test

April 27, 2015

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In [16]:
In [1]: %%file Bayhiecluster.py
                      import numpy as np
                      from scipy.stats import multivariate_normal
                      import scipy
                      from scipy.special import gamma
                      def marginal_likelihood_NIW(points):
                                 n=len(points)
                                 p=len(points[0])
                                 mean_data = np.mean(points, axis=0)
                                  sum_squares = np.sum([np.array(np.matrix(x - mean_data).T * np.matrix(x - mean_data)) for x
                                 k0=3.0
                                  v0=10
                                  sigma0=np.eye(p)
                                  q*[0]=0um
                                 kn=k0+n
                                  vn=v0+n
                                  sigman = sigma0 + sum\_squares + (float(k0*n)/(k0+n))*np.dot((mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-mu0).reshape(p,1),(mean\_data-m
                                  gammadn=reduce(lambda x,y: x*y, map(lambda x:gamma((vn+1-x)*0.5),range(1,p+1)))
                                  gammad0=reduce(lambda x,y: x*y, map(lambda x:gamma((v0+1-x)*0.5),range(1,p+1)))
                                  mar_like=np.pi**(-p*n*0.5)*(np.linalg.det(sigma0))**(v0*0.5)*(k0/kn)**(0.5*p)*gammadn/(floa
                                 return mar_like
                      class bicluster:
                                  def __init__(self, point, left=None,right=None,probability=None,d=None,id=None):
                                            self.left = left
                                            self.right = right
                                            self.point = point
                                             self.id = id
                                             self.probability = probability
                                            self.d=d
                      def yezi(clust):
                                  if clust.left == None and clust.right == None :
                                            return [clust.id]
                                  return yezi(clust.left) + yezi(clust.right)
                      #cluster function
                      def bcluster(data,function) :
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alpha=3
            biclusters = [ bicluster(point = [data[i]], id = i ,probability=0.0000001,d=alpha) for i in
            flag = None;
            currentclusted = -1 #id for the new cluster
            if len(biclusters) == 1:
                clusters = [yezi(biclusters[i]) for i in range(len(biclusters))]
                return biclusters, clusters
            while(len(biclusters) > 1) :
                max_prob = 0;
                biclusters_len = len(biclusters)
                for i in range(biclusters_len-1) :
                    for j in range(i + 1, biclusters_len) :
                        #calculate P_H1: MC method
                        temp_cluster_points= biclusters[i].point + biclusters[j].point
                        P_H1=function(temp_cluster_points)
                        #P_H1=marginal_likelihood_DW(temp_cluster_points)
                        pi=float(scipy.misc.factorial(len(temp_cluster_points)-1))*alpha/(float(scipy.m
                        marginal_prob=pi*P_H1+(1-pi)*biclusters[i].probability*biclusters[j].probabilit
                        r = pi*P_H1/marginal_prob
                        if r > max_prob :
                            max_prob = r
                            flag = (i,j)
                if max_prob<0.5:
                    break
                bic1,bic2 = flag
                newpoint = biclusters[bic1].point + biclusters[bic2].point #combine the points of two c
                P_H1=function(newpoint)
                #P_H1=marginal_likelihood_DW(newpoint)
                newprob=pi*P_H1+(1-pi)*biclusters[bic1].probability*biclusters[bic2].probability
                newd=float(scipy.misc.factorial(len(newpoint)-1))*alpha+biclusters[bic1].d*biclusters[b
                newbic = bicluster(point=newpoint, left=biclusters[bic1], right=biclusters[bic2], proba
                currentclusted -= 1
                del biclusters[bic2]
                del biclusters[bic1]
                biclusters.append(newbic)
                clusters = [yezi(biclusters[i]) for i in range(len(biclusters))]
            return biclusters, clusters
Overwriting Bayhiecluster.py
In [2]: %%file test_Bayhiecluster.py
        import numpy as np
        from scipy.stats import multivariate_normal
        import scipy
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dim=len(data[0])

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from numpy.testing import assert_almost_equal
       from Bayhiecluster import bcluster, marginal_likelihood_NIW, yezi
        #parameters for test data
       mean0=(0,0)
       cov0=np.eye(2)
       mean2=(0,0)
       cov2=0.5*np.eye(2)
       mean3=(2,2)
       cov3=0.5*np.eye(2)
       mean4=(8,8)
       cov4=0.5*np.eye(2)
       def test_atleast_onecluster():
           for i in range(20):
               data2=np.random.multivariate_normal(mean2,cov2,3)
               data3=np.random.multivariate_normal(mean3,cov3,4)
               data4=np.random.multivariate_normal(mean4,cov4,3)
               sample_NIW=np.concatenate((data2, data3,data4), axis=0)
               k,l=bcluster(sample_NIW,marginal_likelihood_NIW)
               assert len(1) >= 1
       def test_max_cluster_number():
           for i in range(20):
               data2=np.random.multivariate_normal(mean2,cov2,3)
               data3=np.random.multivariate_normal(mean3,cov3,4)
               data4=np.random.multivariate_normal(mean4,cov4,3)
               sample_NIW=np.concatenate((data2, data3,data4), axis=0)
               k, l=bcluster(sample_NIW, marginal_likelihood_NIW)
               n=len(sample_NIW)
               assert len(1) <= n
       def test_if_all_the_same():
           data0=np.random.multivariate_normal(mean0,cov0,1)
           sample0=np.concatenate((data0, data0, data0, data0, data0, data0, data0, data0, data0, data
           k,l=bcluster(sample0,marginal_likelihood_NIW)
           assert len(1) == 1
       def test_only_one_point():
           special=[[np.random.multivariate_normal(mean0,cov0,1)]]
           k,l=bcluster(special,marginal_likelihood_NIW)
           assert len(1) == 1
Overwriting test_Bayhiecluster.py
In [3]: ! py.test
======= test session starts ==============
platform linux2 -- Python 2.7.9 -- py-1.4.25 -- pytest-2.6.3
collected 4 items
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from scipy.special import gamma

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