, pdf(x1,x2)
      import matplotlib
      import matplotlib.pyplot as plt
      from matplotlib import cm # Colormaps
      import matplotlib.gridspec as gridspec
      fig, ax = plt.subplots(figsize=(8, 5))
      # Plot bivariate distribution
      x1, x2, p = surface(mean, cov, d)
      con = ax.contourf(x1, x2, p, 100, cmap=cm.YlGnBu)
      # Plot samples
      ax.plot(X[0,:], X[1,:], 'yo', alpha=1,
              markeredgecolor='k', markeredgewidth=0.3)
      ax.set_xlabel('$X_1$', fontsize=13)
      ax.set_ylabel('$X_2$', fontsize=13)
```

```
ax.axis([-2.5, 2.5, -1.5, 3.5])
ax.set_aspect('equal')
ax.set_title('Samples from $X$')
cbar = plt.colorbar(con)
cbar.ax.set_ylabel('$\mathcal{p}(X_1, X_2)$', fontsize=13)
plt.show()
#
```



[]: