

Homework 1

编辑日期：2023/05/11

1 Data Source

- 本数据来源于 Kaggle，包含了 570 种 cancer cell 的 30 个 feature 信息 所有的数据被标记为两类：
 - Benign Cancer (B)
 - Malignant Cancer (M)
- CSV 中各字段类型如下：
 - Col 0 为该记录的 id (String)
 - Col 1 为该记录的 class(B/M)
 - Col 2~31 为记录的 30 个其余 feature 信息 (Float)

```
import csv

data_file = 'Cancer_Data.csv'

try:
    f = open(data_file, encoding='utf-8')
except:
    print(f'Fail to open file "{data_file}"')

reader = csv.reader(f)
header = next(reader)
n_cols = len(header)

print(f"本数据集共有 {n_cols} 个字段: ")
for idx, key in enumerate(header):
    print("    ", '{0:<2}'.format(idx), key)
```

本数据集共有 32 个字段：

```
0  id
1  diagnosis
2  radius_mean
3  texture_mean
4  perimeter_mean
5  area_mean
6  smoothness_mean
7  compactness_mean
8  concavity_mean
9  concave points_mean
10 symmetry_mean
11 fractal_dimension_mean
```

```
12 radius_se
13 texture_se
14 perimeter_se
15 area_se
16 smoothness_se
17 compactness_se
18 concavity_se
19 concave points_se
20 symmetry_se
21 fractal_dimension_se
22 radius_worst
23 texture_worst
24 perimeter_worst
25 area_worst
26 smoothness_worst
27 compactness_worst
28 concavity_worst
29 concave points_worst
30 symmetry_worst
31 fractal_dimension_worst
```

2 Data Processing

2.1 转化数据格式

- 由于 `id` 字段对于构建分类器没有帮助，此处我们不予保留
- 由于本数据集中 `diagnose` 字段仅有 `M / B` 两种取值，为方便计算，此处分别将两种取值替换为 `0 / 1`

```
# 初始化
total = 0 # 数据总量
original_data = [] # 存储原始数据，Col_N 存储在数组 original_data[N] 中
count_null = [0] * (n_cols-1) # 统计空数据信息

for i in range(1, n_cols):
    original_data.append([])

for row in reader:
    total += 1
    flag = True
    for idx in range(1, n_cols):
        if not row[idx]:
            count_null[idx] += 1
            flag = False
    if flag:
        for i in range(1, n_cols):
            if(i==1):
                if (row[1][0] == 'M'):
                    original_data[0].append(0)
            else:
```

```

        original_data[0].append(1)
    else:
        original_data[i-1].append(float(row[i]))

f.close()

```

2.2 Handle Missing Value

数据缺失信息的统计实现在上一个代码块，此处仅展示统计结果。

不幸的是，本数据集的内容非常完整，不需要进行补全

```

print(f"本数据集共有 {len(original_data[0])} / {total} 条完整数据，空缺详情如下")
for idx in range(0, n_cols-1):
    print(" ", '{0:<4}'.format(idx), '{0:<25}'.format(header[idx+1]), count_null[idx])

```

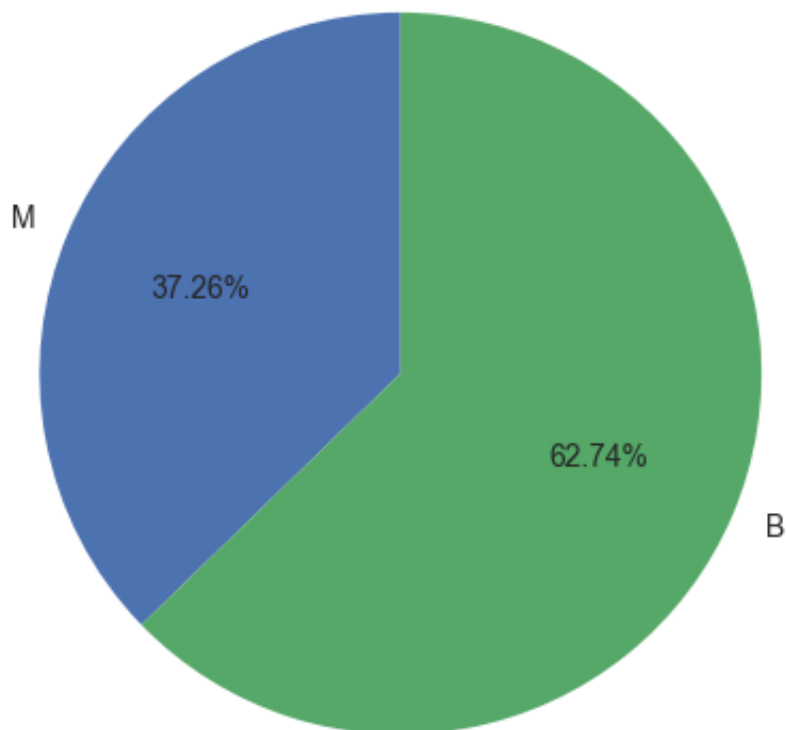
本数据集共有 569 / 569 条完整数据，空缺详情如下

0	diagnosis	0
1	radius_mean	0
2	texture_mean	0
3	perimeter_mean	0
4	area_mean	0
5	smoothness_mean	0
6	compactness_mean	0
7	concavity_mean	0
8	concave points_mean	0
9	symmetry_mean	0
10	fractal_dimension_mean	0
11	radius_se	0
12	texture_se	0
13	perimeter_se	0
14	area_se	0
15	smoothness_se	0
16	compactness_se	0
17	concavity_se	0
18	concave points_se	0
19	symmetry_se	0
20	fractal_dimension_se	0
21	radius_worst	0
22	texture_worst	0
23	perimeter_worst	0
24	area_worst	0
25	smoothness_worst	0
26	compactness_worst	0
27	concavity_worst	0
28	concave points_worst	0
29	symmetry_worst	0
30	fractal_dimension_worst	0

diagnose 字段中各类型数据占比情况如下：

```
import matplotlib.pyplot as plt
plt.style.use('seaborn')
plt.pie([original_data[0].count(0), original_data[0].count(1)], labels=["M", "B"],
autopct='%1.2f%%', startangle=90)
plt.show()
```

```
/var/folders/bv/mc555hkn509dv41d0yq119jr0000gn/T/ipykernel_43189/676113794.py:2:
MatplotlibDeprecationWarning: The seaborn styles shipped by Matplotlib are deprecated since
3.6, as they no longer correspond to the styles shipped by seaborn. However, they will remain
available as 'seaborn-v0_8-<style>'. Alternatively, directly use the seaborn API instead.
  plt.style.use('seaborn')
```

