Half Title Page

Title Page

LOC Page

Vince: to Riggins

Geraint: also, to Riggins

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Foreword

This is the foreword

Preface

This is the preface.

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______ Getting Started

Introduction

HANK you for starting to read this book. This book aims to bring together two fascinating topics:

- Problems that can be solved using mathematics;
- Software that is free to use and change.

What we mean by both of those things will become clear through reading this chapter and the rest of the book.

1.1 WHO IS THIS BOOK FOR?

Anyone who is interested in using mathematics and computers to solve problems will hopefully find this book helpful.

If you are a student of a mathematical discipline, a graduate student of a subject like operational research, a hobbyist who enjoys solving the travelling salesman problem or even if you get paid to do this stuff: this book is for you. We will introduce you to the world of open source software that allows you to do all these things freely.

If you are a student learning to write code, a graduate student using databases for their research, an enthusiast who programmes applications to help coordinate the neighbourhood watch, or even if you get paid to write software: this book is for you. We will introduce you to a world of problems that can be solved using your skill sets.

It would be helpful for the reader of this book to:

- Have access to a computer and be able to connect to the internet (at least once) to be able to download the relevant software.
- Be prepared to read some mathematics. Technically you do not need to understand the specific mathematics to be able to use the tools in this book. The topics covered use some algebra, calculus and probability.

1.2 WHAT DO WE MEAN BY APPLIED MATHEMATICS?

We consider this book to be a book on applied mathematics. This is not however a universal term, for some applied mathematics is the study of mechanics and involves

modelling projectiles being fired out of canons. We will use the term a bit more freely here and mean any type of real world problem that can be tackled using mathematical tools. This is sometimes referred to as operational research, operations research, mathematical modelling or indeed just mathematics.

One of the authors, Vince, used mathematics to plan the sitting plan at his wedding. Using a particular area of mathematics call graph theory he was able to ensure that everyone sat next to someone they liked and/or knew.

The other author, Geraint, used mathematics to find the best team of Pokemon. Using an area of mathematics call linear programming which is based on linear algebra he was able to find the best makeup of pokemon.

Here, applied mathematics is the type of mathematics that helps us answer questions that the real world asks.

1.3 WHAT IS OPEN SOURCE SOFTWARE

Strictly speaking open source software is software with source code that anyone can read, modify and improve. In practice this means that you do not need to pay to use it which is often one of the first attractions. This financial aspect can also be one of the reasons that someone will not use a particular piece of software due to a confusion between cost and value: if something is free is it really going to be any good?

In practice open source software is used all of the world and powers some of the most important infrastructure around. For example, one should never use any cryptographic software that is not open source: if you cannot open up and read things than you should not trust it (this is indeed why most cryptographic systems used are open source).

Today, open source software is a lot more than a licensing agreement: it is a community of practice. Bugs are fixed faster, research is implemented immediately and knowledge is spread more widely thanks to open source software. Bugs are fixed faster because anyone can read and inspect the source code. Most open source software projects also have a clear mechanisms for communicating with the developers and even reviewing and accepting code contributions from the general public. Research is implemented immediately because when new algorithms are discovered they are often added directly to the software by the researchers who found them. This all contributes to the spread of knowledge: open source software is the modern should of giants that we all stand on.

Open source software is software that, like scientific knowledge is not restricted in its use.

1.4 HOW TO GET THE MOST OUT OF THIS BOOK

The book itself is open source. You can find the source files for this book online at github.com/drvinceknight/ampwoss. There will will also find a number of *Jupyter notebooks* and *R markdown files* that include code snippets that let you follow along.

We feel that you can choose to read the book from cover to cover, writing out

the code examples as you go; or it could also be used as a reference text when faced with particular problem and wanting to know where to start.

The book is made up of 10 chapters that are paired in two 4 parts. Each part corresponds to a particular area of mathematics, for example "Emergent Behaviour". Two chapters are paired together for each chapter, usually these two chapters correspond to the same area of mathematics but from a slightly different scale that correspond to different ways of tackling the problem.

Every chapter has the following structure:

- 1. Introduction a brief overview of a given problem type. Here we will describe the problem at hand in general terms.
- 2. An Example problem. This will provide a tangible example problem that offers the reader some intuition for the rest of the discussion.
- 3. Solving with Python. We will describe the mathematical tools available to us in a programming language called Python to solve the problem.
- 4. Solving with R. Here we will do the same with the R programming language.
- 5. Brief theoretic background with pointers to reference texts. Some readers might like to delve in to the mathematics of the problem a bit further, we will include those details here.
- 6. Examples of research using these methods. Finally, some readers might even be interested in finding out a bit more of what mathematicians are doing on these problems. Often this will include some descriptions of the problem considered but perhaps at a much larger scale than the one presented in the example.

For a given reader, not all sections of a chapter will be of interest. Perhaps a reader is only interested in R and finding out more about the research. Please do take from the book what you find useful.

