多因子数据探索

假设检验

```
In [4]:
import numpy as np
import scipy. stats as ss
In [5]:
# 生成正态分布
norm dist = ss. norm. rvs(size=20)
norm dist
Out[5]:
array([ 0.54583651, 0.12684918, 0.4429434, -0.42516543, 0.49901631,
        0. 12287419, -0. 36590323, -0. 97259822, -0. 30700347, 0. 06935185,
        0.79107645, 1.6017795, 0.36359728, -0.54685468, 0.23501521,
       -0. 17216385, 1. 15114342, -0. 26190627, -1. 3299745, 0. 79158252])
In [6]:
# 基于偏度和峰度的正态检验
ss. normaltest (norm_dist)
Out[6]:
NormaltestResult(statistic=0.23319240531955351, pvalue=0.8899444778325278)
In [7]:
# 卡方检验
ss. chi2 contingency([[15, 95], [85, 5]])
Out[7]:
(126.08080808080808, 2.9521414005078985e-29, 1, array([[55., 55.],
        [45., 45.]]))
In [8]:
# 独立t分布检验, 检验两组值的均值是否有比较大的差异性
ss. ttest ind(ss. norm. rvs(size=10), ss. norm. rvs(size=20))
Out[8]:
Ttest indResult(statistic=-0.06670275173565719, pvalue=0.9472923595793121)
In [9]:
ss. ttest ind(ss. norm. rvs(size=100), ss. norm. rvs(size=200))
Out [9]:
```

Ttest indResult(statistic=0.6690022317186542, pvalue=0.5040119878623686)

In [10]:

```
a = [19, 22, 38, 77. 6, 52]
b = [19. 5, 23, 39, 77. 9, 52]
ss. ttest_ind(a, b)
```

Out[10]:

Ttest_indResult(statistic=-0.037019439471572595, pvalue=0.9713765928038277)

In [11]:

```
# f检验,方差检验
ss. f_oneway([49, 50, 39, 40, 43], [28, 32, 30, 26, 34], [38, 40, 45, 42, 48])
```

Out[11]:

F_onewayResult(statistic=17.619417475728156, pvalue=0.0002687153079821641)

QQ图

除了通过假设检验的方法,还可通过QQ图对比一个分布和已知的分布是否一致

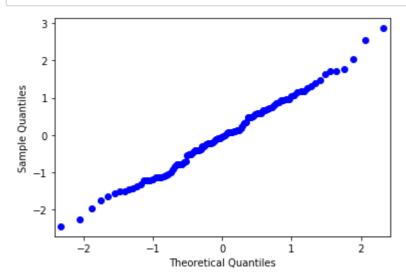
已知一个分布可以找到分位数,对应的分位数找到正态分布的分位数,横轴是正态分布分位数的值,纵轴是已知分布的值,得到一个曲线或散点图,这个图正对着x轴和y轴的角平分线,和平分线重合,就是符合分布的

In [12]:

```
from statsmodels.graphics.api import qqplot
from matplotlib import pyplot as plt
%matplotlib inline
```

In [13]:

```
# QQ图默认对比已知分布和正态分布的一致性
plt. show(qqplot(ss. norm. rvs(size=100)))
```



相关系数

```
In [14]:
```

```
import pandas as pd
```

```
In [15]:
```

```
s1 = pd. Series([0.1,0.2,1.1,2.4,1.3,0.3,0.5])
s2 = pd. Series([0.5,0.4,1.2,2.5,1.1,0.7,0.1])
# 皮尔逊相关系数
s1. corr(s2)
```

Out[15]:

0.9333729600465923

In [16]:

```
# spearman相关系数
sl.corr(s2, method='spearman')
```

Out[16]:

0.7142857142857144

In [17]:

```
df = pd.DataFrame([s1, s2])
df
```

Out[17]:

	0	1	2	3	4	5	6
0	0.1	0.2	1.1	2.4	1.3	0.3	0.5
1	0.5	0.4	1.2	2.5	1.1	0.7	0.1

In [18]:

```
# df是针对列进行相关系数计算,所以需要转化
df.corr()
```

Out[18]:

	0	1	2	3	4	5	6
0	1.0	1.0	1.0	1.0	-1.0	1.0	-1.0
1	1.0	1.0	1.0	1.0	-1.0	1.0	-1.0
2	1.0	1.0	1.0	1.0	-1.0	1.0	-1.0
3	1.0	1.0	1.0	1.0	-1.0	1.0	-1.0
4	-1.0	-1.0	-1.0	-1.0	1.0	-1.0	1.0
5	1.0	1.0	1.0	1.0	-1.0	1.0	-1.0
6	-1 0	-1 0	-1 0	-1 0	1.0	-1 0	1.0

```
In [20]:
```

```
# 转化成np. array(), 求转置,再计算相关系数
df = pd. DataFrame(np. array([s1, s2]). T)
df
```