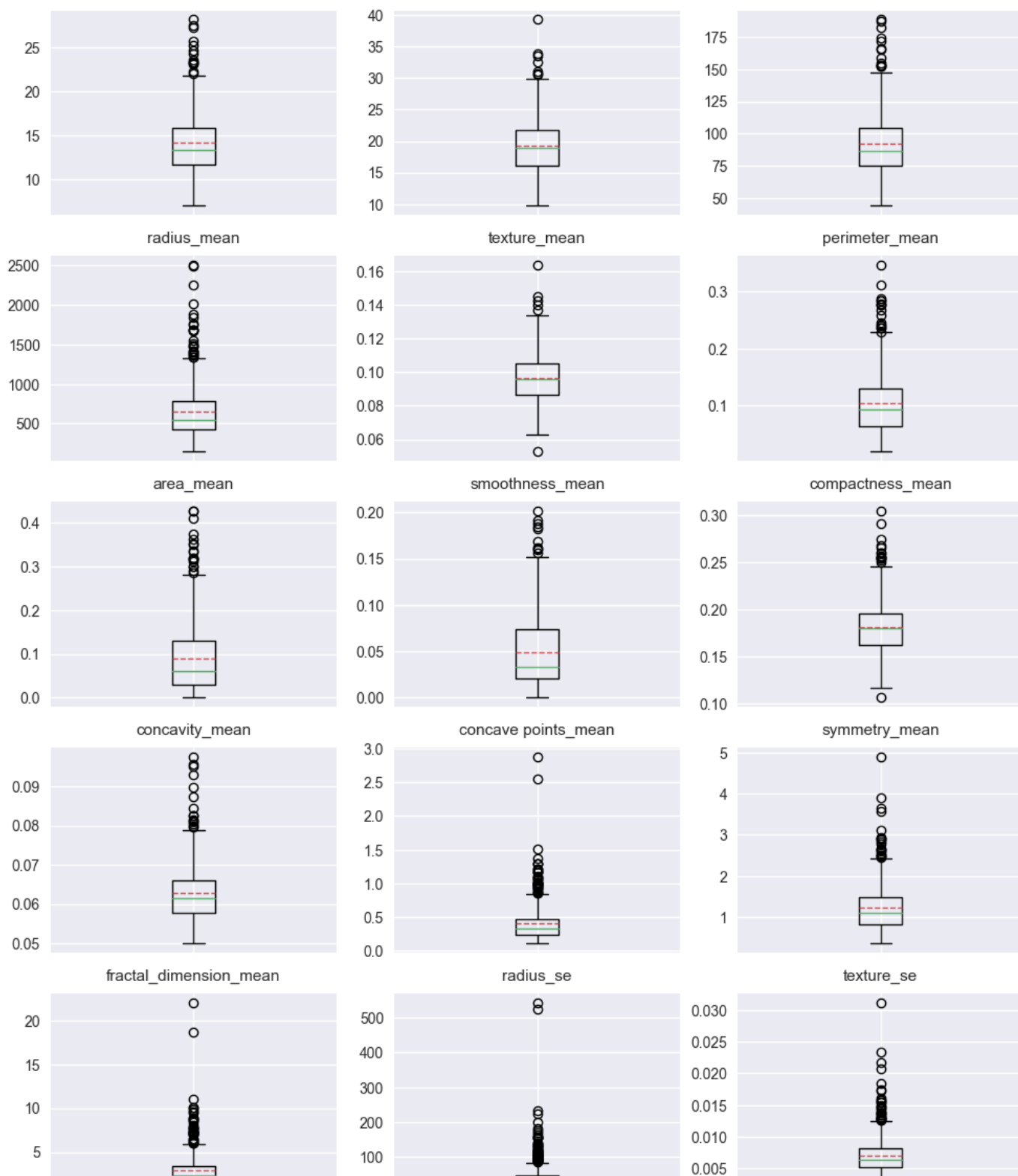


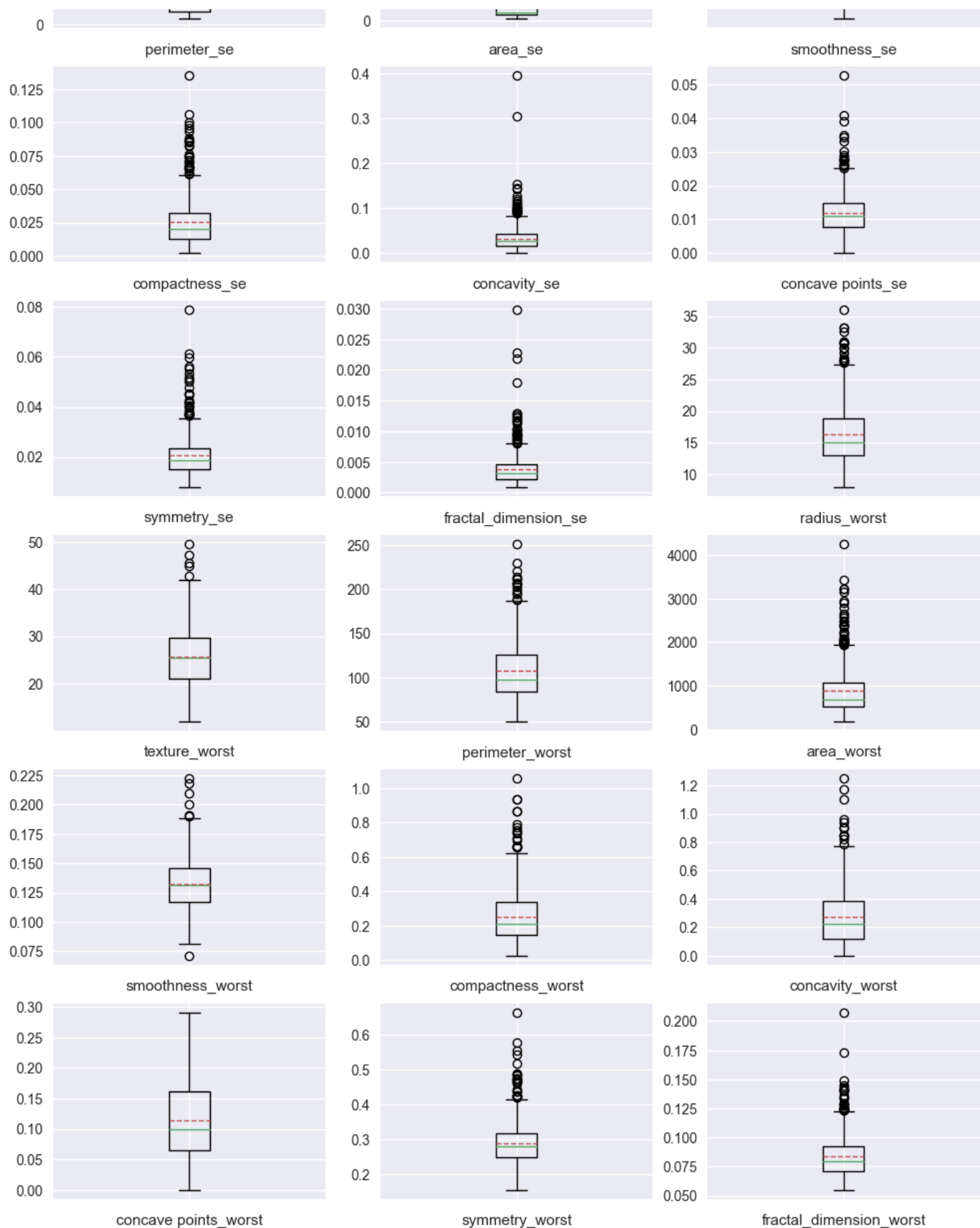
2.3 Remove Outliers

为原始数据绘制箱线图，检查 outliers 的分布状态

原始数据箱线图

```
plt.figure(figsize=(12,30))
for i in range(0, 30):
    ax = plt.subplot(10, 3, i+1)
    ax.set_xlabel(header[i+2])
    ax.set_xticks(range(0,1))
    ax.boxplot(original_data[i+1],
               showmeans=True,
               meanline=True)
```





