# EM算法求解GMM模型参数

```
1 # 导入各个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')
```

# 数据预处理与可视化

### 导入数据

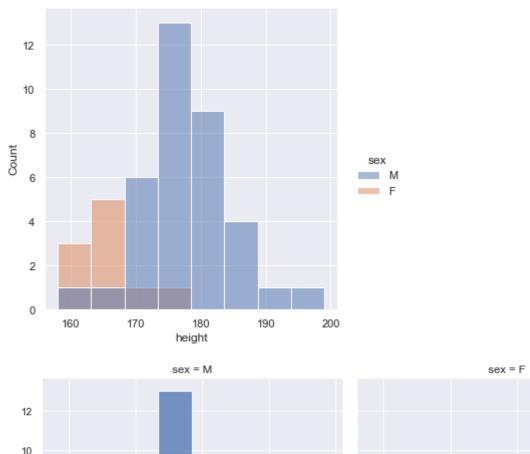
```
2  df = pd.read_excel('hw1_data.xlsx', header=None)
  df.columns = ['sex', 'height']
  male_df = df[df['sex']=='M']
  female_df = df[df['sex']=='F']
```

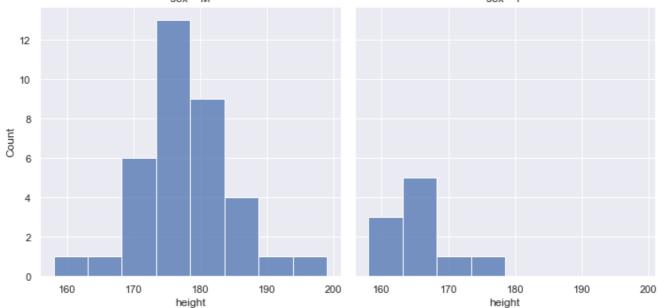
### 可视化

男女生身高的频数直方图

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

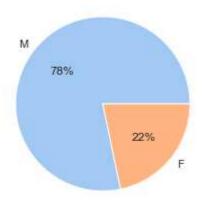
hist_fig_name = 'hist_fig.pdf'
output_fig_path = os.path.join(output_path, hist_fig_name)
hist_fig1.savefig(output_fig_path, dpi=400)
```





### 使用饼图显示男女生比例

```
4 sex_cnt = df['sex'].value_counts()
  #define Seaborn color palette to use
  colors = sns.color_palette('pastel')[0:5]
  #create pie chart
  plt.pie(sex_cnt.values, labels = list(sex_cnt.index), colors = colors, autopct='%.0f%%')
  pie_fig_name = 'pie_fig.pdf'
  output_fig_path = os.path.join(output_path, pie_fig_name)
  plt.savefig(output_fig_path, dpi=400, bbox_inches='tight')
  plt.show()
```



# 计算

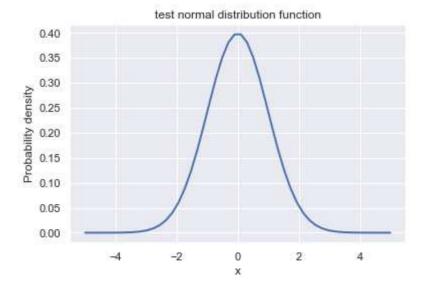
- 1. 取参数的初始值开始迭代
- 2. E步:根据当前模型参数,计算分模型k对观测数据 $y_i$ 的响应度 $\hat{\gamma}_{ik}$

$$\hat{\gamma}_{jk} = rac{lpha_k \phi(y_j | heta_k)}{\sum_{k=1}^K lpha_k \phi(y_j | heta_k)}$$

- 3. M步: 计算新一轮迭代的模型参数
- 4. 重复上面两步直到收敛

#### 正态分布函数

```
def normal_dist(x, mu, sigma):
5
       pdf of normal distribution
       :param x:
       :param mu: mean
       :param sigma: variance
       :return:
       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

```
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):
    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

```
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 calculating new mu")
        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:
        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 calculating new sigma")
        return np.sqrt(numerator/denominator)
    EM算法计算
66 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
        print(test)
        return mu1, mu2, sigma1, sigma2, tau1, tau2, stat
69 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)
   # fig = plt.figure()
   fig, ax = plt.subplots()
   sns.histplot(ax=ax, data=df, x='height', hue='sex', stat="density", common_norm=False)
   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", 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", y="data", marker='o')
   axes[0,1].set_title('$\sigma_1$')
   sns.lineplot(ax=axes[1,1], data=stat.query("label=='sig2'"), x="step", y="data", marker='o')
   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)
```

#### 加载数据

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

## 计算与结果可视化