		_

Probabilistic Modelling

		_

Markov Chains

Many real world situations have some level of unpredictability through randomness: the flip of a coin, the number of orders of coffee in a shop, the winning numbers of the lottery. However, mathematics can in fact let us make predictions about what we expect to happen. One tool used to understand randomness is Markov chains, an area of mathematics sitting at the intersection of probability and linear algebra.

2.1 PROBLEM

Consider a barber shop. The shop owners have noticed that customers will not wait if there is no room in their waiting room and will choose to take their business elsewhere. The Barber shop would like to make an investment so as to avoid this situation. They know the following information:

- They currently have 2 barber chairs (and 2 barbers).
- They have waiting room for 4 people.
- They usually have 10 customers arrive per hour.
- Each Barber takes about 15 minutes to serve a customer so they can serve 4 customers an hour.

This is represented diagrammatically in Figure 2.1.

They are planning on reconfiguring space to either have 2 extra waiting chairs or another barber's chair and barber.

The mathematical tool used to model this situation is a Markov chain.

2.2 THEORY

A Markov chain is a model of a sequence of random events that is defined by a collection of **states** and rules that define how to move between these states.

For example, in the barber shop a single number is sufficient to describe the status of the shop. If that number is 1 this implies that 1 customer is currently having their

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Figure 2.1 Diagrammatic representation of the barber shop as a queuing system.

hair cut. If that number is 5 this implies that 2 customers are being served and 3 are waiting. The entire state space is, in this case a finite set of integers from 0 to 6. If the system is full (all barbers and waiting room occupied) then we are in state 6 and if there is no one at the shop then we are in state 0. This is denoted mathematically as:

$$S = \{0, 1, 2, 3, 4, 5, 6\} \tag{2.1}$$

As customers arrive and leave the system goes between states as shown in Figure 2.2.

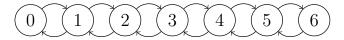


Figure 2.2 Diagrammatic representation of the state space

The rules that govern how to move between these states can be defined in two ways:

- Using probabilities of changing state (or not) in a well defined time period. This is called a discrete Markov chain.
- Using rates of change from one state to another. This is called a continuous time Markov chain.

For our barber shop we will consider it as a continuous Markov chain as shown in Figure 2.3

Note that a Markov chain assumes the rates follow an exponential distribution. One interesting property of this distribution is that it is considered memoryless which means that if a customer has been having their hair cut for 5 minutes this does not change the rate at which their service ends. This distribution is quite common in the real world and therefore a common assumption.

These states and rates can be represented mathematically using a transition matrix Q where Q_{ij} represents the rate of going from state i to state j. In this case we have:

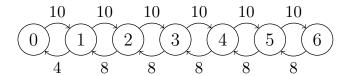


Figure 2.3 Diagrammatic representation of the state space and the transition rates

$$Q = \begin{pmatrix} -10 & 10 & 0 & 0 & 0 & 0 & 0 \\ 4 & -14 & 10 & 0 & 0 & 0 & 0 \\ 0 & 8 & -18 & 10 & 0 & 0 & 0 \\ 0 & 0 & 8 & -18 & 10 & 0 & 0 \\ 0 & 0 & 0 & 8 & -18 & 10 & 0 \\ 0 & 0 & 0 & 0 & 8 & -18 & 10 \\ 0 & 0 & 0 & 0 & 0 & 8 & -8 \end{pmatrix}$$
 (2.2)

You will see that Q_{ii} are negative and ensure the rows of Q sum to 0. This gives the total rate of change leaving state i.

We can use Q to understand the probability of being in a given state after t time unis. This is can be represented mathematically using a matrix P_t where $(P_t)_{ij}$ is the probability of being in state j after t time units having started in state i. We can use Q to calculate P_t using the matrix exponential:

$$P_t = e^{Qt} (2.3)$$

What is also useful is understanding the long run behaviour of the system. This allows us to answer questions such as "what state are we most likely to be in on average?" or "what is the probability of being in the last state on average?".

This long run probability distribution over the state can be represented using a vector π where π_i represents the probability of being in state i. This vector is in fact the solution to the following matrix equation:

$$\pi Q = 0 \tag{2.4}$$

In the upcoming sections we will demonstrate all of the above concepts.

2.3 SOLVING WITH PYTHON

The first step we will take is to write a function to obtain the transition rates between two given states:

```
Python input
    def get_transition_rate(
         in_state, out_state, waiting_room=4, num_barbers=2,
    ):
         """Return the transition rate for two given states.
         Args:
             in_state: an integer
             out_state: an integer
             waiting_room: an integer (default: 4)
9
             num_barbers: an integer (default: 2)
10
11
         Returns:
12
             A real.
13
14
         arrival_rate = 10
15
         service_rate = 4
16
17
         capacity = waiting_room + num_barbers
18
         delta = out_state - in_state
19
20
         if delta == 1 and in_state < capacity:</pre>
21
             return arrival rate
22
23
         if delta == -1:
^{24}
             return min(in_state, num_barbers) * service_rate
25
26
         return 0
27
```

