

analysis

June 10, 2017

1 Evaluation analysis

We will be taking a look at the evaluations from the data folder ../data/ ([notebook](#), [github](#)).

1.1 Setup

Before looking at the data, a list of imports and the version of libraries used is reported.

```
In [1]: # Built-in python libraries
import platform
from glob import glob
from itertools import chain

# 3rd-party libraries
import matplotlib
import matplotlib.pyplot as plt
import numpy as np
import pandas as pd
import IPython
from IPython.utils.coloransi import TermColors

# Print versions.
print('Python version: {}'.format(platform.python_version()))
print('IPython version: {}'.format(IPython.__version__))
print('matplotlib version: {}'.format(matplotlib.__version__))
print('numpy version: {}'.format(np.__version__))
print('pandas version: {}'.format(pd.__version__))

# Initialize the backend for Jupyter
%matplotlib notebook

# Set style-sheet to grayscale.
matplotlib.style.use('ggplot')
colormap = plt.cm.get_cmap('RdYlBu_r')
C = [colormap(x/5) for x in range(5)]
# Set figure font to serif.
```

```
plt.rcParams['font.family'] = 'serif'

# Set how many columns to show in tables.
pd.options.display.max_columns = 50
pd.options.display.max_rows = 400
# Set the format to print float values to 3 decimal points.
pd.options.display.float_format = lambda x: '%.3f' % x
```

```
Python version: 3.6.1
IPython version: 5.3.0
matplotlib version: 2.0.2
numpy version: 1.12.1
pandas version: 0.20.1
```

2 The data

First we load the CSV file into a [pandas DataFrame](#), print the amount of samples and take a look at the column headers of the dataset.

```
In [2]: file = '../data/evaluations.csv'

conversion_dict = {'research_type': lambda x: int(x == 'E')}

evaluation_data = pd.read_csv(file, sep=',', header=0, index_col=0, converters=conversion_dict)

print('Amount of samples: {}'.format(len(evaluation_data.index)))

column_headers = evaluation_data.columns.values
print('\nColumn headers: {}'.format(column_headers))
```

```
Amount of samples: 400
```

```
Column headers: ['title' 'research_type' 'result_outcome' 'affiliation'
'problem_description' 'goal/objective' 'research_method'
'research_question' 'hypothesis' 'prediction' 'contribution' 'pseudocode'
'open_source_code' 'open_experiment_code' 'train' 'validation' 'test'
'results' 'hardware_specification' 'software_dependencies'
'third_party_citation' 'experiment_setup' 'evaluation_criteria' 'authors'
'link' 'comments' 'conference']
```

There are 400 samples with 27 columns in total for each sample. However, some columns are not necessary for further analysis: *title*, *authors*, *link*, *comments*. The *comments* column contains short messages such as “Points to an extended paper” or “Links to appendix which links to code” to give extra information in case an evaluation is unclear. The other three identify which paper was evaluated. These columns are therefore removed from the dataframe.

```
In [3]: evaluation_data.drop(['title', 'link', 'authors', 'comments'], axis=1, inplace=True)
        column_headers = evaluation_data.columns.values
        print('\nColumn headers: {}'.format(column_headers))
```

```
Column headers: ['research_type' 'result_outcome' 'affiliation' 'problem_description'
'goal/objective' 'research_method' 'research_question' 'hypothesis'
'prediction' 'contribution' 'pseudocode' 'open_source_code'
'open_experiment_code' 'train' 'validation' 'test' 'results'
'hardware_specification' 'software_dependencies' 'third_party_citation'
'experiment_setup' 'evaluation_criteria' 'conference']
```

The remaining 23 columns can be placed in more clarifying categories. All data is boolean with the value 0 or 1, unless otherwise specified below.

Miscellaneous Variables describing the research

research_type - Experimental (1) or theoretical (0).

result_outcome - Novel research or not.

affiliation - The affiliation of the authors; academia (0), collaboration (1), industry (2).

conference - The conference the paper was accepted to.

third_party_citation - Is third-party source code or data referenced?

Research Transparency How well documented is the research method?

problem_description - The problem the research seeks to solve.

goal/objective - The objective of the research.

research_method - Research method used.

research_question - Research question(s) asked.

hypothesis - Investigated hypothesis.

prediction - Predictions related to the hypothesis.

contribution - Contribution of the research.

