

# sP Exam Mini-Project

Nikolaj Kofod Krogh

June 14, 2024

## Conclusion

I have included CMakeLists.txt that shows how to build the project and run the tests. It is included as the first listing.

As we can see from the benchmark it is much faster to run the simulation in parallel (22.1 s) compared to running on a single thread (89.7 s), because it is possible to use multiple threads and thereby distribute the load.

```
COVID-19 simulation took 89692.4 ms (89.7 seconds)
Parallel COVID-19 simulation took 22098.6 ms (22.1 seconds)
```

Figure 1: benchmark

```
COVID19 SEIHR_5822763
Peak hospitalized: 1180
```

Figure 2: peak hospitalized NDK

```
COVID19 SEIHR_589755
Peak hospitalized: 115
```

Figure 3: peak hospitalized NNJ

```
Average peak hospitalized over 100 simulations: 3.64
```

Figure 4: average peak over 100 simulations with a population size of 10000

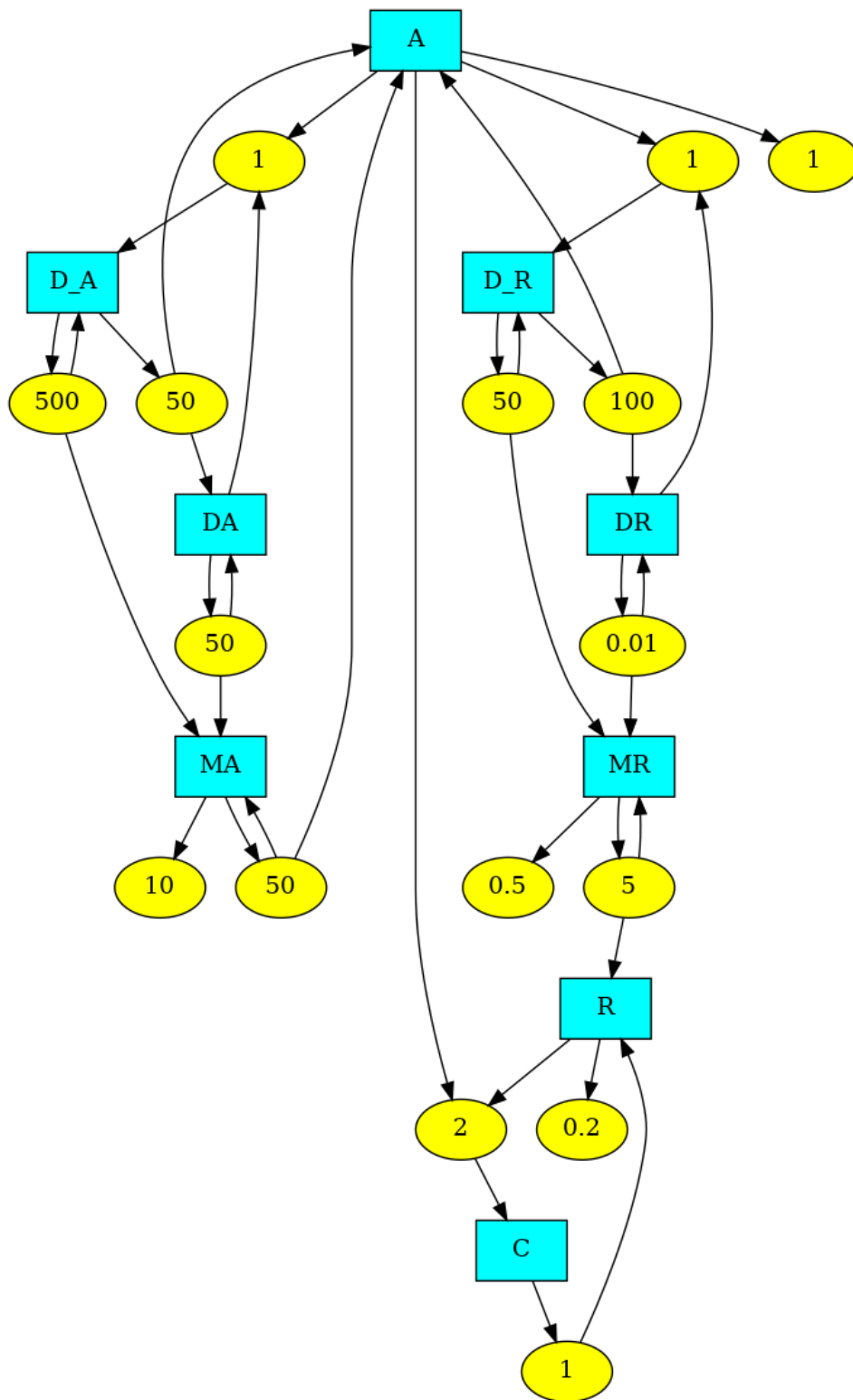


Figure 5: Circadian Rhythm tree

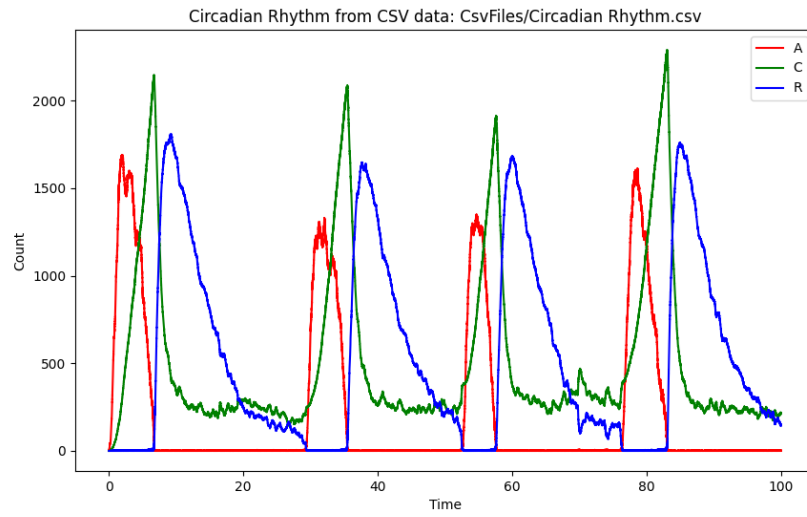


Figure 6: Circadian Rhythm in images

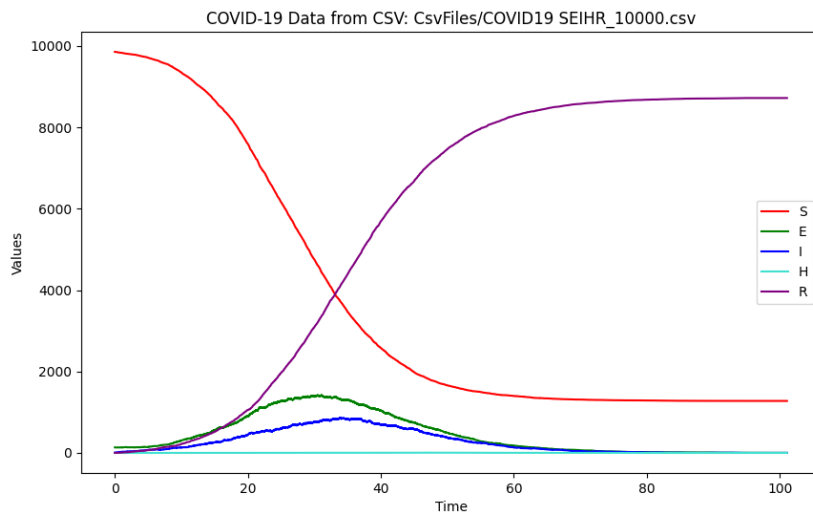


Figure 7: COVID19 population size 10000

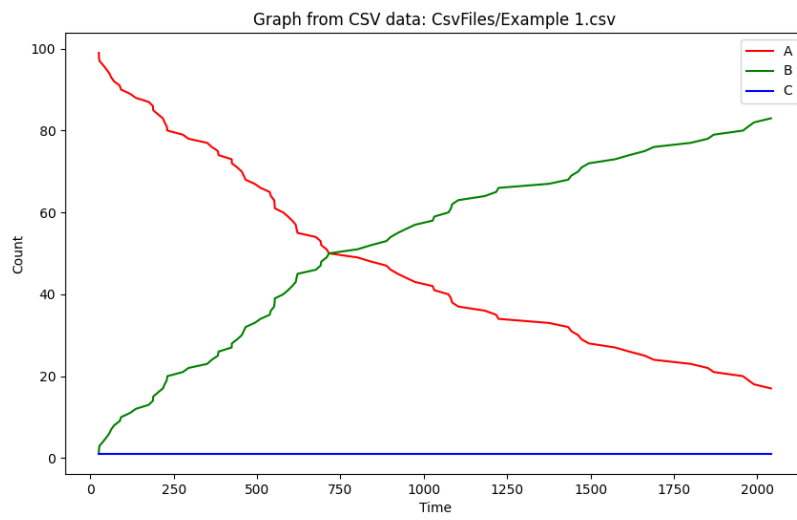


Figure 8: Example 1

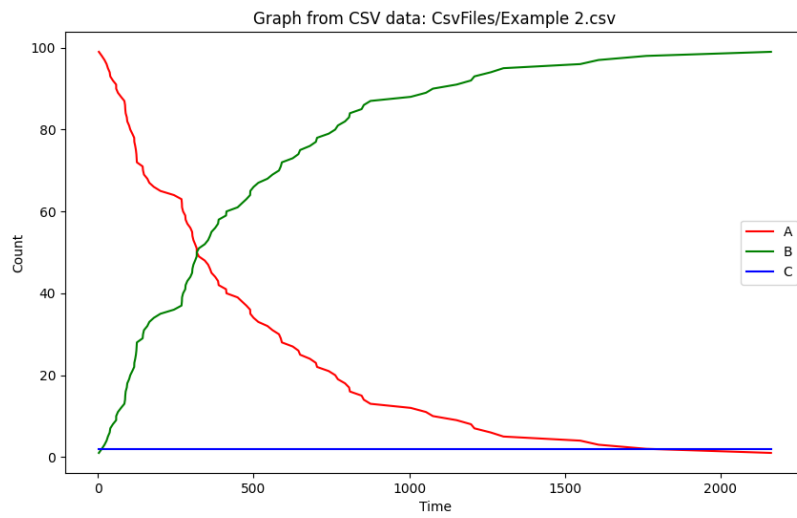


Figure 9: Example 2

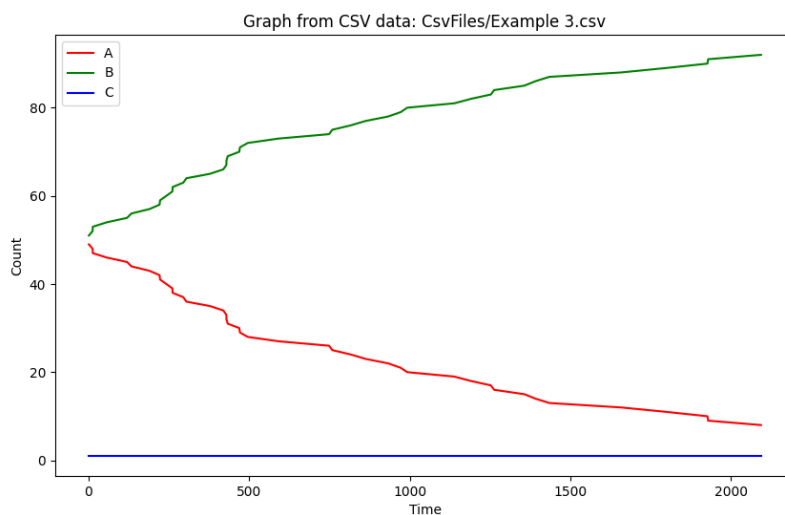


Figure 10: Example 3

Listing 1: ./CMakeLists.txt

```

1  cmake_minimum_required(VERSION 3.27)
2  project(Exam)
3
4  set(CMAKE_CXX_STANDARD 20)
5  enable_testing()
6
7  # Locate GTest
8  find_package(GTest REQUIRED)
9  include_directories(${GTEST_INCLUDE_DIRS})
10
11 # Define common source files
12 set(COMMON_SRC
13     Arrow.cpp
14     Arrow.h
15     benchmark.h
16     benchmark.cpp
17     Environment.cpp
18     Environment.h
19     Molecule.cpp
20     Molecule.h
21     ParallelSimulation.h
22     Reaction.cpp
23     Reaction.h
24     Simulation.h
25     SymbolTable.h
26     Vessel.cpp
27     Vessel.h
28 )
29
30 add_executable(Exam main.cpp ${COMMON_SRC})
31
32 # Define test source files
33 set(TEST_SRC
34     Tests/SymbolTableTests.cpp
35     Tests/VesselTests.cpp
36 )
37
38 # Create a test executable for each test source file
39 foreach(test_src ${TEST_SRC})

```

```

40     get_filename_component(test_name ${test_src} NAME_WE)
41     add_executable(${test_name} ${test_src} ${COMMON_SRC})
42     target_link_libraries(${test_name} ${GTEST_LIBRARIES} ${GTEST_MAIN_LIBRARIES} pthread)
43 endforeach()

```

---

Listing 2: ./main.cpp

```

1  #include <iostream>
2  #include <thread>
3  #include <cmath>
4  #include "Vessel.h"
5  #include "Simulation.h"
6  #include "ParallelSimulation.h"
7  #include "benchmark.h"
8  #include <vector>
9
10
11 stochastic::Vessel simulation_circadian_rhythm() {
12     const auto alphaA = 50;
13     const auto alpha_A = 500;
14     const auto alphaR = 0.01;
15     const auto alpha_R = 50;
16     const auto betaA = 50;
17     const auto betaR = 5;
18     const auto gammaA = 1;
19     const auto gammaR = 1;
20     const auto gammaC = 2;
21     const auto deltaA = 1;
22     const auto deltaR = 0.2;
23     const auto deltaMA = 10;
24     const auto deltaMR = 0.5;
25     const auto thetaA = 50;
26     const auto thetaR = 100;
27
28
29     auto v = stochastic::Vessel{"Circadian Rhythm"};
30     const auto env = v.environment();
31     const auto DA = v.add("DA", 1);
32     const auto D_A = v.add("D_A", 0);
33     const auto DR = v.add("DR", 1);
34     const auto D_R = v.add("D_R", 0);
