

Project Structure:

- data folder
 - All analysis tool output.
 - All csv sheets created with the analysis tool output.
 - The file “correlation_analysis.xlsx”, which stores an overview of data collected including correlation results.
- tool-execution folder
 - All shell scripts for executing different steps of the pipeline.
 - Note: ALL .sh scripts are designed to be run from the project root directory
- simple-datasets
 - Contains the snippets for COG datasets 1, 2, and 3. Also contains the snippets for the fMRI study.
- dataset6
 - Contains the snippets for COG dataset 6
- dataset9
 - contains the snippets for COG dataset 9

Setting Up Project Environment:

- Make sure that you have Python and Java installed on your machine
 - I use Java versions 17.0.3 and 8 and Python version 3.10.5 although other similar versions will probably work fine.
 - You will need to swap between the Java versions when using some of the different analysis tools. On Linux, this can be easily done with the following **command**: `sudo update-alternatives --config java`
- Clone the repository to the current directory.
 - git clone <https://github.com/sea-lab-wm/complexity-verification-project.git>
- Go to project root
 - `cd complexity_verification_project`
- Run the following 3 commands from project root to setup the python environment:
 - Create the virtual environment: `python3 -m venv complexity_verification_project_venv`
 - Note: The command name might differ depending on your version of Python: Could be `python`, `python3`, `python3.10`, etc
 - Activate the virtual environment with bash/zsh: `source complexity_verification_project_venv/bin/activate`
 - Note: activation differs based on platform: <https://docs.python.org/3/library/venv.html>
 - Install the required libraries to the virtual environment: `pip3 install -r requirements.txt`
- Note: you might have to give yourself permission to run the .sh scripts using `chmod`.

Checklist of all things to do when adding a new analysis tool to the project:

- Create new .bat and .sh scripts to run the analysis tool + add any other dependencies, files, etc that are needed to run the analysis tool.

- Create a new function in parser.py to parse the analysis tool's output and convert it to a .csv.
- Create a new sheet in "correlation_analysis.xlsx" named after the analysis tool and fill in its column titles as well as the first 2 columns worth of data (same as the other sheets).
- In correlation.py:
 - Update the 'if __name__ == "__main__":' section at the bottom so that correlations can be done on the new analysis tool isolated from the others.
 - Update the readAnalysisToolOutput() function to read in the .csv file created for this analysis tool.

Checklist of all things to do when adding a new dataset/metric to the project:

- Add the dataset to the project
- Add any relevant metric results from the previous study that used the dataset to the project
- Update the .bat and .sh scripts that are used to run the analysis tools on all the datasets to include the new dataset:
 - Some tools such as the Checker Framework automatically check all .java files within the project
 - Other tools such as the Typestate Checker need to explicitly be given single directories of .java files at a time, thus needing to be updated when adding a new dataset
- Update every sheet in "correlation_analysis.xlsx". Add a new row containing the name of the dataset and metric in the same format as the other rows. Also add to the #of snippets judged column.
- In correlation.py, update the setupCorrelationData() function to read in the relevant metrics that we will correlate our warning counts against.

Collecting the Data:

1. Run the Checker Framework (Checker Framework version 3.21.3, Java version 17.0.3) on all the snippets. Collect all of the warnings generated.
 - a. How To: Run this command from the project root directory:
 - i. ./tool-execution/run_checker_framework.sh
 - b. Output: The Checker Framework outputs all of its warnings to "checker_framework_output.txt"
2. Run the Infer tool (Java version 17.0.3) on all the snippets. Collect all the warnings generated. Note that some warning types such as null dereferences cause the tool to stop analyzing the current method when spotted and move on to the next one.
 - a. How To: Run this command from the project root directory:
 - i. ./tool-execution/run_infer.sh
 - b. Output: Infer outputs all of its warnings to "infer_output.txt"
3. Manually swap to Java version 8.
4. Run the TypeState Checker (Checker Framework version 3.14.0, Java version 8 as recommended on the typestate Github) on a single folder of snippets. Collect all of the warnings generated. This differs from the Checker Framework in that the Checker

Framework checks all snippets simultaneously while the Typestate Checker only operates on a single folder at a time.

