## 数据挖掘作业——马的疝病分析

# 一、 数据摘要及可视化

### 1. 数据摘要

该部分内容均在 python 下通过 pandas 库完成。

### 1.1 标称属性频次统计

在该数据集中包含以下标称属性: surgery、Age、Hospital Number、temperature of extremities、peripheral pulse、mucous membranes、capillary refill time、pain、peristalsis、abdominal distension、nasogastric tube、nasogastric reflux、rectal examination – feces、abdomen、abdominocentesis appearance、outcome、surgical lesion、cp\_data。

其分别的词频统计为:

#### (1) Surgery:

1.0 214 2.0 152

Name: surgery, dtype: int64

#### (2) Age:

1 340

Name: Age, dtype: int64

#### (3) temperature of extremities:

3.0 135

1.0 95

2.0 3

4.0 34

Name: temperature of extremities, dtype: int64

```
(4) peripheral pulse:
     1.0
            151
     3.0
            116
     4.0
             12
     2.0
              6
     Name: peripheral pulse, dtype: int64
(5) mucous membranes:
    1.0
            98
    3.0
            81
            50
    4.0
    2.0
            38
    5.0
            28
            25
    6.0
    Name: mucous membranes, dtype: int64
(6) capillary refill time:
    1.0
            232
    2.0
            96
    3.0
              2
    Name: capillary refill time, dtype: int64
(7) pain:
    3.0
            82
     2.0
            77
    5.0
            50
    1.0
            49
     4.0
            47
    Name: pain, dtype: int64
(8) Peristalsis:
    3.0
            154
            91
    4.0
    1.0
             49
    2.0
             22
    Name: peristalsis, dtype: int64
(9) abdominal distension:
           101
    1.0
    3.0
            85
    2.0
            75
    4.0
            42
    Name: abdominal distension, dtype: int64
```

(10) nasogastric tube:

```
2.0
           121
    1.0
            89
    3.0
            27
    Name: nasogastric tube, dtype: int64
(11) nasogastric reflux:
    1.0
            141
     3.0
             49
     2.0
             45
    Name: nasogastric reflux, dtype: int64
(12) rectal examination – feces:
    4.0
           97
    1.0
           68
    3.0
           61
    2.0
            14
    Name: rectal examination - feces, dtype: int64
(13) abdomen:
    5.0
            96
    4.0
            55
    1.0
            31
    2.0
            24
    3.0
            19
    Name: abdomen, dtype: int64
( 14 ) abdominocentesis appearance :
    2.0
           62
    3.0
           60
           52
    1.0
    Name: abdominocentesis appearance, dtype: int64
( 15 ) outcome :
           225
    1.0
    2.0
            89
            52
    3.0
    Name: outcome, dtype: int64
(16) surgical lesion:
         232
    2
         136
    Name: surgical lesion, dtype: int64
( 17 ) cp_data :
```

2 244 1 124

Name: cp\_ data, dtype: int64

### 1.2 数值型数据统计

在该数据集中包含以下数值属性: rectal temperature、pulse、respiratory rate、nasogastric reflux PH、packed cell volume、total protein、abdomcentesis total protein。

下面分别统计出数值属性的有效总数、平均值、标准差、最小值、二分位数、中位数、四分位数及最大值。

(1) rectal temperature:

299.000000 count 38.134448 mean 0.711684 std min 35.400000 25% 37.800000 50% 38.100000 75% 38.500000 40.800000 max

Name: rectal temperature, dtype: float64

(2) pulse:

count 342.000000
mean 70.757310
std 28.089867
min 30.000000
25% 48.000000
50% 60.000000
75% 88.000000
max 184.000000

Name: pulse, dtype: float64

(3) respiratory rate:

297.000000 count mean 30.521886 17.669651 std 8.000000 min 25% 18.000000 50% 28.000000 75% 36.000000 96.000000 max

Name: respiratory rate, dtype: float64

(4) nasogastric reflux PH:

```
69.000000
count
          4.962319
mean
          2.003901
std
min
          1.000000
25%
          3.500000
50%
          5.400000
75%
          6.500000
          8.500000
max
Name: nasogastric reflux PH, dtype: float64
```

