HW1

April 7, 2022

1 EM GMM

```
[71]: # module
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
import os

sns.set_theme(style="darkgrid")

current_path = os.getcwd()
output_path = os.path.join(current_path, 'output')
```

1.1

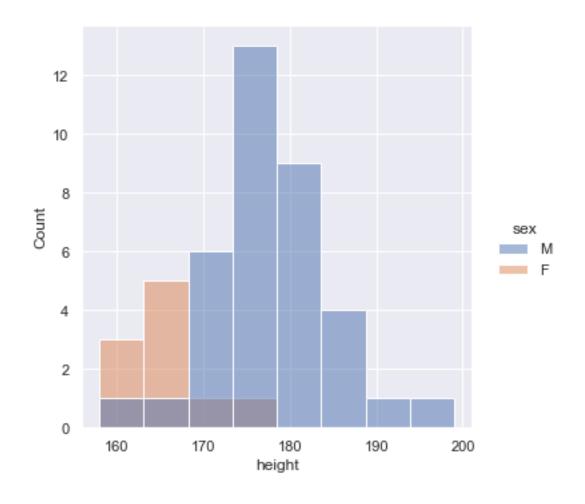
1.1.1

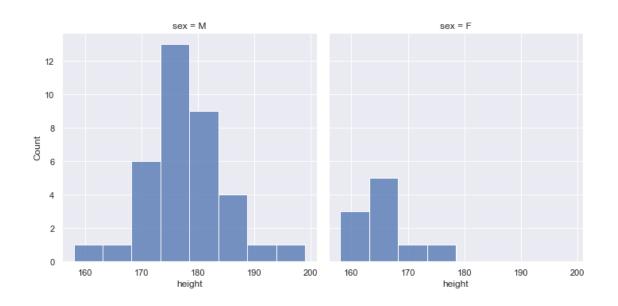
```
[72]: df = pd.read_excel('hw1_data.xlsx', header=None)
    df.columns = ['sex', 'height']
    male_df = df[df['sex'] == 'M']
    female_df = df[df['sex'] == 'F']
```

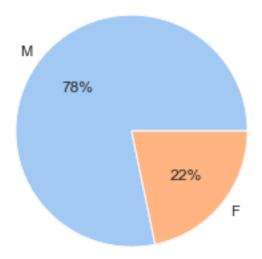
1.1.2

```
[73]: hist_fig1 = sns.displot(df, x='height', hue='sex')
hist_fig2 = sns.displot(df, x='height', col='sex')

hist_fig_name = 'hist_fig.pdf'
output_fig_path = os.path.join(output_path, hist_fig_name)
hist_fig1.savefig(output_fig_path, dpi=400)
```







1.2

1.

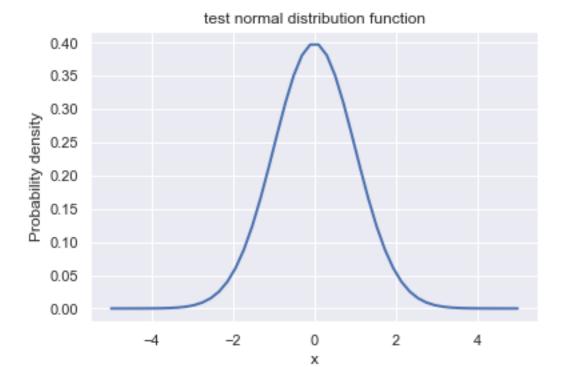
2. E $k y_j \hat{\gamma}_{jk}$

$$\hat{\gamma}_{jk} = \frac{\alpha_k \phi(y_j | \theta_k)}{\sum_{k=1}^K \alpha_k \phi(y_j | \theta_k)}$$

