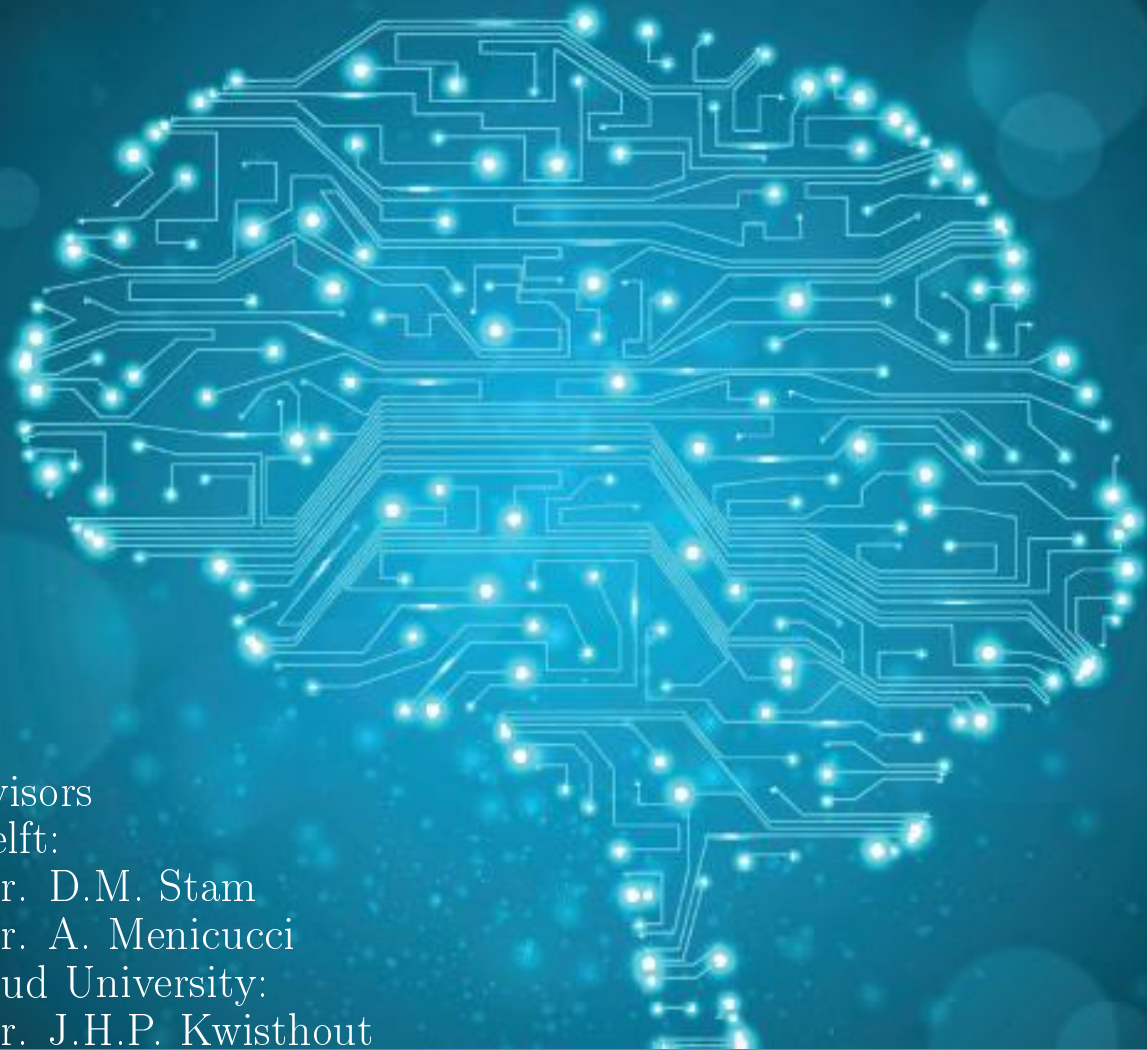


Leveraging brain adaptation to increase radiation resistance of neuromorphic space hardware

AE5810 Thesis Project: baseline report

Akke Toeter



Supervisors

TU Delft:

Dr. D.M. Stam

Dr. A. Menicucci

Radboud University:

Dr. J.H.P. Kwisthout

Leveraging brain adaptation to increase radiation resistance of neuromorphic space hardware

AE5810 Thesis Project: baseline report

by

Akke Toeter

Systems Engineering component of the AE5810 Thesis
at the Delft University of Technology.

Student number: 1507958
Project Planning duration: September 20, 2020 – September 24, 2021
Version: 0.1

Contents

1	Introduction	3
2	Functional Flow Diagram	5
2.1	Detailed Functional Flow Diagram	5
2.1.1	Description	6
3	Functional Breakdown Diagram	9
4	Requirements Discovery Tree	11
4.1	Mission Need Statement	11
4.2	Stakeholder Requirements	11
4.3	Top Level Requirements.	12
4.4	Identification Key Requirements	12
4.5	Requirements Discovery Tree	12
5	Resource Allocation and Budget Breakdown	13
6	Technical Risk assessment	15
7	Design Option Structuring Tree	17
7.1	Pruning	17
7.1.1	Tested Functionality	18
7.1.2	How To Perform Radiation Tests	18
7.1.3	Scope Of Radiation Tests	18
7.1.4	Brain Adaptation Implementation	18
7.1.5	Neuroplasticity Mechanism	18
7.1.6	Neuromorphic Architecture	18
7.1.7	Result Quantification	18
7.1.8	Pruned Design Option Tree	19
7.2	Preliminary Design Option Selection	19
7.2.1	Tested Functionality Preference	19
7.2.2	How To Perform Radiation Tests Preference	19
7.2.3	Scope of Radiation Tests Preference	19
7.2.4	Brain Adaptation Implementation Preference	20
7.2.5	Neuroplasticity Mechanism Preference	20
7.2.6	Neuromorphic Architecture Type Preference.	20
7.2.7	Result Quantification Preference.	20
7.2.8	Preliminary Design Option Tree	20
7.3	Proposed Design options	20
8	Contingency Management	23
9	Market Analysis	25
10	Sustainable Development Management	27
11	Reporting and Quality Control	29
12	Conclusion	31
	Bibliography	33

Acronyms	35
Glossary	37
Nomenclature	39
.1 Appendix	40
.2 Appendix __main__.py	41
.3 Appendix Hardcoded_data.py.	42
.4 Appendix Plot_to_tex.py	43
.5 Appendix arg_parser.py	48
.6 Appendix create_dynamic_diagrams.py.	50
.7 Appendix create_planar_triangle_free_graph.py	51
.8 Appendix create_static_diagrams.py	53
.9 Appendix distributed.py	54
.10 Appendix export_data.py	55
.11 Appendix helper_bash_commands.py	56
.12 Appendix helper_dir_file_edit.py	57
.13 Appendix latex_compile.py	60
.14 Appendix latex_export_code.py.	61
.15 Appendix neumann.py	84
.16 Appendix neumann_a_t_0.py	86
.17 Appendix plantuml_compile.py.	89
.18 Appendix plantuml_generate.py	93
.19 Appendix plantuml_get_package.py	96
.20 Appendix plantuml_to_tex.py.	98

Chapter 1

Introduction

This document presents the baseline for the AE5810 Thesis Project of the Space Flight Master at the Faculty of Aerospace Engineering of Delft University of Technology and the SOW-MKI92 Research Project of the Master in Artificial Intelligence at the faculty of Social Sciences of Radboud University. Its purpose is to identify the 2-5 most feasible design options that can be used to determine whether the principle of brain adaptation can be leveraged in neuromorphic space hardware.

The baseline report presents the Functional Flow Diagram (FFD) and Functional Break-down Diagram (FBD) in chapter 2 and chapter 3 respectively. These function descriptions of the system that is to be designed, is then used to generate the Requirements Discovery Tree (RDT) in chapter 4. Next, the resource allocation and budget breakdown presented in chapter 5. This is followed by the technical risk assessment in chapter 6. From the RDT, the Design Options Structuring Tree (DOT) is generated in chapter 4. Contingency management is applied in chapter 8. A market analysis is presented in chapter 9, and the sustainable development strategy is presented in chapter 10. To ensure this work is performed with sufficient quality, the reporting and quality control is presented in chapter 11. The baseline is concluded in chapter 12.

Chapter 2

Functional Flow Diagram

To gain insight in the system that is to be designed, a Functional Flow Diagram is generated. This FFD presents the high level functions that the system should be able to perform, in chronological order. These functions are presented in the flow diagram of fig. 2.1.

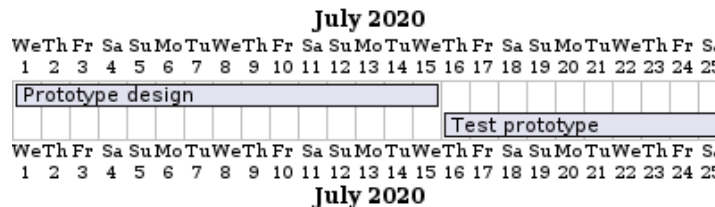


Figure 2.1: A functional flow diagram with the high-level functions of the system that is to be designed.

1. Initialise & start space-related function on neuromorphic architecture without brain adaptation implementation.
2. Initialise & start space-related function on neuromorphic architecture with brain adaptation implementation.
3. Endure modelled space radiation on neuromorphic architecture.
4. Measure space-related function performance without brain adaptation implementation.
5. Measure space-related function performance with brain adaptation implementation.
6. Report any performance difference between with- and without brain adaptation.
7. Determine significance of difference.

From these high level functions, a more detailed functional breakdown diagram is generated.

2.1. Detailed Functional Flow Diagram

1. Initialise & start space-related function on neuromorphic architecture without brain adaptation implementation.
 - (a) Boot/initialise neuromorphic architecture
 - (b) Load function.
 - (c) Load function data.
2. Initialise & start space-related function on neuromorphic architecture with brain adaptation implementation.
 - (a) Boot/initialise neuromorphic architecture

- (b) Load brain adaptation.
 - (c) Load space-related function.
 - (d) Load space-related function data.
3. Render and endure modelled space radiation on neuromorphic architecture.
 - (a) Determine simulated space radiation pattern of neuromorphic space architecture.
 - (b) Expose neuromorphic architecture to simulated space radiation pattern.
 - (c) Complete space-related function.
 - (d) Return results of space-related performance.
4. Measure space-related function performance without brain adaptation implementation.
 - (a) Retrieve space-related function output.
 - (b) Convert space-related function output to score.
5. Measure space-related function performance with brain adaptation implementation.
6.
 - (a) Retrieve space-related function output.
 - (b) Convert space-related function output to score.
7. Report difference between the scores of the architectures with- and without brain adaptation.
8. Determine significance of difference.

2.1.1. Description

1. Initialise & start space-related function on neuromorphic architecture without brain adaptation implementation.
 - To run a function on the neuromorphic architecture, it will have to be booted and initialised. The function that will be ran on the neuromorphic architecture is space related, to increase the level of representativeness of this study, in terms of space applications. The initialisation allows for loading the space related function that is to be executed. Additionally, the space related function may require (training) data on which it is ran. For example, a Martian rover that has a function that identifies rocks in its environment, may be partially simulated by loading a (labelled) dataset of Martian images.
To run the function without brain adaptation implementation, allows for the creation of a baseline to which the brain adaptation performance can be compared.
2. Initialise & start space-related function on neuromorphic architecture with brain adaptation implementation.
 - The intialisation of the brain adaptation implementation can occur, before, during or after the loading of the space related function. Which of these options is selected depends on the more detailed design process.
3. Endure modelled space radiation on neuromorphic architecture.
 - The radiation robustness of the neuromorphic architecture can be tested by exposing the neuro-morphic architecture to the radiation that it would experience in a space application. To determine what this radiation is, a relevant space mission and space function are selected. The time, position and orientation of the spacecraft in such a mission is then used to derive the radiation pattern to which the radiation may be exposed. This radiation is pattern is then used to determine to which (simulated) radiation the neuromorphic architecture will be exposed.
4. Measure space-related function performance without brain adaptation implementation.

- The performance of the space related function without brain adaptation implementation on the neuromorphic architecture is measured before, during and/or after radiation exposure. Which of these measuring moments are used, is still to be determined by the detailed design process. This measurement then serves as a comparison baseline to put the impact of the brain adaptation implementation into context.
5. Measure space-related function performance with brain adaptation implementation.
 6.
 - Once a baseline for comparison is established, the space related function can be ran again on the neuromorphic hardware, whilst being exposed to radiation. In this second setting, the brain adaptation implementation is used in an attempt to increase the radiation robustness of the neuromorphic architecture. The performance of the space related function is then measured before, during and/or after radiation exposure. Which of these measuring moments are used, is still to be determined by the detailed design process.
 7. Report any performance difference between with- and without brain adaptation.
 8.
 - If any performance difference is observed between the space related function with- and without brain adaptation implementation, it will be computed and stored.
 9. Determine significance of difference.
 10.
 - An analysis is performed to determine the level of significance of any observed difference.

Chapter 3

Functional Breakdown Diagram

This section presents the functional breakdown diagram of the system that is designed in this thesis project. This FBD is generated using the detailed functional flow diagram of section 2.1. The FBD presents the activities in an hierarchical style.

1. Run space-related function on neuromorphic architecture.
 - (a) Initialise neuromorphic architecture.
 - (b) Optional: Load brain adaptation implementation.
 - (c) Load space related function.
 - (d) Load space related function data.
 - (e) Run space related function.
 - (f) Complete running space related function.
 - (g) Retrieve space related function outputs.
 - (h) Convert space related function outputs to performance score.
 - (i) Report difference between the scores of the architectures with- and without brain adaptation.
 - (j) Determine significance of difference.
2. Render and endure modelled space radiation on neuromorphic architecture.
 - (a) Generate simulated space radiation pattern of neuromorphic space architecture.
 - (b) Expose neuromorphic architecture to simulated space radiation pattern.
 - (c) Optional: model architecture-radiation interaction.
 - (d) Optional: measure architecture-radiation interaction.

Chapter 4

Requirements Discovery Tree

This section presents an overview of the requirements that are identified within this thesis project. Its purpose consists of listing the requirements that drive the design, identifying killer requirements and presenting an overview of the project requirements. section 4.1 contains the mission need statement of this project. Next, the stakeholder requirements are identified in section 4.2. The top level requirements are presented in section 4.3, and the key requirements are presented in section 4.4. From these combined requirements, the Requirements Discovery Tree is drafted in section 4.5.

