device_failure_prediction

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Note that data manupulation and exploration techniques illustrated in this demonstraton does not mean that the audience have to follow the same ways as shown here.

1 Device Failure and Maintenance Prediction Model

It is to build a predictive model or models with diagnoses of telemetry attributes to classify whether maintenance should be performed on devices or not. The column to be use to predict is called "failure" with binary value 0 for "non-failure" and 1 for "failure". *The goal is to minimize false positives and false negatives*.

2 Assumption

As there is no meta data for the description of the current dataset, assumptions can be applied to the current dataset. The dataset consists of diagnoses of telemetry attributes, so it might be rational that some variables are assumed to consist of *categorical nominal type values* while other variables would consist of *continuous type values*.

3 Roadmap

This demonstration is to show how to handle datasets which are imbalanced. The provided dataset has no description for variables. The dataset will be analyzed by using various statistical approaches. The main issue with the current dataset is that it is highly imblanaced. Various sampling techniques usually applied to the imbalanced dataset. There will be detailed explanations with each sampling techniques as the analyses progresses further. In this demonstration, data exploration will only be extensively explained. In the following successive sections, first, no sampling technique will be applied in this demonstration to understand how sampling techniques will affect performances of the predictive model on an imblanaces dataset. Although data preprocessing is done for most datasets but there will be no data preprocessing for this demonstration.

Additive variable generations such as day, day of the week, month, device operation days, or season from the 'date' variable could be an option to enhance the model performance. However, additivie variable generations does not always have to be applied to introduce new variables to the original dataset, unless there is any acceptable model performance with the original dataset. One could introduce additive variables before any model development but there is no guarantee that those additive variables are going to improve the model performances.

However, it should be noted that normalizations for continuous values shoud not be confused with additive variable generations. One might be also tempted to apply binning to continuous values but it sould be applied when there is some group of values that are significantly different from other groups of values in a variable. Otherwise, binning would not be much help and it will only cause lose of information of continuity in the values. Distribution transformations, for example, log-normal transformation, could be used but it is not necessary to be applied for Randome Forest learning algorithm. It all depends on what the analyzer intend to do with the dataset.

Logsitic Regression and Neural Network based algorithms could also have been used but assuming that values of some variables are categorical nominal type values, to properly use those algorithms, creating dummy or one-hot-encoded variables for those algorithms simply overwhelms contraints on the given task. Therefore, for the predictive model development, instead of using Losigitic Regression or Neural Network based algorithms, Random Forest predictive model learning algorithm will be adopted because Random Forest can handle very well for both categorical nominal type values and continuous type values at the same time. Random Forest does not require continuous type values to be normalized at all.

4 Data Loading

import necessary python modules.

```
[25]: import pandas as pd import numpy as np
```

Load the dataset into memory.

```
[26]: datafile = r'/home/thomas/Downloads/device_failure.csv'
dataset = pd.read_csv(datafile, sep=',', engine='python')
```

5 Data Exploration

See if the dataset has been loaded correctly. There should be 12 columns meaning 12 variables.

					-						
3]:	da	taset.head()									
3]:		date	device	failure	attribute1		attr	ibute2	attr	ibute3	\
	0	2015-01-01	S1F01085	0	215	630672	56			0	
	1	2015-01-01	S1F0166B	0	61	370680	0			3	
	2	2015-01-01	S1F01E6Y	0	173	295968		0		0	
	3	2015-01-01	S1F01JE0	0	79	694024		0		0	
	4	2015-01-01	S1F01R2B	0	135	970480		0		0	
		attribute4	attribute5	attrib	ute6	attrib	ute7	te7 attribut		attrib	ute9
	0	52	6	40	7438		0		0		7
	1	0	6	40	403174		0	0			0
	2	0	12	23	7394		0		0		0
	3	0	6	41	0186		0		0		0
	4	0	15	31	3173		0	0			3

Take a look at the number of total records.

```
[4]: dataset.failure.count()
```

[4]: 124494

Take a look at the data structure.

