

# Homework7

June 16, 2016

```
In [3]: import numpy as np
import matplotlib.pyplot as plt
from scipy.io import loadmat as loadmat
% matplotlib inline
```

## 1 7.1 Natural Gradient (3 points)

- (a) Extend your code from the previous problem sheet to get an ICA-learning scheme based on the natural gradient with a learning rate  $\eta$  that decays slowly to 0 (e.g.  $\eta_{t+1} = \lambda \eta_t$  with  $\lambda \approx 1$ ,  $\lambda < 1$ ). Note that depending on  $\lambda$  you have to iterate over the (shuffled) data more than once for proper convergence.

```
In [15]: def fhat(y):
    return 1 / (1 + np.exp(-y))

def fhatpp_fhatp(y):
    return 1 - 2*fhat(y)

def dWnatural(eta, W, x):
    #x0 x1
    #x0 x1
    #xv = np.vstack( (x.T, x.T) )
    xv = np.vstack( (x.T, x.T, x.T) )
    #x0 x0
    #x1 x1
    #xh = np.hstack( (x.reshape(-1,1), x.reshape(-1,1)) )
    xh = np.hstack( (x.reshape(-1,1), x.reshape(-1,1), x.reshape(-1,1)) )
    #W-1 + f(W . xh) * xv
    subtotal = (np.eye(W.shape[0]) + np.multiply( fhatpp_fhatp(np.dot(W, xh)), xv))
    #return eta * np.dot(subtotal, np.dot(W.T, W))
    return eta * np.dot(subtotal, W)

def buildcorrelations(N, s, x):
    p = np.zeros((N, N))
    for i in range(N):
        for j in range(N):
            #print("i: "+str(i)+" j: "+str(j)+" corr:\n"+str((np.cov(s[i], x[j])) / (np.std(s[i]
            p[i, j] = (np.cov(s[i], x[j]) / (np.std(s[i]) * np.std(x[j])))[0,1]
    return p

def gradientdescent(A, W, s, eta, l):
```

```

print("Mixing Matrix:\n"+str(A)+"\nDet: "+str(np.linalg.det(A)))
x = np.dot(A, s)

#(c) Remove the temporal structure by permuting the columns of the  $N \times p$  matrix  $X$  randomly
xs = np.random.permutation(x)

#(e) Center the data to zero mean.
#print( np.mean(x, axis=1).shape )
x_mean = np.mean(xs, axis=1).reshape(-1,1)
xsc = xs - x_mean

#(f) Initialize the unmixing matrix  $W$  with random values.
#s = W.x, s: Nx p, x: Nx p -> W: Nx N
#W = np.random.uniform(0, 1, (N, N))

#(c) Choose a suitable learning rate  $\eta$  and apply both versions to the data to unmix the so
epsilon = .000000001
Wb = W.copy()
Wbbs = W.copy()
BConverged = False
BbsConverged = False
etan = eta

print("Initial W:\n"+str(W))
for n in range(10):
    for t in range(0, xsc.shape[1]):
        xa = xsc[:,t]
        #xa = xsc[:,np.random.randint(xsc.shape[1])]
        etan = etan*l

        #Natural
        if not BConverged:
            natural = dWnatural(etan, Wb, xa)
            if np.absolute(natural).sum() > epsilon:
                Wb = Wb + natural
            else:
                BConverged = True
                print("(b) converged at: "+str(n*t))

        #Natural with Bell-Sejnowski regularization
        if not BbsConverged:
            naturalbs = dWnatural(etan, Wbbs, xa)
            for n in range(N):
                naturalbs[n, n] = 0
            if np.absolute(naturalbs).sum() > epsilon:
                Wbbs = Wbbs + naturalbs
                for n in range(N):
                    Wbbs[n, n] = 1
            else:
                BbsConverged = True
                print("(b.bs) Bell-Sejnowski converged at: "+str(n*t))
        else:
            pass
return Wb, Wbbs, x, x_mean