-----, ---, ---, ---, ---,

(5) packed cell volume:

```
count 331.000000
mean 45.656798
std 10.865663
min 4.000000
25% 37.250000
50% 44.000000
75% 52.000000
max 75.000000
```

Name: packed cell volume, dtype: float64

(6) total protein:

```
325.000000
count
          24.771077
mean
          27.704880
std
           3.300000
min
25%
           6.500000
           7.500000
50%
75%
          58.000000
          89.000000
max
```

Name: total protein, dtype: float64

(7) abdomcentesis total protein:

```
133.000000
count
mean
           2.948120
           1.927064
std
min
           0.100000
25%
           2.000000
50%
           2.100000
75%
           3.900000
          10.100000
max
```

Name: abdomcentesis total protein, dtype: float64

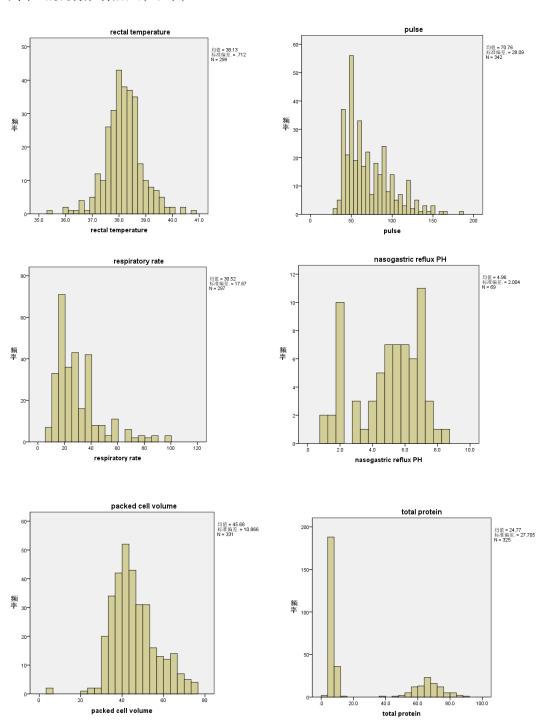
## 2. 数据可视化

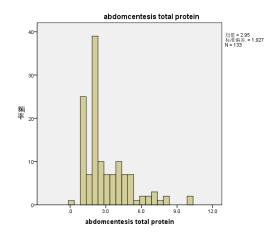
该部分内容通过 spss 工具完成。

# 2.1 数值属性的直方图及 Q\_Q 图

# 2.1.1 直方图

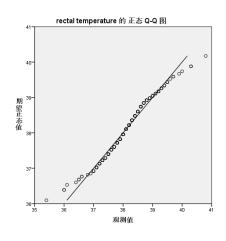
### 下面分别为数值数据的直方图:

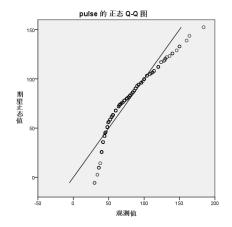


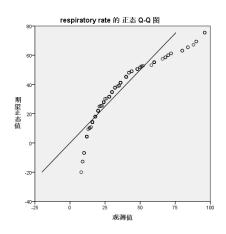


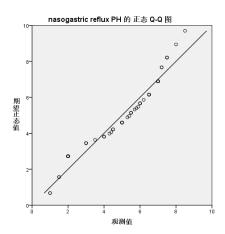
# 2.1.2 Q\_Q 图

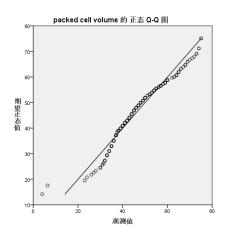
## 下面分别为数值数据的直方图:

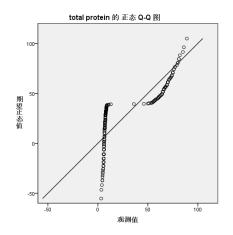


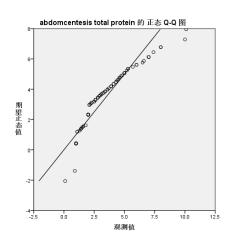






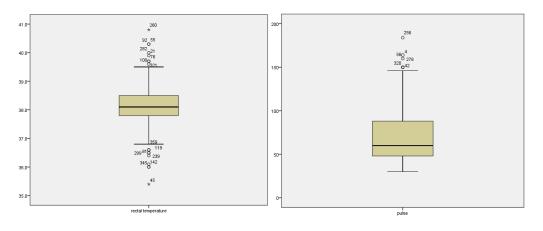


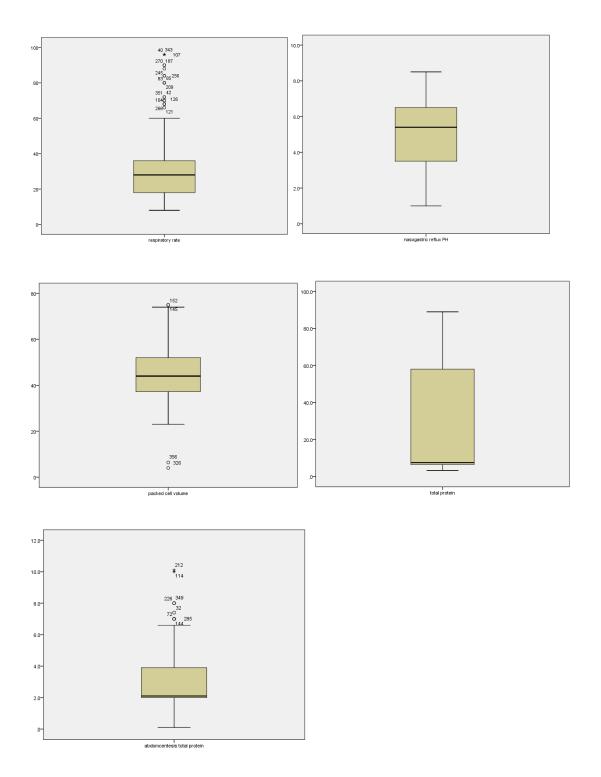




# 2.2 数值属性的盒图

### 下面分别为数值数据的盒图:





# 二、数据缺失的处理

该部分内容先通过 python 对数据进行处理,后通过 spss 制作绘图

# 1. 将缺失部分剔除

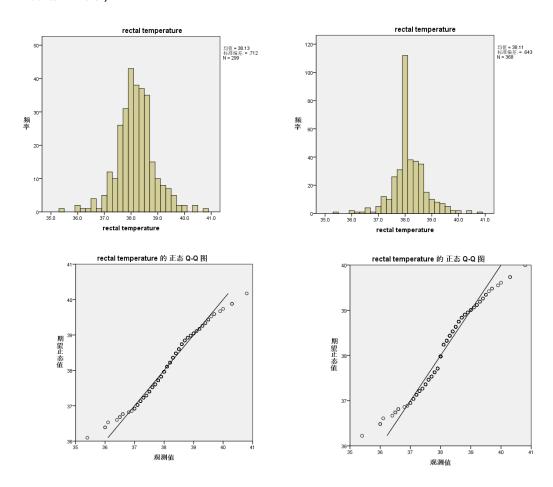
该部分图例已在上一部分展示,不做重复处理。

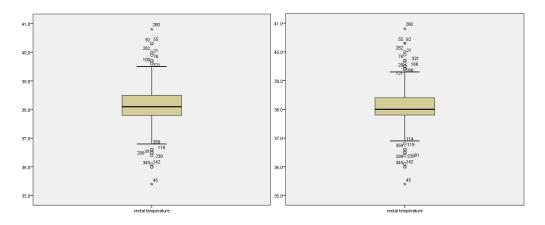
# 2. 用最高频率值来填补缺失值

首先通过对每组数值属性进行词频统计,找出该组数值属性的最高频值,并用该值填补缺失值。

