analysis

June 11, 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.

```python
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'
```

```python
        # 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.

```python
In [ ]: 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=convers:

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

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

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.

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

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.

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

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

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 [4]: 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?

```python
In [5]: 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) Removed in favor of captions in report.

            # 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', 'Collaboratio
        plot_full_series(evaluation_data.research_type, 'Research Type', ['Theoretical', 'Exper
        plot_full_series(evaluation_data.result_outcome, 'Result Outcome', ['Negative', 'Positi
        plot_full_series(experimental_data.third_party_citation, 'Third-party Citation', ['Not
```

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.

```python
In [ ]: plot_full_series(evaluation_data.contribution, 'Contribution', ['Not present', 'Present
        plot_full_series(evaluation_data['goal/objective'], 'Goal or Objective', ['Not 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 pres
    plot_full_series(evaluation_data.research_method, 'Research Method', ['Not present', 'P
    plot_full_series(evaluation_data.research_question, 'Research Question', ['Not 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
    plot_full_series(experimental_data.experiment_setup, 'Experiment Setup', ['False', 'Tr
    plot_full_series(experimental_data.hardware_specification, 'Hardware Specification', [
    plot_full_series(experimental_data.open_experiment_code, 'Open Experiment Code', ['Fals
    plot_full_series(experimental_data.software_dependencies, 'Software Dependencies', ['Fa
```

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', 'Tr
```

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', 'Tr
    plot_full_series(experimental_data.test, 'Test Data', ['False', 'True'])
    plot_full_series(experimental_data.results, 'Results Data', ['False', 'True'])
    all_sets = experimental_data[['train', 'validation', 'test']].all(axis=1)
    plot_full_series(all_sets, 'Data_Sets', ['False', 'True'])
```

2.6   Analasys patterns

The analysis patterns will be examined for variables related to open data and source code.

2.6.1 Author affiliation

```
In [ ]: labels_of_interest = ['open_source_code', 'open_experiment_code',
                              'train', 'validation', 'test', 'results']

        for label in labels_of_interest:
            print(experimental_data.groupby('affiliation')[label].value_counts())
```

2.6.2 Conference differences

```
In [ ]: for label in labels_of_interest:
            print(experimental_data.groupby('conference')[label].value_counts())
```

2.6.3 Novelty of research

This analysis pattern has been discarded due to problems with the evaluation of Result outcome.

2.7 Reproducibility

```
In [ ]: methods_reproducible = experimental_data[labels_of_interest[0:-1]].all(axis=1)
        results_reproducible = experimental_data[labels_of_interest[0:2]].all(axis=1)

        plot_full_series(methods_reproducible, 'Methods Reproducible', ['False', 'True'])
        plot_full_series(results_reproducible, 'Results Reproducible', ['False', 'True'])
```