## GERARDIUM RUSH - MINERAL CIRCUIT OPTIMIZER v1.0.0

Delivered by Pentlandite

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### **Chapter 1**

# GERARDIUM RUSH - MINERAL CIRCUIT OPTIMIZER

This is an implementation for the ACS - Gerardium Rush project delivered by group Pentlandite.

This project is designed to optimize circuit configurations using genetic algorithms to enhance the recovery of valuable materials in separation technologies. The goal is to create an efficient and effective tool for optimizing circuits in minerals processing, particularly for the recovery of "gerardium". This project combines the power of genetic algorithms with mass balance simulations to find the optimal circuit configuration.

### 1.0.1 Overview

The project is organized into several key components, each with specific responsibilities:

### 1. Circuit Simulation (CSimulator.cpp):

- · Core simulation logic for evaluating circuit performance
- Simulates the mass balance for given circuit configurations.
- Utilizes classes like CCircuit and CUnit to represent and manage circuit units and their interactions.

### 2. Genetic Algorithm (Genetic\_Algorithm.cpp):

- Implements the genetic algorithm for optimizing circuit configurations.
- Includes various operators for selection, crossover, and mutation.
- Parallel processing enabled using OpenMP for efficient computation.

### 1.0.2 Features

- **Genetic Algorithm Optimization**: Efficiently explore numerous circuit configurations using genetic algorithms to find the most effective setups.
- Mass Balance Simulation: Simulate the mass balance for each circuit to ensure accurate recovery and grade calculations, ensuring the reliability of the proposed solutions.
- Parallel Processing: Leverage OpenMP for faster computations and simulations, enhancing performance and reducing computation time.
- Comprehensive Unit Tests: Ensure the correctness of the circuit simulator and genetic algorithm through extensive unit tests, providing a robust and reliable tool.
- **Visualization Tools**: Utilize Python scripts and Jupyter notebooks for post-processing and visualizing the results, helping to analyze and interpret the outcomes effectively.
- **Documentation**: Detailed API documentation and project reports to facilitate understanding and usage, making it easier for users to get started and for developers to extend the project.

### 1.0.3 Requirements

The project uses CMake for building and requires several tools and libraries. Below are the required tools and libraries:

Tool/Library	Purpose	Version
CMake	For managing the build process	3.10 or higher
GCC and G++	For compiling the code	13.0 or higher
OpenMP	For parallel processing	4.5 or higher
Python	For post-processing and visualization	3.10 or higher
Graphviz	For generating diagrams	2.50.0 or higher
Matplotlib	For plotting and visualizing data	3.8.2 or higher
Pandas	For data manipulation and analysis	2.1.3 or higher

To install these requirements on a Debian-based system, you can use the following commands:

```
$ sudo apt update
$ sudo apt install cmake gcc g++ libomp-dev python3 python3-pip graphviz
$ pip3 install matplotlib pandas
```

### 1.0.4 Installation

To build the project, you need to have CMake and GCC installed on your system.

1. Clone the repository:

```
$ git clone https://github.com/yourusername/circuit-optimizer.git
```

2. Create a build directory and navigate to it:

```
$ mkdir build
$ cd build
```

3. Configure the project using CMake:

```
$ cmake .. -DCMAKE_C_COMPILER=gcc -DCMAKE_CXX_COMPILER=g++
```

4. Build the project:

```
$ cmake --build .
```

### 1.0.5 Usage

### Run Executable

To run the Circuit Optimizer, execute the following commands from the build directory:

```
$ export OMP_NUM_THREADS=10
$ ./bin/Circuit_Optimizer
```

### **Run Tests**

The project includes a set of unit tests to verify the functionality of the circuit simulator and the genetic algorithm. The tests are located in the tests directory.

To run the tests, use the following command:

```
$ ctest --output-on-failure
```

### **Run Visualization**

To run the visualization tool, use the following command:

```
$ python3 ../post_process/visualize.py <vector> <performance> <recovery> <grade>
```

### **Run Scripts (Recommended)**

Alternatively, you can use the provided shell scripts:

- To run the optimizer, go to the root directory and run the following command:
   \$ ./scripts/run.sh
- To run the tests, go to the root directory and run the following command:

```
$ ./scripts/run_test.sh
```

You can modify the OMP\_NUM\_THREADS in corresponding shell scripts.

### 1.0.6 Parameter Setting

In src/main.cpp, you can set the Genetic Algorithm parameters and Circuit Simulator parameters. Here is an example:

```
int generations = 1000;
int populationSize = 100;
std::string selection = "NaryTournamentSelection";
// std::string selection = "RankSelection";
// std::string selection = "RouletteWheelSelection";
int tournamentSize = 5;
std::string crossover = "TwoPointCrossover";
// std::string crossover = "UniformCrossover";
// std::string crossover = "PureSinglePointCrossover";
double crossRate = 0.8;
std::string mutation = "GuidedMutation";
// std::string mutation = "GenProgMutation";
double mutationRate = 0.1;
double eliteRate = 0.2;
double initialTemp = 1000.0;
double deltT = 0.85;
unsigned int randomSeed = 0;
double tolerance = 1e-6;
int maxIteration = 1000;
CircuitParameters c_params(tolerance, maxIteration);
```

### **Genetic Algorithm Operators**

#### · Selection:

- NaryTournamentSelection: Selects the best individuals based on tournament competition.
- RankSelection: Selects individuals based on their rank within the population.
- RouletteWheelSelection: Selects individuals based on their fitness proportionally.

### · Crossover:

- TwoPointCrossover: Combines genes from two parents at two crossover points.
- UniformCrossover: Combines genes from two parents based on a uniform distribution.
- PureSinglePointCrossover: Combines genes from two parents at a single crossover point.

### Mutation

- GuidedMutation: Performs guided mutation based on specific rules or heuristics.
- GenProgMutation: Performs genetic programming mutation by changing operations and parameters.

### 1.0.7 Project Structure

```
CMakeLists.txt
                                              # Main CMake configuration file for the entire project
                                              # Configuration file for generating documentation with Doxygen
# License information for the project
Doxvaen.confia
LICENSE.txt
README.md
                                                Project's readme file with instructions and documentation
                                              # Build directory containing compiled binaries and test
     executables
                                           # Directory for main binary executables
# Main executable for the Circuit Optimizer project
   bin
      Circuit_Optimizer
   tests
                                             # Directory for test executables and related files
                                             # Binaries for the tests
       bin
          test_circuit_simulator
                                            # Test executable for circuit simulator
          test_genetic_algorithm
                                            # Test executable for genetic algorithm
          test_validity_checker
                                            # Test executable for validity checker
                                             # Other test-related files
circuit_diagrams
                                              # Directory for circuit diagrams
                                             # Example circuit diagram image
   circuit.png
                                              # Documentation files
                                             # API documentation in PDF format
   API_Document.pdf
   CCircuit_Document_Analysis.md
                                             # Markdown document with circuit design analysis
   CCircuit_Document_Analysis.pdf
                                             # PDF document with circuit design analysis
   CSimulator_Document_and_Analysis.md
                                             # Markdown document with simulator performance analysis
   CSimulator_Document_and_Analysis.pdf
                                             # PDF document with simulator performance analysis
   GeneticAlgorithm_Design_Document.md
                                             # Markdown document with algorithm design illstration
```

```
GeneticAlgorithm_Design_Document.pdf
                                            # PDF document with algorithm design illstration
                                            # Markdown document with algorithm & circuit params analysis
   GeneticAlgorithm_Params_Analysis.md
   GeneticAlgorithm_Params_Analysis.pdf
                                            # Markdown document with algorithm & circuit params analysis
   Problem_Statement.pdf
                                            # Project problem statement
   doxygen
                                            # Directory for Doxygen-generated documentation files
# Header files for the project
include
   CSimulator.h
                                            # Header for the circuit simulator
   Genetic_Algorithm.h
                                            # Header for the genetic algorithm
   circuit
                                            # Directory for circuit-related headers
      CCircuit.h
                                           # Header for the circuit class
                                           # Header for the unit class
      CUnit.h
   operators
                                            \# Directory for operator-related headers
                                           # Crossover operator headers
      Crossover
         PureSinglePointCrossover.h
                                          # Header for pure single point crossover
         TwoPointCrossover.h
                                          # Header for two-point crossover
         UniformCrossover.h
                                          # Header for uniform crossover
      Mutation
                                           # Mutation operator headers
                                          # Header for genetic programming mutation
# Header for guided mutation
         GenProgMutation.h
         GuidedMutation.h
                                           # Selection operator headers
          NaryTournamentSelection.h
                                           # Header for N-ary tournament selection
          RankSelection.h
                                           # Header for rank selection
          RouletteWheelSelection.h
                                           # Header for roulette wheel selection
                                            # Utility headers
   utils
                                            # Header for helper functions
       Helper.h
                                            # Header for individual class
       Individual.h
       Parameter.h
                                            # Header for parameter handling
       tqdm.hpp
                                            # Header for progress bar utility
investigate
                                             # Directory for investigation scripts and logs
   CCircuit.cpp
                                            # Circuit source file for investigation
                                            # Jupyter notebook for evaluating circuits
   evaluate circuit.ipvnb
   evaluate_circuit_time.ipynb
                                            # Jupyter notebook for evaluating circuit times
   execution_times.txt
                                            # Log file for execution times
   timings_thread.log
                                            # Log file for thread timings
                                            # Directory for log files
# Log directory with timestamp
   2024-05-23_21-28-04
                                            # Log file for circuit information
      circuit_info.log
                                             # Directory for post-processing scripts
post_process
   visualize.ipynb
                                            # Jupyter notebook for visualization
                                            # Python script for visualization
   visualize.py
scripts
                                             # Shell scripts for running the project
                                            \ensuremath{\text{\#}} Script to run the main optimizer
   run.sh
                                            # Script to run the tests
   run test.sh
                                             # Source code for the project
                                            # CMake configuration file for source code
   CMakeLists.txt
   CSimulator.cpp
                                            # Source file for the circuit simulator
   Genetic_Algorithm.cpp
                                            # Source file for the genetic algorithm
   circuit
                                            # Directory for circuit-related source files
                                           \ensuremath{\text{\#}} Source file for the circuit class
      CCircuit.cpp
                                           # Source file for the unit class
      CUnit.cpp
   main.cpp
                                            # Main source file for the project
                                            # Directory for operator-related source files
   operators
      Crossover
                                           # Crossover operator source files
         PureSinglePointCrossover.cpp  # Source file for pure single point crossover
         TwoPointCrossover.cpp
                                          # Source file for two-point crossover
         UniformCrossover.cpp
                                        # Source file for uniform crossover
      Mutation
                                           # Mutation operator source files
         GenProgMutation.cpp
                                          # Source file for genetic programming mutation
         GuidedMutation.cpp
                                          # Source file for guided mutation
      Selection
                                           # Selection operator source files
                                           # Source file for N-ary tournament selection
          NarvTournamentSelection.cpp
                                           # Source file for rank selection
          RankSelection.cpp
          RouletteWheelSelection.cpp
                                           # Source file for roulette wheel selection
                                            # Utility source files
   utils
       Helper.cpp
                                            # Source file for helper functions
                                             # Test files
tests
                                           # Test IIIes
# CMake configuration file for tests
    CMakeLists.txt
    test_circuit_simulator.cpp
test_circuit_time.cpp
                                            # Test file for circuit simulator
                                            # Test file for circuit timing
    test_genetic_algorithm.cpp
                                             # Test file for genetic algorithm
    test_validity_checker.cpp
                                             # Test file for validity checker
```

### 1.0.8 Contributing

Contributions are welcome! Please follow these steps:

- 1. Fork the repository.
- 2. Create a new branch (git checkout -b feature-branch).
- Make your changes.
- 4. Commit your changes ('git commit -m 'Add new feature') .

- 5. Push to the branch (git push origin feature-branch`).
- 6. Create a new Pull Request.

### 1.0.9 License

This project is licensed under the MIT License. See the LICENSE.txt file for more details.

### **Chapter 2**

### Optimised Circuit Configuration vs Economic Factors

### 2.1 Introduction

To get a heuristic circuit design we utilised our base case solution and varied 3 economic factors (Number of units in the circuit, Price paid per gerardium relative to waste, Purity of the input feed) independently to observe how the optimal circuit configuration changes with each factor. This allows us to identify common patterns in the optimal circuit.

### 2.2 Base Case (10 Units with Specified Economic Factors)

From the base case circuit solution, we observed several features:

- 1. A row of units which handles further filtering of the tail streams. (Units 5, 0, 4, 7, 3)
- 2. A unit which takes in the concentrate streams from the row of units. (Unit 1)
- 3. A unit to handle one last filtering of the material before sending to the concentrate (Unit 8)
- 4. A unit which takes in the intermediate streams from this row of units, and recycles its tailing to the start of the tail stream unit row and directs its concentrate stream towards the concentrate unit (Unit 6)
- 5. A unit which allows recycling of material midway through the circuit (Unit 9)

### 2.3 Economic Factors:

### 2.3.1 1) Number of Units

Investigated circuits with a number of units from 5 to 15:

There are common features among the optimal configuration of circuits with a different number of units

- The concentrate and tailing streams are connected to only one other unit each
- · Overall they have a similar structure to the base case as described above, these features includes:
  - A row of units to handle the tailing streams (This row is longer and more obvious as number of units increase)
  - 1 unit which handles the intermediate streams of these units
  - 1 unit which handles the concentrate stream of these units
- Circuits with more units, have units to handle recycling and further filtering between these structures, as seen in the base case and the circuit with 15 units below (2 units, Unit 6, 7, handles filtering before depositing in concentrate)

Circuit with 5 units: Circuit with 15 units:

### 2.3.2 2) Price of Reward and Penalty

Investigated 4 different cases utilising circuits of 5 units, 10 units:

- 1. Reward of £100 \* 5 per kg
- 2. Reward of £100 \* 10 per kg
- 3. Penalties of £750 \* 5 per kg
- 4. Penalties of £750 \* 10 per kg

#### With 5 units:

- At higher penalty, the optimal circuit's intermediate streams showed tendency to recycle (Unit 3, 4, 6 conducts further filtering), following the route of the tail stream, as opposed to continuing forward in the circuit towards the concentrate
- For higher rewards, the circuit is comparable to the base case for 5 units, just with an additional unit to filter for the tailing stream (to extract as much value from this stream)

### With 10 units:

- The same general structure mentioned multiple times above was observed for various degrees of penalities and rewards
- A similar trend to the circuit of 5 units was observed
- For higher rewards, there are more filters for tailing stream:

### £1000 reward, 10 units circuit

- For a high penalty there is an increased number of units to filter out the material before depositing into the concentrate (Units 8, 4, 3), and fewer units to filter out tailing stream
- This is expected as we want to prevent as much waste in the concentrate as possible, therefore prioritise this over collecting material from tailing stream

### £7500 penalty, 10 units circuit

### 2.3.3 3) Purity

Investigated a range of gerardium feed purity, from 10-90%:

- There was a general trend where an increase in purity of gerardium corresponded to more intermediate flow streams directly connected to the concentrate.
  - For an extreme case of 90% geranium and 10% waste for a circuit of 5 units it connects all unit's concentrate and intermediate streams directly to the concentrate.

### 90% gerardium feed, 5 units circuit:

### 90% gerardium feed, 10 units circuit:

- A similar trend, to a lesser degree, can be seen in the 10 units circuit, where earlier on in the row of units (Units 0, 8, 7) the intermediate streams directly connect to the concentrate, but later on in the row (with increasing waste content) the intermediate streams is recycled to the start of the tailing row. Note that there is no unit designated for recycling like in the base case.
- Overall the general design for having a row of units to filter out the tailing stream holds true for the range of purities explored.
- The circuit design differs at lower purities where it is closer to the base case with fewer units in a row and units utilised for further filtering prior to deposition to the concentrate.

### 2.4 Heuritic Circuit Design

Based on the above observations we can see some similarities in the design which could provide an insight into a heuristic circuit design to consider:

- 1. Having a row of units to filter out the tailing streams leading to the tail.
- 2. A unit to handle concentrate streams from this row of units.
- 3. Given sufficient units, the intermediate streams from these rows will be fed into another unit before either being sent to the concentrate unit/recycled to the top of the row of units

### Additional Considerations:

- 1. Depending on the penalty cost, we can also arrange more units (if higher penalty) or less units (if lower penalty) before material reaches concentrate unit to further filter out the waste material from concentrate
- 2. Purity can also be taken into consideration when optimising a circuit. We can arrange more units for the tailing stream row (to extract as much valuable material from waste before deposal) the more pure the feed, the more units we can assign for this purpose

Optimised Circuit Configuration vs Economic Factors

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### **Chapter 3**

### **Document & Analysis for Circuit Simulator**

### 3.1 Introduction

This document provides a comprehensive design framework for the CSimulator class and its associated components. The CSimulator class is designed to simulate the mass balance of a separation circuit composed of multiple units, evaluate the performance of the circuit, and ensure that the circuit operates correctly under specified constraints.

### 3.2 1. Overview of CSimulator

### 3.2.1 1.1 Purpose

The purpose of the CSimulator class is to simulate the mass balance of a circuit that separates valuable material (gerardium) from waste. It calculates the mass flows through each unit in the circuit and determines the overall performance of the circuit.

### 3.2.2 1.2 Scope

The scope of the CSimulator includes:

- Parsing the circuit vector into a structured representation.
- · Simulating the mass balance for the entire circuit.
- · Evaluating the performance of the circuit based on the mass balance.
- · Ensuring the circuit reaches a steady-state solution.

### 3.3 2. Class Design

### 3.3.1 2.1 UML Class Diagram

```
kC: double, kI: double): void|
                        Attributes
                            | Attributes
                               + concNum: int
+ concNum: int
                           | + interNum: int
+ interNum: int
+ tailsNum: int
                            | + tailsNum: int
+ feedGerardium: double | + feedGerardium: double + feedWaste: double | + feedWaste: double
+ newFeedGerardium: double | + newFeedGerardium: double |
+ newFeedWaste: double | + newFeedWaste: double + differenceG: double | + differenceG: double
                          | + differenceW: double
+ differenceW: double
+ concPtr: CUnit*
                           | + concPtr: CUnit*
+ interPtr: CUnit*
                           | + interPtr: CUnit*
+ tailsPtr: CUnit*
                               + tailsPtr: CUnit*
+ mark: bool
                              + mark: bool
```

### 3.3.2 2.2 CSimulator Class

#### 3.3.2.1 2.2.1 Attributes

- circuit\_vector: A vector of integers representing the configuration of the circuit.
- units: A vector of CUnit objects, each representing a unit in the circuit.
- feed\_rate: A double representing the feed rate of the input stream.
- **tolerance**: A double specifying the tolerance for convergence in the simulation.
- max\_iterations: An integer representing the maximum number of iterations allowed for achieving convergence.

### 3.3.2.2 2.2.2 Methods

**Constructor:** The constructor initializes the CSimulator with a given circuit vector, feed rate, tolerance, and maximum number of iterations. It also calls a method to parse the circuit vector into a structured format.

**EvaluateCircuit:** This method evaluates the entire circuit and returns a performance score. It checks if the circuit reaches a steady-state solution within the specified maximum iterations. If convergence is not achieved, it returns a very negative value to indicate failure.

**SimulateUnit:** This method calculates the mass balance for a single unit in the circuit. It uses the feed rate and rate constants to compute the flow rates of the high-grade concentrate, intermediate stream, and tailings stream. **SolveSteadyState:** This method iteratively solves for the steady-state mass balance of the entire circuit. It repeatedly updates the feed rates of all units and checks for convergence within the specified tolerance. If all units' feed rates stabilize within the tolerance, it returns true; otherwise, it continues until the maximum iterations are reached.

**ParseCircuitVector:** This method parses the circuit vector into a structured representation by creating CUnit objects for each unit in the circuit. It sets the connections between units based on the vector.

### 3.3.3 2.3 CUnit Class

### 3.3.4 2.3.1 Attributes

- concNum: An integer representing the index of the unit for the high-grade concentrate stream.
- interNum: An integer representing the index of the unit for the intermediate stream.
- tailsNum: An integer representing the index of the unit for the tailings stream.
- **feedGerardium:** A double representing the feed rate of gerardium to the unit.
- feedWaste: A double representing the feed rate of waste to the unit.
- newFeedGerardium: A double representing the new feed rate of gerardium after processing.
- newFeedWaste: A double representing the new feed rate of waste after processing.
- differenceG: A double representing the relative change in feed rate of gerardium.
- differenceW: A double representing the relative change in feed rate of waste.

3.4 3. Function Definitions

- concPtr: A pointer to the CUnit receiving the concentrate stream.
- interPtr: A pointer to the CUnit receiving the intermediate stream.
- tailsPtr: A pointer to the CUnit receiving the tailings stream.
- mark: A boolean indicating whether the unit has been marked for traversal.

### 3.3.5 2.3.2 Methods

**Constructor:** The constructor initializes a CUnit with the indices for its concentrate, intermediate, and tailings streams. It also initializes the feed rates and other attributes.

**CalculateFlows:** This method calculates the mass flows for the unit based on the feed rate and given rate constants for the high-grade and intermediate streams. It computes the residence time and uses it to determine the flow rates for the concentrate, intermediate, and tailings streams.

### 3.4 3. Function Definitions

### 3.4.1 3.1 CSimulator Methods

### Constructor:

- The constructor initializes the CSimulator with the provided circuit vector, feed rate, tolerance, and maximum iterations.
- It calls the ParseCircuitVector method to convert the circuit vector into a structured format, creating
  a collection of CUnit objects representing the units in the circuit.

### **EvaluateCircuit:**

- This method first calls SolveSteadyState to iteratively solve for the steady-state mass balance.
- · If the steady-state solution is not achieved, it returns a very negative value to indicate failure.
- If convergence is achieved, it calculates the overall performance score by summing the contributions from all units based on their concentrate and tailings rates.

### SimulateUnit:

- · This method calculates the mass flows for a single unit.
- It uses the feed rate and rate constants to compute the recovery rates for the high-grade concentrate and intermediate streams.
- It then determines the flow rates for the concentrate, intermediate, and tailings streams based on these recovery rates.

### SolveSteadyState:

- This method iteratively updates the feed rates of all units and checks for convergence within the specified tolerance.
- It stores the old feed rates and compares them with the new feed rates to check for convergence.
- If all units' feed rates stabilize within the tolerance, it returns true; otherwise, it continues until the maximum number of iterations is reached.

### ParseCircuitVector:

- This method converts the circuit vector into a structured representation by creating CUnit objects for each unit.
- It sets the connections between units based on the vector, ensuring each unit is correctly linked to its concentrate, intermediate, and tailings streams.

### 3.4.2 3.2 CUnit Methods

### Constructor:

- The constructor initializes a CUnit with the indices for its concentrate, intermediate, and tailings streams.
- · It sets the initial feed rates and other attributes.

### CalculateFlows:

- This method calculates the mass flows for the unit based on the feed rate and given rate constants.
- It computes the residence time and uses it to determine the flow rates for the concentrate, intermediate, and tailings streams.
- The calculated flow rates are then assigned to the respective attributes of the CUnit.

### 3.5 4. Validation and Testing

### 3.5.1 4.1 Unit Tests

### · CSimulator Constructor:

Test initialization with valid and invalid circuit vectors to ensure proper setup and error handling.

### EvaluateCircuit:

 Validate performance evaluation by comparing the calculated performance score with expected values for known valid circuits.

### SimulateUnit:

 Verify mass flow calculations by checking the output flow rates for various input feed rates and rate constants.

### · SolveSteadyState:

 Check convergence behavior for different circuit configurations and feed rates, ensuring the method stops iterating once convergence is achieved.

### · ParseCircuitVector:

 Test the parsing of circuit vectors to ensure that the resulting unit configurations match the expected structure.

### 3.5.2 4.2 Integration Tests

### Complete Circuit Simulation:

 Validate end-to-end simulation for sample circuits, ensuring correct performance score and convergence for the entire circuit.

### Boundary Conditions:

 Test simulator behavior with edge cases such as zero feed rate, maximum allowable iterations, and extreme values for tolerance.

#### 3.6 5. Performance Considerations

#### · Efficiency:

 Optimize the iterative solver to achieve faster convergence by improving the algorithm and reducing computational overhead.

#### · Scalability:

 Ensure the simulator can handle larger circuits with more units without significant performance degradation, potentially through parallel processing.

#### · Parallelization:

Consider parallelizing unit simulations to leverage multi-core processors, which can improve performance and reduce simulation time.

### 3.7 6. Usage

#### · User Guide:

- Provide detailed instructions for setting up the environment, running the simulator, and interpreting the output. Include examples and usage scenarios to help users understand how to use the tool effectively.
- Ensure that the user guide includes steps to compile and run the simulator, required dependencies, and configuration options.
- Example:

#### 3.7.0.1 Setup

- 1. Clone the repository.
- 2. Navigate to the project directory.
- 3. Install required dependencies:

```
shell sudo apt-get install g++ sudo apt-get install cmake
```

#### 3.7.0.2 Build the project

```
shell mkdir build cd build cmake .. make
```

#### 3.7.0.3 Running the Simulator

To run the simulator, use the following command:  $shell ./simulator < input_file>$  Replace  $< input_file>$  with the path to your input configuration file.

#### 3.7.0.4 Example

```
shell ./simulator config.txt
```

### 3.7.0.5 Interpreting the Output

The simulator will output the performance score and the detailed mass flow rates for each unit in the circuit.

#### · Code Comments:

 Ensure all methods and key sections of the code are well-documented with comments explaining their functionality, parameters, and return values.

#### - Example:

```
// Constructor for CSimulator
// Initializes the simulator with the given circuit vector, feed rate, tolerance, and max iterations.
CSimulator::CSimulator(std::vector<int>& circuit_vector, double feed_rate, double tolerance, int max_iterations) {
    // Initialization code
}
```

#### · Examples:

- Include example circuit vectors and expected outputs to help users understand the simulator's behavior and validate its performance.
- Example:

#### 3.7.1 Example Circuit Vector

```
0, 1, 3, 3, 2, 2, 0, 4, 1, 1, 1, 0, 5
```

#### 3.7.2 Expected Output

```
Performance Score: 110.25
Unit 0: Concentrate Rate: 1.5, Intermediate Rate: 0.3, Tailings Rate: 8.2
```

By following this detailed design framework, the CSimulator class can be developed to provide robust and efficient simulation of circuit configurations, meeting the requirements of the project and delivering accurate and reliable performance evaluations.

# **Genetic Algorithm Design**

#### 4.1 Overview

This project employs a genetic algorithm to optimize circuit configurations. Genetic algorithms are heuristic search algorithms inspired by the process of natural selection. They use operations such as selection, crossover, and mutation to find the optimal solution. This document provides a detailed explanation of the design structure and optimization strategies implemented in the genetic algorithm.

### 4.2 UML Graph

The following documentation provides an in-depth look at the design and implementation of the genetic algorithm components within our project. The structure of the project is depicted in the accompanying diagram, which illustrates the relationships between various header and implementation files.

#### Relationships

- The central file Operator.h is included by all specific operator header files (Crossover, Mutation, Selection), establishing a common base or interface for different types of genetic algorithm operators.
- Each operator header (. h) file is paired with its corresponding implementation (. cpp) file, where the declared methods and classes are defined.
- Genetic\_Algorithm.h includes various operator headers to utilize the defined crossover, mutation, and selection strategies within the genetic algorithm.
- The main program (main.cpp) and test file (test\_genetic\_algorithm.cpp) include Genetic\_Algorithm.h to execute and test the genetic algorithm functionalities.

The diagram effectively illustrates the modular structure of the genetic algorithm, showing clear separations between declarations and implementations, and the hierarchical inclusion of various components. This modularity facilitates easy maintenance, testing, and extension of the genetic algorithm with new operators or strategies. The project is organized to ensure that each component can be developed and tested independently, making it easier to manage and extend the genetic algorithm framework.

## 4.3 Directory Structure

The following is the directory structure relevant to the genetic algorithm:

```
include
  Genetic_Algorithm.h
                                        # Main genetic algorithm components and functions.
     Crossover
        PureSinglePointCrossover.h
                                      # Header for PureSinglePointCrossover class.
        TwoPointCrossover.h
                                      # Header for TwoPointCrossover class.
        UniformCrossover.h
                                      # Header for UniformCrossover class.
     Mutation
                                      # Header for GenProgMutation class.
        GuidedMutation.h
                                      # Header for GuidedMutation class.
     Selection
         NaryTournamentSelection.h
                                       # Header for NaryTournamentSelection class.
```

```
RankSelection.h
                                       # Header for RankSelection class.
       RouletteWheelSelection.h
                                       # Header for RouletteWheelSelection class.
utils
    Helper.h
                                        \ensuremath{\text{\#}} Utility functions for logging and current time.
                                        # Definition of the Individual class used in the genetic
    Individual.h
  algorithm.
    Parameter.h
                                        # Definition of the parameter structures for the algorithm.
 Genetic_Algorithm.cpp
                                         # Implementation of the main genetic algorithm components and
  functions.
 operators
    Crossover
       PureSinglePointCrossover.cpp # Implementation of PureSinglePointCrossover class.
       TwoPointCrossover.cpp # Implementation of IworoIncolocity
UniformCrossover.cpp # Implementation of UniformCrossover class.
                                         Implementation of TwoPointCrossover class.
       UniformCrossover.cpp
    Mutation
                                       # Implementation of GenProgMutation class.
       GenProgMutation.cpp
       GuidedMutation.cpp
                                      # Implementation of GuidedMutation class.
    Selection
        NaryTournamentSelection.cpp # Implementation of NaryTournamentSelection class.
                                         # Implementation of RankSelection class.
        RankSelection.cpp
                                      # Implementation of RouletteWheelSelection class.
        RouletteWheelSelection.cpp
```

### 4.4 Major Components

#### 4.4.1 1. Genetic Algorithm.h and Genetic Algorithm.cpp

These files define the core module of the genetic algorithm, including the main flow and operations of the algorithm, such as population initialization, parent selection, crossover, mutation, and generation of new populations.

#### 4.4.1.1 Key Functions

- initializeRandomSeed: Initializes the random seed.
- initializePopulation: Initializes the population.
- elitism: Implements the elitism strategy, preserving the best individuals.
- optimize: The main optimization function, executing the main loop of the genetic algorithm.

#### 4.4.2 2. Selection Operators

Selection operators are responsible for selecting individuals from the current population to serve as parents for the next generation.

#### 4.4.2.1 Files

- NaryTournamentSelection.h / NaryTournamentSelection.cpp: n-ary tournament selection, randomly selects n individuals to compete, and selects the best one.
- RankSelection.h / RankSelection.cpp: Rank-based selection, individuals are sorted by fitness, and selection probability is based on rank.
- RouletteWheelSelection.h/RouletteWheelSelection.cpp: Roulette wheel selection, selection probability is proportional to fitness.

#### 4.4.3 3. Crossover Operators

Crossover operators combine the genes of parent individuals to create new offspring.

#### 4.4.3.1 Files

- PureSinglePointCrossover.h / PureSinglePointCrossover.cpp: Single-point crossover, swaps genes between two parents at a random point.
- TwoPointCrossover.h / TwoPointCrossover.cpp: Two-point crossover, swaps genes between two random points.

 UniformCrossover.h/UniformCrossover.cpp: Uniform crossover, randomly selects genes from each parent.

#### 4.4.4 4. Mutation Operators

Mutation operators introduce random changes to the genes of an individual, increasing the diversity of the population.

#### 4.4.4.1 Files

- GenProgMutation.h / GenProgMutation.cpp: General-purpose mutation, randomly modifies the genes of an individual.
- GuidedMutation.h / GuidedMutation.cpp: Guided mutation, modifies genes based on specific rules or heuristics

#### 4.4.5 5. Utilities

Auxiliary functions and structures definitions.

#### 4.4.5.1 Files

- Helper.h / Helper.cpp: Contains auxiliary functions like logging results and getting the current time.
- Individual.h: Defines the individual structure, including the gene sequence and fitness.
- Parameter.h: Defines the structure for algorithm parameters, including population size, crossover rate, mutation rate, etc.

# 4.5 Optimization Strategies

#### 4.5.1 1. Elitism Strategy

The elitism strategy in genetic algorithms ensures that the best-performing individuals from the current generation are preserved and carried over to the next generation. This helps in maintaining the quality of solutions over generations and prevents the loss of the best solutions found so far.

#### 4.5.1.1 Implementation Details

The implementation of the elitism strategy is done in the elitism function in Genetic\_Algorithm.cpp. This function sorts the population based on fitness and selects the top individuals to be directly transferred to the new population.

#### 4.5.1.2 Code Explanation

```
**elitism**
```

The elitism function takes the current population and a reference to a new population vector where the elite individuals will be stored. It also takes the algorithm parameters which include the proportion of the population to be preserved as elites.

#### 4.5.1.3 Steps in the elitism Function

 Copy the Population: A copy of the current population is created to avoid modifying the original population directly.

```
std::vector<Individual> sortedPopulation = population;
```

2. **Sort the Population**: The copied population is sorted based on the fitness values of the individuals in descending order. This means the individuals with the highest fitness values come first.

```
std::sort(sortedPopulation.begin(), sortedPopulation.end(), [](const Individual &a, const Individual
&b) {
    return a.fitness > b.fitness;
});
```

3. **Calculate the Number of Elites**: The number of elite individuals to be preserved is calculated based on the elite percentage parameter.

```
int numElites = static_cast<int>(params.populationSize * params.elitePercentage);
```

4. **Add Elite Individuals to the New Population**: The top elite individuals are added to the new population. If an individual's fitness is invalid, the process stops to avoid adding unfit individuals.

```
for (int i = 0; i < numElites; i++) {
   if (sortedPopulation[i].fitness == INVALID_FITNESS) {
      break;
   }
   newPopulation.push_back(sortedPopulation[i]);
}</pre>
```

#### 4.5.1.4 Integration with the Main Optimization Loop

In the main optimization function (optimize), the elitism function is called to add the best individuals from the current population to the new population before proceeding with the selection, crossover, and mutation operations. std::vector<Individual> newPopulation; elitism(population, newPopulation, params);

By preserving the best individuals, the elitism strategy ensures that the genetic algorithm does not lose the highest quality solutions found in each generation, thereby maintaining or improving the overall quality of the population over time.

#### 4.5.2 2. Selection Strategies

Selection strategies are critical in genetic algorithms as they determine how individuals are chosen to create the next generation. The implemented selection strategies in this project include Nary Tournament Selection, Rank Selection, and Roulette Wheel Selection. Each strategy has its advantages in maintaining diversity and preventing premature convergence.

#### 4.5.2.1 Nary Tournament Selection

**Nary Tournament Selection** is a method where 'n' individuals are randomly chosen from the population, and the one with the highest fitness among them is selected as a parent. This process helps in maintaining a healthy diversity in the population and prevents premature convergence to suboptimal solutions.

#### 4.5.2.2 Rank Selection

**Rank Selection** assigns selection probabilities to individuals based on their rank rather than their raw fitness values. This ensures that even individuals with lower fitness have a chance of being selected, thereby maintaining diversity in the population.

#### 4.5.2.3 Roulette Wheel Selection

**Roulette Wheel Selection** (also known as fitness proportionate selection) assigns selection probabilities proportional to the individuals' fitness values. Individuals with higher fitness have a higher chance of being selected, but there is still a probability for lower fitness individuals to be chosen, thus maintaining diversity.

### 4.5.3 3. Crossover Strategies

Crossover strategies play a vital role in genetic algorithms by combining the genetic information of parent individuals to create offspring for the next generation. This process helps in exploring the search space and finding better solutions.

#### 4.5.3.1 PureSinglePointCrossover

**PureSinglePointCrossover** involves selecting a random crossover point and exchanging the subsequences after this point between two parent individuals. This strategy helps to combine different features from the parents, potentially leading to better offspring.

#### 4.5.3.2 TwoPointCrossover

**TwoPointCrossover** is similar to the single-point crossover but involves selecting two random points. The segments between these two points are exchanged between the parents, leading to potentially more diverse offspring.

#### 4.5.3.3 UniformCrossover

**UniformCrossover** randomly selects genes from each parent, independent of their positions, resulting in a higher level of genetic diversity in the offspring. This method can be particularly effective in preventing premature convergence.

#### 4.5.4 4. Mutation Strategies

Mutation strategies introduce variability in the genetic algorithm population, helping to explore the search space more thoroughly and avoid local optima.

#### 4.5.4.1 GenProgMutation

**GenProgMutation** introduces random changes to the genes of an individual. This randomness helps the algorithm to avoid being trapped in local optima by exploring new areas of the search space.

#### 4.5.4.2 GuidedMutation

**GuidedMutation** uses heuristic information to guide the mutation process. This approach increases the effectiveness of the mutations by focusing on areas more likely to improve the individual's fitness.

#### 4.5.5 5. Simulated Annealing

Simulated Annealing (SA) is a probabilistic technique used for approximating the global optimum of a given function. In the context of genetic algorithms, SA is utilized to accept or reject new individuals (offspring) based on a temperature-controlled probability. This helps in balancing the exploration and exploitation of the search space. Simulated Annealing helps the genetic algorithm escape local optima by occasionally accepting worse solutions with a certain probability. This probability decreases as the algorithm progresses, controlled by a parameter called "temperature."

#### 4.5.5.1 Key Components of Simulated Annealing in Genetic Algorithms

- 1. **Acceptance Probability**: Determines whether to accept a new solution based on the difference in fitness and the current temperature.
- 2. **Temperature Schedule**: Controls how the temperature decreases over time.
- 3. **Integration with Genetic Algorithm**: Used in conjunction with crossover and mutation to guide the search process.

#### 4.5.5.2 Code Explanation

#### **Acceptance Probability**

The acceptance probability is calculated using the formula:  $P = \exp(\frac{D}{E}T)$  \$\\$ where:

- is the change in fitness (new fitness current fitness).
- · is the current temperature.

```
// Genetic_Algorithm.cpp

// Calculate acceptance probability of Simulated Annealing functions
double acceptanceProbability(double currentEnergy, double newEnergy, double temperature) {
   if (newEnergy > currentEnergy) {
      return 1.0;
   }
   return std::exp((newEnergy - currentEnergy) / temperature);
}
```

If the new solution is better (>0), the probability is greater than 1, and the new solution is accepted. If the new solution is worse (<0), it might still be accepted based on the calculated probability, allowing the algorithm to escape local optima.

The temperature is gradually decreased according to a cooling schedule. This reduces the likelihood of accepting worse solutions as the algorithm progresses, focusing more on exploitation of the best solutions found so far. The cooling schedule can be a simple linear decrement or a more complex function, but in our implementation, a straightforward decrement is used: T = T - Delta T where is a small decrement value.

#### **Simulated Annealing Process**

During each iteration of the genetic algorithm, after generating offspring through crossover and mutation, the applySimulatedAnnealing function is applied to decide whether to accept the offspring. This mechanism ensures that even suboptimal solutions can occasionally be accepted, promoting diversity and preventing premature convergence to suboptimal solutions.

// Genetic\_Algorithm.cpp

```
bool applySimulatedAnnealing(Individual &offspring, Individual &parent1, Individual &parent2, int
      vector_size,
                             double (&func) (int, int *, struct CircuitParameters), bool (&validity) (int, int
      *),
                             double &Temp, const Algorithm_Parameters &params, const CircuitParameters
      c_params,
                             std::mt19937 &generator) {
    \ensuremath{//} Calculate the fitness (energy) of the offspring
    if (validity(vector_size, offspring.genes.data())) {
       offspring.fitness = func(vector_size, offspring.genes.data(), c_params);
       offspring.fitness = INVALID_FITNESS; // Give a negative fitness for invalid individuals
        return false; // Reject invalid offspring
    double parents_fitness = (parent1.fitness + parent2.fitness) / 2;
    // Simulated annealing acceptance criterion
    std::uniform_real_distribution<double> distribution(0.0, 1.0);
    // Check the threshold
    if (Temp > 1) {
        if (acceptanceProbability(parents_fitness, offspring.fitness, Temp) > distribution(generator)) {
           return true; // Accept the offspring
        // Cool down the temperature
        Temp -= params.deltT;
    } else {
        // Keep the original parents
        if (parent1.fitness > parent2.fitness) {
            offspring = parent1;
           offspring = parent2;
        return true:
    return false; // Reject the offspring
```

#### **Explanation of the Code**

• Fitness Calculation: The fitness (energy) of the offspring is calculated using the provided fitness function func. If the offspring is valid, its fitness is evaluated; otherwise, it is assigned a negative fitness value, and the offspring is rejected.

- Parents' Fitness: The average fitness of the two parents is computed. This serves as the baseline for comparing the fitness of the offspring.
- Acceptance Criterion: A uniform random number generator determines whether the offspring is accepted based on the calculated acceptance probability.
- Temperature Check and Cooling: The temperature is checked to see if it is above a certain threshold. If so, the acceptance probability is compared with a random number to decide if the offspring should be accepted. The temperature is then decreased according to the cooling schedule. If the temperature is too low, the offspring is not accepted unless it is better than the worse parent.
- **Updating Offspring**: If the offspring is accepted, it becomes part of the population. If not, the better of the two parents is retained.

#### Integration with the Genetic Algorithm

In the main loop of the genetic algorithm, simulated annealing is applied after the crossover and mutation steps. This ensures that the offspring are accepted based on the temperature-controlled probability.

// Genetic\_Algorithm.cpp

```
// Main function to optimize the solution using a genetic algorithm
int optimize(int vector_size, int *vector, double (&func)(int, int *, struct CircuitParameters),
             bool (&validity)(int, int *), Algorithm_Parameters params, CircuitParameters c_params) {
                // Parameters for Simulated Annealing
                double Temp = params.initialTemp;
                // Create offspring1 with Simulated Annealing Acceptance
                if (localNewPopulation.size() < params.populationSize) {</pre>
                        crossover->setParents(offspring2, selectedParents[parent1_index],
      selectedParents[parent2_index]);
                        crossover->apply(generators[omp_get_thread_num()]);
                        mutation->setIndividual(offspring2):
                        mutation->apply(generators[omp get thread num()]);
                    } while (!applySimulatedAnnealing(offspring2, selectedParents[parent1_index],
      selectedParents[parent2_index], vector_size, func, validity, Temp, params, c_params,
      generators[omp_get_thread_num()]));
                    localNewPopulation.push_back(offspring2);
                }
            }
        // ...
```

In summary, simulated annealing is an essential component of the genetic algorithm in this project, allowing the algorithm to balance between exploration and exploitation by accepting new individuals based on a temperature-controlled probability. This helps the algorithm to escape local optima and find better solutions over time.

#### 4.5.6 6. Progress Tracking with tqdm

In our genetic algorithm implementation, we use the self-implemented tqdm.hpp to provide real-time progress tracking and visualization of the optimization process. This helps developers and users monitor the progress of the algorithm, gain insights into the computational workload, and estimate the time remaining for completion.

#### 4.5.6.1 Integration of tgdm in Genetic Algorithm

Tqdm is integrated into the main loop of the genetic algorithm to track and display the progress of each generation. This provides a dynamic and user-friendly way to observe the algorithm's performance.

1. #### Initialization of tqdm

We initialize tqdm to set up the progress bar before entering the main loop of the genetic algorithm. The progress bar is configured to track the number of generations.

1. #### Progress Update

Within the main loop, tqdm updates the progress bar at each iteration, showing the current generation number and other relevant metrics, such as the maximum fitness of the population.

Here's the code snippet demonstrating the integration of tqdm:

#### 4.5.6.2 Detailed Explanation

- Initialization: The tqdm progress bar is initialized using tqdm::trange with the maximum number of generations as the parameter. The prefix "Iterating" is set to make it clear what the progress bar represents.
- Progress Update: Inside the main loop of the genetic algorithm, the progress bar is updated at each generation. The current maximum fitness value is also displayed, providing a real-time metric for the algorithm's performance.
- Benefits: This integration provides the following benefits:
  - Real-time Monitoring: Users can monitor the progress and performance of the algorithm in real-time.
  - User-Friendly Visualization: The progress bar offers a user-friendly way to track the computational process.
  - Performance Insights: Displaying metrics like maximum fitness helps in understanding the algorithm's behavior and performance trends.

The use of tqdm significantly enhances the usability and transparency of the genetic algorithm, making it easier to track and understand the optimization process.

#### 4.5.7 7. Optimization with OpenMP

In addition to tqdm, the genetic algorithm implementation leverages OpenMP for parallel processing. OpenMP is used to parallelize the evaluation of the population's fitness and other computationally intensive tasks, significantly improving the algorithm's performance on multi-core systems.

#### 4.5.7.1 Integration of OpenMP in Genetic Algorithm

OpenMP is integrated into the main loop of the genetic algorithm to parallelize the evaluation of fitness and other operations. This provides a substantial performance boost by utilizing multiple CPU cores.

Here's the code snippet demonstrating the integration of OpenMP: #include <omp.h> // Include the OpenMP library

```
}

// Selection, crossover, and mutation operations
// ...
#pragma omp parallel for
for (int i = 0; i < params.populationSize; i++) {
    // ... Genetic operations code ...
}

// Update progress bar
iter « "Max Fitness: " « maxFitness;
}
// ...
}</pre>
```

#### 4.5.7.2 Detailed Explanation

- Parallel Initialization: The population initialization is parallelized using #pragma omp parallel for, allowing multiple individuals to be initialized simultaneously.
- Parallel Fitness Evaluation: The evaluation of the population's fitness is parallelized, significantly speeding up the process, especially for large populations.
- Parallel Genetic Operations: Selection, crossover, and mutation operations are also parallelized to maximize performance.

#### 4.5.7.3 Benefits of Using OpenMP

- **Improved Performance**: Leveraging multiple CPU cores significantly reduces the time required for fitness evaluation and genetic operations.
- Scalability: The algorithm can handle larger populations and more generations within a reasonable time frame.
- **Efficiency**: Parallel processing makes the genetic algorithm more efficient, enabling faster convergence to optimal solutions.

The integration of OpenMP with tqdm provides a powerful combination of performance and usability, making the genetic algorithm both fast and user-friendly.

# **Genetic Algorithm Parameters Analysis**

This document presents a detailed analysis of the various parameters used in the Genetic Algorithm for optimizing circuit configurations. The goal is to determine the optimal settings for each parameter to ensure efficient and effective performance of the algorithm.

#### 5.1 Introduction

Genetic Algorithms (GAs) are powerful optimization tools inspired by the principles of natural selection and genetics. The performance of GAs is highly dependent on the proper tuning of their parameters. This analysis focuses on two main categories of parameters:

- · Algorithm Parameters
- Circuit Parameters

## 5.2 Algorithm Parameters Analysis

#### 5.2.1 Population Size

Population size determines the number of individuals in each generation. A larger population size increases the genetic diversity but also requires more computational resources.

#### 5.2.1.1 Analysis Results

- Population sizes of 600, 800, and 1000 start with relatively high initial fitness values, indicating that they quickly find high-quality solutions.
- The fitness values increase rapidly in the early generations.
- A population size of 400 shows the fastest increase in fitness, reaching 167.378 by generation 20.
- All population sizes eventually stabilize at a fitness value of 167.378, demonstrating effective convergence of the algorithm.

#### 5.2.1.2 Parameter Choice

A population size of 400 is optimal as it balances computational cost and convergence speed effectively.

#### 5.2.2 Tournament Size

Tournament size determines the number of individuals randomly selected from the population for each tournament. The winner of each tournament, being the individual with the highest fitness, is selected to be a parent for the next generation.

#### 5.2.2.1 Analysis Results

- All tournament sizes show a rapid increase in fitness within the first 10 generations.
- Tournament Size 5 reaches the highest fitness value (around 165) by generation 20 and remains stable.
- Sizes 6 and 7 also achieve high fitness values but are slightly lower than Tournament Size 5.

#### 5.2.2.2 Parameter Choice

Choosing a tournament size of 5 ensures that the genetic algorithm efficiently explores the solution space and quickly converges to high-quality solutions, making it suitable for most optimization problems.

#### 5.2.3 Crossover Rate

The crossover rate is a probability value (usually between 0 and 1) that dictates the likelihood of applying the crossover operation to generate new offspring from two parent individuals. It helps control the genetic diversity in the population by mixing genes from different parents.

#### 5.2.3.1 Analysis Results

- Crossover rates of 0.7 and 0.8 demonstrate the fastest convergence, achieving the maximum fitness value the quickest.
- All crossover rates eventually reach the same maximum fitness value, indicating robustness across different rates.

#### 5.2.3.2 Parameter Choice

For future runs, consider using a crossover rate of 0.8 to achieve the fastest improvement and robust performance.

#### 5.2.4 Mutation Rate

Mutation rate is a crucial parameter in genetic algorithms, representing the probability of randomly altering genes in an individual's genome. It helps maintain genetic diversity within the population, preventing premature convergence to local optima and enabling the exploration of the solution space.

#### 5.2.4.1 Analysis Results

- Lower mutation rates (0.01, 0.05, and 0.1) demonstrate quicker convergence and higher final fitness values compared to higher mutation rates (0.2 and 0.3).
- Mutation rate 0.01 shows the fastest and most stable improvement.

#### 5.2.4.2 Parameter Choice

Use a mutation rate of 0.01 to achieve rapid convergence and robust performance.

#### 5.2.5 Elite Percentage

The elitePercentage parameter in a genetic algorithm controls the elitism strategy. This strategy directly preserves the best-performing individuals from the current generation into the next generation without any mutation or crossover. This ensures that the top-quality genes are passed on, enhancing the algorithm's efficiency and stability.

#### 5.2.5.1 Analysis Results

- elitePercentage values of 0.01 show the highest fitness values consistently, reaching a fitness of 836.835 by generation 250.
- elitePercentage values of 0.05, 0.1, and 0.2 show slower fitness increases, with all stabilizing around 829.175 by generation 250.

#### 5.2.5.2 Parameter Choice

0.01 ensures rapid convergence to high fitness values while maintaining the robustness of the genetic algorithm.

## 5.3 Circuit Parameters Analysis

#### 5.3.1 Number of Units

The number of units in the circuit configuration can significantly affect performance metrics like elapsed time, performance, recovery, and grade.

#### 5.3.1.1 Analysis Results

Num Units	Elapsed Time (s)	Performance	Recovery	Grade	Final Circuit Configuration
4	1.35789	110.25	0.150332	0.965671	2112400033025
6	2.45652	232.583	0.280938	0.977566	1655523051524527
					002
8	4.5209	341.632	0.37187	0.989275	4771703036014035
					032039800
10	6.82153	437.72	0.471582	0.990517	08162701088814819
					8011813281227815

#### 5.3.1.2 Visualization

#### 5.3.1.3 Trends

- Elapsed Time: Increases almost linearly with the number of units, reflecting increased complexity and processing time.
- **Performance**: Improves significantly with an increasing number of units, indicating better processing capabilities and efficiency.
- **Recovery**: Increases with the number of units, suggesting improved system capacity to recover from errors or faults.
- **Grade**: Slight improvement with the number of units, indicating overall better performance and efficiency in larger configurations.

#### 5.3.2 Purity of Input Feed (Gerardium: Waste Feed Ratio)

The purity of the input feed, represented as the ratio of gerardium feed to waste feed, can significantly affect performance metrics like elapsed time, performance, recovery, and grade.

#### 5.3.2.1 Analysis Results

Gerardium:Waste Feed	Elapsed Time (s)	Performance	Recovery	Grade	Final Circuit Configuration
10:90	6.64633	437.72	0.471582	0.990517	57162351077
					27171071871
					92237147511
20:80	6.44538	862.07	0.494251	0.983232	8369364536
					105236111033
					52832052736
					1
30:70	6.97441	1320.43	0.498122	0.984718	722828910103
					102028138112
					04206237285

Gerardium:Waste Feed	Elapsed Time (s)	Performance	Recovery	Grade	Final Circuit Configuration
40:60	6.51499	1838.62	0.522236	0.984274	5 10 10 7 0 5 6 0 5
					11 10 0 5 0 3 9 0 3 8
					05210030340
					3 1
50:50	6.95478	2369.51	0.534505	0.985107	7 10 8 4 8 4 6 8 4 1
					108010378098
					4 11 8 0 5 10 10 3 8
					02
60:40	6.10315	3029.69	0.652294	0.970762	610839841081
					108298596710
					909611109610
					10 8
70:30	5.96057	4044.07	0.758418	0.969211	5 10 5 3 10 5 0 10 5
					1 10 5 8 10 5 2 10 4
					455752111059
					10 5 6
80:20	5.83807	5417.92	0.816378	0.97778	210211026103
					3 10 2 7 2 5 11 10 2
					8 10 2 4 10 2 5 10 2
					9 10 2 0
90:10	5.46021	7134.92	0.902775	0.984013	7 10 7 5 10 2 9 10 7
					3 10 7 0 10 7 8 10 7
					4 10 10 2 10 10 6 10
					7 1 10 5 11

#### 5.3.2.2 Visualization

#### 5.3.2.3 Trends

- Performance (red line, circles) shows a significant increase as the proportion of gerardium feed increases.
- Recovery (blue line, dashed) improves with higher gerardium feed, indicating better extraction efficiency.
- Grade (green line, dotted) remains relatively stable with slight fluctuations.
- **Elapsed Time** (orange line, squares) shows a decreasing trend, suggesting improved computational efficiency with higher purity feeds.

#### 5.3.3 Price Ratio of Gerardium to Waste Disposal

The economic parameters, specifically the reward for gerardium and the penalties for waste disposal, significantly affect the performance metrics of the circuit. This analysis examines how changes in these economic parameters impact the optimum circuit configuration, performance, recovery, and grade.

#### 5.3.3.1 Analysis Results

Default reward and penalty is 100 and -750.

Reward/← Penalty Coefficient	Num Units	Elapsed Time (s)	Performance	Recovery	Grade	Final Circuit Configuration
Reward of £100 * 5 per kg	5	1.99066	167.379	0.202835	0.977224	31462245
Reward of £100 * 5 per kg	10	6.86852	437.72	0.471582	0.990517	16986926 941066690 36933561 11697351

Reward/← Penalty Coefficient	Num Units	Elapsed Time (s)	Performance	Recovery	Grade	Final Circuit Configuration
Reward of £100 * 10 per kg	5	2.08692	167.379	0.202835	0.977224	2346533340114332
Reward of £100 * 10 per kg	10	6.76027	437.72	0.471582	0.990517	0 7 5 3 7 0 11 7 5 1 7 5 9 7 5 2 6 8 0 10 7 7 6 6 8 6 7 5 7 5 4
Penalty of £750 * 5 per kg	5	1.94702	167.379	0.202835	0.977224	15444324 36441003
Penalty of £750 * 5 per kg	10	6.7229	437.72	0.471582	0.990517	76916945 69698693 106655269 06711527
Penalty of £750 * 10 per kg	5	1.87025	162.125	0.196688	0.977106	03445333
Penalty of £750 * 10 per kg	10	7.42508	437.72	0.471582	0.990517	2 9 2 11 9 8 7 9 8 1 6 9 8 9 8 0 9 8 4 10 9 9 9 8 5 6 3 2 6 6 3

#### 5.3.3.2 Visualization

The visualizations above show the impact of different economic parameters on the circuit performance metrics for configurations with 5 and 10 units:

- **Performance**: Remains constant at 167.379 for lower penalty coefficients but drops to 162.125 when the penalty increases significantly. For 10 units, the performance remains stable at 437.72 across all economic parameters.
- **Elapsed Time**: Shows slight variations but generally increases with higher penalties. For 5 units, elapsed time decreases slightly with increased reward but rises with higher penalties.
- **Recovery**: Remains relatively constant for both 5 and 10 units, indicating that recovery is less sensitive to changes in economic coefficients.
- Grade: Remains stable for both 5 and 10 units, showing minimal fluctuation.

#### 5.3.4 Recommendations

- Optimize for Higher Units: Configurations with more units (10 units) are less sensitive to economic parameter changes, maintaining stable performance and grade.
- **Penalty Management**: When penalties for waste disposal are high, consider optimizing circuit configurations to minimize waste, as higher penalties can negatively impact performance.
- Reward Adjustments: Increasing rewards has minimal impact on performance and recovery, suggesting that optimizing for penalties might be more critical in achieving better economic outcomes.

# References

### 6.1 Research Papers

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- 2. **Yuan, Y., & Banzhaf, W. (2018).** "Arja: Automated repair of Java programs via multi-objective genetic programming." *IEEE Transactions on Software Engineering*, 46(10), 1040-1067. doi:10.1109/TSE.2018. 

  ∠2876535. Link
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#### 6.2 Official Documentation

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- 2. **Doxygen Documentation**. (2024). *Doxygen: The Standard for Code Documentation*. Retrieved from https://www.doxygen.nl/manual/
- 3. **CMake Documentation**. (2024). *CMake: Build, Test, and Package Software*. Retrieved from https←://cmake.org/documentation/

#### 6.3 Al Tools

- 1. **OpenAl ChatGPT**. (2024). *Conversation with GPT-4 on Genetic Algorithm Implementation*. Retrieved from https://chatgpt.com/share/02d1db48-b452-425d-9303-9183f7f1841a
- 2. **OpenAl ChatGPT**. (2024). *Conversation with GPT-40 on Elite Reservation Strategy*. Retrieved from https://chatgpt.com/share/ac8b9b73-cd17-4677-9677-daa1d8151d66

#### 6.4 Books

1. Henderson, D., Jacobson, S. H., & Johnson, A. W. (2003). "The theory and practice of simulated annealing." *Handbook of Metaheuristics*, 287-319. doi:10.1007/978-1-4757-3482-7\_10. Link

## 6.5 Code Repositories

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- 2. GitHub Repository. (2019). arja. Retrieved from https://github.com/yyxhdy/arja

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# 6.6 Miscellaneous

1. Author's Notes. (2024). Project notes and internal documentation.

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# **Namespace Documentation**

### 11.1 tqdm Namespace Reference

#### **Data Structures**

· class Chronometer

A simple chronometer to measure elapsed time.

class int\_iterator

A random access iterator for integers.

· class iter\_wrapper

A wrapper class for iterators to update the progress bar.

class progress\_bar

A class representing a progress bar.

· class range

A range of integers represented by iterators.

struct timer

A timer that can be used to create a timing iterator.

· class timing\_iterator

A forward iterator that returns the elapsed time.

class timing\_iterator\_end\_sentinel

An end sentinel for timing iterators.

• class tqdm\_for\_lvalues

A class for progress bars that iterate over Ivalues.

• class tqdm\_for\_rvalues

A class for progress bars that iterate over rvalues.

· class tqdm\_timer

A progress bar for a timer.

#### **Typedefs**

- using index = std::ptrdiff\_t
- using time\_point\_t = std::chrono::time\_point<std::chrono::steady\_clock>

#### **Functions**

double elapsed\_seconds (time\_point\_t from, time\_point\_t to)

Calculate the elapsed seconds between two time points.

void clamp (double &x, double a, double b)

Clamp a value between two bounds.

• int get\_terminal\_width ()

Get the width of the terminal window.

```
• template < class Container >
  tqdm_for_lvalues (Container &) -> tqdm_for_lvalues < typename Container::iterator >
• template<class Container >
  tqdm_for_lvalues (const Container &) -> tqdm_for_lvalues < typename Container::const_iterator >
• template < class Container >
  tqdm_for_rvalues (Container &&) -> tqdm_for_rvalues< Container >

    template < class ForwardIter >

  auto tqdm (const ForwardIter &first, const ForwardIter &last)
      Create a tqdm progress bar for a range of iterators.

    template < class ForwardIter >

  auto tqdm (const ForwardIter &first, const ForwardIter &last, index total)
      Create a tqdm progress bar for a range of iterators with a specified total size.
• template < class Container >
  auto tqdm (const Container &C)
      Create a tqdm progress bar for a container.

    template < class Container >

  auto tqdm (Container &C)
      Create a tqdm progress bar for a container.

    template < class Container >

  auto tqdm (Container &&C)
      Create a tqdm progress bar for a container (rvalue reference).

    template < class IntType >

  auto trange (IntType first, IntType last)
      Create a tgdm progress bar for a range of integers.

    template < class IntType >

  auto trange (IntType last)
      Create a tqdm progress bar for a range of integers.
• auto tqdm (timer t)
      Create a tqdm progress bar for a timer.
```

#### 11.1.1 Typedef Documentation

#### 11.1.1.1 index

```
using tqdm::index = std::ptrdiff_t
```

#### 11.1.1.2 time\_point\_t

using tqdm::time\_point\_t = std::chrono::time\_point<std::chrono::steady\_clock>

#### 11.1.2 Function Documentation

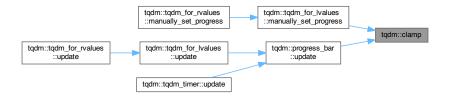
#### 11.1.2.1 clamp()

Clamp a value between two bounds.

#### **Parameters**

X	The value to clamp.
а	The lower bound.
b	The upper bound.

Here is the caller graph for this function:



### 11.1.2.2 elapsed\_seconds()

Calculate the elapsed seconds between two time points.

This function calculates the elapsed time in seconds between two time points.

#### **Parameters**

from	The starting time point.
to	The ending time point.

#### Returns

The elapsed time in seconds.

Here is the caller graph for this function:



#### 11.1.2.3 get\_terminal\_width()

```
int tqdm::get_terminal_width ( ) [inline]
Get the width of the terminal window.
```

#### Returns

The width of the terminal window in columns.

Here is the caller graph for this function:



#### 11.1.2.4 tqdm() [1/6]

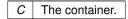
```
template<class Container > auto tqdm::tqdm (  {\rm const~Container~\&~C~)}
```

Create a tqdm progress bar for a container.

#### **Template Parameters**

Container	The type of the container.
-----------	----------------------------

#### **Parameters**



#### Returns

A tqdm progress bar.

Here is the call graph for this function:



#### 11.1.2.5 tqdm() [2/6]

Create a tqdm progress bar for a range of iterators.

#### **Template Parameters**

Forwardltor	The type of the forward iterator.

#### **Parameters**

first	The beginning of the range.
last	The end of the range.

#### Returns

A tqdm progress bar.

Here is the call graph for this function:



#### 11.1.2.6 tqdm() [3/6]

Create a tqdm progress bar for a range of iterators with a specified total size.

#### **Template Parameters**

ForwardIter	The type of the forward iterator.
-------------	-----------------------------------

#### **Parameters**

first	The beginning of the range.
last	The end of the range.
total	The total size of the range.

#### Returns

A tqdm progress bar.

Here is the call graph for this function:



### 11.1.2.7 tqdm() [4/6]

Create a tqdm progress bar for a container (rvalue reference).

#### **Template Parameters**

e container.
e container.

#### **Parameters**

C The container.

#### Returns

A tqdm progress bar.

Here is the call graph for this function:



## 11.1.2.8 tqdm() [5/6]

Create a tqdm progress bar for a container.

### **Template Parameters**

Container	The type of the container.
Comamer	The type of the container.

#### **Parameters**

C The container.

Returns

A tqdm progress bar.

Here is the call graph for this function:



#### 11.1.2.9 tqdm() [6/6]

Create a tqdm progress bar for a timer.

#### **Parameters**

t The timer.

#### Returns

A tqdm progress bar.

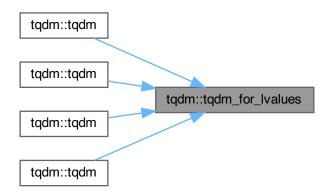
Here is the call graph for this function:



#### 11.1.2.10 tqdm\_for\_lvalues() [1/2]

#### 11.1.2.11 tqdm\_for\_lvalues() [2/2]

Here is the caller graph for this function:



#### 11.1.2.12 tqdm\_for\_rvalues()

Here is the caller graph for this function:



#### 11.1.2.13 trange() [1/2]

Create a tqdm progress bar for a range of integers.

#### **Template Parameters**

IntType	The type of the integer.
,,	,,

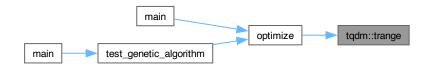
#### **Parameters**

first	The first integer.
last	The last integer.

#### Returns

A tqdm progress bar.

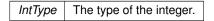
Here is the caller graph for this function:



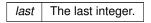
#### 11.1.2.14 trange() [2/2]

Create a tqdm progress bar for a range of integers.

#### **Template Parameters**



#### **Parameters**



#### Returns

A tqdm progress bar.

# 11.2 visualize Namespace Reference

#### **Functions**

• parse\_list (string)

#### **Variables**

- parser
- type
- args = parser.parse\_args()
- x = args.vector
- p = args.perf
- r = args.recovery
- g = args.grade
- graph = graphviz.Digraph()
- rankdir
- nodesep
- ranksep
- splines

- overlap
- dpi
- str feed node = 'Feed'
- str concentrate = 'Concentrate'
- str tailings = 'Tailings'
- shape
- color = 1 else 'red'
- tuple num\_nodes = (len(x) 1) // 3
- str from\_node = f'Unit {i // 3}'
- tuple to\_node = (num\_nodes + 1) else f'Unit {x[i + 1]}'
- cleanup
- True
- format
- view
- dict legend\_labels
- · list legend\_handles
- list columns = ['Feed'] + [f'Unit {i}' for i in range(0, num\_nodes)]
- list reshaped\_list
- df = pd.DataFrame(reshaped\_list).transpose()
- · figsize
- handles
- labels
- loc
- bbox\_to\_anchor
- fontsize
- cellText
- values
- · colLabels
- cellLoc
- bbox
- ha
- va
- transform
- bbox\_inches

#### 11.2.1 Function Documentation

#### 11.2.1.1 parse list()

```
visualize.parse_list (
     string )
```

#### 11.2.2 Variable Documentation

#### 11.2.2.1 args

```
visualize.args = parser.parse_args()
```

#### 11.2.2.2 bbox

visualize.bbox

#### 11.2.2.3 bbox\_inches

visualize.bbox\_inches

#### 11.2.2.4 bbox\_to\_anchor

visualize.bbox\_to\_anchor

#### 11.2.2.5 cellLoc

visualize.cellLoc

#### 11.2.2.6 cellText

visualize.cellText

#### 11.2.2.7 cleanup

visualize.cleanup

#### 11.2.2.8 colLabels

visualize.colLabels

#### 11.2.2.9 color

str visualize.color = 1 else 'red'

#### 11.2.2.10 columns

visualize.columns = ['Feed'] + [f'Unit {i}' for i in range(0, num\_nodes)]

#### 11.2.2.11 concentrate

visualize.concentrate = 'Concentrate'

#### 11.2.2.12 df

visualize.df = pd.DataFrame(reshaped\_list).transpose()

#### 11.2.2.13 dpi

visualize.dpi

#### 11.2.2.14 feed\_node

visualize.feed\_node = 'Feed'

#### 11.2.2.15 figsize

visualize.figsize

#### 11.2.2.16 fontsize

visualize.fontsize

#### 11.2.2.17 format

visualize.format

#### 11.2.2.18 from\_node

visualize.from\_node = f'Unit {i // 3}'

#### 11.2.2.19 g

visualize.g = args.grade

#### 11.2.2.20 graph

visualize.graph = graphviz.Digraph()

#### 11.2.2.21 ha

visualize.ha

#### 11.2.2.22 handles

visualize.handles

#### 11.2.2.23 labels

visualize.labels

#### 11.2.2.24 legend handles

visualize.legend\_handles

#### Initial value:

#### 11.2.2.25 legend\_labels

dict visualize.legend\_labels

#### Initial value:

```
00001 = {'Concentrate': 'blue', 'Intermediate': 'purple', 00002 'Tailings': 'red'}
```

#### 11.2.2.26 loc

visualize.loc

#### 11.2.2.27 nodesep

visualize.nodesep

#### 11.2.2.28 num\_nodes

tuple visualize.num\_nodes = (len(x) - 1) // 3

#### 11.2.2.29 overlap

visualize.overlap

#### 11.2.2.30 p

visualize.p = args.perf

#### 11.2.2.31 parser

visualize.parser

#### Initial value:

```
00001 = argparse.ArgumentParser(description='Command line arguments for \ 00002 visualizing the circuit diagram.')
```

```
11.2.2.32 r
```

visualize.r = args.recovery

#### 11.2.2.33 rankdir

visualize.rankdir

#### 11.2.2.34 ranksep

visualize.ranksep

#### 11.2.2.35 reshaped list

list visualize.reshaped\_list

#### Initial value:

#### 11.2.2.36 shape

visualize.shape

#### 11.2.2.37 splines

visualize.splines

#### 11.2.2.38 tailings

visualize.tailings = 'Tailings'

#### 11.2.2.39 to\_node

```
visualize.to_node = (num_nodes + 1) else f'Unit \{x[i + 1]\}'
```

#### 11.2.2.40 transform

 $\verb|visualize.transform||\\$ 

#### 11.2.2.41 True

visualize.True

#### 11.2.2.42 type

visualize.type

#### 11.2.2.43 va

visualize.va

#### 11.2.2.44 values

visualize.values

#### 11.2.2.45 view

visualize.view

#### 11.2.2.46 x

visualize.x = args.vector

# **Chapter 12**

# **Data Structure Documentation**

## 12.1 Algorithm Parameters Struct Reference

Configures the parameters of a genetic algorithm.

#include <Parameter.h>

Collaboration diagram for Algorithm\_Parameters:

#### Algorithm\_Parameters

- + maxGenerations
- + populationSize
- + selection
- + tournamentSize
- + crossover
- + crossoverRate
- + mutation
- + mutationRate
- + elitePercentage
- + initialTemp
- + deltT
- + randomSeed
- + Algorithm\_Parameters()

#### **Public Member Functions**

 Algorithm\_Parameters (int gens, int popSize, std::string selection, int tourSize, std::string crossover, double crossRate, std::string mutation, double mutRate, double elitePerc, double initialTemp, double deltT, double randomSeed)

Constructor to initialize the parameters with custom values.

#### **Data Fields**

- int maxGenerations = 1000
- int populationSize = 100

- std::string selection = "NaryTournamentSelection"
- int tournamentSize = 3
- std::string crossover = "TwoPointCrossover"
- double crossoverRate = 0.8
- std::string mutation = "GuidedMutation"
- double mutationRate = 0.1
- double elitePercentage = 0.2
- double initialTemp = 1000.0
- double deltT = 1.0
- unsigned int randomSeed = 0

#### 12.1.1 Detailed Description

Configures the parameters of a genetic algorithm.

The Algorithm\_Parameters structure contains various parameters used to configure the behavior of a genetic algorithm, including the number of generations, population size, tournament size, crossover rate, mutation rate, elite percentage, and random seed.

#### 12.1.2 Constructor & Destructor Documentation

#### 12.1.2.1 Algorithm Parameters()

```
Algorithm_Parameters::Algorithm_Parameters (
    int gens,
    int popSize,
    std::string selection,
    int tourSize,
    std::string crossover,
    double crossRate,
    std::string mutation,
    double mutRate,
    double elitePerc,
    double initialTemp,
    double deltT,
    double randomSeed) [inline]
```

Constructor to initialize the parameters with custom values.

#### **Parameters**

gens	Maximum number of generations.
popSize	Size of the population.
tourSize	Number of individuals to select in each tournament.
crossRate	Rate at which crossover occurs.
mutRate	Rate at which mutation occurs.
elitePerc	Percentage of elite individuals to keep in the next generation.
seed	Random seed, 0 means using current time.

#### 12.1.3 Field Documentation

#### 12.1.3.1 crossover

```
std::string Algorithm_Parameters::crossover = "TwoPointCrossover"
Crossover method
```

#### 12.1.3.2 crossoverRate

double Algorithm\_Parameters::crossoverRate = 0.8
Rate at which crossover occurs

#### 12.1.3.3 deltT

double Algorithm\_Parameters::deltT = 1.0

The temperature decrement of Simulated Annealing algorithm

#### 12.1.3.4 elitePercentage

double Algorithm\_Parameters::elitePercentage = 0.2
Percentage of elite individuals to keep in the next generation

#### 12.1.3.5 initialTemp

double Algorithm\_Parameters::initialTemp = 1000.0
Initial temperature of Simulated Annealing algorithm

#### 12.1.3.6 maxGenerations

int Algorithm\_Parameters::maxGenerations = 1000
Maximum number of generations

#### 12.1.3.7 mutation

std::string Algorithm\_Parameters::mutation = "GuidedMutation"
Mutation method

#### 12.1.3.8 mutationRate

double Algorithm\_Parameters::mutationRate = 0.1
Rate at which mutation occurs

#### 12.1.3.9 populationSize

int Algorithm\_Parameters::populationSize = 100
Size of the population

#### 12.1.3.10 randomSeed

unsigned int Algorithm\_Parameters::randomSeed = 0
Random seed, 0 means using current time

#### 12.1.3.11 selection

std::string Algorithm\_Parameters::selection = "NaryTournamentSelection"
Selection method

#### 12.1.3.12 tournamentSize

int Algorithm\_Parameters::tournamentSize = 3
Number of individuals to select in each tournament
The documentation for this struct was generated from the following file:

· include/utils/Parameter.h

# 12.2 tqdm::Chronometer Class Reference

A simple chronometer to measure elapsed time.