```

```

def myplot(Wb, Wbbs, x, x_mean, s):
    #retrieve shats - ^s
    shatb = np.dot(Wb, x)
    shatb_decentered = shatb + x_mean
    shatbbs = np.dot(Wbbs, x)
    shatbbs_decentered = shatbbs + x_mean

    #Check correlations, to check whcih source goes to which output channel
    mixcors = buildcorrelations(N, s, x)
    naturalcors = buildcorrelations(N, shatb_decentered, s)
    naturalbscors = buildcorrelations(N, shatbbs_decentered, s)

    isflippedmixes = (mixcors[0,0] < mixcors[0,1]) and (mixcors[1,0] > mixcors[1,1])
    isflippednatural = (naturalcors[0,0] < naturalcors[0,1]) and (naturalcors[1,0] > naturalco
    isflippednaturalbs = (naturalbscors[0,0] < naturalbscors[0,1]) and (naturalbscors[1,0] > n

    ys = [(s, 'Original Sources', False),
           (x, 'Mixed Sources', isflippedmixes),
           (shatb_decentered, '(b) - Natural Gradient Descent', isflippednatural),
           (shatbbs_decentered, '(b) - Natural Gradient Descent Bell-Sejnowski', isflippednatur

    ylimits = [-14,14]
    f, axarr = plt.subplots(len(ys), 3)
    f.set_size_inches(30, len(ys)*10, forward=False)

    axarr[0, 0].set_title("Source 1")
    axarr[0, 1].set_title("Source 2")
    axarr[0, 2].set_title("Source 3")

    for r in range(len(ys)):
        #axarr[r, 0].set_ylim(ylimits)
        #axarr[r, 1].set_ylim(ylimits)
        #axarr[r, 2].set_ylim(ylimits)
        axarr[r, 0].set_ylabel(ys[r][1])

        axarr[r, 0].plot(range(ys[r][0].shape[1]), ys[r][0][0])
        axarr[r, 1].plot(range(ys[r][0].shape[1]), ys[r][0][1])
        axarr[r, 2].plot(range(ys[r][0].shape[1]), ys[r][0][2])
        axarr[r, 0].set_title("Output 1")
        axarr[r, 1].set_title("Output 2")
        axarr[r, 2].set_title("Output 3")

    plt.show()

```

- (b) Use the two sound signals from the last problem sheet and add (as third source s3) an additional “noise” source (normally distributed random numbers with a standard deviation similar to the two signals). Mix the signals using a mixing matrix of your choice and apply your ICA-algorithm. Plot the Mixed Sounds and recovered Sources

```

In [68]: s1 = np.loadtxt("sounds/sound1.dat")
         s2 = np.loadtxt("sounds/sound2.dat")
         s3 = np.random.normal(0, 1, len(s1))
         s = np.array([s1, s2, s3])
         print("Standard Deviation s1: "+str(np.std(s[0])))
         print("Standard Deviation s2: "+str(np.std(s[1])))