4.1. Mission Need Statement

The mission need statement is generated in the project plan phase, and is included in this section again to provide the context of the requirement derivation process. The MSN is:

Increase the radiation robustness of neuromorphic space hardware by leveraging the principle of brain adaptation in neuromorphic hardware.

4.2. Stakeholder Requirements

The following stakeholder requirements are identified:

- **STKH-UNI-01** - The research that is performed shall be reproducible.
- **STKH-UNI-02** - The sustainability of the design concepts shall be taken into account in the design trade-off process.
- **STKH-SPACEBRAINS-01** - Achieve results towards the REACH research by Q2 2022.
- **STKH-SPACEBRAINS-01-a** - Achieve results towards the REACH research by either: 2022-03-01, 2022-04-07, 2022-07-01.
- **STKH-ICONS-01** - Submit paper documenting research results before April 15th, 2022.
- **STKH-ICONS-01-a** - Submit full paper of 6-8 pages, presenting original research, or submit short paper of 3-4 pages that has preliminary results.
- **STKH-ICONS-02** - Upon acceptance for a presentation, submit presentation before July 27th, 2022.
- **STKH-RADBOUD-01** - The research proceedings shall be documented and submitted in a format accepted by Dr. J.H.P. Kwisthout before the SOW-MKI92 Research Project is completed.
- **STKH-RADBOUD-02** - The research for the SOW-MKI92 Research Project shall be performed using at least 28 EC of work.
- **STKH-Delft-01** - The research proceedings shall be documented and submitted in a format accepted by Dr. D.M. Stam and Dr. A. Menicucci before the AE5810 Thesis Project is completed.
- **STKH-Delft-02** - The research for the AE5810 Thesis Project shall be performed using at least 42 EC of work.

4.3. Top Level Requirements

The following requirements for this thesis project are identified:

- **TECH-01** - Technology Readiness Level (TRL) of the used technology shall be at least TRL 4 [-].
- **TEST-01** - Radiation robustness shall be tested in terms of algorithmic performance.
- **TEST-02** - The radiation tests shall be technically and economically feasible.
- **TEST-03** - The radiation tests results shall be generated before September 2022.
- **SUS-01** - Sustainability management shall be integrated in each phase of this project.
- **SUS-02** - Sustainability shall be assessed in the design trade-off process.
- **SAF-01** - All participants involved in testing, integrating and operations shall not be exposed to serious danger.
- **ESA-01** - Throughout this project quarterly reports shall be provided to the SpaceBrains foundation and the European Space Agency (ESA).

4.4. Identification Key Requirements

The key requirements are requirements that can render the project infeasible, requirements that drive the design, and/or requirements that induce high risk to project success. Within this thesis project, the following key requirements are identified:

1. **STKH-SPACEBRAINS-01** - Achieve results towards the REACH research by Q2 2022.
2. **TECH-01** - Technology Readiness Level (TRL) of the used technology shall be at least TRL 4 [-].
3. **TEST-03** - The radiation tests results shall be generated before September 2022.

The **STKH-SPACEBRAINS-01** requirement significantly drives the design space, as physical radiation tests are not deemed feasible within the given timeframe and project constraints. **TEST-03** implies a strict time constraint that induces significant risk to project success as it limits the amount of time available for development and scheduling. **TECH-01** significantly drives the design as it limits the design space to concepts that rely only on technologies of TRL 4 and higher.

4.5. Requirements Discovery Tree

A requirements' discovery tree is currently omitted due to time constraints.

Chapter 5

Resource Allocation and Budget Breakdown

Chapter 6

Technical Risk assessment

Chapter 7

Design Option Structuring Tree

The purpose of the design option tree is to find a feasible design option that can satisfy the requirements. This selection process can be an iterative process in case no feasible design option is found in the initial design option tree. By positioning the design options in a tree format, their hierarchical structure becomes visible. The tree is structured such that the most impactful decisions are selected at the top of the tree, whereas more detailed decisions are made at lower levels of the tree. Several subsections are created to Figure 7.1 presents the design option tree for this thesis. The nodes in the tree represent design options and child leafs represent design options within a parent design option. For example, a choice may be made to use digital neuromorphic hardware, and within that design space, a particular chip may be selected. Multiple (parallel) design options may be selected. For example, a softwarematic and hardwarematic implementation may be chosen.

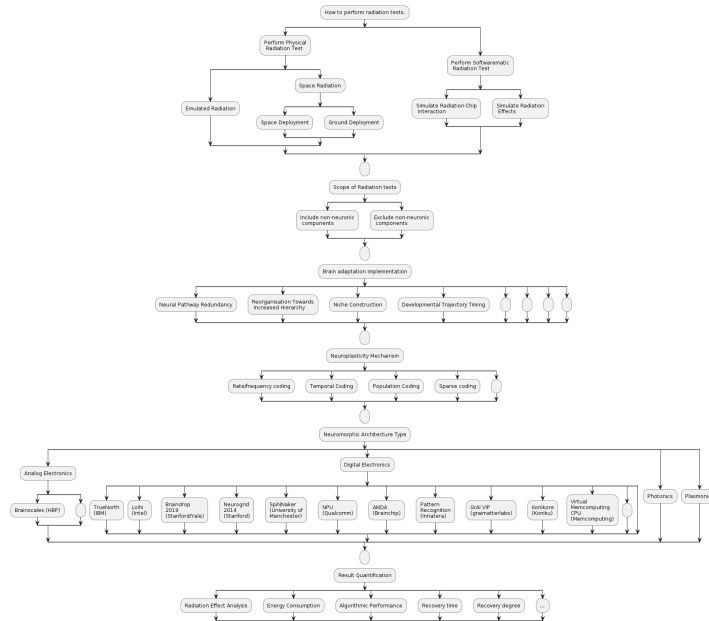


Figure 7.1: A functional flow diagram with the high-level functions of the system that is to be designed.

7.1. Pruning

With the design option tree generated, it can be pruned of options that are considered infeasible. This is done using the knowledge base that was generated in the literature study and using the requirements identified in chapter 4. The pruning process will be performed from top to bottom, which matches from high- to low hierarchical design choices.

7.1.1. Tested Functionality

At the time of writing, no specific function that is used for testing, can be eliminated.

7.1.2. How To Perform Radiation Tests

1. Starting with key requirement: **STKH-SPACEBRAINS-01** and **STKH-ICONS-01**, it is possible to eliminate the physical radiation test design option(along with its children), before the ICONS deadline of April 15th, 2022. Given the full scope of the thesis, and the **TEST-03** requirement which implies a test deadline before September 2022, a physical radiation test is still considered feasible before that time. Hence, instead of a complete termination (red), it is turned orange.
2. Continuing with the **STKH-SPACEBRAINS-01** requirement, it is considered infeasible to do a full simulation of radiation effects on the hardware components, before the ICONS deadline of April 15th, 2022. Hence, also this option will be coloured orange. Most neuromorphic chips in the DOT are proprietary, with many of the chip designs not being publically available. Since that makes it difficult to determine what the radiation effects will be on the hardware components of the chip, and how those effects, such as single-event upsets, would propagate towards influencing neurons and/or synapses. Therefore, this option is not considered feasible before April 15th 2022. It may be possible to contact manufacturers to ask how the radiation influences the neuronal- and synaptic properties. If such research is performed, it may be applied to simulate the neuromorphic hardware-radiation interaction softwarematically with sufficient accuracy to produce meaningful results.

7.1.3. Scope Of Radiation Tests

1. For the same as the last enumerated point of section 7.1.2, including the non-neuromorphic components is not considered feasible for before the ICONS deadline of April 15th, 2022. Hence, this element is also coloured orange.

7.1.4. Brain Adaptation Implementation

At the time of writing, no brain adaptation mechanisms can be eliminated.

7.1.5. Neuroplasticity Mechanism

At the time of writing, no neuroplasticity mechanisms can be eliminated.

7.1.6. Neuromorphic Architecture

1. Some neuromorphic architectures can be eliminated based on logistical reasons. Currently, the only direct access within this thesis project is to the Loihi. Furthermore, it may be expected that access to Pattern Recognition Chip by Innatera may be realised after the ICONS deadline. Similarly, the Spinnaker device may become available later-on in the project. An economic feasibility assessment needs to be made on whether they should be used in physical radiation testing or not.
2. Some of the neuromorphic chip manufacturers have been contacted in the past, these contacts may allow for access to their respective chips for physical radiation testing, if the intermediate results at ICONS are promising. Hence, they are kept orange.

7.1.7. Result Quantification

1. Since physical radiation testing is not deemed possible, and since the hardware diagrams of the respective neuromorphic architectures are not available, it is not deemed feasible to include a radiation effect analysis before ICONS. Therefore, this option is turned orange.

7.1.8. Pruned Design Option Tree

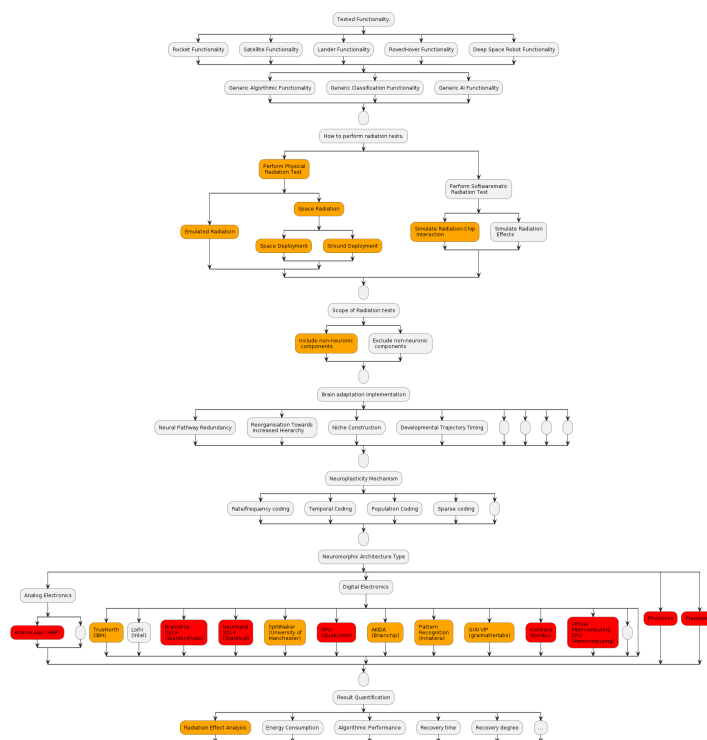


Figure 7.2: A functional flow diagram with the high-level functions of the system that is to be designed.

7.2. Preliminary Design Option Selection

With some design options eliminated, an analysis can be performed to see whether some design options are considered to be more feasible than others. This analysis is started by taking the **STKH-SPACEBRAINS-01** requirement into account, which implies results need to be produced by April 15th. This allows for expressing some design option preferences as listed in section 7.2.1 to section 7.2.7. Taking the full scope of the thesis project into account allows identification of design options that seem most feasible for a follow-up with physical radiation testing.

7.2.1. Tested Functionality Preference

For performing a generic algorithmic functionality for the following two reasons:

1. No additional dependencies such as datasets, panda packages, tensorflow etc. is required. This lowers the probability of allocating time on work that does not directly support the objective of this thesis project.
2. No preprocessing work, such as loading and/or cropping images etc., is required. This increases the amount of time that can be allocated to implementing the brain adaptation and testing its functionality.
3. Graph algorithms are typically used in space applications [?].
4. Thorough testing can quickly be set up for graph algorithms.

7.2.2. How To Perform Radiation Tests Preference

Using softwarematic radiation tests that simulate radiation effects.

7.2.3. Scope of Radiation Tests Preference

Excluding non-neural components from radiation effects allows for a complete focus using the expected radiation effects on the neural and synaptic properties, without having to model additional radiation interactions with (other) hardware elements. This increases the feasibility of producing results by April 15th.

neural networks. For this first softwarematic test, the radiation effects on non-neural/traditional hardware components (such as a memory bus, CPU etc.) of the neuromorphic architecture are ignored. The Loihi platform will be used for the simulation development and radiation tests. The test results will be interpreted in terms of performance of the graph algorithm. The most feasible brain-adaptation method and neuroplasticity mechanism will be determined using hands-on experimentation. For the second tests, a physical radiation test is proposed. For this test, the Innatera chips are currently considered as the most feasible option. Based on the results of the softwarematic testing, a decision can be made to include the traditional hardware components of the respective chips. An alternative option could be to apply extra shielding to those components, whilst saving mass on the non-shielded components through the use of brain adaptation inspired implementations.

Chapter 8

Contingency Management

Chapter 9

Market Analysis

Chapter 10

Sustainable Development Management

Chapter 11

Reporting and Quality Control

The following quality control compliance checklist can currently be generated for this project plan:

- Language Tools Grammar check applied:✓
- Language Tools Spelling check applied:✓
- CI is ran on code base used in generating this Project Plan:✗
 - Python Black Formatting Compliance:✗
 - ShellCheck Compliance:✗
 - Latex Prettier Formatting Compliance:✗
 - Python Unit Test Passing: 2 failed, 7 passed [tests]
 - Python Test Coverage %:✗[%]
 - Shell Unit Test Passing:✗[tests]
 - Shell Unit Test Coverage %:✗[%]
- Manual Quality check:
 1. Citations still have to be compiled correctly.
 2. Reproduction instructions in Appendix A should be updated.
 3. Word wrap should be applied to .uml appendices.

