In [274]:

```
import pandas as pd
import numpy as np
import scipy.stats as stats
import pylab

#读取数据
df=pd.read_csv('horse-colic.data.txt',sep=' ')
attrname=['surgery','Age','Hospital_Number','rectal_temperature','pulse','respirato:
labelattr=[1,2,3,7,8,9,10,11,12,13,14,15,17,18,21,23,24,25,26,27,28]
valueattr=[4,5,6,16,19,20,22]
```

```
In [196]:
#分析标签属性的出现频数
for attrid in labelattr:
    print(df[attrname[attrid-1]].value counts())
    print()
1
     180
2
     119
       1
Name: surgery, dtype: int64
     276
1
9
      24
Name: Age, dtype: int64
529796
            2
            2
529424
5279822
            2
            2
527916
530526
            2
            2
528996
528729
            2
            2
528151
            2
529461
528931
            2
            2
528890
528469
            2
            2
530693
532349
            2
            2
528904
            2
527544
533697
            1
533696
            1
533736
            1
            1
534719
530101
            1
529399
            1
533692
            1
            1
528570
527957
            1
534157
            1
521399
            1
529045
            1
530612
            1
528047
            1
           . .
533887
           1
535031
            1
530301
            1
            1
5294369
530297
            1
529272
            1
            1
535415
530294
            1
534899
            1
            1
529777
```

```
53485/
            1
530276
            1
535392
            1
535338
            1
527709
            1
527706
            1
            1
533847
            1
528214
535381
            1
            1
533750
            1
527698
530255
            1
            1
530254
533836
            1
530251
            1
            1
535043
Name: Hospital Number, dtype: int64
3
     109
1
      78
?
      56
2
      30
4
      27
Name: temperature_of_extremities, dtype: int64
1
     115
3
     103
?
      69
4
       8
2
       5
Name: peripheral_pulse, dtype: int64
1
     79
3
     58
?
     47
4
     41
     30
2
5
     25
6
Name: mucous_membranes, dtype: int64
1
     188
2
      78
?
      32
3
       2
Name: capillary_refill_time, dtype: int64
3
     67
2
     59
?
     55
5
     42
4
     39
     38
1
Name: pain, dtype: int64
3
     128
4
      73
?
      44
1
      39
2
      16
Name: peristalsis, dtype: int64
```

```
1
     76
2
     65
3
     65
?
     56
4
     38
Name: abdominal_distension, dtype: int64
?
     104
2
     102
1
      71
3
      23
Name: nasogastric_tube, dtype: int64
1
     120
?
     106
3
      39
2
      35
Name: nasogastric_reflux, dtype: int64
?
     102
4
      79
1
      57
3
      49
2
      13
Name: rectal_examination, dtype: int64
?
     118
5
      79
4
      43
      28
1
2
      19
      13
3
Name: abdomen, dtype: int64
?
     165
2
      48
3
      46
1
Name: abdominocentesis_appearance, dtype: int64
1
     178
2
      77
3
      44
?
        1
Name: outcome, dtype: int64
     191
1
     109
Name: surgical_lesion, dtype: int64
0
          56
3111
          33
3205
          29
2208
          20
          13
2205
4205
          11
2209
          11
           9
2124
           8
1400
           7
31110
```

```
2113
           6
           5
2112
           5
400
3209
           4
           4
4300
           4
2206
           4
5400
           3
3112
           3
4124
           3
2111
           3
2207
7209
           3
           2
5206
           2
5124
3124
           2
           2
5111
9400
           2
           2
6111
2322
           2
          . .
3025
           2
8400
           2
           2
6112
11300
           1
4122
           1
           1
7113
6209
           1
3115
           1
5000
           1
           1
3133
4111
           1
3400
           1
300
           1
12208
           1
           1
9000
           1
5205
           1
1111
1124
           1
8300
           1
2305
           1
4206
           1
4207
           1
           1
21110
2300
           1
           1
3207
11400
           1
           1
7400
           1
3113
3300
           1
41110
           1
Name: #1_lesion, dtype: int64
0
        293
           3
3111
           1
6112
7111
           1
           1
1400
           1
3112
Name: #2_lesion, dtype: int64
         299
```