[5]: dataset.dtypes

```
[5]: date
                   object
    device
                   object
    failure
                    int64
                    int64
    attribute1
    attribute2
                    int64
                    int64
    attribute3
                    int64
    attribute4
    attribute5
                    int64
                    int64
    attribute6
    attribute7
                    int64
    attribute8
                    int64
    attribute9
                    int64
    dtype: object
```

There are 12 variables in the dataset. All of the variables in the dataset are all integer values except date and device variables. Let's take a look at the distribution of the dataset in terms of failure and non-failure.

Non-failure: 124388 [99.91%] Failure: 106 [0.09%]

```
[8]: value_distribution.plot(kind='bar')
```

[8]: <matplotlib.axes._subplots.AxesSubplot at 0x7f50c88322b0>

The number of device failures is less than 0.1% and the dataset appears to be highly imbalanced. As shown above, there are 12 variables. Failure is the target variable and 11 variables are considered to be independent variables. All the independent variables seems to be integers. The next things to do is to see if there are any null values or missing values in columns of the dataset. The following two lines will figure out if there any null and missing values in the dataset.

```
[30]: dataset.isnull().any()
```

```
[30]: date False device False failure False attribute1 False attribute2 False attribute3 False
```

```
False
     attribute4
                    False
     attribute5
     attribute6
                    False
     attribute7
                    False
                    False
     attribute8
     attribute9
                    False
     dtype: bool
[31]: dataset.isna().any()
[31]: date
                    False
     device
                    False
     failure
                    False
     attribute1
                    False
     attribute2
                    False
     attribute3
                    False
     attribute4
                    False
                    False
     attribute5
                    False
     attribute6
     attribute7
                    False
                    False
     attribute8
     attribute9
                    False
     dtype: bool
```

In order to confirm that there are no missing and null values in every variable, the following can be executed for a list of all the summations of each column:

```
[32]: dataset.isnull().sum()
[32]: date
                     0
     device
                     0
     failure
                     0
     attribute1
                     0
     attribute2
                     0
     attribute3
                     0
     attribute4
                     0
     attribute5
                     0
                     0
     attribute6
     attribute7
                     0
     attribute8
                     0
     attribute9
                     0
     dtype: int64
```