Chapter 12

Conclusion

Bibliography

Acronyms

ALU Arithmetic Logic Unit. 32

ANN Artificial Neural Network. 32

BIOS Basic Input/Output System. 32

BISER Built-in Soft Error Resilience. 32

BIT Binary Digit. 32

BJT Bipolar Junction Transistor. 32

CME Coronal Mass Ejection. 32

CMOS Complementary Metalâ€Oxideâ€Semiconductor. 32

CPU Central Processing Unit. 21, 32

DEC Double-error Correcting Code. 32

DICE Dual Interlocked Storage Cell. 32

DNN Deep Neural Network. 32

DNU Dual-node Upset. 32

DOD Department of Defence. 32

DOT Design Options Structuring Tree. 3, 18, 32

DRAM Dynamic Random Access Memory. 32

ECC Error Correction Code. 32

ELT Enclosed Layout Transistor. 32

ESA European Space Agency. 12, 32

FBD Functional Break-down Diagram. 3, 9, 32

FET Field-Effect Transistor. 32

FFD Functional Flow Diagram. 3, 5, 32

GCD Greatest Common Divisor. 32

GCR Galactic Cosmic Ray. 32

IC Integrated Circuit. 32

INRC Intel Neuromorphic Research Community. 32

LCM Least Common Multiple. 32

LET Linear Energy Transfer. 32

MOS Metalâ€šOxideâ€šSemiconductor. 32

MOSFET Metalâ€šOxideâ€šSemiconductor Field-Effect Transistor. 32

MPNN Multilayer Perceptron Neural Network. 32

MSN Mission Need Statement. 11, 32

NVM Non-Volatile Memory. 32

POS Project Objective Statement. 32

RAM Random Access Memory. 32

RDT Requirements Discovery Tree. 3, 11, 32

ROM Read Only Memory. 32

SEC Single-error Correcting Code. 32

SEE Single-event Effects. 32

SEGR Single-event Gate Rupture. 32

SEIB Single-event Induced Burnout. 32

SEL Single-event Latch-up. 32

SER Soft-error Rate. 32

SERL Soft-error Resilient Latch. 32

SES Single-event Snapback. 32

SET Single-event Transient. 32

SEU Single-event Upset. 32

SNN Spiking Neural Network. 32

SNU Single-node Upset. 32

SPE Solar Particle Events. 32

SPENVIS Space Environment Information System. 32

SRAM Static Random Access Memory. 32

STDP Spike-Timing-Dependent Plasticity. 32

TLB Translation Lookaside Buffer. 32

TMR Tripple-mode Redundancy. 32

VLSI Very-Large-Scale Integration. 32

WBS Work Break-down Structure. 32

WFD Work Flow Diagram. 32

Glossary

(electron) holes In the context of doped semiconductors, holes are positions where electron acceptors are located, in other words, they are holes at which the electron can go.. 32

CPU cache A small hardware memory unit that is faster than the main memory and closer to the Arithmetic Logic Unit.. 32

data cache A cache that is designed to increase the speed with which data is fetched and stored.. 32

formula A mathematical expression. 32

instruction cache A cache that is designed to increase the speed with which instructions are fetched.. 32

latex Is a mark up language specially suited for scientific documents. 32

mathematics Mathematics is what mathematicians do. 32

memory cell A fundamental/basic unit in computing that is used to store information.. 32

n-type semiconductor A semiconductor that is doped with electron donors. 32

p-n junction A boundary interface of two a p-type semiconductor and a n-type semiconductor. 32

p-type semiconductor A semiconductor that is doped with electron acceptors. 32

primary memory A form of memory that is only accessible to the Central Processing Unit which reads instructions from it and executes those instructions.. 32

prompt charge The charge that is collected by means of funnelling.. 32

random-access A memory type that has access times that are independent of its physical location.. 32

unipolar transistors Unipolar transistors are transistors that use either electrons or electron holes as charge carriers and not the combination of the two.. 32

Nomenclature

c Speed of light in a vacuum inertial frame

h Planck constant

.1. Appendix

This is a manual appendix.

.2. Appendix __main__.py

```
## Entry point for this project , runs the project code and
    ↳ exports data if
# export commands are given to the cli command that invokes
    ↳ this script.

## Import used functions.
# Project code imports.
from .create_planar_triangle_free_graph import get_graph
from .neumann import compute_mtds
from .neumann_a_t_0 import compute_mtds_a_t_0
from .arg_parser import parse_cli_args

# Export data import.
from .export_data.export_data import export_data

## Parse command line interface arguments to determine what
    ↳ this script does.
args = parse_cli_args()

## Run main code.
G = get_graph(args, False)
# compute_mtds(G)
compute_mtds_a_t_0(G)

## Run data export code if any argument is given.
if not all(arg is None for arg in [args.l, args.dd, args.sd,
    ↳ args.c2l, args.ec2l]):
    export_data(args)
```

.3. Appendix Hardcoded_data.py

```
# Specify hardcoded output data.
from .latex_export_code import export_code_to_latex
from .latex_compile import compile_latex
from .plantuml_generate import generate_all_dynamic_diagrams
from .plantuml_compile import
    ↪ compile_diagrams_in_dir_relative_to_root
from .plantuml_to_tex import export_diagrams_to_latex

class Hardcoded_data:
    """ """

    def __init__(self):

        # Specify code configuration details
        # TODO: include as optional arguments.
        self.await_compilation = True
        self.verbose = True
        self.gantt_extension = ".uml"
        self.diagram_extension = ".png"

        # Filenames.
        self.main_latex_filename = "report.tex"
        self.export_data_dirname = "export_data"
        self.diagram_dir = "Diagrams"
        self.plantuml_java_filename = "plantuml.jar"

        # Folder names.
        self.dynamic_diagram_dir = "Dynamic_diagrams"
        self.static_diagram_dir = "Static_diagrams"

        # Specify paths relative to root.
        self.path_to_export_data_from_root = f"src/{self.
            ↪ export_data_dirname}"
        self.jar_path_relative_from_root = (
            f"{self.path_to_export_data_from_root}/{self.
            ↪ plantuml_java_filename}"
        )
        self.diagram_output_dir_relative_to_root = f"latex/
            ↪ Images/{self.diagram_dir}"

        # Path related variables
        self.relative_src_filepath_from_root = f"src/"
        self.append_export_code_to_latex = True
        self.path_to_dynamic_gantts = f"{self.
            ↪ path_to_export_data_from_root}/{self.diagram_dir
            ↪}/{self.dynamic_diagram_dir}"
        self.path_to_static_gantts = f"{self.
            ↪ path_to_export_data_from_root}/{self.diagram_dir
            ↪}/{self.static_diagram_dir}"
```

.4. Appendix Plot_to_tex.py

```

### Call this from another file , for project 11, question 3b:
### from Plot_to_tex import Plot_to_tex as plt_tex
### multiple_y_series = np.zeros((nrOfDataSeries ,
    ↳ nrOfDataPoints), dtype=int); # actually fill with data
### lineLabels = [] # add a label for each dataseries
### plt_tex.plotMultipleLines(plt_tex , single_x_series ,
    ↳ multiple_y_series , "x-axis label [units]" , "y-axis label [
    ↳ units]" , lineLabels , "3 b" , 4 , 11)
### 4b=filename
### 4 = position of legend , e.g. top right.
###
### For a single line , use:
### plt_tex.plotSingleLine(plt_tex , range(0 , len(dataseries)) ,
    ↳ dataseries , "x-axis label [units]" , "y-axis label [units]" ,
    ↳ lineLabel , "3 b" , 4 , 11)

### You can also plot a table directly into latex , see
    ↳ example_create_a_table(..)
###
### Then put it in latex with for example:
### \begin{table}[H]
###     \centering
###     \caption{Results some computation.}\label{tab:
    ↳ some_computation}
###     \begin{tabular}{|c|c|} % remember to update this to
    ↳ show all columns of table
###         \hline
###         \input{latex/project3/tables/q2.txt}
###     \end{tabular}
### \end{table}
import random
from matplotlib import lines
import matplotlib.pyplot as plt
import numpy as np
import os

class Plot_to_tex:
    """ """

    def __init__(self):
        self.script_dir = self.get_script_dir()
        print("Created main")

    # plot graph (legendPosition = integer 1 to 4)
    def plotSingleLine(
        self,
        x_path,
        y_series,
        x_axis_label,
        y_axis_label,
        label,
        filename,
        legendPosition,

```

```

        project_name,
    ):
        """

        :param x_path: param y_series:
        :param x_axis_label: param y_axis_label:
        :param label: param filename:
        :param legendPosition: param project_name:
        :param y_series: param y_axis_label:
        :param filename: param project_name:
        :param y_axis_label: param project_name:
        :param project_name:

        """
        fig = plt.figure()
        ax = fig.add_subplot(111)
        ax.plot(x_path, y_series, c="b", ls="-", label=label,
            ↪ fillstyle="none")
        plt.legend(loc=legendPosition)
        plt.xlabel(x_axis_label)
        plt.ylabel(y_axis_label)
        plt.savefig(
            os.path.dirname(__file__)
            + f"/../../../latex/{project_name}"
            + "/Images/"
            + filename
            + ".png"
        )

#         plt.show() ;

# plot graphs
def plotMultipleLines(
    self,
    x,
    y_series,
    x_label,
    y_label,
    label,
    filename,
    legendPosition,
    project_name,
):
    """

    :param x: param y_series:
    :param x_label: param y_label:
    :param label: param filename:
    :param legendPosition: param project_name:
    :param y_series: param y_label:
    :param filename: param project_name:
    :param y_label: param project_name:
    :param project_name:

    """
    fig = plt.figure()

```

```

ax = fig.add_subplot(111)

# generate colours
cmap = self.get_cmap(len(y_series[:, 0]))

# generate line types
lineTypes = self.generateLineTypes(y_series)

for i in range(0, len(y_series)):
    # overwrite linetypes to single type
    lineTypes[i] = "-"
    ax.plot(
        x,
        y_series[i, :],
        ls=lineTypes[i],
        label=label[i],
        fillstyle="none",
        c=cmap(i),
    )
    # color

# configure plot layout
plt.legend(loc=legendPosition)
plt.xlabel(x_label)
plt.ylabel(y_label)
plt.savefig(
    os.path.dirname(__file__)
    + f"/../../../latex/{project_name}"
    + "/Images/"
    + filename
    + ".png"
)

print(f"plotted lines")

# Generate random line colours
# Source: https://stackoverflow.com/questions/14720331/how
# -to-generate-random-colors-in-matplotlib
def get_cmap(n, name="hsv"):
    """Returns a function that maps each index in 0, 1,
    ..., n-1 to a distinct
    RGB color; the keyword argument name must be a
    standard mpl colormap name.

    :param n:
    :param name: (Default value = "hsv")
    :param name: Default value = "hsv")

    """
    return plt.cm.get_cmap(name, n)

def generateLineTypes(y_series):
    """

    :param y_series:

    """

```

```

# generate varying linetypes
typeOfLines = list(lines.lineStyles.keys())

while len(y_series) > len(typeOfLines):
    typeOfLines.append("-.")

# remove void lines
for i in range(0, len(y_series)):
    if typeOfLines[i] == "None":
        typeOfLines[i] = "-"
    if typeOfLines[i] == ":":
        typeOfLines[i] = ":"
    if typeOfLines[i] == " ":
        typeOfLines[i] = "--"
return typeOfLines

# Create a table with: table_matrix = np.zeros((4,4), dtype
→ =object) and pass it to this object
def put_table_in_tex(self, table_matrix, filename,
→ project_name):
    """

    :param table_matrix: param filename:
    :param project_name: param filename:
    :param filename:

    """
    cols = np.shape(table_matrix)[1]
    format = "%s"
    for col in range(1, cols):
        format = format + " & %s"
    format = format + ""
    plt.savetxt(
        os.path.dirname(__file__)
        + f"/../../../latex/{project_name}"
        + "/tables/"
        + filename
        + ".txt",
        table_matrix,
        delimiter=" & ",
        fmt=format,
        newline=" \\ \\ \\ \\ \\hline \\n",
    )

# replace this with your own table creation and then pass
→ it to put_table_in_tex(..)
def example_create_a_table(self):
    """ """
    project_name = "1"
    table_name = "example_table_name"
    rows = 2
    columns = 4
    table_matrix = np.zeros((rows, columns), dtype=object)
    table_matrix[:, :] = "" # replace the standard zeros
    → with empty cell
    print(table_matrix)

```

```
    for column in range(0, columns):
        for row in range(0, rows):
            table_matrix[row, column] = row + column
    table_matrix[1, 0] = "example"
    table_matrix[0, 1] = "grid sizes"

    self.put_table_in_tex(table_matrix, table_name,
        ↪ project_name)

def get_script_dir(self):
    """returns the directory of this script regardless of
        ↪ from which level the code is executed"""
    return os.path.dirname(__file__)

if __name__ == "__main__":
    main = Plot_to_tex()
    main.example_create_a_table()
```

.5. Appendix arg_parser.py

```

# This is the main code of this project nr, and it manages
    ↪ running the code and
# outputting the results to LaTeX.
import argparse

def parse_cli_args():
    # Instantiate the parser
    parser = argparse.ArgumentParser(description="Optional app
    ↪ description")

    ## Include argument parsing for default code.
    # Allow user to load a graph from file.
    parser.add_argument(
        "--g",
        dest="graph_from_file",
        action="store_true",
        help="boolean flag, determines whether energy analysis
        ↪ graph is created from file ",
    )

    # Allow user to specify an infile.
    parser.add_argument("infile", nargs="?", type=argparse.
    ↪ FileType("r"))

    # Specify default argument values for the parser.
    parser.set_defaults(
        infile=None, graph_from_file=False,
    )

    ## Include argument parsing for data exporting code.
    # Compile LaTeX
    parser.add_argument(
        "--l", action="store_true", help="Boolean indicating
        ↪ if code compiles LaTeX"
    )

    # Generate, compile and export Dynamic PlantUML diagrams
    ↪ to LaTeX.
    parser.add_argument(
        "--dd",
        action="store_true",
        help="A boolean indicating if code generated diagrams
        ↪ are compiled and exported.",
    )

    # Generate, compile and export Static PlantUML diagrams to
    ↪ LaTeX.
    parser.add_argument(
        "--sd",
        action="store_true",
        help="A boolean indicating if static diagrams are
        ↪ compiled and exported.",
    )

```

```
# Export the project code to LaTeX.
parser.add_argument(
    "--c2l",
    action="store_true",
    help="A boolean indicating if project code is exported
    ↪ to LaTeX.",
)

# Export the exporting code, and the project code to LaTeX
↪ .
parser.add_argument(
    "--ec2l",
    action="store_true",
    help="A boolean indicating if code that exports code
    ↪ is exported to LaTeX.",
)

# Load the arguments that are given.
args = parser.parse_args()
return args
```

.6. Appendix create_dynamic_diagrams.py

```
# Data export imports.
from .plantuml_generate import generate_all_dynamic_diagrams
from .plantuml_compile import
    ↪ compile_diagrams_in_dir_relative_to_root
from .plantuml_to_tex import export_diagrams_to_latex

def create_dynamic_diagrams(args, hd):
    ## PlantUML
    # Generate PlantUML diagrams dynamically (using code).
    if args.dd:
        generate_all_dynamic_diagrams(
            f"{hd.path_to_export_data_from_root}/Diagrams/
            ↪ Dynamic"
        )

    # Compile dynamically generated PlantUML diagrams to
    ↪ images.
    compile_diagrams_in_dir_relative_to_root(
        hd.await_compilation,
        hd.gantt_extension,
        hd.jar_path_relative_from_root,
        hd.path_to_dynamic_gantts,
        hd.verbose,
    )

    # Export dynamic PlantUML text files to LaTeX.
    export_diagrams_to_latex(
        hd.path_to_dynamic_gantts,
        hd.gantt_extension,
        hd.diagram_output_dir_relative_to_root,
    )

    # Export dynamic PlantUML diagram images to LaTeX.
    export_diagrams_to_latex(
        hd.path_to_dynamic_gantts,
        hd.diagram_extension,
        hd.diagram_output_dir_relative_to_root,
    )
```