#include <tqdm.hpp>

Collaboration diagram for tqdm::Chronometer:

# tqdm::Chronometer

- start\_
- + Chronometer()
- + reset()
- + peek()
- + get\_start()

#### **Public Member Functions**

· Chronometer ()

Construct a new Chronometer object.

• double reset ()

Reset the chronometer and return the elapsed time since the last reset.

• double peek () const

Get the elapsed time without resetting the chronometer.

time\_point\_t get\_start () const

Get the starting time point.

#### **Private Attributes**

time\_point\_t start\_

#### 12.2.1 Detailed Description

A simple chronometer to measure elapsed time.

#### 12.2.2 Constructor & Destructor Documentation

#### 12.2.2.1 Chronometer()

```
tqdm::Chronometer::Chronometer ( ) [inline]
Construct a new Chronometer object.
```

#### 12.2.3 Member Function Documentation

#### 12.2.3.1 get\_start()

```
time_point_t tqdm::Chronometer::get_start ( ) const [inline]
Get the starting time point.
```

#### Returns

The starting time point.

#### 12.2.3.2 peek()

double tqdm::Chronometer::peek ( ) const [inline]

Get the elapsed time without resetting the chronometer.

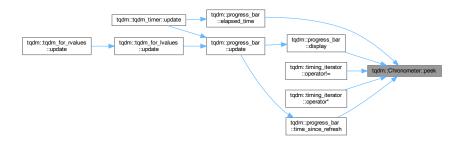
#### Returns

The elapsed time in seconds.

Here is the call graph for this function:



Here is the caller graph for this function:



#### 12.2.3.3 reset()

double tqdm::Chronometer::reset ( ) [inline]

Reset the chronometer and return the elapsed time since the last reset.

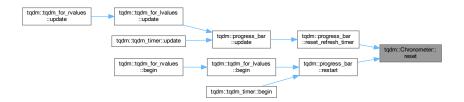
#### Returns

The elapsed time in seconds.

Here is the call graph for this function:



Here is the caller graph for this function:



#### 12.2.4 Field Documentation

#### 12.2.4.1 start\_

time\_point\_t tqdm::Chronometer::start\_ [private]

The documentation for this class was generated from the following file:

• include/utils/tqdm.hpp

#### 12.3 Circuit Class Reference

#include <CCircuit.h>

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#### Collaboration diagram for Circuit:

# Circuit + concentrateG + concentrateW + tailingsG + tailingsW + wasteFeed + gerardiumFeed - units - numUnits - feed - check + Circuit() + ~Circuit() + print\_info() + connected() + initialize\_feed\_rates() + mark\_units() + check\_convergence() + calculate\_flows() + Check\_Validity()

- reset\_new\_feeds()

#### **Public Member Functions**

Circuit (int num\_units)

Constructs a Circuit object with a specified number of units.

• ∼Circuit ()

Destructs the Circuit object.

· void print\_info () const

Prints information about the circuit.

void connected (int \*circuitVector)

Connects the units in the circuit based on the circuit vector.

• void initialize\_feed\_rates (double gerardium\_=10, double waste\_=90)

Initializes the feed rates for gerardium and waste.

void mark\_units (int unitNum, std::vector< CUnit \* > &units, bool check[3])

Marks a unit in the circuit and recursively marks its connected units.

bool check\_convergence (double threshold)

Checks if the circuit has converged based on a threshold value.

void calculate\_flows ()

Calculates the flows within the circuit.

#### **Static Public Member Functions**

static bool Check\_Validity (int vectorSize, int \*circuitVector)
 Checks the validity of the circuit vector.

#### **Data Fields**

- · double concentrateG
- · double concentrateW
- double tailingsG
- double tailingsW
- · double wasteFeed
- double gerardiumFeed

#### **Private Member Functions**

• void reset new feeds ()

Resets the new feeds in the circuit.

#### **Private Attributes**

- std::vector< CUnit \* > units
- · int numUnits
- · int feed

#### **Static Private Attributes**

• static bool check [3]

#### 12.3.1 Constructor & Destructor Documentation

#### 12.3.1.1 Circuit()

```
Circuit::Circuit (
          int num_units )
```

Constructs a Circuit object with a specified number of units.

#### **Parameters**

num units	The number of units in the circuit.

#### 12.3.1.2 $\sim$ Circuit()

```
Circuit::~Circuit ()

Destructs the Circuit object.
```

#### 12.3.2 Member Function Documentation

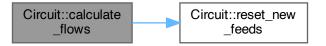
#### 12.3.2.1 calculate\_flows()

```
void Circuit::calculate_flows ( )
```

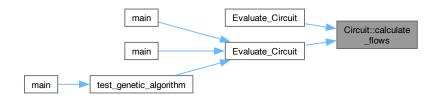
Calculates the flows within the circuit.

This member function calculates the flow rates of gerardium and waste through the circuit's units. Here is the call

graph for this function:



Here is the caller graph for this function:



#### 12.3.2.2 check\_convergence()

Checks if the circuit has converged based on a threshold value.

This member function checks whether the circuit's calculations have converged by comparing the changes in feed rates to a specified threshold.

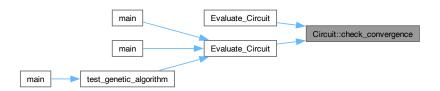
#### **Parameters**

threshold	The convergence threshold.

#### Returns

True if the circuit has converged, otherwise false.

Here is the caller graph for this function:



#### 12.3.2.3 Check\_Validity()

Checks the validity of the circuit vector.

This static member function checks whether a given circuit vector is valid. It verifies the connections and configurations of the units in the circuit.

#### **Parameters**

vectorSize	The size of the circuit vector.
circuitVector	A pointer to the circuit vector.

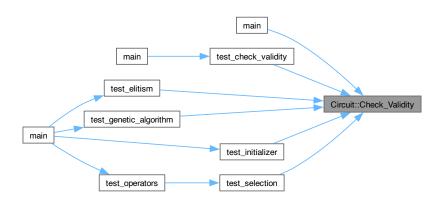
#### Returns

True if the circuit vector is valid, otherwise false.

Here is the call graph for this function:



Here is the caller graph for this function:



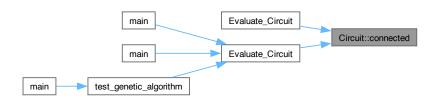
#### 12.3.2.4 connected()

Connects the units in the circuit based on the circuit vector.

This member function establishes connections between the units in the circuit according to the specified circuit vector.

#### **Parameters**

Here is the caller graph for this function:



#### 12.3.2.5 initialize\_feed\_rates()

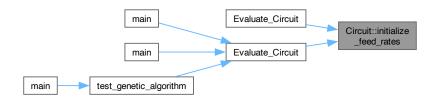
Initializes the feed rates for gerardium and waste.

This member function initializes the feed rates for gerardium and waste to the specified values.

#### **Parameters**

gerardium← _	The feed rate of gerardium (default is 10).
waste_	The feed rate of waste (default is 90).

Here is the caller graph for this function:



#### 12.3.2.6 mark\_units()

```
void Circuit::mark_units (
    int unitNum,
    std::vector< CUnit * > & units,
    bool check[3] )
```

Marks a unit in the circuit and recursively marks its connected units.

This function marks a specified unit in the circuit as visited. It then recursively marks all units connected to this unit through its concNum, interNum, and tailsNum connections. If a connection points to an outlet, it updates the corresponding check flag to indicate an exit has been seen.

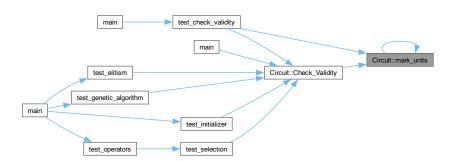
#### **Parameters**

unit_num	The index of the unit to be marked.
units	A vector of pointers to CUnit objects representing the units in the circuit.
check	An array of three booleans that track if the concNum, interNum, or tailsNum connections lead to a circuit outlet.

Here is the call graph for this function:



Here is the caller graph for this function:



#### 12.3.2.7 print\_info()

void Circuit::print\_info ( ) const

Prints information about the circuit.

This member function prints detailed information about the circuit, including the number of units, feed rates, and concentrations.

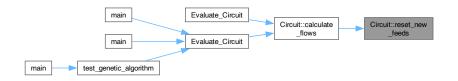
#### 12.3.2.8 reset\_new\_feeds()

void Circuit::reset\_new\_feeds ( ) [private]

Resets the new feeds in the circuit.

This helper function resets the new feed rates in the circuit to their initial values. Here is the caller graph for this

function:



#### 12.3.3 Field Documentation

#### 12.3.3.1 check

bool Circuit::check[3] [static], [private]
Static array to check the units

#### 12.3.3.2 concentrateG

double Circuit::concentrateG
Concentration of gerardium in the circuit

#### 12.3.3.3 concentrateW

double Circuit::concentrateW
Concentration of waste in the circuit

#### 12.3.3.4 feed

int Circuit::feed [private]
The feed rate for the circuit

#### 12.3.3.5 gerardiumFeed

double Circuit::gerardiumFeed
Feed rate of gerardium in the circuit

#### 12.3.3.6 numUnits

int Circuit::numUnits [private]
The number of units in the circuit

#### 12.3.3.7 tailingsG

double Circuit::tailingsG

Tailings of gerardium in the circuit

#### 12.3.3.8 tailingsW

 $\begin{tabular}{ll} \begin{tabular}{ll} \beg$ 

#### 12.3.3.9 units

std::vector<CUnit \*> Circuit::units [private]
Vector of pointers to the units in the circuit

#### 12.3.3.10 wasteFeed

double Circuit::wasteFeed

Feed rate of waste in the circuit

The documentation for this class was generated from the following files:

- · include/circuit/CCircuit.h
- investigate/CCircuit.cpp
- src/circuit/CCircuit.cpp

#### 12.4 CircuitParameters Struct Reference

Defines the parameters for the circuit simulator.

#include <Parameter.h>

Collaboration diagram for CircuitParameters:

# CircuitParameters + tolerance + maxIterations + CircuitParameters()

#### **Public Member Functions**

• CircuitParameters (double tol, int maxIter)

Constructor to initialize the parameters with custom values.

#### **Data Fields**

- double tolerance = 1e-6
- int maxIterations = 1000

#### 12.4.1 Detailed Description

Defines the parameters for the circuit simulator.

The CircuitParameters structure contains various parameters used for simulating the circuit, such as tolerance and maximum iterations.

#### 12.4.2 Constructor & Destructor Documentation

#### 12.4.2.1 CircuitParameters()

Constructor to initialize the parameters with custom values.

#### **Parameters**

tol	The tolerance level for the simulation.
maxIter	The maximum number of iterations for the simulation.

12.5 CUnit Class Reference 71

#### 12.4.3 Field Documentation

#### 12.4.3.1 maxIterations

int CircuitParameters::maxIterations = 1000

The maximum number of iterations for the simulation.

#### 12.4.3.2 tolerance

double CircuitParameters::tolerance = 1e-6

The tolerance level for the simulation.

The documentation for this struct was generated from the following file:

• include/utils/Parameter.h

#### 12.5 CUnit Class Reference

#include <CUnit.h>

Collaboration diagram for CUnit:

## **CUnit** + concNum + interNum + tailsNum - mark - feedGerardium - feedWaste - newFeedGerardium - newFeedWaste - differenceG -concPtr - differenceW -interPtr + CUnit() -tailsPtr + CUnit() + isMarked() + setMarked() + getFeedGerardium() + setFeedGerardium() + getFeedWaste() + setFeedWaste() + getNewFeedGerardium() + setNewFeedGerardium() and 11 more...

#### **Public Member Functions**

• CUnit ()

Default constructor for CUnit.

· CUnit (int concNum, int interNum, int tailsNum)

Parameterized constructor for CUnit.

• bool isMarked () const

Checks if the unit is marked.

void setMarked (bool mark)

Sets the mark status of the unit.

• double getFeedGerardium () const

Gets the feed rate of gerardium.

· void setFeedGerardium (double feedGerardium)

Sets the feed rate of gerardium.

• double getFeedWaste () const

Gets the feed rate of waste.

void setFeedWaste (double feedWaste)

Sets the feed rate of waste.

double getNewFeedGerardium () const

Gets the new feed rate of gerardium.

void setNewFeedGerardium (double newFeedGerardium)

Sets the new feed rate of gerardium.

· double getNewFeedWaste () const

Gets the new feed rate of waste.

void setNewFeedWaste (double newFeedWaste)

Sets the new feed rate of waste.

• double getDifferenceG () const

Gets the difference in feed rate of gerardium.

• double getDifferenceW () const

Gets the difference in feed rate of waste.

• void setDifference ()

Sets the differences in feed rates for gerardium and waste.

CUnit \* getConcPtr () const

Gets the pointer to the concentrate connection.

void setConcPtr (CUnit \*ptr)

Sets the pointer to the concentrate connection.

• CUnit \* getInterPtr () const

Gets the pointer to the intermediate connection.

void setInterPtr (CUnit \*ptr)

Sets the pointer to the intermediate connection.

CUnit \* getTailsPtr () const

Gets the pointer to the tailings connection.

void setTailsPtr (CUnit \*ptr)

Sets the pointer to the tailings connection.

#### **Data Fields**

- · int concNum
- int interNum
- · int tailsNum

12.5 CUnit Class Reference 73

#### **Private Attributes**

- bool mark
- · double feedGerardium
- · double feedWaste
- · double newFeedGerardium
- double newFeedWaste
- double differenceG
- double differenceW
- CUnit \* concPtr
- CUnit \* interPtr
- · CUnit \* tailsPtr

#### 12.5.1 Constructor & Destructor Documentation

#### 12.5.1.1 CUnit() [1/2]

```
CUnit::CUnit ()
```

Default constructor for CUnit.

Constructs a CUnit object with default values for concentrate, intermediate, and tailings numbers.

#### 12.5.1.2 CUnit() [2/2]

```
CUnit::CUnit (
    int concNum,
    int interNum,
    int tailsNum )
```

Parameterized constructor for CUnit.

Constructs a CUnit object with specified values for concentrate, intermediate, and tailings numbers.

#### **Parameters**

concNum	The concentrate number.
interNum	The intermediate number.
tailsNum	The tailings number.

#### 12.5.2 Member Function Documentation

#### 12.5.2.1 getConcPtr()

```
CUnit * CUnit::getConcPtr ( ) const
```

Gets the pointer to the concentrate connection.

#### Returns

The pointer to the concentrate connection.

#### 12.5.2.2 getDifferenceG()

```
double CUnit::getDifferenceG ( ) const
```

Gets the difference in feed rate of gerardium.

#### Returns

The difference in feed rate of gerardium.

#### 12.5.2.3 getDifferenceW()

double CUnit::getDifferenceW ( ) const
Gets the difference in feed rate of waste.

Returns

The difference in feed rate of waste.

#### 12.5.2.4 getFeedGerardium()

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Returns

The feed rate of gerardium.

#### 12.5.2.5 getFeedWaste()

double CUnit::getFeedWaste ( ) const
Gets the feed rate of waste.

Returns

The feed rate of waste.

#### 12.5.2.6 getInterPtr()

 $\begin{array}{lll} & \texttt{CUnit} * & \texttt{CUnit::getInterPtr} & ( \ ) & \texttt{const} \\ & \textbf{Gets the pointer to the intermediate connection.} \end{array}$ 

Returns

The pointer to the intermediate connection.

#### 12.5.2.7 getNewFeedGerardium()

 $\begin{tabular}{ll} \begin{tabular}{ll} \beg$ 

Returns

The new feed rate of gerardium.

#### 12.5.2.8 getNewFeedWaste()

double CUnit::getNewFeedWaste ( ) const
Gets the new feed rate of waste.

Returns

The new feed rate of waste.

#### 12.5.2.9 getTailsPtr()

CUnit \* CUnit::getTailsPtr ( ) const
Gets the pointer to the tailings connection.

Returns

The pointer to the tailings connection.

12.5 CUnit Class Reference 75

#### 12.5.2.10 isMarked()

```
bool CUnit::isMarked ( ) const Checks if the unit is marked.
```

Returns

True if the unit is marked, otherwise false.

#### 12.5.2.11 setConcPtr()

Sets the pointer to the concentrate connection.

#### **Parameters**

*ptr* The pointer to the concentrate connection to set.

#### 12.5.2.12 setDifference()

```
void CUnit::setDifference ( )
```

Sets the differences in feed rates for gerardium and waste.

#### 12.5.2.13 setFeedGerardium()

Sets the feed rate of gerardium.

#### **Parameters**

feedGerardium	The feed rate of gerardium to set.

#### 12.5.2.14 setFeedWaste()

Sets the feed rate of waste.

#### **Parameters**

feedWaste	The feed rate of waste to set.
ICCUVVASIC	The leed rate of waste to set.

#### 12.5.2.15 setInterPtr()

Sets the pointer to the intermediate connection.

#### **Parameters**

*ptr* The pointer to the intermediate connection to set.

#### 12.5.2.16 setMarked()

Sets the mark status of the unit.

#### **Parameters**

mark The mark status to set.

#### 12.5.2.17 setNewFeedGerardium()

Sets the new feed rate of gerardium.

#### **Parameters**

#### 12.5.2.18 setNewFeedWaste()

Sets the new feed rate of waste.

#### **Parameters**

newFeedWaste	The new feed rate of waste to set.
HEWI CCUVVASIC	I THE HEW ICCUTATE OF WASTE TO SET.

#### 12.5.2.19 setTailsPtr()

Sets the pointer to the tailings connection.

#### **Parameters**

*ptr* The pointer to the tailings connection to set.

#### 12.5.3 Field Documentation

#### 12.5.3.1 concNum

int CUnit::concNum
The concentrate number

#### 12.5.3.2 concPtr

CUnit\* CUnit::concPtr [private]
Pointer to the concentrate connection

#### 12.5.3.3 differenceG

double CUnit::differenceG [private]

Difference in feed rate of gerardium

#### 12.5.3.4 differenceW

double CUnit::differenceW [private]
Difference in feed rate of waste

#### 12.5.3.5 feedGerardium

double CUnit::feedGerardium [private]
Feed rate of gerardium

#### 12.5.3.6 feedWaste

double CUnit::feedWaste [private]
Feed rate of waste

#### 12.5.3.7 interNum

int CUnit::interNum
The intermediate number

#### 12.5.3.8 interPtr

CUnit\* CUnit::interPtr [private]
Pointer to the intermediate connection

#### 12.5.3.9 mark

bool CUnit::mark [private]
Mark status of the unit

#### 12.5.3.10 newFeedGerardium

double CUnit::newFeedGerardium [private]
New feed rate of gerardium

#### 12.5.3.11 newFeedWaste

double CUnit::newFeedWaste [private]
New feed rate of waste

#### 12.5.3.12 tailsNum

int CUnit::tailsNum
The tailings number

#### 12.5.3.13 tailsPtr

CUnit\* CUnit::tailsPtr [private]

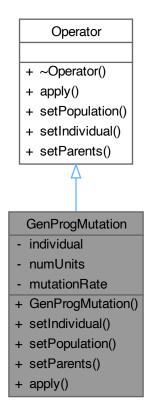
Pointer to the tailings connection

The documentation for this class was generated from the following files:

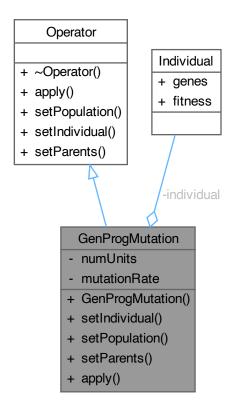
- include/circuit/CUnit.h
- src/circuit/CUnit.cpp

# 12.6 GenProgMutation Class Reference

#include <GenProgMutation.h>
Inheritance diagram for GenProgMutation:



Collaboration diagram for GenProgMutation:



#### **Public Member Functions**

- GenProgMutation (int vectorSize, const Algorithm\_Parameters &params)
   Constructs a GenProgMutation object with specified vector size and algorithm parameters.
- · void setIndividual (Individual &individual) override

Sets the individual to be mutated.

- void setPopulation (const std::vector< Individual > &population, std::vector< Individual > &selected)

  Sets the population and selected vectors for the mutation operation.
- void setParents (Individual &offspring, const Individual &parent1, const Individual &parent2)

  Sets the parents and offspring for the mutation operation.
- void apply (std::mt19937 &generator)

Performs the mutation operation on the individual.

#### **Public Member Functions inherited from Operator**

virtual ~Operator ()=default
 Virtual destructor for the Operator class.

#### **Private Attributes**

- Individual \* individual
- int numUnits
- · double mutationRate

#### 12.6.1 Constructor & Destructor Documentation

#### 12.6.1.1 GenProgMutation()

Constructs a GenProgMutation object with specified vector size and algorithm parameters.

#### Parameters 4 8 1

vectorSize	The size of the vector representing the individual's genetic material.
params	The algorithm parameters including the mutation rate.

#### 12.6.2 Member Function Documentation

#### 12.6.2.1 apply()

Performs the mutation operation on the individual.

This method performs mutation on the individual's genetic material based on the specified mutation rate. The mutation modifies the genes of the individual to introduce variations.

#### **Parameters**

Implements Operator.

#### 12.6.2.2 setIndividual()

Sets the individual to be mutated.

#### **Parameters**

individual The individual to be mutated	d.
---	----

Implements Operator.

#### 12.6.2.3 setParents()

Sets the parents and offspring for the mutation operation.

This method is not used in the mutation operation.

#### Parameters

offspring	The offspring individual to be generated.
parent1	The first parent individual.
parent2	The second parent individual.

Implements Operator.

#### 12.6.2.4 setPopulation()

Sets the population and selected vectors for the mutation operation.

This method is not used in the mutation operation.

#### **Parameters**

population	The vector of individuals in the population.
selected	The vector of selected individuals.

Implements Operator.

#### 12.6.3 Field Documentation

#### 12.6.3.1 individual

```
Individual* GenProgMutation::individual [private]
Pointer to the individual to be mutated
```

#### 12.6.3.2 mutationRate

```
double GenProgMutation::mutationRate [private]
The mutation rate used to determine if mutation should occur
```

#### 12.6.3.3 numUnits

```
int GenProgMutation::numUnits [private]
```

Represents the number of units or types of operations/nodes

The documentation for this class was generated from the following files:

- include/operators/Mutation/GenProgMutation.h
- src/operators/Mutation/GenProgMutation.cpp

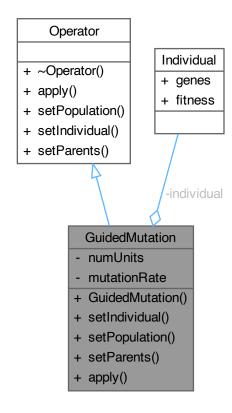
#### 12.7 GuidedMutation Class Reference

#include <GuidedMutation.h>

Inheritance diagram for GuidedMutation:

# Operator + ~Operator() + apply() + setPopulation() + setIndividual() + setParents() GuidedMutation - individual - numUnits - mutationRate + GuidedMutation() + setIndividual() + setPopulation() + setParents() + apply()

Collaboration diagram for GuidedMutation:



#### **Public Member Functions**

- GuidedMutation (int vectorSize, const Algorithm\_Parameters &params)
  - Constructs a GuidedMutation object with specified vector size and algorithm parameters.
- · void setIndividual (Individual &individual) override

Sets the individual to be mutated.

- void setPopulation (const std::vector< Individual > &population, std::vector< Individual > &selected)

  Performs the guided mutation operation on the individual.
- void setParents (Individual &offspring, const Individual &parent1, const Individual &parent2)

  Sets the parents and offspring for the crossover operation.
- · void apply (std::mt19937 &generator) override

Performs the guided mutation operation on the individual.

#### **Public Member Functions inherited from Operator**

virtual ~Operator ()=default
 Virtual destructor for the Operator class.

#### **Private Attributes**

- Individual \* individual
- int numUnits
- · double mutationRate

#### 12.7.1 Constructor & Destructor Documentation

#### 12.7.1.1 GuidedMutation()

Constructs a GuidedMutation object with specified vector size and algorithm parameters.

#### Parameters 4 8 1

vectorSize	The size of the vector representing the individual's genetic material.
params	The algorithm parameters including the mutation rate.

#### 12.7.2 Member Function Documentation

#### 12.7.2.1 apply()

Performs the guided mutation operation on the individual.

This method performs mutation on the individual's genetic material based on the specified mutation rate. The mutation modifies the genes of the individual to introduce variations according to specific rules or heuristics.

#### **Parameters**

Implements Operator.

#### 12.7.2.2 setIndividual()

Sets the individual to be mutated.

#### **Parameters**

individual	The individual to be mutated.
maividadi	The marriada to be matated.

Implements Operator.

#### 12.7.2.3 setParents()

Sets the parents and offspring for the crossover operation.

This method is not used by the GuidedMutation operator.

#### Parameters

offspring	The offspring individual to be generated.
parent1	The first parent individual.
parent2	The second parent individual.

Implements Operator.

#### 12.7.2.4 setPopulation()

Performs the guided mutation operation on the individual.

This method performs mutation on the individual's genetic material based on the specified mutation rate. The mutation modifies the genes of the individual to introduce variations according to specific rules or heuristics.

#### **Parameters**

generator The random number generator used to determine if mutation should on	ccur.
---	-------

Implements Operator.

#### 12.7.3 Field Documentation

#### 12.7.3.1 individual

```
Individual* GuidedMutation::individual [private]
Pointer to the individual to be mutated
```

#### 12.7.3.2 mutationRate

```
double GuidedMutation::mutationRate [private]
The mutation rate used to determine if mutation should occur
```

#### 12.7.3.3 numUnits

```
int GuidedMutation::numUnits [private]
```

Represents the number of units or types of operations/nodes

The documentation for this class was generated from the following files:

- include/operators/Mutation/GuidedMutation.h
- src/operators/Mutation/GuidedMutation.cpp

#### 12.8 Individual Struct Reference

Represents an individual in a genetic algorithm.

#include <Individual.h>

Collaboration diagram for Individual:



#### **Data Fields**

- std::vector< int > genes
- · double fitness

#### 12.8.1 Detailed Description

Represents an individual in a genetic algorithm.

The Individual structure contains a vector of genes that represent the circuit configuration and a fitness value that indicates the quality or performance of the circuit configuration.

#### 12.8.2 Field Documentation

#### 12.8.2.1 fitness

double Individual::fitness

Stores the fitness of the circuit configuration as a double

#### 12.8.2.2 genes

std::vector<int> Individual::genes

Stores the circuit configuration as a vector of integers

The documentation for this struct was generated from the following file:

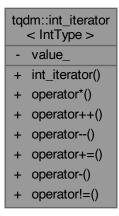
• include/utils/Individual.h

# 12.9 tqdm::int\_iterator< IntType > Class Template Reference

A random access iterator for integers.

#include <tqdm.hpp>

Collaboration diagram for tqdm::int\_iterator< IntType >:



#### **Public Types**

- using iterator\_category = std::random\_access\_iterator\_tag
- using value\_type = IntType
- using difference\_type = IntType
- using pointer = IntType \*

• using reference = IntType &

#### **Public Member Functions**

- int\_iterator (IntType val)
- IntType & operator\* ()
- int\_iterator & operator++ ()
- int iterator & operator-- ()
- int\_iterator & operator+= (difference\_type d)
- difference\_type operator- (const int\_iterator &other) const
- bool operator!= (const int\_iterator &other) const

#### **Private Attributes**

IntType value

#### 12.9.1 Detailed Description

```
template < class IntType > class tqdm::int_iterator < IntType >
```

A random access iterator for integers.

#### **Template Parameters**

```
IntType The type of the integer.
```

#### 12.9.2 Member Typedef Documentation

#### 12.9.2.1 difference\_type

```
template<class IntType >
using tqdm::int_iterator< IntType >::difference_type = IntType
```

#### 12.9.2.2 iterator\_category

```
template<class IntType >
using tqdm::int_iterator< IntType >::iterator_category = std::random_access_iterator_tag
```

#### 12.9.2.3 pointer

```
template<class IntType >
using tqdm::int_iterator< IntType >::pointer = IntType *
```

#### 12.9.2.4 reference

```
template<class IntType >
using tqdm::int_iterator< IntType >::reference = IntType &
```

#### 12.9.2.5 value\_type

```
template<class IntType >
using tqdm::int_iterator< IntType >::value_type = IntType
```

#### 12.9.3 Constructor & Destructor Documentation

#### 12.9.3.1 int\_iterator()

#### 12.9.4 Member Function Documentation

#### 12.9.4.1 operator"!=()

#### 12.9.4.2 operator\*()

```
template<class IntType >
IntType & tqdm::int_iterator< IntType >::operator* ( ) [inline]
```

#### 12.9.4.3 operator++()

```
template<class IntType >
int_iterator & tqdm::int_iterator< IntType >::operator++ ( ) [inline]
```

#### 12.9.4.4 operator+=()

#### 12.9.4.5 operator-()

#### 12.9.4.6 operator--()

```
template<class IntType >
int_iterator & tqdm::int_iterator< IntType >::operator-- ( ) [inline]
```

#### 12.9.5 Field Documentation

#### 12.9.5.1 value\_

```
template < class IntType >
IntType tqdm::int_iterator < IntType >::value_ [private]
```

The documentation for this class was generated from the following file:

include/utils/tqdm.hpp

### 12.10 tqdm::iter\_wrapper< ForwardIter, Parent > Class Template Reference

A wrapper class for iterators to update the progress bar.

```
#include <tqdm.hpp>
```

Collaboration diagram for tqdm::iter\_wrapper< ForwardIter, Parent >:

# tqdm::iter\_wrapper < ForwardIter, Parent > - Parent - current\_ - parent\_ + iter\_wrapper() + operator\*() + operator++() + operator!=() + operator!=() + get()

#### **Public Types**

- using iterator\_category = typename ForwardIter::iterator\_category
- using value\_type = typename ForwardIter::value\_type
- using difference\_type = typename ForwardIter::difference\_type
- using pointer = typename ForwardIter::pointer
- using reference = typename ForwardIter::reference

#### **Public Member Functions**

- iter\_wrapper (ForwardIter it, Parent \*parent)
- auto operator\* ()
- void operator++ ()
- template < class Other >
  bool operator!= (const Other & other) const
- bool operator!= (const iter\_wrapper &other) const
- const ForwardIter & get () const

#### **Private Attributes**

- friend Parent
- ForwardIter current\_
- Parent \* parent\_

#### 12.10.1 Detailed Description

template < class ForwardIter, class Parent > class tqdm::iter\_wrapper < ForwardIter, Parent >

A wrapper class for iterators to update the progress bar.

#### **Template Parameters**

ForwardIter	The type of the underlying iterator.
Parent	The type of the parent progress bar.

#### 12.10.2 Member Typedef Documentation

#### 12.10.2.1 difference type

```
template<class ForwardIter , class Parent >
using tqdm::iter_wrapper< ForwardIter, Parent >::difference_type = typename ForwardIter←
::difference_type
```

#### 12.10.2.2 iterator\_category

```
template<class ForwardIter , class Parent >
using tqdm::iter_wrapper< ForwardIter, Parent >::iterator_category = typename ForwardIter←
::iterator_category
```

#### 12.10.2.3 pointer

```
template<class ForwardIter , class Parent >
using tqdm::iter_wrapper< ForwardIter, Parent >::pointer = typename ForwardIter::pointer
```

#### 12.10.2.4 reference

```
template<class ForwardIter , class Parent >
using tqdm::iter_wrapper< ForwardIter, Parent >::reference = typename ForwardIter::reference
```

#### 12.10.2.5 value\_type

```
template<class ForwardIter , class Parent >
using tqdm::iter_wrapper< ForwardIter, Parent >::value_type = typename ForwardIter::value_type
```

#### 12.10.3 Constructor & Destructor Documentation

#### 12.10.3.1 iter\_wrapper()

#### 12.10.4 Member Function Documentation

#### 12.10.4.1 get()

```
template<class ForwardIter , class Parent >
const ForwardIter & tqdm::iter_wrapper< ForwardIter, Parent >::get () const [inline]
```

#### 12.10.4.2 operator"!=() [1/2]

#### 12.10.4.3 operator"!=() [2/2]

#### 12.10.4.4 operator\*()

```
template < class ForwardIter , class Parent >
auto tqdm::iter_wrapper < ForwardIter, Parent >::operator* ( ) [inline]

12.10.4.5 operator++()

template < class ForwardIter , class Parent >
void tqdm::iter_wrapper < ForwardIter, Parent >::operator++ ( ) [inline]

12.10.5 Field Documentation

12.10.5.1 current_
template < class ForwardIter , class Parent >
ForwardIter tqdm::iter_wrapper < ForwardIter, Parent >::current_ [private]

12.10.5.2 Parent
template < class ForwardIter , class Parent >
friend tqdm::iter_wrapper < ForwardIter, Parent >::Parent [private]

12.10.5.3 parent_
template < class ForwardIter , class Parent >
```

• include/utils/tqdm.hpp

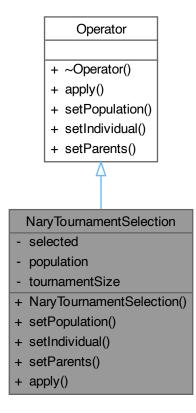
#### 12.11 NaryTournamentSelection Class Reference

The documentation for this class was generated from the following file:

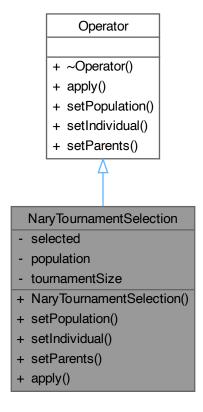
Parent\* tqdm::iter\_wrapper< ForwardIter, Parent >::parent\_ [private]

#include <NaryTournamentSelection.h>

Inheritance diagram for NaryTournamentSelection:



Collaboration diagram for NaryTournamentSelection:



#### **Public Member Functions**

• NaryTournamentSelection (const Algorithm\_Parameters &params)

Constructs an NaryTournamentSelection object with specified algorithm parameters.

 void setPopulation (const std::vector < Individual > &population, std::vector < Individual > &selected) override

Sets the population and selected vectors for the selection operation.

· void setIndividual (Individual &individual)

Sets the individual to be mutated.

• void setParents (Individual &offspring, const Individual &parent1, const Individual &parent2)

Sets the parents and offspring for the crossover operation.

void apply (std::mt19937 &generator) override

Performs the n-ary tournament selection operation.

#### Public Member Functions inherited from Operator

virtual ∼Operator ()=default

Virtual destructor for the Operator class.

#### **Private Attributes**

• std::vector< Individual > \* selected

- const std::vector< Individual > \* population
- · int tournamentSize

#### 12.11.1 Constructor & Destructor Documentation

#### 12.11.1.1 NaryTournamentSelection()

Constructs an NaryTournamentSelection object with specified algorithm parameters.

#### **Parameters**

params The algorithm parameters including the tournament si	ze.
---	-----

#### 12.11.2 Member Function Documentation

#### 12.11.2.1 apply()

Performs the n-ary tournament selection operation.

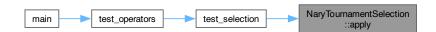
This method selects individuals from the population based on n-ary tournament competition and stores the selected individuals in the provided vector.

#### **Parameters**

	generator	The random number generator used for selecting individuals.	
--	-----------	---	--

Implements Operator.

Here is the caller graph for this function:



#### 12.11.2.2 setIndividual()

Sets the individual to be mutated.

This method sets the individual to be mutated by the operator.

#### **Parameters**

individual	The individual to be mutated.

Implements Operator.

#### 12.11.2.3 setParents()

void NaryTournamentSelection::setParents (

```
Individual & offspring,
const Individual & parent1,
const Individual & parent2 ) [inline], [virtual]
```

Sets the parents and offspring for the crossover operation.