It is clear to say that there are no null values in the dataset. Currently the meaning of each variable is unknown and it is definitely required to explore each variable to find out what each variable would mean. There are 106 samples, approaximately 0.0852%, indicating failure. The above distribution shows that the dataset is perfectly imbalanced. In order to grasp the overall representation of the dataset, it is favorable to look into the characteristics of each variable, first. The following lines will first reveal the number of unique values in each variable and will give an overall idea of how each variable contributes to failure and non-failure.

```
[169]: # Show distinct values of each variable.
columns = dataset.columns
no_unique_values = []
for column in columns:
    unique_values = np.unique(dataset[column])
    print('{}: {} distinct values - {}'.format(column, len(unique_values),
    unique_values))
    no_unique_values.append(len(unique_values))
```

```
date: 304 distinct values - ['2015-01-01' '2015-01-02' '2015-01-03'
'2015-01-04' '2015-01-05'
 '2015-01-06' '2015-01-07' '2015-01-08' '2015-01-09' '2015-01-10'
 '2015-01-11' '2015-01-12' '2015-01-13' '2015-01-14' '2015-01-15'
 '2015-01-16' '2015-01-17' '2015-01-18' '2015-01-19' '2015-01-20'
 '2015-01-21' '2015-01-22' '2015-01-23' '2015-01-24' '2015-01-25'
 '2015-01-26' '2015-01-27' '2015-01-28' '2015-01-29' '2015-01-30'
 '2015-01-31' '2015-02-01' '2015-02-02' '2015-02-03' '2015-02-04'
 '2015-02-05' '2015-02-06' '2015-02-07' '2015-02-08' '2015-02-09'
 '2015-02-10' '2015-02-11' '2015-02-12' '2015-02-13' '2015-02-14'
 '2015-02-15' '2015-02-16' '2015-02-17' '2015-02-18' '2015-02-19'
 '2015-02-20' '2015-02-21' '2015-02-22' '2015-02-23' '2015-02-24'
 '2015-02-25' '2015-02-26' '2015-02-27' '2015-02-28' '2015-03-01'
 '2015-03-02' '2015-03-03' '2015-03-04' '2015-03-05' '2015-03-06'
 '2015-03-07' '2015-03-08' '2015-03-09' '2015-03-10' '2015-03-11'
 '2015-03-12' '2015-03-13' '2015-03-14' '2015-03-15' '2015-03-16'
 '2015-03-17' '2015-03-18' '2015-03-19' '2015-03-20' '2015-03-21'
 '2015-03-22' '2015-03-23' '2015-03-24' '2015-03-25' '2015-03-26'
 '2015-03-27' '2015-03-28' '2015-03-29' '2015-03-30' '2015-03-31'
 '2015-04-01' '2015-04-02' '2015-04-03' '2015-04-04' '2015-04-05'
 '2015-04-06' '2015-04-07' '2015-04-08' '2015-04-09' '2015-04-10'
 '2015-04-11' '2015-04-12' '2015-04-13' '2015-04-14' '2015-04-15'
 '2015-04-16' '2015-04-17' '2015-04-18' '2015-04-19' '2015-04-20'
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 '2015-04-26' '2015-04-27' '2015-04-28' '2015-04-29' '2015-04-30'
 '2015-05-01' '2015-05-02' '2015-05-03' '2015-05-04' '2015-05-05'
 '2015-05-06' '2015-05-07' '2015-05-08' '2015-05-09' '2015-05-10'
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 '2015-06-25' '2015-06-26' '2015-06-27' '2015-06-28' '2015-06-29'
 '2015-06-30' '2015-07-01' '2015-07-02' '2015-07-03' '2015-07-04'
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 '2015-08-24' '2015-08-25' '2015-08-26' '2015-08-27' '2015-08-28'
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 '2015-09-08' '2015-09-09' '2015-09-10' '2015-09-11' '2015-09-12'
 '2015-09-13' '2015-09-14' '2015-09-15' '2015-09-16' '2015-09-17'
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 '2015-10-03' '2015-10-04' '2015-10-05' '2015-10-06' '2015-10-07'
 '2015-10-08' '2015-10-09' '2015-10-10' '2015-10-11' '2015-10-12'
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 '2015-10-18' '2015-10-19' '2015-10-20' '2015-10-21' '2015-10-22'
 '2015-10-23' '2015-10-24' '2015-10-25' '2015-10-26' '2015-10-27'
 '2015-10-29' '2015-10-30' '2015-10-31' '2015-11-02']
device: 1169 distinct values - ['S1F01085' 'S1F013BB' 'S1F0166B' ... 'Z1F26YZB'
'Z1F282ZV' 'Z1F2PBHX']
failure: 2 distinct values - [0 1]
attribute1: 123877 distinct values - [
                                                 0
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244136552 244138600 244140480]
attribute2: 558 distinct values - [
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 14856 14904 14920 15304 15328 15336 16408 16864 17408 17704 18072 19200
 21200 21528 21544 21816 21928 21944 22816 23208 23400 23936 24656 24680
 24792 24904 24920 24976 24984 27856 28344 32936 33000 33016 33024 33528
 34912 35616 37936 41232 43712 44008 44024 44152 44176 44224 44232 46264
 46280 46296 46304 46328 46400 46408 46496 46520 46584 46848 46872 47944
49768 49840 51976 54752 54896 56584 56736 61592 62296 64464 64472 64584
 64728 64736 64776 64784 64792 64968]
attribute3: 47 distinct values - [
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attribute4: 115 distinct values - [
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                                  431
                                             487
                                                  521
                                                        529
                                                             533
                                                                   763
                                                                        841
                                       486
 1033 1074 1666]
attribute5: 60 distinct values - [ 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15
16 17 18 19 20 21 22 23 24
 25 29 30 31 32 33 34 35 36 37 38 39 40 41 42 57 58 59 60 61 62 63 64 65
66 67 68 70 78 89 90 91 92 94 95 98]
```