.7. Appendix create_planar_triangle_free_graph.py

```

# This code creates a planar graph without triangles.
# Planar implies that it fits on a 2D plane, without any edges
  ↪ crossing.
# Triangle free means there are no triangles in the graph.

import networkx as nx

import matplotlib.pyplot as plt

def get_graph(args, show_graph):
    if args.infile and not args.graph_from_file:
        try:
            graph = nx.read_graphml(args.infile.name)
        except Exception as exc:
            raise Exception(
                "Supplied input file is not a gml networkx
                 ↪ graph object."
            ) from exc
    else:
        graph = create_manual_test_graph()
    if show_graph:
        plot_graph(graph)
    return graph

def create_manual_test_graph():
    """
    creates manual test graph with 7 undirected nodes.
    """

    graph = nx.Graph()
    graph.add_nodes_from(
        ["a", "b", "c", "d", "e", "f", "g"],
        x=0,
        color="w",
        dynamic_degree=0,
        delta_two=0,
        p=0,
        xds=0,
    )
    graph.add_edges_from(
        [
            ("a", "b"),
            ("a", "c"),
            # ("b", "c"),
            # ("b", "d"),
            ("c", "d"),
            ("d", "e"),
            ("e", "g"),
            ("b", "e"),
            ("b", "f"),
            ("f", "g"),
        ]
    )

```

```
)  
    return graph  
  
# TODO: move to helper  
def plot_graph(G):  
    options = {"with_labels": True, "node_color": "white", "  
        ↪ edgecolors": "blue"}  
    nx.draw_networkx(G, **options)  
    plt.show()  
    plt.clf()
```

.8. Appendix create_static_diagrams.py

```
# Data export imports.
from .plantuml_generate import generate_all_dynamic_diagrams
from .plantuml_compile import
    ↪ compile_diagrams_in_dir_relative_to_root
from .plantuml_to_tex import export_diagrams_to_latex

def create_static_diagrams(args, hd):

    ## PlantUML
    if args.sd:
        # Compile statically generated PlantUML diagrams to
        ↪ images.
        compile_diagrams_in_dir_relative_to_root(
            hd.await_compilation,
            hd.gantt_extension,
            hd.jar_path_relative_from_root,
            hd.path_to_static_gantts,
            hd.verbose,
        )

        # Export static PlantUML text files to LaTeX.
        export_diagrams_to_latex(
            hd.path_to_static_gantts,
            hd.gantt_extension,
            hd.diagram_output_dir_relative_to_root,
        )

        # Export static PlantUML diagram images to LaTeX.
        export_diagrams_to_latex(
            hd.path_to_static_gantts,
            hd.diagram_extension,
            hd.diagram_output_dir_relative_to_root,
        )
```

.9. Appendix distributed.py

```
# 1.a Each vertex v_i chooses random float r_i in range 0<r_i
    ↳ <1

# 1.b Each vertex v_i computes d_i. d_i=degree of vertex v_i
# 1.c Each vertex v_i computes weight: w_i=d_i+r_i
# 1.d Each vertex v_i sends w_i to each of its neighbours.

# 2.a Each vertex v_i gets the index of the
# neighbouring vertex v_j_(w_max) that has the heighest w_i,
    ↳ with i!=j.
# 2.b Each vertex v_i adds a mark to that neighbour vertex
    ↳ v_j_(w_max).

# 3 for k in range [0,m] rounds , do:

# 4.a Each node v_i computes how many marks it has received ,
    ↳ as (x_i)_k.
# 4.b Each node v_i computes (w_i)_k=(x_i)_k+ r_i

# 5. Reset marked vertices: for each vertex v_i , (x_i)_k=0

# 6.a Each vertex v_i computes d_i. d_i=degree of vertex v_i
# 6.b Each vertex v_i sends w_i to each of its neighbours.
# 6.c Each vertex v_i gets the index of the
# neighbouring vertex v_j_(w_max) that has the heighest w_i,
    ↳ with i!=j.
# 6.d Each vertex v_i adds a mark to that neighbour vertex
    ↳ v_j_(w_max).
```

.10. Appendix export_data.py

```
# Data export imports.
from .Hardcoded_data import Hardcoded_data

from .create_dynamic_diagrams import create_dynamic_diagrams
from .create_static_diagrams import create_static_diagrams
from .latex_export_code import export_code_to_latex
from .latex_compile import compile_latex

def export_data(args):

    hd = Hardcoded_data()

    ## PlantUML
    create_dynamic_diagrams(args, hd)
    create_static_diagrams(args, hd)

    ## Plotting
    # Generate plots.
    # Export plots to LaTeX.

    ## Export code to LaTeX.
    if args.c2l:
        # TODO: verify whether the latex /{project_name}/
        #       ↳ Appendices folder exists before exporting.
        # TODO: verify whether the latex /{project_name}/ Images
        #       ↳ folder exists before exporting.
        export_code_to_latex(hd.main_latex_filename, False)
    elif args.ec2l:
        # TODO: verify whether the latex /{project_name}/
        #       ↳ Appendices folder exists before exporting.
        # TODO: verify whether the latex /{project_name}/ Images
        #       ↳ folder exists before exporting.
        export_code_to_latex(hd.main_latex_filename, hd.
        #       ↳ append_export_code_to_latex)

    ## Compile the accompanying LaTeX report.
    if args.l:
        compile_latex(True, True)
        print("")
    print(f"\n\nDone exporting data.")
```

.11. Appendix helper_bash_commands.py

```
import subprocess
```

```
def run_bash_command(await_compilation, bash_command, verbose)
    :
    if await_compilation:
        if verbose:
            subprocess.call(bash_command, shell=True)
        else:
            subprocess.call(
                bash_command,
                shell=True,
                stderr=subprocess.DEVNULL,
                stdout=subprocess.DEVNULL,
            )
    else:
        if verbose:
            subprocess.Popen(bash_command, shell=True)
        else:
            subprocess.Popen(
                bash_command,
                shell=True,
                stderr=subprocess.DEVNULL,
                stdout=subprocess.DEVNULL,
            )
```

.12. Appendix helper_dir_file_edit.py

```

import os
import shutil
import glob

def delete_directory_contents(self, folder):
    """
    :param folder: Directory that is being deleted.

    """
    for filename in os.listdir(folder):
        file_path = os.path.join(folder, filename)
        try:
            if os.path.isfile(file_path) or os.path.islink(
                ↪ file_path):
                os.unlink(file_path)
            elif os.path.isdir(file_path):
                shutil.rmtree(file_path)
        except Exception as e:
            print("Failed to delete %s. Reason: %s" % (
                ↪ file_path, e))

def file_contains(filepath, substring):
    with open(filepath) as f:
        if substring in f.read():
            return True

def get_dir_filelist_based_on_extension(dir_relative_to_root,
    ↪ extension):
    """
    :param dir_relative_to_root: A relative directory as seen
        ↪ from the root dir of this project.
    :param extension: The file extension that is used/searched
        ↪ in this function.

    """
    selected_filenames = []
    # TODO: assert directory exists
    for filename in os.listdir(dir_relative_to_root):
        if filename.endswith(extension):
            selected_filenames.append(filename)
    return selected_filenames

def create_dir_relative_to_root_if_not_exists(
    ↪ dir_relative_to_root):
    """

    :param dir_relative_to_root: A relative directory as seen
        ↪ from the root dir of this project.

```

```

"""
    if not os.path.exists(dir_relative_to_root):
        os.makedirs(dir_relative_to_root)

def delete_dir_relative_to_root_if_not_exists(
    ↪ dir_relative_to_root):
    """

    :param dir_relative_to_root: A relative directory as seen
        ↪ from the root dir of this project.

    """
    if os.path.exists(dir_relative_to_root):
        # Remove directory and its content.
        shutil.rmtree(dir_relative_to_root)

def dir_relative_to_root_exists(dir_relative_to_root):
    """

    :param dir_relative_to_root: A relative directory as seen
        ↪ from the root dir of this project.

    """
    if not os.path.exists(dir_relative_to_root):
        return False
    elif os.path.exists(dir_relative_to_root):
        return True
    else:
        raise Exception(
            "Directory relative to root: {dir_relative_to_root}
            ↪ did not exist, nor did it exist."
        )

def get_all_files_in_dir_and_child_dirs(extension, path,
    ↪ excluded_files=None):
    """ Returns a list of the relative paths to all files
        ↪ within the some path that match
        the given file extension. Also includes files in child
        ↪ directories.

    :param extension: The file extension that is used/searched
        ↪ in this function. The file extension of the file
        ↪ that is sought in the appendix line. Either ".py" or
        ↪ ".pdf".
    :param path: Absolute filepath in which files are being
        ↪ sought.
    :param excluded_files: Default value = None) Files that
        ↪ will not be included even if they are found.

    """
    filepaths = []
    for r, d, f in os.walk(path):

```

```

        for file in f:
            if file.endswith(extension):
                if (excluded_files is None) or (
                    (not excluded_files is None) and (not file
                     ↪ in excluded_files)
                ):
                    filepaths.append(r + "/" + file)
    return filepaths

def get_filepaths_in_dir(extension, path, excluded_files=None)
↪ :
    """ Returns a list of the relative paths to all files
        ↪ within the some path that match
        the given file extension. Does not include files in
        ↪ child_directories.

    :param extension: The file extension that is used/ searched
        ↪ in this function. The file extension of the file
        ↪ that is sought in the appendix line. Either ".py" or
        ↪ ".pdf".
    :param path: Absolute filepath in which files are being
        ↪ sought.
    :param excluded_files: Default value = None) Files that
        ↪ will not be included even if they are found.

    """
    filepaths = []
    current_path = os.getcwd()
    os.chdir(path)
    for file in glob.glob(f"*.{extension}"):
        print(file)
        if (excluded_files is None) or (
            (not excluded_files is None) and (not file in
             ↪ excluded_files)
        ):
            filepaths.append(f"{path}/{file}")
    os.chdir(current_path)
    return filepaths

```

.13. Appendix latex_compile.py

```
# Compiles the latex report using the compile script.
import subprocess

from .helper_bash_commands import run_bash_command

def compile_latex(await_compilation, verbose):
    """
    Compiles the LaTeX report of this project using its
    ↪ compile script.

    :param await_compilation: Make python wait untill the
    ↪ PlantUML compilation is completed.
    :param project_name: The name of the project that is being
    ↪ executed/ran.
    :param verbose: True, ensures compilation output is
    ↪ printed to terminal, False means compilation is
    ↪ silent.

    Returns:
        Nothing.

    Raises:
        Nothing.
    """

    # Ensure compile script is runnable.
    bash_make_compile_script_runnable_command = f"chmod +x
    ↪ latex/compile_script.sh"
    run_bash_command(
        await_compilation,
        ↪ bash_make_compile_script_runnable_command,
        ↪ verbose
    )

    # Run latex compilation script to compile latex project.
    bash_compilation_command = f"latex/compile_script.sh"
    run_bash_command(await_compilation,
        ↪ bash_compilation_command, verbose)
```

.14. Appendix latex_export_code.py

```

# runs a jupyter notebook and converts it to pdf
import os
import shutil
import ntpath

from .helper_dir_file_edit import
    ↪ get_all_files_in_dir_and_child_dirs
from .helper_dir_file_edit import get_filepaths_in_dir
from .helper_dir_file_edit import file_contains

def export_code_to_latex(main_latex_filename,
    ↪ include_export_code):
    """ This function exports the python files and compiled
        ↪ pdfs of jupyter notebooks into the
        latex of the same project number. First it scans which
        ↪ appendices (without code, without
        notebooks) are already manually included in the main latex
        ↪ code. Next, all appendices
        that contain the python code are either found or created in
        ↪ the following order:
        First, the __main__.py file is included, followed by the
        ↪ main.py file, followed by all
        python code files in alphabetic order. After this, all the
        ↪ pdfs of the compiled notebooks
        are added in alphabetic order of filename. This order of
        ↪ appendices is overwritten in the
        main tex file.

    :param main_latex_filename: Name of the main latex
        ↪ document of this project number
    :param project_name: The name of the project that is being
        ↪ executed/ran. The number indicating which project
        ↪ this code pertains to.

    """
    script_dir = get_script_dir()
    latex_dir = script_dir + "/../..//latex/"
    appendix_dir = f"{latex_dir}Appendices/"
    path_to_main_latex_file = f"{latex_dir}{
        ↪ main_latex_filename}"
    normalised_root_dir = script_dir + "/../..//"
    normalised_root_dir = os.path.normpath(normalised_root_dir
        ↪ )
    src_dir = script_dir + "/../"

    # Verify the latex file supports auto-generated python
    ↪ appendices.
    verify_latex_supports_auto_generated_appendices(
        ↪ path_to_main_latex_file)
    print(f"1={os.getcwd()}")
    # Get paths to files containing project python code.
    python_filepaths = get_filepaths_in_dir("py", src_dir, ["
        ↪ __init__.py"])

```

```

print(f"2={os.getcwd()}")
compiled_notebook_pdf_filepaths =
    ↪ get_compiled_notebook_paths(script_dir)
print(f"Before, python_filepaths={python_filepaths}")

# Get paths to the files containing the latex export code
if include_export_code:
    python_filepaths.extend(get_filepaths_in_dir("py",
    ↪ script_dir, ["__init__.py"]))
print(f"After, python_filepaths={python_filepaths}")
print(f"3={os.getcwd()}")

# Check which files are already included in the latex
    ↪ appendices.
python_files_already_included_in_appendices =
    ↪ get_code_files_already_included_in_appendices(
        python_filepaths, appendix_dir, ".py",
        ↪ normalised_root_dir
    )
notebook_pdf_files_already_included_in_appendices =
    ↪ get_code_files_already_included_in_appendices(
        compiled_notebook_pdf_filepaths, appendix_dir, ".ipynb",
        ↪ "", normalised_root_dir,
    )
print(
    f"python_files_already_included_in_appendices={
    ↪ python_files_already_included_in_appendices}"
)
print(f"4={os.getcwd()}")
# Get which appendices are still missing.
missing_python_files_in_appendices =
    ↪ get_code_files_not_yet_included_in_appendices(
        python_filepaths,
        ↪ python_files_already_included_in_appendices, ".py",
        ↪ ""
    )
missing_notebook_files_in_appendices =
    ↪ get_code_files_not_yet_included_in_appendices(
        compiled_notebook_pdf_filepaths,
        notebook_pdf_files_already_included_in_appendices,
        ↪ ".pdf",
    )
print(f"5={os.getcwd()}")
print(f"missing_python_files_in_appendices={
    ↪ missing_python_files_in_appendices}")
# Create the missing appendices.
created_python_appendix_filenames =
    ↪ create_appendices_with_code(
        appendix_dir, missing_python_files_in_appendices, ".py",
        ↪ "", normalised_root_dir
    )
created_notebook_appendix_filenames =
    ↪ create_appendices_with_code(
        appendix_dir,
        missing_notebook_files_in_appendices,
        ↪ ".ipynb",
    )