Out[20]:

		0	1	
•	0	0.1	0.5	
	1	0.2	0.4	
	2	1.1	1.2	
	3	2.4	2.5	
	4	1.3	1.1	
	5	0.3	0.7	
	6	0.5	0.1	

In [21]:

df.corr()

Out[21]:

	U	1
0	1.000000	0.933373
1	0.933373	1.000000

In [22]:

```
df.corr(method='spearman')
```

Out[22]:

```
        0
        1

        0
        1.000000
        0.714286

        1
        0.714286
        1.000000
```

线性回归 y = wx + b

```
In [23]:
x = np. arange(10). astype(np. float). reshape((10, 1))
X
Out[23]:
array([[0.],
       [1.],
       [2.],
       [3.],
       [4.],
       [5.],
       [6.],
       [7.],
       [8.],
       [9.]
In [24]:
y = x * 3 + 4 + np. random. random((10, 1))
У
Out[24]:
array([[ 4.28641058],
       [ 7.94062331],
       [10.38510338],
       [13. 28458232],
       [16.82820008],
       [19. 32953052],
       [22.31209976],
       [25. 5777032],
       [28. 10107974],
       [31. 76226065]])
In [25]:
from sklearn.linear_model import LinearRegression
reg = LinearRegression()
# 拟合
res = reg. fit(x, y)
# 求估计值
y_pred = res. predict(x)
y_pred
Out[25]:
array([[ 4.50932133],
       [7.50297422],
       [10.49662712],
       [13.49028001],
       [16.48393291],
       [19.4775858],
       [22.4712387],
       [25.46489159],
       [28.45854448],
       [31. 45219738]])
```

```
In [26]:
# 参数w
reg.coef_
Out [26]:
array([[2.99365289]])
In [27]:
# 截距b
reg.intercept_
Out[27]:
array([4.50932133])
PCA主成分分析
In [28]:
data = np. array([np. array([2.5, 0.5, 2.2, 1.9, 3.1, 2.3, 2, 1, 1.5, 1.1]),
                 np. array([2.4, 0.7, 2.9, 2.2, 3, 2.7, 1.6, 1.1, 1.6, 0.9])]). T
data
Out[28]:
array([[2.5, 2.4],
       [0.5, 0.7],
       [2.2, 2.9],
       [1.9, 2.2],
       [3.1, 3.],
       [2.3, 2.7],
       [2., 1.6],
       [1., 1.1],
       [1.5, 1.6],
       [1.1, 0.9]
In [29]:
# sklearn中的PCA方法,使用的是奇异值分解SVD的方法
from sklearn. decomposition import PCA
# 降维-降成1维
lower_dim = PCA(n_components=1)
lower dim. fit (data)
# 降维后维度的属性
lower_dim. explained_variance_ratio_
Out[29]:
array([0.96318131])
```

```
In [30]:
```

```
# 直接得到转化后的数值
lower_dim.fit_transform(data)
Out[30]:
array([[-0.82797019],
      [ 1.77758033],
      [-0.99219749],
      [-0.27421042],
      [-1.67580142],
      [-0.9129491],
      [0.09910944],
      [ 1.14457216],
      [ 0.43804614],
      [ 1. 22382056]])
In [31]:
# 手写实现PCA方法,求协方差矩阵,求特征值和特征向量,得到转化后的值
def myPCA(data, n_components=1000000):
    # 对每列求均值
   mean_vals = np. mean(data, axis=0)
    # 每个数减去均值
   mid = data - mean_vals
    # 得到协方差矩阵
    cov_mat = np. cov (mid, rowvar=False)
    # 导入线性计算包
    from scipy import linalg
    # 求特征值和特征向量
    eig_vals, eig_vects = linalg.eig(np.mat(cov_mat))
    # 对特征值排序,记录索引
    eig_val_index = np. argsort(eig_vals)
    # 取k个最大的特征值
    eig val index = eig val index[:-(n components+1):-1]
    eig_vects = eig_vects[:,eig_val_index]
    # 点乘求降维后的矩阵
    low dim mat = np. dot(mid, eig vects)
   return low dim mat, eig vals
myPCA(data, n components=1)
Out[31]:
(array([[-0.82797019],
       [ 1.77758033],
       [-0.99219749],
       [-0.27421042],
       \lfloor -1.67580142 \rfloor,
       [-0.9129491],
       [0.09910944],
       [ 1.14457216],
       [0.43804614],
       [ 1.22382056]]), array([0.0490834 +0.j, 1.28402771+0.j]))
In [ ]:
```