

pract6

May 23, 2017

```
In [105]: import numpy as np
import matplotlib.pyplot as plt
import networkx as nx
import seaborn
```

1 Practica 6 Redes Neuronales

1.1 Miguel Angel Carvajal

1.1.1 Ejercicio 1

Defino la matrix de covarianza Σ

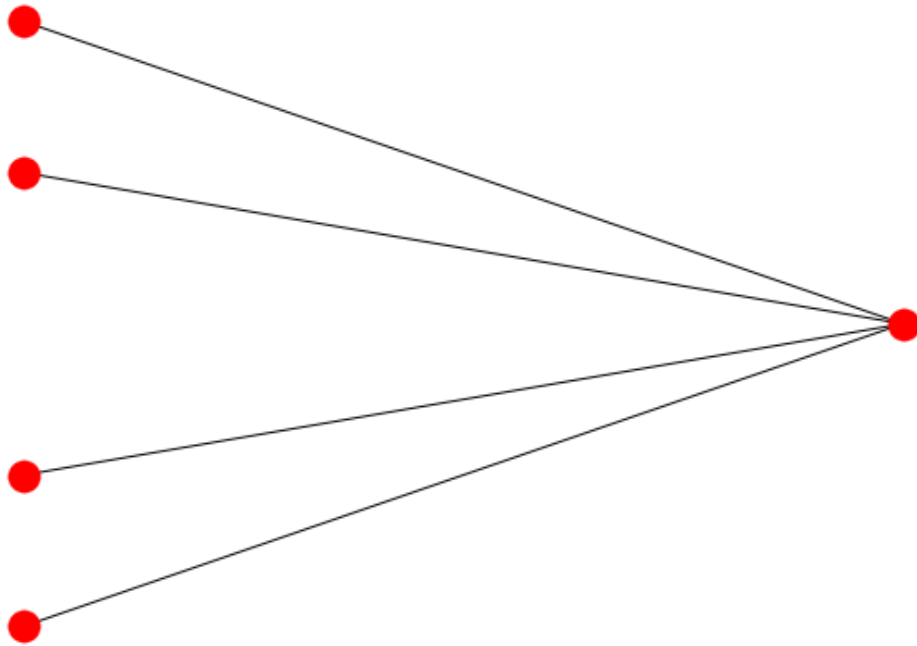
```
In [6]: cov_mat = np.ones((4,4))
np.fill_diagonal(cov_mat,2)
print cov_mat
```

```
[[ 2.  1.  1.  1.]
 [ 1.  2.  1.  1.]
 [ 1.  1.  2.  1.]
 [ 1.  1.  1.  2.]]
```

```
In [7]: # learning speed
eta = 0.0001
```

Draw the network achitecture

```
In [106]: nn = nx.Graph()
nn.add_nodes_from([1,2,3,4,5])
nn.add_edges_from([(1,5),(2,5),(3,5),(4,5)])
nx.draw(nn, pos={1:(0,2),2:(0,1),3:(0,-1),4:(0,-2),5:(1,0)})
plt.plot()
plt.show()
```



1.2 Run the simulation

The input data is generated with a multigaussian probability given by

$$P(\bar{\xi}) = \frac{1}{(2\pi)^2 \sqrt{\det(\xi)}} \exp\left(-\frac{1}{2} \bar{\xi}^T \Sigma^{-1} \bar{\xi}\right)$$

```

In [9]: np.random.seed(1)
        N_in = 4 # input size
        N_out = 1 # output size
        # init weights
        weights = (2*np.random.rand(4,1)-1) /10
        mean_val = np.zeros(N_in)
        p = 30000
        # print weights.shape
        D_in = np.random.multivariate_normal(mean_val, cov_mat, p)
        for i in range(0,p):
            D_in_i = D_in[i:i+1,:]
            # print D_in_i.shape
            V = np.dot(weights.T,D_in_i.T)
            error = eta * V * (D_in_i.T - V*weights)
            # print error.shape
            # print np.mean(np.abs(error))
            weights += error
        print weights
  