显然，除 `concave points_worst` 外，各 Feature 均在不同程度上存在 outlier。

下面标记存在 outlier 的记录，并进行移除。

```
import math
```

```

q1_pos, q3_pos = math.floor(total*0.25), math.ceil(total*0.75)

lowers, uppers = [], []
for i in range(0,30):
    q1 = sorted(original_data[i+1])[q1_pos]
    q3 = sorted(original_data[i+1])[q3_pos]
    iqr = q3 - q1
    lowers.append(q1-1.5*iqr)
    uppers.append(q3+1.5*iqr)

outlier_rm = []
for i in range(0,31):
    outlier_rm.append([])

removed = []
for i in range(0, total):
    flag = True
    for j in range(0, 30):
        val = original_data[j+1][i]
        lower = lowers[j]
        upper = uppers[j]
        if val < lower or val > upper:
            flag = False
            break
    if flag:
        for j in range(0, 31):
            outlier_rm[j].append(original_data[j][i])
    else:
        removed.append(i)

print(f"{len(removed)} ({'{:.2f}'.format(len(removed)/total*100)}%) lines has been removed,
detail is as follows:")
print(removed)

```

```

169 (29.70%) lines has been removed, detail is as follows:
[0, 1, 2, 3, 4, 5, 8, 9, 12, 14, 15, 18, 22, 23, 24, 25, 26, 27, 30, 31, 33, 34, 35, 38, 41,
42, 53, 56, 60, 62, 63, 68, 70, 71, 72, 76, 77, 78, 82, 83, 95, 105, 108, 110, 111, 112, 116,
118, 119, 121, 122, 136, 138, 145, 146, 147, 150, 151, 152, 156, 161, 162, 164, 168, 173, 176,
180, 181, 185, 190, 192, 196, 199, 202, 203, 210, 212, 213, 214, 218, 219, 229, 232, 236, 239,
242, 245, 250, 252, 254, 256, 257, 258, 259, 262, 265, 272, 273, 275, 288, 290, 300, 302, 314,
318, 323, 329, 332, 335, 337, 339, 343, 345, 351, 352, 366, 368, 369, 370, 372, 376, 379, 388,
389, 391, 393, 400, 416, 417, 424, 430, 433, 443, 449, 450, 455, 460, 461, 465, 468, 469, 471,
473, 485, 489, 492, 498, 503, 504, 505, 507, 520, 521, 528, 533, 535, 537, 539, 553, 556, 557,
559, 561, 562, 563, 564, 565, 567, 568]

```

3 Data Reduction

3.1 Feature Selection

使用 Variance Threshold Filtering 进行数据降维

```
# 仅对后 30 个 Feature 进行筛选
vairances = []
for i in range(1, len(original_data)):
    avg = sum(original_data[i])/total
    tot = 0
    for j in range(0, total):
        tot += (original_data[i][j] - avg)*(original_data[i][j] - avg)
    vairances.append(tot/total)

# 输出不同阈值取值对筛选结果的影响
thresholds = [0.0001, 0.001, 0.01, 0.1, 0.5, 1, 2, 4, 8, 10, 15, 650]
percentages = []

print('{0:^11}'.format("Threshold"), '{0:^10}'.format('N-Feature'),
      '{0:^12}'.format("Percentage"), "   ", "Feature List")
print("-----")

for i in range(0, len(thresholds)):
    threshold = thresholds[i]
    selected = []
    for j in range(0, len(vairances)):
        if vairances[j] > threshold:
            selected.append(j+1)
    percentages.append(len(selected)/30*100)
    print('{0:^11}'.format(threshold), '{0:^10}'.format(len(selected)),
          '{0:^12}'.format('{0:.2f}'.format(percentages[i]) + "%"), "   ", selected)
```