This method is not used by the NaryTournamentSelection operator.

#### **Parameters**

offspring	The offspring individual to be generated.
parent1	The first parent individual.
parent2	The second parent individual.

Implements Operator.

#### 12.11.2.4 setPopulation()

Sets the population and selected vectors for the selection operation.

#### **Parameters**

population	The vector of individuals in the population.
selected	The vector of selected individuals.

Implements Operator.

Here is the caller graph for this function:



#### 12.11.3 Field Documentation

#### 12.11.3.1 population

const std::vector<Individual>\* NaryTournamentSelection::population [private]
Pointer to the vector of individuals in the population

#### 12.11.3.2 selected

std::vector<Individual>\* NaryTournamentSelection::selected [private]
Pointer to the vector of selected individuals

#### 12.11.3.3 tournamentSize

int NaryTournamentSelection::tournamentSize [private]
Size of the tournament (number of individuals competing)

The documentation for this class was generated from the following files:

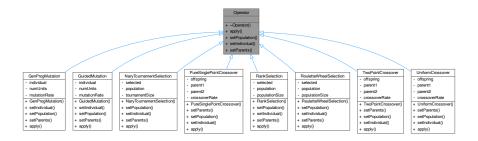
- include/operators/Selection/NaryTournamentSelection.h
- src/operators/Selection/NaryTournamentSelection.cpp

#### 12.12 Operator Class Reference

Abstract base class for genetic algorithm operators.

#include <Operator.h>

Inheritance diagram for Operator:



Collaboration diagram for Operator:

Operator
+ ~Operator()
+ apply()
+ setPopulation()
+ setIndividual()
+ setParents()

#### **Public Member Functions**

virtual ∼Operator ()=default

Virtual destructor for the Operator class.

virtual void apply (std::mt19937 &generator)=0

Pure virtual function to apply the operator.

virtual void setPopulation (const std::vector< Individual > &population, std::vector< Individual > &selected)=0

Sets the population and selected individuals for the operator.

• virtual void setIndividual (Individual &individual)=0

Sets the individual for the operator.

virtual void setParents (Individual &offspring, const Individual &parent1, const Individual &parent2)=0
 Sets the parents for the operator.

#### 12.12.1 Detailed Description

Abstract base class for genetic algorithm operators.

The Operator class provides an abstract interface for genetic algorithm operators. It defines a pure virtual function <code>apply()</code>, which must be implemented by derived classes. This class serves as a base for various types of genetic algorithm operators, such as selection, crossover, and mutation.

#### 12.12.2 Constructor & Destructor Documentation

#### 12.12.2.1 ∼Operator()

```
virtual Operator::~Operator ( ) [virtual], [default]
```

Virtual destructor for the Operator class.

The virtual destructor ensures that derived class destructors are called correctly when an object is deleted through a base class pointer.

#### 12.12.3 Member Function Documentation

#### 12.12.3.1 apply()

Pure virtual function to apply the operator.

This function must be implemented by derived classes to perform the specific operation of the genetic algorithm operator.

Implemented in GenProgMutation, PureSinglePointCrossover, TwoPointCrossover, UniformCrossover, GuidedMutation, NaryTournamentSelection, RankSelection, and RouletteWheelSelection.

#### 12.12.3.2 setIndividual()

Sets the individual for the operator.

This function sets the individual for the operator. It is used to provide the operator with the necessary data to perform its operation on a single individual.

#### **Parameters**

individual	The individual to operate on.
iiiuiviuuai	The individual to operate on.

Implemented in PureSinglePointCrossover, TwoPointCrossover, UniformCrossover, NaryTournamentSelection, RankSelection, RouletteWheelSelection, GenProgMutation, and GuidedMutation.

#### 12.12.3.3 setParents()

Sets the parents for the operator.

This function sets the parents for the operator. It is used to provide the operator with the necessary data to perform its operation on two parent individuals.

#### **Parameters**

parent1	The first parent individual.
parent2	The second parent individual.

Implemented in GenProgMutation, GuidedMutation, NaryTournamentSelection, RankSelection, RouletteWheelSelection, PureSinglePointCrossover, TwoPointCrossover, and UniformCrossover.

#### 12.12.3.4 setPopulation()

Sets the population and selected individuals for the operator.

This function sets the population and selected individuals for the operator. It is used to provide the operator with the necessary data to perform its operation.

#### **Parameters**

population	The population of individuals to operate on.
selected	The selected individuals to store the results of the operation.

Implemented in PureSinglePointCrossover, TwoPointCrossover, UniformCrossover, GenProgMutation, GuidedMutation, RouletteWheelSelection, NaryTournamentSelection, and RankSelection.

The documentation for this class was generated from the following file:

• include/operators/Operator.h

#### 12.13 tqdm::progress\_bar Class Reference

A class representing a progress bar.

#include <tqdm.hpp>

Collaboration diagram for tqdm::progress\_bar:



#### **Public Member Functions**

• void restart ()

Restart the progress bar.

• void update (double progress)

Update the progress bar with the current progress.

void set\_ostream (std::ostream &os)

Set the output stream for the progress bar.

void set\_prefix (std::string s)

Set the prefix string for the progress bar.

void set\_bar\_size (int size)

Set the size of the progress bar.

• void set\_min\_update\_time (double time)

Set the minimum time between updates.

template<class T >

```
progress_bar & operator<< (const T &t)</pre>
```

Append a value to the progress bar suffix.

• double elapsed\_time () const

Get the elapsed time since the progress bar started.

#### **Private Member Functions**

• void display (double progress)

Display the progress bar.

• void print\_bar (std::stringstream &ss, double filled) const

Print the progress bar with symbols.

· double time since refresh () const

Get the time since the last refresh.

void reset\_refresh\_timer ()

Reset the refresh timer.

#### **Private Attributes**

- Chronometer chronometer {}
- Chronometer refresh\_ {}
- double min\_time\_per\_update\_ {0.15}
- std::ostream \* os\_ {&std::cerr}
- index bar\_size\_ {40}
- index term cols {1}
- index symbol\_index\_ {0}
- std::string prefix\_{}
- std::stringstream suffix\_{}

#### 12.13.1 Detailed Description

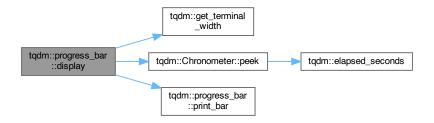
A class representing a progress bar.

#### 12.13.2 Member Function Documentation

#### 12.13.2.1 display()

Display the progress bar.

#### **Parameters**



Here is the caller graph for this function:



#### 12.13.2.2 elapsed\_time()

double tqdm::progress\_bar::elapsed\_time ( ) const [inline]
Get the elapsed time since the progress bar started.

#### Returns

The elapsed time in seconds.

Here is the call graph for this function:



Here is the caller graph for this function:



#### 12.13.2.3 operator<<()

Append a value to the progress bar suffix.

#### **Parameters**

```
t The value to append.
```

#### Returns

A reference to the progress bar.

#### 12.13.2.4 print\_bar()

Print the progress bar with symbols.

#### **Parameters**

ss	The stringstream to print the bar into.
filled	The fraction of the bar that is filled.

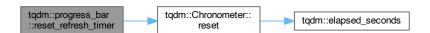
Here is the caller graph for this function:



#### 12.13.2.5 reset\_refresh\_timer()

```
\label{lem:progress_bar::reset_refresh\_timer ( ) [inline], [private]} \\ \textbf{Reset the refresh timer}.
```

Here is the call graph for this function:





#### 12.13.2.6 restart()

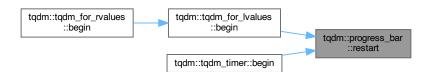
```
void tqdm::progress_bar::restart ( ) [inline]
```

Restart the progress bar.

Here is the call graph for this function:



Here is the caller graph for this function:

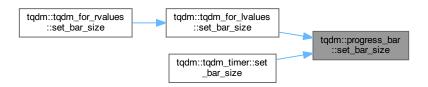


#### 12.13.2.7 set\_bar\_size()

Set the size of the progress bar.

#### **Parameters**

size The size of the progress bar in characters.



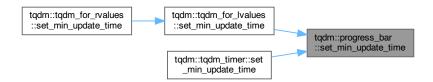
#### 12.13.2.8 set\_min\_update\_time()

Set the minimum time between updates.

#### **Parameters**

*time* The minimum time between updates in seconds.

Here is the caller graph for this function:



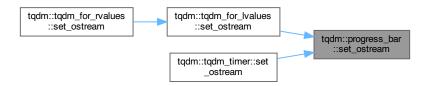
#### 12.13.2.9 set\_ostream()

```
void tqdm::progress_bar::set_ostream (  std::ostream \ \& \ os \ ) \quad [inline] \\
```

Set the output stream for the progress bar.

#### **Parameters**

os The output stream.



#### 12.13.2.10 set\_prefix()

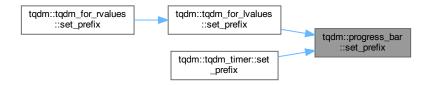
```
void tqdm::progress_bar::set_prefix (  std::string \ s \ ) \ [inline]
```

Set the prefix string for the progress bar.

#### **Parameters**

```
s The prefix string.
```

Here is the caller graph for this function:



#### 12.13.2.11 time\_since\_refresh()

double tqdm::progress\_bar::time\_since\_refresh ( ) const [inline], [private] Get the time since the last refresh.

#### Returns

The time since the last refresh in seconds.

Here is the call graph for this function:





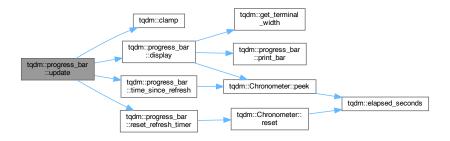
#### 12.13.2.12 update()

Update the progress bar with the current progress.

#### **Parameters**

progress	The current progress (a value between 0 and 1).
----------	---

Here is the call graph for this function:



Here is the caller graph for this function:



#### 12.13.3 Field Documentation

#### 12.13.3.1 bar\_size\_

```
index tqdm::progress_bar::bar_size_ {40} [private]
```

#### 12.13.3.2 chronometer\_

```
Chronometer tqdm::progress_bar::chronometer_ {} [private]
```

```
12.13.3.3 min_time_per_update_

double tqdm::progress_bar::min_time_per_update_ {0.15} [private]

12.13.3.4 os_

std::ostream* tqdm::progress_bar::os_ {&std::cerr} [private]

12.13.3.5 prefix_

std::string tqdm::progress_bar::prefix_ {} [private]

12.13.3.6 refresh_

Chronometer tqdm::progress_bar::refresh_ {} [private]

12.13.3.7 suffix_

std::stringstream tqdm::progress_bar::suffix_ {} [private]

12.13.3.8 symbol_index_
index tqdm::progress_bar::symbol_index_ {0} [private]

12.13.3.9 term_cols_
index tqdm::progress_bar::term_cols_ {1} [private]

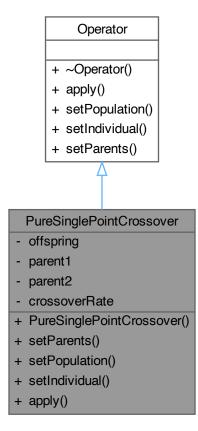
The documentation for this class was generated from the following file:
```

• include/utils/tqdm.hpp

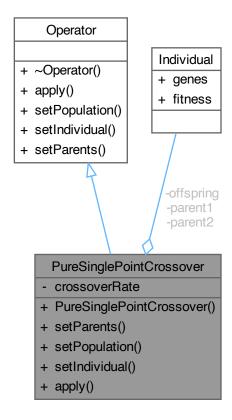
#### 12.14 PureSinglePointCrossover Class Reference

#include <PureSinglePointCrossover.h>

Inheritance diagram for PureSinglePointCrossover:



Collaboration diagram for PureSinglePointCrossover:



#### **Public Member Functions**

- PureSinglePointCrossover (const Algorithm\_Parameters &params)
  - Constructs a PureSinglePointCrossover object with specified algorithm parameters.
- void setParents (Individual &offspring, const Individual &parent1, const Individual &parent2) override Sets the parents and offspring for the crossover operation.
- void setPopulation (const std::vector< Individual > &population, std::vector< Individual > &selected)

  Sets the population and selected vectors for the selection operation.
- · void setIndividual (Individual &individual)
  - Sets the individual for the selection operation.
- void apply (std::mt19937 &generator) override
  - Performs the single-point crossover operation.

#### Public Member Functions inherited from Operator

virtual ∼Operator ()=default

Virtual destructor for the Operator class.

#### **Private Attributes**

- Individual \* offspring
- const Individual \* parent1

- const Individual \* parent2
- · double crossoverRate

#### 12.14.1 Constructor & Destructor Documentation

#### 12.14.1.1 PureSinglePointCrossover()

Constructs a PureSinglePointCrossover object with specified algorithm parameters.

#### **Parameters**

params	The algorithm parameters including the crossover rate.
--------	--

#### 12.14.2 Member Function Documentation

#### 12.14.2.1 apply()

Performs the single-point crossover operation.

This method performs a single-point crossover on the two parent individuals to generate an offspring individual. The crossover point is chosen randomly, and the genetic material from the two parents is combined to create the offspring.

#### **Parameters**

generator	The random number generator used to determine the crossover point.
-----------	--

Implements Operator.

#### 12.14.2.2 setIndividual()

Sets the individual for the selection operation.

This method is not used in the PureSinglePointCrossover class.

#### **Parameters**

```
individual
```

Implements Operator.

#### 12.14.2.3 setParents()

Sets the parents and offspring for the crossover operation.

#### **Parameters**

offspring	The offspring individual to be generated.
parent1	The first parent individual.
parent2	The second parent individual.

Implements Operator.

#### 12.14.2.4 setPopulation()

Sets the population and selected vectors for the selection operation.

This method is not used in the PureSinglePointCrossover class.

#### **Parameters**

population	
selected	

Implements Operator.

#### 12.14.3 Field Documentation

#### 12.14.3.1 crossoverRate

double PureSinglePointCrossover::crossoverRate [private]
The crossover rate used to determine if crossover should occur

#### 12.14.3.2 offspring

```
Individual* PureSinglePointCrossover::offspring [private]
Pointer to the offspring individual
```

#### 12.14.3.3 parent1

```
const Individual* PureSinglePointCrossover::parent1 [private]
Pointer to the first parent individual
```

#### 12.14.3.4 parent2

```
const Individual* PureSinglePointCrossover::parent2 [private]
Pointer to the second parent individual
```

The documentation for this class was generated from the following files:

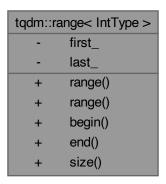
- include/operators/Crossover/PureSinglePointCrossover.h
- src/operators/Crossover/PureSinglePointCrossover.cpp

#### 12.15 tqdm::range< IntType > Class Template Reference

A range of integers represented by iterators.

```
#include <tqdm.hpp>
```

Collaboration diagram for tqdm::range < IntType >:



#### **Public Types**

- using iterator = int\_iterator < IntType >
- using const\_iterator = iterator
- using value\_type = IntType

#### **Public Member Functions**

- range (IntType first, IntType last)
- range (IntType last)
- iterator begin () const
- iterator end () const
- index size () const

#### **Private Attributes**

- · iterator first\_
- iterator last\_

#### 12.15.1 Detailed Description

template<class IntType> class tqdm::range< IntType >

A range of integers represented by iterators.

#### **Template Parameters**

IntType	The type of the integer.
---------	--------------------------

#### 12.15.2 Member Typedef Documentation

#### 12.15.2.1 const\_iterator

```
template<class IntType >
using tqdm::range< IntType >::const_iterator = iterator
```

#### 12.15.2.2 iterator

```
template<class IntType >
using tqdm::range< IntType >::iterator = int_iterator<IntType>
12.15.2.3 value_type
```

```
template<class IntType >
using tqdm::range< IntType >::value_type = IntType
```

#### 12.15.3 Constructor & Destructor Documentation

#### 12.15.3.1 range() [1/2]

#### 12.15.3.2 range() [2/2]

#### 12.15.4 Member Function Documentation

#### 12.15.4.1 begin()

```
template<class IntType >
iterator tqdm::range< IntType >::begin ( ) const [inline]
```

#### 12.15.4.2 end()

```
template<class IntType >
iterator tqdm::range< IntType >::end ( ) const [inline]
```

#### 12.15.4.3 size()

```
template<class IntType >
index tqdm::range< IntType >::size ( ) const [inline]
```

#### 12.15.5 Field Documentation

#### 12.15.5.1 first

```
template<class IntType >
iterator tqdm::range< IntType >::first_ [private]
```

#### 12.15.5.2 last\_

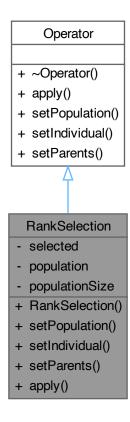
```
template<class IntType >
iterator tqdm::range< IntType >::last_ [private]
```

The documentation for this class was generated from the following file:

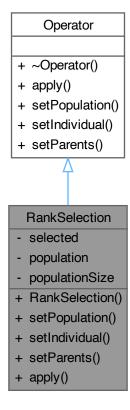
• include/utils/tqdm.hpp

#### 12.16 RankSelection Class Reference

#include <RankSelection.h>
Inheritance diagram for RankSelection:



Collaboration diagram for RankSelection:



#### **Public Member Functions**

• RankSelection (const Algorithm\_Parameters &params)

Constructs a RankSelection object with specified algorithm parameters.

 void setPopulation (const std::vector< Individual > &population, std::vector< Individual > &selected) override

Sets the population and selected vectors for the selection operation.

void setIndividual (Individual &individual)

Sets the individual to be selected.

• void setParents (Individual &offspring, const Individual &parent1, const Individual &parent2)

Sets the parents and offspring for the selection operation.

void apply (std::mt19937 &generator) override

Performs the rank-based selection operation.

#### Public Member Functions inherited from Operator

virtual ∼Operator ()=default

Virtual destructor for the Operator class.

#### **Private Attributes**

• std::vector< Individual > \* selected

- const std::vector< Individual > \* population
- · int populationSize

#### 12.16.1 Constructor & Destructor Documentation

#### 12.16.1.1 RankSelection()

Constructs a RankSelection object with specified algorithm parameters.

#### **Parameters**

params	The algorithm parameters including the population size.
--------	---

#### 12.16.2 Member Function Documentation

#### 12.16.2.1 apply()

Performs the rank-based selection operation.

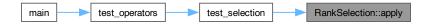
This method selects individuals from the population based on their rank and stores the selected individuals in the provided vector.

#### **Parameters**

	generator	The random number generator used for selection.	ĺ
--	-----------	---	---

Implements Operator.

Here is the caller graph for this function:



#### 12.16.2.2 setIndividual()

Sets the individual to be selected.

This method sets the individual to be selected from the population.

#### **Parameters**

individual The individual to be selected.

Implements Operator.

#### 12.16.2.3 setParents()

```
const Individual & parent1,
const Individual & parent2 ) [inline], [virtual]
```

Sets the parents and offspring for the selection operation.

This method is not used by the RankSelection operator.

#### **Parameters**

offspring	The offspring individual to be generated.
parent1	The first parent individual.
parent2	The second parent individual.

Implements Operator.

#### 12.16.2.4 setPopulation()

Sets the population and selected vectors for the selection operation.

#### **Parameters**

population	The vector of individuals in the population.
selected	The vector of selected individuals.

Implements Operator.

Here is the caller graph for this function:



#### 12.16.3 Field Documentation

#### 12.16.3.1 population

```
const std::vector<Individual>* RankSelection::population [private]
Pointer to the vector of individuals in the population
```

#### 12.16.3.2 populationSize

```
int RankSelection::populationSize [private]
Size of the population
```

#### 12.16.3.3 selected

```
std::vector<Individual>* RankSelection::selected [private]
```

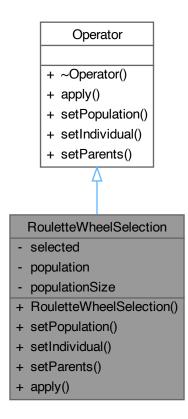
Pointer to the vector of selected individuals

The documentation for this class was generated from the following files:

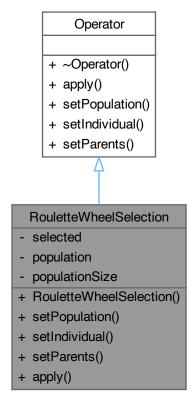
- include/operators/Selection/RankSelection.h
- src/operators/Selection/RankSelection.cpp

#### 12.17 RouletteWheelSelection Class Reference

#include <RouletteWheelSelection.h>
Inheritance diagram for RouletteWheelSelection:



Collaboration diagram for RouletteWheelSelection:



#### **Public Member Functions**

• RouletteWheelSelection (const Algorithm\_Parameters &params)

Constructs a RouletteWheelSelection object with specified algorithm parameters.

• void setPopulation (const std::vector< Individual > &population, std::vector< Individual > &selected)

Sets the population and selected vectors for the selection operation.

• void setIndividual (Individual &individual)

Sets the individual to be mutated.

• void setParents (Individual &offspring, const Individual &parent1, const Individual &parent2)

Sets the parents and offspring for the crossover operation.

· void apply (std::mt19937 &generator) override

Performs the roulette wheel selection operation.

#### Public Member Functions inherited from Operator

virtual ∼Operator ()=default

Virtual destructor for the Operator class.

#### **Private Attributes**

- std::vector< Individual > \* selected
- const std::vector< Individual > \* population
- · int populationSize

#### 12.17.1 Constructor & Destructor Documentation

#### 12.17.1.1 RouletteWheelSelection()

Constructs a RouletteWheelSelection object with specified algorithm parameters.

#### **Parameters**

params	The algorithm parameters.
--------	---------------------------

#### 12.17.2 Member Function Documentation

#### 12.17.2.1 apply()

Performs the roulette wheel selection operation.

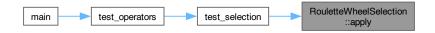
This method selects individuals from the population based on fitness-proportional selection and stores the selected individuals in the provided vector.

#### **Parameters**

generator	The random number generator used for selection.
-----------	---

Implements Operator.

Here is the caller graph for this function:



#### 12.17.2.2 setIndividual()

Sets the individual to be mutated.

#### **Parameters**

individual	The individual to be mutated.
individual	The individual to be mutated.

Implements Operator.

#### 12.17.2.3 setParents()

Sets the parents and offspring for the crossover operation.

This method is not used by the RouletteWheelSelection operator.

#### **Parameters**

offspring	The offspring individual to be generated.
parent1	The first parent individual.
parent2	The second parent individual.

Implements Operator.

#### 12.17.2.4 setPopulation()

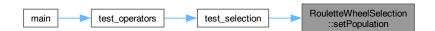
Sets the population and selected vectors for the selection operation.

#### **Parameters**

population	The vector of individuals in the population.
selected	The vector of selected individuals.

Implements Operator.

Here is the caller graph for this function:



#### 12.17.3 Field Documentation

#### 12.17.3.1 population

const std::vector<Individual>\* RouletteWheelSelection::population [private]
Pointer to the vector of individuals in the population

#### 12.17.3.2 populationSize

```
\begin{tabular}{ll} \end{tabular} int $\tt RouletteWheelSelection::populationSize & [private] \\ \end{tabular} \end{tabular} Size of the population
```

#### 12.17.3.3 selected

```
std::vector<Individual>* RouletteWheelSelection::selected [private]
```

Pointer to the vector of selected individuals

The documentation for this class was generated from the following files:

- include/operators/Selection/RouletteWheelSelection.h
- src/operators/Selection/RouletteWheelSelection.cpp

#### 12.18 tqdm::timer Struct Reference

A timer that can be used to create a timing iterator.

#include <tqdm.hpp>
Collaboration diagram for tqdm::timer:

## tqdm::timer - num\_seconds\_ + timer() + end() + num\_seconds() + begin()

#### **Public Types**

- using iterator = timing\_iterator
- using end\_iterator = timing\_iterator\_end\_sentinel
- using const\_iterator = iterator
- using value\_type = double

#### **Public Member Functions**

- timer (double num\_seconds)
- end\_iterator end () const
- double num\_seconds () const

#### **Static Public Member Functions**

• static iterator begin ()

#### **Private Attributes**

• double num\_seconds\_

#### 12.18.1 Detailed Description

A timer that can be used to create a timing iterator.

#### 12.18.2 Member Typedef Documentation

#### 12.18.2.1 const\_iterator

using tqdm::timer::const\_iterator = iterator

#### 12.18.2.2 end\_iterator

using tqdm::timer::end\_iterator = timing\_iterator\_end\_sentinel

#### 12.18.2.3 iterator

using tqdm::timer::iterator = timing\_iterator

#### 12.18.2.4 value\_type

```
using tqdm::timer::value_type = double
```

### 12.18.3 Constructor & Destructor Documentation

#### 12.18.3.1 timer()

#### 12.18.4 Member Function Documentation

#### 12.18.4.1 begin()

```
static iterator tqdm::timer::begin ( ) [inline], [static]
```

#### 12.18.4.2 end()

```
end_iterator tqdm::timer::end ( ) const [inline]
```

#### 12.18.4.3 num\_seconds()

double tqdm::timer::num\_seconds ( ) const [inline]
Here is the caller graph for this function:



### 12.18.5 Field Documentation

#### 12.18.5.1 num\_seconds\_

```
double tqdm::timer::num_seconds_ [private]
```

The documentation for this struct was generated from the following file:

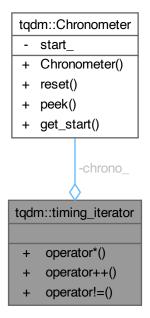
• include/utils/tqdm.hpp

## 12.19 tqdm::timing\_iterator Class Reference

A forward iterator that returns the elapsed time.

```
#include <tqdm.hpp>
```

Collaboration diagram for tqdm::timing\_iterator:



#### **Public Types**

- using iterator\_category = std::forward\_iterator\_tag
- using value\_type = double
- using difference\_type = double
- using pointer = double \*
- using reference = double &

#### **Public Member Functions**

- double operator\* () const
- timing\_iterator & operator++ ()
- bool operator!= (const timing\_iterator\_end\_sentinel &other) const

### **Private Attributes**

• tqdm::Chronometer chrono\_

### 12.19.1 Detailed Description

A forward iterator that returns the elapsed time.

### 12.19.2 Member Typedef Documentation

#### 12.19.2.1 difference\_type

using tqdm::timing\_iterator::difference\_type = double

#### 12.19.2.2 iterator\_category

```
using tqdm::timing_iterator::iterator_category = std::forward_iterator_tag
```

#### 12.19.2.3 pointer

```
using tqdm::timing_iterator::pointer = double *
```

### 12.19.2.4 reference

```
using tqdm::timing_iterator::reference = double &
```

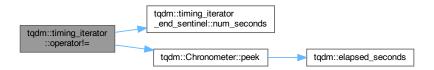
#### 12.19.2.5 value\_type

using tqdm::timing\_iterator::value\_type = double

#### 12.19.3 Member Function Documentation

#### 12.19.3.1 operator"!=()

Here is the call graph for this function:



#### 12.19.3.2 operator\*()

double tqdm::timing\_iterator::operator\* ( ) const [inline]
Here is the call graph for this function:



### 12.19.3.3 operator++()

```
timing_iterator & tqdm::timing_iterator::operator++ ( ) [inline]
```

### 12.19.4 Field Documentation

### 12.19.4.1 chrono\_

```
tqdm::Chronometer tqdm::timing_iterator::chrono_ [private]
```

The documentation for this class was generated from the following file:

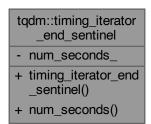
include/utils/tqdm.hpp

### 12.20 tqdm::timing iterator end sentinel Class Reference

An end sentinel for timing iterators.

#include <tqdm.hpp>

Collaboration diagram for tqdm::timing\_iterator\_end\_sentinel:



#### **Public Member Functions**

- timing\_iterator\_end\_sentinel (double num\_seconds)
- double num\_seconds () const

#### **Private Attributes**

• double num\_seconds\_

### 12.20.1 Detailed Description

An end sentinel for timing iterators.

#### 12.20.2 Constructor & Destructor Documentation

#### 12.20.2.1 timing\_iterator\_end\_sentinel()

#### 12.20.3 Member Function Documentation

### 12.20.3.1 num\_seconds()

double tqdm::timing\_iterator\_end\_sentinel::num\_seconds ( ) const [inline] Here is the caller graph for this function:



### 12.20.4 Field Documentation

12.20.4.1 num\_seconds\_

double tqdm::timing\_iterator\_end\_sentinel::num\_seconds\_ [private] The documentation for this class was generated from the following file:

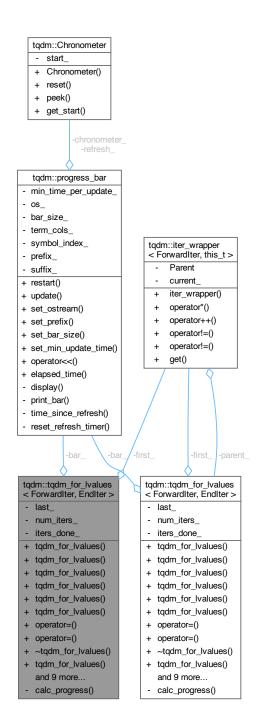
• include/utils/tqdm.hpp

# 12.21 tqdm::tqdm\_for\_lvalues< ForwardIter, EndIter > Class Template Reference

A class for progress bars that iterate over Ivalues.

#include <tqdm.hpp>

Collaboration diagram for tqdm::tqdm\_for\_lvalues< ForwardIter, EndIter >:



#### **Public Types**

- using this\_t = tqdm\_for\_lvalues<ForwardIter, EndIter>
- using iterator = iter\_wrapper<ForwardIter, this\_t>
- using value\_type = typename ForwardIter::value\_type
- using size\_type = index
- using difference\_type = index

### **Public Member Functions**

- tqdm\_for\_lvalues (ForwardIter begin, EndIter end)
- tqdm for Ivalues (ForwardIter begin, EndIter end, index total)
- template < class Container > tqdm\_for\_lvalues (Container &C)
- template<class Container >
  - tgdm for Ivalues (const Container &C)
- tqdm\_for\_lvalues (const tqdm\_for\_lvalues &)=delete
- tqdm for Ivalues (tqdm for Ivalues &&)=delete
- tqdm\_for\_lvalues & operator= (tqdm\_for\_lvalues &&)=delete
- tqdm\_for\_lvalues & operator= (const tqdm\_for\_lvalues &)=delete
- ~tqdm\_for\_lvalues ()=default
- $\bullet \ \ \text{template}{<} \text{class Container} >$ 
  - tgdm for Ivalues (Container &&)=delete
- iterator begin ()
- EndIter end () const
- void update ()
- void set\_ostream (std::ostream &os)
- void set prefix (std::string s)
- void set\_bar\_size (int size)
- void set\_min\_update\_time (double time)
- template<class T >
  - tqdm\_for\_lvalues & operator<< (const T &t)
- void manually\_set\_progress (double to)

#### **Private Member Functions**

• double calc\_progress () const

#### **Private Attributes**

- · iterator first\_
- EndIter last\_
- index num\_iters\_ {0}
- index iters\_done\_ {0}
- progress\_bar bar\_

### 12.21.1 Detailed Description

template<class ForwardIter, class EndIter = ForwardIter> class tqdm::tqdm\_for\_lvalues< ForwardIter, EndIter>

A class for progress bars that iterate over Ivalues.

### **Template Parameters**

ForwardIter Th		The type of the forward iterator.
	EndIter	The type of the end iterator.

### 12.21.2 Member Typedef Documentation

### 12.21.2.1 difference\_type

```
template<class ForwardIter , class EndIter = ForwardIter>
using tqdm::tqdm_for_lvalues< ForwardIter, EndIter >::difference_type = index
```

#### 12.21.2.2 iterator

```
template<class ForwardIter , class EndIter = ForwardIter>
using tqdm::tqdm_for_lvalues< ForwardIter, EndIter >::iterator = iter_wrapper<ForwardIter,
this_t>
```

#### 12.21.2.3 size\_type

```
template<class ForwardIter , class EndIter = ForwardIter>
using tqdm::tqdm_for_lvalues< ForwardIter, EndIter >::size_type = index
```

#### 12.21.2.4 this\_t

```
template<class ForwardIter , class EndIter = ForwardIter>
using tqdm::tqdm_for_lvalues< ForwardIter, EndIter >::this_t = tqdm_for_lvalues<ForwardIter,
EndIter>
```

#### 12.21.2.5 value\_type

```
template<class ForwardIter , class EndIter = ForwardIter>
using tqdm::tqdm_for_lvalues< ForwardIter, EndIter >::value_type = typename ForwardIter←
::value_type
```

#### 12.21.3 Constructor & Destructor Documentation

### 12.21.3.1 tqdm\_for\_lvalues() [1/7]

#### 12.21.3.2 tqdm\_for\_lvalues() [2/7]

#### 12.21.3.3 tqdm for Ivalues() [3/7]

#### 12.21.3.4 tqdm\_for\_lvalues() [4/7]

#### 12.21.3.5 tqdm for Ivalues() [5/7]

#### 12.21.3.6 tqdm\_for\_lvalues() [6/7]

#### 12.21.3.7 ~tqdm\_for\_lvalues()

```
template<class ForwardIter , class EndIter = ForwardIter>
tqdm::tqdm_for_lvalues
ForwardIter, EndIter >::~tqdm_for_lvalues ( ) [default]
```

#### 12.21.3.8 tqdm\_for\_lvalues() [7/7]

#### 12.21.4 Member Function Documentation

#### 12.21.4.1 begin()

template<class ForwardIter , class EndIter = ForwardIter>
iterator tqdm::tqdm\_for\_lvalues< ForwardIter, EndIter >::begin ( ) [inline]
Here is the call graph for this function:



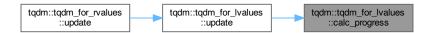
Here is the caller graph for this function:



### 12.21.4.2 calc\_progress()

```
template<class ForwardIter , class EndIter = ForwardIter>
double tqdm::tqdm_for_lvalues< ForwardIter, EndIter >::calc_progress ( ) const [inline],
[private]
```

Here is the caller graph for this function:



### 12.21.4.3 end()

```
template<class ForwardIter , class EndIter = ForwardIter>
EndIter tqdm::tqdm_for_lvalues< ForwardIter, EndIter >::end ( ) const [inline]
Here is the caller graph for this function:
```

```
tqdm::tqdm_for_rvalues::end tqdm::tqdm_for_lvalues::end
```

### 12.21.4.4 manually\_set\_progress()

Here is the call graph for this function:



Here is the caller graph for this function:



#### 12.21.4.5 operator<<()

```
template<class ForwardIter , class EndIter = ForwardIter>
template < class T >
tqdm_for_lvalues & tqdm::tqdm_for_lvalues< ForwardIter, EndIter >::operator<< (
             {\tt const T \& t ) \quad [inline]}
12.21.4.6 operator=() [1/2]
template<class ForwardIter , class EndIter = ForwardIter>
tqdm_for_lvalues & tqdm::tqdm_for_lvalues< ForwardIter, EndIter >::operator= (
             const tqdm_for_lvalues< ForwardIter, EndIter > \& ) [delete]
12.21.4.7 operator=() [2/2]
```

```
template<class ForwardIter , class EndIter = ForwardIter>
tqdm_for_lvalues & tqdm::tqdm_for_lvalues< ForwardIter, EndIter >::operator= (
            tqdm_for_lvalues< ForwardIter, EndIter > && ) [delete]
```

#### 12.21.4.8 set bar size()

```
template<class ForwardIter , class EndIter = ForwardIter>
void tqdm::tqdm_for_lvalues< ForwardIter, EndIter >::set_bar_size (
            int size ) [inline]
```

Here is the call graph for this function:



Here is the caller graph for this function:

```
tqdm::tqdm_for_rvalues
                                   tqdm::tqdm_for_lvalues
    ::set_bar_size
                                       ::set_bar_size
```

### 12.21.4.9 set\_min\_update\_time()

```
template<class ForwardIter , class EndIter = ForwardIter>
void tqdm::tqdm_for_lvalues< ForwardIter, EndIter >::set_min_update_time (
            double time ) [inline]
```

Here is the call graph for this function:



Here is the caller graph for this function:

```
tqdm::tqdm_for_rvalues
::set_min_update_time tqdm::tqdm::tqdm_for_lvalues
::set_min_update_time
```

### 12.21.4.10 set\_ostream()

Here is the call graph for this function:



Here is the caller graph for this function:



#### 12.21.4.11 set\_prefix()

Here is the call graph for this function:

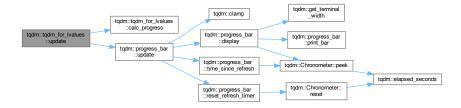


Here is the caller graph for this function:



#### 12.21.4.12 update()

```
template<class ForwardIter , class EndIter = ForwardIter>
void tqdm::tqdm_for_lvalues< ForwardIter, EndIter >::update ( ) [inline]
Here is the call graph for this function:
```



Here is the caller graph for this function:



#### 12.21.5 Field Documentation

### 12.21.5.1 bar\_

```
template<class ForwardIter , class EndIter = ForwardIter>
progress_bar tqdm::tqdm_for_lvalues< ForwardIter, EndIter >::bar_ [private]
```

#### 12.21.5.2 first

```
template<class ForwardIter , class EndIter = ForwardIter>
iterator tqdm::tqdm_for_lvalues< ForwardIter, EndIter >::first_ [private]
```

#### 12.21.5.3 iters\_done\_

```
template<class ForwardIter , class EndIter = ForwardIter>
index tqdm::tqdm_for_lvalues< ForwardIter, EndIter >::iters_done_ {0} [private]
```

#### 12.21.5.4 last\_

```
template<class ForwardIter , class EndIter = ForwardIter>
EndIter tqdm::tqdm_for_lvalues< ForwardIter, EndIter >::last_ [private]
```

### 12.21.5.5 num\_iters\_

```
template<class ForwardIter , class EndIter = ForwardIter>
index tqdm::tqdm_for_lvalues< ForwardIter, EndIter >::num_iters_ {0} [private]
The documentation for this class was generated from the following file:
```

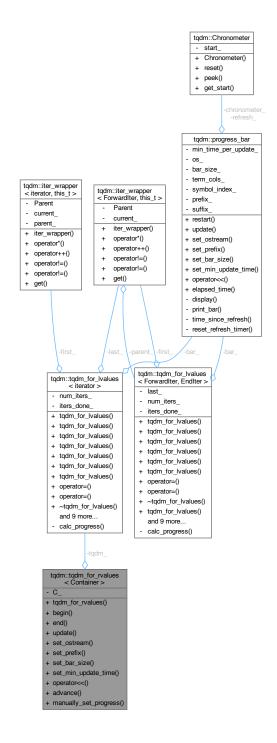
• include/utils/tqdm.hpp

### 12.22 tqdm::tqdm\_for\_rvalues< Container > Class Template Reference

A class for progress bars that iterate over rvalues.

```
#include <tqdm.hpp>
```

Collaboration diagram for tqdm::tqdm\_for\_rvalues< Container >:



### **Public Types**

- using iterator = typename Container::iterator
- using const\_iterator = typename Container::const\_iterator
- using value\_type = typename Container::value\_type

### **Public Member Functions**

• tqdm\_for\_rvalues (Container &&C)

- · auto begin ()
- auto end ()
- void update ()
- void set ostream (std::ostream &os)
- void set\_prefix (std::string s)
- void set\_bar\_size (int size)
- void set\_min\_update\_time (double time)
- template < class T >
   auto & operator << (const T &t)</li>
- void advance (index amount)
- void manually\_set\_progress (double to)

#### **Private Attributes**

- Container C
- tqdm\_for\_lvalues< iterator > tqdm\_

#### 12.22.1 Detailed Description

```
template < class Container > class tqdm::tqdm for rvalues < Container >
```

A class for progress bars that iterate over rvalues.