attribu	te6: 4	8	ç)	12	689035 689062									
689161]															
attribu	te7: 2	8 dist	tinct v	values ·	- [0 6	8	16	22	24	32	40	48	56	72
80 96 104 112 128 136 152															
176 216 240 272 312 424 496 736 744 832]															
attribu	te8: 2	8 dist	tinct v	values ·	- [0 6	8	16	22	24	32	40	48	56	72
80 96	104 11	2 128	136 1	52											
176 216 240 272 312 424 496 736 744 832]															
attribu	te9: 6	5 dist	tinct v	values -	- [0	1		2	3	4	4	5		6
7 8	9	10) 1:	1											
12	13	14	15	18	19	20	2	1	22	2	3	24	2	5	
26	30	33	34	38	41	42	5	1	52	5	7	65	7	0	
98	104	120	145	155	164	177	20	5	222	23	3 2	241	24	8:	
255	263	269	400	898	1080	1150	116	5 1	1864	2269	9 22	270	252	2	
2637	2794	7226	10137	18701]											

Summary of unique values

- date: 304 unique values.
- devce: 1,169 unique values.
- attribute1: 123,877 unique values.
- attribute2: 558 unique values.
- attribute3: 47 unique values.
- attribute4: 115 unique values.
- attribute5: 60 unique values.
- attribute6: 44,838 unique values.
- attribute7: 28 unique values.
- attribute9: 65 unique values.

For a dataset with a large number of variables, it would be extremly hard to manually look at all the values of each variable in the dataset. Nonetheless, data exploration for understanding characteristics of individual variables is mandatory to undergo when a story telling about the current dataset is required. However, the current dataset only has 9 variables excluding 'date', 'device', and 'failure' variables, so it is reasonble to look at individual characteristics of unique values of each variable.

- 'attribute1' and 'attribute6' have relatively high numbers of unique values while numbers of unique values for other values are small. Those two variables must have a lot of ways for the predictive model to learn non-failures of devices. The predictive model must not only know how to tell devices with failure but it also has to learn ways to failure, too. A brief unique value summary will help grasp a further picture of the dataset.
- The ranges of unique values of 'attribute1' and 'attribute6' are much broader than the ranges of unique values of other variables.
- 'attribute7' and 'attribute8' seem to be idential to one another. Either of the variables, 'attribute7' or 'attribute8', should be removed from analysis if those variables are used by Logistic Regression or Ordinal Regression since those two analyses strongly assume little or no multicollinearity.

- 'attribute1' appears to be multiples of 'attribute2', either of those two variables should also be remove from the dataset unless only one of them should be used. Otherwise, they will cause a multicollinearity problem if Logistic Regression or Ordinal Regression analyses are implemented for the predictive model. 'attribute1' seems to express the amount of daily data transmission handled by each device.
- Although Logistics Regression and Ordinary Regression are not applied here, but those
 multi-collinearity issues should be noted. When Logitics Regression or Ordinary Regression is undertaken, multi-collinearity tests must be performed for a reliable multicollinearity assiciation to ensure the predictive model performance.

Let's first check if 'attribute7' and 'attribute8' are identical to one another.

```
[23]: # Create two different dataframes for attribute7 and attribute8.

attribute_7 = pd.DataFrame(dataset, columns = ['attribute7'])

attribute_8 = pd.DataFrame(dataset, columns = ['attribute8'])

attribute_test = pd.DataFrame(dataset, columns = ['matched'])

# Compare attribute7 and attribute8 by their values for each record.

attribute_test['matched'] = pd.DataFrame(np.where(attribute_7.attribute7 == □

→attribute_8.attribute8, 'True', 'False'))

[24]: # See the number of matched values.

attribute_test['matched'].value_counts()

[24]: True 124494
```