```

```

        normalised_root_dir,
    )
    print(f"created_python_appendix_filenames={
        ↪ created_python_appendix_filenames}")
    print(f"6={os.getcwd()}")
    appendices = get_list_of_appendix_files(
        appendix_dir, compiled_notebook_pdf_filepaths,
        ↪ python_filepaths
    )
    print(f"appendices={appendices}")
    main_tex_code, start_index, end_index, appendix_tex_code =
        ↪ get_appendix_tex_code(
            path_to_main_latex_file
        )
    print(f"7={os.getcwd()}")
    # assumes non-included non-code appendices should not be
    ↪ included:
    # overwrite the existing appendix lists with the current
    ↪ appendix list.
    (
        non_code_appendices,
        main_non_code_appendix_inclusion_lines,
    ) = get_order_of_non_code_appendices_in_main(appendices,
        ↪ appendix_tex_code)
    print(f"8={os.getcwd()}")
    python_appendix_filenames = list(
        map(
            lambda x: x.appendix_filename,
            filter_appendices_by_type(appendices, "python"),
        )
    )
    print(f"9={os.getcwd()}")
    print(f"python_appendix_filenames={
        ↪ python_appendix_filenames}")
    # exit ()
    sorted_created_python_appendices = sort_python_appendices(
        filter_appendices_by_type(appendices, "python")
    )
    sorted_python_appendix_filenames = list(
        map(lambda x: x.appendix_filename,
            ↪ sorted_created_python_appendices)
    )
    print(f"10={os.getcwd()}")

    notebook_appendix_filenames = list(
        map(
            lambda x: x.appendix_filename,
            filter_appendices_by_type(appendices, "notebook"),
        )
    )
    sorted_created_notebook_appendices =
        ↪ sort_notebook_appendices_alphabetically(
            filter_appendices_by_type(appendices, "notebook")
        )
    sorted_notebook_appendix_filenames = list(

```

```

        map(lambda x: x.appendix_filename,
             ↪ sorted_created_notebook_appendices)
    )
    print(f"11={os.getcwd()}")
    appendix_latex_code = create_appendices_latex_code(
        main_latex_filename,
        main_non_code_appendix_inclusion_lines,
        sorted_created_notebook_appendices,
        sorted_created_python_appendices,
    )
    print(f"12={os.getcwd()}")
    updated_main_tex_code = substitute_appendix_code(
        end_index, main_tex_code, start_index,
        ↪ appendix_latex_code
    )
    # print(f"\n\n")
    # for line in updated_main_tex_code:
    #     print(line)
    # print(f"updated_main_tex_code={updated_main_tex_code}")
    # print(f"\n\n")
    print(f"13={os.getcwd()}")
    overwrite_content_to_file(updated_main_tex_code,
        ↪ path_to_main_latex_file)
    print(f"14={os.getcwd()}")

def verify_latex_supports_auto_generated_appendices(
    ↪ path_to_main_latex_file):
    print("hi")
    determining_overleaf_home_line = "\def\overleafhome{/tmp}%"
    ↪ change as appropriate"
    begin_apendices_line = "\\begin{appendices}"
    print(f"determining_overleaf_home_line={
        ↪ determining_overleaf_home_line}")
    print(f"begin_apendices_line={begin_apendices_line}")

    if not file_contains(path_to_main_latex_file,
        ↪ determining_overleaf_home_line):
        raise Exception(
            f"Error, {path_to_main_latex_file} does not
            ↪ contain:\n\n{determining_overleaf_home_line}\n\n
            ↪ so this Python code cannot export the
            ↪ code as latex appendices."
        )
    if not file_contains(path_to_main_latex_file,
        ↪ determining_overleaf_home_line):
        raise Exception(
            f"Error, {path_to_main_latex_file} does not
            ↪ contain:\n\n{begin_apendices_line}\n\n so
            ↪ this Python code cannot export the code as
            ↪ latex appendices."
        )

def create_appendices_latex_code(
    main_latex_filename,

```

```

    main_non_code_appendix_inclusion_lines,
    notebook_appendices,
    python_appendices,
):
    """ Creates the latex code that includeds the appendices in
        ↳ the main latex file .

    :param main_non_code_appendix_inclusion_lines: latex code
        ↳ that includes the appendices that do not contain
        ↳ python code nor notebooks
    :param notebook_appendices: List of Appendix objects
        ↳ representing appendices that include the pdf files of
        ↳ compiled Jupiter notebooks
    :param project_name: The name of the project that is being
        ↳ executed / ran. The number indicating which project
        ↳ this code pertains to.
    :param python_appendices: List of Appendix objects
        ↳ representing appendices that include the python code
        ↳ files .
    :param main_latex_filename :

    """
    main_appendix_inclusion_lines =
        ↳ main_non_code_appendix_inclusion_lines
    print(f"BEFORE main_appendix_inclusion_lines={
        ↳ main_appendix_inclusion_lines}")

    appendices_of_all_types = [python_appendices,
        ↳ notebook_appendices]

    print(f"\n\n")
    main_appendix_inclusion_lines.append("\ifhidesourcecode{}\
        ↳ else{")
    main_appendix_inclusion_lines.append(
        f"\IfFileExists{{\latex/{main_latex_filename}}}{{"
    )
    main_appendix_inclusion_lines =
        ↳ append_latex_inclusion_command(
            appendices_of_all_types, True,
            ↳ main_appendix_inclusion_lines,
        )
    main_appendix_inclusion_lines.append(f"}}{{{"")
    main_appendix_inclusion_lines =
        ↳ append_latex_inclusion_command(
            appendices_of_all_types, False,
            ↳ main_appendix_inclusion_lines,
        )
    print(f"AFTER main_appendix_inclusion_lines={
        ↳ main_appendix_inclusion_lines}")
    return main_appendix_inclusion_lines

def append_latex_inclusion_command(
    appendices_of_all_types, is_from_normalised_root_dir,
        ↳ main_appendix_inclusion_lines,
):

```

```

"""

:param appendices_of_all_types: param
    ↳ is_from_normalised_root_dir:
:param main_appendix_inclusion_lines: param project_name:
:param is_from_normalised_root_dir: param project_name:
:param project_name: The name of the project that is being
    ↳ executed / ran.

"""
for appendix_type in appendices_of_all_types:
    for appendix in appendix_type:
        line = update_appendix_tex_code(
            appendix.appendix_filename,
            ↳ is_from_normalised_root_dir
        )
        print(f"appendix.appendix_filename={appendix.
            ↳ appendix_filename}")
        main_appendix_inclusion_lines.append(line)
return main_appendix_inclusion_lines

def filter_appendices_by_type(appendices, appendix_type):
    """Returns the list of all appendices of a certain
        ↳ appendix type, from the incoming list of Appendix
        ↳ objects.

    :param appendices: List of Appendix objects
    :param appendix_type: Can consist of "no_code", "python",
        ↳ or "notebook" and indicates different appendix types

    """
    return_appendices = []
    for appendix in appendices:
        if appendix.appendix_type == appendix_type:
            return_appendices.append(appendix)
    return return_appendices

def sort_python_appendices(appendices):
    """First puts __main__.py, followed by main.py followed by
        ↳ a-z code files.

    :param appendices: List of Appendix objects

    """
    return_appendices = []
    for appendix in appendices: # first get appendix
        ↳ containing __main__.py
        if (appendix.code_filename == "__main__.py") or (
            appendix.code_filename == "__Main__.py"
        ):
            return_appendices.append(appendix)
            appendices.remove(appendix)
    for appendix in appendices: # second get appendix
        ↳ containing main.py

```



```

        if (appendix.code_filename == "main.py") or (
            appendix.code_filename == "Main.py"
        ):
            return_appendices.append(appendix)
            appendices.remove(appendix)
    return_appendices

# Filter remaining appendices in order of a-z
filtered_remaining_appendices = [
    i for i in appendices if i.code_filename is not None
]
appendices_sorted_a_z = sort_appendices_on_code_filename(
    filtered_remaining_appendices
)
return return_appendices + appendices_sorted_a_z

def sort_notebook_appendices_alphabetically(appendices):
    """ Sorts notebook appendix objects alphabetic order of
        ↳ their pdf filenames.

    :param appendices: List of Appendix objects

    """
    return_appendices = []
    filtered_remaining_appendices = [
        i for i in appendices if i.code_filename is not None
    ]
    appendices_sorted_a_z = sort_appendices_on_code_filename(
        filtered_remaining_appendices
    )
    return return_appendices + appendices_sorted_a_z

def sort_appendices_on_code_filename(appendices):
    """ Returns a list of Appendix objects that are sorted and
        ↳ based on the property: code_filename.
    Assumes the incoming appendices only contain python files.

    :param appendices: List of Appendix objects

    """
    attributes = list(map(lambda x: x.code_filename,
        ↳ appendices))
    sorted_indices = sorted(range(len(attributes)), key=lambda
        ↳ k: attributes[k])
    sorted_list = []
    for i in sorted_indices:
        sorted_list.append(appendices[i])
    return sorted_list

def get_order_of_non_code_appendices_in_main(appendices,
    ↳ appendix_tex_code):
    """ Scans the lines of appendices in the main code, and
        ↳ returns the lines

```

of the appendices that do not contain code, in the order
 ↳ in which they were
 included in the main latex file.

```
:param appendices: List of Appendix objects
:param appendix_tex_code: latex code from the main latex
  ↳ file that includes the appendices

"""
non_code_appendices = []
non_code_appendix_lines = []
appendix_tex_code = list(dict.fromkeys(appendix_tex_code))
for line in appendix_tex_code:
    appendix_filename =
        ↳ get_filename_from_latex_appendix_line(appendices,
        ↳ line)

    # Check if line is not commented
    if not appendix_filename is None:
        if not line_is_commented(line, appendix_filename):
            appendix = get_appendix_from_filename(
                ↳ appendices, appendix_filename)
            if appendix.appendix_type == "no_code":
                non_code_appendices.append(appendix)
                non_code_appendix_lines.append(line)
return non_code_appendices, non_code_appendix_lines
```

```
def get_filename_from_latex_appendix_line(appendices,
  ↳ appendix_line):
    """Returns the first filename from a list of incoming
        ↳ filenames that
    occurs in a latex code line.

    :param appendices: List of Appendix objects
    :param appendix_line: latex code (in particular expected
        ↳ to be the code from main that is used to include
        ↳ appendix latex files.)

    """
    for filename in list(map(lambda appendix: appendix.
        ↳ appendix_filename, appendices)):
        if filename in appendix_line:
            if not line_is_commented(appendix_line, filename):
                return filename
```

```
def get_appendix_from_filename(appendices, appendix_filename):
    """Returns the first Appendix object with an appendix
        ↳ filename that matches the incoming appendix_filename.
    The Appendix objects are selected from an incoming list of
        ↳ Appendix objects.

    :param appendices: List of Appendix objects
    :param appendix_filename: name of a latex appendix file ,
        ↳ ends in .tex ,
```

```

"""
for appendix in appendices:
    if appendix_filename == appendix.appendix_filename:
        return appendix

def get_compiled_notebook_paths(script_dir):
    """ Returns the list of jupyter notebook filepaths that
        ↳ were compiled successfully and that are
        included in the same dir as this script (the src directory).

    :param script_dir: absolute path of this file.

    """
    notebook_filepaths = get_all_files_in_dir_and_child_dirs("
        ↳ .ipynb", script_dir)
    compiled_notebook_filepaths = []

    # check if the jupyter notebooks were compiled
    for notebook_filepath in notebook_filepaths:

        # swap file extension
        notebook_filepath = notebook_filepath.replace(".ipynb"
            ↳ , ".pdf")

        # check if file exists
        if os.path.isfile(notebook_filepath):
            compiled_notebook_filepaths.append(
                ↳ notebook_filepath)
    return compiled_notebook_filepaths

def get_list_of_appendix_files(
    appendix_dir, absolute_notebook_filepaths,
    ↳ absolute_python_filepaths
):
    """ Returns a list of Appendix objects that contain all the
        ↳ appendix files with .tex extension.

    :param appendix_dir: Absolute path that contains the
        ↳ appendix .tex files.
    :param absolute_notebook_filepaths: List of absolute paths
        ↳ to the compiled notebook pdf files.
    :param absolute_python_filepaths: List of absolute paths
        ↳ to the python files.

    """
    appendices = []
    appendices_paths = get_all_files_in_dir_and_child_dirs("
        ↳ .tex", appendix_dir)
    print(f"appendix_dir={appendix_dir}")
    print(f"appendices_paths={appendices_paths}")
    # exit()
    for appendix_filepath in appendices_paths:
        appendix_type = "no_code"

```

```

appendix_filecontent = read_file(appendix_filepath)
line_nr_python_file_inclusion =
    ↪ get_line_of_latex_command(
        appendix_filecontent, "\pythonexternal{"
    )
line_nr_notebook_file_inclusion =
    ↪ get_line_of_latex_command(
        appendix_filecontent, "\includepdf[pages="
    )
if line_nr_python_file_inclusion > -1:
    appendix_type = "python"
    # get python filename
    line = appendix_filecontent[
        ↪ line_nr_python_file_inclusion]
    filename =
        ↪ get_filename_from_latex_inclusion_command(
            line, ".py", "\pythonexternal{"
        )
    appendices.append(
        Appendix(
            appendix_filepath,
            appendix_filecontent,
            appendix_type,
            filename,
            line,
        )
    )
if line_nr_notebook_file_inclusion > -1:
    appendix_type = "notebook"
    line = appendix_filecontent[
        ↪ line_nr_notebook_file_inclusion]
    filename =
        ↪ get_filename_from_latex_inclusion_command(
            line, ".pdf", "\includepdf[pages="
        )
    appendices.append(
        Appendix(
            appendix_filepath,
            appendix_filecontent,
            appendix_type,
            filename,
            line,
        )
    )
else:
    appendices.append(
        Appendix(appendix_filepath,
        ↪ appendix_filecontent, appendix_type)
    )
return appendices

def get_filename_from_latex_inclusion_command(
    appendix_line, extension, start_substring
):