```

```

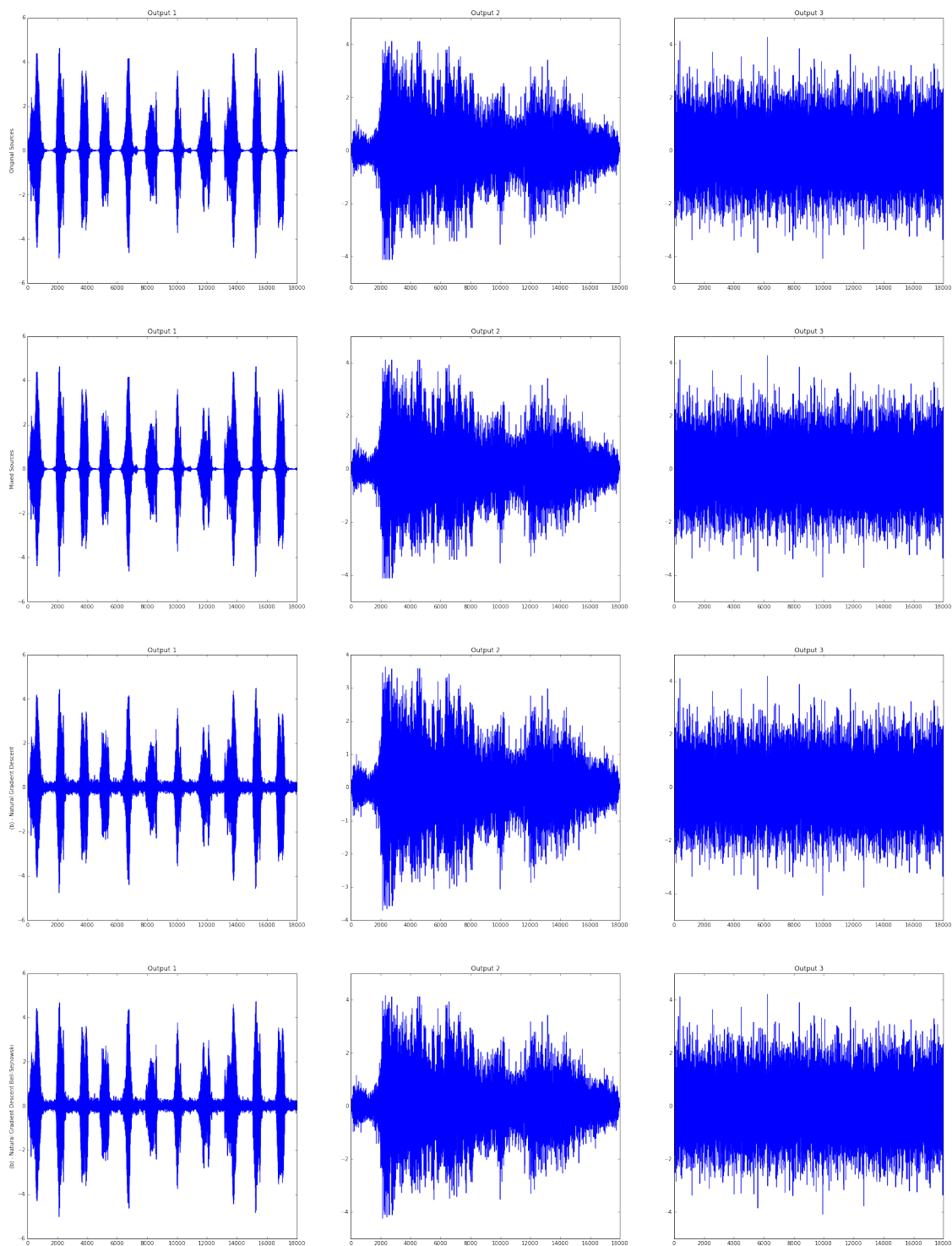
print("Standard Deviation s3: "+str(np.std(s[2])))
N = s.shape[0]

A = np.eye(N)
W = np.random.uniform(0, .1, (N, N))
W[0,0] = np.random.uniform(.8, 1)
W[1,1] = np.random.uniform(.8, 1)
W[2,2] = np.random.uniform(.8, 1)

Wb, Wbbs, x, x_mean = gradientdescent(A, W, s, .0000001, .9999)
print("Wb:")
print(Wb)
print("Wbbs:")
print(Wbbs)
myplot(Wb, Wbbs, x, x_mean, s)

Standard Deviation s1: 0.998545937615
Standard Deviation s2: 0.998854345166
Standard Deviation s3: 0.986100852986
Mixing Matrix:
[[ 1.  0.  0.]
 [ 0.  1.  0.]
 [ 0.  0.  1.]]
Det: 1.0
Initial W:
[[ 0.94736096  0.06200152  0.09854062]
 [ 0.04283364  0.86946837  0.00626526]
 [ 0.05738817  0.05842622  0.99659339]]
(b.bs) Bell-Sejnowski converged at: 12422
(b) converged at: 35552
Wb:
[[ 0.94790721  0.06201358  0.09855416]
 [ 0.04284101  0.87002072  0.00626386]
 [ 0.05739633  0.05844015  0.99721645]]
Wbbs:
[[ 1.          0.06200926  0.0985515 ]
 [ 0.04283859  1.          0.00626381]
 [ 0.05739381  0.05843595  1.          ]]