http://10.108.20.54:9999/notebooks/datamining_ex1/datamining.ipynb

2209 1

Name: #3_lesion, dtype: int64

2 2011 99

Name: cp_data, dtype: int64

In [263]:

```
#数值属性的最小值,1/4分位数,中位数,均值,3/4分位数,最大值
for attrid in valueattr:
    print(attrname[attrid - 1])
    series=df[attrname[attrid - 1]].apply(pd.to numeric, errors='coerce')
    series=series(series.notnull())
    print('min:',series.min())
    print('1/4 quantile:', series.quantile(0.25))
    print('mean:',series.mean())
    print('median:',series.median())
    print('3/4 quantile:', series.quantile(0.75))
    print('max:',series.max())
    print()
rectal temperature
min: 35.4
1/4 quantile: 37.8
mean: 38.16791666666669
median: 38.2
3/4 quantile: 38.5
max: 40.8
pulse
min: 30.0
1/4 quantile: 48.0
mean: 71.91304347826087
median: 64.0
3/4 quantile: 88.0
max: 184.0
respiratory_rate
min: 8.0
1/4 quantile: 18.5
mean: 30.417355371900825
median: 24.5
3/4 quantile: 36.0
max: 96.0
nasogastric reflux PH
min: 1.0
1/4 quantile: 3.0
mean: 4.707547169811321
median: 5.0
3/4 quantile: 6.5
max: 7.5
packed_cell_volume
min: 23.0
1/4 quantile: 38.0
mean: 46.29520295202952
median: 45.0
3/4 quantile: 52.0
max: 75.0
total protein
min: 3.3
1/4 quantile: 6.5
mean: 24.456928838951317
median: 7.5
3/4 quantile: 57.0
```

max: 89.0

abdomcentesis_total_protein

min: 0.1

1/4 quantile: 2.0

mean: 3.0196078431372553

median: 2.25
3/4 quantile: 3.9

max: 10.1

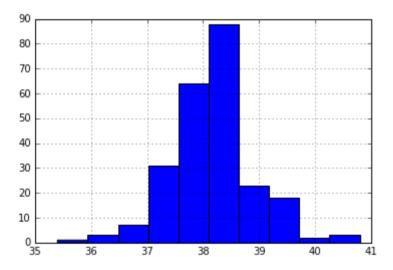
In [203]:

```
#直方图
attrid=4
print(attrname[attrid - 1])
series=df[attrname[attrid - 1]]
series=series[series != '?'].apply(pd.to_numeric, errors='coerce')
series.hist()
```

rectal_temperature

Out[203]:

<matplotlib.axes._subplots.AxesSubplot at 0x7f2507e6eb70>



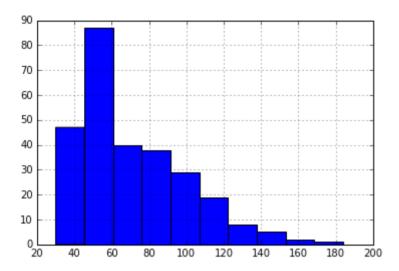
In [204]:

```
#直方图
attrid=5
print(attrname[attrid - 1])
series=df[attrname[attrid - 1]]
series=series[series != '?'].apply(pd.to_numeric, errors='coerce')
series.hist()
```

pulse

Out[204]:

<matplotlib.axes._subplots.AxesSubplot at 0x7f2507df9128>



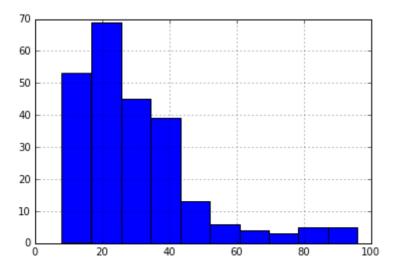
In [205]:

```
#直方图
attrid=6
print(attrname[attrid - 1])
series=df[attrname[attrid - 1]]
series=series[series != '?'].apply(pd.to_numeric, errors='coerce')
series.hist()
```

respiratory_rate

Out[205]:

<matplotlib.axes._subplots.AxesSubplot at 0x7f2507d03400>



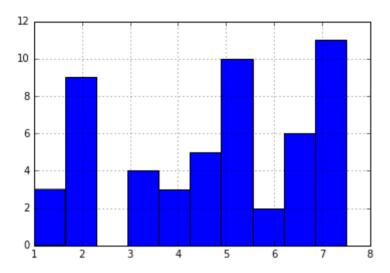
In [206]:

```
#直方图
attrid=16
print(attrname[attrid - 1])
series=df[attrname[attrid - 1]]
series=series[series != '?'].apply(pd.to_numeric, errors='coerce')
series.hist()
```

nasogastric_reflux_PH

Out[206]:

<matplotlib.axes._subplots.AxesSubplot at 0x7f2507ca0748>



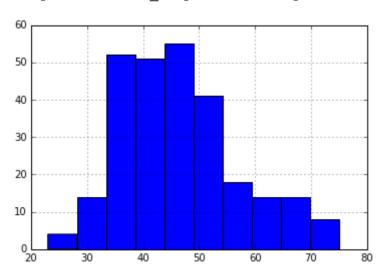
In [207]:

```
#直方图
attrid=19
print(attrname[attrid - 1])
series=df[attrname[attrid - 1]]
series=series[series != '?'].apply(pd.to_numeric, errors='coerce')
series.hist()
```

packed_cell_volume

Out[207]:

<matplotlib.axes._subplots.AxesSubplot at 0x7f2507c26240>



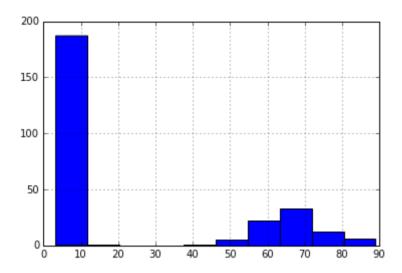
In [208]:

```
#直方图
attrid=20
print(attrname[attrid - 1])
series=df[attrname[attrid - 1]]
series=series[series != '?'].apply(pd.to_numeric, errors='coerce')
series.hist()
```

total_protein

Out[208]:

<matplotlib.axes._subplots.AxesSubplot at 0x7f2507c16ef0>



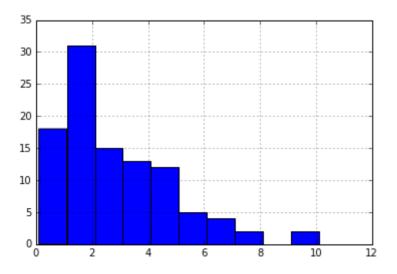
In [209]:

```
#直方图
attrid=22
print(attrname[attrid - 1])
series=df[attrname[attrid - 1]]
series=series[series != '?'].apply(pd.to_numeric, errors='coerce')
series.hist()
```

 $\verb|abdomcentesis_total_protein|\\$

Out[209]:

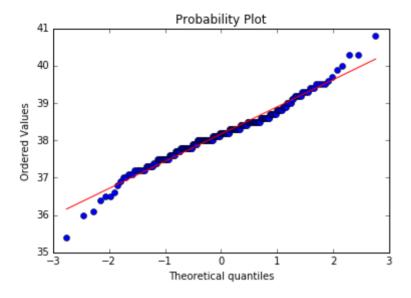
<matplotlib.axes._subplots.AxesSubplot at 0x7f2507d7b550>



In [231]:

```
#qq
attrid=4
print(attrname[attrid - 1])
series=df[attrname[attrid - 1]]
series=series[series != '?'].apply(pd.to_numeric, errors='coerce')
_=stats.probplot(series, dist="norm", plot=pylab)
```