3. M

4.

```
[75]: def normal_dist(x, mu, sigma):
          pdf of normal distribution
          :param x:
          :param mu: mean
          :param sigma: variance
          :return:
          11 11 11
          a = 1.0 / sigma / np.sqrt(2*np.pi)
          exp = -1.0/2.0*((x-mu)/(sigma))**2
          exp = np.exp(exp)
          return a*exp
      test_x = np.linspace(-5, 5)
      test_y = normal_dist(test_x, 0.0, 1.0)
      # plot
      fig, ax = plt.subplots()
      ax.plot(test_x, test_y, linewidth=2.0)
      ax.set_title('test normal distribution function')
      ax.set_xlabel('x')
      ax.set_ylabel('Probability density')
      plt.show()
```



Membership probabilities

```
[76]: def membership_prob(x,
                          mu:list,
                          sigma:list,
                          tau:list,
                          idx):
          For the i th data, compute the probability that it belongs to
          the j th model, based on current parameters
          :param x: value of the i th data
          :param mu: list of mean values of all models
          :param sigma: list of variant values of all models
          :param tau: list of weights
          :param idx: index of the model
          :return:
          if len(mu)==len(sigma) and len(sigma)==len(tau):
              pass
          else:
              raise ValueError("The length of the input values is not identical")
          if idx >= len(mu):
              raise IndexError("The index is larger than the number of models")
          denominator = 0
          for j in range(len(mu)):
              denominator += tau[j]*normal_dist(x, mu[j], sigma[j])
          numerator = tau[idx] * normal_dist(x, mu[idx], sigma[idx])
          return numerator/denominator
```

New parameters

```
[77]: def new_tau(gamma:np.ndarray):
    return np.mean(gamma)

def new_mu(x:np.ndarray, gamma:np.ndarray):
    """
    Calculate the mean of j th model, using all the data
    :param x:
    :param gamma:
    :return:
    """

if x.size == gamma.size:
    numerator = x * gamma # do multiply element wise
    numerator = np.sum(numerator)
    denominator = np.sum(gamma)
    else:
```

```
raise ValueError("The length of x and gamma should be the same when_
return numerator/denominator
def new_sigma(x:np.ndarray, gamma:np.ndarray, mu):
   Calculate the variant of j th model, using all the data
   :param x:
   :param gamma:
   :param mu:
   :return:
   11 11 11
   if x.size == gamma.size:
       numerator = x - mu
       numerator = numerator**2
       numerator = numerator * gamma
       numerator = np.sum(numerator)
       denominator = np.sum(gamma)
   else:
       raise ValueError("The length of x and gamma should be the same when ⊔
return np.sqrt(numerator/denominator)
```