#### **Template Parameters**

```
Container The type of the container.
```

### 12.22.2 Member Typedef Documentation

#### 12.22.2.1 const\_iterator

```
template<class Container >
using tqdm::tqdm_for_rvalues< Container >::const_iterator = typename Container::const_iterator
```

#### 12.22.2.2 iterator

```
template<class Container >
using tqdm::tqdm_for_rvalues< Container >::iterator = typename Container::iterator
```

#### 12.22.2.3 value\_type

```
template<class Container >
using tqdm::tqdm_for_rvalues< Container >::value_type = typename Container::value_type
```

### 12.22.3 Constructor & Destructor Documentation

#### 12.22.3.1 tqdm\_for\_rvalues()

#### 12.22.4 Member Function Documentation

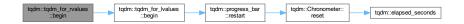
#### 12.22.4.1 advance()

```
template<class Container >
void tqdm::tqdm_for_rvalues< Container >::advance (
```

```
index amount ) [inline]
```

#### 12.22.4.2 begin()

```
template<class Container >
auto tqdm::tqdm_for_rvalues< Container >::begin ( ) [inline]
Here is the call graph for this function:
```



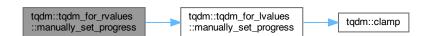
#### 12.22.4.3 end()

```
template<class Container >
auto tqdm::tqdm_for_rvalues< Container >::end ( ) [inline]
Here is the call graph for this function:
```



### 12.22.4.4 manually\_set\_progress()

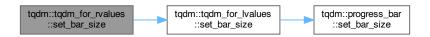
Here is the call graph for this function:



### 12.22.4.5 operator<<()

### 12.22.4.6 set\_bar\_size()

Here is the call graph for this function:



#### 12.22.4.7 set\_min\_update\_time()

Here is the call graph for this function:



### 12.22.4.8 set\_ostream()

Here is the call graph for this function:



### 12.22.4.9 set\_prefix()

Here is the call graph for this function:



#### 12.22.4.10 update()

```
template < class Container >
void tqdm::tqdm_for_rvalues < Container >::update ( ) [inline]
Here is the call graph for this function:
```



#### 12.22.5 Field Documentation

#### 12.22.5.1 C

```
template<class Container >
Container tqdm::tqdm_for_rvalues< Container >::C_ [private]
```

#### 12.22.5.2 tqdm\_

```
template<class Container >
tqdm_for_lvalues<iterator> tqdm::tqdm_for_rvalues< Container >::tqdm_ [private]
The documentation for this class was generated from the following file:
```

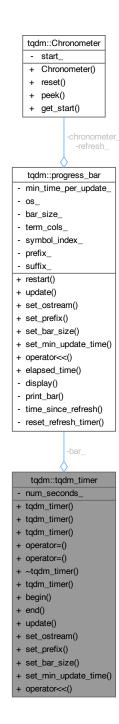
• include/utils/tqdm.hpp

### 12.23 tqdm::tqdm\_timer Class Reference

### A progress bar for a timer.

#include <tqdm.hpp>

Collaboration diagram for tqdm::tqdm\_timer:



### **Public Types**

- $\bullet \ \ using \ iterator = iter\_wrapper < timing\_iterator, \ tqdm\_timer >$
- using end\_iterator = timer::end\_iterator
- using value\_type = typename timing\_iterator::value\_type
- using size\_type = index
- using difference\_type = index

#### **Public Member Functions**

- tqdm\_timer (double num\_seconds)
- tqdm timer (const tqdm timer &)=delete
- tqdm\_timer (tqdm\_timer &&)=delete
- tqdm\_timer & operator= (tqdm\_timer &&)=delete
- tqdm timer & operator= (const tqdm timer &)=delete
- ~tqdm\_timer ()=default
- template < class Container > tqdm\_timer (Container &&)=delete
- iterator begin ()
- end\_iterator end () const
- void update ()
- void set ostream (std::ostream &os)
- void set\_prefix (std::string s)
- void set\_bar\_size (int size)
- void set\_min\_update\_time (double time)
- template < class T > tqdm\_timer & operator << (const T &t)</li>

#### **Private Attributes**

- double num seconds
- · progress\_bar bar\_

### 12.23.1 Detailed Description

A progress bar for a timer.

#### 12.23.2 Member Typedef Documentation

#### 12.23.2.1 difference type

```
using tqdm::tqdm_timer::difference_type = index
```

#### 12.23.2.2 end\_iterator

```
using tqdm::tqdm_timer::end_iterator = timer::end_iterator
```

#### 12.23.2.3 iterator

```
using tqdm::tqdm_timer::iterator = iter_wrapper<timing_iterator, tqdm_timer>
```

#### 12.23.2.4 size\_type

```
using tqdm::tqdm_timer::size_type = index
```

#### 12.23.2.5 value type

```
using tqdm::tqdm_timer::value_type = typename timing_iterator::value_type
```

### 12.23.3 Constructor & Destructor Documentation

#### 12.23.3.1 tqdm\_timer() [1/4]

#### 12.23.3.2 tqdm\_timer() [2/4]

#### 12.23.4 Member Function Documentation

Container && ) [delete]

#### 12.23.4.1 begin()

iterator tqdm::tqdm\_timer::begin ( ) [inline]
Here is the call graph for this function:



### 12.23.4.2 end()

```
end_iterator tqdm::tqdm_timer::end ( ) const [inline]
```

#### 12.23.4.3 operator<<()

### **12.23.4.4** operator=() [1/2]

#### 12.23.4.5 operator=() [2/2]

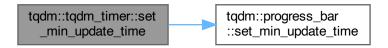
### 12.23.4.6 set\_bar\_size()

Here is the call graph for this function:



### 12.23.4.7 set\_min\_update\_time()

Here is the call graph for this function:



#### 12.23.4.8 set\_ostream()

Here is the call graph for this function:



#### 12.23.4.9 set\_prefix()

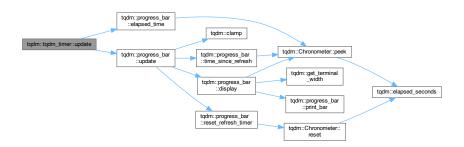
```
void tqdm::tqdm_timer::set_prefix (  std::string \ s \ ) \quad [inline] \\
```

Here is the call graph for this function:



### 12.23.4.10 update()

void tqdm::tqdm\_timer::update ( ) [inline]
Here is the call graph for this function:



### 12.23.5 Field Documentation

### 12.23.5.1 bar\_

progress\_bar tqdm::tqdm\_timer::bar\_ [private]

### 12.23.5.2 num\_seconds\_

double tqdm::tqdm\_timer::num\_seconds\_ [private]

The documentation for this class was generated from the following file:

• include/utils/tqdm.hpp

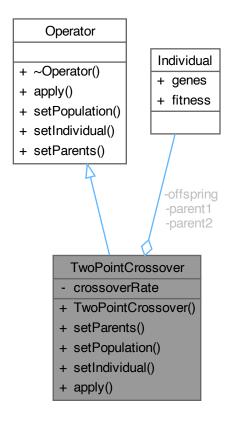
### 12.24 TwoPointCrossover Class Reference

#include <TwoPointCrossover.h>

Inheritance diagram for TwoPointCrossover:

# Operator + ~Operator() + apply() + setPopulation() + setIndividual() + setParents() TwoPointCrossover - offspring - parent1 - parent2 - crossoverRate + TwoPointCrossover() + setParents() + setPopulation() + setIndividual() + apply()

Collaboration diagram for TwoPointCrossover:



#### **Public Member Functions**

- TwoPointCrossover (const Algorithm\_Parameters &params)
  - Constructs a TwoPointCrossover object with specified algorithm parameters.
- void setParents (Individual &offspring, const Individual &parent1, const Individual &parent2) override Sets the parents and offspring for the crossover operation.
- void setPopulation (const std::vector< Individual > &population, std::vector< Individual > &selected)

  Sets the population and selected vectors for the selection operation.
- · void setIndividual (Individual &individual)

Sets the individual for the operator.

• void apply (std::mt19937 &generator) override

Performs the two-point crossover operation.

### **Public Member Functions inherited from Operator**

virtual ~Operator ()=default
 Virtual destructor for the Operator class.

#### **Private Attributes**

- Individual \* offspring
- const Individual \* parent1

- const Individual \* parent2
- · double crossoverRate

#### 12.24.1 Constructor & Destructor Documentation

#### 12.24.1.1 TwoPointCrossover()

Constructs a TwoPointCrossover object with specified algorithm parameters.

#### **Parameters**

	params	The algorithm parameters including the crossover rate.
--	--------	--

#### 12.24.2 Member Function Documentation

#### 12.24.2.1 apply()

Performs the two-point crossover operation.

This method performs a two-point crossover on the two parent individuals to generate an offspring individual. Two crossover points are chosen randomly, and the genetic material from the two parents is combined to create the offspring.

#### **Parameters**

generator	The random number generator used to determine the crossover points.
-----------	---

Implements Operator.

#### 12.24.2.2 setIndividual()

Sets the individual for the operator.

This method is not used by the TwoPointCrossover operator.

#### **Parameters**

	individual	The individual to be set.

Implements Operator.

### 12.24.2.3 setParents()

Sets the parents and offspring for the crossover operation.

#### **Parameters**

offspring	The offspring individual to be generated.
	, ,
parent1	The first parent individual.
parent2	The second parent individual.

Implements Operator.

#### 12.24.2.4 setPopulation()

Sets the population and selected vectors for the selection operation.

This method is not used by the TwoPointCrossover operator.

#### **Parameters**

population	The vector of individuals in the population.
selected	The vector of selected individuals.

Implements Operator.

#### 12.24.3 Field Documentation

#### 12.24.3.1 crossoverRate

double TwoPointCrossover::crossoverRate [private]
The crossover rate used to determine if crossover should occur

### 12.24.3.2 offspring

```
Individual* TwoPointCrossover::offspring [private]
Pointer to the offspring individual
```

#### 12.24.3.3 parent1

```
const Individual* TwoPointCrossover::parent1 [private]
Pointer to the first parent individual
```

#### 12.24.3.4 parent2

```
const Individual* TwoPointCrossover::parent2 [private]
```

Pointer to the second parent individual

The documentation for this class was generated from the following files:

- include/operators/Crossover/TwoPointCrossover.h
- src/operators/Crossover/TwoPointCrossover.cpp

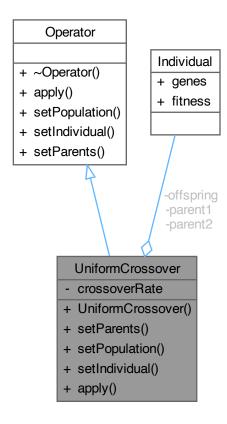
### 12.25 UniformCrossover Class Reference

#include <UniformCrossover.h>

Inheritance diagram for UniformCrossover:

# Operator + ~Operator() + apply() + setPopulation() + setIndividual() + setParents() UniformCrossover - offspring - parent1 - parent2 - crossoverRate + UniformCrossover() + setParents() + setPopulation() + setIndividual() + apply()

Collaboration diagram for UniformCrossover:



#### **Public Member Functions**

- UniformCrossover (const Algorithm\_Parameters &params)
  - Constructs a UniformCrossover object with specified algorithm parameters.
- void setParents (Individual &offspring, const Individual &parent1, const Individual &parent2) override Sets the parents and offspring for the crossover operation.
- void setPopulation (const std::vector< Individual > &population, std::vector< Individual > &selected)

  Sets the population and selected vectors for the selection operation.
- · void setIndividual (Individual &individual)
  - Sets the individual for the selection operation.
- · void apply (std::mt19937 &generator) override
  - Performs the uniform crossover operation.

Virtual destructor for the Operator class.

### **Public Member Functions inherited from Operator**

virtual ∼Operator ()=default

#### **Private Attributes**

- Individual \* offspring
- const Individual \* parent1

- const Individual \* parent2
- · double crossoverRate

#### 12.25.1 Constructor & Destructor Documentation

#### 12.25.1.1 UniformCrossover()

Constructs a UniformCrossover object with specified algorithm parameters.

#### **Parameters**

params	The algorithm parameters including the crossover rate.
--------	--

#### 12.25.2 Member Function Documentation

#### 12.25.2.1 apply()

Performs the uniform crossover operation.

This method performs a uniform crossover on the two parent individuals to generate an offspring individual. Each gene in the offspring is chosen randomly from one of the two parents.

#### **Parameters**

	generator	The random number generator used to determine gene selection.	
--	-----------	---	--

Implements Operator.

### 12.25.2.2 setIndividual()

Sets the individual for the selection operation.

This method is not used in the UniformCrossover class.

#### **Parameters**

individ	<i>ual</i> The ir	ndividual to be set.

Implements Operator.

#### 12.25.2.3 setParents()

Sets the parents and offspring for the crossover operation.

#### **Parameters**

offspring	The offspring individual to be generated.
, ,	
parent1	The first parent individual.
parent2	The second parent individual.

Implements Operator.

### 12.25.2.4 setPopulation()

Sets the population and selected vectors for the selection operation.

This method is not used in the UniformCrossover class.

#### **Parameters**

population	The vector of individuals in the population.
selected	The vector of selected individuals.

Implements Operator.

#### 12.25.3 Field Documentation

#### 12.25.3.1 crossoverRate

```
double UniformCrossover::crossoverRate [private]
The crossover rate used to determine if crossover should occur
```

### 12.25.3.2 offspring

```
Individual* UniformCrossover::offspring [private]
Pointer to the offspring individual
```

#### 12.25.3.3 parent1

```
const Individual* UniformCrossover::parent1 [private]
Pointer to the first parent individual
```

### 12.25.3.4 parent2

```
const Individual* UniformCrossover::parent2 [private]
```

Pointer to the second parent individual

The documentation for this class was generated from the following files:

- include/operators/Crossover/UniformCrossover.h
- src/operators/Crossover/UniformCrossover.cpp

# **Chapter 13**

# **File Documentation**

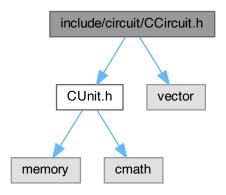
- 13.1 docs/CCircuit Document Analysis.md File Reference
- 13.2 docs/CSimulator\_Document\_and\_Analysis.md File Reference
- 13.3 docs/GeneticAlgorithm\_Design\_Document.md File Reference
- 13.4 docs/GeneticAlgorithm Params Analysis.md File Reference

### 13.5 include/circuit/CCircuit.h File Reference

Defines the Circuit class for modeling and simulating a circuit system.

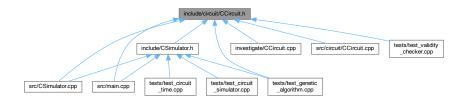
#include "CUnit.h"
#include <vector>

Include dependency graph for CCircuit.h:



156 File Documentation

This graph shows which files directly or indirectly include this file:



#### **Data Structures**

· class Circuit

#### **Macros**

#define CIRCUIT\_MODELLING\_CCIRCUIT\_H

### 13.5.1 Detailed Description

Defines the Circuit class for modeling and simulating a circuit system.

The Circuit class provides methods to initialize, configure, and simulate a circuit with multiple units. It includes methods for checking validity, printing information, connecting units, initializing feed rates, checking convergence, and calculating flows. The Circuit class is designed to facilitate the modeling and simulation of circuits in various applications, including industrial processes and scientific research.

Date

Created on May 20, 2024

#### Authors

ACS Gerardium Rush - Pentlandite:

- · Alex N Njeumi
- Geyu Ji
- Melissa Y S Sim
- · Mingsheng Cai
- · Peifeng Tan
- · Yongwen Chen
- Zihan Li

### 13.5.2 Macro Definition Documentation

### 13.5.2.1 CIRCUIT\_MODELLING\_CCIRCUIT\_H

#define CIRCUIT\_MODELLING\_CCIRCUIT\_H

### 13.6 CCircuit.h

### Go to the documentation of this file.

```
00001
00024 #pragma once
00025
00026 #ifndef CIRCUIT_MODELLING_CCIRCUIT_H
00027 #define CIRCUIT_MODELLING_CCIRCUIT_H
00028
00028 #include "CUnit.h"
```

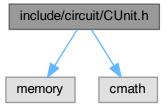
```
00030 #include <vector>
00031
00032 class Circuit {
00033 public:
          // Concentrates
00034
00035
          double concentrateG:
          double concentrateW;
00038
          // Tailings
00039
          double tailingsG;
00040
          double tailingsW;
00042
          // Feed rates
          double wasteFeed;
00043
00044
          double gerardiumFeed;
00051
          Circuit(int num_units);
00052
00056
          ~Circuit();
00057
00068
          static bool Check_Validity(int vectorSize, int *circuitVector);
00069
00076
          void print_info() const;
00077
00086
          void connected(int *circuitVector);
00087
00096
          void initialize_feed_rates(double gerardium_ = 10, double waste_ = 90);
00097
00110
          void mark_units(int unitNum, std::vector<CUnit *> &units, bool check[3]);
00111
00121
          bool check_convergence(double threshold);
00122
00128
          void calculate flows();
00129
00130 private:
00136
          void reset_new_feeds();
00137
00138
          std::vector<CUnit *> units;
          int numUnits;
static bool check[3];
00139
00140
00141
          int feed;
00142 };
00143
00144 #endif // CIRCUIT_MODELLING_CCIRCUIT_H
```

### 13.7 include/circuit/CUnit.h File Reference

Defines the CUnit class for representing a unit within a circuit system.

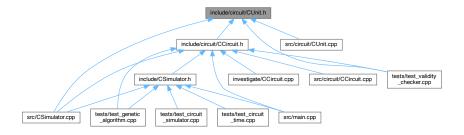
```
#include <memory>
#include <cmath>
```

Include dependency graph for CUnit.h:



158 File Documentation

This graph shows which files directly or indirectly include this file:



#### **Data Structures**

· class CUnit

#### **Macros**

• #define CIRCUIT\_MODELLING\_CUNIT\_H

### 13.7.1 Detailed Description

Defines the CUnit class for representing a unit within a circuit system.

The CUnit class provides methods to manage and simulate individual units within a circuit. It includes methods for setting and getting various properties of the unit, such as feed rates, mark status, and connection pointers. This class is designed to facilitate the modeling and simulation of circuits in various applications, including industrial processes and scientific research.

Date

Created on May 20, 2024

#### **Authors**

ACS Gerardium Rush - Pentlandite:

- · Alex N Njeumi
- · Geyu Ji
- · Melissa Y S Sim
- · Mingsheng Cai
- · Peifeng Tan
- · Yongwen Chen
- · Zihan Li

### 13.7.2 Macro Definition Documentation

### 13.7.2.1 CIRCUIT\_MODELLING\_CUNIT\_H

#define CIRCUIT\_MODELLING\_CUNIT\_H

### 13.8 CUnit.h

### Go to the documentation of this file.

```
00001
00024 #pragma once
00025
00026 #ifndef CIRCUIT_MODELLING_CUNIT_H
```

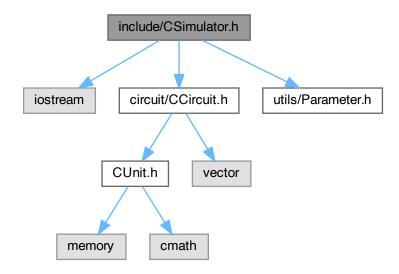
```
00027 #define CIRCUIT_MODELLING_CUNIT_H
00028
00029 #include <memory> //
00030 #include <cmath>
00031
00032 class CUnit {
00033 public:
00039
          CUnit();
00040
00050
          CUnit(int concNum, int interNum, int tailsNum);
00051
00052
          int concNum;
00053
          int interNum;
00054
          int tailsNum;
00061
          bool isMarked() const;
00062
00068
          void setMarked(bool mark);
00069
00075
          double getFeedGerardium() const;
00076
00082
          void setFeedGerardium(double feedGerardium);
00083
00089
          double getFeedWaste() const;
00090
00096
          void setFeedWaste(double feedWaste);
00097
00103
          double getNewFeedGerardium() const;
00104
00110
          void setNewFeedGerardium(double newFeedGerardium);
00111
00117
          double getNewFeedWaste() const;
00118
00124
          void setNewFeedWaste(double newFeedWaste);
00125
00131
          double getDifferenceG() const;
00132
00138
          double getDifferenceW() const;
00139
00143
          void setDifference();
00144
00145
          CUnit *getConcPtr() const;
00151
00152
00158
          void setConcPtr(CUnit *ptr);
00159
00165
          CUnit *getInterPtr() const;
00166
00172
          void setInterPtr(CUnit *ptr);
00173
00179
          CUnit *getTailsPtr() const;
00180
00186
          void setTailsPtr(CUnit *ptr);
00187
00188 private:
          bool mark;
00189
00190
          double feedGerardium;
          double feedWaste;
00192
          double newFeedGerardium;
00193
          double newFeedWaste;
00194
          double differenceG;
00195
          double differenceW;
          CUnit *concPtr;
CUnit *interPtr;
00197
00198
00199
          CUnit *tailsPtr;
00200 };
00201
00202 #endif // CIRCUIT_MODELLING_CUNIT_H
```

# 13.9 include/CSimulator.h File Reference

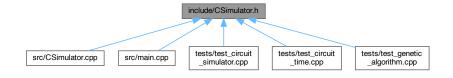
Defines the functions and structures for simulating and evaluating circuits in a circuit modeling framework.

```
#include <iostream>
#include "circuit/CCircuit.h"
#include "utils/Parameter.h"
```

Include dependency graph for CSimulator.h:



This graph shows which files directly or indirectly include this file:



### **Macros**

• #define CIRCUIT\_MODELLING\_CSIMULATOR\_H

### **Functions**

- double Evaluate\_Circuit (int vectorSize, int \*circuitVector, struct CircuitParameters parameters)

  Evaluates the performance of a circuit.
- double Evaluate\_Circuit (int vectorSize, int \*circuitVector, double &Recovery, double &Grade, struct CircuitParameters parameters)

Evaluates the performance, recovery, and grade of a circuit.

- double performance (double concentrateG, double concentrateW)
  - Computes the performance metric of a circuit.
- double grade (double concentrateG, double concentrateW)

Computes the grade of a circuit.

· double recovery (double concentrateG, double gerardiumFeed)

Computes the recovery of a circuit.

# 13.9.1 Detailed Description

Defines the functions and structures for simulating and evaluating circuits in a circuit modeling framework.

The CSimulator module provides functions for evaluating the performance, grade, and recovery of circuits in a circuit modeling framework. It includes definitions for circuit parameters and multiple overloads for evaluating circuits with or without additional parameters.

Date

Created on May 20, 2024

#### **Authors**

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- · Mingsheng Cai
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### 13.9.2 Macro Definition Documentation

### 13.9.2.1 CIRCUIT\_MODELLING\_CSIMULATOR\_H

#define CIRCUIT\_MODELLING\_CSIMULATOR\_H

### 13.9.3 Function Documentation

### 13.9.3.1 Evaluate\_Circuit() [1/2]

```
double Evaluate_Circuit (
                int vectorSize,
                int * circuitVector,
                double & Recovery,
                 double & Grade,
                 struct CircuitParameters parameters )
```

Evaluates the performance, recovery, and grade of a circuit.

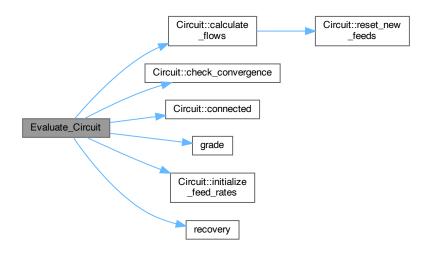
This function evaluates the performance, recovery, and grade of a circuit based on the given circuit vector and additional circuit parameters.

vectorSize	The size of the circuit vector.
circuitVector	Pointer to the array representing the circuit configuration.
Recovery	Reference to a double variable to store the recovery value.
Grade	Reference to a double variable to store the grade value.
parameters	Additional parameters for the circuit simulation.

### Returns

The performance metric of the circuit.

Here is the call graph for this function:



# 13.9.3.2 Evaluate\_Circuit() [2/2]

Evaluates the performance of a circuit.

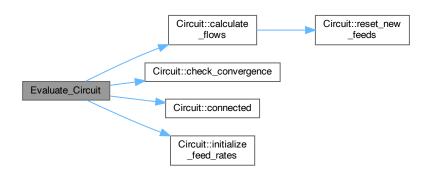
This function evaluates the performance of a circuit based on the given circuit vector and additional circuit parameters.

vectorSize	The size of the circuit vector.
circuitVector	Pointer to the array representing the circuit configuration.
parameters	Additional parameters for the circuit simulation.

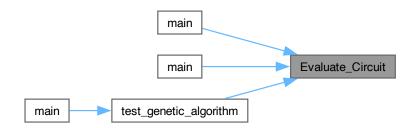
### Returns

The performance metric of the circuit.

Here is the call graph for this function:



Here is the caller graph for this function:



### 13.9.3.3 grade()

Computes the grade of a circuit.

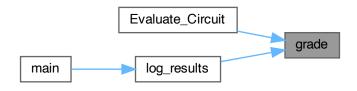
This function computes the grade of a circuit based on the given concentrate gerardium and waste values.

concentrateG	The amount of gerardium in the concentrate.
concentrateW	The amount of waste in the concentrate.

### Returns

The grade of the circuit.

Here is the caller graph for this function:



### 13.9.3.4 performance()

Computes the performance metric of a circuit.

This function computes the performance metric of a circuit based on the given concentrate gerardium and waste values.

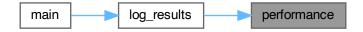
### **Parameters**

concentrateG	The amount of gerardium in the concentrate.
concentrateW	The amount of waste in the concentrate.

### Returns

The performance metric of the circuit.

Here is the caller graph for this function:



# 13.9.3.5 recovery()

Computes the recovery of a circuit.

This function computes the recovery of a circuit based on the given concentrate gerardium and gerardium feed values.

13.10 CSimulator.h

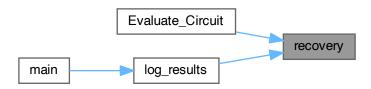
#### **Parameters**

concentrateG	The amount of gerardium in the concentrate.
gerardiumFeed	The amount of gerardium fed into the circuit.

### Returns

The recovery of the circuit.

Here is the caller graph for this function:



# 13.10 CSimulator.h

### Go to the documentation of this file.

```
00001
00023 #pragma once
00024
00025 #ifndef CIRCUIT_MODELLING_CSIMULATOR_H
00026 #define CIRCUIT_MODELLING_CSIMULATOR_H
00027
00028 #pragma once
00029
00030 #include <iostream>
00031 #include "circuit/CCircuit.h"
00032 #include "utils/Parameter.h"
00033
00045 double Evaluate_Circuit(int vectorSize, int *circuitVector, struct CircuitParameters parameters);
00046
00060 double Evaluate_Circuit(int vectorSize, int *circuitVector, double& Recovery, double& Grade, struct
     CircuitParameters parameters);
00061
00072 double performance(double concentrateG, double concentrateW);
00073
00083 double grade(double concentrateG, double concentrateW);
00084
00095 double recovery (double concentrateG, double gerardiumFeed);
00097 #endif //CIRCUIT_MODELLING_CSIMULATOR_H
```

# 13.11 include/Genetic\_Algorithm.h File Reference

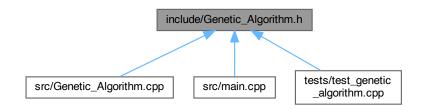
Defines the functions and structures for implementing a genetic algorithm.

```
#include <vector>
#include <stdio.h>
#include <cmath>
#include <array>
#include <cstdlib>
#include <ctime>
#include <algorithm>
#include <optional>
#include <random>
```

```
#include <memory>
#include "utils/Individual.h"
#include "utils/Parameter.h"
#include "utils/tqdm.hpp"
#include "operators/Selection/NaryTournamentSelection.h"
#include "operators/Selection/RankSelection.h"
#include "operators/Selection/RouletteWheelSelection.h"
#include "operators/Mutation/GuidedMutation.h"
#include "operators/Mutation/GenProgMutation.h"
#include "operators/Crossover/PureSinglePointCrossover.h"
#include "operators/Crossover/TwoPointCrossover.h"
#include dependency graph for Genetic_Algorithm.h:
```



This graph shows which files directly or indirectly include this file:



# Macros

#define GENETIC\_ALGORITHM\_H

### **Functions**

void initializeRandomSeed (const Algorithm\_Parameters &params, std::vector< std::mt19937 > &generators)

Initializes the random seed based on algorithm parameters.

• std::vector< Individual > initializePopulation (const Algorithm\_Parameters &params, int vectorSize, bool(&validity)(int, int \*), std::vector< std::mt19937 > &generators)

Initializes the population for the genetic algorithm.

• void elitism (const std::vector< Individual > &population, std::vector< Individual > &newPopulation, const Algorithm\_Parameters &params)

Apply elitism to preserve the best individuals in a new population.

- double acceptanceProbability (double currentEnergy, double newEnergy, double temperature)
  - Calculate the acceptance probability for a new state in Simulated Annealing.
- bool applySimulatedAnnealing (Individual &offspring, Individual &parent1, Individual &parent2, int vector
   \_size, double(&func)(int, int \*, struct CircuitParameters), bool(&validity)(int, int \*), double &Temp, const
   Algorithm\_Parameters &params, const CircuitParameters c\_params, std::mt19937 &generator)

Applies simulated annealing technique to decide whether to accept an offspring in a genetic algorithm.

int optimize (int vector\_size, int \*vector, double(&func)(int, int \*, struct CircuitParameters), bool(&validity)(int, int \*), struct Algorithm\_Parameters params, struct CircuitParameters c\_params)

Optimizes a solution using the genetic algorithm.

# 13.11.1 Detailed Description

Defines the functions and structures for implementing a genetic algorithm.

The Genetic\_Algorithm module provides functions for initializing populations, setting random seeds, and optimizing solutions using a genetic algorithm framework. This includes selection, crossover, and mutation operators.

Date

Created on May 20, 2024

### **Authors**

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# 13.11.2 Macro Definition Documentation

### 13.11.2.1 GENETIC\_ALGORITHM\_H

#define GENETIC\_ALGORITHM\_H

# 13.11.3 Function Documentation

# 13.11.3.1 acceptanceProbability()

Calculate the acceptance probability for a new state in Simulated Annealing.

This function calculates the probability with which a new state should be accepted over the current state, based on their respective energies and the current temperature of the system. If the new energy is better (higher value), the function returns 1.0, meaning the new state is always accepted. Otherwise, it returns a value calculated using the Boltzmann probability distribution which considers both the difference in energy and the current temperature.

currentEnergy	The energy of the current state.
newEnergy	The energy of the new state being considered.
temperature	The current temperature of the system (controls the probability threshold).

### Returns

double The probability of accepting the new state.

Here is the caller graph for this function:



### 13.11.3.2 applySimulatedAnnealing()

Applies simulated annealing technique to decide whether to accept an offspring in a genetic algorithm.

This function performs the simulated annealing decision process on a newly generated offspring. It compares the offspring's fitness to the average fitness of its parents using the simulated annealing acceptance probability, which depends on the current temperature of the system. The function also handles the cooling of the system temperature and validates the offspring using a provided validity function.

### **Parameters**

offspring	A reference to the offspring individual whose acceptance is being determined.
parent1	A reference to the first parent of the offspring.
parent2	A reference to the second parent of the offspring.
vector_size	The size of the individual's gene vector.
func	A function that calculates the fitness of an individual based on its genes.
validity	A function that checks the validity of an individual's gene configuration.
Temp	A reference to the current temperature used for the simulated annealing process.
params	A structure containing parameters relevant to the annealing process, such as the temperature decrement.
c_params	The circuit parameters used in the fitness evaluation function.
generator	A random number generator used for probabilistic decisions.

### Returns

bool Returns true if the offspring is accepted, false otherwise. The function can also modify the offspring directly, setting it to one of the parents if the temperature is low and the acceptance criteria are not met.

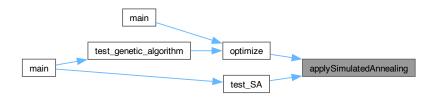
Note

The function directly modifies the temperature, decrementing it according to deltT from params. If the offspring is invalid (as determined by the validity function), it is immediately rejected and assigned a fitness value of INVALID\_FITNESS.

Here is the call graph for this function:



Here is the caller graph for this function:



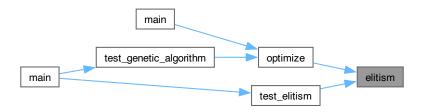
### 13.11.3.3 elitism()

Apply elitism to preserve the best individuals in a new population.

