Data Cleaning Part 2

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Data Cleansing involves the following aspects:

- missing values
- data formatting
- data normalization
- data standardization
- data binning
- remove duplicates

Data Standardization

Standardization transforms data to have a mean of zero and a standard deviation of 1.

Techniques for standardization

- z-score
- z-map

z-score

The new value is calculated as the difference between the current value and the average value, divided by the standard deviation.

We can use the zscore() function of the scipy.stats library.

Example

from scipy.stats import zscore
df['Value'] = zscore(df['Value'])



MEAN: 2.66 STD: 1.25

z-map

The new value is calculated as the difference between the current value and the average value of a comparison array, divided by the standard deviation of a comparison array.

We can use the **zmap()** function of the **scipy.stats** library.

Example

from scipy.stats import zmap
df['Value'] = zmap(df['Value'], df['Count'])

Value	Count	Value	Count
1	3	-3.67	3
3	4	-1.22	4
4	5	0	5

Data Binning

Data binning (or bucketing) groups data in bins (or buckets), in the sense that it replaces values contained into a small interval with a single representative value for that interval.

Binning

Binning can be applied to convert numeric values to categorical or to sample (quantize) numeric values.

Binning is a technique for data smoothing. Data smoothing is employed to remove noise from data. Three techniques for data smoothing:

- binning
- regression
- outlier analysis

Techniques for binning

- convert numeric to categorical
 - binning by distance
 - binning by frequency
- reduce numeric values
 - \circ sampling

Binning by distance - cut()

- Define the bin edges
- Convert numeric into categorical variables
- Define the number of bins and the associated labels



Example

import numpy as np

bins = [0, 50, 100, 500, 1000]

labels = ['small', 'medium', 'large','very large']

df['Size'] = pd.cut(df['Size'] , bins=bins, labels=labels, include lowest=True)

Example 2 - Linear Space among ranges

```
min value = df['Size'].min()
max value = df['Size'].max()
n bins = 4
bins = np.linspace(min value,max value,n bins+1)
array([ 5., 336.66666667, 668.33333333, 1000.])
labels = ['small', 'medium', 'large','very large']
df['Size'] = pd.cut(df['Size'] , bins=bins, labels=labels,
include lowest=True)
```

Size	# bins = 4	Size
1000	Label Ranges	very large
5	small 0 - 5	small
500	medium 5 - 336.67	medium
100	large 336.67-668.33	small
250	very large 668.33 - 1000	small
400		medium

Binning by frequency - qcut()

- Quantile-based discretization function
- Calculate the size of each bin so that each bin contains (almost) the same number of observations, but the bin range will vary.

Example



labels = ['small', 'medium', 'large','very large']

n bins = 4

df['Size'] = pd.qcut(df['Size'], q=n_bins,precision=1, labels=labels)

We can set the precision parameter to define the number of decimal points.

Sampling

It permits to reduce the number of samples, by grouping similar values or contiguous values. There are three approaches to perform sampling:

- by bin means: each value in a bin is replaced by the mean value of the bin.
- by bin median: each bin value is replaced by its bin median value.
- by bin boundary: each bin value is replaced by the closest boundary value, i.e. maximum or minimum value of the bin.

binned_statistics()

- We exploit the binned_statistic() function of the scipy.stats package can be used.
- This function receives two arrays as input, x_data and y_data, as well as the statistics to be used (e.g. median or mean) and the number of bins to be created.
- The function returns the values of the bins as well as the edges of each bin.

Example



from scipy.stats import binned_statistic

```
x data = np.arange(0, len(df))
```

```
y_data = df['Size']
```

```
x_bins,bin_edges, misc = binned_statistic(y_data,x_data,
statistic="median", bins=4)
```

bin_intervals = pd.IntervalIndex.from_arrays(bin_edges[:-1], bin_edges[1:],closed='both')

```
IntervalIndex([[5.0, 253.75], [253.75, 502.5], [502.5,
751.25], [751.25, 1000.0]])
```

def set_to_median(x, bin_intervals):

for interval in bin intervals:

if x in interval:

return interval.mid

```
df['sampled_size''] = df['Size'].apply(lambda x:
set to median(x, bin intervals))
```



Natural breaks in data

We can use the package jenkspy, which contains a single function, called jenks_breaks(), which calculates the natural breaks of an array, exploiting the Fisher-Jenks algorithm.

We can install the package by running pip3 install jenkspy.

Example

import jenkspy

breaks = jenkspy.jenks_breaks(df['Size'], nb_class=3)

df['size_break'] = pd.cut(df['Size'] , bins=breaks, labels=labels, include_lowest=True)

Remove Duplicates

Remove all rows that appear at least twice.

The concept of duplicate

Index	Name	Surname	Value
1	Mark	Grenn	3
2	Mark	Grenn	3
3	Mark	Grenn	4

Rows 1 and 2 are duplicates

Row 1, 2 and 3 are duplicates in column Name and Surname

Drop duplicates on the basis of all columns

keep just one row for each duplicate

Name	Surname	Value
Mark	Grenn	3

Do not maintain any row for the duplicate

Name	Surname	Value
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Drop duplicates on the basis of the Name and Surname Columns

Keep just one value for column

Name	Surname	Value
Mark	Grenn	3
Mark	Grenn	4

Do not maintain any row for the duplicate

Name	Surname	Value
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drop_duplicates()

- df1 = df.drop_duplicates()
- df2 = df.drop_duplicates(keep=False)
- df3 = df.drop duplicates(subset=["Name", "Surname"])

df4 = df.drop_duplicates(subset=["Name", "Surname"],
keep=False)