

Project Plovdiv 3.1 user manual

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1 General

Project Plovdiv is based on two separate pieces of software. The first one was developed under the York Transit Bursary Scheme under the supervision of Dr Daniel Franks, and the second at the University of Sussex under Dr Istvan Kiss. It is not intended for detailed modelling or efficient calculations but rather as a supplementary course resource. Our main goals throughout the development process were simplicity and ease of use as well as mathematical correctness.

Plovdiv is hosted on Github and is distributed under a three-clause BSD license. Please submit bug reports and feature requests to the our Github issue tracker¹. We welcome contributions from the community.

2 User interface

2.1 Graph visualisation

The overall appearance of the application is shown in Figure 1. The window is divided into two areas. A visual representation of the currently loaded graph is on the left and a number of information panels are on the right. Each of these is discussed separately below.

TODO

- vertices and edges, colour coding of infection state and selection
- vertex icon types, icon autodetermination, changing types
- changing background image and colour

2.2 Mouse modes

The mouse is can be used to control how the graph is visualised. It can operate in four different modes:

1. Selection mode allows the user to select vertices and change their characteristics (see Vertex data).

¹<https://github.com/mbatchkarov/Plovdiv/issues>

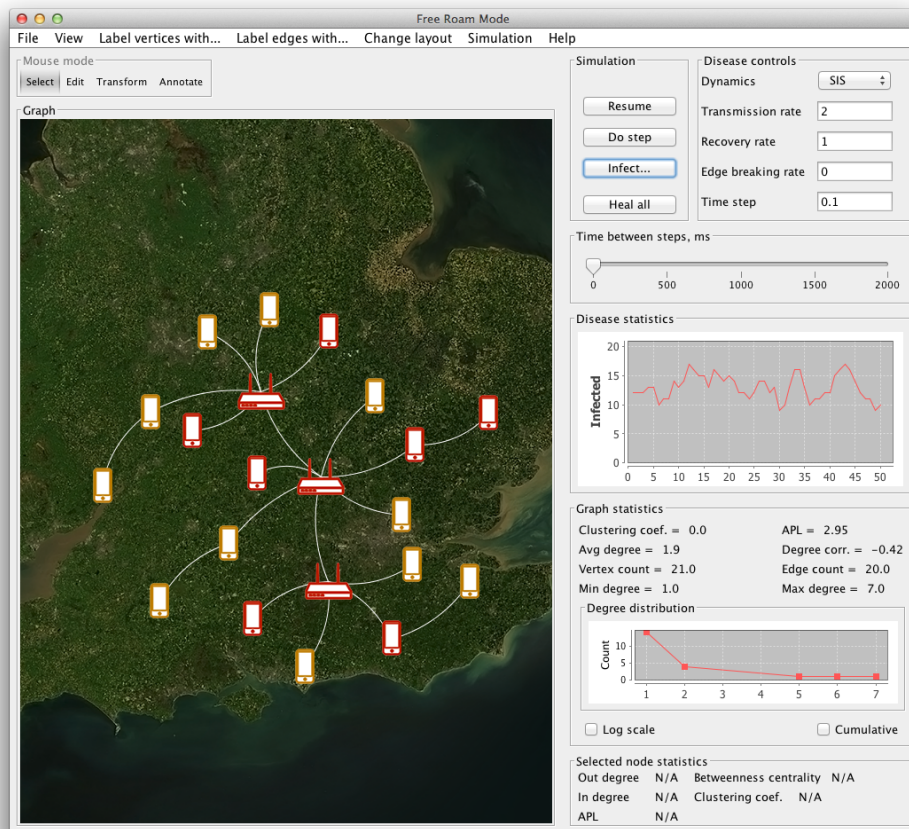


Figure 1: Overall appearance

- (a) Left- clicking a vertex will select it
 - (b) Left- clicking the background will deselect all vertices
 - (c) Ctrl + left click over a vertex will select it and move it the the centre of the screen
 - (d) Shift + left click will select multiple vertices Click and drag will draw a selection rectangle- when the mouse button is released, all vertices inside the rectangle are selected. Selection by dragging is also supported.
2. Editing mode: allows the user to create vertices and edges
- (a) Left clicking anywhere on the screen will create a new vertex at that location
 - (b) Dragging a line from a vertex to another will connect them with an undirected edge, unless they are already connected or you are trying to connect a vertex to itself
 - (c) Holding Shift while dragging will create a directed edge
3. Transforming mode: changes the way the graph is drawn (also see Layouts)
- (a) Dragging will move the portion of the graph that is displayed on screen
 - (b) Ctrl + drag will skew the graph
 - (c) Shift + drag will rotate the graph
4. Annotation mode: this mode allows the user to draw annotations on the screen, for example for teaching or demonstration purposes. In this mode an additional tool bar will appear at the bottom of the screen, allowing control over the way the annotations are created
- (a) Dragging will create a new annotation with the selected shape, colour and position relative to the other elements in the graph
 - (b) Shift + left click over an annotation will delete it
 - (c) Right click anywhere on the screen will let the user enter some text, which will be drawn on the scree. Please note: clicking the right mouse button in annotation mode overrides the default behaviour of the right mouse button (see below)

The additional panel that appears when the mouse is in Annotation mode has four adjustable parameters:

1. Shape of the annotation (rectangle by default)
2. Colour of the annotation

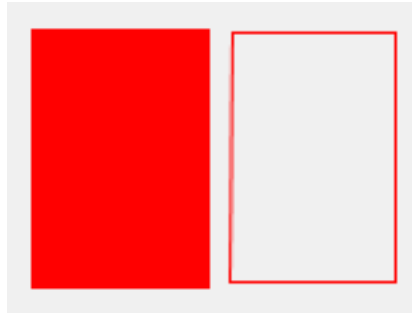


Figure 2: Annotations with and without fill

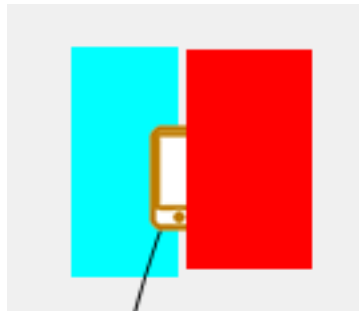


Figure 3: Annotations in lower (blue) and upper (red) position

3. Fill. If this button is not toggled, an annotation will appear as a coloured frame. If it is toggled, the annotation area will be filled with the chosen colour. See Figure 2.
4. Z-order (lower or upper). When lower is selected, the annotation appear behind the graph. When upper is selected, the annotation appear in front of the graph. See Figure 3.

The following actions work in all mouse modes:

1. Scrolling the mouse wheel up will zoom out and scrolling down will zoom in
2. Right clicking a non- selected vertex will allow you to remove it or modify its characteristics (see Vertex data)
3. Right clicking a selected vertex will pop up a context- sensitive menu, which allows you to create edges as well as to modify the vertex characteristics (see Vertex data)

2.3 Information panel

A close-up of the information panel normally located on the right hand side of the application window is shown in Figure 4. This consists of six parts, labelled A to F in the figure. Each of these is described separately below.

2.3.1 A: Simulation controls

This toolbar control the simulation of disease transmission. When the application starts, a simulation is initialised with default settings, and is paused. The **Do step** (keyboard shortcut **S**) button will perform a single step of the simulation. The **Resume** (**R**) button will start the simulation and will then change to **Pause**. The **Pause** button can be used to temporarily pause the simulation—it can be resumed by clicking **Resume**. The **Heal all** (**H**) resets the simulation to its initial state and heals all vertices.

The **Infect** (**I**) button brings up the Infect window (Figure 5), which is used to randomly infect a number of vertices. You can either specify a proportion of the vertices that should be infected by using the top slider, or enter a number in the bottom input field.

2.3.2 B: Infection parameters

This panel controls various parameters of the infection, such as its type (SI, SIS, SIR), transmissions rate, recovery rate (where applicable), etc. Input validation is performed on the fly and the text will change to red if invalid data is entered (e.g. string instead of a number).

2.3.3 C: Simulation speed

When the simulation is running, this slider controls how much time should pass before the next simulation step. Use the left and right arrow keys to adjust this value.

2.3.4 D: Infection statistics

Displays a graph of the number of infected vertices over the past 100 time steps. This updates dynamically as the simulation runs.

2.3.5 E: Graph statistics

Displays statistics for the whole graph. These include average clustering coefficient, the average, min and max degree, average weighted path length, etc. There is also a chart of the degree distribution of the graph, which can be switched to cumulative degree distribution. The y-axis of the chart can be switched to log scale.

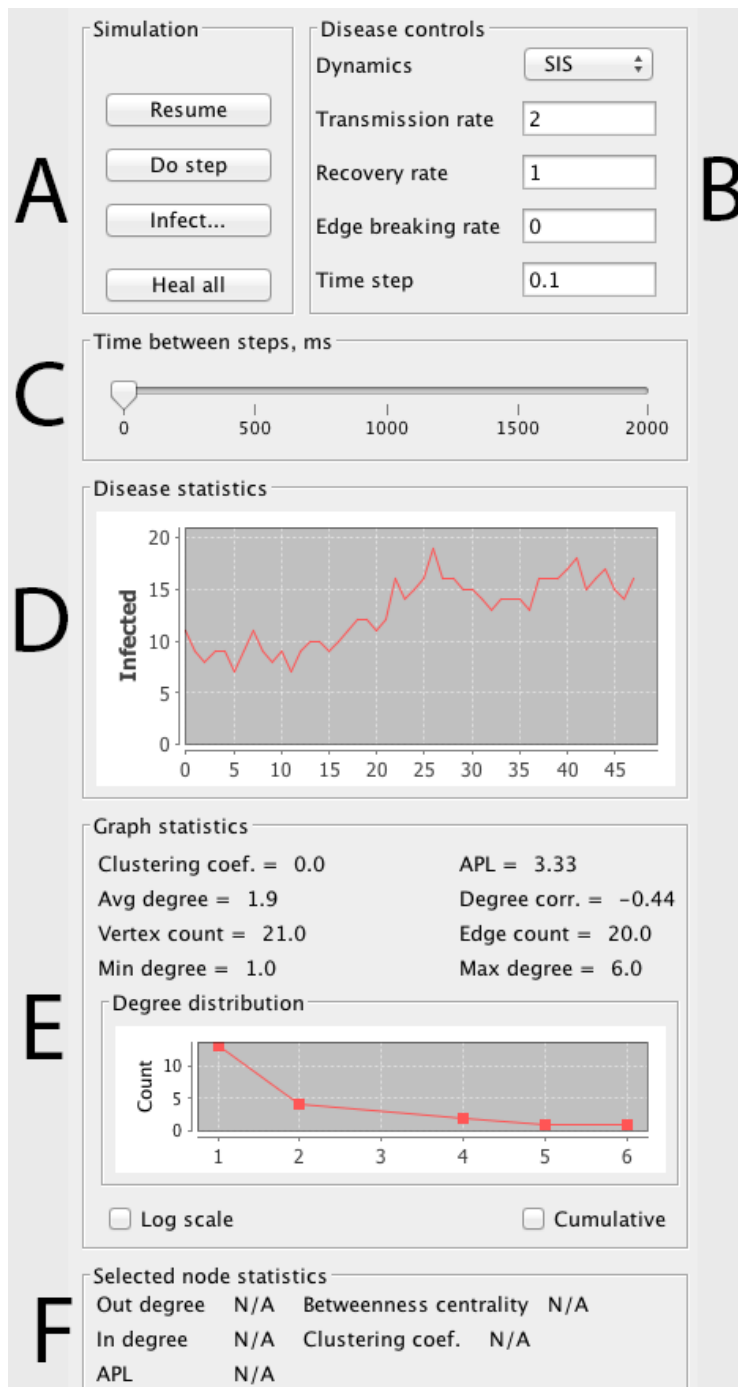


Figure 4: The information panel and its components

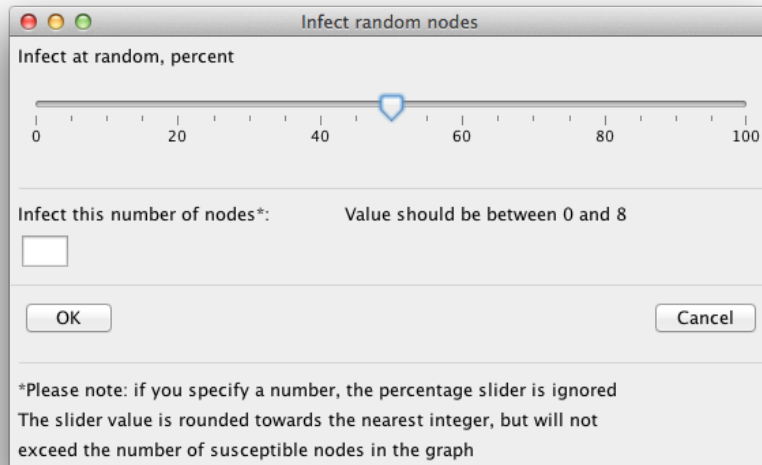


Figure 5: The information panel and its components

2.3.6 F: Selected node statistics

This tool bar displays information about the selected node, such as in and out degree, betweenness centrality, clustering coefficient and average path length. If you select a number of vertices other than one, all fields will read **N/A**.

3 Using the menu options

3.1 The File menu

Contains the following commands:

New Creates a new graph with no vertices and no edges. Keyboard shortcut is **CTRL+N** (or **CMD+N** on OS X)

Save Saves the graph to a file. This creates two files, one with a .graph extension and one with a .layout. The first one contains a description of the graph in Pajek plain text format. Information about the state of each vertex (infected, susceptible or resistant) is also encoded in this file. This is editable in any text editor. The second file is binary (not editable) and describes where each vertex of the graph should be positioned on screen. The two files should be thought of as one for all intents and purposes. Keyboard shortcut is **CTRL+S**.

Load Loads a .graph file that has been previously saved. If a .layout file is present, it will be loaded and all vertices will be restored to their previous position. Otherwise the graph will be laid out using one of the built-in algorithms. Keyboard shortcut is **CTRL+O**.

Generate Generates a new graph. Keyboard shortcut is **CTRL+G**. The program currently employs the following algorithms:

1. Rectangular lattice: a lattice in which each vertex has four local connections
2. Hexagonal lattice: a lattice with six local connections per vertex
3. Scale-free generator: Starts with a single vertex and creates the specified number of vertices, attaching them to existing ones based on the principle of preferential attachment.
4. Small world generator- starts with a circular lattice of the specified size, connecting each vertex to its k closest neighbours on each side. Then traverses all edges, and rewires some of them with probability p .
5. Random, with a pre-defined number of vertices and edges.

Note on performance: depending on your machine's specification, working with large graphs may be very slow. For demonstration purposes it is recommended that you do not generate a graph with more than **100** vertices.

An example of each kind of graph Plovdin can generate is shown in Figure 6.

3.2 The View menu

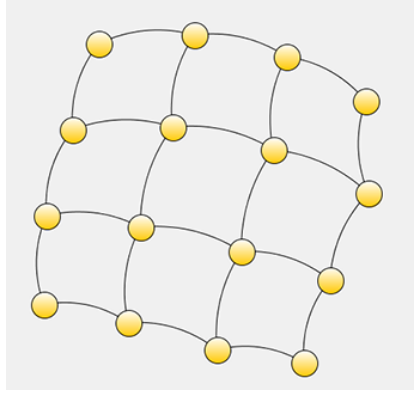
This menu currently contains a single command. To save the what is currently on the screen to a jpg file, click View> Save to .jpg.

3.3 The Label edges/vertices menu

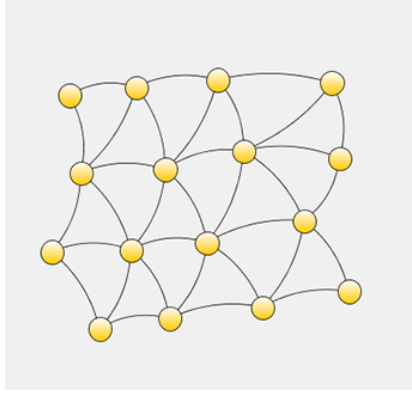
By default edges and vertices are not labelled. Vertex features include degree, local clustering coefficient, betweenness centrality, distance from the currently selected vertex. Edge features include weight and centrality.

3.4 Layouts: The Change Layout menu

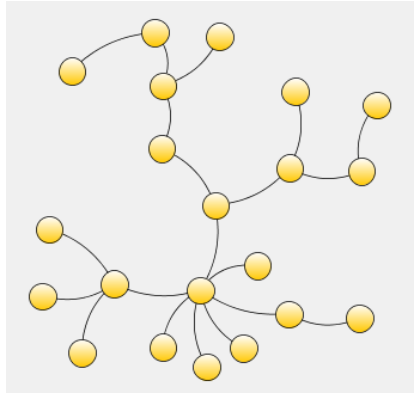
If the graph looks cluttered or unordered, try changing the layout. Different algorithms will produce different results, and sometimes dynamic algorithms, when applied on the same graph more than once, can produce significantly different results. Some algorithms will work better than others on certain types of graphs. This is still an open research area, so there is no optimal layout- try which one looks best for you.



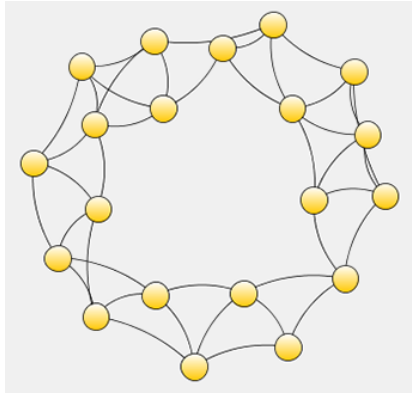
(a) Rectangular lattice



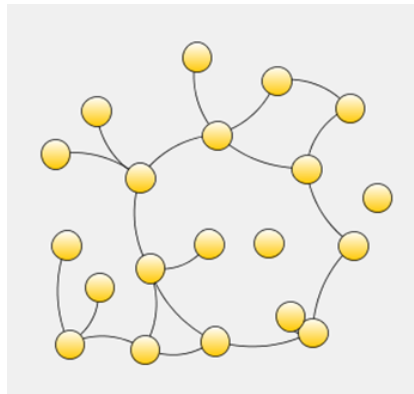
(b) Hexagonal lattice



(c) Scale free



(d) Small world



(e) Random

Figure 6: Example generated graphs

3.5 The Simulation menu

This menu provides another way of accessing the buttons in the Simulation panel (A in Figure 4).

3.6 The Help menu

The Help menu contains a copy of the mouse modes documentation (Section 2.2) as well as information about the version of the software you are running.