

Practical No.-4

Question 4:

1. Design and Develop a System Dynamic Model
 - i. Create a Stock flow diagram.
 - ii. Adding a plot to Visualize Dynamics.
 - iii. Parameter Variation
 - iv. Calibration

SEIR Model: Susceptible Exposed Infectious Recovered

Aim:

To develop a System Dynamics model using AnyLogic to simulate the spread of a contagious disease based on the SEIR (Susceptible Exposed Infectious Recovered) model.

Theory:

System Dynamics is a methodology for understanding the dynamic behaviour of complex systems over time. It involves creating stock-flow diagrams to represent the system's structure and behaviour, along with equations to describe the flow of entities between stocks. In the context of disease spread, the SEIR model divides the population into four compartments: Susceptible (S), Exposed (E), Infectious (I), and Recovered (R). The model tracks the flow of individuals between these compartments over time.

Procedure:

Step 1: Create a Stock Flow Diagram

- Identify the key components of the SEIR model: Susceptible, Exposed, Infectious, and Recovered.
- Use AnyLogic's graphical interface to create stocks representing each compartment.
- Define flows between the compartments to represent the dynamics of disease transmission and recovery.

Step 2: Adding a Plot to Visualize Dynamics

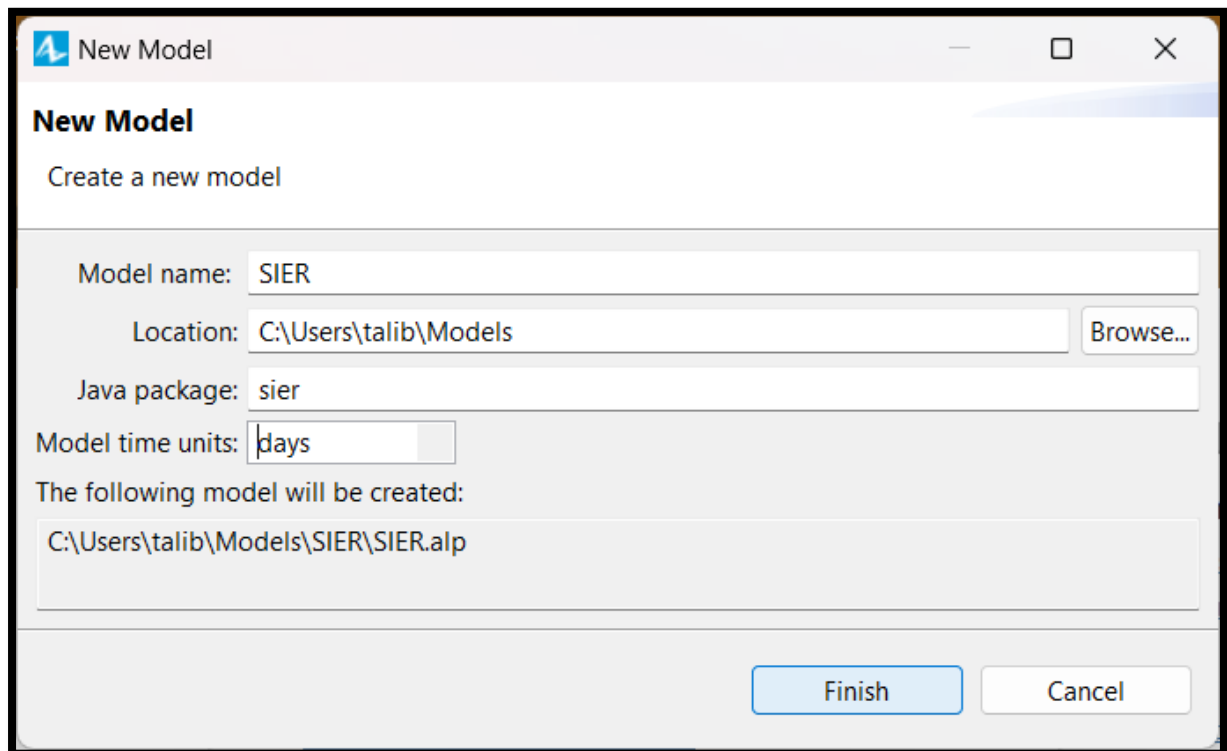
- Add a plot to the model to visualize the changes in the population of each compartment over time.
- Configure the plot to display the population of Susceptible, Exposed, Infectious, and Recovered individuals.

Step 3: Parameter Variation

- Define parameters such as transmission rate, incubation period, and recovery rate.
- Implement parameter variation to observe the effects of different parameter values on the dynamics of disease spread.
- Use sliders or input fields in the model interface to allow users to adjust parameter values interactively.

Step 4: Calibration

- Calibrate the model using real-world data or estimates for parameters such as transmission rate and recovery rate.
- Adjust the model's parameters to match the observed patterns of disease spread in the calibration data.
- Validate the calibrated model by comparing its predictions to historical data or independent sources.



New Model

Create a new model

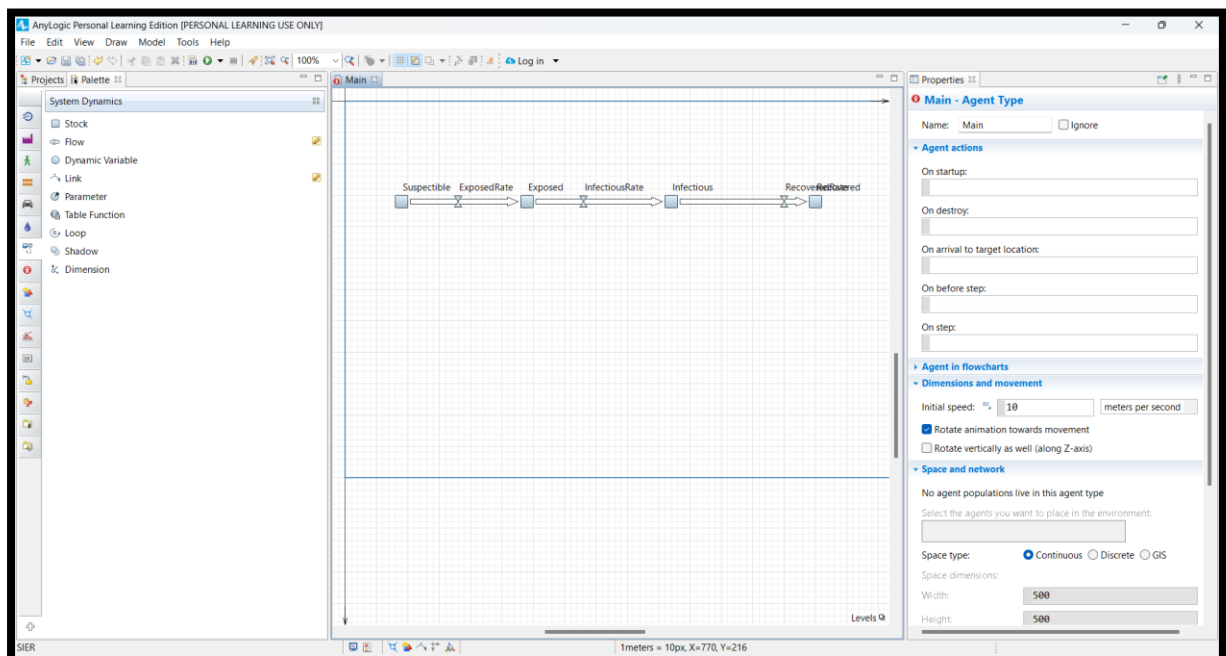
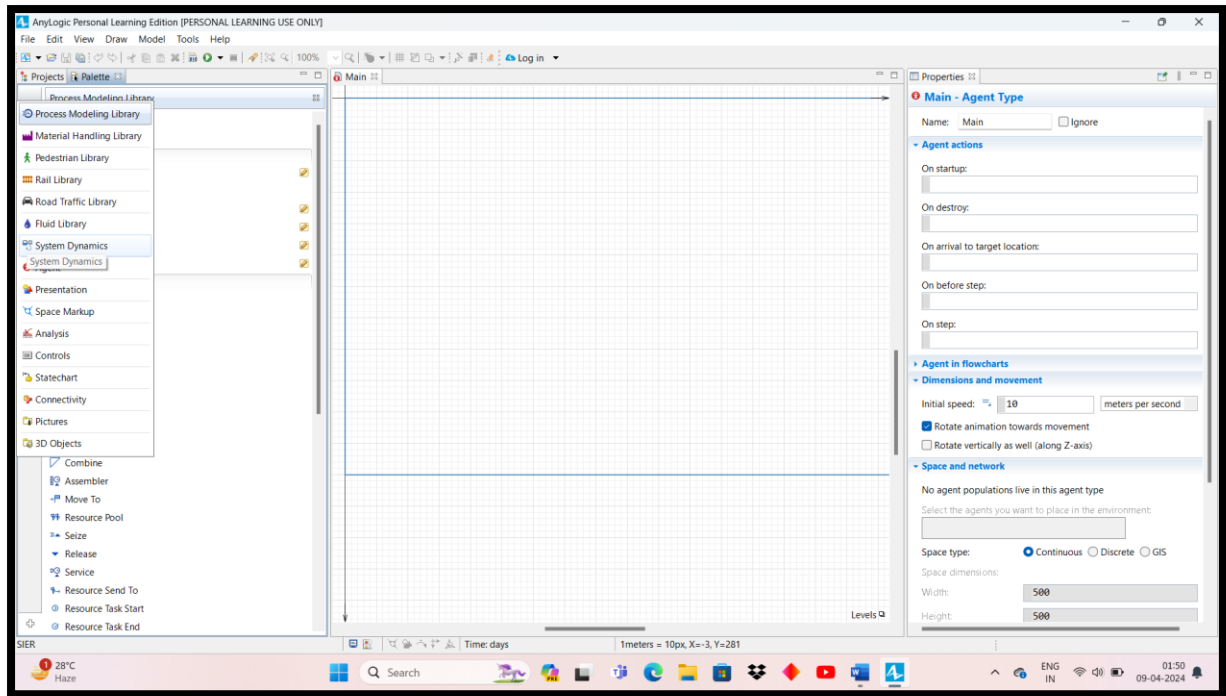
Model name:

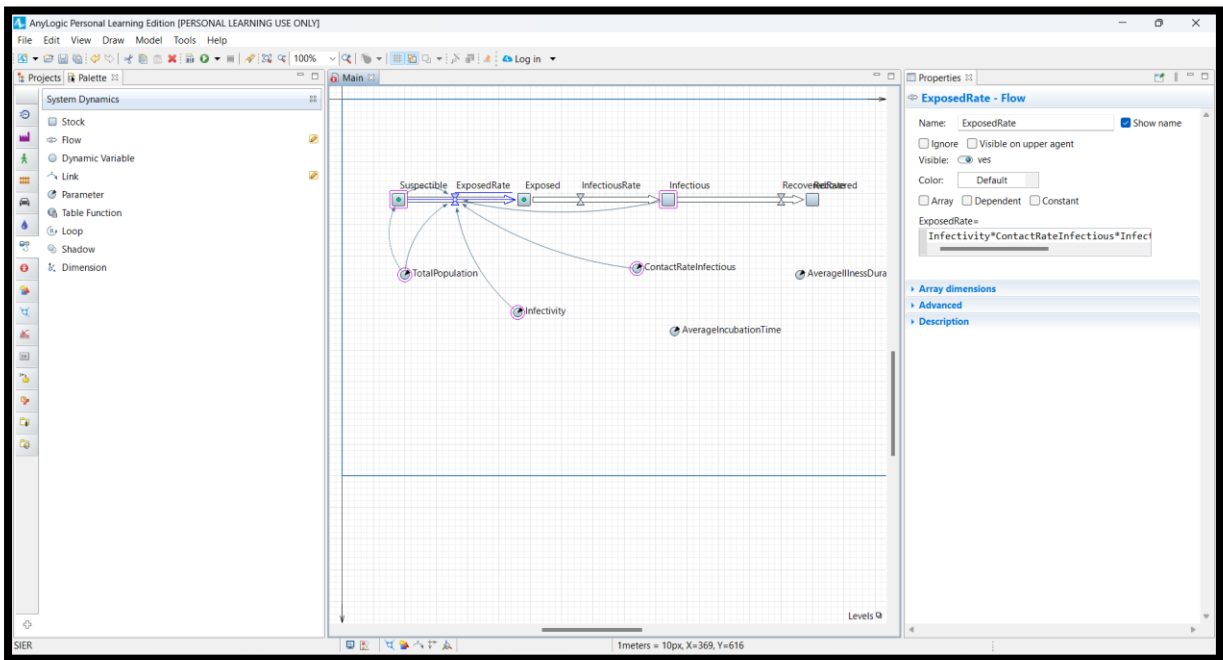
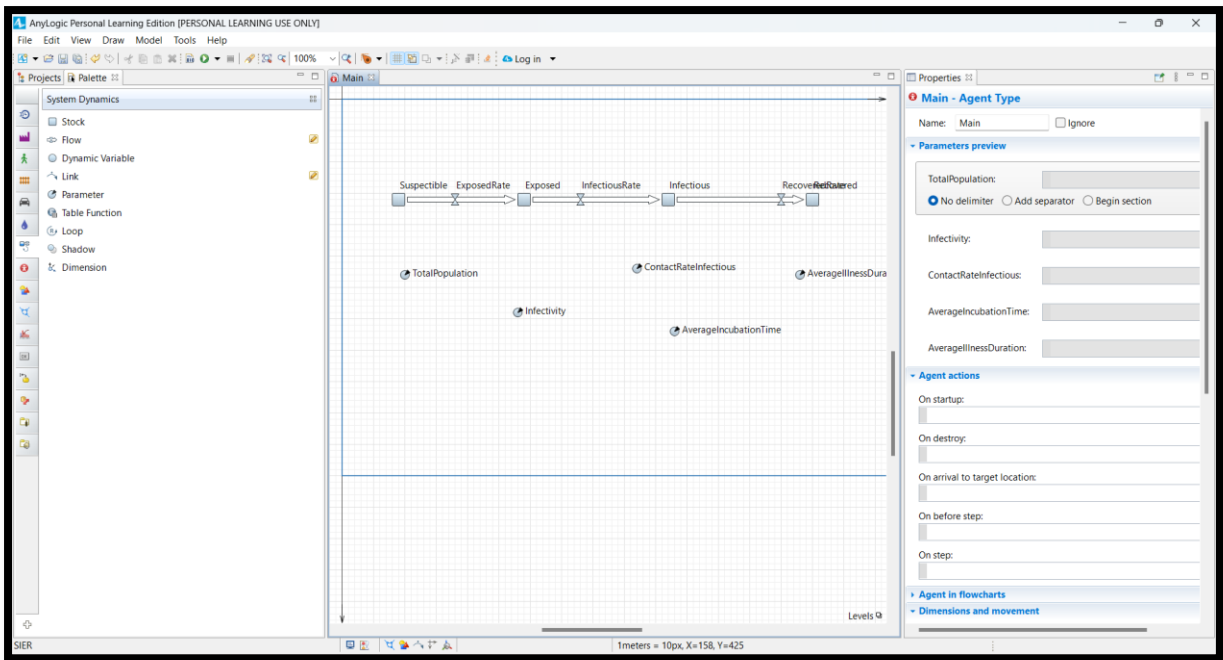
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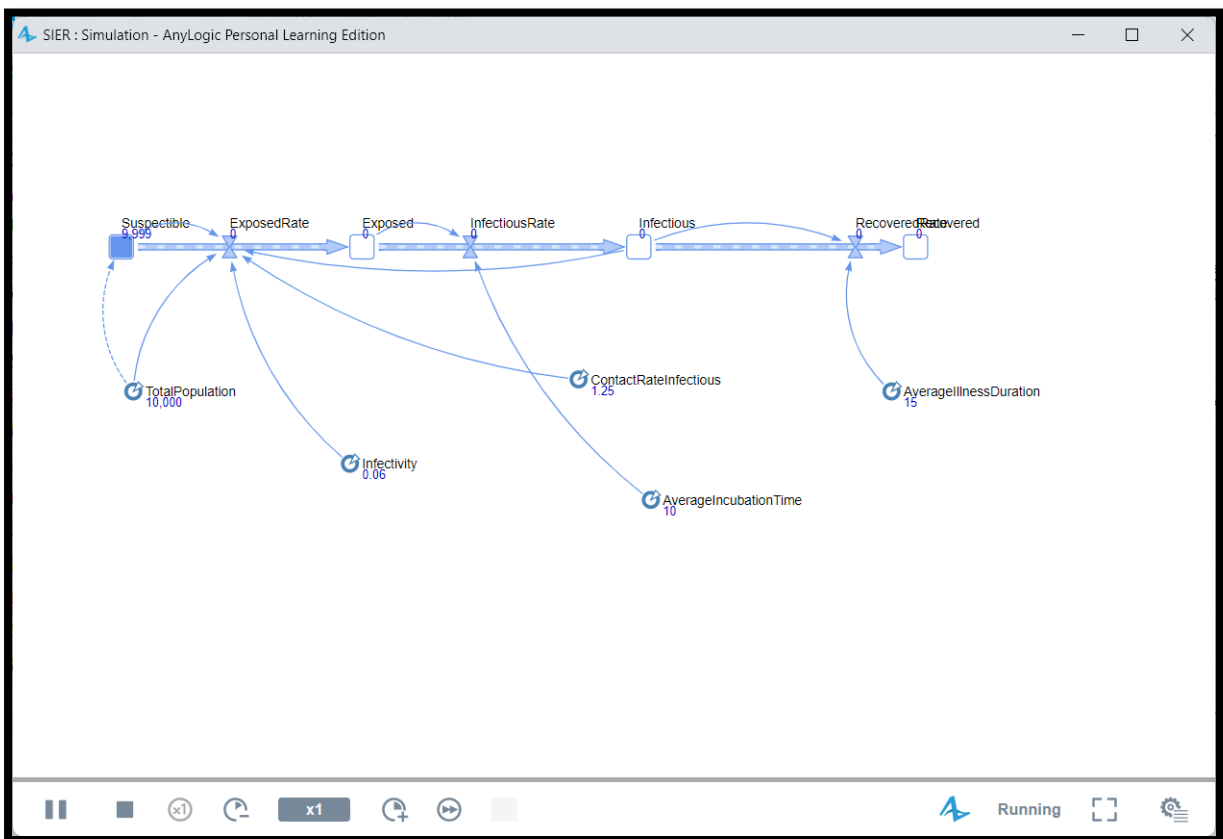
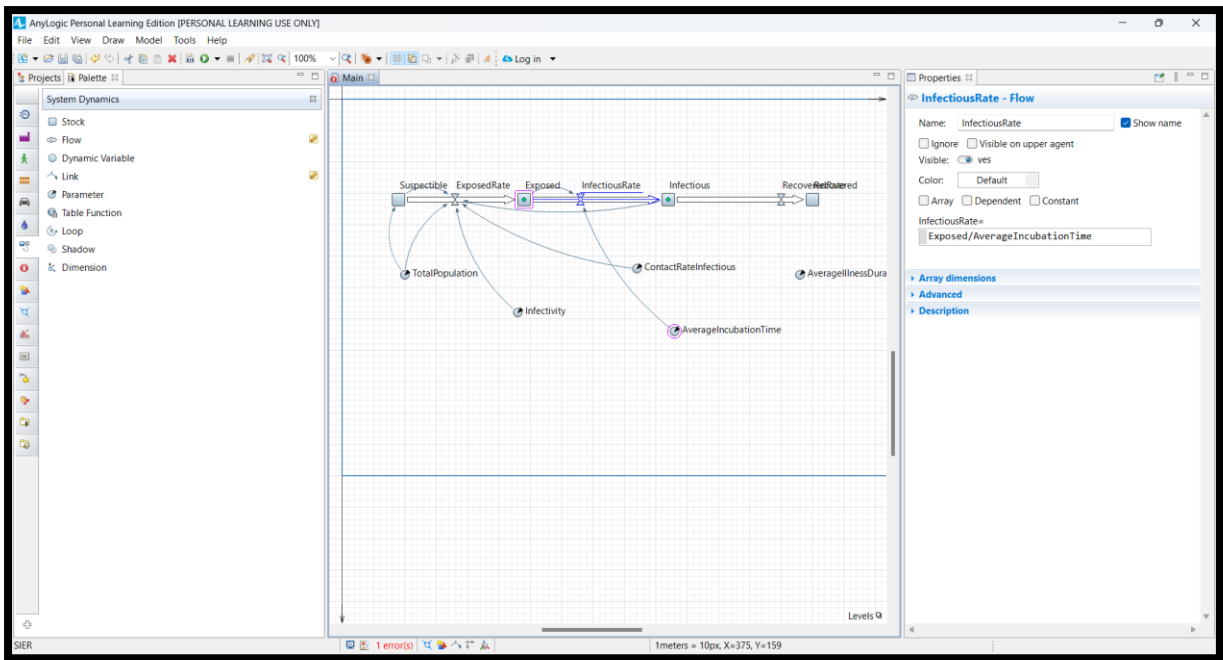
Java package:

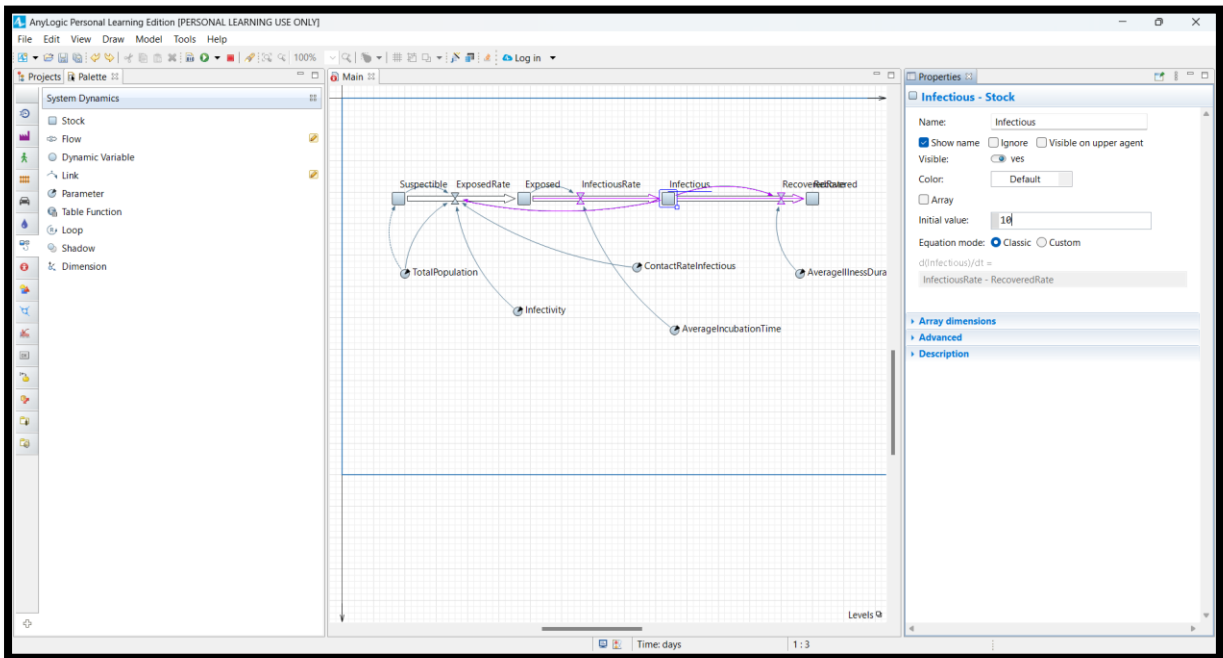
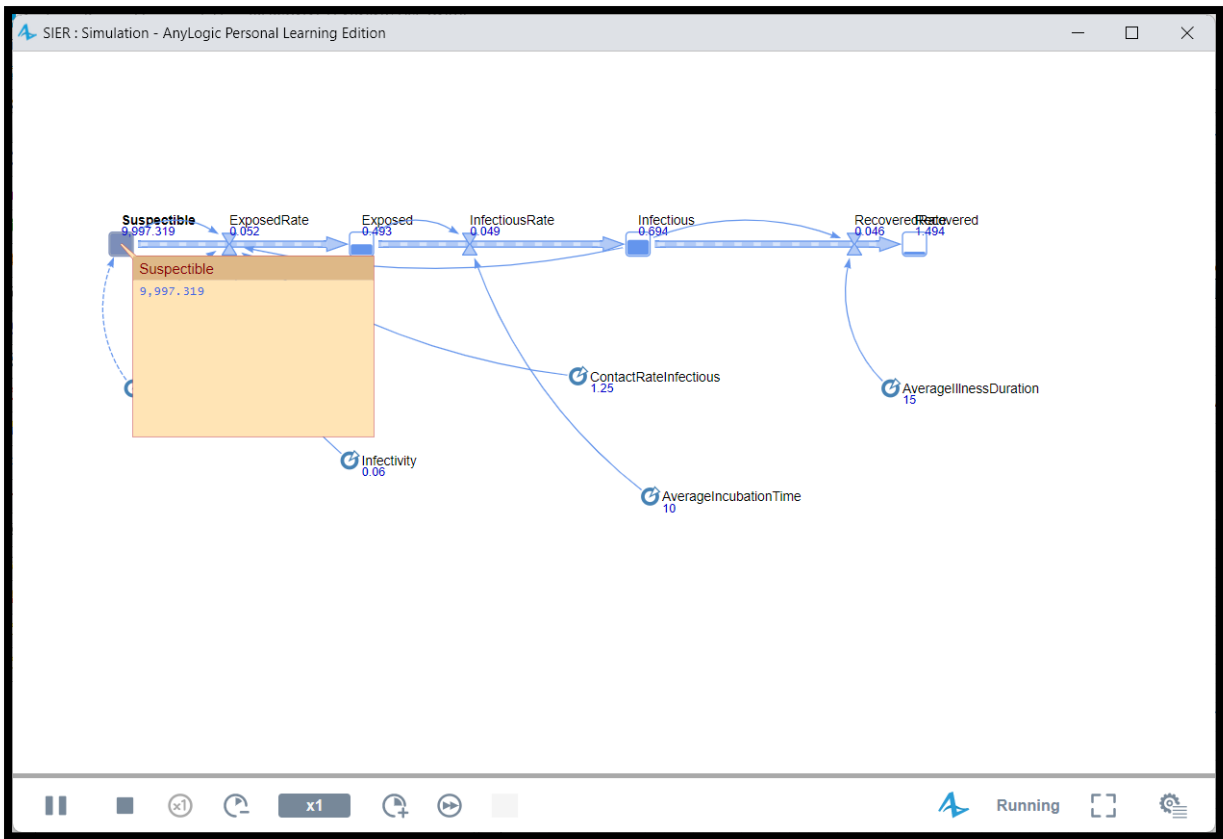
Model time units:

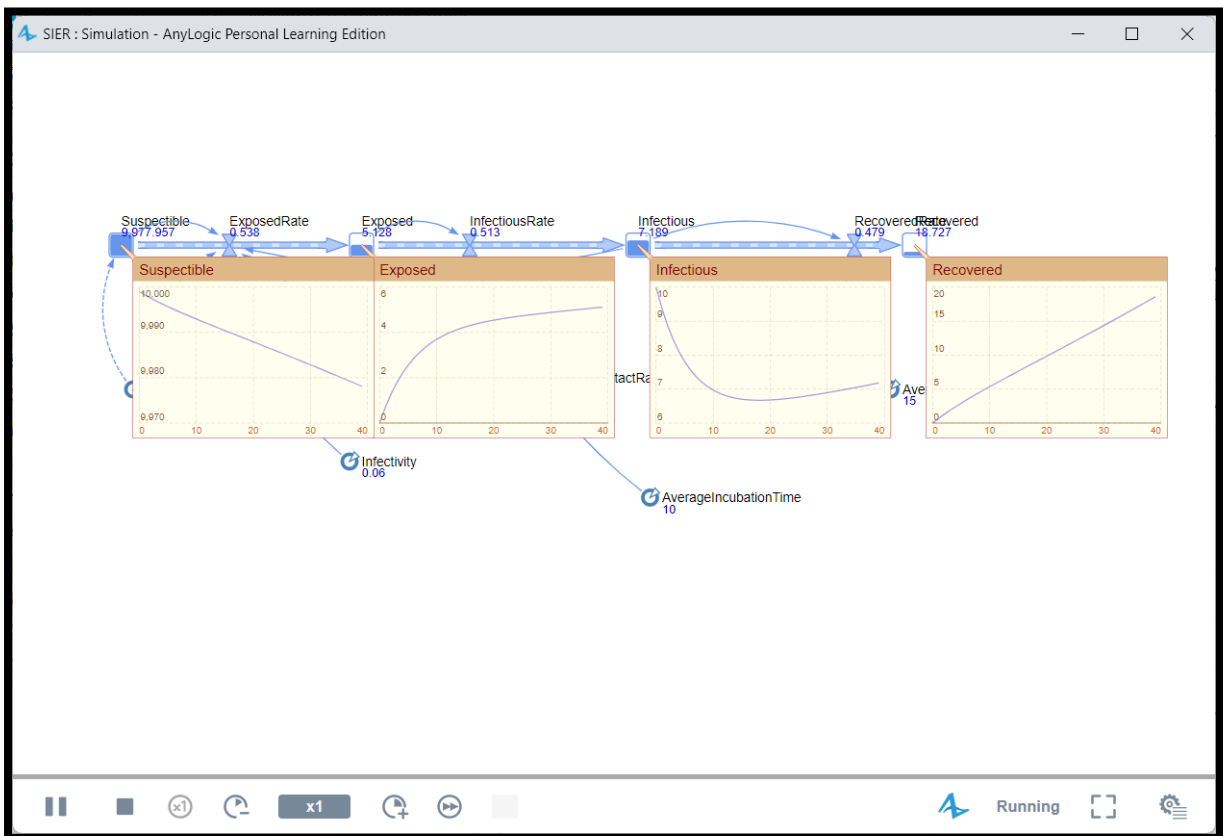
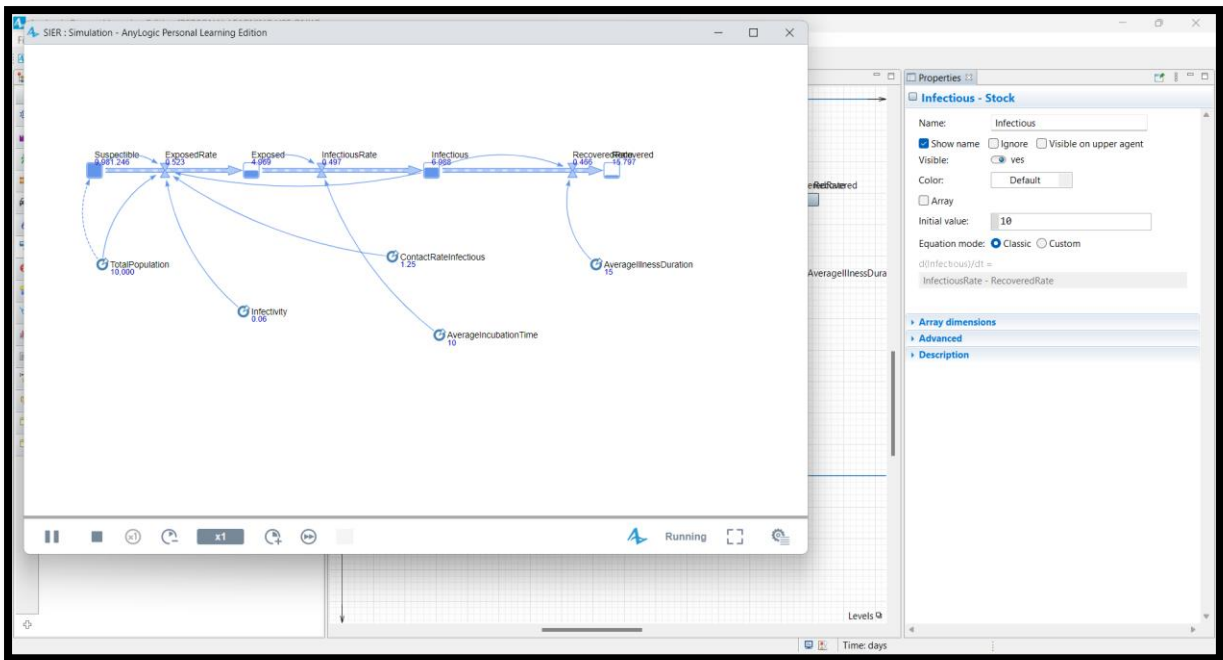
The following model will be created:











▼ Data

☒ Value ☐ Data set

Title: Susceptible People

Value: Susceptible

Point style: 


Line width:  1 pt

Color: dodgerBlue

☒ Value ☐ Data set

Title: Exposed People

Value: Exposed

Point style: 

Line width:  1 pt

Color: darkOrange

☒ Value ☐ Data set

Title: Infectious People

Value: Infectious

Point style: 

Line width:  1 pt

Color: magenta

☒ Value ☐ Data set

Title: Recovered People

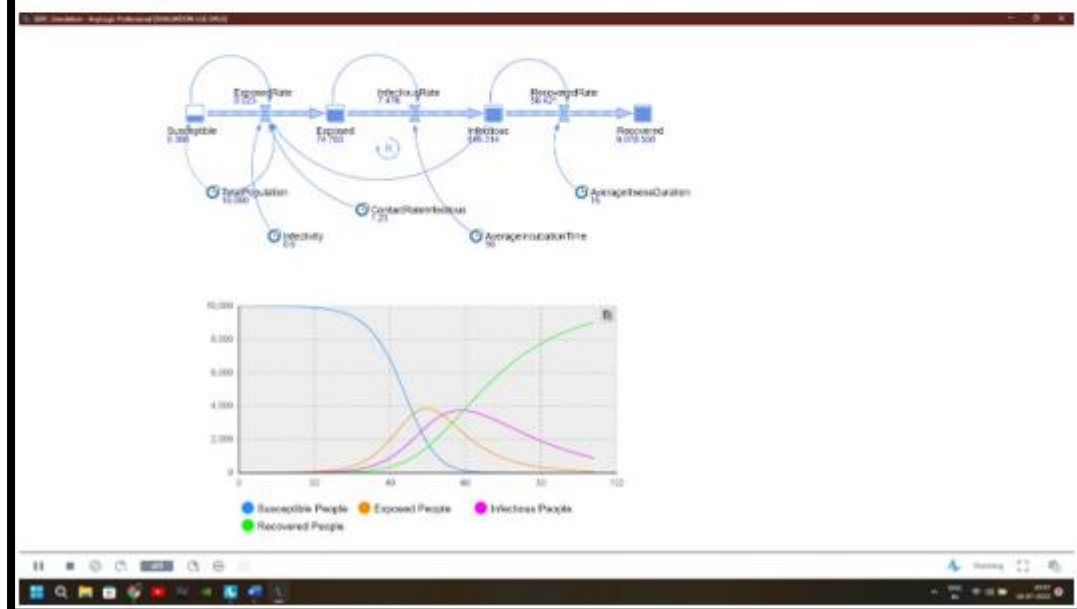
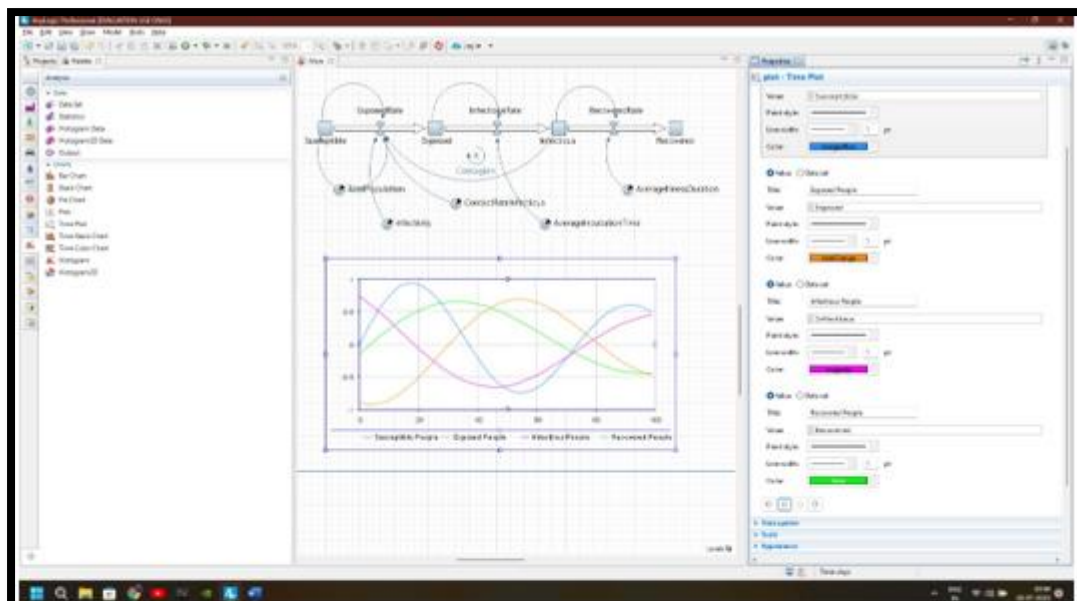
Value: Recovered

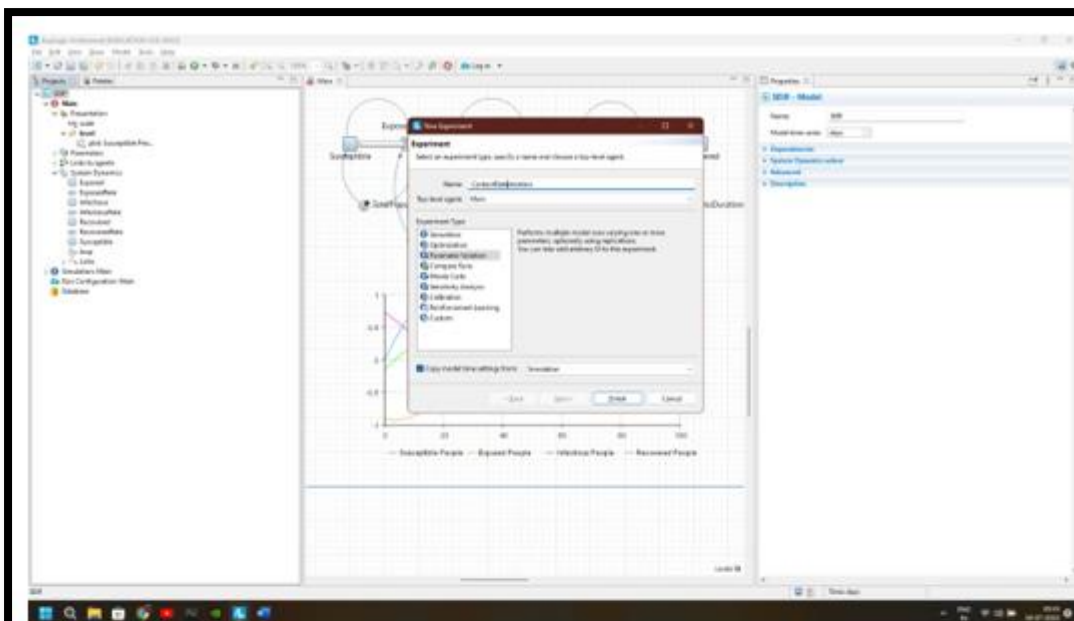
Point style: 

Line width:  1 pt

Color: lime







Parameters

Parameters: ☒ Varied in range ☐ Freeform

Number of runs: 10

Parameter	Type	Value		
		Min	Max	Step
TotalPopulation	Fixed	10000		
Infectivity	Fixed	0.6		
Contact...ectious	Fixed	1.25		
Averag...onTime	Fixed	10		
Average...ration	Fixed	15		

Parameters

Parameters: ☒ Varied in range ☐ Freeform

Number of runs: 10

Parameter	Type	Value		
		Min	Max	Step
TotalPopulation	Fixed	10000		
Infectivity	Fixed	0.6		
Contact...ectious	Range	0.3	2	0.1
Averag...onTime	Fixed	10		
Average...ration	Fixed	15		

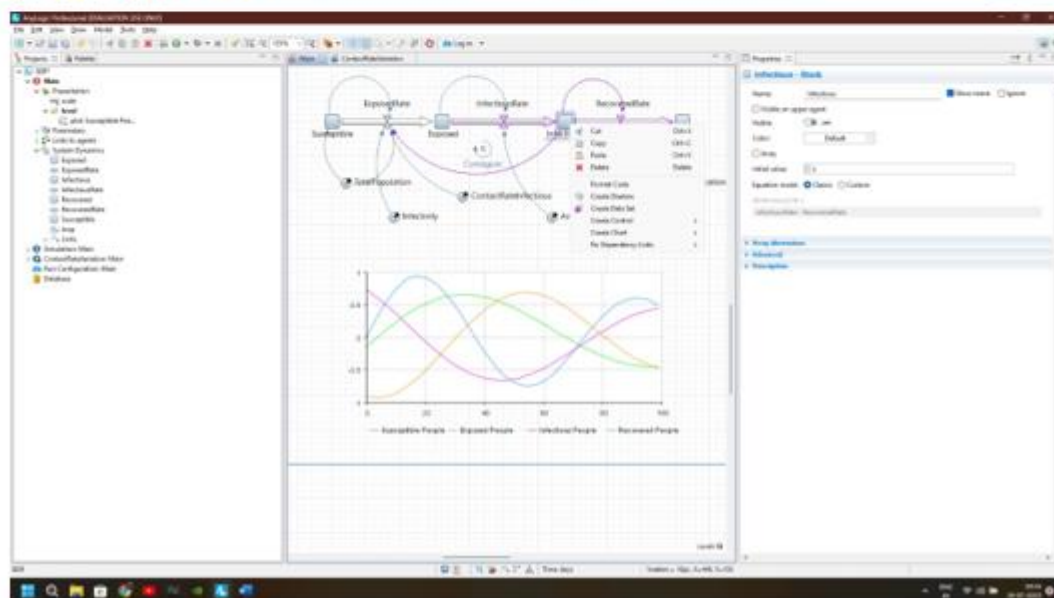
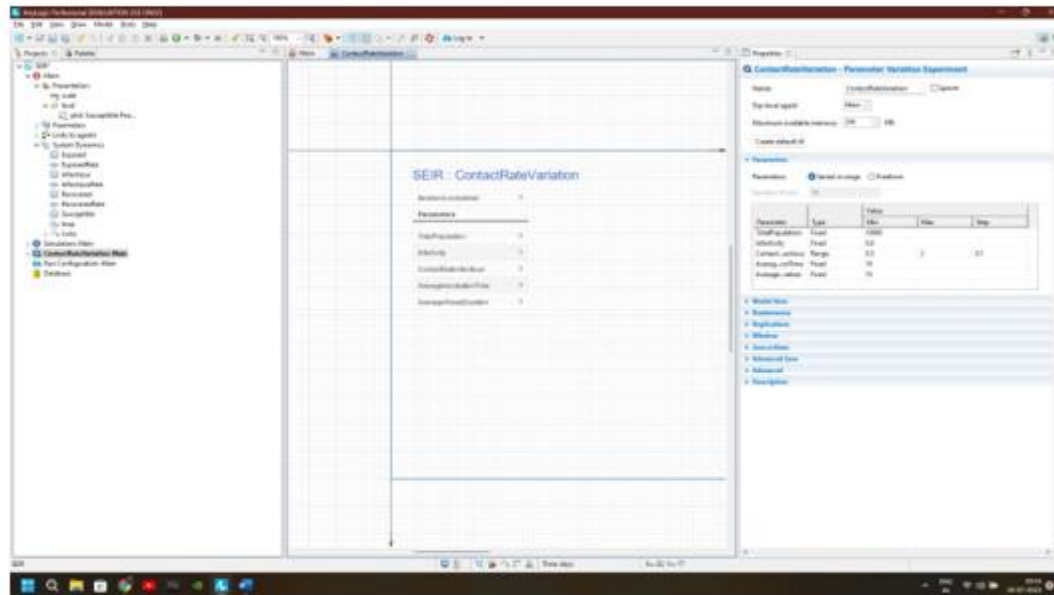
ContactRateVariation - Parameter Variation Experiment

Name: ☐ Ignore

Top-level agent:

Maximum available memory: Mb

Clears experiment's presentation and creates new interface according to its properties



InfectiousDS - Data Set

Name: ☒ Show name ☐ Ignore

Visible: ☒ yes

☒ Use time as horizontal axis value

Horizontal axis value:

Vertical axis value:

Keep up to latest samples

☒ Update data automatically

☐ Do not update data automatically

☒ Use model time ☐ Use calendar dates

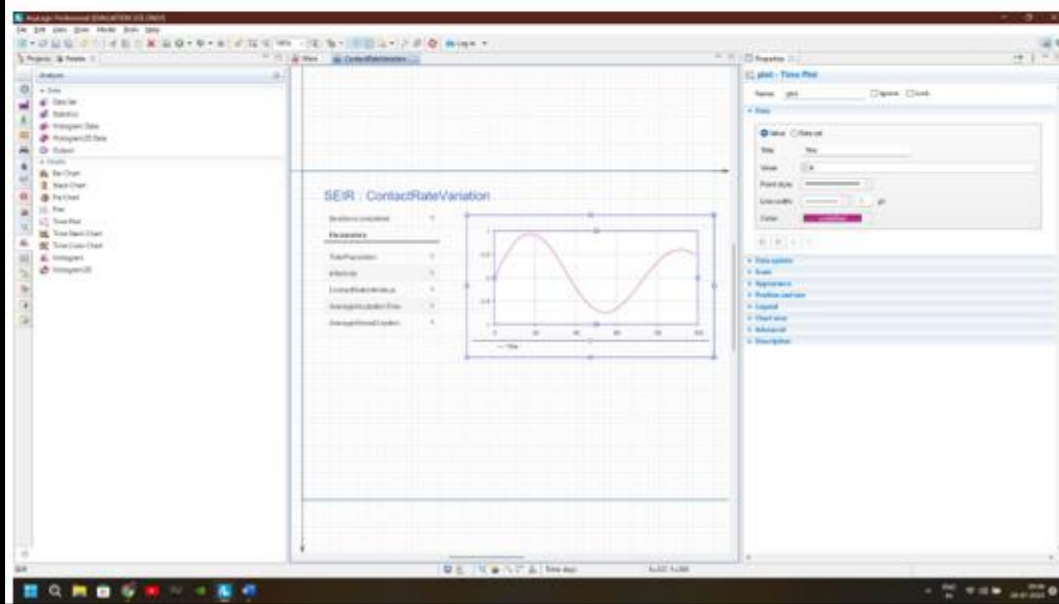
First update time:

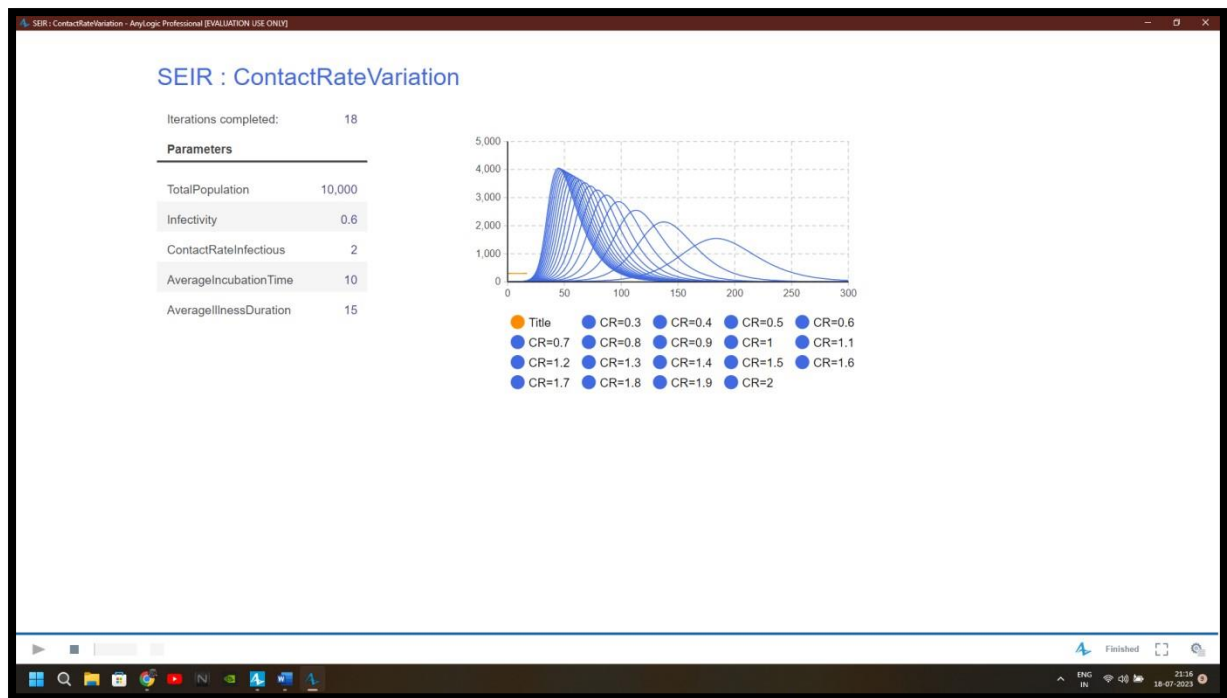
Update date:

Recurrence time:

☒ Log to database

[Turn on model execution logging](#)





Conclusion:

In this practical, we successfully designed and developed a System Dynamics model using AnyLogic to simulate the spread of a contagious disease based on the SEIR model. By creating a stock-flow diagram, adding plots to visualize dynamics, implementing parameter variation, and calibrating the model, we gained insights into the dynamics of disease spread and explored the effects of different parameters on the epidemic trajectory. This model can be further refined and used to inform public health decision-making and interventions aimed at controlling infectious diseases.