```



(c) Do the same analysis but adding a different “noise”-source (e.g. Laplace distributed) instead of the normal one.

```
In [69]: s1 = np.loadtxt("sounds/sound1.dat")
         s2 = np.loadtxt("sounds/sound2.dat")
```

```

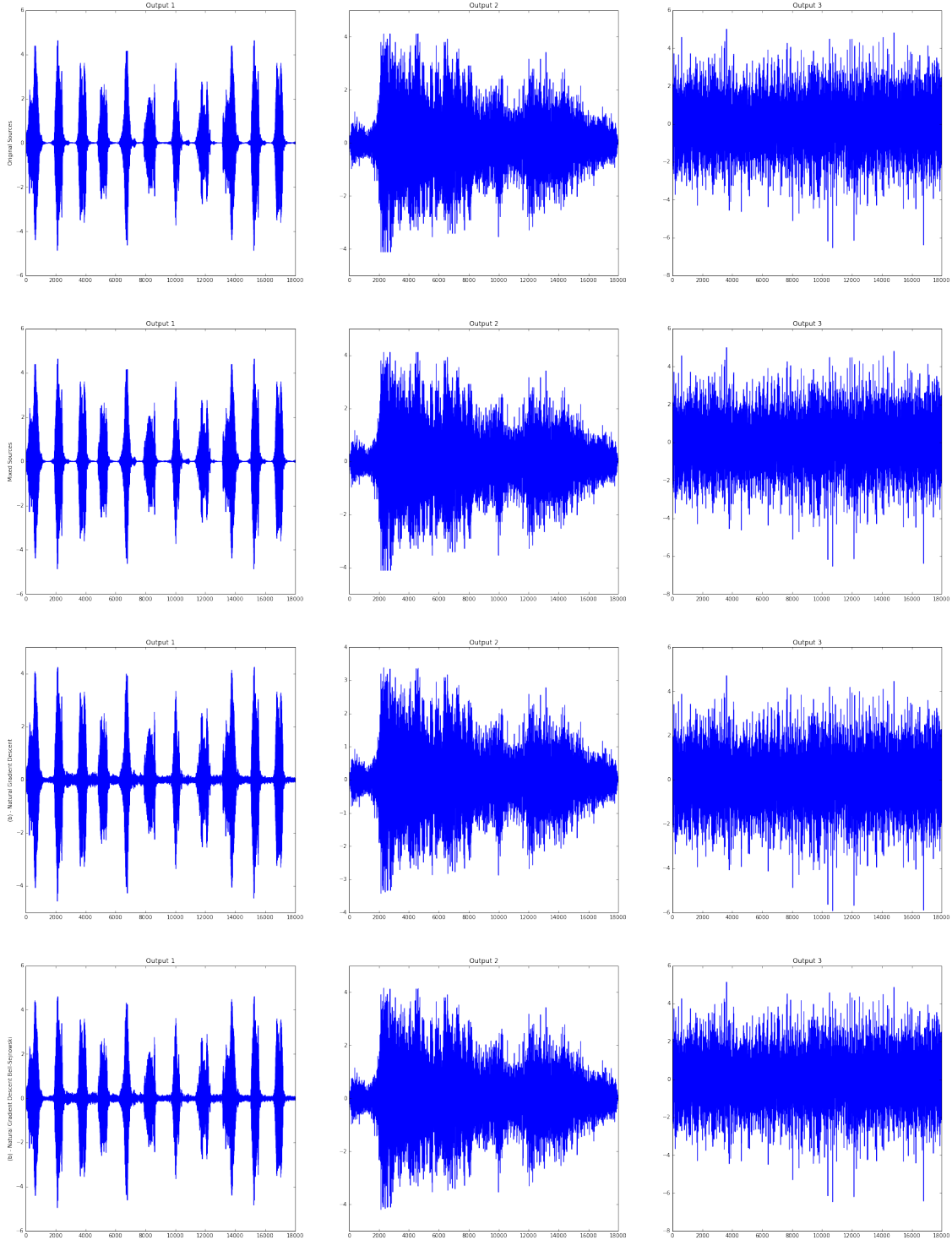
s3 = np.random.laplace(0.0, 0.7, len(s1))
s = np.array([s1, s2, s3])

print("Standard Deviation s1: "+str(np.std(s[0])))
print("Standard Deviation s2: "+str(np.std(s[1])))
print("Standard Deviation s3: "+str(np.std(s[2])))
N = s.shape[0]
A = np.eye(N)
W = np.random.uniform(0, .1, (N, N))
W[0,0] = np.random.uniform(.8, 1)
W[1,1] = np.random.uniform(.8, 1)
W[2,2] = np.random.uniform(.8, 1)

Wb, Wbbs, x, x_mean = gradientdescent(A, W, s, .0000001, .9999)
print("Wb:")
print(Wb)
print("Wbbs:")
print(Wbbs)
myplot(Wb, Wbbs, x, x_mean, s)

Standard Deviation s1: 0.998545937615
Standard Deviation s2: 0.998854345166
Standard Deviation s3: 0.980715089168
Mixing Matrix:
[[ 1.  0.  0.]
 [ 0.  1.  0.]
 [ 0.  0.  1.]]
Det: 1.0
Initial W:
[[ 0.92268291  0.06539629  0.02809297]
 [ 0.02891274  0.81202717  0.01726697]
 [ 0.07903204  0.09890617  0.91645418]]
(b.bs) Bell-Sejnowski converged at: 17768
(b) converged at: 32670
Wb:
[[ 0.92328144  0.06540724  0.02809883]
 [ 0.028915    0.81256614  0.01726954]
 [ 0.0790489   0.09892605  0.91699215]]
Wbbs:
[[ 1.          0.06539828  0.02809692]
 [ 0.02891187  1.          0.01726796]
 [ 0.07904427  0.09891487  1.          ]]