[24]: True 124494

Name: matched, dtype: int64

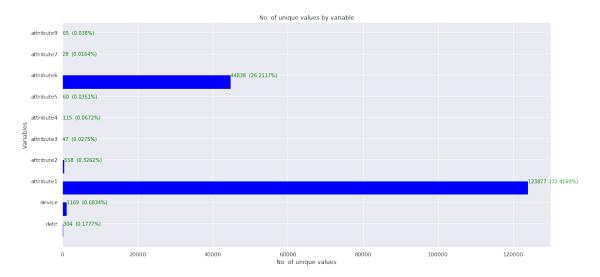
It is safe to say 'attribute7' is a duplicate of 'attribute8' or the other way around. As mentioned earlier above, if a variable is a copy of another variable, it could be problematic when the predictive model is developed using Logistic Regression Classifier since Logistic Regression assumes little or no multi-collinearity among independent variables, so either of them should be removed from the dataset or they both can stay in the dataset as long as only one of them is applied to the model development.

If the model is developed by using a ensemble classfier such as Random Forest, the effect of multi-collinearity would not have much impact on the preditive model since Random Forest will create the predictive model with random selection of features at each node creation, but in general the effect is not removed completely. It could be experimented during the model development.

It would be useful if the overall number of unique values in each variable could be presented with graphical representations. A pie chart and a bar chart would do the trick to quickly understand the overall characteristics of distributions of unique values in the dataset.

```
no_unique_values.append(no_unique_value)
         return no_unique_values
     111
         Plot a bar chart for the number of unique values in each variable.
         Numbers of unique values of each variable will also be displayed on top of \Box
      \rightarrow each bar,
         otherwise those small values would have no way to display themselves.
     def plot_no_unique_values_bar_chart(columns):
         no_unique_values = get_no_unique_values(columns)
         total_no_unique_value = sum(no_unique_values)
         percentages = []
         for value in no_unique_values:
             percentages.append(round((value/total_no_unique_value)*100, 4))
         fig, ax = plt.subplots()
         percent idx = 0
         for i, v in enumerate(no_unique_values):
             ax.text(v + 3, i + .25, str(v)+str(" 
      →(")+str(percentages[percent_idx])+str("%)"), color='green')
             percent_idx += 1
         width = 0.65 # the width of the bars
         ind = np.arange(len(no_unique_values))
         ax.barh(ind, no unique values, width, color="blue")
         ax.set yticks(ind+width/2)
         ax.set_yticklabels(columns, minor=False)
         plt.title('No. of unique values by variable')
         plt.xlabel('No. of unique values')
         plt.ylabel('Variables')
         plt.show()
[31]: '''
         attribute8 is removed from the column list since it is a duplicate of |
      \rightarrow attribute7.
     , , ,
     columns = ['date',
                              'device', 'attribute1',
                'attribute2', 'attribute3', 'attribute4',
                'attribute5', 'attribute6', 'attribute7',
                'attribute9'l
     111
         Generate a bar chart for more details with counts of unique values in each ⊔
      \rightarrow variable.
```

```
plt.rcParams['figure.figsize'] = (18.0, 8.5)
plot_no_unique_values_bar_chart(columns)
```