#### (1) rectal temperature:

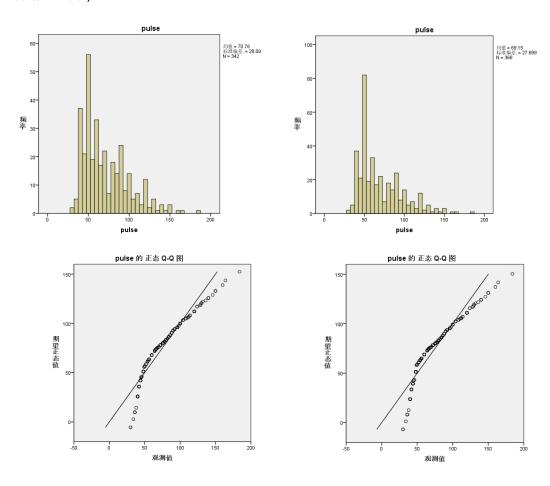
该属性的最高频值为 38.0,用该值填充缺失值后,与省略缺失值图的对比如下(左侧 为省略缺失图):

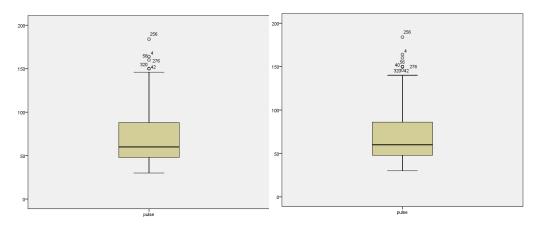




### (2) pulse:

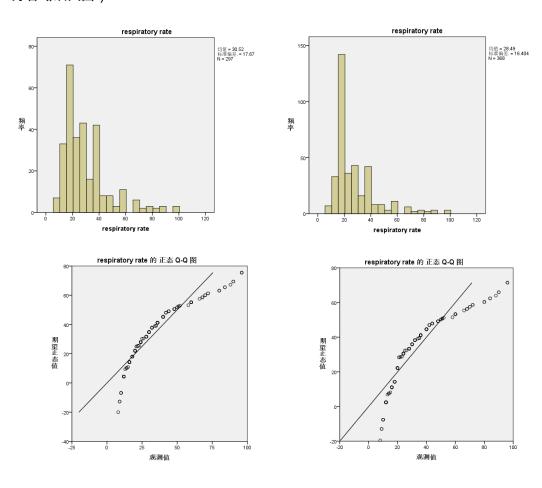
该属性的最高频值为 48,用该值填充缺失值后,与省略缺失值图的对比如下(左侧为省略缺失图):

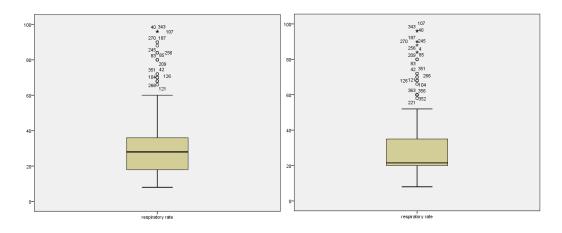




### (3) respiratory rate:

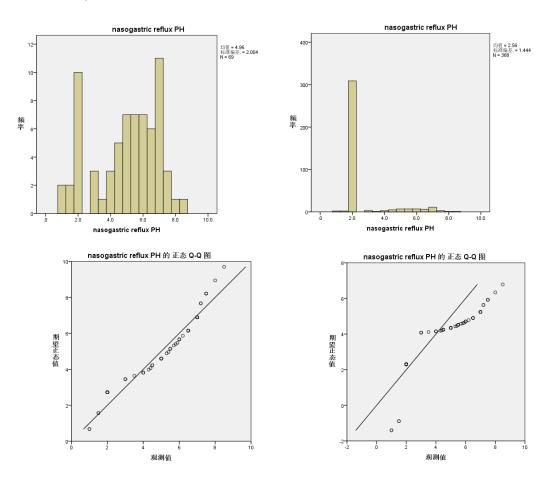
该属性的最高频值为 20.0,用该值填充缺失值后,与省略缺失值图的对比如下(左侧 为省略缺失图):