35     const auto MA = v.add("MA", 0);
36     const auto MR = v.add("MR", 0);
37     const auto A = v.add("A", 1);
38     const auto R = v.add("R", 0);
39     const auto C = v.add("C", 0);
40     v.add((A + DA) >> gammaA >>= D_A);
41     v.add(D_A >> thetaA >>= DA + A);
42     v.add((A + DR) >> gammaR >>= D_R);
43     v.add(D_R >> thetaR >>= DR + A);
44     v.add(D_A >> alpha_A >>= MA + D_A);
45     v.add(DA >> alphaA >>= MA + DA);
46     v.add(D_R >> alpha_R >>= MR + D_R);
47     v.add(DR >> alphaR >>= MR + DR);
48     v.add(MA >> betaA >>= MA + A);
49     v.add(MR >> betaR >>= MR + R);
50     v.add((A + R) >> gammaC >>= C);
51     v.add(C >> deltaA >>= R);
52     v.add(A >> deltaA >>= env);
53     v.add(R >> deltaR >>= env);
54     v.add(MA >> deltaMA >>= env);
55     v.add(MR >> deltaMR >>= env);

```

```

56
57
58     return v;
59 };
60
61 stochastic::Vessel simulation_covid19(uint32_t population_size) {
62     auto v = stochastic::Vessel{"COVID19 SEIHR_" + std::to_string(population_size)};
63     const auto eps = 0.0009; // initial fraction of infectious
64     const auto I0 = size_t(std::round(eps * population_size)); // initial infectious
65     const auto E0 = size_t(std::round(eps * population_size * 15)); // initial exposed
66     const auto S0 = population_size - I0 - E0; // initial susceptible
67     const auto R0 = 2.4; // initial basic reproductive number
68     const auto alpha = 1.0 / 5.1; // incubation rate (E -> I) ~5.1 days
69     const auto gamma = 1.0 / 3.1; // recovery rate (I -> R) ~3.1 days
70     const auto beta = R0 * gamma; // infection/generation rate (S+I -> E+I)
71     const auto P_H = 0.9e-3; // probability of hospitalization
72     const auto kappa = gamma * P_H * (1.0 - P_H); // hospitalization rate (I -> H)
73     const auto tau = 1.0 / 10.12; // removal rate in hospital (H -> R) ~10.12 days
74     const auto S = v.add("S", S0); // susceptible
75     const auto E = v.add("E", E0); // exposed
76     const auto I = v.add("I", I0); // infectious
77     const auto H = v.add("H", 0); // hospitalized
78     const auto R = v.add("R", 0); // removed/immune (recovered + dead)
79     v.add((S + I) >> beta / population_size >>= E + I); // susceptible becomes exposed by infectious
80     v.add(E >> alpha >>= I); // exposed becomes infectious
81     v.add(I >> gamma >>= R); // infectious becomes removed
82     v.add(I >> kappa >>= H); // infectious becomes hospitalized
83     v.add(H >> tau >>= R); // hospitalized becomes removed
84
85     return v;
86
87 }
88
89 stochastic::Vessel simulation_example1() {
90     auto v = stochastic::Vessel{"Example 1"};
91     const auto env = v.environment();
92     const auto A = v.add("A", 100);
93     const auto B = v.add("B", 0);
94     const auto C = v.add("C", 1);
95     const auto lambda = 0.001;
96     v.add((A + C) >> lambda >>= B + C);
97     return v;
98 }
99
100 stochastic::Vessel simulation_example2() {
101     auto v = stochastic::Vessel{"Example 2"};
102     const auto env = v.environment();
103     const auto A = v.add("A", 100);
104     const auto B = v.add("B", 0);
105     const auto C = v.add("C", 2);
106     const auto lambda = 0.001;
107     v.add((A + C) >> lambda >>= B + C);
108
109
110     return v;
111 }
112
113 stochastic::Vessel simulation_example3() {
114     auto v = stochastic::Vessel{"Example 3"};
115     const auto env = v.environment();
116     const auto A = v.add("A", 50);