```

```

""" returns the code/notebook filename in a latex command
    ↳ which includes that code in an appendix.
The inclusion command includes a python code or jupyter
    ↳ notebook pdf.

:param appendix_line: Line of latex code (in particular
    ↳ expected to be the latex code from an appendix.).
:param extension: The file extension that is used/searched
    ↳ in this function. The file extension of the file
    ↳ that is sought in the appendix line. Either ".py" or
    ↳ ".pdf".
:param start_substring: The substring that characterises
    ↳ the latex inclusion command.

"""
start_index = appendix_line.index(start_substring)
end_index = appendix_line.index(extension)
return get_filename_from_dir(
    appendix_line[start_index : end_index + len(extension)
    ↳ ]
)

def get_code_files_already_included_in_appendices(
    absolute_code_filepaths, appendix_dir, extension,
    ↳ normalised_root_dir
):
    """ Returns a list of code filepaths that are already
        ↳ properly included the latex appendix files of this
        ↳ project.

    :param absolute_code_filepaths: List of absolute paths to
        ↳ the code files (either python files or compiled
        ↳ jupyter notebook pdfs).
    :param appendix_dir: Absolute path that contains the
        ↳ appendix .tex files.
    :param extension: The file extension that is used/searched
        ↳ in this function. The file extension of the file
        ↳ that is sought in the appendix line. Either ".py" or
        ↳ ".pdf".
    :param project_name: The name of the project that is being
        ↳ executed/ran. The number indicating which project
        ↳ this code pertains to.
    :param normalised_root_dir: The root directory of this
        ↳ repository.

    """
    appendix_files = get_all_files_in_dir_and_child_dirs(".tex
    ↳ ", appendix_dir)
    contained_codes = []
    for code_filepath in absolute_code_filepaths:
        for appendix_filepath in appendix_files:
            appendix_filecontent = read_file(appendix_filepath
            ↳ )
            line_nr = check_if_appendix_contains_file(

```

```

        appendix_filecontent, code_filepath, extension
        ↪ , normalised_root_dir
    )
    if line_nr > -1:
        # add filepath to list of files that are
        ↪ already in the appendices
        contained_codes.append(
            Appendix_with_code(
                code_filepath,
                appendix_filepath,
                appendix_filecontent,
                line_nr,
                ".py",
            )
        )
    return contained_codes

def check_if_appendix_contains_file(
    appendix_content, code_filepath, extension,
    ↪ normalised_root_dir
):
    """ Scans an appendix content to determine whether it
        ↪ contains a substring that
        includes a code file (of either python or compiled
        ↪ notebook=pdf extension).

    :param appendix_content: content in an appendix latex file
        ↪ .
    :param code_filepath: Absolute path to a code file (either
        ↪ python files or compiled jupyter notebook pdfs).
    :param extension: The file extension that is used/searched
        ↪ in this function. The file extension of the file
        ↪ that is sought in the appendix line. Either ".py" or
        ↪ ".pdf".
    :param project_name: The name of the project that is being
        ↪ executed/ran. The number indicating which project
        ↪ this code pertains to.
    :param normalised_root_dir: The root directory of this
        ↪ repository.

    """
    # convert code_filepath to the inclusion format in latex
    ↪ format

    code_filepath_relative_from_latex_dir = (
        f"latex/../../{code_filepath[len(normalised_root_dir)
        ↪ :]}"
    )
    print(f"code_filepath={code_filepath}")
    print(
        f"code_filepath_relative_from_latex_dir={
        ↪ code_filepath_relative_from_latex_dir}"
    )
    latex_command = get_latex_inclusion_command(
        extension, code_filepath_relative_from_latex_dir
    )

```

```

    )
    return get_line_of_latex_command(appendix_content,
        ↪ latex_command)

def get_line_of_latex_command(appendix_content, latex_command)
    ↪ :
    """ Returns the line number of a latex command if it is
        ↪ found. Returns -1 otherwise.

    :param appendix_content: content in an appendix latex file
        ↪ .
    :param latex_command: A line of latex code. (Expected to
        ↪ come from some appendix)

    """
    # check if the file is in the latex code
    line_nr = 0
    for line in appendix_content:
        if latex_command in line:
            if line_is_commented(line, latex_command):
                commented = True
            else:
                return line_nr
        line_nr = line_nr + 1
    return -1

def line_is_commented(line, target_substring):
    """ Returns True if a latex code line is commented, returns
        ↪ False otherwise

    :param line: A line of latex code that contains a relevant
        ↪ command (target substring).
    :param target_substring: Used to determine whether the
        ↪ command that is found is commented or not.

    """
    left_of_command = line[: line.rfind(target_substring)]
    if "%" in left_of_command:
        return True
    return False

def get_latex_inclusion_command(extension,
    ↪ code_filepath_relative_from_latex_dir):
    """ Creates and returns a latex command that includes
        ↪ either a python file or a compiled jupyter
        notebook pdf (wherever the command is placed). The
        ↪ command is intended to be placed in the appendix.

    :param extension: The file extension that is used/searched
        ↪ in this function. The file extension of the file
        ↪ that is sought in the appendix line. Either ".py" or
        ↪ ".pdf".

```

```

:param code_filepath_relative_from_latex_dir: The latex
    ↳ compilation requires a relative path towards code
    ↳ files
that are included. Therefore, a relative path towards the
    ↳ code is given.

"""
if extension == ".py":
    left = "\pythonexternal{"
    right = "}"
    latex_command = f"{left}{
        ↳ code_filepath_relative_from_latex_dir}{right}"
elif extension == ".ipynb":
    left = "\includepdf[pages=-]{
    right = "}"
    latex_command = f"{left}{
        ↳ code_filepath_relative_from_latex_dir}{right}"
return latex_command

def read_file(filepath):
    """Reads content of a file and returns it as a list of
        ↳ strings, with one string per line.

:param filepath: path towards the file that is being read.

"""
with open(filepath) as f:
    content = f.readlines()
return content

def get_code_files_not_yet_included_in_appendices(
code_filepaths, contained_codes, extension
):
    """Returns a list of filepaths that are not yet properly
        ↳ included in some appendix of this project.

:param code_filepath: Absolute path to all the code files
    ↳ in this project (source directory).
(either python files or compiled jupyter notebook pdfs).
:param contained_codes: list of Appendix objects that
    ↳ include either python files or compiled jupyter
    ↳ notebook pdfs, which
are already included in the appendix tex files. (Does not
    ↳ care whether those appendices are also actually
included in the main or not.)
:param extension: The file extension that is used/searched
    ↳ in this function. The file extension of the file
    ↳ that is sought in the appendix line. Either ".py" or
    ↳ ".pdf".
:param code_filepaths:

"""
contained_filepaths = list(

```



```

        map(lambda contained_file: contained_file.
             ↪ code_filepath, contained_codes)
    )
    not_contained = []
    for filepath in code_filepaths:
        if not filepath in contained_filepaths:
            not_contained.append(filepath)
    return not_contained

def create_appendices_with_code(
    appendix_dir, code_filepaths, extension,
    ↪ normalised_root_dir
):
    """ Creates the latex appendix files in with relevant codes
    ↪ included.

    :param appendix_dir: Absolute path that contains the
    ↪ appendix .tex files.
    :param code_filepaths: Absolute path to code files that
    ↪ are not yet included in an appendix
    (either python files or compiled jupyter notebook pdfs).
    :param extension: The file extension that is used/searched
    ↪ in this function. The file extension of the file
    ↪ that is sought in the appendix line. Either ".py" or
    ↪ ".pdf".
    :param project_name: The name of the project that is being
    ↪ executed/ran. The number indicating which project
    ↪ this code pertains to.
    :param normalised_root_dir: The root directory of this
    ↪ repository.

    """
    appendix_filenames = []
    appendix_reference_index = (
        get_index_of_auto_generated_appendices(appendix_dir,
        ↪ extension) + 1
    )
    print(f"\n\ncode_filepaths={code_filepaths}")
    for code_filepath in code_filepaths:
        normalised_code_filepath = os.path.normpath(
        ↪ code_filepath)
        code_filepath_relative_from_root =
        ↪ normalised_code_filepath[
            len(normalised_root_dir) :
        ]
        code_filepath_relative_from_latex_dir = (
            f"latex/..{code_filepath_relative_from_root}"
        )
        print(f"normalised_code_filepath={
        ↪ normalised_code_filepath}")
        print(f"code_filepath_relative_from_root={
        ↪ code_filepath_relative_from_root}")
        print(
            f"code_filepath_relative_from_latex_dir={
            ↪ code_filepath_relative_from_latex_dir}"

```

```

    )

    # code_filename=get_filename_from_filepath (
    ↪ code_filepath)
    content = []
    filename = get_filename_from_dir(
    ↪ normalised_code_filepath)

    content = create_section(appendix_reference_index,
    ↪ filename, content)
    content = add_include_code_in_appendix(
        content,
        code_filepath_relative_from_root,
        extension,
        code_filepath_relative_from_latex_dir,
        normalised_root_dir,
    )

    print(f"content={content}")

    overwrite_content_to_file(
        content,
        f"{appendix_dir}Auto_generated_{extension[1:]}_App{
        ↪ {appendix_reference_index}.tex",
        False,
    )
    appendix_filenames.append(
        f"Auto_generated_{extension[1:]}_App{
        ↪ appendix_reference_index}.tex"
    )
    appendix_reference_index = appendix_reference_index +
    ↪ 1
    return appendix_filenames

def get_filename_from_filepath(path):
    head, tail = ntpath.split(path)
    return tail or ntpath.basename(head)

def add_include_code_in_appendix(
    content,
    code_filepath_relative_from_root,
    extension,
    code_filepath_relative_from_latex_dir,
    normalised_root_dir,
):
    """Includes the latex code that includes code in the
    ↪ script.

    :param content: The latex content that is being written to
    ↪ an appendix.
    :param code_path_from_latex_main_path: the path to the
    ↪ code as seen from the folder that contains main.tex.
    :param extension: The file extension that is used/searched
    ↪ in this function. The file extension of the file

```

```

    ↪ that is sought in the appendix line. Either ".py" or
    ↪ ".pdf".
:param code_filepath_relative_from_latex_dir: The latex
    ↪ compilation requires a relative path towards code
    ↪ files
that are included. Therefore, a relative path towards the
    ↪ code is given.
:param code_filepath_relative_from_root: param
    ↪ code_filepath_relative_from_latex_dir:
:param project_name: The name of the project that is being
    ↪ executed/ran. param normalised_root_dir:
:param code_filepath_relative_from_latex_dir: param
    ↪ normalised_root_dir:
:param normalised_root_dir:

"""
print(f"before={content}")

# TODO: append if exists}
content.append("%TESTCOMMENT")
latex_path_to_python_file = "src/filename"
print(f"code_filepath_relative_from_root={
    ↪ code_filepath_relative_from_root}")
line = f"\IfFileExists{{{
    ↪ code_filepath_relative_from_latex_dir}}}{{"
print(f"line={line}")
content.append(line)
# append current line
content.append(
    get_latex_inclusion_command(extension,
    ↪ code_filepath_relative_from_latex_dir)
)
# TODO: append {}
content.append(f"}}{{")
# TODO: code_path_from latex line
content.append(
    get_latex_inclusion_command(extension,
    ↪ code_filepath_relative_from_latex_dir)
)
# TODO: add closing bracket }
content.append(f"}}")
content.append("%TESTCOMMENTCLOSING")
print(f"after={content}")
return content

def get_index_of_auto_generated_appendices(appendix_dir,
    ↪ extension):
    """Returns the maximum index of auto generated appendices
    ↪ of
    a specific extension type.

:param extension: The file extension that is used/searched
    ↪ in this function. The file extension of the file
    ↪ that is sought in the appendix line. Either ".py" or
    ↪ ".pdf".

```

```

:param appendix_dir: Absolute path that contains the
    ↳ appendix .tex files.

"""
max_index = -1
appendices =
    ↳ get_auto_generated_appendix_filenames_of_specific_extension
    ↳ (
        appendix_dir, extension
    )
for appendix in appendices:
    substring = f"Auto_generated_{extension[1:]}_App"
    # remove left of index
    remainder = appendix[appendix.rfind(substring) + len(
        ↳ substring) :]
    # remove right of index
    index = int(remainder[:-4])
    if index > max_index:
        max_index = index
return max_index

def
    ↳ get_auto_generated_appendix_filenames_of_specific_extension
    ↳ (
        appendix_dir, extension
    ):
    """Returns the list of auto generated appendices of
    a specific extension type.

    :param extension: The file extension that is used/searched
        ↳ in this function. The file extension of the file
        ↳ that is sought in the appendix line. Either ".py" or
        ↳ ".pdf".
    :param appendix_dir: Absolute path that contains the
        ↳ appendix .tex files.

    """
    appendices_of_extension_type = []

    # get all appendices
    appendix_files = get_all_files_in_dir_and_child_dirs(".tex
        ↳ ", appendix_dir)

    # get appendices of particular extension type
    for appendix_filepath in appendix_files:
        right_of_slash = appendix_filepath[appendix_filepath.
            ↳ rfind("/") + 1 :]
        if (
            right_of_slash[: 15 + len(extension) - 1]
            == f"Auto_generated_{extension[1:]}_App"
        ):
            appendices_of_extension_type.append(
                ↳ appendix_filepath)
    return appendices_of_extension_type

```

```

def create_section(appendix_reference_index, code_filename,
    ↪ content):
    """ Creates the header of a latex appendix file , such that
        ↪ it contains a section that
        indicates the section is an appendix , and indicates which
        ↪ python or notebook file is
        being included in that appendix.

    :param appendix_reference_index: A counter that is used in
        ↪ the label to ensure the appendix section labels are
        ↪ unique.
    :param code_filename: file name of the code file that is
        ↪ included
    :param content: A list of strings that make up the
        ↪ appendix , with one line per element.

    """
    # write section
    left = "\section{Appendix "
    middle = code_filename.replace("_", "\_")
    right = "}\label{app:"
    end = "}" # TODO: update appendix reference index
    content.append(f"{left}{middle}{right}{"
        ↪ appendix_reference_index}{end}")
    return content


def overwrite_content_to_file(content, filepath,
    ↪ content_has_newlines=True):
    """ Writes a list of lines of tex code from the content
        ↪ argument to a .tex file
        using overwriting method. The content has one line per
        ↪ element.

    :param content: The content that is being written to file.
    :param filepath: Path towards the file that is being read.
    :param content_has_newlines: Default value = True)

    """
    with open(filepath, "w") as f:
        for line in content:
            if content_has_newlines:
                f.write(line)
            else:
                f.write(line + "\n")


def get_appendix_tex_code(main_latex_filename):
    """ gets the latex appendix code from the main tex file .

    :param main_latex_filename: Name of the main latex
        ↪ document of this project number

    """
    main_tex_code = read_file(main_latex_filename)