Threshold	N-Feature	Percentage	Feature List

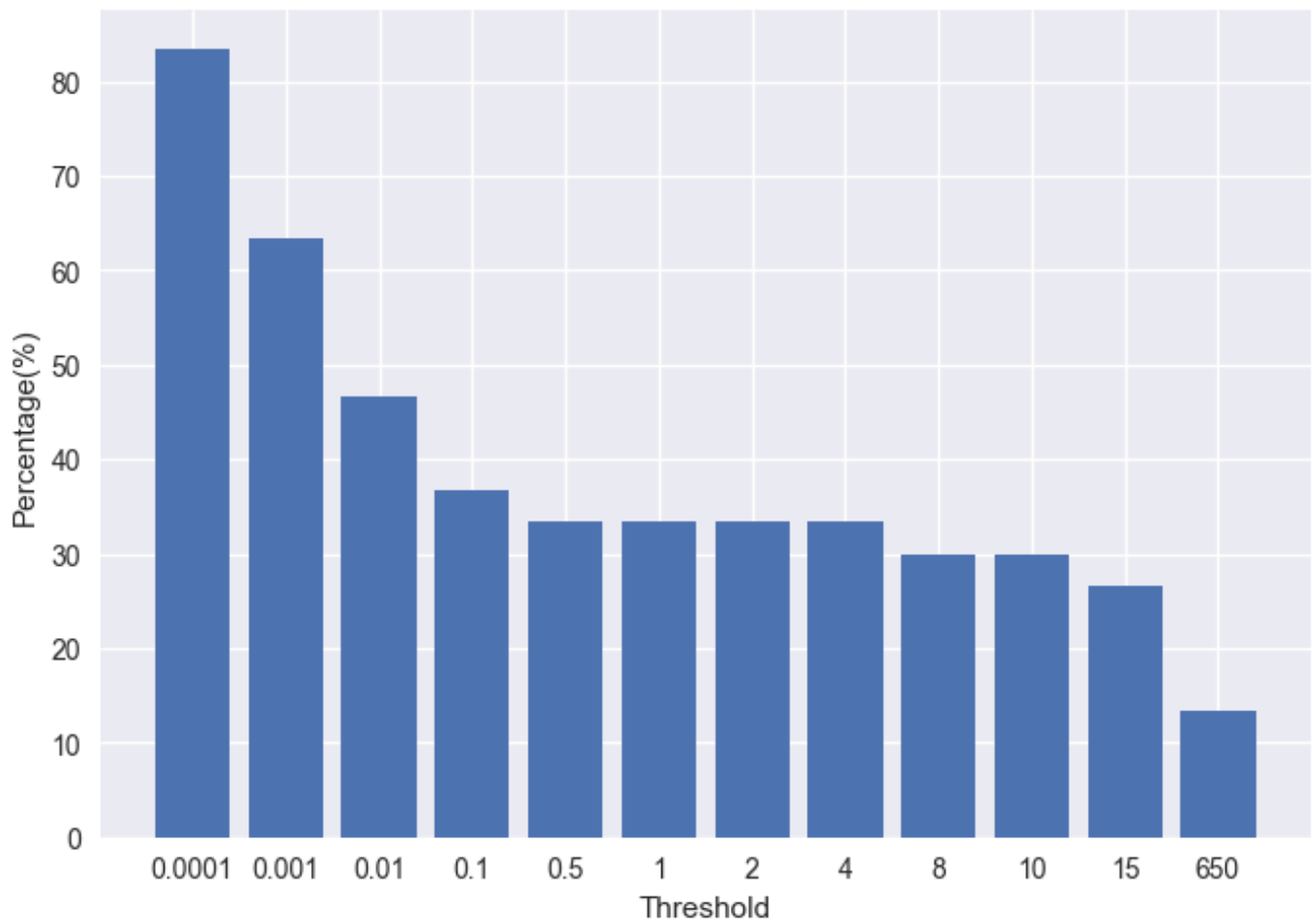
0.0001	25	83.33%	[1, 2, 3, 4, 5, 6, 7, 8, 9, 11, 12, 13, 14, 16, 17, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30]
0.001	19	63.33%	[1, 2, 3, 4, 6, 7, 8, 11, 12, 13, 14, 21, 22, 23, 24, 26, 27, 28, 29]
0.01	14	46.67%	[1, 2, 3, 4, 11, 12, 13, 14, 21, 22, 23, 24, 26, 27]
0.1	11	36.67%	[1, 2, 3, 4, 12, 13, 14, 21, 22, 23, 24]
0.5	10	33.33%	[1, 2, 3, 4, 13, 14, 21, 22, 23, 24]
1	10	33.33%	[1, 2, 3, 4, 13, 14, 21, 22, 23, 24]
2	10	33.33%	[1, 2, 3, 4, 13, 14, 21, 22, 23, 24]
4	10	33.33%	[1, 2, 3, 4, 13, 14, 21, 22, 23, 24]
8	9	30.00%	[1, 2, 3, 4, 14, 21, 22, 23, 24]
10	9	30.00%	[1, 2, 3, 4, 14, 21, 22, 23, 24]
15	8	26.67%	[2, 3, 4, 14, 21, 22, 23, 24]
650	4	13.33%	[4, 14, 23, 24]

Threshold 取值与 Feature 筛选数量的关系如下图所示：


```

ax = plt.subplot()
ax.bar(range(len(thresholds)), percentages)
ax.set_xlabel('Threshold')
ax.set_xticks(range(0, len(thresholds)))
ax.set_xticklabels([str(i) for i in thresholds])
ax.set_ylabel('Percentage(%)')
plt.show()

```



以上是对于原始数据的筛选结果，下面尝试对去除 outlier 的数据进行筛选：

```

# 计算方差
vairances = []
total = len(outlier_rm[0])
for i in range(1, 31):
    avg = sum(outlier_rm[i])/total
    tot = 0
    for j in range(0, total):
        tot += (outlier_rm[i][j] - avg)*(outlier_rm[i][j] - avg)
    vairances.append(tot/total)

# 不同阈值取值对筛选结果的影响
percentages = []

```

```

print('{0:^11}'.format("Threshold"), '{0:^10}'.format('N-Feature'),
      '{0:^12}'.format("Percentage"), " ", "Feature List")
print("-----")

for i in range(0, len(thresholds)):
    threshold = thresholds[i]
    selected = []
    for j in range(0, len(vairances)):
        if vairances[j] > threshold:
            selected.append(j+1)
    percentages.append(len(selected)/30*100)
    print('{0:^11}'.format(threshold), '{0:^10}'.format(len(selected)),
          '{0:^12}'.format('{0:.2f}'.format(percentages[i]) + "%"), " ", selected)

```

Threshold	N-Feature	Percentage	Feature List

0.0001	25	83.33%	[1, 2, 3, 4, 5, 6, 7, 8, 9, 11, 12, 13, 14, 16, 17, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30]
0.001	18	60.00%	[1, 2, 3, 4, 6, 7, 11, 12, 13, 14, 21, 22, 23, 24, 26, 27, 28, 29]
0.01	14	46.67%	[1, 2, 3, 4, 11, 12, 13, 14, 21, 22, 23, 24, 26, 27]
0.1	11	36.67%	[1, 2, 3, 4, 12, 13, 14, 21, 22, 23, 24]
0.5	10	33.33%	[1, 2, 3, 4, 13, 14, 21, 22, 23, 24]
1	9	30.00%	[1, 2, 3, 4, 14, 21, 22, 23, 24]
2	9	30.00%	[1, 2, 3, 4, 14, 21, 22, 23, 24]
4	9	30.00%	[1, 2, 3, 4, 14, 21, 22, 23, 24]
8	8	26.67%	[2, 3, 4, 14, 21, 22, 23, 24]
10	8	26.67%	[2, 3, 4, 14, 21, 22, 23, 24]
15	7	23.33%	[2, 3, 4, 14, 22, 23, 24]
650	2	6.67%	[4, 24]