```

```

117     const auto B = v.add("B", 50);
118     const auto C = v.add("C", 1);
119     const auto lambda = 0.001;
120     v.add((A + C) >> lambda >>= B + C);
121
122     return v;
123 }
124
125 void single_simulation_test() {
126     auto covid19 = simulation_covid19(10000);
127     auto circadian_rhythm = simulation_circadian_rhythm();
128     auto example1 = simulation_example1();
129     auto example2 = simulation_example2();
130     auto example3 = simulation_example3();
131     std::string path = stochastic::Simulation::assign_unique_filename(covid19.name);
132     stochastic::Simulation::simulate(path, covid19, [(stochastic::Vessel &v, double ↗
↪current_time) {}], 100);
133 }
134
135 void parallel_simulation_test() {
136     auto covid19 = simulation_covid19(10000);
137     auto circadian_rhythm = simulation_circadian_rhythm();
138     auto example1 = simulation_example1();
139     auto example2 = simulation_example2();
140     auto example3 = simulation_example3();
141     auto observer = [(stochastic::Vessel &v, double current_time) {}];
142     stochastic::ParallelSimulation::parallelize_simulations(
143         {example1, example2, example3}, observer, 2000);
144 }
145
146 void peak_hospitalization_NNJ_NDK() {
147
148     auto observer = [(stochastic::Vessel &vessel, double current_time) {
149         static int max_hospitalized = 0;
150         int current_hospitalized = vessel.molecules.get("H")->quantity;
151         if (current_hospitalized > max_hospitalized) {
152             max_hospitalized = current_hospitalized;
153         }
154         if (current_time > 100) {
155             std::cout << vessel.name << "\nPeak hospitalized: " << max_hospitalized << std::endl;
156         }
157     }];
158     auto NDK = 5822763;
159     auto NNJ = 589755;
160     std::vector<int> population_sizes = {NNJ};
161     for (int population_size: population_sizes) {
162         auto vessel = simulation_covid19(population_size);
163         stochastic::Simulation::simulate("", vessel, observer, 100, false);
164     }
165 }
166
167 void average_peak_hospitalized_over_100_simulations() {
168     int sum_max_hospitalized = 0;
169     int number_of_simulations = 100;
170
171     // Observer to keep track of the maximum number of hospitalized
172     auto observer = [&sum_max_hospitalized](stochastic::Vessel &vessel, double current_time) {
173         static int max_hospitalized = 0;
174         int current_hospitalized = vessel.molecules.get("H")->quantity;
175         if (current_hospitalized > max_hospitalized) {
176             max_hospitalized = current_hospitalized;

```



```

177     }
178     if (current_time > 100) {
179         sum_max_hospitalized += max_hospitalized;
180         max_hospitalized = 0; // reset for next simulation
181     }
182 };
183
184 std::vector<stochastic::Vessel> vessels;
185 for (int i = 0; i < number_of_simulations; ++i) {
186     vessels.push_back(simulation_covid19(10000));
187 }
188 stochastic::ParallelSimulation::parallelize_simulations(vessels, observer, 100, false);
189 double average_peak_hospitalized = static_cast<double>(sum_max_hospitalized) / ↵
↵number_of_simulations;
190 std::cout << "Average peak hospitalized over " << number_of_simulations << " simulations: " ↵
↵<< average_peak_hospitalized
191         << std::endl;
192 }
193
194 void pretty_print_circadian_rhythm() {
195     const auto alphaA = 50;
196     const auto alpha_A = 500;
197     const auto alphaR = 0.01;
198     const auto alpha_R = 50;
199     const auto betaA = 50;
200     const auto betaR = 5;
201     const auto gammaA = 1;
202     const auto gammaR = 1;
203     const auto gammaC = 2;
204     const auto deltaA = 1;
205     const auto deltaR = 0.2;
206     const auto deltaMA = 10;
207     const auto deltaMR = 0.5;
208     const auto thetaA = 50;
209     const auto thetaR = 100;
210
211
212     auto v = stochastic::Vessel{"Circadian_Rhythm"};
213     const auto env = v.environment();
214     const auto DA = v.add("DA", 1);
215     const auto D_A = v.add("D_A", 0);
216     const auto DR = v.add("DR", 1);
217     const auto D_R = v.add("D_R", 0);
218     const auto MA = v.add("MA", 0);
219     const auto MR = v.add("MR", 0);
220     const auto A = v.add("A", 1);
221     const auto R = v.add("R", 0);
222     const auto C = v.add("C", 0);
223     v.add((A + DA) >> gammaA >>= D_A);
224     v.add(D_A >> thetaA >>= DA + A);
225     v.add((A + DR) >> gammaR >>= D_R);
226     v.add(D_R >> thetaR >>= DR + A);
227     v.add(D_A >> alpha_A >>= MA + D_A);
228     v.add(DA >> alphaA >>= MA + DA);
229     v.add(D_R >> alpha_R >>= MR + D_R);
230     v.add(DR >> alphaR >>= MR + DR);
231     v.add(MA >> betaA >>= MA + A);
232     v.add(MR >> betaR >>= MR + R);
233     v.add((A + R) >> gammaC >>= C);
234     v.add(C >> deltaA >>= R);
235     v.add(A >> deltaA >>= env);

```

```

236     v.add(R >> deltaR >>= env);
237     v.add(MA >> deltaMA >>= env);
238     v.add(MR >> deltaMR >>= env);
239
240     v.pretty_print();
241     v.get_digraph();
242 }
243
244 void symbol_table_test() {
245     // Create a SymbolTable instance
246     stochastic::SymbolTable<std::string, double> symbolTable;
247
248     // Insert some key-value pairs
249     symbolTable.insert("DA", 1);
250     symbolTable.insert("D_A", 0);
251     symbolTable.insert("DR", 1);
252     symbolTable.insert("D_R", 0);
253     symbolTable.insert("MA", 0);
254     symbolTable.insert("MR", 0);
255     symbolTable.insert("A", 1);
256     symbolTable.insert("R", 0);
257     symbolTable.insert("C", 0);
258
259     // Check if a key exists
260     if (symbolTable.contains("DA")) {
261         std::cout << "DA exists in the symbol table." << std::endl;
262     } else {
263         std::cout << "DA does not exist in the symbol table." << std::endl;
264     }
265
266     // Get the value associated with a key
267     double value = symbolTable.get("DR");
268     std::cout << "The value of DR is: " << value << std::endl;
269
270     // Try to insert a key that already exists
271     symbolTable.insert("DA", 1);
272
273     // Update the value associated with a key
274     symbolTable.update("DA", 2);
275
276     // Update the value associated with a key that does not exist
277     symbolTable.update("test", 2);
278
279 }
280
281 void benchmark() {
282     // Make 100 vessels for single simulation
283     std::vector<stochastic::Vessel> single_simulation_vessels;
284     for (int i = 0; i < 100; ++i) {
285         single_simulation_vessels.push_back(simulation_covid19(10000));
286     }
287     // Make 100 vessels for parallel simulation
288     std::vector<stochastic::Vessel> parallel_simulation_vessels;
289     for (int i = 0; i < 100; ++i) {
290         parallel_simulation_vessels.push_back(simulation_covid19(10000));
291     }
292
293     stochastic::Benchmark benchmark;
294
295     benchmark.start("COVID-19 simulation");
296     for (int i = 0; i < 100; ++i) {

```

```

297         stochastic::Simulation::simulate("", single_simulation_vessels[i], [](stochastic::Vessel
298         ↪ &v, double current_time) {}, 100,
299                                     false);
300     }
301     benchmark.stop("COVID-19 simulation");
302
303     benchmark.start("Parallel COVID-19 simulation");
304     auto observer = [](stochastic::Vessel &v, double current_time) {
305     };
306     stochastic::ParallelSimulation::parallelize_simulations(parallel_simulation_vessels,
307     ↪ observer, 100, false);
308     benchmark.stop("Parallel COVID-19 simulation");
309
310     benchmark.report();
311 }
312
313 int main() {
314     // symbol_table_test();
315     // pretty_print_circadian_rhythm();
316     // single_simulation_test();
317     // parallel_simulation_test();
318     // peak_hospitalization_NNJ_NDK();
319     // average_peak_hospitalized_over_100_simulations();
320     // benchmark();
321     return 0;
322 }