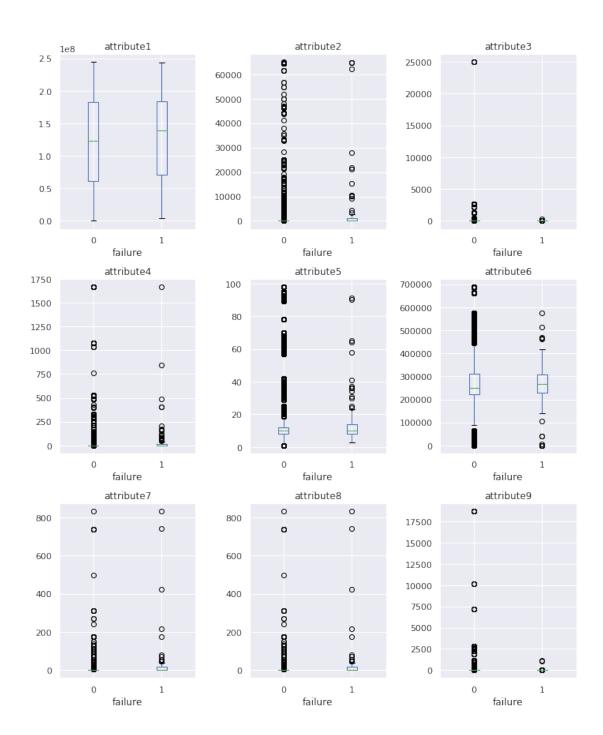


- 'attribute1' and 'attribute6' have large numbers of unique values while other variables have relatively small numbers of unique values.
- Unique values of 'attribute1' are assumed to be the amount of daily data exchange. 'attribute6' also a large number of unique values but it is not really conclude what 'attribute6' means.
- It seems reasonable to consider unique values of 'attribute1' and 'attribute6' as continuous values.
- 'attribute2' has 558 unique values and 'attribute4' has 115 unique values.
- Treating unique values of 'attribute2' as a continuous values seems to be rational.
- 'attribute3', 'attribute4', 'attribute5', 'attribute7', and 'attribute9' are assumed to be categorical nominal type values. discrete values of those variables could be treated as continuous type values but in this demonstration, they are forced to be categorical nominal type values to apply a variety of modeling approaches.

'attribute1', 'attribute2', and 'attribute6' are considered to be continuous type values while 'attribute3', 'attribute4', 'attribute5', 'attribute7', and 'attribute9' are assumed to be categorical nominal value types. Recall that 'attribute8' is not included because 'attribute8' is identical to 'attribute7'. One can use both 'attribute7' and 'attribute8' if Random Forest is the learning algorithm but for this demonstration, 'attribute8' will not be used. Before precedeing further, let's take a look at ranges of values in each variable for overall characteristics using boxplots. Although boxplots are for continuous values, they can be utilized to picture overall characteristics of variables even for categorical values, too.

```
[230]: fig = plt.figure(figsize=(10,12))
ax1 = plt.subplot(331)
ax2 = plt.subplot(332)
```

```
ax3 = plt.subplot(333)
ax4 = plt.subplot(334)
ax5 = plt.subplot(335)
ax6 = plt.subplot(336)
ax7 = plt.subplot(337)
ax8 = plt.subplot(338)
ax9 = plt.subplot(339)
dataset.boxplot(column='attribute1',by='failure',ax=ax1)
dataset.boxplot(column='attribute2',by='failure',ax=ax2)
dataset.boxplot(column='attribute3',by='failure',ax=ax3)
dataset.boxplot(column='attribute4',by='failure',ax=ax4)
dataset.boxplot(column='attribute5',by='failure',ax=ax5)
dataset.boxplot(column='attribute6',by='failure',ax=ax6)
dataset.boxplot(column='attribute7',by='failure',ax=ax7)
dataset.boxplot(column='attribute8',by='failure',ax=ax8)
dataset.boxplot(column='attribute9',by='failure',ax=ax9)
plt.suptitle('')
plt.tight_layout()
```



5.1 Categorical Variable Exploration

Unique values, contributing to failure, of each variable will be studied. The following uitility functions will help plot the distribution of contributions of unique values in each variable.