对比前表可知，在相同的 Threshold 下，对去除 outlier 的数据集进行筛选将保留 **更多的 Feature**。

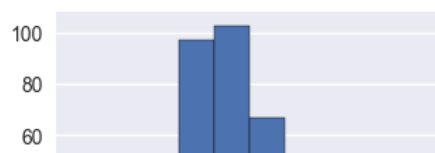
3.2 Histogram Analysis

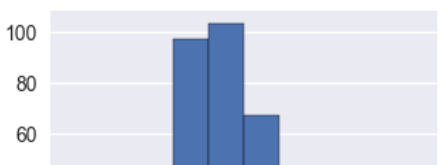
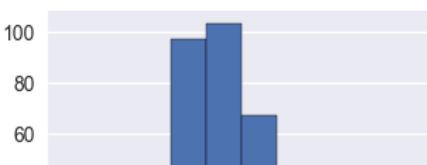
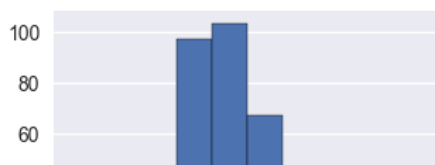
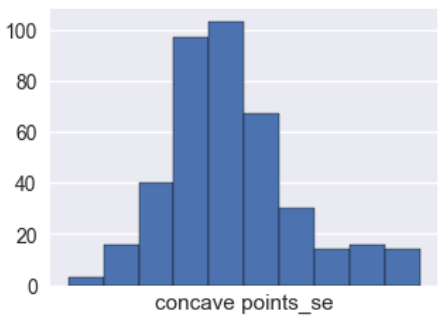
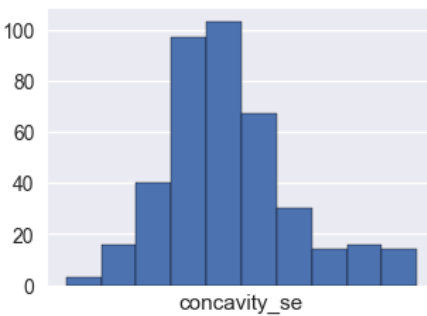
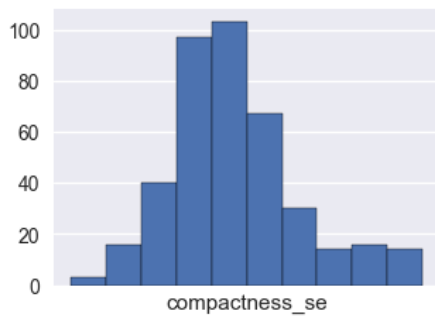
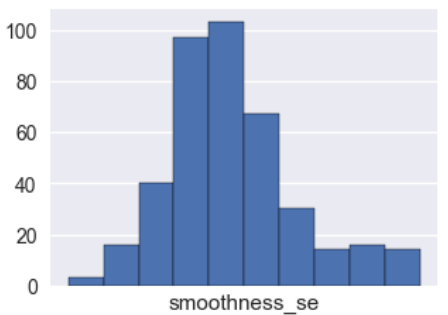
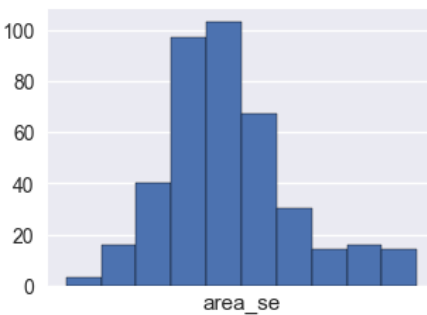
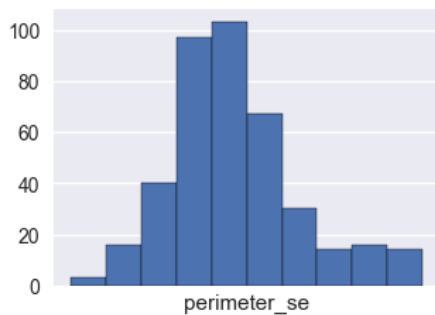
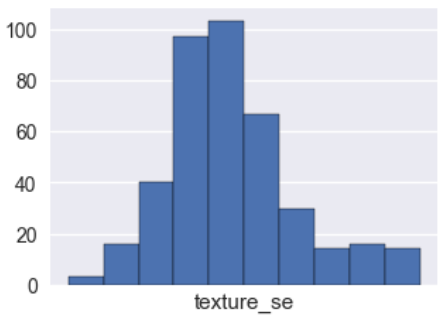
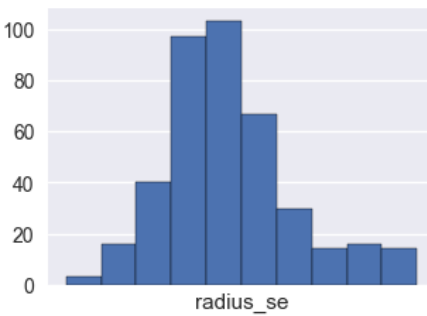
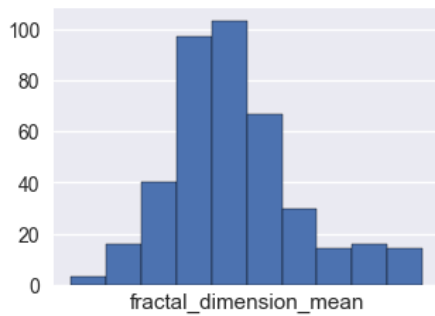
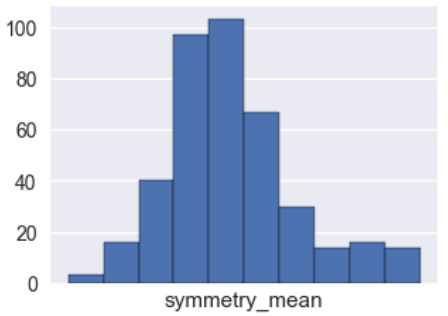
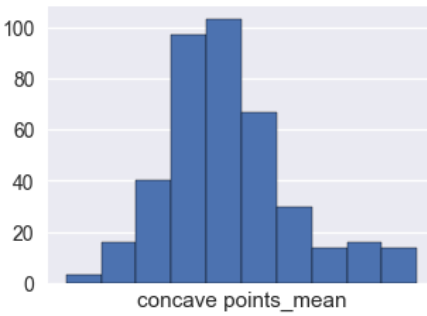
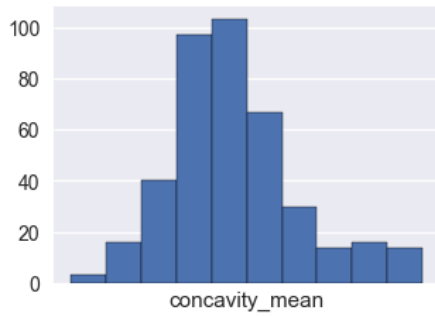
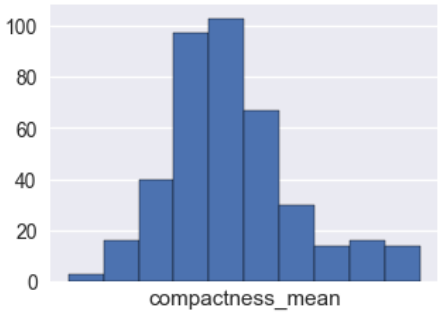
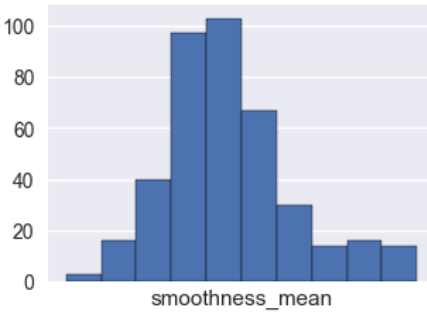
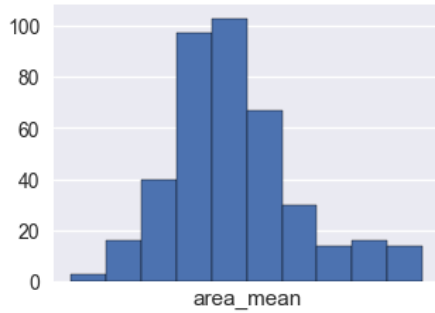
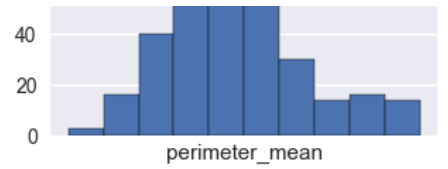
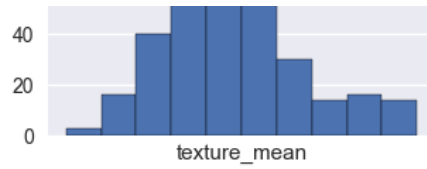
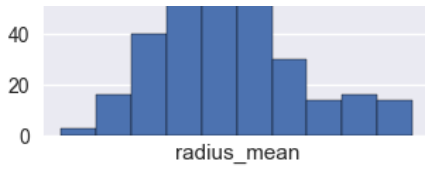
通过绘制去除 outlier 数据的 equi-width 直方图，观察数据分布情况。