```

Listing 3: ./Molecule.h

```

1  #pragma once
2
3  #include <string>
4  #include "Reaction.h"
5
6  namespace stochastic {
7
8      class Reaction;
9
10     class Molecule {
11     public:
12         std::string name;
13         double quantity;
14         std::string graphviz_tag;
15
16         Molecule(const std::string &name, double quantity);
17
18         ~Molecule() = default;
19
20         void operator-=(double quantity);
21
22         void operator+=(double quantity);
23
24         Reaction operator+(const Molecule &molecule) const;
25
26         Reaction operator+(const Reaction &reaction);
27
28         Reaction operator>>(double reaction_rate) const;
29     };
30
31 } // stochastic

```

```

1  #include "Molecule.h"
2
3  namespace stochastic {
4      Molecule::Molecule(const std::string &name, double quantity) {
5          this->name = name;
6          this->quantity = quantity;
7      }
8
9      /**
10       * ----- Exercise 1 -----
11       * @brief Overloads the '-=' operator to decrease the quantity of the molecule.
12       *
13       * @param q The amount by which the quantity of the molecule should be decreased.
14       */
15     void Molecule::operator-=(double q) {
16         Molecule::quantity -= q;
17     };
18
19     /**
20      * ----- Exercise 1 -----
21      * @brief Overloads the '+=' operator to increase the quantity of the molecule.
22      *
23      * @param q The amount by which the quantity of the molecule should be increased.
24      */
25     void Molecule::operator+=(double q) {
26         Molecule::quantity += q;
27     }
28
29     /**
30      * ----- Exercise 1 -----
31      * @brief Overloads the '+' operator to create a reaction with two molecules as reactants.
32      *
33      * This function creates a new reaction and adds the current molecule.
34      *
35      * @param molecule The molecule that reacts with the current molecule.
36      * @return The created reaction with the two molecules as reactants.
37      */
38     Reaction Molecule::operator+(const Molecule &molecule) const {
39         auto r = Reaction();
40         // dereference the current molecule and add it to the reactants of the reaction
41         r.reactants.push_back(*this);
42         r.reactants.push_back((molecule));
43         return r;
44     }
45
46     /**
47      * ----- Exercise 1 -----
48      * @brief Overloads the '+' operator to create a reaction with the current molecule and the
49      * → reactants of another reaction.
50      *
51      * This function creates a new reaction and adds the current molecule and the reactants of
52      * → the reaction.
53      *
54      * @param reaction The reaction whose reactants will react with the current molecule.
55      * @return The created reaction with the current molecule and the reactants of the passed
56      * → reaction as reactants.
57      */
58     Reaction Molecule::operator+(const Reaction &reaction) {

```

```

56     auto r = Reaction();
57     r.reactants.push_back(*this);
58     for (auto reactant: reaction.reactants) {
59         r.reactants.push_back(reactant);
60     }
61
62     return r;
63 }
64
65 /**
66  * ----- Exercise 1 -----
67  * @brief Overloads the '>>' operator to create a reaction with the current molecule as a
68  * reactant and a specified rate.
69  * This function creates a new reaction and adds the current molecule as a reactant.
70  *
71  * @param rate The rate at which the current molecule reacts.
72  * @return The created reaction with the current molecule as a reactant and the specified rate.
73  */
74 Reaction Molecule::operator>>(double rate) const {
75     auto reaction = Reaction();
76     reaction.reactants.push_back(*this);
77     reaction.rate = rate;
78     return reaction;
79 }
80 } // stochastic

```

Listing 5: ./Reaction.h

```

1  #pragma once
2
3  #include <string>
4  #include <vector>
5  #include "Molecule.h"
6  #include "Environment.h"
7
8  namespace stochastic {
9
10     // Required to avoid circular dependencies
11     class Molecule;
12
13     // Use enum to check if we should add the reactants to the left or right side of the reaction
14     typedef enum {
15         left,
16         right
17     } reaction_side;
18
19     class Reaction {
20     public:
21         std::string name;
22         double rate;
23         reaction_side side = left;
24         std::vector<Molecule> products;
25         std::vector<Molecule> reactants;
26         double delay;
27
28         Reaction(const std::string &name, double rate, const std::vector<Molecule> &reactants,
29                 const std::vector<Molecule> &products);
30
31         Reaction();
32
33         ~Reaction() = default;

```

```

34
35     Reaction operator>>(double reaction_rate);
36
37     Reaction operator+(const Molecule &molecule);
38
39     Reaction operator>>=(const Molecule &products);
40
41     Reaction operator>>=(const Environment &env);
42
43     Reaction operator>>=(const Reaction &reaction);
44
45 };
46
47
48 } // stochastic

```

Listing 6: ./Reaction.cpp

```

1  #include "Reaction.h"
2
3  namespace stochastic {
4      Reaction::Reaction(const std::string &name, double rate, const std::vector<Molecule> &r,
5                          const std::vector<Molecule> &p) {
6          this->name = name;
7          this->rate = rate;
8          this->reactants = r;
9          this->products = p;
10     }
11
12     Reaction::Reaction() = default;
13
14
15     /**
16      * ----- Exercise 1 -----
17      * @brief Overloads the '>>' operator to set the rate of the reaction.
18      *
19      * The rate of a reaction is a measure of how quickly the reactants turn into products.
20      *
21      * @param r The new rate of the reaction.
22      * @return The modified reaction with the updated rate.
23      */
24     Reaction Reaction::operator>>(double r) {
25         Reaction::rate = r;
26         return *this;
27     }
28
29
30     /**
31      * ----- Exercise 1 -----
32      * @brief Overloads the '+' operator to add a molecule to the reaction.
33      *
34      * This function adds a molecule to either the reactants or the products of the reaction,
35      * depending on the current side. If the current side is 'left', the molecule is added to
36      * the reactants.
37      * If the current side is 'right', the molecule is added to the products.
38      *
39      * @param m The molecule to be added to the reaction.
40      * @return The modified reaction with the added molecule.
41      */
42     Reaction Reaction::operator+(const Molecule &m) {
43         if (Reaction::side == left) {
44             Reaction::reactants.push_back(m);

```

```

44     } else {
45         Reaction::products.push_back(m);
46     }
47     return *this;
48 }
49
50 /**
51  * ----- Exercise 1 -----
52  * @brief Overloads the '>=>' operator to add a product molecule to the reaction.
53  *
54  * This function sets the current side to 'right' and adds a molecule to the products of the
55  * reaction.
56  *
57  * @param p The product molecule to be added to the reaction.
58  * @return The modified reaction with the added product molecule.
59  */
60 Reaction Reaction::operator>=>(const Molecule &p) {
61     Reaction::side = right;
62     Reaction::products.push_back(p);
63     return *this;
64 }
65
66 /**
67  * ----- Exercise 1 -----
68  * @brief Overloads the '>=>' operator to clear the products of the reaction.
69  *
70  * This function clears all the product molecules from the reaction when an environment is
71  * passed.
72  *
73  * @param env The environment, which when passed, triggers the clearing of the products.
74  * @return The modified reaction with the cleared products.
75  */
76 Reaction Reaction::operator>=>(const Environment &env) {
77     this->products.clear();
78     return *this;
79 }
80
81 /**
82  * ----- Exercise 1 -----
83  * @brief Overloads the '>=>' operator to replace the products of the current reaction with
84  * the reactants of another reaction.
85  *
86  * This function replaces the products of the current reaction with the reactants of the
87  * passed reaction.
88  *
89  * @param reaction The reaction whose reactants will replace the products of the current
90  * reaction.
91  * @return The modified reaction with the replaced products.
92  */
93 Reaction Reaction::operator>=>(const Reaction &reaction) {
94     Reaction::products = reaction.reactants;
95     return *this;
96 }
97 } // stochastic

```

Listing 7: ./Environment.h

```

1 #pragma once
2
3 namespace stochastic {
4
5     class Environment {

```

```

6     private:
7
8     public:
9         Environment();
10
11         ~Environment() = default;
12     };
13
14 } // stochastic

```

Listing 8: ./Environment.cpp

```

1 #include "Environment.h"
2
3 namespace stochastic {
4     /**
5      * @brief Environment should be empty because the reactions decay into the environment
6      */
7     Environment::Environment() = default;
8
9 } // stochastic

```

Listing 9: ./Vessel.h

```

1 #pragma once
2
3 #include "Reaction.h"
4 #include "SymbolTable.h"
5 #include "Arrow.h"
6
7 namespace stochastic {
8
9     class Vessel {
10     public:
11         stochastic::SymbolTable<std::string, Molecule *> molecules;
12         std::vector<Reaction> reactions;
13         std::string name;
14
15         Vessel(const std::string &name);
16
17         ~Vessel() = default;
18
19
20         Molecule &add(const std::string &name, double quantity);
21
22         void add(const Reaction &reaction);
23
24         Environment environment();
25
26         void pretty_print();
27
28         void get_digraph();
29
30         void assign_tags(SymbolTable <std::string, std::string> &table, int &counter);
31
32         std::vector<Arrow> create_arrows(const SymbolTable <std::string, std::string> &table);
33
34         void write_to_file(const std::vector<Arrow> &arrows, const SymbolTable <std::string,
35         ↪std::string> &table);
36     };
37 } // stochastic