[247]:

```
total_record = dataset['failure'].count() # Get a count for total records.
      failure_summary_table = [["Variable", "Total", "Failure", "Non-failure", "
       →"Intersected"]] # a summary table.
      111
          This will plot statistics on unique values with failure contribution.
      def plot_unique_value_contribution_device_failure(attr):
          . . .
              Prepare dataframe from the original dataset for required statistics.
          plt.rcParams['figure.figsize'] = (18.0, 3.5)
          attr_failure_info = dataset['failure'] == 1
          df1 = pd.DataFrame(dataset, columns=[attr])
          df2 = pd.DataFrame(dataset[attr_failure_info], columns=[attr])
          attr_failure_info = dataset[attr_failure_info].groupby(attr).failure.sum()
          111
              Compute necessary information to generate summary statistics on unique\sqcup
       \rightarrow values with
              contributions to both failure and non-failure.
          total_no_uniq_values = dataset[attr].nunique()
          unique_value_ratio = round((total_no_uniq_values/total_record)*100, 4)
          failures = attr_failure_info.count()
          failure ratio = round((failures/total no uniq values)*100, 4)
          non_failures = total_no_uniq_values - failures
          non_failure_ratio = round((non_failures/total_no_uniq_values)*100, 4)
          intersection_df = pd.merge(df1, df2, on=[attr], how='inner')
          intersection = intersection_df[attr].nunique()
          intersection_ratio = round((intersection/failures)*100, 4)
              Store summaries on unique values with failure contribution.
          tr = ['{}'.format(attr),
                '{:,}({:.2f}%)'.format(total_no_uniq_values, unique_value_ratio),
                '{:,}({:.2f}%)'.format(failures, failure_ratio),
                '{:,}({:.2f}%)'.format(non_failures, non_failure_ratio),
                '{:,}({:.2f}%)'.format(intersection, intersection_ratio)
          failure_summary_table.append(tr)
[248]: # See how unique values in attribute3 contributes to failure.
      plot_unique_value_contribution_device_failure('attribute3')
      plot_unique_value_contribution_device_failure('attribute4')
```

```
plot_unique_value_contribution_device_failure('attribute5')
plot_unique_value_contribution_device_failure('attribute7')
plot_unique_value_contribution_device_failure('attribute9')
```

Let's take a look at the summary of category variable analyses.

```
[249]: from IPython.display import HTML, display import tabulate

[250]: display(HTML(tabulate.tabulate(failure_summary_table, tablefmt='html')))
```

<IPython.core.display.HTML object>

Those variables in the summary are all assumed to have categorical nominal type values. All the category variables have high cardinarities, meaning that those category variables have a large number of unique categorical values.

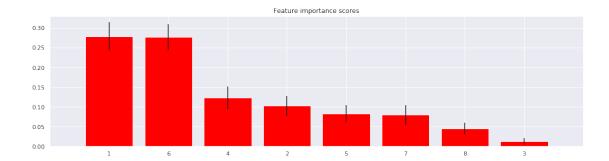
The predictive model needs to classify devices into either failure or non-failure. Looking at the summary above, 'attribute3' will be the most importance variable for the predictive model to learn that a device is either in failure or non-failure since it has the highest number of intersected unique values. Having a higher number of intersected unique values for a categorical variable means that the predictive model knows many ways to tell if a device is in failure or non-failure. The rank of variable importances will follow high numbers of intersected unique values in each variable.

In addition to the summary above, it might a good idea to look at numerical categorical variable importance measures by using *'ExtraTreesClassifer'* in *'sklearn'* and compare the numerical feature importances to the summary above.

```
[217]: # import ExtraTreesClassifier.
      from sklearn.ensemble import ExtraTreesClassifier
[223]: # Test features for importances.
      def examine_features(x_train, y_train):
          # Build a forest and compute the feature importances
          forest = ExtraTreesClassifier(n_estimators=250, random_state=0)
          forest.fit(x_train, y_train)
          importances = forest.feature_importances_
          std = np.std([tree.feature_importances_ for tree in forest.estimators_],
                       axis=0)
          indices = np.argsort(importances)[::-1]
          # Print the feature ranking
          print("Feature ranking:")
          for f in range(X.shape[1]):
              print("%d. attribute %d (%f)" % (f + 1, indices[f] + 1, u
       →importances[indices[f]]))
          # Plot the feature importances of the forest
          plt.rcParams['figure.figsize'] = (18.0, 4.5)
          plt.figure()
          plt.title("Feature importance scores")
```