Note: The variables under Research Transparency are 1 if explicitly mentioned in the paper, otherwise 0.

Experiment Documentation How well is the experiment documented?

open_experiment_code - Is the experiment code available?

hardware_specification - Hardware used.

software_dependencies - For method or experiment.

experiment_setup - Is the experiment setup described with parameters etc.?

evaluation_criteria - Specification of evaluation criteria.

Method Documentation How well is the method under investigation documented?

pseudocode - Method described in pseudocode.

open_source_code - Is the method code available?

Open Data How well is the data documented, and is it available?

train - Training set specification.

validation - Validation set specification.

test - Test set specification.

results - Raw results data.

Note: If no data is open sourced all will be 0. If data is open source but the sets are not specified train or test will be set to 1 depending on whether the research requires training or not. If the research does not require training, train and validation does not have a value set.

```
In [4]: category_headers = {
    'Miscellaneous': np.append(column_headers[0:3], column_headers[[19, 22]]),
    'Research Transparency': column_headers[3:10],
    'Method Documentation': column_headers[10:12],
    'Open Data': column_headers[13:17],
    'Experiment Documentation': column_headers[[12, 17, 18, 20, 21]]
}
```

A look at the first two samples of the dataset show the difference between experimental and theoretical papers.

```
In [5]: evaluation_data.head(2)
```

```
Out[5]:
```

| | research_type | result_outcome | affiliation | problem_description | \ |
|-------|---------------|----------------|-------------|---------------------|---|
| index | | | | | |
| 1 | 1 | 1 | 0 | 1 | |
| 2 | 0 | 1 | 0 | 0 | |

| | goal/objective | research_method | research_question | hypothesis | \ |
|-------|----------------|-----------------|-------------------|------------|---|
| index | | | | | |
| 1 | 0 | 0 | 0 | 0 | |
| 2 | 0 | 0 | 0 | 0 | |

| | prediction | contribution | pseudocode | open_source_code | \ |
|-------|------------|--------------|------------|------------------|---|
| index | | | | | |
| 1 | 0 | 1 | 1.000 | 0.000 | |
| 2 | 0 | 0 | nan | nan | |

| | open_experiment_code | train | validation | test | results | \ |
|-------|----------------------|-------|------------|-------|---------|---|
| index | | | | | | |
| 1 | 0.000 | 1.000 | 1.000 | 0.000 | 0.000 | |
| 2 | nan | nan | nan | nan | nan | |

| | hardware_specification | software_dependencies | third_party_citation | \ |
|-------|------------------------|-----------------------|----------------------|---|
| index | | | | |
| 1 | 0.000 | 0.000 | 0.000 | |
| 2 | nan | nan | nan | |

| | experiment_setup | evaluation_criteria | conference |
|-------|------------------|---------------------|------------|
| index | | | |
| 1 | 1.000 | 1.000 | IJCAI 16 |
| 2 | nan | nan | IJCAI 16 |

The first sample is an experimental paper (**research_type=1**) and has values set for all the columns. The second paper, however, is a theoretical paper (**research_type=0**) and only has values set for the *Miscellaneous*, and *Research Transparency* categories, excluding the *third_part_citation* column. Note that the datafile has Experimental noted as E and theoretical noted as T.

Cells with missing values are represented as NaN in pandas and can be seen for all the columns exclusive to experimental papers in the second sample above. For experimental papers where

training is not relevant, both the *train* and *validation* columns will show as NaN. To add NaN to visualisations below, we fill them out with the value -1.

Additionally, we split the experimental papers into a separate dataframe for plotting later.

```
In [6]: evaluation_data = evaluation_data.fillna(-1)
        experimental_data = evaluation_data[evaluation_data.research_type == 1]
```

2.1 Miscellaneous

We start with the miscellaneous category, defining the plot function which will be used for all categories. The only variable not plotted is the conference variable, which has its frequencies printed out instead.