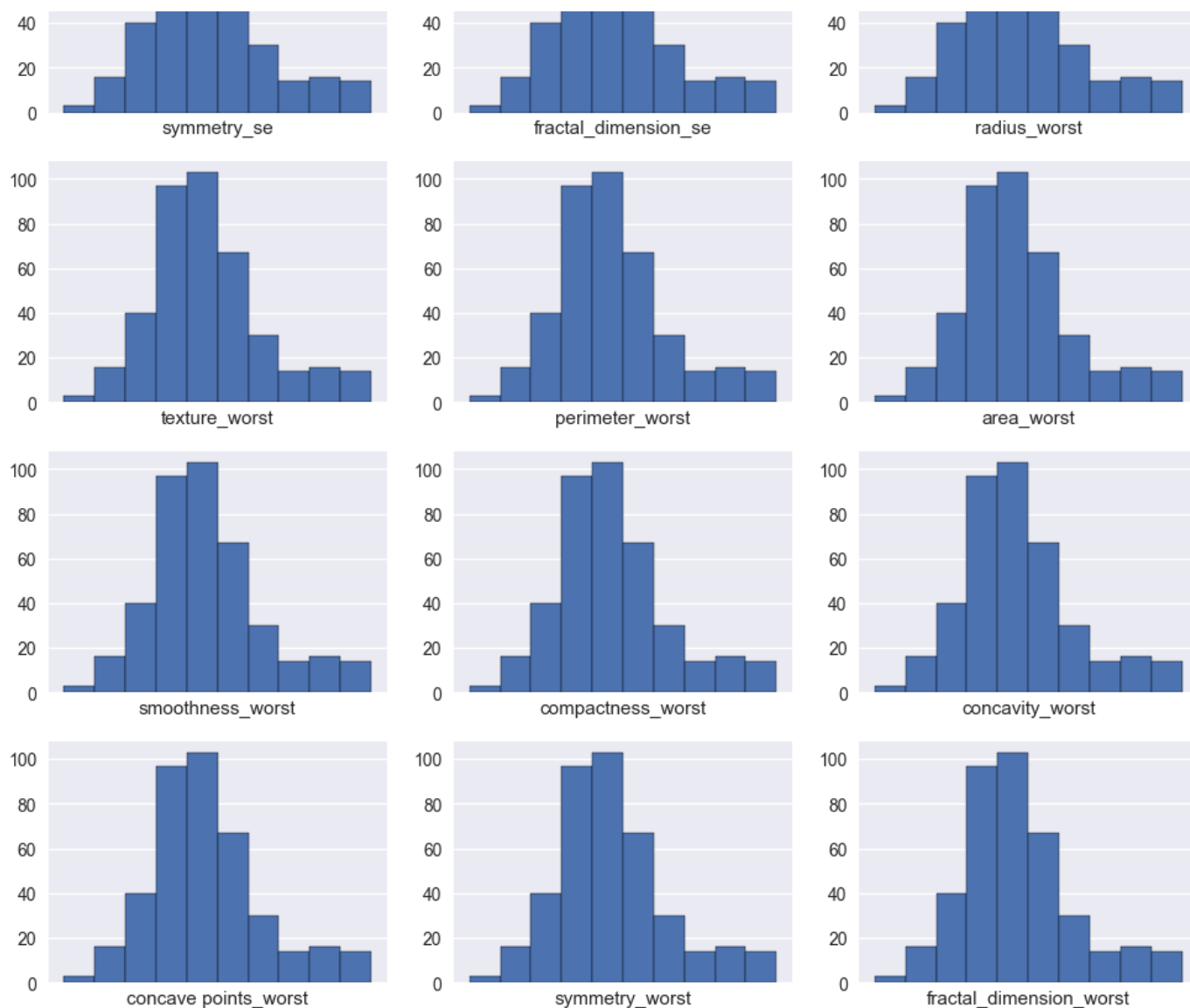
```

# 去除 outlier 数据的直方图
plt.figure(figsize=(12,30))
for i in range(0, 30):
    ax = plt.subplot(10, 3, i+1)
    ax.set_xlabel(header[i+2])
    ax.set_xticks(range(0,1))
    ax.hist(outlier_rm[1], edgecolor="black")