This function sorts the given population in descending order based on their fitness, then selectively copies the top percentage of individuals as specified by the elitePercentage in the Algorithm\_Parameters. This is used to ensure that the best individuals are preserved for the next generation, promoting genetic diversity and preventing loss of the best found solutions.

population	The current generation of individuals (const reference).
newPopulation	The next generation of individuals where elites will be added (reference).
params	The parameters of the algorithm including population size and elite percentage.

Here is the caller graph for this function:



# 13.11.3.4 initializePopulation()

Initializes the population for the genetic algorithm.

This function initializes the population for the genetic algorithm, ensuring each individual meets the validity criteria specified by the user.

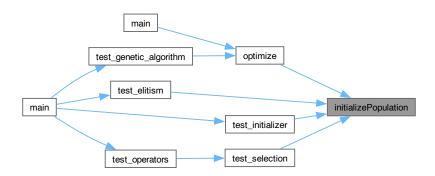
### **Parameters**

params	The algorithm parameters containing population size and other configurations.
vectorSize	The size of the gene vector for each individual.
validity	A function pointer to the validity checking function.
generators	A vector of random number generators for each thread.

### Returns

A vector of individuals representing the initial population.

Here is the caller graph for this function:



### 13.11.3.5 initializeRandomSeed()

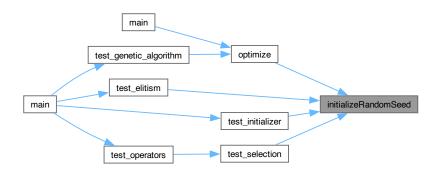
Initializes the random seed based on algorithm parameters.

This function initializes the random seed using the specified seed in the algorithm parameters. If the seed is zero, it uses the current time as the seed.

#### **Parameters**

params	The algorithm parameters containing the random seed.
generators	A vector of random number generators for each thread.

Here is the caller graph for this function:



### 13.11.3.6 optimize()

```
int optimize (
    int vector_size,
    int * vector,
    double(&)(int, int *, struct CircuitParameters) func,
    bool(&)(int, int *) validity,
    struct Algorithm_Parameters params,
    struct CircuitParameters c_params)
```

Optimizes a solution using the genetic algorithm.

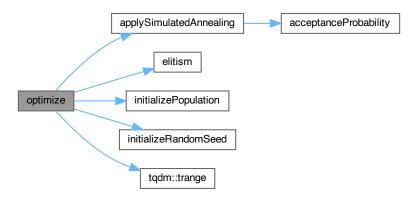
This function performs optimization using a genetic algorithm. It initializes the population, assesses the fitness of each individual, and iterates through generations applying selection, crossover, and mutation operators.

vector_size	The size of the gene vector.
vector	Pointer to the array representing the solution vector.
func	A function pointer to the fitness evaluation function.
validity	A function pointer to the validity checking function.
parameters	The algorithm parameters containing various configurations.

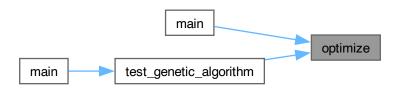
### Returns

An integer indicating the success (0) or failure (-1) of the optimization.

Here is the call graph for this function:



Here is the caller graph for this function:



# 13.12 Genetic\_Algorithm.h

### Go to the documentation of this file.

```
00001
00023 #pragma once
00024
00025 #ifndef GENETIC_ALGORITHM_H
00026 #define GENETIC_ALGORITHM_H
00028 #include <vector>
00029 #include <stdio.h>
00030 #include <cmath>
00031 #include <array>
00032 #include <cstdlib>
00033 #include <ctime>
00034 #include <algorithm>
00035 #include <optional>
00036 #include <random>
00037 #include <memory>
00038
00039 #include "utils/Individual.h"
00040 #include "utils/Parameter.h"
00041 #include "utils/tqdm.hpp"
00042 #include "operators/Selection/NaryTournamentSelection.h"
00042 #Include "operators/Selection/RaryFournamentSelection.n"
00043 #include "operators/Selection/RankSelection.h"
00045 #include "operators/Mutation/GuidedMutation.h"
00046 #include "operators/Mutation/GenProgMutation.h"
```

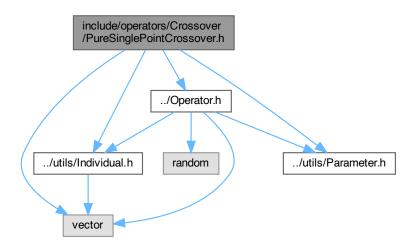
```
00047 #include "operators/Crossover/PureSinglePointCrossover.h"
00048 #include "operators/Crossover/TwoPointCrossover.h"
00049 #include "operators/Crossover/UniformCrossover.h"
00050
00060 void initializeRandomSeed(const Algorithm_Parameters &params, std::vector<std::mt19937> &generators);
00061
00074 std::vector<Individual>
00075 initializePopulation(const Algorithm_Parameters &params, int vectorSize, bool (&validity)(int, int *),
     std::vector<std::mt19937> &generators);
00076
00089 void elitism(const std::vector<Individual> &population, std::vector<Individual> &newPopulation,
00090
                   const Algorithm_Parameters &params);
00091
00092
00108 double acceptanceProbability(double currentEnergy, double newEnergy, double temperature);
00109
00137 bool applySimulatedAnnealing(Individual &offspring, Individual &parent1, Individual &parent2, int
      vector_size,
                                   double (&func)(int, int *, struct CircuitParameters), bool
      (&validity)(int, int *),
00139
                                   double &Temp, const Algorithm_Parameters &params, const CircuitParameters
     c_params,
00140
                                   std::mt19937 &generator);
00141
00156 int optimize(int vector_size, int *vector,
                 double (&func)(int, int *, struct CircuitParameters),
00158
                   bool (&validity)(int, int *),
00159
                   struct Algorithm_Parameters params,
00160
                   struct CircuitParameters c_params);
00161
00162 #endif //GENETIC_ALGORITHM_H
```

# 13.13 include/operators/Crossover/PureSinglePointCrossover.h File Reference

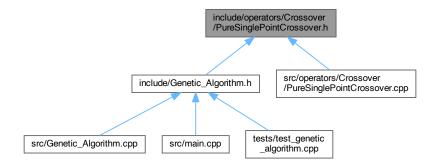
Defines the PureSinglePointCrossover class for performing single-point crossover in a genetic algorithm.

```
#include <vector>
#include "../Operator.h"
#include "../../utils/Individual.h"
#include "../../utils/Parameter.h"
```

Include dependency graph for PureSinglePointCrossover.h:



This graph shows which files directly or indirectly include this file:



### **Data Structures**

· class PureSinglePointCrossover

### **Macros**

• #define GENETIC ALGORITHM PURESINGLEPOINTCROSSOVER H

# 13.13.1 Detailed Description

Defines the PureSinglePointCrossover class for performing single-point crossover in a genetic algorithm. The PureSinglePointCrossover class provides methods to perform a single-point crossover on two parent individuals to generate an offspring individual. This class is designed to be used in genetic algorithms to combine genetic material from two parents to create new individuals, which can then be used in subsequent generations of the algorithm.

Date

Created on May 20, 2024

### **Authors**

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- · Melissa Y S Sim
- · Mingsheng Cai
- · Peifeng Tan
- · Yongwen Chen
- · Zihan Li

# 13.13.2 Macro Definition Documentation

# 13.13.2.1 GENETIC ALGORITHM PURESINGLEPOINTCROSSOVER H

#define GENETIC\_ALGORITHM\_PURESINGLEPOINTCROSSOVER\_H

# 13.14 PureSinglePointCrossover.h

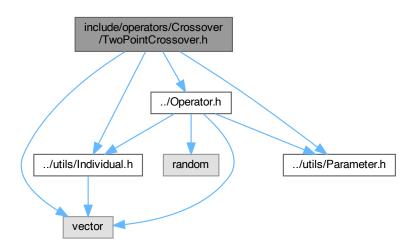
### Go to the documentation of this file.

```
00023 #pragma once
00024
00025 #ifndef GENETIC ALGORITHM PURESINGLEPOINTCROSSOVER H
00026 #define GENETIC_ALGORITHM_PURESINGLEPOINTCROSSOVER_H
00028 #include <vector>
00029 #include "../Operator.h"
00030 #include "../../utils/Individual.h"
00031 #include "../../utils/Parameter.h"
00032
00033 class PureSinglePointCrossover : public Operator {
00034 private:
00035
           Individual *offspring;
          const Individual *parent1;
const Individual *parent2;
00036
00037
00038
          double crossoverRate;
00040 public:
           PureSinglePointCrossover(const Algorithm_Parameters &params);
00047
00055
           void setParents(Individual &offspring, const Individual &parent1, const Individual &parent2)
00056
00065
           void setPopulation(const std::vector<Individual> &population, std::vector<Individual> &selected)
      { } ;
00066
00074
           void setIndividual(Individual &individual) {};
00075
00084
           void apply(std::mt19937 &generator) override;
00085 };
00086
00087 #endif //GENETIC_ALGORITHM_PURESINGLEPOINTCROSSOVER_H
```

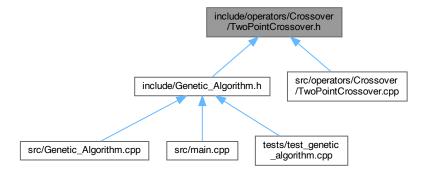
# 13.15 include/operators/Crossover/TwoPointCrossover.h File Reference

Defines the TwoPointCrossover class for performing two-point crossover in a genetic algorithm.

```
#include <vector>
#include "../Operator.h"
#include "../../utils/Individual.h"
#include "../../utils/Parameter.h"
Include dependency graph for TwoPointCrossover.h:
```



This graph shows which files directly or indirectly include this file:



### **Data Structures**

class TwoPointCrossover

### **Macros**

• #define GENETIC\_ALGORITHM\_TWOPOINTCROSSOVER\_H

# 13.15.1 Detailed Description

Defines the TwoPointCrossover class for performing two-point crossover in a genetic algorithm.

The TwoPointCrossover class provides methods to perform a two-point crossover on two parent individuals to generate an offspring individual. This class is designed to be used in genetic algorithms to combine genetic material from two parents to create new individuals, which can then be used in subsequent generations of the algorithm.

Date

Created on May 20, 2024

# **Authors**

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- · Melissa Y S Sim
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- · Peifeng Tan
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# 13.15.2 Macro Definition Documentation

### 13.15.2.1 GENETIC\_ALGORITHM\_TWOPOINTCROSSOVER\_H

#define GENETIC\_ALGORITHM\_TWOPOINTCROSSOVER\_H

13.16 TwoPointCrossover.h 177

# 13.16 TwoPointCrossover.h

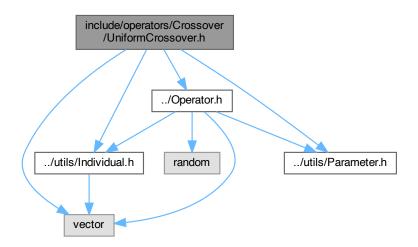
### Go to the documentation of this file.

```
00024 #pragma once
00025
00026 #ifndef GENETIC ALGORITHM TWOPOINTCROSSOVER H
00027 #define GENETIC_ALGORITHM_TWOPOINTCROSSOVER_H
00029 #include <vector>
00030 #include "../Operator.h"
00031 #include "../../utils/Individual.h"
00032 #include "../../utils/Parameter.h"
00033
00034 class TwoPointCrossover : public Operator {
00035 private:
00036
           Individual *offspring;
           const Individual *parent1;
const Individual *parent2;
00037
00038
00039
           double crossoverRate;
00041 public:
00047
           TwoPointCrossover(const Algorithm_Parameters &params);
00048
00056
           void setParents(Individual &offspring, const Individual &parent1, const Individual &parent2)
00057
00066
           void setPopulation(const std::vector<Individual> &population, std::vector<Individual> &selected)
      { } ;
00067
00075
           void setIndividual(Individual &individual) {};
00076
00085
           void apply(std::mt19937 &generator) override;
00086 };
00087
00088 #endif //GENETIC_ALGORITHM_TWOPOINTCROSSOVER_H
```

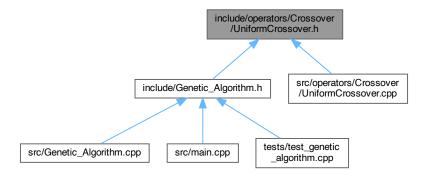
# 13.17 include/operators/Crossover/UniformCrossover.h File Reference

Defines the UniformCrossover class for performing uniform crossover in a genetic algorithm.

```
#include <vector>
#include "../Operator.h"
#include "../../utils/Individual.h"
#include "../../utils/Parameter.h"
Include dependency graph for UniformCrossover.h:
```



This graph shows which files directly or indirectly include this file:



### **Data Structures**

· class UniformCrossover

#### **Macros**

• #define GENETIC\_ALGORITHM\_UNIFORMCROSSOVER\_H

# 13.17.1 Detailed Description

Defines the UniformCrossover class for performing uniform crossover in a genetic algorithm.

The UniformCrossover class provides methods to perform a uniform crossover on two parent individuals to generate an offspring individual. This class is designed to be used in genetic algorithms to combine genetic material from two parents to create new individuals, which can then be used in subsequent generations of the algorithm.

Date

Created on May 20, 2024

# **Authors**

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### 13.17.2 Macro Definition Documentation

### 13.17.2.1 GENETIC ALGORITHM UNIFORMCROSSOVER H

#define GENETIC\_ALGORITHM\_UNIFORMCROSSOVER\_H

13.18 UniformCrossover.h 179

# 13.18 UniformCrossover.h

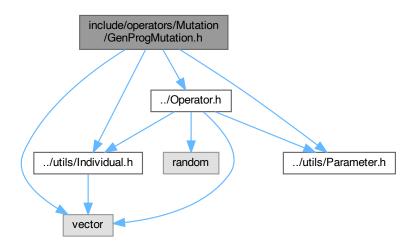
### Go to the documentation of this file.

```
00024 #pragma once
00025
00026 #ifndef GENETIC ALGORITHM UNIFORMCROSSOVER H
00027 #define GENETIC ALGORITHM UNIFORMCROSSOVER H
00029 #include <vector>
00030 #include "../Operator.h"
00031 #include "../../utils/Individual.h"
00032 #include "../../utils/Parameter.h"
00033
00034 class UniformCrossover : public Operator {
00035 private:
00036
           Individual *offspring;
           const Individual *parent1;
const Individual *parent2;
00037
00038
00039
           double crossoverRate;
00041 public:
00047
           UniformCrossover(const Algorithm_Parameters &params);
00048
00056
           void setParents(Individual &offspring, const Individual &parent1, const Individual &parent2)
00057
00066
           void setPopulation(const std::vector<Individual> &population, std::vector<Individual> &selected)
      { } ;
00067
00075
           void setIndividual(Individual &individual) {};
00076
00085
           void apply(std::mt19937 &generator) override;
00086 };
00087
00088 #endif //GENETIC_ALGORITHM_UNIFORMCROSSOVER_H
```

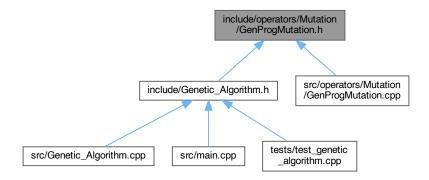
# 13.19 include/operators/Mutation/GenProgMutation.h File Reference

Defines the GenProgMutation class for performing mutation operations in a genetic programming algorithm.

```
#include <vector>
#include "../Operator.h"
#include "../../utils/Individual.h"
#include "../../utils/Parameter.h"
Include dependency graph for GenProgMutation.h:
```



This graph shows which files directly or indirectly include this file:



### **Data Structures**

· class GenProgMutation

### **Macros**

• #define GENETIC\_ALGORITHM\_GENPROGMUTATION\_H

# 13.19.1 Detailed Description

Defines the GenProgMutation class for performing mutation operations in a genetic programming algorithm. The GenProgMutation class provides methods to perform mutation operations on individuals in a genetic programming algorithm. This class is designed to introduce variations into the population by modifying individuals' genetic material, which can then be used in subsequent generations of the algorithm.

Date

Created on May 20, 2024

# **Authors**

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### 13.19.2 Macro Definition Documentation

### 13.19.2.1 GENETIC ALGORITHM GENPROGMUTATION H

#define GENETIC\_ALGORITHM\_GENPROGMUTATION\_H

# 13.20 GenProgMutation.h

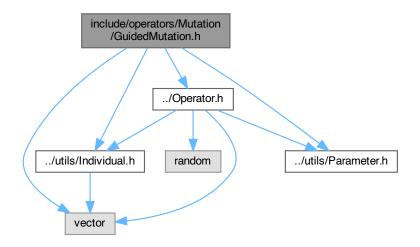
Go to the documentation of this file.

```
00023 #pragma once
00024
00025 #ifndef GENETIC ALGORITHM GENPROGMUTATION H
00026 #define GENETIC ALGORITHM GENPROGMUTATION H
00027
00028 #include <vector>
00029 #include "../Operator.h"
00030 #include "../../utils/Individual.h"
00031 #include "../../utils/Parameter.h"
00032
00033 class GenProgMutation : public Operator {
00034 public:
00041
            GenProgMutation(int vectorSize, const Algorithm_Parameters &params);
00042
00048
            void setIndividual(Individual &individual) override;
00049
            \verb|void| \textbf{setPopulation}| (\texttt{const}| \textbf{std}:: \texttt{vector} < \texttt{Individual} > \texttt{\&population}|, \textbf{std}:: \texttt{vector} < \texttt{Individual} > \texttt{\&selected}|) \\
00058
      { };
00059
00069
            void setParents(Individual &offspring, const Individual &parent1, const Individual &parent2) {};
00070
            void apply(std::mt19937 &generator);
00079
00080
00081 private:
00082
            Individual *individual;
00083
            int numUnits;
00084
            double mutationRate;
00085 };
00086
00087 #endif // GENPROG_MUTATION_H
```

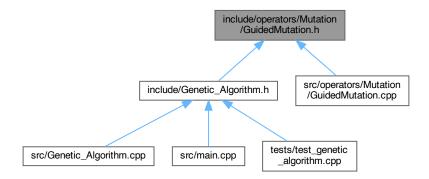
# 13.21 include/operators/Mutation/GuidedMutation.h File Reference

Defines the GuidedMutation class for performing guided mutation operations in a genetic algorithm.

```
#include <vector>
#include "../Operator.h"
#include "../../utils/Individual.h"
#include "../../utils/Parameter.h"
Include dependency graph for GuidedMutation.h:
```



This graph shows which files directly or indirectly include this file:



### **Data Structures**

· class GuidedMutation

### **Macros**

• #define GENETIC\_ALGORITHM\_GUIDEDMUTATION\_H

# 13.21.1 Detailed Description

Defines the GuidedMutation class for performing guided mutation operations in a genetic algorithm.

The GuidedMutation class provides methods to perform guided mutation operations on individuals in a genetic algorithm. This class is designed to introduce targeted variations into the population by modifying individuals' genetic material based on specific rules or heuristics, which can then be used in subsequent generations of the algorithm.

Date

Created on May 20, 2024

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### 13.21.2 Macro Definition Documentation

# 13.21.2.1 GENETIC\_ALGORITHM\_GUIDEDMUTATION\_H

#define GENETIC\_ALGORITHM\_GUIDEDMUTATION\_H

13.22 GuidedMutation.h

# 13.22 GuidedMutation.h

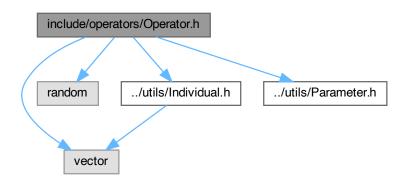
Go to the documentation of this file.

```
00023 #pragma once
00024
00025 #ifndef GENETIC ALGORITHM GUIDEDMUTATION H
00026 #define GENETIC_ALGORITHM_GUIDEDMUTATION_H
00028 #include <vector>
00029 #include "../Operator.h"
00030 #include "../../utils/Individual.h"
00031 #include "../../utils/Parameter.h"
00032
00033 class GuidedMutation : public Operator {
00034 private:
00035
           Individual *individual;
00036
           int numUnits;
00037
          double mutationRate;
00039 public:
          GuidedMutation(int vectorSize, const Algorithm_Parameters &params);
00046
00047
00053
           void setIndividual(Individual &individual) override;
00054
00063
          void setPopulation(const std::vector<Individual> &population, std::vector<Individual> &selected)
      { } ;
00064
00074
           void setParents(Individual &offspring, const Individual &parent1, const Individual &parent2) {};
00075
00084
           void apply(std::mt19937 &generator) override;
00085 };
00086
00087
00088 #endif //GENETIC_ALGORITHM_GUIDEDMUTATION_H
```

# 13.23 include/operators/Operator.h File Reference

Defines the abstract base class for genetic algorithm operators.

```
#include <vector>
#include <random>
#include "../utils/Individual.h"
#include "../utils/Parameter.h"
Include dependency graph for Operator.h:
```



This graph shows which files directly or indirectly include this file:



### **Data Structures**

· class Operator

Abstract base class for genetic algorithm operators.

### **Macros**

• #define GENETIC\_ALGORITHM\_OPERATOR\_H

# 13.23.1 Detailed Description

Defines the abstract base class for genetic algorithm operators.

The Operator class serves as an abstract base class for various genetic algorithm operators, such as selection, crossover, and mutation. Each specific operator must inherit from this class and implement the pure virtual function apply(). This design allows for a consistent interface and enables polymorphism, making it easier to manage and apply different operators within a genetic algorithm framework.

Date

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### 13.23.2 Macro Definition Documentation

### 13.23.2.1 GENETIC\_ALGORITHM\_OPERATOR\_H

#define GENETIC\_ALGORITHM\_OPERATOR\_H

# 13.24 Operator.h

### Go to the documentation of this file.

```
00001
00025 #pragma once
00026
00027 #ifndef GENETIC_ALGORITHM_OPERATOR_H
00028 #define GENETIC_ALGORITHM_OPERATOR_H
00029
00030 #include <vector>
00031 #include <random>
```

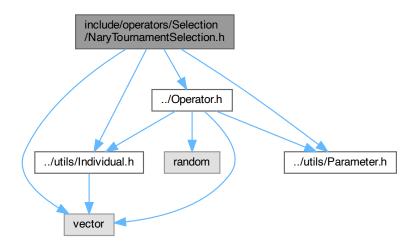
```
00032
00033 #include "../utils/Individual.h"
00034 #include "../utils/Parameter.h"
00035
00046 class Operator {
00047 public:
          virtual ~Operator() = default;
00055
00062
          virtual void apply(std::mt19937 &generator) = 0;
00063
          virtual void setPopulation(const std::vector<Individual> &population, std::vector<Individual>
00073
     &selected) = 0;
00074
00083
          virtual void setIndividual(Individual &individual) = 0;
00084
00094
          virtual void setParents(Individual &offspring, const Individual &parent1, const Individual
      &parent2) = 0;
00095 };
00096
00097 #endif //GENETIC_ALGORITHM_OPERATOR_H
```

# 13.25 include/operators/Selection/NaryTournamentSelection.h File Reference

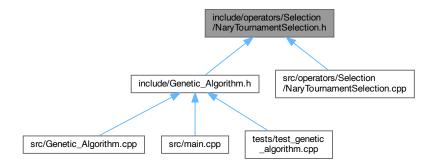
Defines the NaryTournamentSelection class for performing n-ary tournament selection in a genetic algorithm.

```
#include <vector>
#include "../Operator.h"
#include "../../utils/Individual.h"
#include "../../utils/Parameter.h"
```

Include dependency graph for NaryTournamentSelection.h:



This graph shows which files directly or indirectly include this file:



### **Data Structures**

· class NaryTournamentSelection

### **Macros**

#define GENETIC\_ALGORITHM\_NARYTOURNAMENTSELECTION\_H

# 13.25.1 Detailed Description

Defines the NaryTournamentSelection class for performing n-ary tournament selection in a genetic algorithm. The NaryTournamentSelection class provides methods to perform n-ary tournament selection on a population of individuals to select individuals for the next generation. This class is designed to be used in genetic algorithms to select the fittest individuals based on tournament competition among a subset of the population.

Date

Created on May 20, 2024

# **Authors**

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- Peifeng Tan
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- · Zihan Li

# 13.25.2 Macro Definition Documentation

# 13.25.2.1 GENETIC\_ALGORITHM\_NARYTOURNAMENTSELECTION\_H

#define GENETIC\_ALGORITHM\_NARYTOURNAMENTSELECTION\_H

# 13.26 NaryTournamentSelection.h

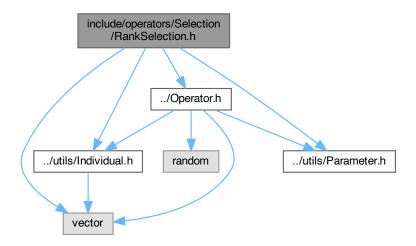
### Go to the documentation of this file.

```
00023 #pragma once
00024
00025 #ifndef GENETIC_ALGORITHM_NARYTOURNAMENTSELECTION_H
00026 #define GENETIC_ALGORITHM_NARYTOURNAMENTSELECTION_H
00029 #include <vector>
00030 #include "../Operator.h"
00031 #include "../../utils/Individual.h"
00032 #include "../../utils/Parameter.h"
00033
00034 class NaryTournamentSelection : public Operator {
00035 private:
00036
          std::vector<Individual> *selected;
00037
           const std::vector<Individual> *population;
00038
           int tournamentSize;
00040 public:
00046
          NaryTournamentSelection(const Algorithm_Parameters &params);
00047
00054
           void setPopulation(const std::vector<Individual> &population, std::vector<Individual> &selected)
      override;
00055
00063
          void setIndividual(Individual &individual) {};
00064
00074
           void setParents(Individual &offspring, const Individual &parent1, const Individual &parent2) {};
00075
00084
           void apply(std::mt19937 &generator) override;
00085 };
00086
00087 #endif //GENETIC_ALGORITHM_NARYTOURNAMENTSELECTION_H
```

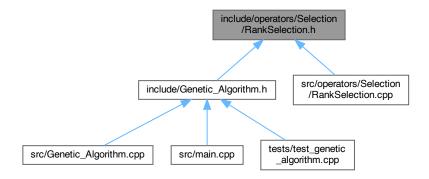
# 13.27 include/operators/Selection/RankSelection.h File Reference

Defines the RankSelection class for performing rank-based selection in a genetic algorithm.

```
#include <vector>
#include "../Operator.h"
#include "../../utils/Individual.h"
#include "../../utils/Parameter.h"
Include dependency graph for RankSelection.h:
```



This graph shows which files directly or indirectly include this file:



### **Data Structures**

· class RankSelection

### Macros

#define GENETIC\_ALGORITHM\_RANKSELECTION\_H

# 13.27.1 Detailed Description

Defines the RankSelection class for performing rank-based selection in a genetic algorithm.

The RankSelection class provides methods to perform rank-based selection on a population of individuals to select individuals for the next generation. This class is designed to be used in genetic algorithms to select the fittest individuals based on their rank within the population.

Date

Created on May 21, 2024

### **Authors**

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# 13.27.2 Macro Definition Documentation

# 13.27.2.1 GENETIC\_ALGORITHM\_RANKSELECTION\_H

#define GENETIC\_ALGORITHM\_RANKSELECTION\_H

13.28 RankSelection.h 189

# 13.28 RankSelection.h

# Go to the documentation of this file.

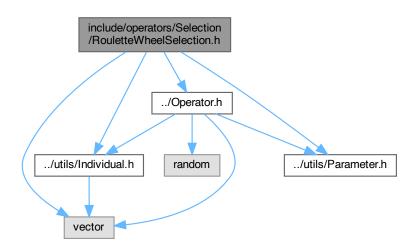
```
00023 #pragma once
00024
00025 #ifndef GENETIC ALGORITHM RANKSELECTION H
00026 #define GENETIC_ALGORITHM_RANKSELECTION_H
00029 #include <vector>
00030 #include "../Operator.h"
00031 #include "../../utils/Individual.h"
00032 #include "../../utils/Parameter.h"
00033
00034 class RankSelection : public Operator {
00035 private:
00036
          std::vector<Individual> *selected;
00037
           const std::vector<Individual> *population;
00038
           int populationSize;
00040 public:
00046
           RankSelection(const Algorithm_Parameters &params);
00047
00054
           void setPopulation(const std::vector<Individual> &population, std::vector<Individual> &selected)
      override;
00055
00063
          void setIndividual(Individual &individual) {};
00064
00074
           void setParents(Individual &offspring, const Individual &parent1, const Individual &parent2) {};
00075
00084
           void apply(std::mt19937 &generator) override;
00085 };
00086
00087 #endif //GENETIC_ALGORITHM_NARYTOURNAMENTSELECTION_H
```

# 13.29 include/operators/Selection/RouletteWheelSelection.h File Reference

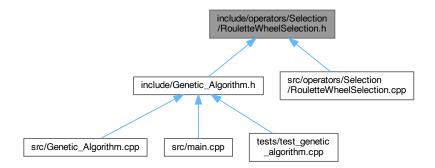
Defines the RouletteWheelSelection class for performing roulette wheel selection in a genetic algorithm.

```
#include <vector>
#include "../Operator.h"
#include "../../utils/Individual.h"
#include "../../utils/Parameter.h"
```

Include dependency graph for RouletteWheelSelection.h:



This graph shows which files directly or indirectly include this file:



### **Data Structures**

· class RouletteWheelSelection

### **Macros**

• #define GENETIC\_ALGORITHM\_ROULETTEWHEELSELECTION\_H

# 13.29.1 Detailed Description

Defines the RouletteWheelSelection class for performing roulette wheel selection in a genetic algorithm. The RouletteWheelSelection class provides methods to perform roulette wheel selection on a population of individuals to select individuals for the next generation. This class is designed to be used in genetic algorithms to select individuals based on their fitness proportionally.

Date

Created on May 21, 2024

### **Authors**

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# 13.29.2 Macro Definition Documentation

# 13.29.2.1 GENETIC\_ALGORITHM\_ROULETTEWHEELSELECTION\_H

#define GENETIC\_ALGORITHM\_ROULETTEWHEELSELECTION\_H

# 13.30 RouletteWheelSelection.h

Go to the documentation of this file.

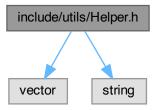
```
00023 #pragma once
00024
00025 #ifndef GENETIC ALGORITHM ROULETTEWHEELSELECTION H
00026 #define GENETIC_ALGORITHM_ROULETTEWHEELSELECTION_H
00028 #include <vector>
00029 #include "../Operator.h"
00030 #include "../../utils/Individual.h"
00031 #include "../../utils/Parameter.h"
00032
00033 class RouletteWheelSelection: public Operator {
00034 private:
00035
          std::vector<Individual> *selected;
00036
           const std::vector<Individual> *population;
00037
           int populationSize;
00039 public:
00045
          RouletteWheelSelection(const Algorithm_Parameters &params);
00053
           void setPopulation(const std::vector<Individual> &population, std::vector<Individual> &selected);
00054
00062
           void setIndividual(Individual &individual) {};
00063
00073
           void setParents(Individual &offspring, const Individual &parent1, const Individual &parent2) {};
00074
00083
           void apply(std::mt19937 &generator) override;
00084 };
00085
00086 #endif //GENETIC_ALGORITHM_ROULETTEWHEELSELECTION_H
```

# 13.31 include/utils/Helper.h File Reference

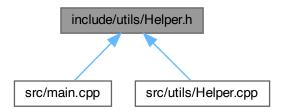
Header file containing helper functions for logging and time retrieval.

```
#include <vector>
#include <string>
```

Include dependency graph for Helper.h:



This graph shows which files directly or indirectly include this file:



### **Macros**

• #define HELPER\_H

### **Functions**

• std::string get\_current\_time\_str ()

Retrieves the current time as a formatted string.

• void log\_results (const std::vector< int > &vector, double elapsed\_time, double performance, double recovery, double grade)

Logs the results of the genetic algorithm optimization to a file.

# 13.31.1 Detailed Description

Header file containing helper functions for logging and time retrieval.

This header file declares functions for retrieving the current time as a string and logging the results of a genetic algorithm optimization process. The logging function saves performance metrics and the final configuration of the optimized circuit to a log file.

Date

Created on May 23, 2024

### **Authors**

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### 13.31.2 Macro Definition Documentation

# 13.31.2.1 HELPER\_H

#define HELPER\_H

# 13.31.3 Function Documentation

# 13.31.3.1 get\_current\_time\_str()

```
std::string get_current_time_str ()
```

Retrieves the current time as a formatted string.

This function returns the current time formatted as "YYYY-MM-DD HH-MM-SS".

### Returns

A string representing the current time.

Here is the caller graph for this function:



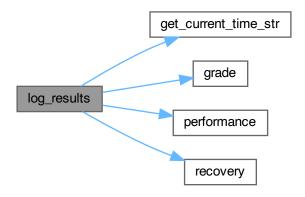
# 13.31.3.2 log\_results()

Logs the results of the genetic algorithm optimization to a file.

This function creates a new log folder with the current timestamp, and saves the performance metrics, recovery, grade, and the final configuration of the optimized circuit to a log file.

vector	The final configuration vector of the optimized circuit.
elapsed_time	The time taken to complete the optimization process, in seconds.
performance	The performance metric of the optimized circuit.
recovery	The recovery metric of the optimized circuit.
grade	The grade metric of the optimized circuit.

Here is the call graph for this function:



Here is the caller graph for this function:



# 13.32 Helper.h

Go to the documentation of this file.

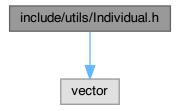
```
00001
00023 #pragma once
00024
00025 #ifndef HELPER_H
00026 #define HELPER_H
00027
00028 #include <vector>
00029 #include <string>
00030
00030
00038 std::string get_current_time_str();
00039
00052 void log_results(const std::vector<int>& vector, double elapsed_time, double performance, double recovery, double grade);
00053
00054 #endif // HELPER_H
```

# 13.33 include/utils/Individual.h File Reference

Defines the Individual structure for representing individuals in a genetic algorithm.

#include <vector>

Include dependency graph for Individual.h:



This graph shows which files directly or indirectly include this file:



# **Data Structures**

struct Individual

Represents an individual in a genetic algorithm.

# Macros

#define GENETIC\_ALGORITHM\_INDIVIDUAL\_H

# 13.33.1 Detailed Description

Defines the Individual structure for representing individuals in a genetic algorithm.

The Individual structure is used to represent a single individual in a genetic algorithm population. It contains a vector of genes that represent the circuit configuration and a fitness value that represents the quality or performance of the circuit configuration.

Date

Created on May 20, 2024

# Authors

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#### 13.33.2 Macro Definition Documentation

# 13.33.2.1 GENETIC\_ALGORITHM\_INDIVIDUAL\_H

#define GENETIC\_ALGORITHM\_INDIVIDUAL\_H

# 13.34 Individual.h

#### Go to the documentation of this file.

```
00001
00023 #pragma once
00024
00025 #ifndef GENETIC_ALGORITHM_INDIVIDUAL_H
00026 #define GENETIC_ALGORITHM_INDIVIDUAL_H
00028 #include <vector>
00029
00030
00039 struct Individual {
00040
         std::vector<int> genes;
00041
          double fitness;
00042 };
00043
00044 #endif //GENETIC_ALGORITHM_INDIVIDUAL_H
```

# 13.35 include/utils/Parameter.h File Reference

Defines the Algorithm\_Parameters structure for configuring the parameters of a genetic algorithm. This graph shows which files directly or indirectly include this file:



#### **Data Structures**

• struct Algorithm\_Parameters

Configures the parameters of a genetic algorithm.

• struct CircuitParameters

Defines the parameters for the circuit simulator.

#### Macros

• #define GENETIC\_ALGORITHM\_PARAMETER\_H

# 13.35.1 Detailed Description

Defines the Algorithm\_Parameters structure for configuring the parameters of a genetic algorithm.

The Algorithm\_Parameters structure is used to configure various parameters of a genetic algorithm, such as the number of generations, population size, tournament size, crossover rate, mutation rate, elite percentage, and random seed. These parameters control the behavior of the genetic algorithm and can be customized to optimize performance.