```

```

# print(f"main_tex_code={main_tex_code}")
start = "\\begin{appendices}"
end = "\\end{appendices}"
# TODO: if last 4 characters before \\end{appendices} match
#       }}\\fi then don't prepend it,
# otherwise, do prepend it
start_index = get_index_of_substring_in_list(main_tex_code
#       , start) + 1
end_index = get_index_of_substring_in_list(main_tex_code,
#       end)
return main_tex_code, start_index, end_index,
#       main_tex_code[start_index:end_index]

def get_index_of_substring_in_list(lines, target_substring):
    """Returns the index of the line in which the first
    character of a latex substring if it is found
    uncommented in the incoming list.

    :param lines: List of lines of latex code.
    :param target_substring: Some latex command/code that is
    sought in the incoming text.

    """
    for i in range(0, len(lines)):
        if target_substring in lines[i]:
            if not line_is_commented(lines[i],
#               target_substring):
                return i

def update_appendix_tex_code(appendix_filename,
# is_from_normalised_root_dir):
    """Returns the latex command that includes an appendix .
    text file in an appendix environment
    as can be used in the main tex file.

    :param appendix_filename: Name of the appendix that is
    included by the generated command.
    :param project_name: The name of the project that is being
    executed/ran. The number indicating which project
    this code pertains to.
    :param is_from_normalised_root_dir:

    """
    if is_from_normalised_root_dir:
        left = f"\\input{{latex/"
    else:
        left = "\\input{"
    middle = "Appendices/"
    right = "} \\newpage\\n"
    return f"{left}{middle}{appendix_filename}{right}"

def substitute_appendix_code(

```

```

    end_index, main_tex_code, start_index,
    ↪ updated_appendices_tex_code
):
    """ Replaces the old latex code that included the
    ↪ appendices in the main.tex file with the new latex
    commands that include the appendices in the latex report.

    :param end_index: Index at which the appendix section ends
    ↪ right before the latex \end{appendix} line ,
    :param main_tex_code: The code that is saved in the main .
    ↪ tex file .
    :param start_index: Index at which the appendix section
    ↪ starts right after the latex \begin{appendix} line ,
    :param updated_appendices_tex_code: The newly created code
    ↪ that includes all the relevant appendices .
    (relevant being (in order): manually created appendices ,
    ↪ python codes , pdfs of compiled jupyter notebooks).

    """
    start_of_main_tex_code_till_appendices = main_tex_code[0:
    ↪ start_index]
    tex_code_after_appendices =
    ↪ inject_closing_hide_source_conditional_close(
        main_tex_code[end_index:]
    )
    updated_main_tex_code = (
        start_of_main_tex_code_till_appendices
        + updated_appendices_tex_code
        + tex_code_after_appendices
    )
    print(f"start_index={start_index}")
    print(f"main_tex_code[end_index:] = {main_tex_code[
    ↪ end_index:]}")
    return updated_main_tex_code

def inject_closing_hide_source_conditional_close(
    ↪ tex_code_after_appendices):
    injected_string_to_close_conditional = "}}\\fi"

    # Get the line directly after the (autogenerated)
    ↪ appendices .
    line_closing_appendices = tex_code_after_appendices[0]
    # Get the first characters of that closing line to see if
    ↪ they match the
    # injected string that needs to be in.
    start_of_close = line_closing_appendices[
        : len(injected_string_to_close_conditional)
    ]

    # If the first 4 characters do not start with the close of
    ↪ the hide source
    # code conditional , then inject that close . Othwerise
    ↪ return tex code as is .
    if start_of_close == f"{
    ↪ injected_string_to_close_conditional}":

```

```

        return tex_code_after_appendices
    else:
        # Inject the string that closes the hideshowcode
        ↪ conditional.
        tex_code_after_appendices[
            0
        ] = f"{injected_string_to_close_conditional}{"
        ↪ tex_code_after_appendices[0]}"
        return tex_code_after_appendices

def get_filename_from_dir(path):
    """Returns a filename from an absolute path to a file.

    :param path: path to a file of which the name is queried.

    """
    return path[path.rfind("/") + 1 :]

def get_script_dir():
    """returns the directory of this script regardless of from
    ↪ which level the code is executed"""
    return os.path.dirname(__file__)

class Appendix_with_code:
    """stores in which appendix file and accompanying line
    ↪ number in the appendix in which a code file is
    already included. Does not take into account whether this
    ↪ appendix is in the main tex file or not

    """

    def __init__(
        self,
        code_filepath,
        appendix_filepath,
        appendix_content,
        file_line_nr,
        extension,
    ):
        self.code_filepath = code_filepath
        self.appendix_filepath = appendix_filepath
        self.appendix_content = appendix_content
        self.file_line_nr = file_line_nr
        self.extension = extension

class Appendix:
    """stores in appendix files and type of appendix."""

    def __init__(
        self,
        appendix_filepath,
    ):

```



```
        appendix_content ,
        appendix_type ,
        code_filename=None,
        appendix_inclusion_line=None,
    ):
        self.appendix_filepath = appendix_filepath
        self.appendix_filename = get_filename_from_dir(self.
            ↳ appendix_filepath)
        self.appendix_content = appendix_content
        self.appendix_type = appendix_type # TODO: perform
            ↳ validation of input values
        self.code_filename = code_filename
        self.appendix_inclusion_line = appendix_inclusion_line
```

.15. Appendix neumann.py

```

# This code computes a minimum total dominating set
  → approximation.
import random

def compute_mtds(G, m=1):

    # No directed graphs supported in this algorithm.
    if G.is_directed():
        raise Exception("Parameter graph should be undirected"
            → )

    # 1.a Each vertex v_i chooses random float r_i in range 0<
    → r_i < 1
    for node in G.nodes:
        # print(f'label={G.nodes[node]["label"]}')
        G.nodes[node]["rand_val"] = random.uniform(0, 1)
        print(f'rand_val={G.nodes[node]["rand_val"]}')
    # 1.b Each vertex v_i computes d_i. d_i=degree of vertex
    → v_i
    for node in G.nodes:
        G.nodes[node]["degree"] = G.degree(node)
        print(f'degree={G.nodes[node]["degree"]}')
    # 1.c Each vertex v_i computes weight: w_i=d_i+r_i
    for node in G.nodes:
        G.nodes[node]["weight"] = G.nodes[node]["degree"] + G.
            → nodes[node]["rand_val"]
        print(f'weight={G.nodes[node]["weight"]}')

    # 1.d Each vertex v_i sends w_i to each of its neighbours.
    # TODO: paradigm: Either each neuron sends some value to
    → another neuron.
    # OR: each neuron asks/demands some value of each of its
    → neighbours.
    store_index_max_neighbour_weight(G)

    # 2.a Each vertex v_i gets the index of the
    # neighbouring vertex v_j_ (w_max) that has the heighest
    → w_i, with i!=j.
    # 2.b Each vertex v_i adds a mark to that neighbour vertex
    → v_j_ (w_max).
    for node in G.nodes:
        G.nodes[node]["mark"] = 0
    for node in G.nodes:
        neighbor_with_max_weight = G.nodes[node]["
            → neighbor_with_max_weight"]
        G.nodes[neighbor_with_max_weight]["mark"] = (
            G.nodes[neighbor_with_max_weight]["mark"] + 1
        )

    # 3 for k in range [0,m] rounds, do:
    for _ in range(0, m):

```

```

# 4.a Each node v_i computes how many marks it has
#     ↳ received, as (x_i)_k.
# 4.b Each node v_i computes (w_i)_k=(x_i)_k+ r_i
for node in G.nodes:
    G.nodes[node]["weight"] = G.nodes[node]["mark"] +
    #     ↳ G.nodes[node]["rand_val"]
    print(f'weight={G.nodes[node]["weight"]}')

# 5. Reset marked vertices: for each vertex v_i, (x_i)
#     ↳ _k=0

# 6.a Each vertex v_i computes d_i. d_i=degree of
#     ↳ vertex v_i
# 6.b Each vertex v_i sends w_i to each of its
#     ↳ neighbours.
# 6.c Each vertex v_i gets the index of the
#     neighbouring vertex v_j_ (w_max) that has the
#     ↳ heighest w_i, with i!=j.
# 6.d Each vertex v_i adds a mark to that neighbour
#     ↳ vertex v_j_ (w_max).

def store_index_max_neighbour_weight(G):
    for node in G.nodes:
        G.nodes[node]["neighbor_with_max_weight"] = None
        max = 0 # Assumes positive weights
        for neighbor in G.neighbors(node):
            if G.nodes[node]["neighbor_with_max_weight"] is
            #     ↳ None:
                G.nodes[node][
                    "neighbor_with_max_weight"
                ] = neighbor # G.nodes[neighbor]
                max = G.nodes[neighbor]["weight"]
            elif G.nodes[neighbor]["weight"] > max:
                G.nodes[node][
                    "neighbor_with_max_weight"
                ] = neighbor # G.nodes[neighbor]
                max = G.nodes[neighbor]["weight"]
        print(
            f'node={node}G.nodes[node][
            #     ↳ neighbor_with_max_weight]={G.nodes[node][
            #     ↳ neighbor_with_max_weight}]'
        )
    return G

```

.16. Appendix `neumann_a_t_0.py`

```

# This code computes a minimum total dominating set
  → approximation.
import random

def compute_mtds_a_t_0(G, m=2):

    # No directed graphs supported in this algorithm.
    if G.is_directed():
        raise Exception("Parameter graph should be undirected"
            → )

    # 1.a Each vertex v_i chooses random float r_i in range 0<
    → r_i < 1
    for node in G.nodes:
        # print(f' label={G.nodes[node]["label"]} ')
        G.nodes[node]["rand_val"] = random.uniform(0, 1)
        print(f'rand_val={G.nodes[node]["rand_val"]} ')
    # 1.b Each vertex v_i computes d_i. d_i=degree of vertex
    → v_i
    for node in G.nodes:
        G.nodes[node]["degree"] = G.degree(node)
        print(f'degree={G.nodes[node]["degree"]} ')
    # 1.c Each vertex v_i computes weight: w_i=d_i+r_i
    for node in G.nodes:
        G.nodes[node]["weight"] = G.nodes[node]["degree"] + G.
            → nodes[node]["rand_val"]
        print(f'weight={G.nodes[node]["weight"]} ')

    # 1.d Each vertex v_i sends w_i to each of its neighbours.
    # TODO: paradigm: Either each neuron sends some value to
    → another neuron.
    # OR: each neuron asks/demands some value of each of its
    → neighbours.
    store_index_max_neighbour_weight(G)

    # 2.a Each vertex v_i gets the index of the
    # neighbouring vertex v_j_ (w_max) that has the heighest
    → w_i, with i!=j.
    # 2.b Initialise the mark counter x_i
    for node in G.nodes:
        G.nodes[node]["mark"] = 0
    # 2.c Each vertex v_i adds a mark to that neighbour vertex
    → v_j_ (w_max).
    for node in G.nodes:
        neighbor_with_max_weight = G.nodes[node]["
            → neighbor_with_max_weight"]
        G.nodes[neighbor_with_max_weight]["mark"] = (
            G.nodes[neighbor_with_max_weight]["mark"] + 1
        )

    # 3 for k in range [0,m] rounds, do:
    for _ in range(0, m):

```

```

# 4.a Each node v_i computes how many marks it has
#     ↳ received, as (x_i)_k.
# 4.b Each node v_i computes (w_i)_k=(x_i)_k+ r_i
for node in G.nodes:
    G.nodes[node]["weight"] = G.nodes[node]["mark"] +
    #     ↳ G.nodes[node]["rand_val"]
    print(f'weight={G.nodes[node]["weight"]}')
```

5. Reset marked vertices: for each vertex v_i, (x_i)

```

#     ↳ _k=0
for node in G.nodes:
    G.nodes[node]["mark"] = 0
```

6.a Each vertex v_i computes d_i. d_i=degree of

```

#     ↳ vertex v_i
# 6.b Each vertex v_i sends w_i to each of its
#     ↳ neighbours.
# 6.c Each vertex v_i gets the index of the
#     neighbouring vertex v_j_ (w_max) that has the
#     ↳ heighest w_i, with i!=j.
store_index_max_neighbour_weight(G)
# 6.d Each vertex v_i adds a mark to that neighbour
#     ↳ vertex v_j_ (w_max).
for node in G.nodes:
    neighbor_with_max_weight = G.nodes[node]["
    #     ↳ neighbor_with_max_weight"]
    G.nodes[neighbor_with_max_weight]["mark"] = (
        G.nodes[neighbor_with_max_weight]["mark"] + 1
    )
```

Print Minimum Total Dominating Set approximation.

```

mtds = []
for node in G.nodes:
    print(f'node {node} has {G.nodes[node]["mark"]}
    #     ↳ marks')
    if 0 < G.nodes[node]["mark"]:
        mtds.append(node)
print(f"mtds={mtds}")
```

```

def store_index_max_neighbour_weight(G):
    for node in G.nodes:
        G.nodes[node]["neighbor_with_max_weight"] = None
        max = 0 # Assumes positive weights
        for neighbor in G.neighbors(node):
            if G.nodes[node]["neighbor_with_max_weight"] is
            #     ↳ None:
                G.nodes[node][
                    "neighbor_with_max_weight"
                ] = neighbor # G.nodes[neighbor]
                max = G.nodes[neighbor]["weight"]
            elif G.nodes[neighbor]["weight"] > max:
                G.nodes[node][
                    "neighbor_with_max_weight"
                ] = neighbor # G.nodes[neighbor]
                max = G.nodes[neighbor]["weight"]
```

```
    print(  
        f'node={node}G.nodes[node]["  
            ↳ neighbor_with_max_weight"]={G.nodes[node]["  
            ↳ neighbor_with_max_weight"]}'  
    )  
    return G
```

.17. Appendix plantuml_compile.py

```

# This script automatically compiles the text files
    ↳ representing a PlantUML
# diagram into an actual figure.

# To compile locally manually:
# pip install plantuml
# export PLANTUML_LIMIT_SIZE=8192
# java -jar plantuml.jar -verbose sequenceDiagram.txt

import os
import subprocess
from os.path import abspath

from .helper_dir_file_edit import
    ↳ get_dir_filelist_based_on_extension
from .plantuml_get_package import got_java_file

def compile_diagrams_in_dir_relative_to_root(
    await_compilation,
    extension,
    jar_path_relative_from_root,
    input_dir_relative_to_root,
    verbose,
):
    """
    Loops through the files in a directory and exports them to
        ↳ the latex /Images
    directory.

    Args:
    :param await_compilation: Make python wait untill the
        ↳ PlantUML compilation is completed. param extension:
        ↳ The filetype of the text file that is converted to
        ↳ image.
    :param jar_path_relative_from_root: The path as seen from
        ↳ root towards the PlantUML .jar file that compiles .
        ↳ uml files to .png files.
    :param verbose: True, ensures compilation output is
        ↳ printed to terminal, False means compilation is
        ↳ silent.
    :param extension: The file extension that is used/searched
        ↳ in this function.
    :param input_dir_relative_to_root: The directory as seen
        ↳ from root containing files that are modified in this
        ↳ function.

    Returns:
        Nothing

    Raises:
        Nothing
    """
    # Verify the PlantUML .jar file is gotten.