EM

```
[78]: def run(max_step, mu1, mu2, sigma1, sigma2, tau1):
          tau2 = 1 - tau1
          stat_label = ['mu1', 'mu2', 'sig1', 'sig2', 'tau1', 'tau2']
          stat_data = [mu1, mu2, sigma1, sigma2, tau1, tau2]
          stat_cnt = [0]*6
          gamma1 = np.zeros(shape=height_data.values.shape)
          gamma2 = np.zeros(shape=height_data.values.shape)
          for i in range(max_step):
              # E STEP
              for j in range(len(height_data.values)):
                  gamma1[j] = membership_prob(height_data[j], [mu1, mu2], [sigma1,__
       →sigma2], [tau1, tau2], 0)
                  gamma2[j] = membership_prob(height_data[j], [mu1, mu2], [sigma1,__
       →sigma2], [tau1, tau2], 1)
              # M step
              mu1 = new_mu(height_data.values, gamma1)
              mu2 = new_mu(height_data.values, gamma2)
              sigma1 = new_sigma(height_data.values, gamma1, mu1)
              sigma2 = new_sigma(height_data.values, gamma2, mu2)
              tau1 = new_tau(gamma1)
              tau2 = new_tau(gamma2)
```

```
# record the parameter
              stat_label = stat_label+['mu1', 'mu2', 'sig1', 'sig2', 'tau1', 'tau2']
              stat_data = stat_data + [mu1, mu2, sigma1, sigma2, tau1, tau2]
              stat_cnt = stat_cnt + [i+1]*6
          stat = pd.DataFrame({'step':stat_cnt, 'label':stat_label, 'data':stat_data})
          print("EM algorithm for GMM model -- Rsult:")
          print("mu1=",mu1,"mu2=",mu2)
          print("sigma1=",sigma1, "sigma2=",sigma2)
          print("tau1=",tau1, "tau2=",tau2)
          test = pd.DataFrame({"$\mu_1$":[mu1],"$\mu_2$":[mu2], "$\sigma_1$":
       →[sigma1], "$\sigma_2$":[sigma2], "$p_1$":[tau1], "$p_2$":[tau2]})
          print(test)
          return mu1, mu2, sigma1, sigma2, tau1, tau2, stat
[79]: def draw_result(mu1, mu2, sigma1, sigma2, filename):
          x_M = np.linspace(155, 200)
          x_F = np.linspace(150, 200)
          pdf_M = normal_dist(x_M, mu1, sigma1)
          pdf_F = normal_dist(x_F, mu2, sigma2)
          # fiq = plt.figure()
          fig, ax = plt.subplots()
          sns.histplot(ax=ax, data=df, x='height', hue='sex', stat="density", u
       sns.lineplot(ax=ax, x=x_M, y=pdf_M)
          sns.lineplot(ax=ax, x=x F,y=pdf F)
          ax.set title("Result")
          filename = os.path.join(output_path, filename)
          fig.savefig(filename, dpi=400)
      def draw_history(stat, filename):
          fig, axes = plt.subplots(2, 3, figsize=(15, 8), sharex=True)
          fig.suptitle('History of params during iteration')
          sns.lineplot(ax=axes[0,0], data=stat.query("label=='mu1'"), x="step", u

    y="data", marker='o')
          axes[0,0].set_title('$\mu_1$')
          sns.lineplot(ax=axes[1,0], data=stat.query("label=='mu2'"), x="step", __

    y="data", marker='o')
          axes[1,0].set_title('\$\mu_2\$')
          sns.lineplot(ax=axes[0,1], data=stat.query("label=='sig1'"), x="step", __
```

sns.lineplot(ax=axes[1,1], data=stat.query("label=='sig2'"), x="step", __

y="data", marker='o')

→y="data", marker='o')

axes[0,1].set_title('\$\sigma_1\$')

axes[1,1].set_title('\$\sigma_2\$')

```
sns.lineplot(ax=axes[0,2], data=stat.query("label=='tau1'"), x="step",
y="data", marker='o')
axes[0,2].set_title('$p_1$')
sns.lineplot(ax=axes[1,2], data=stat.query("label=='tau2'"), x="step",
y="data", marker='o')
axes[1,2].set_title('$p_2$')

filename = os.path.join(output_path, filename)
fig.savefig(filename, dpi=400)
```

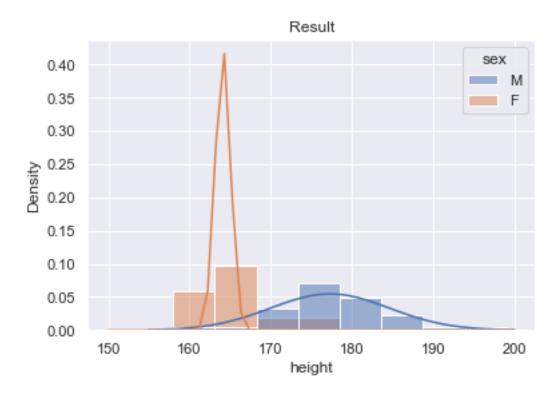
```
[80]: df = pd.read_csv('hw1_data.csv', header=None)
    df.columns = ['sex', 'height']
    height_data = df['height']
```

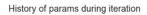
1.3

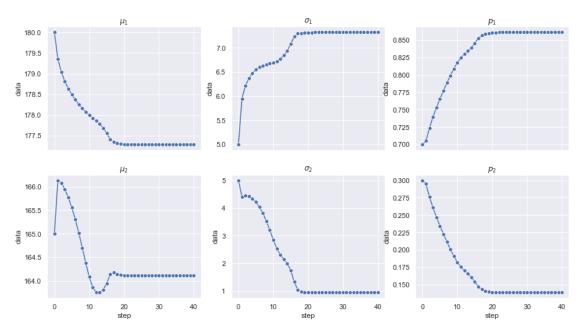
mu1= 177.28427949660465 mu2= 164.11589378931535 sigma1= 7.319264255059287 sigma2= 0.9428005047027699 tau1= 0.8612010767225255 tau2= 0.13879892327747453