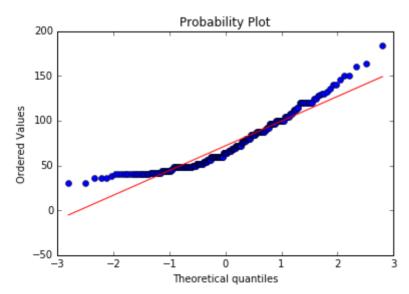
rectal_temperature



In [230]:

```
#qq
attrid=5
print(attrname[attrid - 1])
series=df[attrname[attrid - 1]]
series=series[series != '?'].apply(pd.to_numeric, errors='coerce')
_=stats.probplot(series, dist="norm", plot=pylab)
```

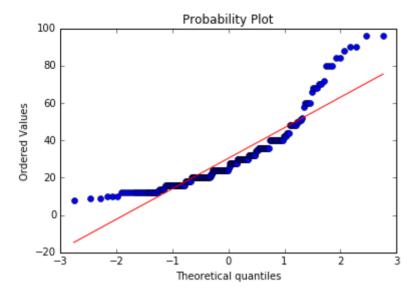
pulse



In [229]:

```
#qq
attrid=6
print(attrname[attrid - 1])
series=df[attrname[attrid - 1]]
series=series[series != '?'].apply(pd.to_numeric, errors='coerce')
_=stats.probplot(series, dist="norm", plot=pylab)
```

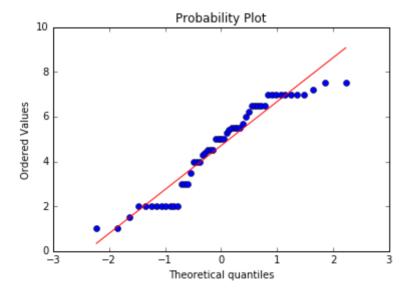
respiratory_rate



In [228]:

```
#qq
attrid=16
print(attrname[attrid - 1])
series=df[attrname[attrid - 1]]
series=series[series != '?'].apply(pd.to_numeric, errors='coerce')
_=stats.probplot(series, dist="norm", plot=pylab)
```

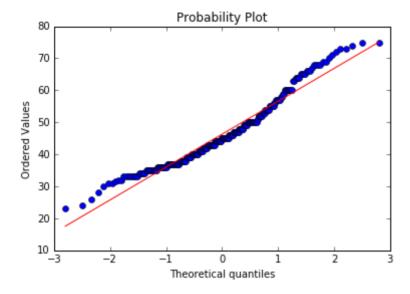
nasogastric_reflux_PH



In [227]:

```
#qq
attrid=19
print(attrname[attrid - 1])
series=df[attrname[attrid - 1]]
series=series[series != '?'].apply(pd.to_numeric, errors='coerce')
_=stats.probplot(series, dist="norm", plot=pylab)
```

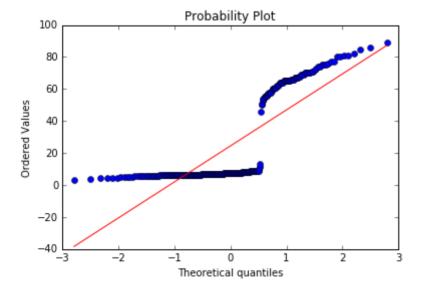
packed_cell_volume



In [226]:

```
#qq
attrid=20
print(attrname[attrid - 1])
series=df[attrname[attrid - 1]]
series=series[series != '?'].apply(pd.to_numeric, errors='coerce')
_=stats.probplot(series, dist="norm", plot=pylab)
```