```



## 2 Assignment 7.3

The file `distrib.mat` contains three toy datasets (uniform, normal, laplacian), each 10000 samples of 2 sources. Do the following for each dataset (which can be read for example using Python with `loadmat` from `scipy.io`):

7.2 The moment generating function is defined as:

$$M_X(t) = E(e^{tX}) = \int_{-\infty}^{\infty} e^{tx} f(x) dx \quad \left. \begin{array}{l} f(x) \text{ as} \\ \text{probability} \\ \text{density} \end{array} \right\}$$

$$\frac{d^k}{dt^k} M_X(t) = \frac{d^k}{dt^k} E(e^{tX}) = E(X^k e^{tX})$$

$$E(X^k e^{tX}) \Big|_{t=0} = E(X^k) \quad \left| \begin{array}{l} k\text{-th moment} \end{array} \right.$$

moment generating functions:

$$\text{Laplace: } M_X(t) = \int_{-\infty}^{\infty} e^{tx} \frac{1}{\sqrt{2\sigma}} e^{-\frac{|x-\mu|}{\sigma}} dx = \frac{e^{\mu t}}{1 - \sigma^2 t^2}$$

$$\text{Gauß: } M_X(t) = \int_{-\infty}^{\infty} e^{tx} \frac{1}{\sqrt{2\sigma^2\pi}} e^{-\frac{(x-\mu)^2}{2\sigma^2}} dx = e^{\mu t + \frac{\sigma^2 t^2}{2}}$$

$$\text{Uniform: } M_X(t) = \int_a^b e^{tx} \frac{1}{b-a} dx = \left[ \frac{1}{t(b-a)} e^{tx} \right]_a^b = \frac{e^{bt} - e^{at}}{t(b-a)}$$