Date

Created on May 20, 2024

13.36 Parameter.h

#### **Authors**

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- · Geyu Ji
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- · Mingsheng Cai
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- · Zihan Li

#### 13.35.2 Macro Definition Documentation

# 13.35.2.1 GENETIC\_ALGORITHM\_PARAMETER\_H

#define GENETIC\_ALGORITHM\_PARAMETER\_H

# 13.36 Parameter.h

#### Go to the documentation of this file.

```
00001
00024 #pragma once
00025
00026 #ifndef GENETIC_ALGORITHM_PARAMETER_H
00027 #define GENETIC_ALGORITHM_PARAMETER_H
00028
00038 struct Algorithm_Parameters
        int maxGenerations = 1000;
00039
          int populationSize = 100;
std::string selection = "NaryTournamentSelection";
00040
00041
00042
          int tournamentSize = 3;
          std::string crossover = "TwoPointCrossover";
double crossoverRate = 0.8;
00043
00044
          double crossoverkate - v.o,
std::string mutation = "GuidedMutation";
double mutationRate = 0.1;
double elitePercentage = 0.2;
00045
00046
00048
          double initialTemp = 1000.0;
00049
          double deltT = 1.0;
00050
          unsigned int randomSeed = 0;
00063
          Algorithm_Parameters(int gens, int popSize, std::string selection, int tourSize,
00064
                                  std::string crossover, double crossRate, std::string mutation, double
00065
                                  double elitePerc, double initialTemp, double deltT, double randomSeed)
00066
                    : maxGenerations(gens), populationSize(popSize), selection(selection),
      tournamentSize(tourSize),
00067
                     crossover(crossover), crossoverRate(crossRate), mutation(mutation),
      mutationRate(mutRate).
00068
                     elitePercentage(elitePerc), initialTemp(initialTemp), deltT(deltT),
      randomSeed(randomSeed) {}
00069 };
00070
00079 struct CircuitParameters {
00080
          double tolerance = 1e-6;
           int maxIterations = 1000;
00081
          CircuitParameters(double tol, int maxIter) : tolerance(tol), maxIterations(maxIter) {}
00090 };
00091
00092 #endif //GENETIC_ALGORITHM_PARAMETER_H
```

# 13.37 include/utils/tqdm.hpp File Reference

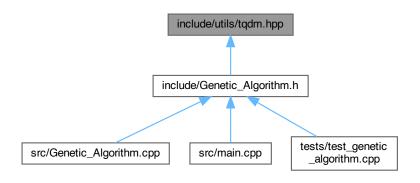
```
#include <chrono>
#include <cmath>
#include <iomanip>
#include <iostream>
#include <iterator>
#include <sstream>
#include <type_traits>
```

#include <sys/ioctl.h>
#include <unistd.h>

Include dependency graph for tqdm.hpp:



This graph shows which files directly or indirectly include this file:



#### **Data Structures**

· class tqdm::Chronometer

A simple chronometer to measure elapsed time.

· class tqdm::progress\_bar

A class representing a progress bar.

class tqdm::iter\_wrapper< ForwardIter, Parent >

A wrapper class for iterators to update the progress bar.

class tqdm::tqdm\_for\_lvalues< ForwardIter, EndIter >

A class for progress bars that iterate over Ivalues.

class tqdm::tqdm\_for\_rvalues< Container >

A class for progress bars that iterate over rvalues.

class tqdm::int\_iterator< IntType >

A random access iterator for integers.

class tqdm::range< IntType >

A range of integers represented by iterators.

• class tqdm::timing\_iterator\_end\_sentinel

An end sentinel for timing iterators.

· class tqdm::timing\_iterator

A forward iterator that returns the elapsed time.

· struct tqdm::timer

A timer that can be used to create a timing iterator.

· class tqdm::tqdm\_timer

A progress bar for a timer.

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#### **Namespaces**

namespace tqdm

#### **Typedefs**

```
using tgdm::index = std::ptrdiff t
```

```
using tqdm::time_point_t = std::chrono::time_point<std::chrono::steady_clock>
```

#### **Functions**

• double tqdm::elapsed\_seconds (time\_point\_t from, time\_point\_t to)

Calculate the elapsed seconds between two time points.

• void tqdm::clamp (double &x, double a, double b)

Clamp a value between two bounds.

int tqdm::get\_terminal\_width ()

Get the width of the terminal window.

template < class Container >

```
tqdm::tqdm_for_lvalues (Container &) -> tqdm_for_lvalues < typename Container::iterator >
```

• template < class Container >

```
tqdm::tqdm_for_lvalues (const Container &) -> tqdm_for_lvalues < typename Container::const_iterator >
```

template < class Container >

```
tgdm::tgdm for rvalues (Container &&) -> tgdm for rvalues < Container >
```

• template<class ForwardIter >

auto tqdm::tqdm (const ForwardIter &first, const ForwardIter &last)

Create a tqdm progress bar for a range of iterators.

• template<class ForwardIter >

auto tqdm::tqdm (const ForwardIter &first, const ForwardIter &last, index total)

Create a tqdm progress bar for a range of iterators with a specified total size.

• template<class Container >

```
auto tqdm::tqdm (const Container &C)
```

Create a tqdm progress bar for a container.

template < class Container >

```
auto tqdm::tqdm (Container &C)
```

Create a tqdm progress bar for a container.

template < class Container >

```
auto tqdm::tqdm (Container &&C)
```

Create a tqdm progress bar for a container (rvalue reference).

template < class IntType >

```
auto tqdm::trange (IntType first, IntType last)
```

Create a tqdm progress bar for a range of integers.

 $\bullet \ \ \mathsf{template}{<}\mathsf{class} \ \mathsf{IntType}>$ 

```
auto tqdm::trange (IntType last)
```

Create a tqdm progress bar for a range of integers.

• auto tqdm::tqdm (timer t)

Create a tqdm progress bar for a timer.

# 13.38 tqdm.hpp

Go to the documentation of this file.

```
00001

00053  #pragma once

00054

00055  #include <chrono>

00056  #include <cmath>

00057  #include <iomanip>

00058  #include <iostream>
```

```
00059 #include <iterator>
00060 #include <sstream>
00061 #include <type_traits>
00062 #include <sys/ioctl.h>
00063 #include <unistd.h>
00064
                     ----- chrono stuff -----
00065 //
00066
00067 namespace tqdm {
         using index = std::ptrdiff_t; // maybe std::size_t, but I hate unsigned types.
00068
00069
         using time_point_t = std::chrono::time_point<std::chrono::steady_clock>;
00070
08000
          inline double elapsed_seconds(time_point_t from, time_point_t to) {
00081
             using seconds = std::chrono::duration<double>;
00082
              return std::chrono::duration_cast<seconds>(to - from).count();
00083
00084
00088
         class Chronometer {
00089
         public:
00093
             Chronometer() : start_(std::chrono::steady_clock::now()) {}
00094
00100
              double reset() {
00101
                 auto previous = start_;
00102
                  start_ = std::chrono::steady_clock::now();
00103
00104
                 return elapsed_seconds(previous, start_);
00105
              }
00106
00112
              [[nodiscard]] double peek() const {
00113
                 auto now = std::chrono::steady_clock::now();
00114
00115
                  return elapsed seconds(start , now);
00116
00117
00123
             [[nodiscard]] time_point_t get_start() const { return start_; }
00124
00125
         private:
00126
             time_point_t start_;
00127
00128
00129 // ----- progress_bar -----
00130
00138
          inline void clamp (double &x, double a, double b) {
00139
              if (x < a) x = a;
00140
              if (x > b) x = b;
00141
00142
00148
         inline int get_terminal_width() {
00149
              struct winsize w;
              ioctl(STDOUT_FILENO, TIOCGWINSZ, &w);
00150
00151
             return w.ws_col;
00152
00153
00157
         class progress_bar {
         public:
00158
00162
             void restart() {
00163
                 chronometer_.reset();
00164
                  refresh_.reset();
                 symbol_index_ = 0;
00165
00166
             }
00167
             void update(double progress) {
00173
00174
                 clamp(progress, 0, 1);
00175
                  symbol_index_++;
00176
00177
                  if (time_since_refresh() > min_time_per_update_ || progress == 0 ||
00178
                     progress == 1) {
                      reset_refresh_timer();
00179
00180
                     display(progress);
00181
00182
                  suffix_.str("");
00183
              }
00184
00190
              void set_ostream(std::ostream &os) { os = &os; }
00191
00197
              void set_prefix(std::string s) { prefix_ = std::move(s); }
00198
00204
              void set_bar_size(int size) { bar_size_ = size; }
00205
00211
              void set_min_update_time(double time) { min_time_per_update_ = time; }
00212
00219
              template<class T>
00220
              progress_bar &operator«(const T &t) {
00221
                  suffix_ « t;
00222
                  return *this;
00223
              }
00224
```

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```
00230
              double elapsed_time() const { return chronometer_.peek(); }
00231
          private:
00232
00238
              void display(double progress) {
00239
                  auto flags = os ->flags();
00240
                  double t = chronometer_.peek();
00242
                  double eta = t / progress - t;
00243
00244
                  std::stringstream bar;
00245
                  // Lengthy prefix
00246
00247
                   std::stringstream temp;
00248
                   temp « prefix_ « '{' « std::fixed « std::setprecision(1)
                        « std::setw(5) « 100 * progress « "%}
00249
00250
                        « " (" « t « "s < " « eta « "s) ";
00251
00252
                   // Dynamic bar size
00253
                   int terminal_width = get_terminal_width();
                   int fixed_length = temp.str().size() + suffix_.str().size();
00254
00255
                  bar_size_ = terminal_width - fixed_length - 10; // Safety margin
00256
                  00257
00258
00259
00260
                  print_bar(bar, progress);
00261
00262
                  bar « " (" « t « "s < " « eta « "s) ";
00263
00264
                   std::string sbar = bar.str();
00265
                   std::string suffix = suffix .str();
00266
00267
                   index out_size = sbar.size() + suffix.size();
00268
                   term_cols_ = std::max(term_cols_, out_size);
00269
                  index num_blank = term_cols_ - out_size;
00270
00271
                   (*os_) « sbar « suffix « std::string(num_blank, ' ') « std::flush;
00272
00273
                  os_->flags(flags);
00274
              }
00275
00282
              void print bar(std::stringstream &ss, double filled) const {
                  auto num_filled = static_cast<index>(std::round(filled * bar_size_));
static const char symbols[] = {'!', '@', '$', '%', '^', '&', '*', '+'
int num_symbols = sizeof(symbols) / sizeof(symbols[0]);
00283
00284
00285
00286
00287
                   ss « '[';
                   for (index i = 0; i < bar_size_; ++i) {</pre>
00288
                      if (i < num_filled) {</pre>
00289
00290
                           ss « '#';
                       } else if (i == num_filled && num_filled > 0) {
00291
00292
                          ss « symbols[symbol_index_ % num_symbols]; // Use symbol_index_ for frequency
00293
                       } else {
                           ss « '.';
00294
00295
00296
00297
                   ss « ']';
00298
00299
00305
               double time_since_refresh() const { return refresh_.peek(); }
00306
00310
              void reset refresh timer() { refresh .reset(); }
00311
00312
              Chronometer chronometer_{{}};
00313
              Chronometer refresh_{{}};
00314
              double min_time_per_update_{0.15}; // found experimentally
00315
00316
              std::ostream *os {&std::cerr};
00317
00318
              index bar_size_{40};
00319
              index term_cols_{1};
00320
              index symbol_index_{0};
00321
00322
              std::string prefix_{};
00323
              std::stringstream suffix {};
00324
00325
00326 // ----- iter_wrapper -----
00327
          template<class ForwardIter, class Parent>
00334
00335
          class iter_wrapper {
00336
          public:
00337
              using iterator_category = typename ForwardIter::iterator_category;
00338
              using value_type = typename ForwardIter::value_type;
00339
              using difference_type = typename ForwardIter::difference_type;
              using pointer = typename ForwardIter::pointer;
using reference = typename ForwardIter::reference;
00340
00341
```

```
00342
00343
              iter_wrapper(ForwardIter it, Parent *parent) : current_(it), parent_(parent) {}
00344
00345
              auto operator*() { return *current_; }
00346
00347
              void operator++() { ++current_; }
00348
00349
              template<class Other>
00350
              bool operator!=(const Other &other) const {
00351
                  parent_->update(); // here and not in ++ because I need to run update
                  // before first advancement!
return current_ != other;
00352
00353
00354
              }
00355
00356
              bool operator!=(const iter_wrapper &other) const {
00357
                  parent_->update(); // here and not in ++ because I need to run update
00358
                  //\ {\tt before\ first\ advancement!}
                  return current_ != other.current_;
00359
00360
00361
00362
              [[nodiscard]] const ForwardIter &get() const { return current_; }
00363
00364
         private:
              friend Parent:
00365
00366
              ForwardIter current_;
00367
              Parent *parent_;
00368
00369
00370 // ------ tqdm_for_lvalues -----
00371
00378
          template<class ForwardIter, class EndIter = ForwardIter>
00379
          class tqdm_for_lvalues {
00380
          public:
00381
              using this_t = tqdm_for_lvalues<ForwardIter, EndIter>;
              using iterator = iter_wrapper<ForwardIter, this_t>;
00382
00383
              using value_type = typename ForwardIter::value_type;
              using size_type = index;
00384
00385
              using difference_type = index;
00386
00387
              tqdm_for_lvalues(ForwardIter begin, EndIter end)
00388
                      : first_(begin, this), last_(end), num_iters_(std::distance(begin, end)) {}
00389
              tqdm_for_lvalues(ForwardIter begin, EndIter end, index total)
00390
00391
                      : first_(begin, this), last_(end), num_iters_(total) {}
00392
00393
              template<class Container>
00394
              explicit tqdm_for_lvalues(Container &C)
00395
                      : first_(C.begin(), this), last_(C.end()), num_iters_(C.size()) {}
00396
00397
              template<class Container>
00398
              explicit tqdm_for_lvalues(const Container &C)
00399
                      : first_(C.begin(), this), last_(C.end()), num_iters_(C.size()) {}
00400
00401
              tqdm_for_lvalues(const tqdm_for_lvalues &) = delete;
00402
00403
              tgdm for lvalues(tgdm for lvalues &&) = delete;
00404
00405
              tqdm_for_lvalues &operator=(tqdm_for_lvalues &&) = delete;
00406
00407
              tqdm_for_lvalues &operator=(const tqdm_for_lvalues &) = delete;
00408
00409
              ~tgdm for lvalues() = default;
00410
00411
              template<class Container>
00412
              tqdm_for_lvalues(Container &&) = delete; // prevent misuse!
00413
00414
              iterator begin() {
00415
                  bar_.restart();
00416
                  iters_done_ = 0;
00417
                  return first_;
00418
00419
00420
              EndIter end() const { return last_; }
00421
00422
              void update() {
00423
                  ++iters_done_;
00424
                  bar_.update(calc_progress());
00425
00426
00427
              void set ostream(std::ostream &os) { bar .set ostream(os); }
00428
00429
              void set_prefix(std::string s) { bar_.set_prefix(std::move(s)); }
00430
00431
              void set_bar_size(int size) { bar_.set_bar_size(size); }
00432
00433
              void set_min_update_time(double time) { bar_.set_min_update_time(time); }
00434
```

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```
00435
              template<class T>
00436
              tqdm_for_lvalues &operator (const T &t) {
00437
                 bar_ « t;
00438
                  return *this;
00439
00440
              void manually_set_progress(double to) {
   clamp(to, 0, 1);
00441
00442
00443
                  iters_done_ = std::round(to * num_iters_);
00444
00445
00446
         private:
00447
             double calc_progress() const {
00448
                 double denominator = num_iters_;
00449
                  if (num_iters_ == 0) denominator += 1e-9;
00450
                 return iters_done_ / denominator;
00451
00452
00453
              iterator first_;
00454
              EndIter last_;
00455
              index num_iters_{0};
00456
              index iters_done_{0};
00457
             progress_bar bar_;
00458
         };
00459
00460
          template<class Container>
00461
          tqdm_for_lvalues(Container &) -> tqdm_for_lvalues<typename Container::iterator>;
00462
00463
          template<class Container>
00464
          tqdm_for_lvalues(const Container &)
00465
          -> tgdm for lyalues<typename Container::const iterator>;
00466
00467 //
         ----- tqdm_for_rvalues ------
00468
00474
         template<class Container>
00475
         class tqdm_for_rvalues {
00476
         public:
00477
             using iterator = typename Container::iterator;
00478
              using const_iterator = typename Container::const_iterator;
00479
             using value_type = typename Container::value_type;
00480
00481
              explicit tqdm_for_rvalues(Container &&C)
00482
                      : C (std::forward<Container>(C)), tgdm (C) {}
00483
00484
              auto begin() { return tqdm_.begin(); }
00485
00486
              auto end() { return tqdm_.end(); }
00487
00488
              void update() { return tqdm_.update(); }
00489
00490
              void set_ostream(std::ostream &os) { tqdm_.set_ostream(os); }
00491
00492
              void set_prefix(std::string s) { tqdm_.set_prefix(std::move(s)); }
00493
00494
              void set_bar_size(int size) { tqdm_.set_bar_size(size); }
00495
00496
              void set_min_update_time(double time) { tqdm_.set_min_update_time(time); }
00497
00498
              template<class T>
00499
              auto &operator«(const T &t) {
00500
                 return tqdm_ « t;
00501
00502
00503
              void advance(index amount) { tqdm_.advance(amount); }
00504
00505
             void manually_set_progress(double to) { tqdm_.manually_set_progress(to); }
00506
00507
         private:
00508
              Container C_;
00509
              tqdm_for_lvalues<iterator> tqdm_;
00510
00511
00512
          template<class Container>
          tqdm_for_rvalues(Container &&) -> tqdm_for_rvalues<Container>;
00513
00514
00515 //
             ----- tqdm -----
00516
00525
          template<class ForwardIter>
00526
          auto tqdm(const ForwardIter &first, const ForwardIter &last) {
00527
             return tqdm_for_lvalues(first, last);
00528
00529
00539
          template<class ForwardIter>
00540
          auto tqdm(const ForwardIter &first, const ForwardIter &last, index total) {
00541
             return tqdm_for_lvalues(first, last, total);
00542
00543
```

```
template<class Container>
00552
         auto tqdm(const Container &C) {
00553
             return tqdm_for_lvalues(C);
00554
00555
00563
         template<class Container>
00564
         auto tqdm(Container &C) {
00565
             return tqdm_for_lvalues(C);
00566
00567
00575
         template<class Container>
00576
         auto tgdm (Container &&C) {
00577
             return tqdm_for_rvalues(std::forward<Container>(C));
00578
00579
00580 // ----- int_iterator ------
00581
00587
         template<class IntType>
00588
         class int_iterator {
00589
         public:
00590
             using iterator_category = std::random_access_iterator_tag;
00591
             using value_type = IntType;
             using difference_type = IntType;
00592
             using pointer = IntType *;
00593
00594
             using reference = IntType &;
00595
00596
             explicit int_iterator(IntType val) : value_(val) {}
00597
00598
             IntType &operator*() { return value_; }
00599
00600
             int iterator & operator ++ () {
00601
                 ++value_;
00602
                 return *this;
00603
00604
             int_iterator &operator--() {
00605
00606
                  --value ;
00607
                 return *this;
00608
00609
00610
             int_iterator &operator+=(difference_type d) {
00611
                 value_ += d;
00612
                 return *this;
00613
             }
00614
00615
             difference_type operator-(const int_iterator &other) const {
00616
                 return value_ - other.value_;
00617
             }
00618
00619
             bool operator!=(const int_iterator &other) const {
00620
                 return value_ != other.value_;
00621
00622
         private:
00623
             IntType value_;
00624
00625
00626
00627 //
             ------ range ------
00628
00634
         template<class IntType>
00635
         class range {
00636
         public:
00637
             using iterator = int_iterator<IntType>;
00638
             using const_iterator = iterator;
00639
             using value_type = IntType;
00640
00641
             range(IntType first, IntType last) : first_(first), last_(last) {}
00642
00643
             explicit range(IntType last) : first_(0), last_(last) {}
00644
00645
             [[nodiscard]] iterator begin() const { return first_; }
00646
00647
             [[nodiscard]] iterator end() const { return last_; }
00648
00649
             [[nodiscard]] index size() const { return last - first; }
00650
00651
         private:
00652
             iterator first_;
             iterator last_;
00653
00654
00655
00664
          template<class IntType>
00665
          auto trange(IntType first, IntType last) {
00666
             return tqdm(range(first, last));
00667
00668
00676
         template<class IntTvpe>
```

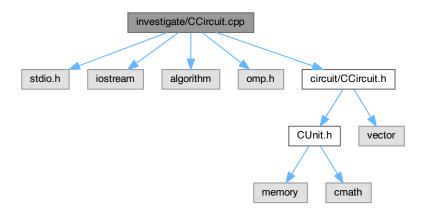
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```
00677
          auto trange(IntType last)
00678
             return tqdm(range(last));
00679
00680
00681 // ----- timing iterator -----
00682
00686
          class timing_iterator_end_sentinel {
00687
          public:
             explicit timing_iterator_end_sentinel(double num_seconds)
00688
00689
                      : num_seconds_(num_seconds) {}
00690
00691
              [[nodiscard]] double num seconds() const { return num seconds ; }
00692
00693
00694
             double num_seconds_;
00695
00696
00700
          class timing_iterator {
00701
          public:
00702
              using iterator_category = std::forward_iterator_tag;
00703
              using value_type = double;
00704
              using difference_type = double;
00705
              using pointer = double *;
00706
              using reference = double &;
00707
00708
              double operator*() const { return chrono_.peek(); }
00709
00710
              timing_iterator &operator++() { return *this; }
00711
00712
              bool operator!=(const timing_iterator_end_sentinel &other) const {
00713
                  return chrono_.peek() < other.num_seconds();</pre>
00714
00715
00716
          private:
00717
              tqdm::Chronometer chrono_;
00718
00719
00720 //
                       ----- timer -----
00721
00725
          struct timer {
00726
          public:
00727
              using iterator = timing_iterator;
              using end_iterator = timing_iterator_end_sentinel;
using const_iterator = iterator;
00728
00729
              using value_type = double;
00730
00731
00732
              explicit timer(double num_seconds) : num_seconds_(num_seconds) {}
00733
00734
              [[nodiscard]] static iterator begin() { return iterator(); }
00735
00736
              [[nodiscard]] end_iterator end() const {
00737
                  return end_iterator(num_seconds_);
00738
00739
00740
              [[nodiscard]] double num_seconds() const { return num_seconds_; }
00741
00742
         private:
00743
              double num_seconds_;
00744
00745
00749
          class tqdm_timer {
00750
          public:
00751
              using iterator = iter_wrapper<timing_iterator, tqdm_timer>;
00752
              using end_iterator = timer::end_iterator;
00753
              using value_type = typename timing_iterator::value_type;
              using size_type = index;
00754
00755
              using difference_type = index;
00756
00757
              explicit tgdm timer(double num seconds) : num seconds (num seconds) {}
00758
00759
              tqdm_timer(const tqdm_timer &) = delete;
00760
00761
              tqdm_timer(tqdm_timer &&) = delete;
00762
00763
              tgdm timer &operator=(tgdm timer &&) = delete;
00764
00765
              tqdm_timer &operator=(const tqdm_timer &) = delete;
00766
00767
              ~tqdm_timer() = default;
00768
00769
              template<class Container>
00770
              tqdm_timer(Container &&) = delete; // prevent misuse!
00771
00772
              iterator begin() {
00773
                  bar_.restart();
00774
                  return iterator(timing_iterator(), this);
00775
              }
```

```
00777
              end_iterator end() const { return end_iterator(num_seconds_); }
00778
              void update() {
   double t = bar_.elapsed_time();
00779
00780
00781
00782
                  bar_.update(t / num_seconds_);
00783
00784
00785
              void set_ostream(std::ostream &os) { bar_.set_ostream(os); }
00786
00787
              void set_prefix(std::string s) { bar_.set_prefix(std::move(s)); }
00788
00789
              void set_bar_size(int size) { bar_.set_bar_size(size); }
00790
00791
              void set_min_update_time(double time) { bar_.set_min_update_time(time); }
00792
00793
              template<class T>
              tqdm_timer &operator (const T &t) {
00795
                  bar_ « t;
00796
                  return *this;
00797
00798
00799
          private:
00800
              double num_seconds_;
00801
              progress_bar bar_;
00802
00803
00810
          inline auto tqdm(timer t) { return tqdm_timer(t.num_seconds()); }
00811
00812 } // namespace tqdm
```

# 13.39 investigate/CCircuit.cpp File Reference

```
#include <stdio.h>
#include <iostream>
#include <algorithm>
#include <omp.h>
#include "circuit/CCircuit.h"
Include dependency graph for CCircuit.cpp:
```



# Variables

- const double k\_G\_C = 0.004
- const double k\_G\_I = 0.001
- const double k w C = 0.0002
- const double  $k_w_l = 0.0003$
- const double V = 10.0

```
const double phi = 0.1
const double rho = 3000.0
13.39.1 Variable Documentation
13.39.1.1 k_G_C
const double k_G_C = 0.004
13.39.1.2 k_G_I
const double k_G_I = 0.001
```

13.39.1.3 k\_w\_C

const double  $k_w_C = 0.0002$ 

13.39.1.4 k\_w\_l

const double  $k_w_I = 0.0003$ 

13.39.1.5 phi

const double phi = 0.1

13.39.1.6 rho

const double rho = 3000.0

13.39.1.7 V

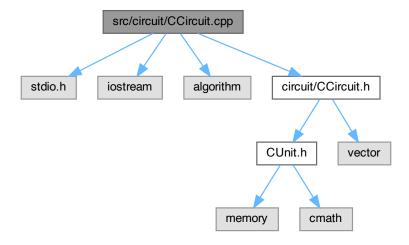
const double V = 10.0

# 13.40 src/circuit/CCircuit.cpp File Reference

Defines the Circuit class for modeling and simulating a circuit system.

```
#include <stdio.h>
#include <iostream>
#include <algorithm>
#include "circuit/CCircuit.h"
```

Include dependency graph for CCircuit.cpp:



#### **Variables**

- const double **k\_G\_C** = 0.004
- const double **k\_G\_I** = 0.001
- const double k\_w\_C = 0.0002
- const double k\_w\_l = 0.0003
- const double V = 10.0
- const double phi = 0.1
- const double rho = 3000.0

# 13.40.1 Detailed Description

Defines the Circuit class for modeling and simulating a circuit system.

The Circuit class provides methods to initialize, configure, and simulate a circuit with multiple units. It includes methods for checking validity, printing information, connecting units, initializing feed rates, checking convergence, and calculating flows. The Circuit class is designed to facilitate the modeling and simulation of circuits in various applications, including industrial processes and scientific research.

Date

Created on May 20, 2024

#### **Authors**

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- · Peifeng Tan
- · Yongwen Chen
- · Zihan Li

# 13.40.2 Variable Documentation

# 13.40.2.1 k\_G\_C

const double  $k_G_C = 0.004$ 

#### 13.40.2.2 k\_G\_I

const double  $k_G_I = 0.001$ Constants for Gerardium recovery

# 13.40.2.3 k\_w\_C

const double  $k_w_C = 0.0002$ 

# 13.40.2.4 k\_w\_l

const double  $k_w_I = 0.0003$ Constants for waste recovery

#### 13.40.2.5 phi

const double phi = 0.1

#### 13.40.2.6 rho

const double rho = 3000.0 Constants for the circuit

# 13.40.2.7 V

const double V = 10.0

# 13.41 post\_process/visualize.py File Reference

# Namespaces

· namespace visualize

# **Functions**

• visualize.parse\_list (string)

# Variables

- · visualize.parser
- · visualize.type
- visualize.args = parser.parse\_args()
- visualize.x = args.vector
- visualize.p = args.perf
- visualize.r = args.recovery
- visualize.g = args.grade
- visualize.graph = graphviz.Digraph()
- · visualize.rankdir
- · visualize.nodesep
- · visualize.ranksep
- · visualize.splines
- visualize.overlap
- visualize.dpi

- str visualize.feed\_node = 'Feed'
- str visualize.concentrate = 'Concentrate'
- str visualize.tailings = 'Tailings'
- · visualize.shape
- visualize.color = 1 else 'red'
- tuple visualize.num\_nodes = (len(x) 1) // 3
- str visualize.from\_node = f'Unit {i // 3}'
- tuple visualize.to\_node = (num\_nodes + 1) else f'Unit {x[i + 1]}'
- visualize.cleanup
- · visualize.True
- · visualize.format
- · visualize.view
- · dict visualize.legend\_labels
- · list visualize.legend\_handles
- list visualize.columns = ['Feed'] + [f'Unit {i}' for i in range(0, num\_nodes)]
- · list visualize.reshaped list
- visualize.df = pd.DataFrame(reshaped\_list).transpose()
- · visualize.figsize
- · visualize.handles
- · visualize.labels
- · visualize.loc
- visualize.bbox\_to\_anchor
- · visualize.fontsize
- · visualize.cellText
- · visualize.values
- · visualize.colLabels
- · visualize.cellLoc
- visualize.bbox
- visualize.ha
- visualize.va
- · visualize.transform
- · visualize.bbox inches

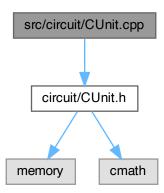
# 13.42 README.md File Reference

# 13.43 REFERENCE.md File Reference

# 13.44 src/circuit/CUnit.cpp File Reference

Defines the CUnit class for representing a unit within a circuit system.

#include "circuit/CUnit.h"
Include dependency graph for CUnit.cpp:



# 13.44.1 Detailed Description

Defines the CUnit class for representing a unit within a circuit system.

The CUnit class provides methods to manage and simulate individual units within a circuit. It includes methods for setting and getting various properties of the unit, such as feed rates, mark status, and connection pointers. This class is designed to facilitate the modeling and simulation of circuits in various applications, including industrial processes and scientific research.

Date

Created on May 20, 2024

# **Authors**

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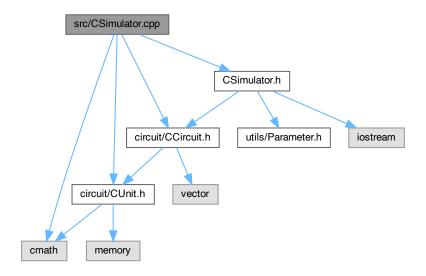
- · Alex N Njeumi
- · Geyu Ji
- Melissa Y S Sim
- Mingsheng Cai
- Peifeng Tan
- · Yongwen Chen
- · Zihan Li

# 13.45 src/CSimulator.cpp File Reference

Defines the functions and structures for simulating and evaluating circuits in a circuit modeling framework.

```
#include "circuit/CUnit.h"
#include "circuit/CCircuit.h"
#include "CSimulator.h"
#include <cmath>
```

Include dependency graph for CSimulator.cpp:



#### **Functions**

double performance (double concentrateG, double concentrateW)

Computes the performance metric of a circuit.

• double grade (double concentrateG, double concentrateW)

Computes the grade of a circuit.

• double recovery (double concentrateG, double gerardiumFeed)

Computes the recovery of a circuit.

• double Evaluate\_Circuit (int vectorSize, int \*circuitVector, struct CircuitParameters parameters)

Evaluates the performance of a circuit.

• double Evaluate\_Circuit (int vectorSize, int \*circuitVector, double &Recovery, double &Grade, struct CircuitParameters parameters)

Evaluates the performance, recovery, and grade of a circuit.

# 13.45.1 Detailed Description

Defines the functions and structures for simulating and evaluating circuits in a circuit modeling framework.

The CSimulator module provides functions for evaluating the performance, grade, and recovery of circuits in a circuit modeling framework. It includes definitions for circuit parameters and multiple overloads for evaluating circuits with or without additional parameters.

Date

Created on May 20, 2024

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# 13.45.2 Function Documentation

# 13.45.2.1 Evaluate\_Circuit() [1/2]

```
double Evaluate_Circuit (
                int vectorSize,
                int * circuitVector,
                double & Recovery,
                 double & Grade,
                 struct CircuitParameters parameters )
```

Evaluates the performance, recovery, and grade of a circuit.

This function evaluates the performance, recovery, and grade of a circuit based on the given circuit vector and additional circuit parameters.

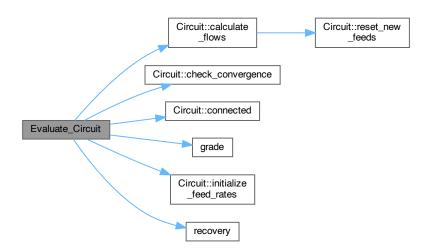
#### **Parameters**

vectorSize	The size of the circuit vector.
circuitVector	Pointer to the array representing the circuit configuration.
Recovery	Reference to a double variable to store the recovery value.
Grade	Reference to a double variable to store the grade value.
parameters	Additional parameters for the circuit simulation.

# Returns

The performance metric of the circuit.

Here is the call graph for this function:



# 13.45.2.2 Evaluate\_Circuit() [2/2]

```
int * circuitVector,
struct CircuitParameters parameters )
```

Evaluates the performance of a circuit.

This function evaluates the performance of a circuit based on the given circuit vector and additional circuit parameters.

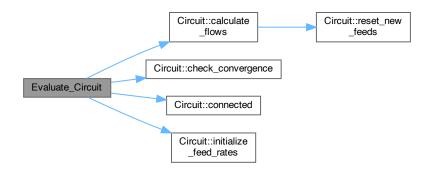
#### **Parameters**

vectorSize	The size of the circuit vector.
circuitVector	Pointer to the array representing the circuit configuration.
parameters	Additional parameters for the circuit simulation.

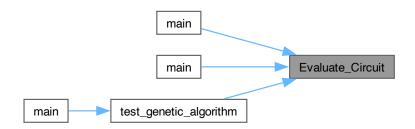
# Returns

The performance metric of the circuit.

Here is the call graph for this function:



Here is the caller graph for this function:



# 13.45.2.3 grade()

Computes the grade of a circuit.

This function computes the grade of a circuit based on the given concentrate gerardium and waste values.

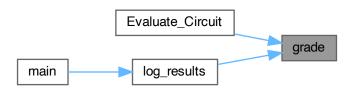
#### **Parameters**

concentrateG	The amount of gerardium in the concentrate.
concentrateW	The amount of waste in the concentrate.

#### Returns

The grade of the circuit.

Here is the caller graph for this function:



# 13.45.2.4 performance()

Computes the performance metric of a circuit.

This function computes the performance metric of a circuit based on the given concentrate gerardium and waste values.

#### **Parameters**

concentrateG	The amount of gerardium in the concentrate.
concentrateW	The amount of waste in the concentrate.

#### Returns

The performance metric of the circuit.

Here is the caller graph for this function:



# 13.45.2.5 recovery()

```
double recovery (
```

```
double concentrateG,
double gerardiumFeed )
```

Computes the recovery of a circuit.

This function computes the recovery of a circuit based on the given concentrate gerardium and gerardium feed values.

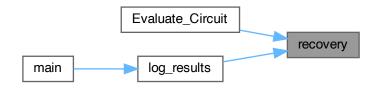
#### **Parameters**

concentrateG	The amount of gerardium in the concentrate.
gerardiumFeed	The amount of gerardium fed into the circuit.

#### Returns

The recovery of the circuit.

Here is the caller graph for this function:



# 13.46 src/Genetic\_Algorithm.cpp File Reference

Defines the functions and structures for implementing a genetic algorithm.