```

```

got_java_file(jar_path_relative_from_root)

diagram_text_filenames =
    ↪ get_dir_filelist_based_on_extension(
        input_dir_relative_to_root, extension
    )

for diagram_text_filename in diagram_text_filenames:
    diagram_text_filepath_relative_from_root = (
        ↪ f"{input_dir_relative_to_root}/{
            ↪ diagram_text_filename}"
    )

    execute_diagram_compilation_command(
        await_compilation,
        jar_path_relative_from_root,
        diagram_text_filepath_relative_from_root,
        verbose,
    )

def execute_diagram_compilation_command(
    await_compilation,
    jar_path_relative_from_root,
    relative_filepath_from_root,
    verbose,
):
    """
    Compiles a .uml/text file containing a PlantUML diagram to
    ↪ a .png image
    using the PlantUML .jar file.

    Args:
    :param await_compilation: Make python wait untill the
        ↪ PlantUML compilation is completed. param
        ↪ jar_path_relative_from_root:
    :param relative_filepath_from_root: Relative filepath as
        ↪ seen from root of file that is used in this function.
    :param jar_path_relative_from_root: The path as seen from
        ↪ root towards the PlantUML .jar file that compiles .
        ↪ uml files to .png files.
    :param verbose: True, ensures compilation output is
        ↪ printed to terminal, False means compilation is
        ↪ silent.

    Returns:
        Nothing

    Raises:
        Nothing
    """
    # Verify the files required for compilation exist, and
    ↪ convert the paths
    # into absolute filepaths.
    abs_diagram_filepath, abs_jar_path =
    ↪ assert_diagram_compilation_requirements(

```



```

        jar_path_relative_from_root,
        ↪ relative_filepath_from_root
    )

    # Generate command to compile the PlantUML diagram locally
    ↪ .
    print(
        f"abs_jar_path={abs_jar_path}, abs_diagram_filepath={
        ↪ abs_diagram_filepath}\n\n"
    )
    bash_diagram_compilation_command = (
        f"java -jar {abs_jar_path} -verbose {
        ↪ abs_diagram_filepath}"
    )
    print(f"bash_diagram_compilation_command={
    ↪ bash_diagram_compilation_command}")
    # Generate global variable specifying max image width in
    ↪ pixels , in the
    # shell that compiles.
    os.environ["PLANTUML_LIMIT_SIZE"] = "16192"

    # Perform PlantUML compilation locally.
    if await_compilation:
        if verbose:
            subprocess.call(bash_diagram_compilation_command,
            ↪ shell=True)
        else:
            subprocess.call(
                bash_diagram_compilation_command,
                shell=True,
                stderr=subprocess.DEVNULL,
                stdout=subprocess.DEVNULL,
            )
    else:
        if verbose:
            subprocess.Popen(bash_diagram_compilation_command,
            ↪ shell=True)
        else:
            subprocess.Popen(
                bash_diagram_compilation_command,
                shell=True,
                stderr=subprocess.DEVNULL,
                stdout=subprocess.DEVNULL,
            )

def assert_diagram_compilation_requirements(
    jar_path_relative_from_root, relative_filepath_from_root,
):
    """
    Asserts that the PlantUML .jar file used for compilation
    ↪ exists , and that
    the diagram file with the .uml content for the diagram
    ↪ exists. Throws an
    error if either of two is missing.

```

```

:param relative_filepath_from_root: Relative filepath as
    ↳ seen from root of file that is used in this function.
:param output_dir_from_root: Relative directory as seen
    ↳ from root, to which files are outputted.
:param jar_path_relative_from_root: The path as seen from
    ↳ root towards the PlantUML .jar file that compiles .
    ↳ uml files to .png files.

```

```

Returns:
    Nothing

```

```

Raises:
    Exception if PlantUML .jar file used to compile the .
    ↳ uml to .png files
    is missing.
    Exception if the file with the .uml content is missing
    ↳ .
"""
abs_diagram_filepath = abspath(relative_filepath_from_root
    ↳ )
abs_jar_path = abspath(jar_path_relative_from_root)
if os.path.isfile(abs_diagram_filepath):
    if os.path.isfile(abs_jar_path):
        return abs_diagram_filepath, abs_jar_path
    else:
        raise Exception(
            f"The input diagram file:{abs_diagram_filepath}
            ↳ } does not exist."
        )
else:
    raise Exception(f"The input jar file:{abs_jar_path}
    ↳ does not exist.")

```

.18. Appendix plantuml_generate.py

```

# This script generates PlantUML diagrams and outputs them as
→ .uml files .

import os
import subprocess
from os.path import abspath

from .helper_dir_file_edit import
→ create_dir_relative_to_root_if_not_exists
from .helper_dir_file_edit import dir_relative_to_root_exists

def generate_all_dynamic_diagrams(output_dir_relative_to_root)
→ :
    """
    Manages the generation of all the diagrams created in this
    → file .

    Args :
    :param output_dir_relative_to_root: Relative path as seen
    → from the root dir of this project , to which modified
    → files are outputted .

    Returns :
        Nothing

    Raises :
    """
    # Create a example Gantt output file .
    filename_one, lines_one = create_trivial_gantt("
    → trivial_gantt.uml")
    output_diagram_text_file(filename_one, lines_one,
    → output_dir_relative_to_root)

    # Create another example Gantt output file .
    filename_two, lines_two = create_trivial_gantt("
    → another_trivial_gantt.uml")
    output_diagram_text_file(
        filename_two, lines_two, output_dir_relative_to_root,
    )

def output_diagram_text_file(filename, lines,
→ output_dir_relative_to_root):
    """
    Gets the filename and lines of an PlantUML diagram , and
    → writes these to a
    file at the relative output path .

    Args :
    :param filename: The filename of the PlantUML Gantt file
    → that is being created .
    :param lines: The lines of the Gantt chart PlantUML code
    → that is being written to file .

```

```
:param output_dir_relative_to_root: Relative path as seen
    ↳ from the root dir of this project, to which modified
    ↳ files are outputted.
```

```
Returns:
    Nothing
```

```
Raises:
    Exception if input file does not exist.
"""
```

```
abs_filepath = abspath(f"{output_dir_relative_to_root}/{
    ↳ filename}")

# Ensure output directory is created.
create_dir_relative_to_root_if_not_exists(
    ↳ output_dir_relative_to_root)
if not dir_relative_to_root_exists(
    ↳ output_dir_relative_to_root):
    raise Exception(
        f"Error, the output directory relative to root:{
            ↳ output_dir_relative_to_root} does not exist."
    )

# Delete output file if it already exists.
if os.path.exists(abs_filepath):
    os.remove(abs_filepath)

# Write lines to file.
f = open(abs_filepath, "w")
for line in lines:
    f.write(line)
f.close()

# Assert output file exists.
if not os.path.isfile(abs_filepath):
    raise Exception(f"The input file:{abs_filepath} does
        ↳ not exist.")
```

```
def create_trivial_gantt(filename):
    """
    Creates a trivial Gantt chart.

    Args:
    :param filename: The filename of the PlantUML diagram file
        ↳ that is being
        created.

    Returns:
        The filename of the PlantUML diagram, and the lines of
        ↳ the uml content
        of the diagram

    Raises:
        Nothing
    """
```

```

lines = []
lines.append("@startuml\n")
lines.append("[Prototype design] lasts 15 days\n")
lines.append("[Test prototype] lasts 10 days\n")
lines.append("\n")
lines.append("Project starts 2020-07-01\n")
lines.append("[Prototype design] starts 2020-07-01\n")
lines.append("[Test prototype] starts 2020-07-16\n")
lines.append("@enduml\n")
return filename, lines

```

```

def create_another_trivial_gantt(filename):
    """
    Creates a trivial Gantt chart.

    :param filename: The filename of the PlantUML Gantt file
        ↳ that is being created.

    Returns:
        The filename of the PlantUML diagram, and the lines of
        ↳ the uml content
        of the diagram

    Raises:
        Nothing
    """
    lines = []
    lines.append("@startuml\n")
    lines.append("[EXAMPLE SENTENCE] lasts 15 days\n")
    lines.append("[Test prototype] lasts 10 days\n")
    lines.append("\n")
    lines.append("Project starts 2022-07-01\n")
    lines.append("[Prototype design] starts 2022-07-01\n")
    lines.append("[Test prototype] starts 2022-07-16\n")
    lines.append("@enduml\n")

    return filename, lines

```

.19. Appendix plantuml_get_package.py

```

import os
import subprocess

def check_if_java_file_exists(relative_filepath):
    """
    Safe check to see if file exists or not.

    Args:
    :param relative_filepath: Path as seen from root towards a
        ↪ file.

    Returns:
        True if a file exists.
        False if a file does not exists.

    Raises:
        Nothing
    """
    if os.path.isfile(relative_filepath):
        return True
    else:
        return False

def got_java_file(relative_filepath):
    """
    Asserts if PlantUML .jar file exists. Tries to download is
        ↪ one time if it
    does not exist at the start of the function.

    Args:
    :param relative_filepath: Path as seen from root towards a
        ↪ file.

    Returns:
        True if a file exists.

    Raises:
        Exception if the PlantUML .jar file does not exist
        ↪ after downloading
        it.
    """
    # Check if the jar file exists, curl it if not.
    if not check_if_java_file_exists(relative_filepath):
        # The java file is not found, curl it
        request_file(
            "https://sourceforge.net/projects/plantuml/files/
            ↪ plantuml.jar/download",
            relative_filepath,
        )
    # Check if the jar file exists after curling it. Raise
        ↪ Exception if it is not found after curling.
    if not check_if_java_file_exists(relative_filepath):

```

```

        raise Exception(f"File:{relative_filepath} is not
            ↳ accessible")
    print(f"Got the Java file")
    return True

```

```

def request_file(url, output_filepath):
    """
    Downloads a file or file content.

    Args:
    :param url: Url towards a file that will be downloaded.
    :param relative_filepath: The path as seen from the root
        ↳ of this directory, in which files are outputted.

    Returns:
        Nothing

    Raises:
        Nothing
    """
    import requests

    # Request the file in the url
    response = requests.get(url)
    with open(output_filepath, "wb") as f:
        f.write(response.content)

def run_bash_command(bashCommand):
    """
    Unused method. TODO: verify it is unused and delete it.
    :param bashCommand: A string containing a bash command
        ↳ that can be executed.

    """
    # Verbose call.
    # subprocess.Popen(bashCommand, shell=True)
    # Silent call.
    # subprocess.Popen(bashCommand, shell=True, stderr=
        ↳ subprocess.DEVNULL, stdout=subprocess.DEVNULL)

    # Await completion:
    # Verbose call.
    subprocess.call(bashCommand, shell=True)
    # Silent call.
    # subprocess.call(bashCommand, shell=True, stderr=
        ↳ subprocess.DEVNULL, stdout=subprocess.DEVNULL)

```

.20. Appendix plantuml_to_tex.py

```

from posixpath import abspath
from shutil import copyfile
import shutil
import os.path

from .helper_dir_file_edit import
    ↪ get_dir_filelist_based_on_extension
from .helper_dir_file_edit import
    ↪ create_dir_relative_to_root_if_not_exists

def export_diagrams_to_latex(
    input_dir_relative_to_root, extension,
    ↪ output_dir_relative_to_root
):
    """Loops through the files in a directory and exports them
    ↪ to the latex /Images
    directory.

    :param dir: The directory in which the Gantt charts are
    ↪ being searched.
    :param extension: The file extension that is used/searched
    ↪ in this function. The filetypes that are being
    ↪ exported.
    :param input_dir_relative_to_root: Relative path as seen
    ↪ from the root dir of this project, containing files
    ↪ that modified in this function.
    :param output_dir_relative_to_root: Relative path as seen
    ↪ from the root dir of this project, to which modified
    ↪ files are outputted.

    """
    diagram_filenames = get_dir_filelist_based_on_extension(
        input_dir_relative_to_root, extension
    )
    if len(diagram_filenames) > 0:
        # Ensure output directory is created.
        create_dir_relative_to_root_if_not_exists(
            ↪ output_dir_relative_to_root)

        for diagram_filename in diagram_filenames:
            diagram_filepath_relative_from_root = (
                f"{input_dir_relative_to_root}/{diagram_filename}"
            )

            export_gantt_to_latex(
                diagram_filepath_relative_from_root,
                ↪ output_dir_relative_to_root
            )

def export_gantt_to_latex(relative_filepath_from_root,
    ↪ output_dir_relative_to_root):
    """

```


Takes an input filepath and an output directory as input
 ↳ and copies the
 file towards the output directory.

```
:param relative_filepath_from_root: param
  ↳ output_dir_relative_to_root:
:param output_dir_relative_to_root: Relative path as seen
  ↳ from the root dir of this project, to which modified
  ↳ files are outputted.
```

Returns:
 Nothing.

Raises:
 Exception if the output directory does not exist.
 Exception if the input file is not found.

```
"""
if os.path.isfile(relative_filepath_from_root):
    if os.path.isdir(output_dir_relative_to_root):
        shutil.copy(relative_filepath_from_root,
                    ↳ output_dir_relative_to_root)
    else:
        raise Exception(
            f"The output directory:{
                ↳ output_dir_relative_to_root} does not
                ↳ exist."
        )
else:
    raise Exception(f"The input file:{
        ↳ relative_filepath_from_root} does not exist.")
```