```



```

1  #include <iostream>
2  #include <set>
3  #include <fstream>
4  #include "Vessel.h"
5  #include "Arrow.h"
6
7  namespace stochastic {
8      Vessel::Vessel(const std::string &name) {
9          this->name = name;
10     }
11
12     Environment Vessel::environment() {
13         return Environment();
14     }
15
16     /**
17      * ----- Exercise 3 -----
18      * @brief Adds a molecule to the vessel.
19      *
20      * This function creates a new molecule with the given name and quantity, adds it to the
21      * vessel's collection of molecules, and returns a reference to the molecule. ↗
22      *
23      * @param name The name of the molecule to be added.
24      * @param quantity The quantity of the molecule to be added.
25      * @return A reference to the newly created molecule.
26      */
27     Molecule &Vessel::add(const std::string &name, double quantity) {
28         auto m = new Molecule(name, quantity);
29         molecules.insert(name, m);
30         return *m;
31     }
32
33     /**
34      * ----- Exercise 3 -----
35      * @brief Adds a reaction to the vessel.
36      * @param reaction The reaction to be added.
37      */
38     void Vessel::add(const Reaction &reaction) {
39         reactions.push_back(reaction);
40     }
41
42     /**
43      * ----- Exercise 2 -----
44      * @brief Prints a formatted representation of the reactions in the vessel.
45      *
46      * For each reaction, it prints the names of the reactants, the reaction rate, and the names
47      * of the products. ↗
48      * If a reactant or product is not the last one in its list, a plus sign is printed after
49      * its name. ↗
50      * If there are no products for a reaction, it prints 'none'.
51      */
52     void Vessel::pretty_print() {
53         // Iterate over each reaction
54         for (const auto &reaction: reactions) {
55             // Iterate over each reactant in the reaction
56             for (auto &reactant: reaction.reactants) {
57                 std::cout << reactant.name;

```

```

58
59         if (&reactant != &reaction.reactants.back()) {
60             std::cout << " + ";
61         }
62     }
63
64     // Print the reaction rate
65     std::cout << " -> " << reaction.rate << " -> ";
66
67     // Check if there are any products in the reaction
68     if (reaction.products.empty()) {
69         std::cout << "none";
70     } else {
71         // If there are products, print each one
72         for (size_t i = 0; i < reaction.products.size(); ++i) {
73             std::cout << reaction.products[i].name;
74             if (i != reaction.products.size() - 1) {
75                 std::cout << " + ";
76             }
77         }
78     }
79     std::cout << std::endl;
80 }
81 }
82
83
84 /**
85  * ----- Exercise 2 -----
86  * @brief Assigns unique tags to each molecule involved in the reactions.
87  *
88  * For each reactant and product in the reaction, it checks if the molecule's name is
89  * already in the symbol table.
90  * If it's not, it inserts the molecule's name into the symbol table with a unique tag ("s"
91  * followed by a counter).
92  * The counter is incremented after each insertion.
93  *
94  * @param table A reference to the symbol table where the molecule names and their
95  * corresponding tags are stored.
96  * @param counter A reference to the counter used to generate unique tags for the molecules.
97  */
98 void Vessel::assign_tags(SymbolTable<std::string, std::string> &table, int &counter) {
99     for (auto &r: reactions) {
100         for (auto &m: r.reactants) {
101             if (!table.contains(m.name)) {
102                 table.insert(m.name, ("s" + std::to_string(counter++)));
103             }
104         }
105         for (auto &m: r.products) {
106             if (!table.contains(m.name)) {
107                 table.insert(m.name, ("s" + std::to_string(counter++)));
108             }
109         }
110     }
111 }
112
113 /**
114  * ----- Exercise 2 -----
115  * @brief Creates a vector of Arrows representing the reactions in the vessel.
116  *
117  * For each reaction, it creates an Arrow object.
118  * For each reactant and product in the reaction, it adds their corresponding tags from the

```

```

116     * It also sets the rate of the Arrow to the rate of the reaction.
117     * The created Arrow is then added to a vector of Arrows, which is returned at the end.
118     *
119     * @param table A reference to the symbol table where the molecule names and their ↵
    ↪corresponding tags are stored.
120     * @return A vector of Arrows representing the reactions in the vessel.
121     */
122     std::vector<Arrow> Vessel::create_arrows(const SymbolTable<std::string, std::string> &table) {
123         std::vector<Arrow> arrows;
124         for (const auto &r: reactions) {
125             Arrow arrow = {};
126             for (const auto &reactant: r.reactants) {
127                 arrow.source.push_back(table.get(reactant.name));
128             }
129             for (const auto &product: r.products) {
130                 arrow.target.push_back(table.get(product.name));
131             }
132             arrow.rate = r.rate;
133             arrows.push_back(arrow);
134         }
135         return arrows;
136     }
137
138     /**
139     * ----- Exercise 2 -----
140     * @brief Writes the reactions represented by Arrows to a .dot file.
141     *
142     * The function opens a file with the name of the vessel and writes the .dot representation ↵
    ↪of the reactions to it.
143     * For each molecule in the vessel, it writes a line in the .dot file with the molecule's ↵
    ↪tag, name, and color.
144     * For each Arrow in the vector, it writes a line in the .dot file with the Arrow's tag, ↵
    ↪rate, source, and target.
145     * The function then closes the file and generates a .png file from the .dot file using the ↵
    ↪'.dot' command.
146     *
147     * @param arrows A vector of Arrows representing the reactions in the vessel.
148     * @param table A reference to the symbol table where the molecule names and their ↵
    ↪corresponding tags are stored.
149     */
150     void Vessel::write_to_file(const std::vector<Arrow> &arrows, const SymbolTable<std::string, ↵
    ↪std::string> &table) {
151         std::ofstream file;
152         std::string path = "/home/krogh/CLionProjects/cpp/Exam/" + name + ".dot";
153         file.open(path);
154         if (!file) {
155             std::cerr << "Unable to open file for writing.\n";
156             return;
157         }
158
159         file << "digraph {\n";
160         int counter = 0;
161         // Create a set to store unique tags
162         std::set<std::string> u_sets;
163         for (const auto &r: reactions) {
164             for (const auto &reactant: r.reactants) {
165                 auto pair = u_sets.insert(table.get(reactant.name));
166                 if (pair.second) {
167                     file << table.get(reactant.name) << "[label=\"" << reactant.name
168                         << R"(",shape="box",style="filled",fillcolor="cyan");]" << '\n';

```

```

169     }
170 }
171
172 for (const auto &product: r.products) {
173     auto pair = u_sets.insert(table.get(product.name));
174     if (pair.second) {
175         file << table.get(product.name) << "[label=\" " << product.name
176             << R"(",shape="box",style="filled",fillcolor="cyan");)" << '\n';
177     }
178 }
179
180 }
181
182 for (const auto &arrow: arrows) {
183     file << "r" << counter << "[label=\" " << arrow.rate
184         << R"(",shape="oval",style="filled",fillcolor="yellow");)"
185         << '\n';
186     for (const auto &src: arrow.source) {
187         file << src << " -> " << "r" << counter << ";\n";
188     }
189
190     for (const auto &target: arrow.target) {
191         file << "r" << counter << " -> " << target << ";\n";
192     }
193     counter++;
194 }
195
196 file << "}\n";
197 file.close();
198
199 std::string command =
200     "dot -Tpng " + path + " -o /home/krogh/CLionProjects/cpp/Exam/images/" +
201     name + "_tree" + ".png";
202 system(command.c_str());
203 }
204
205 /**
206  * ----- Exercise 2 -----
207  * @brief Generates a directed graph representation of the reactions in the vessel.
208  *
209  * This function first creates a symbol table and a counter for assigning unique tags to the
210  * → molecules.
211  * It then calls the assign_tags function to assign unique tags to each molecule involved in
212  * → the reactions.
213  * After that, it calls the create_arrows function to create a vector of Arrows representing
214  * → the reactions in the vessel.
215  * Finally, it calls the write_to_file function to write the reactions represented by the
216  * → Arrows to a .dot file.
217  */
218 void Vessel::get_digraph() {
219
220     SymbolTable<std::string, std::string> table;
221     int counter = 0;
222     assign_tags(table, counter);
223     std::vector<Arrow> arrows = create_arrows(table);
224     write_to_file(arrows, table);
225 }
226
227 } // stochastic