Variables describing the research

research_type - Experimental (1) or theoretical (0).

result_outcome - Novel research or not.

affiliation - The affiliation of the authors; academia (0), collaboration (1), industry (2).

conference - The conference the paper was accepted to.

third_party_citation - Is third-party source code or data referenced?

```
In [7]: def plot_full_series(series, title, labels, width=0.4):
        bins=len(labels)
        Y, X = np.histogram(series, bins=bins)
        total_Y = sum(Y)
        fig = plt.figure(figsize=(4,4))
        ax = plt.subplot(111)
        plt.bar(X[:-1], Y, color=C, width=width, axes=ax)
        ax.set_ylim(0, total_Y + 20)
        ax.set_xticks(X[:-1])
        ax.set_xticklabels(labels)
        #ax.set_title(title)

        # Add amount labels to bars
        for y, x in zip(Y, X[:-1]):
            label = '{:3.0f} ({:.1%})'.format(y, y / total_Y)
            ax.text(x, y + 5, label, ha='center', va='bottom')
        plt.show()
        fig.savefig('../doc/report/fig/{}'.format(title.replace(' ', '_')))
```

```
In [ ]: print(evaluation_data.groupby('conference').size(), end='\n\n')
```

```
plot_full_series(evaluation_data.affiliation, 'Affiliation', ['Academia', 'Collaboration', 'Industry'])
plot_full_series(evaluation_data.research_type, 'Research Type', ['Theoretical', 'Experimental'])
plot_full_series(evaluation_data.result_outcome, 'Result Outcome', ['Negative', 'Positive'])
plot_full_series(experimental_data.third_party_citation, 'Third-party Citation', ['Not p
```

2.2 Research Transparency

How well documented is the research method?

problem_description - The problem the research seeks to solve.

goal/objective - The objective of the research.

research_method - Research method used.

research_question - Research question(s) asked.

hypothesis - Investigated hypothesis.

prediction - Predictions related to the hypothesis.

contribution - Contribution of the research.

Note: The variables under Research Transparency are 1 if explicitly mentioned in the paper, otherwise 0.

```
In [ ]: plot_full_series(evaluation_data.contribution, 'Contribution', ['Not present', 'Present'])
        plot_full_series(evaluation_data['goal/objective'], 'Goal or Objective', ['Not present', 'Present'])
        plot_full_series(evaluation_data.hypothesis, 'Hypothesis', ['Not present', 'Present'])
        plot_full_series(evaluation_data.prediction, 'Prediction', ['Not present', 'Present'])
        plot_full_series(evaluation_data.problem_description, 'Problem Description', ['Not present', 'Present'])
        plot_full_series(evaluation_data.research_method, 'Research Method', ['Not present', 'Present'])
        plot_full_series(evaluation_data.research_question, 'Research Question', ['Not present', 'Present'])
```

2.3 Experiment Documentation

How well is the experiment documented?

evaluation_criteria - Specification of evaluation criteria.

experiment_setup - Is the experiment setup described with parameters etc.?

hardware_specification - Hardware used.

open_experiment_code - Is the experiment code available?

software_dependencies - For method or experiment.

```
In [ ]: plot_full_series(experimental_data.evaluation_criteria, 'Evaluation Criteria', ['False', 'True'])
        plot_full_series(experimental_data.experiment_setup, 'Experiment Setup', ['False', 'True'])
        plot_full_series(experimental_data.hardware_specification, 'Hardware Specification', ['False', 'True'])
        plot_full_series(experimental_data.open_experiment_code, 'Open Experiment Code', ['False', 'True'])
        plot_full_series(experimental_data.software_dependencies, 'Software Dependencies', ['False', 'True'])
```

2.4 Method Documentation

How well is the method under investigation documented?

pseudocode - Method described in pseudocode.

open_source_code - Is the method code available?

```
In [ ]: plot_full_series(experimental_data.pseudocode, 'Pseudocode', ['False', 'True'])
        plot_full_series(experimental_data.open_source_code, 'Open Source Code', ['False', 'True'])
```

2.5 Open Data

How well is the data documented, and is it available?

train - Training set specification.

validation - Validation set specification.

test - Test set specification.

results - Raw results data.

```
In [ ]: plot_full_series(experimental_data.train, 'Training Data', ['N/A', 'False', 'True'])
        plot_full_series(experimental_data.validation, 'Validation Data', ['N/A', 'False', 'True'])
        plot_full_series(experimental_data.test, 'Test Data', ['False', 'True'])
        plot_full_series(experimental_data.results, 'Results Data', ['False', 'True'])
```