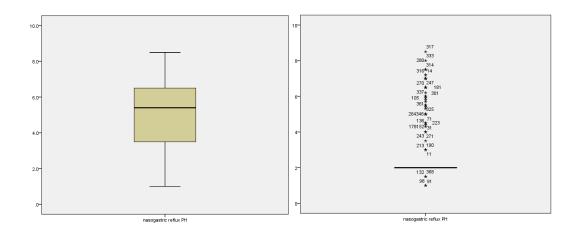




### (4) nasogastric reflux PH:

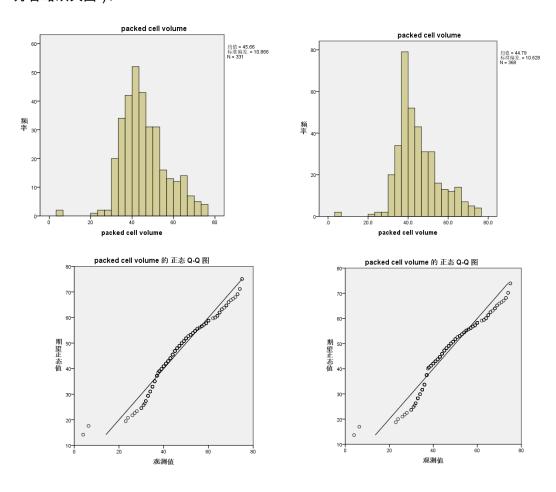
该属性的最高频值为 2.0,用该值填充缺失值后,与省略缺失值图的对比如下(左侧为 省略缺失图):

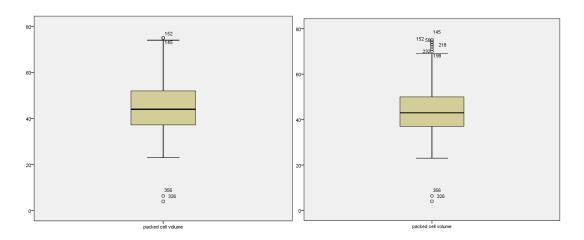




### (5) packed cell volume:

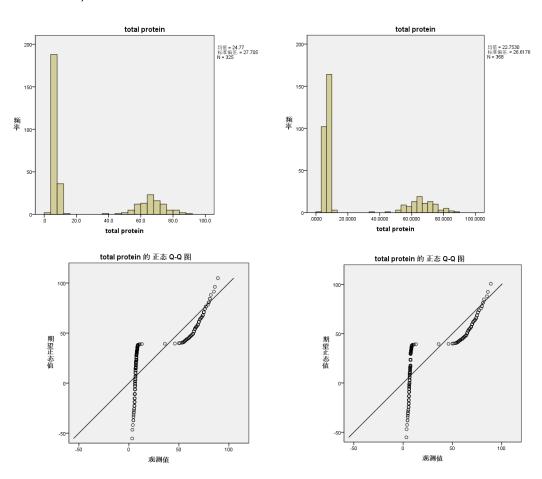
该属性的最高频值为 37.0,用该值填充缺失值后,与省略缺失值图的对比如下(左侧 为省略缺失图):

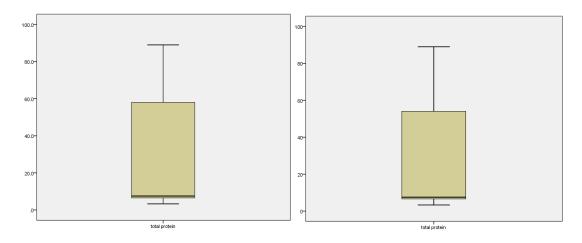




### (6) total protein:

该属性的最高频值为 7.5, 用该值填充缺失值后,与省略缺失值图的对比如下(左侧为省略缺失图):





### (7) abdomcentesis total protein:

该属性的最高频值为 2.0,用该值填充缺失值后,与省略缺失值图的对比如下(左侧为 省略缺失图):

