Network Science Homework 2

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2025-02-09

Assignment Description: Run the random graph model, vary the connection probability p, examine the impact of p on the size of the largest component and the number of components.

Generate Random Graphs

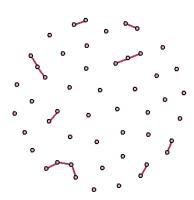
The following code block generates random graphs using the function $sample_gnp()$, which creates random graphs according to the G(n,p) Erdős-Rényi model. I will use this function to create several networks, varying the parameter p which denotes probability for drawing an edge between two arbitrary vertices.

```
set.seed(2025)

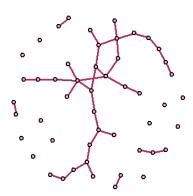
plot_random_graph = function(p, n=50){
   sample_gnp(n, p) |>
      plot(vertex.label = NA, vertex.size = 5, vertex.color = "lightpink", edge.width = 1.5, edge.
color = "maroon")
}
```

Here is a random graph with n=50 and p=0.01.

```
plot_random_graph(0.01)
```

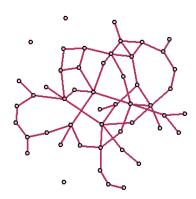


plot_random_graph(0.03)



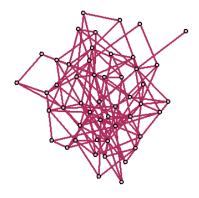
Here is a random graph with n=50 and p=0.05.

plot_random_graph(0.05)



Here is a random graph with n=50 and p=0.1.

plot_random_graph(0.1)



Based on these initial visualizations, I would hypothesize that as the value of p increases, the size of the largest component in a network increases and the number of components decreases. Let's check this out on a larger scale.

Varying p

The code block below generates a dataframe containing random graphs with 50 nodes each and values of p that vary from 0 to 1. For each graph, the dataframe contains the number of components and size of its largest component.

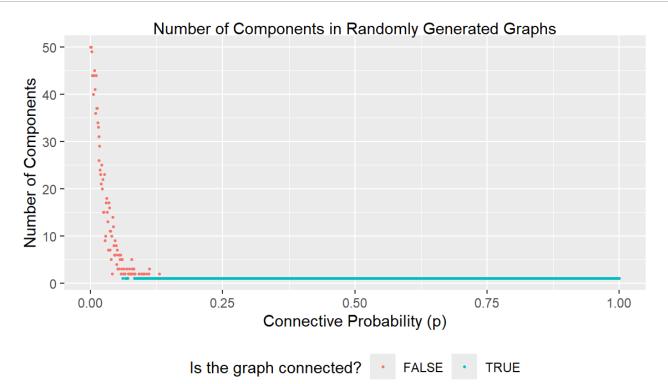
```
# Setting network size at 50 nodes
n = 50

# Generating dataframe of graphs with varying values of p
gnp_df = tibble(
    p = seq(0, 1, 0.001), # varying values of p
    graph = map(p, ~sample_gnp(n, .x)), # generate random graph for each p
    connected = map(graph, is_connected),
    comps = map(graph, decompose),
    n_comp = map(comps, length), # get number of components
    comp_sizes = map(comps, ~sapply(.x, vcount)),
    largest_comp = map(comp_sizes, max) # get size of largest component
) |>
    unnest(c(connected, n_comp, largest_comp)) |>
    select(-c(comps, comp_sizes))
gnp_df |> head(10)
```

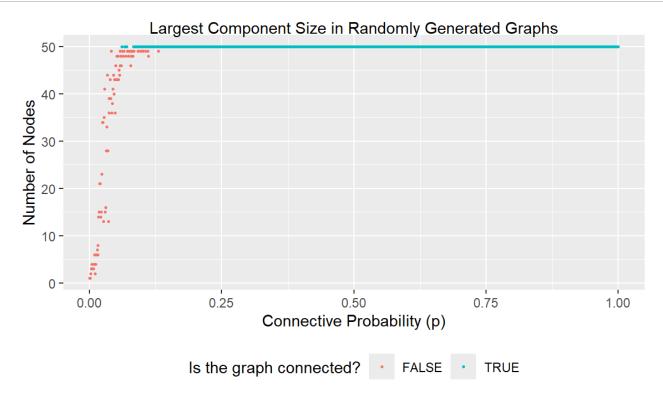
```
## # A tibble: 10 × 5
##
          p graph
                      connected n_comp largest_comp
##
      <dbl> <list>
                      <lgl>
                                 <int>
                                               <dbl>
##
            <igraph> FALSE
                                    50
                                                   1
    2 0.001 <igraph> FALSE
                                    50
                                                   1
##
    3 0.002 <igraph> FALSE
                                    49
                                                   2
##
                                                   3
   4 0.003 <igraph> FALSE
##
                                    44
   5 0.004 <igraph> FALSE
                                    44
                                                   4
##
   6 0.005 <igraph> FALSE
                                    40
                                                   3
##
   7 0.006 <igraph> FALSE
                                    44
                                                   3
##
                                                   3
   8 0.007 <igraph> FALSE
                                    45
   9 0.008 <igraph> FALSE
                                    41
                                                   4
## 10 0.009 <igraph> FALSE
                                     36
                                                   6
```

Now let's visualize how the number of components and size of largest component change with the value of p.

```
gnp_df |>
  ggplot(aes(x = p, y = n_comp, colour = connected)) +
  geom_point(size = 0.5) +
  labs(
    title = "Number of Components in Randomly Generated Graphs",
    x = "Connective Probability (p)",
    y = "Number of Components",
    color = "Is the graph connected?"
)
```

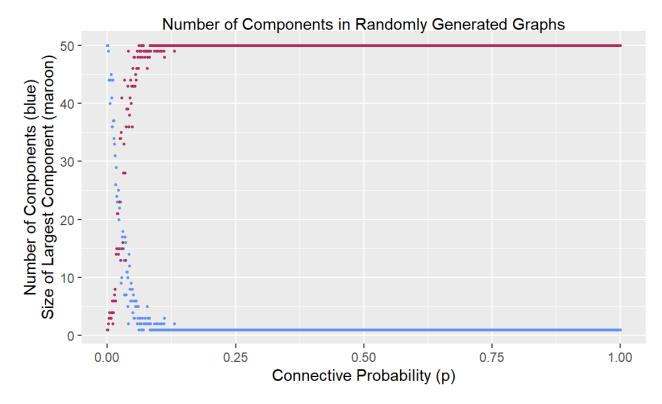


```
gnp_df |>
  ggplot(aes(x = p, y = largest_comp, colour = connected)) +
  geom_point(size = 0.5) +
  labs(
    title = "Largest Component Size in Randomly Generated Graphs",
    x = "Connective Probability (p)",
    y = "Number of Nodes",
    color = "Is the graph connected?"
)
```



Now, just for fun, let's plot both the number of components and size of largest component on the same axes.

```
gnp_df |>
    ggplot(aes(x = p, y = n_comp)) +
    geom_point(size = 0.5, col = "cornflowerblue") +
    geom_point(aes(x = p, y = largest_comp), data = gnp_df, size = 0.5, col = "maroon") +
    labs(
        title = "Number of Components in Randomly Generated Graphs",
        x = "Connective Probability (p)",
        y = "Number of Components (blue)\nSize of Largest Component (maroon)"
    )
```



It seems that increasing the value of p is indeed linked to an increase in the largest component's size and decrease in the number of components of a randomly generated graph.