The centered moment generating function:

$$M_{X_c}(t) = E(e^{t(X-\mu)}) = E(e^{tX}) \cdot E(e^{-t\mu}) = M_X(t) \cdot e^{-t\mu}$$

The standardised moments are then calculated as described in the exercise.



	Laplace ( $\mu, b$ )	Gauß ( $\mu, \sigma$ )	uniform ( $a, b$ )
$\langle x \rangle$	$\mu$	$\mu$	$\frac{1}{2}(a+b)$
$\langle x^2 \rangle_c$	$2\sigma^2$	$\sigma^2$	$\frac{1}{12}(b-a)^2$
$\langle x^3 \rangle_s$	0	0	0
$\langle x^4 \rangle_s$	6	0	$-\frac{6}{5}$

```
In [19]: laplacian = loadmat("distrib.mat")["laplacian"]
        normal = loadmat("distrib.mat")["normal"]
        uniform = loadmat("distrib.mat")["uniform"]
```

(a) Apply the following mixing matrix A to the original data s:

```
4 3
2 1
```

```
In [20]: A = np.array([[4,3],[2,1]])
        x1 = np.dot(A, laplacian)
        xn = np.dot(A, normal)
        xu = np.dot(A, uniform)
```

(b) Center the mixed data to zero mean.

```
In [21]: #keep as pxd
        xlc = (x1 - np.mean(x1, axis=1).reshape(-1,1)).T
        xnc = (xn - np.mean(xn, axis=1).reshape(-1,1)).T
        xuc = (xu - np.mean(xu, axis=1).reshape(-1,1)).T
```

(c) Decorrelate the data by applying principal component analysis (PCA) and project them onto the principal components (PCs).

```
In [46]: #Scatter the centered data, dxd
        S1 = np.dot(xlc.T, xlc)
        Sn = np.dot(xnc.T, xnc)
        Su = np.dot(xuc.T, xuc)

        #Solve the eigensystem
        D1, U1 = np.linalg.eigh(S1)
        Dn, Un = np.linalg.eigh(Sn)
        Du, Uu = np.linalg.eigh(Su)

        #reverse the orders
        D1 = D1[::-1]
        Dn = Dn[::-1]
        Du = Du[::-1]
        U1 = U1.T[::-1]
        Un = Un.T[::-1]
        Uu = Uu.T[::-1]

        #Projections
        xlp = (np.dot(U1, xlc.T)).T
        xnp = (np.dot(Un, xnc.T)).T
        xup = (np.dot(Uu, xuc.T)).T
```

(d) Scale the data to unit variance in each PC direction (now the data is whitened or sphered).

```
In [47]: #scale to unit variance
        xlpuv = xlp / np.std(xlp, axis=0)
        xnpuv = xnp / np.std(xnp, axis=0)
        xupuv = xup / np.std(xup, axis=0)
```

(e) Rotate the data by different angles  $\theta$  and calculate the kurtosis1 empirically for each dimension:

$$x\theta = \begin{vmatrix} \cos(\theta) & -\sin(\theta) \\ \sin(\theta) & \cos(\theta) \end{vmatrix} x$$

$$\theta = 0, \pi, \dots, 2\pi$$

```
In [62]: #compute all rotation amounts
thetas = [(np.pi*x)/50 for x in range(0, 100)]
#compute rotation matrices
rs = [np.array([[np.cos(theta), -np.sin(theta)], [np.sin(theta), np.cos(theta)]]) for theta in thetas]
#compute kurtosises
kurtls = np.array([np.mean(np.power(np.dot(r, xlpuv.T), 4), axis=1)-3 for r in rs])
kurtns = np.array([np.mean(np.power(np.dot(r, xnpuv.T), 4), axis=1)-3 for r in rs])
kurtus = np.array([np.mean(np.power(np.dot(r, xupuv.T), 4), axis=1)-3 for r in rs])
```

```
(100, 2)
```