```

```

1  #include <gtest/gtest.h>
2  #include <filesystem>
3  #include "../Vessel.h"
4
5  using namespace stochastic;
6  /**
7   * Entire file is exercise 9
8   */
9
10 TEST(VesselTest, AddMolecule) {
11     Vessel vessel("TestVessel");
12     Molecule &molecule = vessel.add("H2O", 10.0);
13     ASSERT_EQ(molecule.name, "H2O");
14     ASSERT_EQ(molecule.quantity, 10.0);
15 }
16
17 TEST(VesselTest, AddReaction) {
18     Vessel vessel("TestVessel");
19     Reaction reaction;
20     reaction.reactants.push_back(Molecule("H2O", 10.0));
21     reaction.products.push_back(Molecule("H2", 5.0));
22     reaction.products.push_back(Molecule("O2", 5.0));
23     vessel.add(reaction);
24     ASSERT_EQ(vessel.reactions.size(), 1);
25 }
26
27 TEST(VesselTest, PrettyPrintEmptyVessel) {
28     stochastic::Vessel vessel("TestVessel");
29     testing::internal::CaptureStdout();
30     vessel.pretty_print();
31     std::string output = testing::internal::GetCapturedStdout();
32     EXPECT_EQ(output, "");
33 }
34
35 TEST(VesselTest, PrettyPrintVesselWithSingleReactionNoProducts) {
36     stochastic::Vessel vessel("TestVessel");
37     stochastic::Reaction reaction;
38     reaction.reactants.push_back(stochastic::Molecule("H2O", 2));
39     reaction.rate = 1.0;
40     vessel.add(reaction);
41     testing::internal::CaptureStdout();
42     vessel.pretty_print();
43     std::string output = testing::internal::GetCapturedStdout();
44     EXPECT_EQ(output, "H2O -> 1 -> none\n");
45 }
46
47 TEST(VesselTest, PrettyPrintVesselWithSingleReactionWithProducts) {
48     stochastic::Vessel vessel("TestVessel");
49     stochastic::Reaction reaction;
50     reaction.reactants.push_back(stochastic::Molecule("H2O", 2));
51     reaction.products.push_back(stochastic::Molecule("H2", 1));
52     reaction.products.push_back(stochastic::Molecule("O2", 1));
53     reaction.rate = 1.0;
54     vessel.add(reaction);
55     testing::internal::CaptureStdout();
56     vessel.pretty_print();
57     std::string output = testing::internal::GetCapturedStdout();
58     EXPECT_EQ(output, "H2O -> 1 -> H2 + O2\n");
59 }
60

```

```

61 TEST(VesselTest, PrettyPrintVesselWithMultipleReactions) {
62     stochastic::Vessel vessel("TestVessel");
63     stochastic::Reaction reaction1;
64     reaction1.reactants.push_back(stochastic::Molecule("H2O", 2));
65     reaction1.products.push_back(stochastic::Molecule("H2", 1));
66     reaction1.products.push_back(stochastic::Molecule("O2", 1));
67     reaction1.rate = 1.0;
68     vessel.add(reaction1);
69     stochastic::Reaction reaction2;
70     reaction2.reactants.push_back(stochastic::Molecule("H2", 1));
71     reaction2.reactants.push_back(stochastic::Molecule("O2", 1));
72     reaction2.products.push_back(stochastic::Molecule("H2O", 2));
73     reaction2.rate = 2.0;
74     vessel.add(reaction2);
75     testing::internal::CaptureStdout();
76     vessel.pretty_print();
77     std::string output = testing::internal::GetCapturedStdout();
78     EXPECT_EQ(output, "H2O -> 1 -> H2 + O2\nH2 + O2 -> 2 -> H2O\n");
79 }
80
81 TEST(VesselTest, AssignTagsAssignsUniqueTags) {
82     Vessel vessel("TestVessel");
83     Reaction reaction;
84     reaction.reactants.push_back(Molecule("H2O", 10.0));
85     reaction.products.push_back(Molecule("H2", 5.0));
86     reaction.products.push_back(Molecule("O2", 5.0));
87     vessel.add(reaction);
88     SymbolTable<std::string, std::string> table;
89     int counter = 0;
90     vessel.assign_tags(table, counter);
91     ASSERT_EQ(table.get("H2O"), "s0");
92     ASSERT_EQ(table.get("H2"), "s1");
93     ASSERT_EQ(table.get("O2"), "s2");
94 }
95
96 TEST(VesselTest, CreateArrowsCreatesCorrectNumberOfArrows) {
97     Vessel vessel("TestVessel");
98     Reaction reaction;
99     reaction.reactants.push_back(Molecule("H2O", 10.0));
100    reaction.products.push_back(Molecule("H2", 5.0));
101    reaction.products.push_back(Molecule("O2", 5.0));
102    vessel.add(reaction);
103    SymbolTable<std::string, std::string> table;
104    int counter = 0;
105    vessel.assign_tags(table, counter);
106    std::vector<Arrow> arrows = vessel.create_arrows(table);
107    ASSERT_EQ(arrows.size(), 1);
108 }
109
110 TEST(VesselTest, WriteToFileCreatesFile) {
111     Vessel vessel("TestVessel");
112     Reaction reaction;
113     reaction.reactants.push_back(Molecule("H2O", 10.0));
114     reaction.products.push_back(Molecule("H2", 5.0));
115     reaction.products.push_back(Molecule("O2", 5.0));
116     vessel.add(reaction);
117     SymbolTable<std::string, std::string> table;
118     int counter = 0;
119     vessel.assign_tags(table, counter);
120     std::vector<Arrow> arrows = vessel.create_arrows(table);
121     vessel.write_to_file(arrows, table);

```

```

122     ASSERT_TRUE(std::filesystem::exists("/home/krogh/CLionProjects/cpp/Exam/TestVessel.dot"));
123 }

```

Listing 12: ./Arrow.h

```

1  #pragma once
2
3  #include <vector>
4  #include <string>
5
6  namespace stochastic {
7
8      class Arrow {
9      public:
10         std::vector<std::string> source;
11         std::vector<std::string> target;
12         double rate;
13
14         Arrow(std::vector<std::string> src, std::vector<std::string> tgt, double r);
15
16         Arrow() = default;
17
18         ~Arrow() = default;
19     };
20
21 } // stochastic

```

Listing 13: ./Arrow.cpp

```

1  #include "Arrow.h"
2
3  namespace stochastic {
4
5      /**
6       * ----- Exercise 2 -----
7       * @brief Constructs a new Arrow object.
8       *
9       * The Arrow object represents a reaction in the system, with source molecules, target
10      ↪ molecules, and a reaction rate.
11       *
12       * @param src A vector of source molecules' Graphviz tags, represented as strings.
13       * @param tgt A vector of target molecules' Graphviz tags, represented as strings.
14       * @param r The rate of the reaction, represented as a double.
15       */
16      Arrow::Arrow(std::vector<std::string> src, std::vector<std::string> tgt, double r) {
17
18          // Iterating over the source vector and adding each source to the Arrow's source vector
19          for (auto &s: src) {
20              this->source.push_back(s);
21          }
22
23          // Iterating over the target vector and adding each target to the Arrow's target vector
24          for (auto &t: tgt) {
25              this->target.push_back(t);
26          }
27
28          // Setting the rate of the Arrow (reaction)
29          this->rate = r;
30      }
31 } // stochastic

```

```

1  #pragma once
2
3  #include <map>
4  #include <vector>
5
6  namespace stochastic {
7
8      template<typename K, typename V>
9      class SymbolTable {
10     private:
11         std::map<K, V> table;
12     public:
13         SymbolTable() = default;
14
15         ~SymbolTable() = default;
16
17
18         /**
19          * ----- Exercise 3 -----
20          * @brief Inserts a key-value pair into the symbol table.
21          *
22          * This function attempts to insert a given key-value pair into the symbol table.
23          * If the key already exists in the table, it prints a message indicating that the symbol
24          * →already exists.
25          *
26          * @param key The key to insert into the symbol table.
27          * @param value The value to associate with the given key.
28          */
29         void insert(K key, V value) {
30             // If it is pointing to the end of the table, the key does not exist in the table
31             if (table.find(key) != table.end()) {
32                 std::cout << "Symbol already exists in the table" << std::endl;
33             } else {
34                 table[key] = value;
35             }
36         }
37
38         /**
39          * ----- Exercise 3 -----
40          * @brief Retrieves the value associated with the given key from the symbol table.
41          *
42          * This function attempts to find the given key in the symbol table.
43          * If the key is found, it returns the associated value.
44          * If the key is not found, it prints an error message .
45          *
46          * @param key The key to search for in the symbol table.
47          * @return The value associated with the given key.
48          */
49         V get(const K &key) const {
50             if (table.find(key) == table.end()) {
51                 std::cout << "Symbol not found in the table" << std::endl;
52             }
53             return table.find(key)->second;
54         }
55
56         /**
57          * ----- Exercise 3 -----
58          * @brief Checks if the symbol table contains the given key.
59          *
60          * @param key The key to check for in the symbol table.

```



```

60     * @return True if the key exists in the table, false otherwise.
61     */
62     bool contains(K key) {
63         return table.find(key) != table.end();
64     }
65
66
67 /**
68  * ----- Exercise 3 -----
69  * @brief Retrieves all the values in the symbol table.
70  *
71  * This function iterates over all the key-value pairs in the symbol table,
72  * and collects all the values into a vector. The order of the values in the
73  * vector is the same as the order of the key-value pairs in the symbol table.
74  *
75  * @return A vector containing all the values in the symbol table.
76  */
77     std::vector<V> values() {
78         std::vector<V> values;
79         for (const auto &pair: table) {
80             values.push_back(pair.second);
81         }
82         return values;
83     }
84
85
86 /**
87  * ----- Exercise 3 -----
88  * @brief Updates the value associated with a given key in the symbol table.
89  *
90  * If the key already exists in the table, it updates the value.
91  * If the key does not exist, it prints a message indicating that the key was not found.
92  *
93  * @param key The key for which to update the value.
94  * @param value The new value to associate with the given key.
95  */
96     void update(K key, V value) {
97         if (table.find(key) != table.end()) {
98             table[key] = value;
99         } else {
100             std::cout << ("Key not found in SymbolTable") << std::endl;
101         }
102     }
103
104 };
105
106
107 } // stochastic

```

Listing 15: ./Tests/SymbolTableTests.cpp

```

1  #include <gtest/gtest.h>
2  #include "../SymbolTable.h"
3
4  /**
5   * Entire file is exercise 9
6   */
7
8  TEST(SymbolTableTest, InsertAndRetrieveValue) {
9      stochastic::SymbolTable<int, std::string> table;
10     table.insert(1, "one");
11     EXPECT_EQ(table.get(1), "one");

```

```

12 }
13
14 TEST(SymbolTableTest, InsertDuplicateKey) {
15     stochastic::SymbolTable<int, std::string> table;
16     table.insert(1, "one");
17     table.insert(1, "duplicate");
18     EXPECT_EQ(table.get(1), "one");
19 }
20
21 TEST(SymbolTableTest, RetrieveNonExistentKey) {
22     stochastic::SymbolTable<int, std::string> table;
23     std::string default_value;
24     EXPECT_EQ(table.get(1), default_value);
25 }
26
27
28 TEST(SymbolTableTest, CheckKeyExists) {
29     stochastic::SymbolTable<int, std::string> table;
30     table.insert(1, "one");
31     EXPECT_TRUE(table.contains(1));
32     EXPECT_FALSE(table.contains(2));
33 }
34
35 TEST(SymbolTableTest, RetrieveAllValues) {
36     stochastic::SymbolTable<int, std::string> table;
37     table.insert(1, "one");
38     table.insert(2, "two");
39     std::vector<std::string> values = table.values();
40     EXPECT_EQ(values.size(), 2);
41     EXPECT_EQ(values[0], "one");
42     EXPECT_EQ(values[1], "two");
43 }
44
45 TEST(SymbolTableTest, UpdateValue) {
46     stochastic::SymbolTable<int, std::string> table;
47     table.insert(1, "one");
48     table.update(1, "updated");
49     EXPECT_EQ(table.get(1), "updated");
50 }
51
52 TEST(SymbolTableTest, UpdateNonExistentKey) {
53     stochastic::SymbolTable<int, std::string> table;
54     table.update(1, "updated");
55     std::string defaultValue;
56     EXPECT_EQ(table.get(1), defaultValue);
57 }

```

Listing 16: ./Simulation.h

```

1  #pragma once
2
3  #include <string>
4  #include <filesystem>
5  #include <fstream>
6  #include <random>
7  #include "Vessel.h"
8
9  namespace stochastic {
10
11     class Simulation {
12     private:
13         /**

```

```

14      * ----- Exercise 4 -----
15      * @brief Computes the delay for a given reaction in a vessel.
16      *
17      * If any reactant quantity is zero, the function returns infinity, i.e. reaction cannot
    occur.
18      * Otherwise, it calculates the product of the quantities of all reactants, multiplies it
    by the reaction rate,
19      * and uses this to generate a random delay time from an exponential distribution.
20      *
21      * @param vessel The vessel in which the reaction is occurring.
22      * @param reaction The reaction for which the delay is being computed.
23      * @return The computed delay time for the reaction.
24      */
25      static double compute_delay(stochastic::Vessel &vessel, stochastic::Reaction &reaction) {
26          double product = 1;
27          // Iterate over all reactants in the reaction
28          for (const auto &reactant: reaction.reactants) {
29              // If any reactant quantity is zero, return infinity
30              if (vessel.molecules.get(reactant.name)->quantity <= 0) {
31                  return std::numeric_limits<double>::infinity();
32              }
33              // Multiply the product by the quantity of the current reactant
34              product *= vessel.molecules.get(reactant.name)->quantity;
35          }
36          // Calculate the rate of the reaction
37          double rate = reaction.rate * product;
38          // Initialize a random number generator
39          std::random_device rd;
40          std::mt19937 gen(rd());
41          // Initialize an exponential distribution with the calculated rate
42          std::exponential_distribution<> d(rate);
43          // Return a random delay time from the exponential distribution
44          return d(gen);
45      }
46
47
48      /**
49      * ----- Exercise 4 -----
50      * @brief Checks if a reaction can occur in a given vessel.
51      *
52      * @param vessel The vessel in which the reaction is supposed to occur.
53      * @param reaction The reaction that is supposed to occur.
54      * @return A boolean value indicating whether the reaction can occur (true) or not (false).
55      */
56      static bool can_react(Vessel &vessel, Reaction &reaction) {
57          for (const auto &reactant: reaction.reactants) {
58              if (vessel.molecules.get(reactant.name)->quantity <= 0) {
59                  return false;
60              }
61          }
62          return true;
63      }
64
65
66      /**
67      * ----- Exercise 4 -----
68      * @brief Performs a reaction in a given vessel.
69      *
70      * It iterates over the reactants of the reaction, decreasing their quantity in the
    vessel by one.
71      * Then, it iterates over the products of the reaction, increasing their quantity in the

```

```

72     *
73     * @param vessel The vessel in which the reaction is occurring.
74     * @param reaction The reaction that is being performed.
75     */
76     static void perform_reaction(Vessel &vessel, Reaction &reaction) {
77         for (const auto &reactant: reaction.reactants) {
78             vessel.molecules.get(reactant.name)->quantity -= 1;
79         }
80         for (const auto &product: reaction.products) {
81             vessel.molecules.get(product.name)->quantity += 1;
82         }
83     }
84
85
86     public:
87
88     /**
89     * ----- Exercise 4 + 7 (Observer) -----
90     * @brief Simulates the reactions in a vessel over a given period of time.
91     *
92     * The CSV file includes the time and the quantity of each molecule at each time step.
93     * At each time step, it finds the reaction with the smallest delay and performs that ↵
94     * The function continues until the total time has reached the end time specified by the ↵
95     * user.
96     *
97     * @tparam Observer The type of the observer function that is called at each time step.
98     * @param path The path to the CSV file where the results will be written.
99     * @param vessel The vessel in which the reactions are occurring.
100    * @param observer The observer function that is called at each time step.
101    * @param end_time The end time for the simulation.
102    * @param output_to_file A boolean value indicating whether the results should be ↵
103    * written to a file.
104    */
105    template<class Observer>
106    static void simulate(const std::string &path, Vessel &vessel, Observer observer, double ↵
107    end_time,
108                        bool output_to_file = false) {
109        std::ofstream file(path);
110        if (output_to_file) {
111            file << "Time"; // Write the header for the time column
112            for (const auto &molecule: vessel.molecules.values()) {
113                file << "," << molecule->name; // Write the headers for the molecule columns
114            }
115            file << std::endl;
116        }
117
118        double time = 0;
119        while (time <= end_time) {
120            Reaction min_delay_reaction;
121            double min_delay = std::numeric_limits<double>::infinity();
122
123            // Find the reaction with the smallest delay
124            for (auto &reaction: vessel.reactions) {
125                reaction.delay = compute_delay(vessel, reaction);
126                if (reaction.delay < min_delay) {
127                    // Update the minimum delay and the corresponding reaction
128                    min_delay = reaction.delay;
129                    min_delay_reaction = reaction;
130                }
131            }
132
133            perform_reaction(vessel, min_delay_reaction);
134            time += min_delay;
135            observer(time, vessel);
136        }
137    }

```

```

128     }
129
130     time += min_delay;
131     if (can_react(vessel, min_delay_reaction)) {
132         perform_reaction(vessel, min_delay_reaction);
133     }
134
135     observer(vessel, time);
136     if (output_to_file) {
137         file << time;
138         for (const auto &molecule: vessel.molecules.values()) {
139             file << ", " << molecule->quantity; // Write the quantity of each ↵
            ↪molecule to the file
140         }
141         file << std::endl;
142     }
143 }
144 if (output_to_file) {
145     file.close(); // Close the file
146 }
147 }
148
149
150 /**
151  * ----- Exercise 4 -----
152  * @brief Assigns a unique filename for a given name.
153  *
154  * It appends the name to a predefined path and checks if a file with that name already ↵
            ↪exists.
155  * If a file with that name exists, it appends a counter to the name and increments the ↵
            ↪counter until it finds a filename that does not exist.
156  * Once a unique filename is found, it creates an empty file with that name and returns ↵
            ↪the full path to the file.
157  *
158  * @param name The base name for the file.
159  * @return The full path to the newly created file.
160  */
161 static std::string assign_unique_filename(const std::string &name) {
162     int counter = 1;
163     std::string originalPath = "/home/krogh/CLionProjects/cpp/Exam/CsvFiles/" + name + ↵
            ↪".csv";
164     while (std::filesystem::exists(originalPath)) {
165         originalPath =
166             "/home/krogh/CLionProjects/cpp/Exam/CsvFiles/" + name + "_" + ↵
            ↪std::to_string(counter) + ".csv";
167         counter++;
168     }
169     std::ofstream file;
170     file.open(originalPath);
171     file.close();
172     return originalPath;
173 }
174
175
176 };
177 } // stochastic

```

Listing 17: ./ParallelSimulation.h

```

1 #pragma once
2
3 namespace stochastic {

```

```

4
5 class ParallelSimulation {
6 public:
7     /**
8      * ----- Exercise 8 -----
9      * @brief Runs simulations in parallel for multiple vessels.
10     *
11     * This function runs simulations for multiple vessels in parallel. Each simulation is
12     * If the 'output_to_file' parameter is 'true', a unique filename is assigned to each
13     * The function waits for all simulations to finish before returning.
14     *
15     * @tparam Observer The type of the observer function that is called at each time step
16     * @param vessels A vector of vessels for which simulations will be run.
17     * @param observer The observer function that is called at each time step in the simulation.
18     * @param simulation_time The total time for which the simulation will be run.
19     * @param output_to_file A boolean value indicating whether the results of the
20     */
21     template<class Observer>
22     static void parallelize_simulations(std::vector<Vessel> vessels, Observer observer,
23     double simulation_time, bool output_to_file = false) {
24         std::vector<std::thread> threads;
25         std::string path;
26         // For each vessel, assign a unique filename and start a simulation in a new thread.
27         for (auto &v: vessels) {
28             if (output_to_file) {
29                 path = Simulation::Simulation::assign_unique_filename(v.name);
30             }
31             threads.push_back(std::thread(Simulation::simulate<Observer>, path, std::ref(v),
32     observer, simulation_time, output_to_file));
33         }
34         // Wait for all simulations to finish.
35         for (auto &thread: threads) {
36             thread.join();
37         }
38     };
39
40 } // stochastic

```

Listing 18: ./benchmark.cpp

```

1 #include <iostream>
2 #include <iomanip>
3 #include "benchmark.h"
4
5 namespace stochastic {
6     /**
7      * ----- Exercise 10 -----
8      * @brief Starts a new benchmark.
9      *
10     * This function starts a new benchmark with the given name. It records the current time as the
11     * The benchmark is represented as a 'time_point' object, which is added to the 'time_points'
12     *
13     * @param name The name of the benchmark. This name is used to identify the benchmark when
14     */

```

```

14  */
15  void Benchmark::start(const std::string &name) {
16      time_point tp;
17      tp.name = name;
18      tp.start = std::chrono::high_resolution_clock::now();
19      time_points.push_back(tp);
20  }
21
22  /**
23   * ----- Exercise 10 -----
24   * @brief Stops a running benchmark.
25   *
26   * This function stops a running benchmark with the given name. It records the current time as
27   * the stop time of the benchmark.
28   * The benchmark is represented as a 'time_point' object, which is updated in the 'time_points'
29   * vector.
30   *
31   * @param name The name of the benchmark. This name is used to identify the benchmark when
32   * stopping it and reporting its duration.
33   */
34  void Benchmark::stop(const std::string &name) {
35      for (auto &tp: time_points) {
36          if (tp.name == name) {
37              tp.stop = std::chrono::high_resolution_clock::now();
38              return;
39          }
40      }
41  }
42
43  /**
44   * ----- Exercise 10 -----
45   * @brief Reports the duration of all benchmarks.
46   *
47   * This function iterates over all the 'time_point' objects in the 'time_points' vector,
48   * calculates the duration of each benchmark,
49   * and prints the name of the benchmark and its duration in milliseconds and seconds to the
50   * standard output.
51   */
52  void Benchmark::report() {
53      for (const auto &tp: time_points) {
54          std::chrono::duration<double, std::milli> duration = tp.stop - tp.start;
55          std::cout << tp.name << " took " << duration.count() << " ms ("
56              << std::fixed << std::setprecision(1) << duration.count()/1000 << "
57              << seconds)" << std::endl;
58      }
59  }
60  }

```

Listing 19: ./benchmark.h

```

1  #pragma once
2
3  #include <vector>
4  #include <chrono>
5  #include <string>
6
7  namespace stochastic
8  {
9
10     class Benchmark
11     {

```

```

12 public:
13     void start(const std::string &name);
14
15     void stop(const std::string &name);
16
17     void report();
18
19 private:
20     struct time_point
21     {
22         std::string name;
23         std::chrono::high_resolution_clock::time_point start;
24         std::chrono::high_resolution_clock::time_point stop;
25     };
26     std::vector<time_point> time_points;
27 };
28
29 }

```

Listing 20: ./CsvFiles/graphical\_simulation.py

```

1 import pandas as pd
2 import matplotlib.pyplot as plt
3 import os
4
5 """
6 ----- Exercise 6 -----
7 """
8
9
10 def plot_example(file_path):
11     # Load the data
12     data = pd.read_csv(file_path)
13
14     # Plot the data
15     plt.figure(figsize=(10, 6))
16     plt.plot(data['Time'], data['A'], label='A', color='red')
17     plt.plot(data['Time'], data['B'], label='B', color='green')
18     plt.plot(data['Time'], data['C'], label='C', color='blue')
19
20     # Add labels and title
21     plt.xlabel('Time')
22     plt.ylabel('Count')
23     plt.title('Graph from CSV data: ' + file_path)
24     plt.legend()
25
26     # Ensure the directory exists
27     graphs_dir = 'images/'
28     if not os.path.exists(graphs_dir):
29         os.makedirs(graphs_dir)
30
31     # Save the plot to a PNG file
32     plt.savefig('images/' + file_path.split('/')[-1].replace('.csv', '') + '.png', format='png')
33
34
35 def plot_circadian_rhythm(file_path):
36     # Load the data
37     data = pd.read_csv(file_path)
38
39     # Plot the data
40     plt.figure(figsize=(10, 6))
41     plt.plot(data['Time'], data['A'], label='A', color='red')

```



```

42 plt.plot(data['Time'], data['C'], label='C', color='green')
43 # plt.plot(data['Time'], data['DA'], label='DA', color='gray')
44 # plt.plot(data['Time'], data['DR'], label='DR', color='purple')
45 # plt.plot(data['Time'], data['D_A'], label='D_A', color='orange')
46 # plt.plot(data['Time'], data['D_R'], label='D_R', color='yellow')
47 # plt.plot(data['Time'], data['MA'], label='MA', color='pink')
48 # plt.plot(data['Time'], data['MR'], label='MR', color='brown')
49 plt.plot(data['Time'], data['R'], label='R', color='blue')
50
51 # Add labels and title
52 plt.xlabel('Time')
53 plt.ylabel('Count')
54 plt.title('Circadian Rhythm from CSV data: ' + file_path)
55 plt.legend()
56
57 # Ensure the directory exists
58 graphs_dir = 'images/'
59 if not os.path.exists(graphs_dir):
60     os.makedirs(graphs_dir)
61
62 # Save the plot to a PNG file
63 plt.savefig('images/' + file_path.split('/')[-1].replace('.csv', '') + '.png', format='png')
64
65
66 def plot_covid19(file_path):
67     # Load the data
68     data = pd.read_csv(file_path)
69
70     # Plot the data
71     plt.figure(figsize=(10, 6))
72     plt.plot(data['Time'], data['S'], label='S', color='red')
73     plt.plot(data['Time'], data['E'], label='E', color='green')
74     plt.plot(data['Time'], data['I'], label='I', color='blue')
75     plt.plot(data['Time'], data['H'], label='H', color='turquoise')
76     plt.plot(data['Time'], data['R'], label='R', color='purple')
77
78     # Add labels and title
79     plt.xlabel('Time')
80     plt.ylabel('Values')
81     plt.title('COVID-19 Data from CSV: ' + file_path)
82     plt.legend()
83
84     # Ensure the directory exists
85     graphs_dir = 'images/'
86     if not os.path.exists(graphs_dir):
87         os.makedirs(graphs_dir)
88
89     # Save the plot to a PNG file
90     plt.savefig('images/' + file_path.split('/')[-1].replace('.csv', '') + '.png', format='png')
91
92
93 def main():
94     # Call the function for each file
95     plot_example('CsvFiles/Example 1.csv')
96     plot_example('CsvFiles/Example 2.csv')
97     plot_example('CsvFiles/Example 3.csv')
98     plot_circadian_rhythm('CsvFiles/Circadian Rhythm.csv')
99     plot_covid19('CsvFiles/COVID19 SEIHR_10000.csv')
100
101
102 if __name__ == "__main__":

```