```

```

[[-0.49392035]
 [-0.49219265]
 [-0.51130988]
 [-0.50250419]]

```

Check that the covered weight has norm 1

```

In [10]: norm = np.sum(weights**2)
         print norm

```

```

1.0001591745

```

Diagonalize the covarianza matriz and find the biggest eigenvalue

```

In [11]: vals, vectors = np.linalg.eig(cov_mat)
         print vals

```

```

[ 1.  5.  1.  1.]

```

The biggest eigenvalue is 5. and the eigenvector associated with it is

```

In [12]: y = vectors[:,1]
         print y
         print weights[:,0]

```

```

[ 0.5  0.5  0.5  0.5]
[-0.49392035 -0.49219265 -0.51130988 -0.50250419]

```

Comparando el resultado obtenido con el autovector de la matriz correspondiente al *mayor autovalor* se observa que estos tienen la misma dirección.

1.2.1 Ejercicio 2

Considere una red neuronal de Kohonen con dos neuronas de entrada. Utilice 10 neuronas de salida, dispuestas sobre una línea. Alimente a las neuronas de entrada con una distribución

Define the neighboring function

```

In [13]: def neigh(sigma):
         def neig_impl(i,k):
             return np.exp(-(i - k )**2/(2* sigma**2))
         return neig_impl

f1 = neigh(1)
assert f1(1,1) == 1 # must be one for the same neurons

```

```

In [14]: def scale(initial, final):
    """ Scale an interval (x1,x2) to (y1,y2) """
    x1, x2 = initial
    y1, y2 = final
    slope = (y2 - y1)/(x2 - x1)
    intercept = y1 - slope * x1
    def linf(x):
        assert x >= x1, "invalid input value " + str(x)
        assert x <= x2, "invalid input value " + str(x)
        retval = x * slope + intercept
        assert retval >= y1
        assert retval <= y2
        return retval
    return linf

In [25]: def to_radial_coords(point):
    """ convert a point from linear coordiates to radial coordiantes
        returns: numpy array with[r,theta]
        """
    assert point.shape == (2,), "error on shape: " + str(point.shape)
    x = point[0]
    y = point[1]
    r = np.sqrt(x**2 + y**2)
    assert r <= np.sqrt(2), point
    angle = np.arctan2(y,x)
    return np.array([r,angle])
def to_linear_coords(point):
    r = point[0]
    theta = point[1]
    x = r * np.cos(theta)
    y = r * np.sin(theta)
    return np.array([x,y])

In [115]: p = 3000 # num datapoints
    rmin = 0.9
    rmax = 1.1

    # define the dimensions of the input and output layer
    N1 = 2
    N2 = 10
    # initalize the weights
    weights = 2*np.random.rand(N1,N2)-1
    norms = np.linalg.norm(weights,axis=0)
    # normalize the weights
    for i in range(0,N2):
        weights[:,i] = weights[:,i] / norms[i]

    data_points = (2*np.random.rand(N1,p) -1)