```





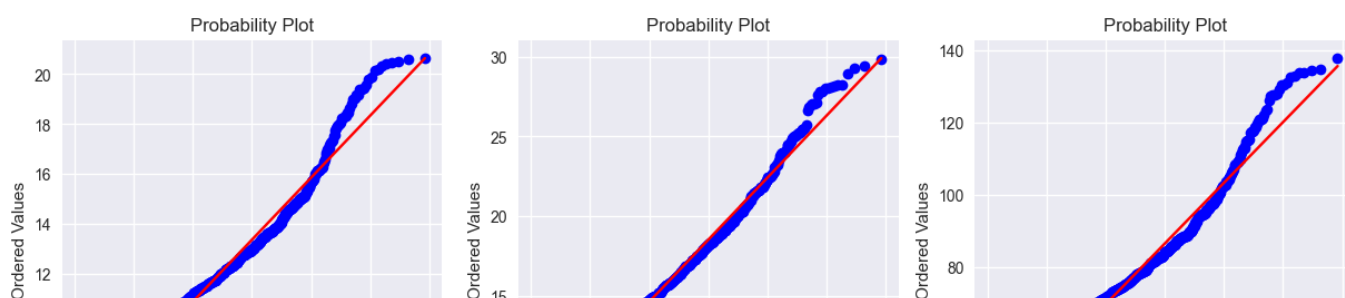


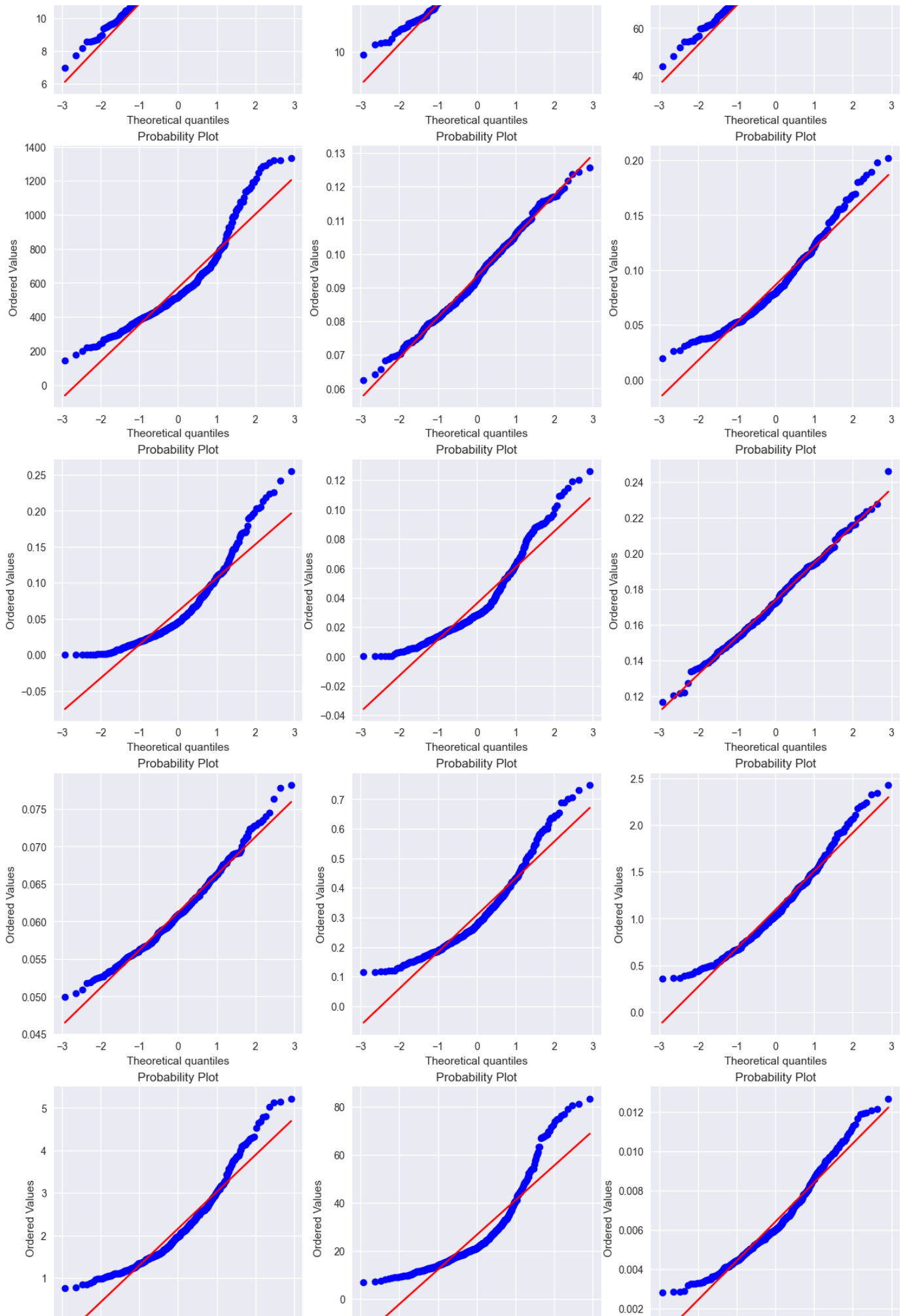
从上图可知，去除 outlier 后的各 Feature 数据基本符合正态分布，下面通过 Q-Q 图进行验证：

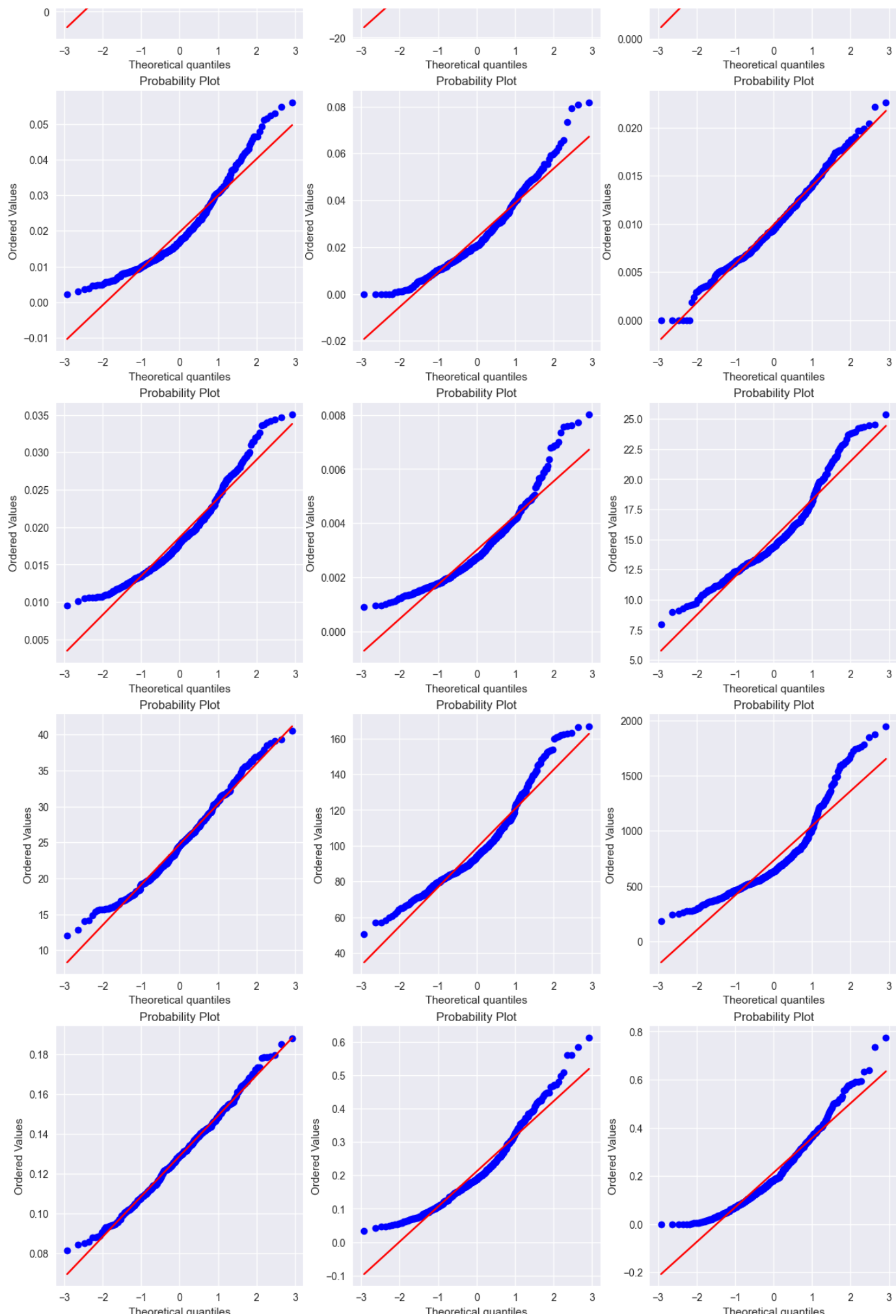
```
# 绘制各 Feature 的 Q-Q 图
from scipy import stats
plt.figure(figsize=(15,55))

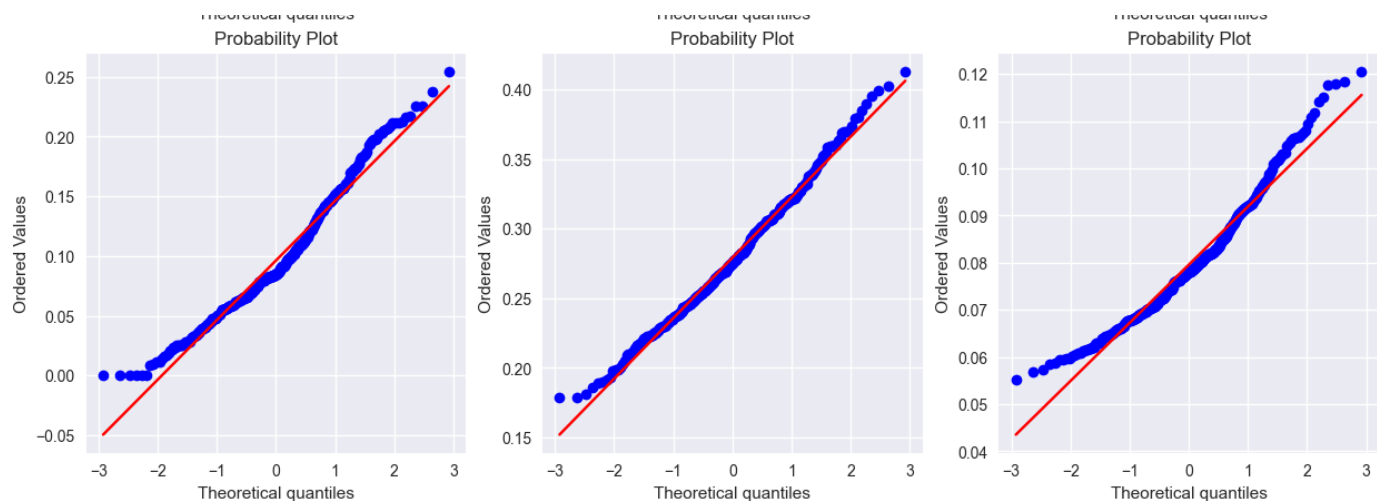
for i in range(0,30):
    ax = plt.subplot(10, 3, i+1)
    # 没办法设小标题（我裂开）
    stats.probplot(outlier_rm[i+1], dist="norm", plot=ax)

plt.show()
```