(f) Find the minimum and maximum kurtosis value for the first dimension and rotate the data accordingly.

```
In [63]: kurtlminidx = np.argmin(kurtls[:,0])
kurtlmaxidx = np.argmax(kurtls[:,0])

kurtnminidx = np.argmin(kurtns[:,0])
kurtnmaxidx = np.argmax(kurtns[:,0])

kurtuminidx = np.argmin(kurtus[:,0])
kurtumaxidx = np.argmax(kurtus[:,0])
```

- Plot the original dataset (sources) and the mixed dataset after the steps (a), (b), (c), (d), and (f) as a scatter plot and display the respective marginal histograms. For step (e) plot the kurtosis value as a function of angle for each dimension.

```
In [65]: bins = 90
f, axarr = plt.subplots(11, 3)
f.set_size_inches(10, 39, forward=False)

r = 0
axarr[r, 0].set_title("Laplacian")
axarr[r, 1].set_title("Normal")
axarr[r, 2].set_title("Uniform")

axarr[r, 0].set_ylabel("Original")
axarr[r, 0].scatter(laplacian[0], laplacian[1], color="firebrick")
axarr[r, 1].scatter(normal[0], normal[1], color="firebrick")
axarr[r, 2].scatter(uniform[0], uniform[1], color="firebrick")

r = r+1
axarr[r, 0].set_ylabel("(a) After mixing")
axarr[r, 0].scatter(xl[0], xl[1], color="slateblue")
axarr[r, 1].scatter(xn[0], xn[1], color="slateblue")
axarr[r, 2].scatter(xu[0], xu[1], color="slateblue")

r = r+1
axarr[r, 0].set_ylabel("(b) After centering")
axarr[r, 0].scatter(xlc.T[0], xlc.T[1], color="slateblue")
axarr[r, 1].scatter(xnc.T[0], xnc.T[1], color="slateblue")
axarr[r, 2].scatter(xuc.T[0], xuc.T[1], color="slateblue")
```

```

r = r+1
axarr[r, 0].set_ylabel("(c) PCs on centered")
axarr[r, 0].scatter(xlc.T[0], xlc.T[1], color="slateblue", alpha=.2)
axarr[r, 0].plot([0, U1[0,0]], [0,U1[0,1]], color="deeppink", linewidth=2)
axarr[r, 0].plot([0, U1[1,0]], [0,U1[1,1]], color="deeppink", linewidth=2)
axarr[r, 1].scatter(xnc.T[0], xnc.T[1], color="slateblue", alpha=.2)
axarr[r, 1].plot([0, U1[0,0]], [0,U1[0,1]], color="deeppink", linewidth=2)
axarr[r, 1].plot([0, U1[1,0]], [0,U1[1,1]], color="deeppink", linewidth=2)
axarr[r, 2].scatter(xuc.T[0], xuc.T[1], color="slateblue", alpha=.2)
axarr[r, 2].plot([0, U1[0,0]], [0,U1[0,1]], color="deeppink", linewidth=2)
axarr[r, 2].plot([0, U1[1,0]], [0,U1[1,1]], color="deeppink", linewidth=2)

r = r+1
axarr[r, 0].set_ylabel("(c) Projections")
axarr[r, 0].scatter(xlp.T[0], xlp.T[1], color="slateblue")
axarr[r, 1].scatter(xnp.T[0], xnp.T[1], color="slateblue")
axarr[r, 2].scatter(xup.T[0], xup.T[1], color="slateblue")

r = r+1
axarr[r, 0].set_ylabel("(d) Scale to unit variance\nWhiten")