```

```

# convert to radial coords
radial_scale = scale((0,np.sqrt(2)), (rmin,rmax))
angular_scale = scale((-np.pi,np.pi), (0, np.pi))
for i in range(0,p):
    data_points[:,i] = to_radial_coords(data_points[:,i])
    data_points[0,i] = radial_scale(data_points[0,i])
    data_points[1,i] = angular_scale(data_points[1,i])

assert np.max(data_points[0, :]) <= np.sqrt(2) ,np.max(data_points[0, :])
assert np.min(data_points[0, :]) >= rmin
assert np.min(data_points[1, :]) >= -np.pi
assert np.max(data_points[0, :]) <= rmax, "error for max boundary " + str
assert np.max(data_points[1, :]) <= np.pi

neighf = neigh(0.1)

for i in range(0,p):
    x = to_linear_coords(data_points[:,i])
    h = np.dot(weights.T,x)
    k = np.argmax(h) # get the pos of the max element
    for j in range(0,N2):
        error = neighf(j,k)*(x - weights[:,j])
        weights[:,j] += error
for i in range(0,N2):
    weights[:,i] = to_radial_coords(weights[:,i])

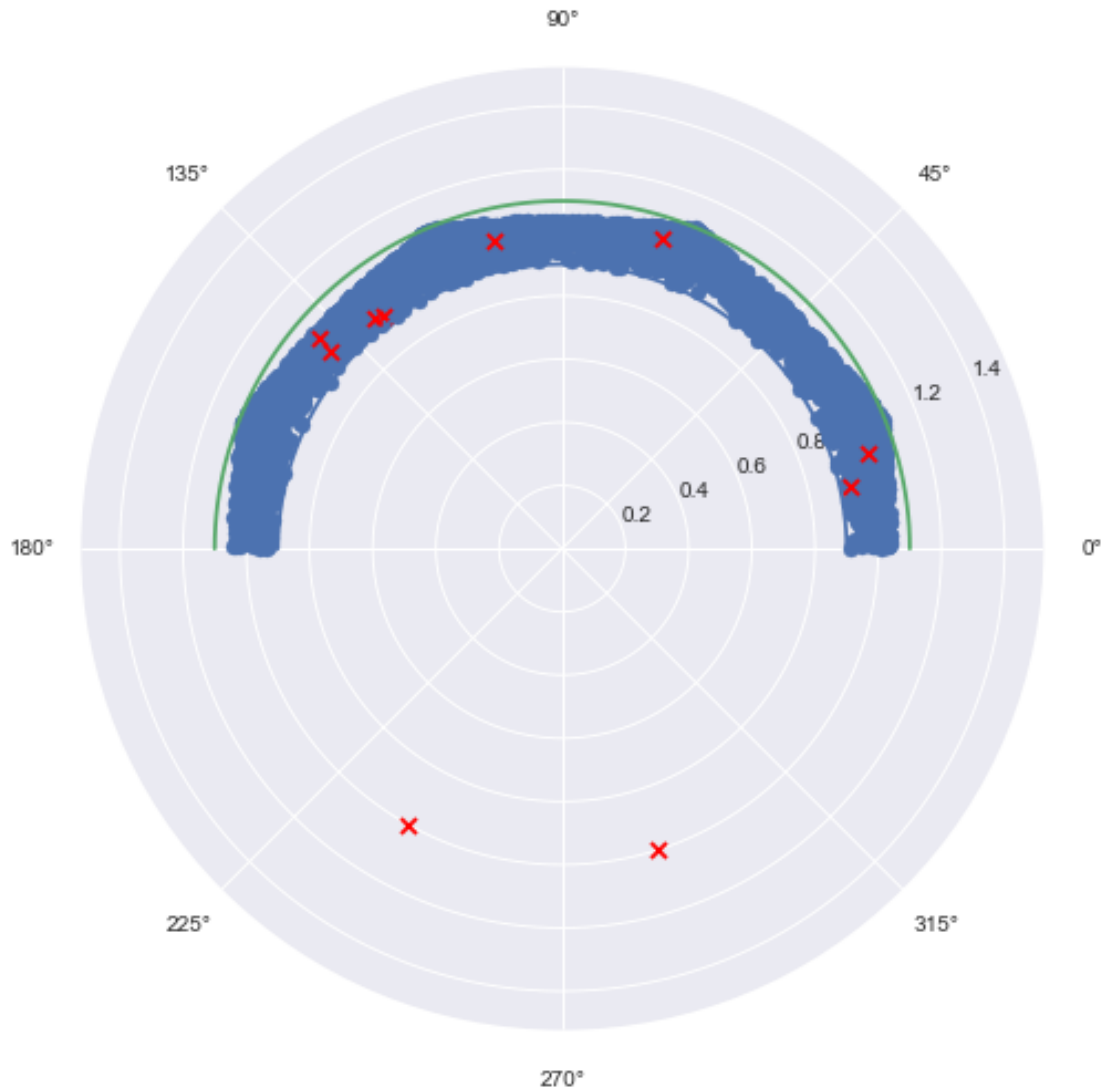
```