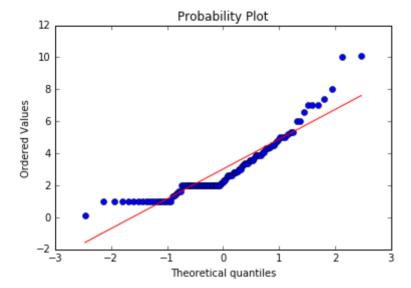
total_protein



In [233]:

```
#qq
attrid=22
print(attrname[attrid - 1])
series=df[attrname[attrid - 1]]
series=series[series != '?'].apply(pd.to_numeric, errors='coerce')
_=stats.probplot(series, dist="norm", plot=pylab)
```

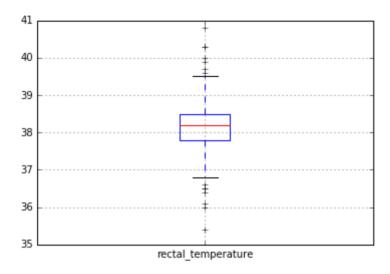
$\verb|abdomcentesis_total_protein|\\$



In [264]:

```
#盒图
attrid=4
print(attrname[attrid - 1])
series=df[attrname[attrid - 1]]
series=series[series != '?'].apply(pd.to_numeric, errors='coerce')
_=pd.DataFrame(series).boxplot()
```

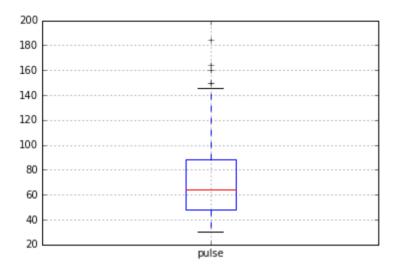
rectal_temperature



In [265]:

```
#qq
attrid=5
print(attrname[attrid - 1])
series=df[attrname[attrid - 1]]
series=series[series != '?'].apply(pd.to_numeric, errors='coerce')
_=pd.DataFrame(series).boxplot()
```

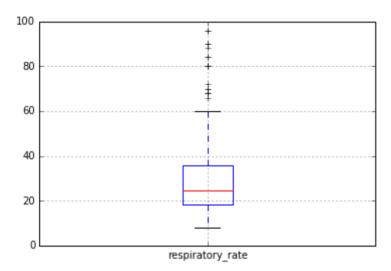
pulse



In [266]:

```
#qq
attrid=6
print(attrname[attrid - 1])
series=df[attrname[attrid - 1]]
series=series[series != '?'].apply(pd.to_numeric, errors='coerce')
_=pd.DataFrame(series).boxplot()
```

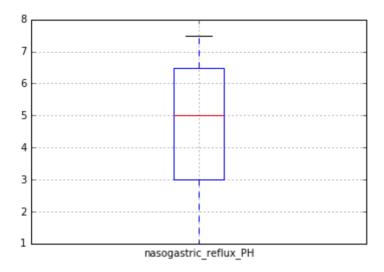
respiratory_rate



In [267]:

```
#qq
attrid=16
print(attrname[attrid - 1])
series=df[attrname[attrid - 1]]
series=series[series != '?'].apply(pd.to_numeric, errors='coerce')
_=pd.DataFrame(series).boxplot()
```

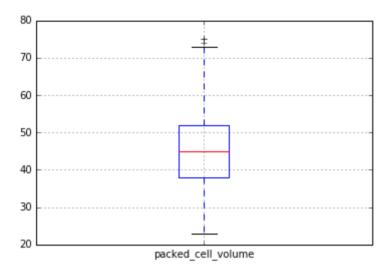
nasogastric_reflux_PH



In [268]:

```
#qq
attrid=19
print(attrname[attrid - 1])
series=df[attrname[attrid - 1]]
series=series[series != '?'].apply(pd.to_numeric, errors='coerce')
_=pd.DataFrame(series).boxplot()
```

packed_cell_volume



In [269]:

```
#qq
attrid=20
print(attrname[attrid - 1])
series=df[attrname[attrid - 1]]
series=series[series != '?'].apply(pd.to_numeric, errors='coerce')
_=pd.DataFrame(series).boxplot()
```

total_protein