```
#include "Genetic_Algorithm.h"
#include <iostream>
#include <algorithm>
#include <ctime>
#include <climits>
#include <unordered_set>
#include <random>
#include <omp.h>
```

Include dependency graph for Genetic\_Algorithm.cpp:



#### **Macros**

#define INVALID\_FITNESS -1000000000.0

#### **Functions**

void initializeRandomSeed (const Algorithm\_Parameters &params, std::vector< std::mt19937 > &generators)

Initializes the random seed based on algorithm parameters.

• std::vector< Individual > initializePopulation (const Algorithm\_Parameters &params, int vectorSize, bool(&validity)(int, int \*), std::vector< std::mt19937 > &generators)

Initializes the population for the genetic algorithm.

 void elitism (const std::vector< Individual > &population, std::vector< Individual > &newPopulation, const Algorithm Parameters &params)

Apply elitism to preserve the best individuals in a new population.

double acceptanceProbability (double currentEnergy, double newEnergy, double temperature)

Calculate the acceptance probability for a new state in Simulated Annealing.

bool applySimulatedAnnealing (Individual &offspring, Individual &parent1, Individual &parent2, int vector
 \_size, double(&func)(int, int \*, struct CircuitParameters), bool(&validity)(int, int \*), double &Temp, const
 Algorithm Parameters &params, const CircuitParameters c params, std::mt19937 &generator)

Applies simulated annealing technique to decide whether to accept an offspring in a genetic algorithm.

• int optimize (int vector\_size, int \*vector, double(&func)(int, int \*, struct CircuitParameters), bool(&validity)(int, int \*), Algorithm\_Parameters params, CircuitParameters c\_params)

Optimizes a solution using the genetic algorithm.

# 13.46.1 Detailed Description

Defines the functions and structures for implementing a genetic algorithm.

The Genetic\_Algorithm module provides functions for initializing populations, setting random seeds, and optimizing solutions using a genetic algorithm framework. This includes selection, crossover, and mutation operators.

Date

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# 13.46.2 Macro Definition Documentation

# 13.46.2.1 INVALID\_FITNESS

#define INVALID\_FITNESS -1000000000.0

#### 13.46.3 Function Documentation

#### 13.46.3.1 acceptanceProbability()

Calculate the acceptance probability for a new state in Simulated Annealing.

This function calculates the probability with which a new state should be accepted over the current state, based on their respective energies and the current temperature of the system. If the new energy is better (higher value), the function returns 1.0, meaning the new state is always accepted. Otherwise, it returns a value calculated using the Boltzmann probability distribution which considers both the difference in energy and the current temperature.

#### **Parameters**

currentEn	ergy	The energy of the current state.
newEnerg	gy	The energy of the new state being considered.
temperatu	ıre	The current temperature of the system (controls the probability threshold).

#### Returns

double The probability of accepting the new state.

Here is the caller graph for this function:



# 13.46.3.2 applySimulatedAnnealing()

Applies simulated annealing technique to decide whether to accept an offspring in a genetic algorithm.

This function performs the simulated annealing decision process on a newly generated offspring. It compares the offspring's fitness to the average fitness of its parents using the simulated annealing acceptance probability, which depends on the current temperature of the system. The function also handles the cooling of the system temperature and validates the offspring using a provided validity function.

# **Parameters**

offspring	A reference to the offspring individual whose acceptance is being determined.
parent1	A reference to the first parent of the offspring.
parent2	A reference to the second parent of the offspring.
vector_size	The size of the individual's gene vector.
func	A function that calculates the fitness of an individual based on its genes.
validity	A function that checks the validity of an individual's gene configuration.
Temp	A reference to the current temperature used for the simulated annealing process.
params	A structure containing parameters relevant to the annealing process, such as the temperature decrement.
c_params	The circuit parameters used in the fitness evaluation function.
generator	A random number generator used for probabilistic decisions.

#### Returns

bool Returns true if the offspring is accepted, false otherwise. The function can also modify the offspring directly, setting it to one of the parents if the temperature is low and the acceptance criteria are not met.

#### Note

The function directly modifies the temperature, decrementing it according to <code>deltT</code> from <code>params</code>. If the offspring is invalid (as determined by the <code>validity</code> function), it is immediately rejected and assigned a fitness value of <code>INVALID\_FITNESS</code>.

Here is the call graph for this function:



Here is the caller graph for this function:



#### 13.46.3.3 elitism()

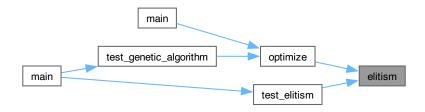
Apply elitism to preserve the best individuals in a new population.

This function sorts the given population in descending order based on their fitness, then selectively copies the top percentage of individuals as specified by the elitePercentage in the Algorithm\_Parameters. This is used to ensure that the best individuals are preserved for the next generation, promoting genetic diversity and preventing loss of the best found solutions.

### **Parameters**

population	The current generation of individuals (const reference).
newPopulation	The next generation of individuals where elites will be added (reference).
params	The parameters of the algorithm including population size and elite percentage.

Here is the caller graph for this function:



# 13.46.3.4 initializePopulation()

Initializes the population for the genetic algorithm.

This function initializes the population for the genetic algorithm, ensuring each individual meets the validity criteria specified by the user.

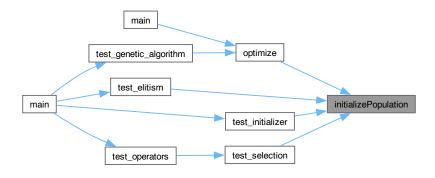
#### **Parameters**

params	The algorithm parameters containing population size and other configurations.
vectorSize	The size of the gene vector for each individual.
validity	A function pointer to the validity checking function.
generators	A vector of random number generators for each thread.

# Returns

A vector of individuals representing the initial population.

Here is the caller graph for this function:



#### 13.46.3.5 initializeRandomSeed()

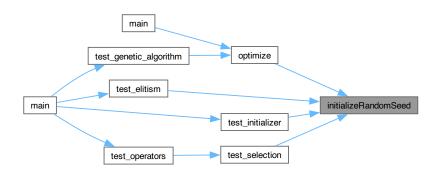
Initializes the random seed based on algorithm parameters.

This function initializes the random seed using the specified seed in the algorithm parameters. If the seed is zero, it uses the current time as the seed.

#### **Parameters**

params	The algorithm parameters containing the random seed.
generators	A vector of random number generators for each thread.

Here is the caller graph for this function:



#### 13.46.3.6 optimize()

```
int optimize (
    int vector_size,
    int * vector,
    double(&)(int, int *, struct CircuitParameters) func,
    bool(&)(int, int *) validity,
    struct Algorithm_Parameters params,
    struct CircuitParameters c_params)
```

Optimizes a solution using the genetic algorithm.

This function performs optimization using a genetic algorithm. It initializes the population, assesses the fitness of each individual, and iterates through generations applying selection, crossover, and mutation operators.

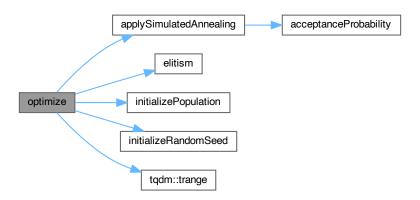
# **Parameters**

vector_size	The size of the gene vector.
vector	Pointer to the array representing the solution vector.
func	A function pointer to the fitness evaluation function.
validity	A function pointer to the validity checking function.
parameters	The algorithm parameters containing various configurations.

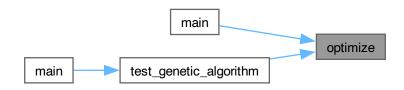
#### Returns

An integer indicating the success (0) or failure (-1) of the optimization.

Here is the call graph for this function:



Here is the caller graph for this function:



# 13.47 src/main.cpp File Reference

Main entry point for the circuit optimization program using genetic algorithms.

```
#include <iostream>
#include <chrono>
#include "circuit/CCircuit.h"
#include "CSimulator.h"
#include "Genetic_Algorithm.h"
#include "utils/Helper.h"
Include dependency graph for main.cpp:
```



#### **Functions**

• int main (int argc, char \*argv[])

# 13.47.1 Detailed Description

Main entry point for the circuit optimization program using genetic algorithms.

Test file for evaluating the functionalities of the Genetic Algorithm module.

Test file for evaluating execution times of the Evaluate Circuit function with different circuit vector sizes.

Test file for evaluating circuit configurations using the Evaluate\_Circuit function.

This program uses a genetic algorithm to optimize a circuit configuration. It sets up the initial circuit parameters and the genetic algorithm parameters, runs the optimization process, and then evaluates the optimized circuit. The results, including performance, recovery, and grade, are printed to the console.

The main steps include:

- · Setting up circuit parameters.
- · Defining the initial circuit vector.
- · Configuring the genetic algorithm parameters.
- · Running the genetic algorithm optimization.
- · Evaluating the optimized circuit and printing the results.

Date

Created on May 20, 2024

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- · Zihan Li

This file contains a series of test cases to evaluate different circuit configurations using the Evaluate\_Circuit function. Each test case checks the performance, recovery, and grade of a given circuit vector against expected values. The results are printed to the console, and the test passes or fails based on the accuracy of the output.

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This file generates circuit vectors of various sizes, evaluates them using the Evaluate\_Circuit function, and records the execution times. The results are printed to the console and also saved to an output file.

Date

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This file contains various test cases to evaluate the correctness and performance of the Genetic Algorithm implementation, including tests for the initializer, crossover, mutation, selection, and simulated annealing.

Date

Created on May 20, 2024

# **Authors**

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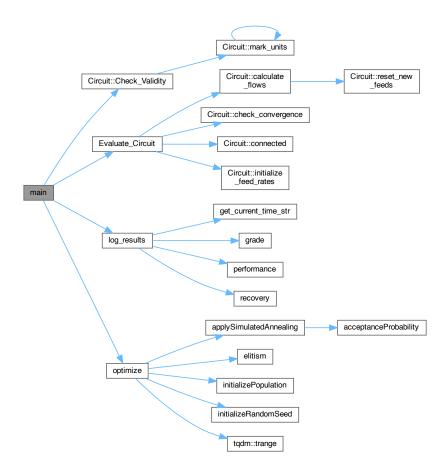
- · Alex N Njeumi
- Geyu Ji
- Melissa Y S Sim
- Mingsheng Cai
- · Peifeng Tan
- · Yongwen Chen
- Zihan Li

# 13.47.2 Function Documentation

# 13.47.2.1 main()

```
int main (
                int argc,
                 char * argv[] )
```

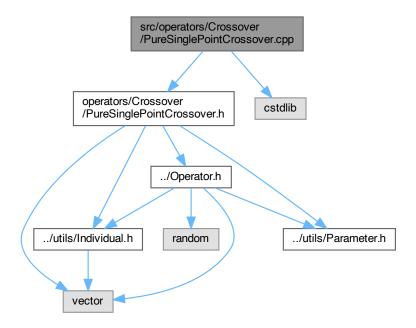
Here is the call graph for this function:



# 13.48 src/operators/Crossover/PureSinglePointCrossover.cpp File Reference

Defines the PureSinglePointCrossover class for performing single-point crossover in a genetic algorithm. #include "operators/Crossover/PureSinglePointCrossover.h" #include <cstdlib>

Include dependency graph for PureSinglePointCrossover.cpp:



# 13.48.1 Detailed Description

Defines the PureSinglePointCrossover class for performing single-point crossover in a genetic algorithm. The PureSinglePointCrossover class provides methods to perform a single-point crossover on two parent individuals to generate an offspring individual. This class is designed to be used in genetic algorithms to combine genetic material from two parents to create new individuals, which can then be used in subsequent generations of the algorithm.

Date

Created on May 20, 2024

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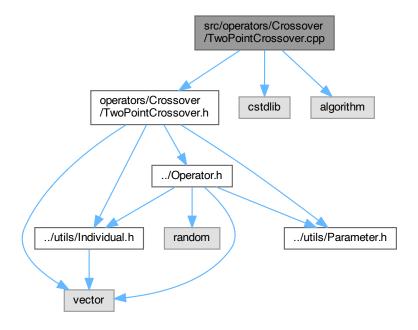
# 13.49 src/operators/Crossover/TwoPointCrossover.cpp File Reference

 $\label{the:constraint} \mbox{Defines the $\mbox{TwoPointCrossover}$ class for performing two-point crossover in a genetic algorithm.}$ 

```
#include "operators/Crossover/TwoPointCrossover.h"
#include <cstdlib>
```

#include <algorithm>

Include dependency graph for TwoPointCrossover.cpp:



# 13.49.1 Detailed Description

Defines the TwoPointCrossover class for performing two-point crossover in a genetic algorithm.

The TwoPointCrossover class provides methods to perform a two-point crossover on two parent individuals to generate an offspring individual. This class is designed to be used in genetic algorithms to combine genetic material from two parents to create new individuals, which can then be used in subsequent generations of the algorithm.

Date

Created on May 20, 2024

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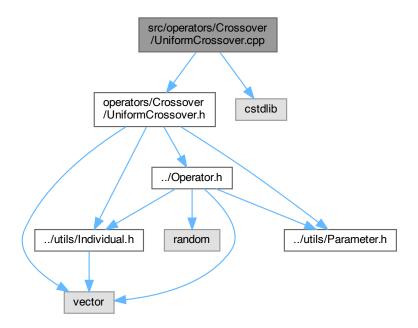
- · Alex N Njeumi
- · Geyu Ji
- · Melissa Y S Sim
- · Mingsheng Cai
- · Peifeng Tan
- Yongwen Chen
- · Zihan Li

# 13.50 src/operators/Crossover/UniformCrossover.cpp File Reference

Defines the UniformCrossover class for performing uniform crossover in a genetic algorithm.

#include "operators/Crossover/UniformCrossover.h"
#include <cstdlib>

Include dependency graph for UniformCrossover.cpp:



# 13.50.1 Detailed Description

Defines the UniformCrossover class for performing uniform crossover in a genetic algorithm.

The UniformCrossover class provides methods to perform a uniform crossover on two parent individuals to generate an offspring individual. This class is designed to be used in genetic algorithms to combine genetic material from two parents to create new individuals, which can then be used in subsequent generations of the algorithm.

Date

Created on May 20, 2024

**Authors** 

ACS Gerardium Rush - Pentlandite:

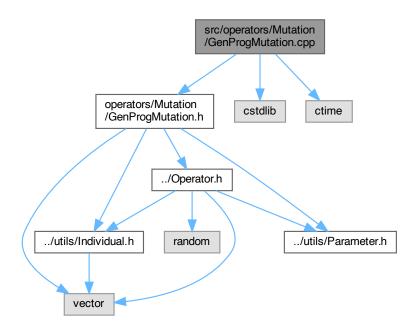
- · Alex N Njeumi
- Geyu Ji
- · Melissa Y S Sim
- · Mingsheng Cai
- · Peifeng Tan
- · Yongwen Chen
- Zihan Li

# 13.51 src/operators/Mutation/GenProgMutation.cpp File Reference

Defines the GenProgMutation class for performing mutation operations in a genetic programming algorithm.

```
#include "operators/Mutation/GenProgMutation.h"
#include <cstdlib>
#include <ctime>
```

Include dependency graph for GenProgMutation.cpp:



# 13.51.1 Detailed Description

Defines the GenProgMutation class for performing mutation operations in a genetic programming algorithm. The GenProgMutation class provides methods to perform mutation operations on individuals in a genetic programming algorithm. This class is designed to introduce variations into the population by modifying individuals' genetic material, which can then be used in subsequent generations of the algorithm.

Date

Created on May 20, 2024

#### Authors

ACS Gerardium Rush - Pentlandite:

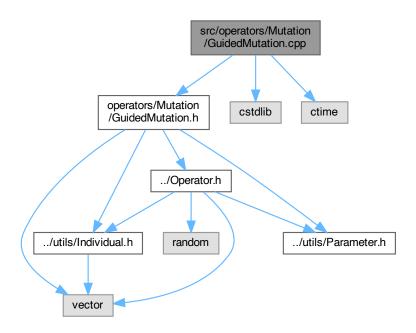
- · Alex N Njeumi
- · Geyu Ji
- Melissa Y S Sim
- · Mingsheng Cai
- Peifeng Tan
- · Yongwen Chen
- · Zihan Li

# 13.52 src/operators/Mutation/GuidedMutation.cpp File Reference

Defines the GuidedMutation class for performing guided mutation operations in a genetic algorithm.

```
#include "operators/Mutation/GuidedMutation.h"
#include <cstdlib>
#include <ctime>
```

Include dependency graph for GuidedMutation.cpp:



# 13.52.1 Detailed Description

Defines the GuidedMutation class for performing guided mutation operations in a genetic algorithm.

The GuidedMutation class provides methods to perform guided mutation operations on individuals in a genetic algorithm. This class is designed to introduce targeted variations into the population by modifying individuals' genetic material based on specific rules or heuristics, which can then be used in subsequent generations of the algorithm.

Date

Created on May 20, 2024

#### **Authors**

ACS Gerardium Rush - Pentlandite:

- · Alex N Njeumi
- Geyu Ji
- · Melissa Y S Sim
- Mingsheng Cai
- · Peifeng Tan
- · Yongwen Chen
- · Zihan Li

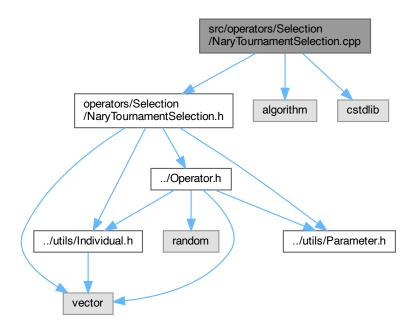
# 13.53 src/operators/Selection/NaryTournamentSelection.cpp File Reference

Defines the NaryTournamentSelection class for performing n-ary tournament selection in a genetic algorithm.

#include "operators/Selection/NaryTournamentSelection.h"
#include <algorithm>

#include <algorithm; #include <cstdlib>

Include dependency graph for NaryTournamentSelection.cpp:



## 13.53.1 Detailed Description

Defines the NaryTournamentSelection class for performing n-ary tournament selection in a genetic algorithm. The NaryTournamentSelection class provides methods to perform n-ary tournament selection on a population of individuals to select individuals for the next generation. This class is designed to be used in genetic algorithms to select the fittest individuals based on tournament competition among a subset of the population.

Date

Created on May 20, 2024

#### **Authors**

ACS Gerardium Rush - Pentlandite:

- Alex N Njeumi
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- · Melissa Y S Sim
- · Mingsheng Cai
- · Peifeng Tan
- · Yongwen Chen
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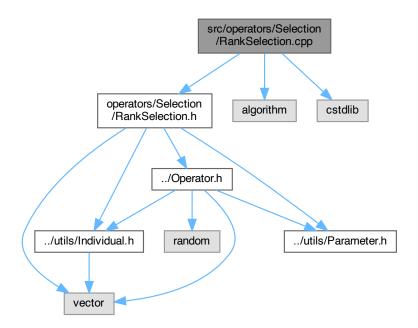
# 13.54 src/operators/Selection/RankSelection.cpp File Reference

Defines the RankSelection class for performing rank-based selection in a genetic algorithm.

#include "operators/Selection/RankSelection.h"

#include <algorithm>
#include <cstdlib>

Include dependency graph for RankSelection.cpp:



## 13.54.1 Detailed Description

Defines the RankSelection class for performing rank-based selection in a genetic algorithm.

The RankSelection class provides methods to perform rank-based selection on a population of individuals to select individuals for the next generation. This class is designed to be used in genetic algorithms to select the fittest individuals based on their rank within the population.

Date

Created on May 21, 2024

**Authors** 

ACS Gerardium Rush - Pentlandite:

- · Alex N Njeumi
- Geyu Ji
- · Melissa Y S Sim
- · Mingsheng Cai
- · Peifeng Tan
- · Yongwen Chen
- · Zihan Li

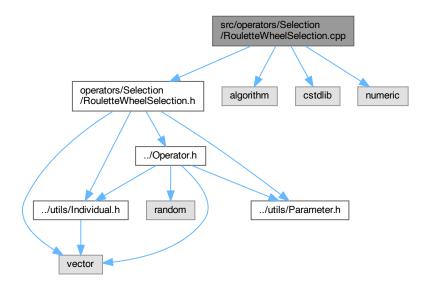
# 13.55 src/operators/Selection/RouletteWheelSelection.cpp File Reference

Defines the RouletteWheelSelection class for performing roulette wheel selection in a genetic algorithm.

#include "operators/Selection/RouletteWheelSelection.h"
#include <algorithm>
#include <cstdlib>

Include dependency graph for RouletteWheelSelection.cpp:

#include <numeric>



## 13.55.1 Detailed Description

Defines the RouletteWheelSelection class for performing roulette wheel selection in a genetic algorithm. The RouletteWheelSelection class provides methods to perform roulette wheel selection on a population of individuals to select individuals for the next generation. This class is designed to be used in genetic algorithms to select individuals based on their fitness proportionally.

Date

Created on May 21, 2024

#### **Authors**

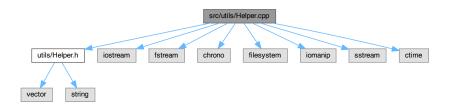
ACS Gerardium Rush - Pentlandite:

- · Alex N Njeumi
- Geyu Ji
- · Melissa Y S Sim
- · Mingsheng Cai
- · Peifeng Tan
- · Yongwen Chen
- Zihan Li

# 13.56 src/utils/Helper.cpp File Reference

```
#include "utils/Helper.h"
#include <iostream>
#include <fstream>
#include <chrono>
#include <filesystem>
#include <iomanip>
#include <sstream>
#include <ctime>
```

Include dependency graph for Helper.cpp:



#### **Functions**

• std::string get\_current\_time\_str ()

Retrieves the current time as a formatted string.

void log\_results (const std::vector< int > &vector, double elapsed\_time, double performance, double recovery, double grade)

Logs the results of the genetic algorithm optimization to a file.

## 13.56.1 Function Documentation

## 13.56.1.1 get\_current\_time\_str()

```
std::string get_current_time_str ()
```

Retrieves the current time as a formatted string.

This function returns the current time formatted as "YYYY-MM-DD\_HH-MM-SS".

Returns

A string representing the current time.

Here is the caller graph for this function:



## 13.56.1.2 log\_results()

```
double elapsed_time,
double performance,
double recovery,
double grade )
```

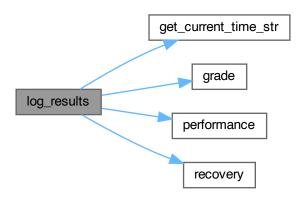
Logs the results of the genetic algorithm optimization to a file.

This function creates a new log folder with the current timestamp, and saves the performance metrics, recovery, grade, and the final configuration of the optimized circuit to a log file.

#### **Parameters**

vector	The final configuration vector of the optimized circuit.
elapsed_time	The time taken to complete the optimization process, in seconds.
performance	The performance metric of the optimized circuit.
recovery	The recovery metric of the optimized circuit.
grade	The grade metric of the optimized circuit.

Here is the call graph for this function:



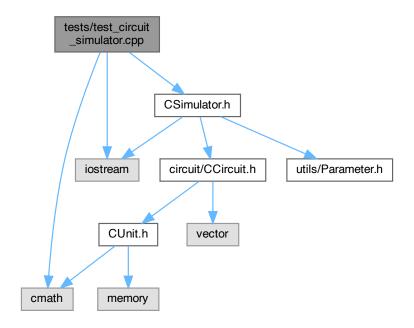
Here is the caller graph for this function:



# 13.57 tests/test\_circuit\_simulator.cpp File Reference

```
#include <cmath>
#include <iostream>
#include "CSimulator.h"
```

Include dependency graph for test\_circuit\_simulator.cpp:



## **Functions**

int main (int argc, char \*argv[])

Main function to run the test cases for evaluating circuit configurations.

## 13.57.1 Function Documentation

## 13.57.1.1 main()

```
int main (
          int argc,
          char * argv[] )
```

Main function to run the test cases for evaluating circuit configurations.

This function initializes several circuit vectors and uses the Evaluate\_Circuit function to check their performance, recovery, and grade. The results are compared against expected values, and the function prints whether each test passes or fails.

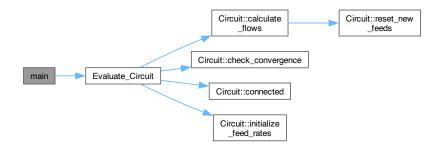
#### **Parameters**

argc	Number of command-line arguments.
argv	Array of command-line arguments.

#### Returns

int Returns 0 if all tests pass, otherwise returns 1.

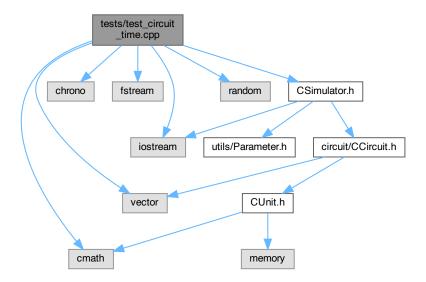
Here is the call graph for this function:



## 13.58 tests/test\_circuit\_time.cpp File Reference

```
#include <cmath>
#include <iostream>
#include <chrono>
#include <fstream>
#include <vector>
#include <random>
#include "CSimulator.h"
```

Include dependency graph for test\_circuit\_time.cpp:



## **Functions**

std::vector < int > generate\_circuit\_vector (int n)
 Generates a circuit vector of a given size.

#### • int main ()

Main function to evaluate execution times of the Evaluate\_Circuit function with different circuit vector sizes.

#### 13.58.1 Function Documentation

#### 13.58.1.1 generate\_circuit\_vector()

Generates a circuit vector of a given size.

This function generates a random circuit vector with the specified number of units. The first element is set to a valid feed unit, and the remaining elements are set to random unit indices.

#### **Parameters**

*n* The number of units in the circuit.

#### Returns

std::vector<int> The generated circuit vector.

Here is the caller graph for this function:



## 13.58.1.2 main()

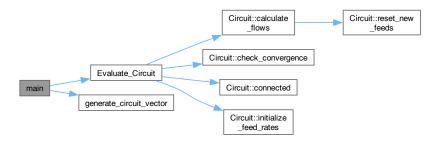
int main ( )

Main function to evaluate execution times of the Evaluate\_Circuit function with different circuit vector sizes. This function generates circuit vectors of various sizes, evaluates them using the Evaluate\_Circuit function, and records the execution times. The results are printed to the console and also saved to an output file.

## Returns

int Returns 0 if the program runs successfully, otherwise returns 1.

Here is the call graph for this function:



## 13.59 tests/test genetic algorithm.cpp File Reference

```
#include <iostream>
#include <cassert>
#include "Genetic_Algorithm.h"
#include "circuit/CCircuit.h"
#include "CSimulator.h"
```

Include dependency graph for test genetic algorithm.cpp:



#### **Functions**

• double test\_function (int vector\_size, int \*vector, CircuitParameters pams)

Test function for the genetic algorithm.

double test\_function\_long (int vector\_size, int \*vector, CircuitParameters pams)

Test function for the genetic algorithm with a long vector.

bool Check\_Validity (int vector\_size, int \*circuit\_vector)

Function to check the validity of a circuit vector.

bool areArraysEqual (const int \*a, const int \*b, size\_t size)

Function to check if two arrays are equal.

• void test genetic algorithm ()

Test the genetic algorithm for various cases.

void test\_initializer ()

Test the initialization of the population.

• void test crossover ()

Test the crossover operators.

void test mutation ()

Test the mutation operators.

void test selection ()

Test the selection operators.

• void test\_operators ()

Test the elitism operator.

· void test\_elitism ()

Test the elitism operator.

void test\_SA ()

Test the simulated annealing operator.

• int main ()

Main function to run all the tests.

## Variables

- int test\_answer [10] = {2, 1, 1, 2, 0, 2, 3, 0, 4, 4}
- int test answer1 [31]

## 13.59.1 Function Documentation

## 13.59.1.1 areArraysEqual()

Function to check if two arrays are equal.

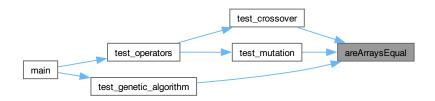
This function checks if two arrays are equal by comparing each element in the arrays.

#### **Parameters**

а	Check if this array is equal to b
b	Check if this array is equal to a
size	Size of the arrays

#### Returns

Here is the caller graph for this function:



## 13.59.1.2 Check\_Validity()

Function to check the validity of a circuit vector.

This function checks the validity of a circuit vector by ensuring that the feed ID and destination ID are within the range of the number of units in the circuit.

#### **Parameters**

vector_size	Size of the vector
circuit_vector	The circuit vector to be checked

Returns

Here is the caller graph for this function:



## 13.59.1.3 main()

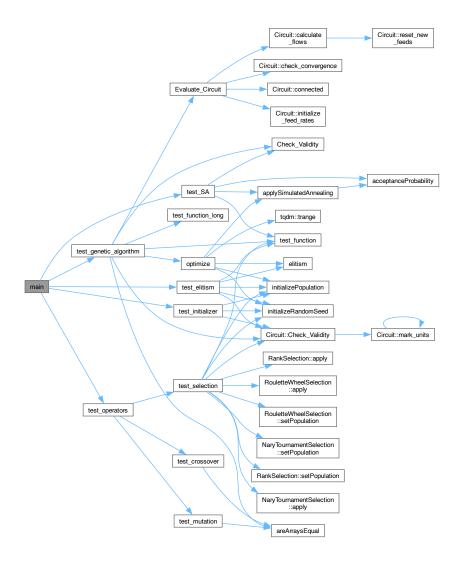
int main ( )

Main function to run all the tests.

This function runs all the tests for the Genetic Algorithm module.

Returns

Here is the call graph for this function:



## 13.59.1.4 test\_crossover()

void test\_crossover ( )

Test the crossover operators.

This function tests various crossover operators including two-point, pure single-point, and uniform crossover. Here is the call graph for this function:



Here is the caller graph for this function:

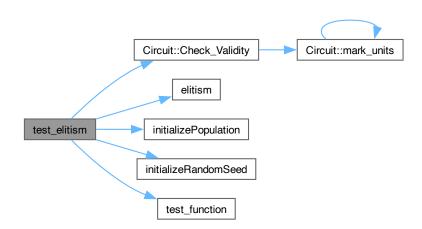


## 13.59.1.5 test\_elitism()

```
void test_elitism ( )
```

Test the elitism operator.

This function tests the elitism operator by checking if the performance of the population is improved after elitism. Here is the call graph for this function:



Here is the caller graph for this function:



## 13.59.1.6 test\_function()

Test function for the genetic algorithm.

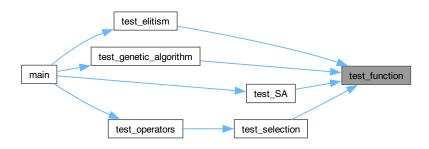
This function is used as the performance function for the genetic algorithm. It calculates the difference between the given vector and the test answer vector, and returns the negative of the sum of the squared differences.

#### **Parameters**

vector_size	Size of the vector
vector	The vector to be evaluated
pams	Circuit parameters

#### Returns

Here is the caller graph for this function:



## 13.59.1.7 test\_function\_long()

Test function for the genetic algorithm with a long vector.

This function is used as the performance function for the genetic algorithm. It calculates the difference between the given vector and the test answer vector, and returns the negative of the sum of the squared differences.

#### **Parameters**

vector_size	Size of the vector
vector	The vector to be evaluated
pams	Circuit parameters

Returns

Here is the caller graph for this function:

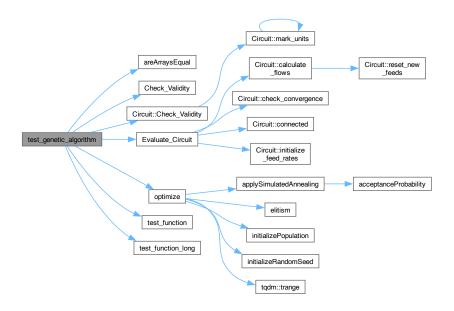


## 13.59.1.8 test\_genetic\_algorithm()

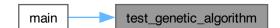
void test\_genetic\_algorithm ( )

Test the genetic algorithm for various cases.

This function sets up the parameters and runs the genetic algorithm on different test cases. It checks the optimized results against expected values. Here is the call graph for this function:



Here is the caller graph for this function:

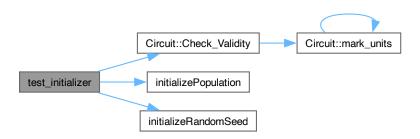


## 13.59.1.9 test\_initializer()

```
void test_initializer ( )
```

Test the initialization of the population.

This function tests if the population is correctly initialized with valid individuals and the correct size. Here is the call graph for this function:



Here is the caller graph for this function:



## 13.59.1.10 test\_mutation()

void test\_mutation ( )

Test the mutation operators.

This function tests various mutation operators including guided mutation and genprog mutation. Here is the call graph for this function:



Here is the caller graph for this function:

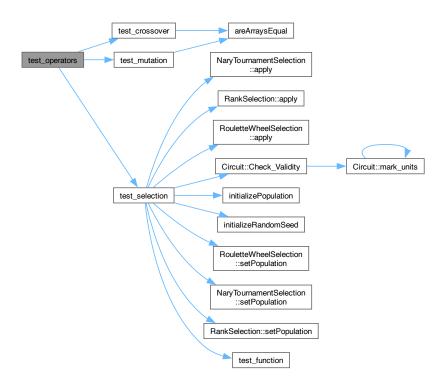


## 13.59.1.11 test\_operators()

```
void test_operators ( )
```

Test the elitism operator.

This function tests the elitism operator by checking if the performance of the population is improved after elitism. Here is the call graph for this function:



Here is the caller graph for this function:

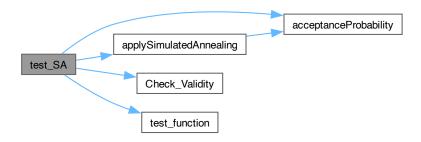


## 13.59.1.12 test\_SA()

```
void test_SA ( )
```

Test the simulated annealing operator.

This function tests the simulated annealing operator by checking if the offspring is accepted when it is better, and if the temperature decreases correctly. Here is the call graph for this function:



Here is the caller graph for this function:



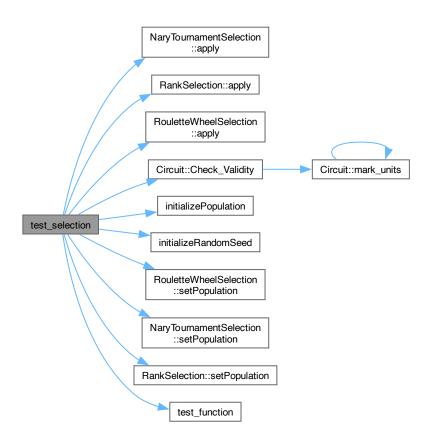
## 13.59.1.13 test\_selection()

void test\_selection ( )

Test the selection operators.

This function tests various selection operators including n-ary tournament, rank, and roulette wheel selection. Here

is the call graph for this function:



Here is the caller graph for this function:



## 13.59.2 Variable Documentation

## 13.59.2.1 test\_answer

```
int test_answer[10] = \{2, 1, 1, 2, 0, 2, 3, 0, 4, 4\}
```

## 13.59.2.2 test\_answer1

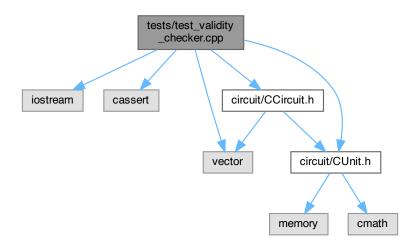
```
int test_answer1[31]
```

## Initial value:

```
= {0, 10, 1, 11, 10, 2, 11, 10, 3, 11, 10, 4, 11, 10, 5, 11, 10, 6, 11, 10, 7, 11, 10, 8, 11, 10, 9, 11, 10, 0, 11}
```

# 13.60 tests/test\_validity\_checker.cpp File Reference

```
#include <iostream>
#include <cassert>
#include <vector>
#include "circuit/CUnit.h"
#include "circuit/CCircuit.h"
Include dependency graph for test_validity_checker.cpp:
```



## **Functions**

void test\_check\_validity ()

Test function for checking the validity of circuit configurations.

• int main (int argc, char \*argv[])

Main function to run the circuit validity tests.

## 13.60.1 Function Documentation

## 13.60.1.1 main()

```
int main (
                      int argc,
                      char * argv[] )
```

Main function to run the circuit validity tests.

This function runs the test\_check\_validity function which includes various test cases for checking the validity of circuit configurations.

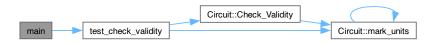
#### **Parameters**

argc	Number of command line arguments
argv	Array of command line arguments

Returns

Exit status of the program

Here is the call graph for this function:

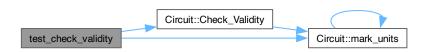


## 13.60.1.2 test\_check\_validity()

void test\_check\_validity ( )

Test function for checking the validity of circuit configurations.

This function runs various test cases to ensure that the circuit validity checks in the CCircuit class are functioning correctly. It includes tests for valid circuits, circuits with incorrect vector sizes, self-recycling circuits, negative feed/destination IDs, and other invalid circuit configurations. Here is the call graph for this function:



Here is the caller graph for this function:



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