- a. How To: Run the script “run_tystate_checker.sh” from the project root with this command. This will run a series of commands to run the Typestate Checker separately for each dataset.
 - i. `./tool-execution/run_tystate_checker.sh`
 - b. Output: The Typestate Checker outputs all of its warnings for each dataset to separate files called: “tystate_checker_output_{dataset name}.txt”
5. At this stage, all warnings have now been collected across all analysis tools and all datasets.

Analyze the Data:

- How To:
 - Run the file “run_correlation_analysis.sh” to complete ALL steps 1-7 automatically.
 - Run this command from the project root directory:
`./tool-execution/run_correlation_analysis.sh`
 - The two scripts, “parser.py” and “correlation.py” can alternatively be run manually and independently of each other if needed.
 - “parser.py” is for Step 1
 - “correlation.py” is for Steps 2-7.
1. Parse the warning output from each analysis tool separately.
 - a. Read the text file output from each analysis tool one by one.
 - i. Note: Some analysis tools such as the tystate checker only run on a single directory at a time so they output multiple text files.
 - b. For each analysis tool, read the text file(s) line by line, looking for keywords that definitively indicate the start of a new warning (as per the structure of each analysis tool’s output which does differ).
 - c. For each warning, get the name of the snippet (format: {dataset} - {snippet #}) as well as the “warning type”, the warning’s definition in the output.
 - d. Store the 2 pieces of data collected for each warning in 2 lists, then create a pivot table that shows the number of each type of warning per snippet
 - e. Output: Pivot table is stored in a csv file named after the analysis tool the warnings came from. Example: “checker_framework_data.csv”.
 - i. Note: If dealing with multiple text files from the same analysis tool, warnings are combined together resulting in ONE csv file output PER analysis tool.
 2. Gather all of the csv files created in Step 1.
 3. Determine the number of snippets that contain warnings within each dataset.
 - a. Using column 1 of “correlation_analysis.xlsx”, get a list of all datasets being used.
 - b. Form a list of all unique snippets that contain warnings by name (across all datasets).
 - c. Using the lists of datasets and unique snippets with warnings, sort those snippets by dataset.

- d. Output: Counts are stored in "correlation_analysis.xlsx".
- 4. Determine the number of warnings per snippet per dataset.
 - a. Get a list of all datasets being used.
 - b. Get a list of the total number of snippets in each dataset.
 - c. Go through each analysis tool's output.
 - i. For each row of data, sum the values in that row, and note the snippet name at the start of the row. This gets the number of warnings for each snippet produced by a specific analysis tool
 - ii. Keep a running count for each snippet to get its total number of warnings across all analysis tools.
- 5. Using the data collected in Step 4, Determine the total number of warnings per dataset
 - a. Since each snippet's individual count of warnings is already sorted by dataset, simply sum those warning counts for each dataset.
 - b. Output: These counts are stored in "correlation_analysis.xlsx".
- 6. Setup the data for conducting a correlation analysis.
 - a. For each dataset/metric, form 2 lists of equal size. The first contains the complexity metric for each snippet used on the dataset in its previous study. The second contains the total number of warnings (across all analysis tools) produced on each snippet. This forms the (x,y) datapoints used for conducting the correlation analysis.
 - b. Update raw_correlation_data.csv
 - c. Note: Some studies used multiple different metrics on a single dataset. These metrics are correlated with our warnings separately.
 - d. Output: The number of datapoints for each dataset are stored in "correlation_analysis.xlsx".
- 7. Conduct Correlation Analysis
 - a. Kendall's Tau:
 - i. For each dataset, take the data points collected in Step 6 where x = complexity metric, y = # of warnings.
 - ii. Perform Kendall's Correlation, getting the correlation coefficient and p-value for each dataset.
 - iii. Output: The correlation coefficients and p-values are stored in "correlation_analysis.xlsx".
 - b. Spearman's Rho:
 - i. Exactly the same process as Kendall's correlation.