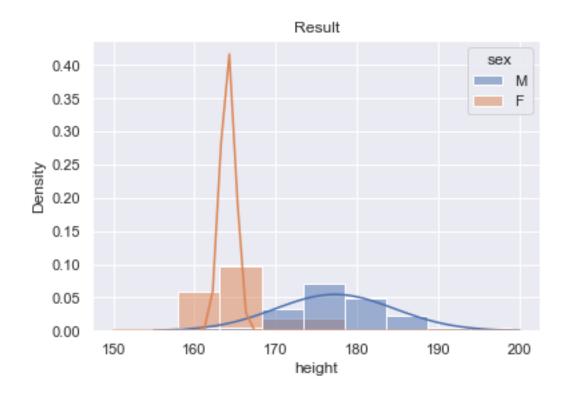
\$\mu_1\$ \$\mu_2\$ \$\sigma_1\$ \$\sigma_2\$ \$p_1\$ \$p_2\$
0 177.284279 164.115894 7.319264 0.942801 0.861201 0.138799
EM algorithm for GMM model -- Rsult:
mu1= 176.74955961795646 mu2= 172.3986702972358
sigma1= 8.23287562075184 sigma2= 7.197744928504974
tau1= 0.7028106707591764 tau2= 0.2971893292408235

\$\mu_1\$ \$\mu_2\$ \$\sigma_1\$ \$\sigma_2\$ \$p_1\$ \$p_2\$
0 176.74956 172.39867 8.232876 7.197745 0.702811 0.297189

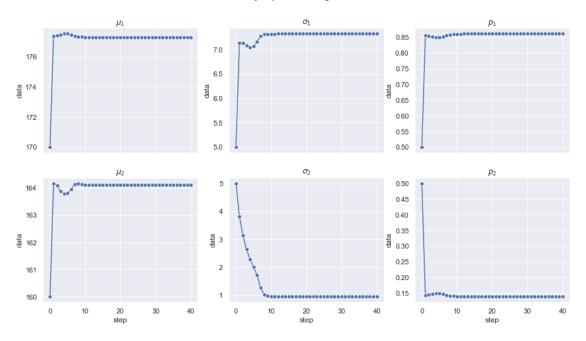


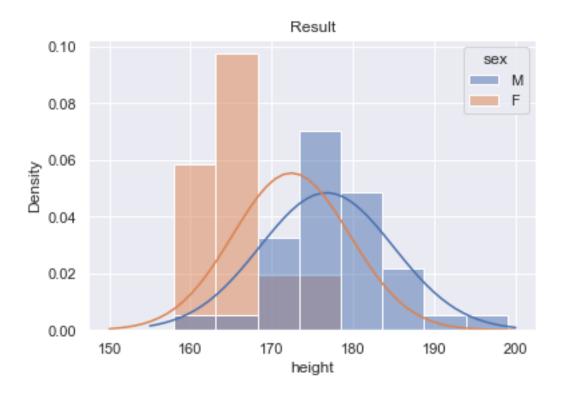




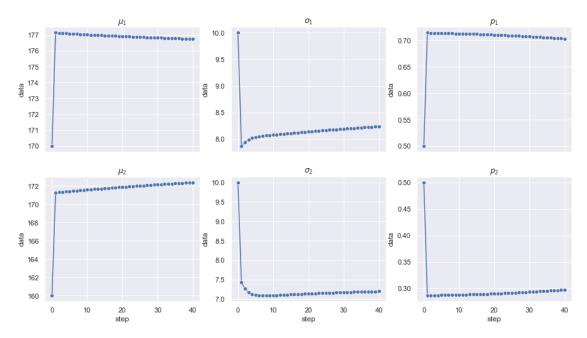












[81]: