

Figure 4.1 Parallel experimental and computational processes.



Figure 4.2 Plot of population size in our fishing pond model with years = 15.

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Figure 4.3 Plot for Exercise 4.2.1.

Courtesy of CRC Press/Taylor & Francis Group



Figure 4.4 Plot for Exercise 4.2.2.



Figure 4.5 The plot produced by calling pond (15, 12000, 1500, 0.01).



Figure 4.6 Plot of carbon-14 decay generated with decayC14(100, 20000, 0.01).

```
import matplotlib.pyplot as pyplot
def SIR(population, days, dt):
    """Simulates the SIR model of infectious disease and
       plots the population sizes over time.
   Parameters:
        population: the population size
        days: number of days to simulate
        dt: the value of "Delta t" in the simulation
            (fraction of a day)
   Return value: None
    .....
    susceptible = population -1 # susceptible count = S(t)
    infected = 1.0
                                  # infected count = I(t)
   recovered = 0.0
                                  \# recovered count = R(t)
   recRate = 0.25
                                  # recovery rate r
    infRate = 0.0004
                                  # infection rate d
   SList = [susceptible]
    IList = [infected]
   RList = [recovered]
   timeList = [0]
   # Loop using the difference equations goes here.
   pyplot.plot(timeList, SList, label = 'Susceptible')
   pyplot.plot(timeList, IList, label = 'Infected')
    pyplot.plot(timeList, RList, label = 'Recovered')
   pyplot.legend(loc = 'center right')
    pyplot.xlabel('Days')
    pyplot.ylabel('Individuals')
   pyplot.show()
```

Figure 4.7 Output from the SIR model with 2,200 individuals over 30 days with $\Delta t = 0.01$.



Figure 4.8 Harmonic series approximation of the natural logarithm (ln).



Figure 4.9 The Leibniz series converging to π .



Figure 4.10 An illustration of linear, quadratic, and exponential growth. The curves are generated by accumulators adding 6, the index variable, and 1.08 times the accumulator, respectively, in each iteration.