Plot the input data The data is distributed in an arc from 0 to π

```

In [117]: fig = plt.figure(figsize=(8,8))
fig.clear()
p1 = fig.add_subplot(111, projection='polar')
p1.scatter(data_points[1,:], data_points[0,:])
p1.scatter(weights[1,:], weights[0,:],color='r', marker='x')
plt.plot(np.pi*np.linspace(0,1,100), rmin*np.ones(100))
plt.plot(np.pi*np.linspace(0,1,100), rmax*np.ones(100))
plt.show()

```



Now wrap everything inside a function with p and σ as parameters

```
In [118]: def kohonen_simulation(sigma,p):
            rmin = 0.9
            rmax = 1.1

            # define the dimensions of the input and output layer
            N1 = 2
            N2 = 10
            # initialize the weights
            weights = 2*np.random.rand(N1,N2)-1
            norms = np.linalg.norm(weights,axis=0)
            # normalize the weights
            for i in range(0,N2):
```

```

        weights[:,i] = weights[:,i] / norms[i]

data_points = (2*np.random.rand(N1,p) -1)
# convert to radial coords
radial_scale = scale((0,np.sqrt(2)), (rmin,rmax))
angular_scale = scale((-np.pi,np.pi), (0, np.pi))
for i in range(0,p):
    data_points[:,i] = to_radial_coords(data_points[:,i])
    data_points[0,i] = radial_scale(data_points[0,i])
    data_points[1,i] = angular_scale(data_points[1,i])

assert np.max(data_points[0, :]) <= np.sqrt(2) ,np.max(data_points[0, :]) >= rmin
assert np.min(data_points[0, :]) >= rmin
assert np.min(data_points[1, :]) >= -np.pi
assert np.max(data_points[0, :]) <= rmax, "error for max boundary " +
assert np.max(data_points[1, :]) <= np.pi

neighf = neigh(sigma)

for i in range(0,p):
    x = to_linear_coords(data_points[:,i])
    h = np.dot(weights.T,x)
    k = np.argmax(h) # get the pos of the max element
    for j in range(0,N2):
        error = neighf(j,k)*(x - weights[:,j])
        weights[:,j] += error

for i in range(0,N2):
    weights[:,i] = to_radial_coords(weights[:,i])
return data_points, weights

def plot_simulation_results(pl,data_points, weights,sigma):
    """ Make a polar graph of the simulation results"""

    pl.scatter(data_points[1,:], data_points[0:],marker= 'o')
    pl.set_title( r'$\sigma= %f$'%(sigma))
    pl.scatter(weights[1,:], weights[0:],color='r', marker= 'x')

    plt.plot(np.pi*np.linspace(0,1,100), rmin*np.ones(100))
    plt.plot(np.pi*np.linspace(0,1,100), rmax*np.ones(100))