4 Data Transformation

由于不同的 `diagnose` 组别间的数据可能满足不同的正态分布，下面通过 MinMax 方式对数据进行标准化：

$$x_{norm} = \frac{x - x_{min}}{x_{max} - x_{min}}$$

```
# 统计每个属性的 min / max 值
mins = [0]
maxs = [1]

print("各属性取值的 min / max 信息如下：")
print('{0:^24}'.format("Attribute Name"), '{0:^10}'.format("min"), '{0:^10}'.format("max"))
print("-----")

for i in range(1, len(outlier_rm)):
    minn = min(outlier_rm[i])
    maxx = max(outlier_rm[i])
    mins.append(minn)
    maxs.append(maxx)
    print('{0:^24}'.format(header[i]), '{0:^10}'.format(minn), '{0:^10}'.format(maxx))
```

各属性取值的 min / max 信息如下：

Attribute Name	min	max
diagnosis	6.981	20.64
radius_mean	9.71	29.81
texture_mean	43.79	137.8
perimeter_mean	143.5	1335.0
area_mean	0.06251	0.1257
smoothness_mean	0.01938	0.2022
compactness_mean	0.0	0.2545
concavity_mean	0.0	0.1259

concave points_mean	0.1167	0.2459
symmetry_mean	0.04996	0.07818
fractal_dimension_mean	0.1144	0.7474
radius_se	0.3602	2.426
texture_se	0.757	5.216
perimeter_se	6.802	83.5
area_se	0.002826	0.01266
smoothness_se	0.002252	0.05592
compactness_se	0.0	0.08158
concavity_se	0.0	0.02258
concave points_se	0.009539	0.03504
symmetry_se	0.0008948	0.008015
fractal_dimension_se	7.93	25.37
radius_worst	12.02	40.54
texture_worst	50.41	166.8
perimeter_worst	185.2	1946.0
area_worst	0.08125	0.1878
smoothness_worst	0.03432	0.611
compactness_worst	0.0	0.7727
concavity_worst	0.0	0.2543
concave points_worst	0.1783	0.4128
symmetry_worst	0.05521	0.1205

对数据进行标准化

```
normalized_data = []
for i in range(0, 31):
    normalized_data.append([])

for i in range(0, total):
    normalized_data[0].append(outlier_rm[0][i])
    for j in range(1, 31):
        normalized_data[j].append((outlier_rm[j][i] - mins[j])/(maxs[j] - mins[j]))
```

下面重新计算输出经过 MinMax 标准化后，各 Feature 字段的最大最小值以验证操作的正确性。

```
print("完成标准化后，各属性取值的 min / max 信息如下：")
print('{0:^24}'.format("Attribute Name"), '{0:^10}'.format("min"), '{0:^10}'.format("max"))
print("-----")

for i in range(1, len(normalized_data)):
    minn = min(normalized_data[i])
    maxx = max(normalized_data[i])
    print('{0:^24}'.format(header[i]), '{0:^10}'.format(minn), '{0:^10}'.format(maxx))
```

完成标准化后，各属性取值的 min / max 信息如下：

Attribute Name	min	max
diagnosis	0.0	1.0
radius_mean	0.0	1.0
texture_mean	0.0	1.0

perimeter_mean	0.0	1.0
area_mean	0.0	1.0
smoothness_mean	0.0	1.0
compactness_mean	0.0	1.0
concavity_mean	0.0	1.0
concave points_mean	0.0	1.0
symmetry_mean	0.0	1.0
fractal_dimension_mean	0.0	1.0
radius_se	0.0	1.0
texture_se	0.0	1.0
perimeter_se	0.0	1.0
area_se	0.0	1.0
smoothness_se	0.0	1.0
compactness_se	0.0	1.0
concavity_se	0.0	1.0
concave points_se	0.0	1.0
symmetry_se	0.0	1.0
fractal_dimension_se	0.0	1.0
radius_worst	0.0	1.0
texture_worst	0.0	1.0
perimeter_worst	0.0	1.0
area_worst	0.0	1.0
smoothness_worst	0.0	1.0
compactness_worst	0.0	1.0
concavity_worst	0.0	1.0
concave points_worst	0.0	1.0
symmetry_worst	0.0	1.0

从上表可知，经过 MinMax 标准化后，所有 Feature 的取值均处于区间 `[0,1]` 内。