axarr[r, 0].scatter(xlpuv.T[0], xlpuv.T[1], color="slateblue")
axarr[r, 1].scatter(xnpuv.T[0], xnpuv.T[1], color="slateblue")
axarr[r, 2].scatter(xupuv.T[0], xupuv.T[1], color="slateblue")

r = r+1
axarr[r, 0].set_ylabel("(e) Kurtosis vs Angle")
axarr[r, 0].plot(range(len(kurtls[:,0])), kurtls[:,0], color="seagreen")
axarr[r, 0].plot(range(len(kurtls[:,1])), kurtls[:,1], color="magenta")
axarr[r, 1].plot(range(len(kurtns[:,0])), kurtns[:,0], color="seagreen")
axarr[r, 1].plot(range(len(kurtns[:,1])), kurtns[:,1], color="magenta")
axarr[r, 2].plot(range(len(kurtus[:,0])), kurtus[:,0], color="seagreen")
axarr[r, 2].plot(range(len(kurtus[:,1])), kurtus[:,1], color="magenta")

r = r+1
axarr[r, 0].set_ylabel("(f) Min Kurtosis")
axarr[r, 0].scatter(np.dot(rs[kurtlminidx], xlpuv.T)[0], np.dot(rs[kurtlminidx], xlpuv.T)[1], color="seagreen")
axarr[r, 1].scatter(np.dot(rs[kurtnminidx], xnpuv.T)[0], np.dot(rs[kurtnminidx], xnpuv.T)[1], color="seagreen")
axarr[r, 2].scatter(np.dot(rs[kurtuminidx], xupuv.T)[0], np.dot(rs[kurtuminidx], xupuv.T)[1], color="seagreen")

r = r+1
axarr[r, 0].set_ylabel("(f) Min Kurtosis")
histodata = np.histogram(np.dot(rs[kurtlminidx], xlpuv.T), bins)
axarr[r, 0].bar(np.arange(len(histodata[0])), histodata[0], color="seagreen")
histodata = np.histogram(np.dot(rs[kurtnminidx], xnpuv.T), bins)
axarr[r, 1].bar(np.arange(len(histodata[0])), histodata[0], color="seagreen")
histodata = np.histogram(np.dot(rs[kurtuminidx], xupuv.T), bins)
axarr[r, 2].bar(np.arange(len(histodata[0])), histodata[0], color="seagreen")

r = r+1
axarr[r, 0].set_ylabel("(f) Max Kurtosis")
axarr[r, 0].scatter(np.dot(rs[kurtlmaxidx], xlpuv.T)[0], np.dot(rs[kurtlmaxidx], xlpuv.T)[1], color="seagreen")
axarr[r, 1].scatter(np.dot(rs[kurtnmaxidx], xnpuv.T)[0], np.dot(rs[kurtnmaxidx], xnpuv.T)[1], color="seagreen")
axarr[r, 2].scatter(np.dot(rs[kurtumaxidx], xupuv.T)[0], np.dot(rs[kurtumaxidx], xupuv.T)[1], color="seagreen")

```

```

r = r+1
axarr[r, 0].set_ylabel("(f) Max Kurtosis")
histodata = np.histogram(np.dot(rs[kurtlmaxidx], xlpuv.T), bins)
axarr[r, 0].bar(np.arange(len(histodata[0])), histodata[0], color="seagreen")
histodata = np.histogram(np.dot(rs[kurtnmaxidx], xnpuv.T), bins)
axarr[r, 1].bar(np.arange(len(histodata[0])), histodata[0], color="seagreen")
histodata = np.histogram(np.dot(rs[kurtumaxidx], xupuv.T), bins)
axarr[r, 2].bar(np.arange(len(histodata[0])), histodata[0], color="seagreen")

plt.show()

```