```

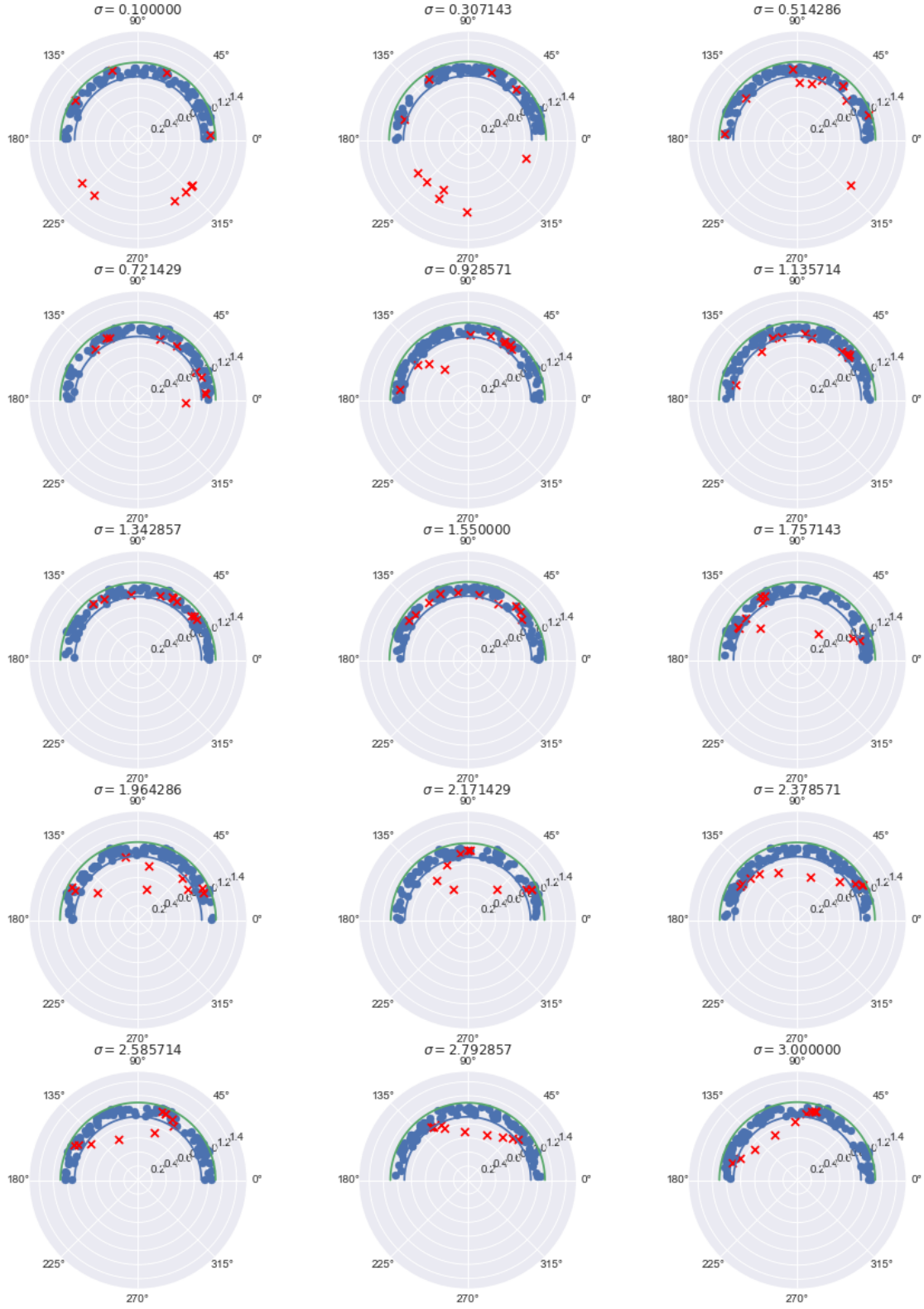
```

In [119]: n = 15
          sigmas = np.linspace(0.1,3,n)
          p = 100
          fig = plt.figure(figsize=(15,25))
          fig.clear()
          for k in range(0,n):
              data_points, weights = kohonen_simulation(sigmas[k],p)

```

```
axes = fig.add_subplot(int(n/3) + 1,3,k+1, projection='polar')
plot_simulation_results(axes,data_points, weights,sigmas[k])

plt.show()
```

En el grafico de arriba se muestra el resultado del aprendizaje para diferentes calores de σ . Se

puede ver que para valores pequeños y grandes se obtiene un resultado peor, mientras que para valores cercanos a 1 se obtienen los mejores resultados.