

# CA – Assignment 2: Argument Mining

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- Group: **FakeNews**
- Group members:
  - Adnan Manzoor
  - Sajjad Pervaiz
  - Kevin Taylor
  - Christoph Schäfer

## Structure

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├── argument-mining-assignment
│   ├── Documentation.pdf
│   ├── README
│   ├── requirements.txt
│   └── code
│       ├── data
│       │   ├── essay_corpus.json
│       │   ├── pred.txt
│       │   ├── test_BIO.txt
│       │   ├── train-test-split.csv
│       │   └── train_BIO.txt
│       ├── convert_to_bio.py
│       ├── convert_to_train_test_bio.py
│       ├── evaluation.py
│       └── model.py
```

## Scripts

**essay\_corpus.json**: Data corpus created in Data Acquisition assignment.

**convert\_to\_bio.py**: provided along with the assignment to convert json corpus to **BIO** format.

**convert\_to\_train\_test\_bio.py**: Our implementation of converting **essay\_corpus.json** to bio format.

- We call **convert\_to\_bio** function from **convert\_to\_bio.py** script in our implementation to create **train\_BIO.txt** and **test\_BIO.txt** based on **train-test-split.csv** scheme. The files are placed in **data/** folder and will be later used to train and test the **ML** model.

**model.py**: The ML model that we use for generating predictions. **evaluation.py**: Script to evaluate the **F1** score of the **ML** model.

## How to run the scripts

- On a venv install the requirements specified in **requirements.txt**

- Make sure you have the same directory structure as above otherwise adjust the paths in the scripts accordingly.
- Run `convert_to_train_test_bio.py`, which will create `train_BIO.txt` and `test_BIO.txt` in `data/`.
- Run `model.py` to generate the predictions in `data/` directory with name `pred.txt`
- Run `evaluation` script with `preds.txt` as predictions and `test_BIO.txt` as ground truth.

## Model Explanation

We choose **Naive Bayes (NB)** for its simplicity and used **Bag of words** as our feature representation to train the model. NB achieves **Macro F1-Score: 0.235** and **Weighted F1-Score: 0.501**.

### Feature Selection

We used **n-grams** to capture the maximum context around each **token** in the training dataset. To implement the model we used **Pipeline** from **Sklearn** and passed it **CountVectorizer()** that calculates word embeddings (bow) for each token. Our **n-gram** logic is the following.

```
For each token:
    if there exists 2 preceeding and succeeding tokens in a sentence`:
        n-gram_token = [preceedingToken1, preceedingToken2, targeToken,
succeedingToken1, succeedingToken2]
    else:
        add `empty` string for corresponding slot.
```

The motivation is to capture as much context as possible around a token.