Feature ranking:

```
1. attribute 1 (0.278196)
2. attribute 6 (0.276792)
3. attribute 4 (0.123269)
4. attribute 2 (0.102534)
5. attribute 5 (0.082676)
6. attribute 7 (0.079533)
7. attribute 8 (0.044559)
8. attribute 3 (0.012441)
```



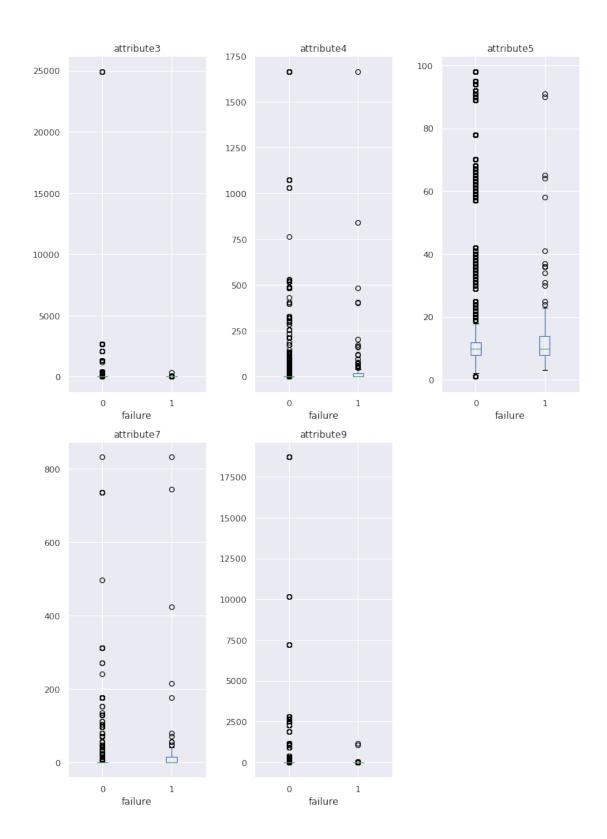
The numeric variable importance test with ExtraTreesClassifier shows the order of important variables according to the variable importance score of each variable. As mentioned above, it is conclusive that variables with a higher number of intersected unique values take higher variable importance scores. Knowing numerical importances of variables gives more confidence in choosing variables to include or exclude when developing the predictive model. With these understandings about variables in the current dataset, diving deeper into the spreads of unique values on failure and non-failure would give more insights to further understand about the variables. Although boxplots are for continuous values, boxplots can be improvised to look at the spreads of unique values of categorical variables, too. Note that one can use *'Logistic Regression'" to display the spread of values separately.

```
[253]: fig = plt.figure(figsize=(10,20))
ax1 = plt.subplot(331)
```

```
ax2 = plt.subplot(332)
ax3 = plt.subplot(333)
ax4 = plt.subplot(334)
ax5 = plt.subplot(335)

dataset.boxplot(column='attribute3',by='failure',ax=ax1)
dataset.boxplot(column='attribute4',by='failure',ax=ax2)
dataset.boxplot(column='attribute5',by='failure',ax=ax3)
dataset.boxplot(column='attribute7',by='failure',ax=ax4)
dataset.boxplot(column='attribute9',by='failure',ax=ax5)

plt.suptitle('')
plt.tight_layout()
```



By looking at the boxplots above, 'attribute4', 'attribute5', and 'attribute7' seem to have more options to contribute to failure than other variables: 'attribute3' and 'attribute9'. Above boxplots

reveal that the unique value, '0', or close to '0' in 'attribute3', and 'attribute9' show that has most frequencies for both failure and non-failure while 'attribute4', 'attribute5', 'attribute7' show similar spread patterns of their unique values except ranges of spreads of their unique values. From the study of variables, 'attribute4', 'attribute5', and 'attribute7' might be good to use for the predictive model developments since they have more options to failure and have different spreads of unique values other than unique values for non-failure.

[]: