- 1. Write a program the draws a *composite* Bezier curve from a list of control points. Let the input of your program be a  $n \times 8$  matrix, where each row consists of the coordinates of the four control points of each piece of the curve. Include the code with your project.
  - (a) Use your code to recreate the letter 'R' pictured in Figure 1.13 on p.68.
  - (b) Find appropriate control points to recreate the following heart shape using a composite Bezier curve. Include the list of control points and the curve created by your program.



2. Consider the points

- (a) (Linear Regression) Use the method of least squares to find the line y = mx + b that best fits the data.
- (b) (Cubic Regression) Use the method of least squares to find the cubic polynomial  $y = ax^3 + bx^2 + cx + d$  that best fits the data.
- (c) (Exponential Regression) Use the method of least squares to find the exponential function  $y = Ce^{kx}$  that best fits the data.
- (d) Plot all three functions on the same graph along with the original points.
- (e) Which function do you think best models the given data?
- 3. Apply the SIR model to the spread of Covid-19 in Greene County, Missouri. (https://www.springfieldmo.gov/5068/Coronavirus)

$$\frac{dS}{dt} = -aSI$$

$$\frac{dI}{dt} = aSI - bI$$

$$\frac{dr}{dt} = bI$$

- (a) Use the data found in 'coviddata.csv' to set up an overdetermined system of equations involving the infection rate a and the recovery rate b, and then use the least squares method to determine the best estimate of a and b. (You will need to find a way to approximate the derivatives.)
- (b) Use your values of a and b to model the spread of the infection. Plot the functions I(t) and R(t) from your model on the same graph as the original data for comparison.
- (c) Make a plot showing the long term behavior of I(t) and R(t).
- (d) Estimate the time when I(t) will reach its maximum value. What is that value?
- (e) Approximately what percentage of the total population does your model predict will have been infected in the long run?
- 4. The file 'wine.csv' contains data about alcohol content and color intensity for 178 different wines of 3 different types. Use k-means clustering with k=3 to see how well you can classify these wines into their types based only on their alcohol content and color intensity. Make a scatter plot of the data that is color coded by cluster. What percentage of wines are clustered correctly? (Data from: http://archive.ics.uci.edu/ml/datasets/Wine)