```
70 mu1, mu2, sigma1, sigma2, tau1, tau2, stat = run(max_step=40,
           mu1=180, mu2=165, sigma1=5, sigma2=5, tau1=0.7)
   draw_result(mu1, mu2, sigma1, sigma2, "result1.pdf")
   # 各个参数在计算过程中的变化
   draw_history(stat, "history1.pdf")
   mu1, mu2, sigma1, sigma2, tau1, tau2, stat = run(max_step=40,
           mu1=170, mu2=160, sigma1=5, sigma2=5, tau1=0.5)
   draw_result(mu1, mu2, sigma1, sigma2, "result2.pdf")
   # 各个参数在计算过程中的变化
   draw_history(stat, "history2.pdf")
   mu1, mu2, sigma1, sigma2, tau1, tau2, stat = run(max_step=40,
           mu1=170, mu2=160, sigma1=10, sigma2=10, tau1=0.5)
   draw_result(mu1, mu2, sigma1, sigma2, "result3.pdf")
   # 各个参数在计算过程中的变化
   draw_history(stat, "history3.pdf")
   EM algorithm for GMM model -- Rsult:
   mu1= 177.28427949725432 mu2= 164.11589378938336
```

sigma1= 7.319264254664316 sigma2= 0.9428005047498634
tau1= 0.8612010766793227 tau2= 0.1387989233206773

 $\mbox{$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:$ 

 $mu1 = \ 177.28427949660465 \ mu2 = \ 164.11589378931535$ 

sigma1= 7.319264255059287 sigma2= 0.9428005047027699

tau1= 0.8612010767225255 tau2= 0.13879892327747453

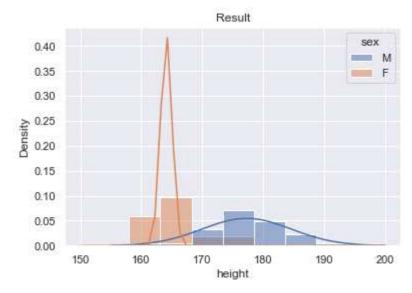
 $\mbox{$mu_1$ $\mu_2$ $\sigma_1$ $\sigma_2$ $p_1$ $p_2$ 0 177.284279 164.115894 7.319264 0.942801 0.861201 0.138799 EM algorithm for GMM model -- Rsult:$ 

mu1= 176.74955961795646 mu2= 172.3986702972358

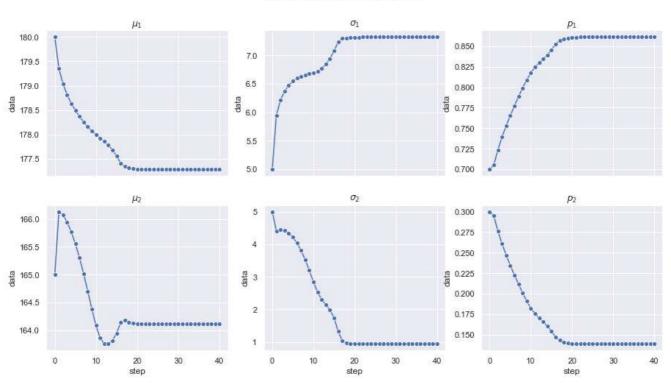
sigma1= 8.23287562075184 sigma2= 7.197744928504974

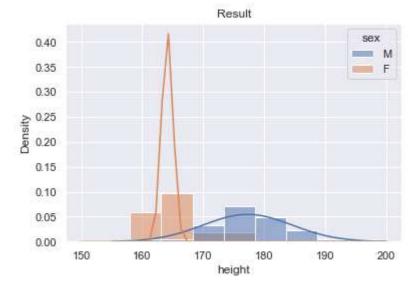
tau1= 0.7028106707591764 tau2= 0.2971893292408235

\$\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

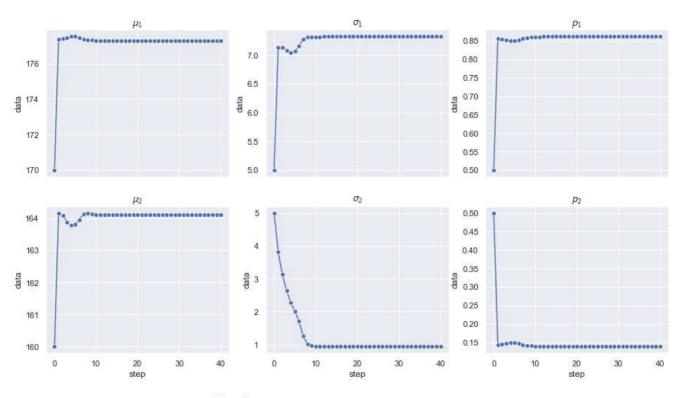


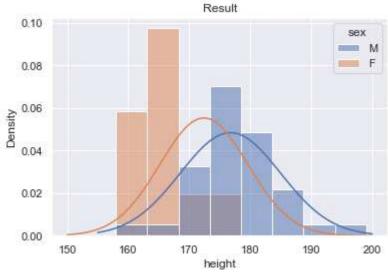
#### History of params during iteration





History of params during iteration





#### History of params during iteration