Next, we write a function that creates an entire transition rate matrix Q for a given problem. We will use the numpy to handle all the linear algebra and the itertools library for some iterations:

```
import itertools
28
     import numpy as np
29
30
31
     def get_transition_rate_matrix(waiting_room=4, num_barbers=2):
32
         """Return the transition matrix Q.
33
34
         Args:
35
             waiting_room: an integer (default: 4)
36
             num_barbers: an integer (default: 2)
37
38
         Returns:
39
             A matrix.
40
41
         capacity = waiting_room + num_barbers
42
         state_pairs = itertools.product(
43
             range(capacity + 1), repeat=2
44
         )
45
46
         flat_transition_rates = [
47
             get_transition_rate(
48
                  in_state=in_state,
49
                  out_state=out_state,
50
                 waiting room=waiting room,
51
                 num_barbers=num_barbers,
52
53
             for in_state, out_state in state_pairs
54
55
         transition_rates = np.reshape(
56
             flat_transition_rates, (capacity + 1, capacity + 1)
57
58
         np.fill_diagonal(
59
             transition_rates, -transition_rates.sum(axis=1)
60
61
62
         return transition_rates
63
```

Using this we can obtain the matrix Q for our default system:

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```
Python input

Q = get_transition_rate_matrix()
print(Q)
```

which gives:

```
Python output
     [[-10
            10
                                    0]
66
         4 - 14
                10
                      0
                                    0]
                          0
67
             8 -18 10
                                    0]
                          0
                               0
68
                                    0]
             0
                 8 -18 10
                               0
69
                      8 -18 10
                                   0]
             0
                  0
70
                                  10
                          8 -18
             0
                  0
                      0
71
                      0
                               8
                                  -8]]
                          0
72
```

We can take the matrix exponential as discussed above. To do this, we need to use the scipy library. To see what would happen after .5 time units we obtain:

```
Python input

import scipy.linalg

print(scipy.linalg.expm(Q * 0.5).round(5))
```

which gives:

```
Python output

[[0.10492 0.21254 0.20377 0.17142 0.13021 0.09564 0.0815 ]
[0.08501 0.18292 0.18666 0.1708 0.14377 0.1189 0.11194]
[0.06521 0.14933 0.16338 0.16478 0.15633 0.14751 0.15346]
[0.04388 0.10931 0.13183 0.15181 0.16777 0.18398 0.21142]
[0.02667 0.07361 0.10005 0.13422 0.17393 0.2189 0.27262]
[0.01567 0.0487 0.07552 0.11775 0.17512 0.24484 0.32239]
[0.01068 0.03668 0.06286 0.10824 0.17448 0.25791 0.34914]]
```

To see what would happen after 500 time units we obtain:

```
Python input

print(scipy.linalg.expm(Q * 500).round(5))
```

which gives:

```
Python output
     [[0.03431 0.08577 0.10722 0.13402 0.16752 0.2094
                                                         0.26176]
84
      [0.03431 0.08577 0.10722 0.13402 0.16752 0.2094
                                                         0.26176]
85
      [0.03431 0.08577 0.10722 0.13402 0.16752 0.2094
                                                         0.26176]
86
      [0.03431 0.08577 0.10722 0.13402 0.16752 0.2094
                                                         0.26176]
87
      [0.03431 0.08577 0.10722 0.13402 0.16752 0.2094
                                                         0.26176]
88
      [0.03431 0.08577 0.10722 0.13402 0.16752 0.2094
                                                         0.26176]
89
      [0.03431 0.08577 0.10722 0.13402 0.16752 0.2094
                                                         0.26176]]
90
```

We see that no matter what state (row) the system is in, after 500 time units the probabilities are all the same. We could in fact stop our analysis here, however our choice of 500 time units was arbitrary and might not be the correct amount for all possible scenarios, as such we will continue to aim to solve the underlying equation 2.4 directly.

To do this we will solve the underlying system using a numerically efficient algorithm called least squares optimisation (available from the numpy library):

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```
Python input
     def get_steady_state_vector(Q):
91
          """Return the steady state vector of any given continuous
92
         time transition rate matrix.
93
94
         Args:
95
            Q: a transition rate matrix
96
97
         Returns:
98
              A vector
99
100
         state space size, = Q.shape
101
         A = np.vstack((Q.T, np.ones(state_space_size)))
102
         b = np.append(np.zeros(state_space_size), 1)
103
         x, _, _ = np.linalg.lstsq(A, b, rcond=None)
104
         return x
105
```

So if we now see the steady state vector for our default system:

```
Python input

print(get_steady_state_vector(Q).round(5))
```

we get:

```
Python output

[0.03431 0.08577 0.10722 0.13402 0.16752 0.2094 0.26176]
```

We can see that the shop is expected to be empty approximately 3.4% of the time and full 26.2% of the time.

The final function we will write is one that uses all of the above to just return the probability of the shop being full.

```
def get_probability_of_full_shop(
108
          waiting_room=4, num_barbers=2
109
     ):
110
          """Return the probability of the barber shop being full.
111
112
          Args:
113
              waiting_room: an integer (default: 4)
114
              num_barbers: an integer (default: 2)
115
116
          Returns:
117
              A real.
119
          Q = get_transition_rate_matrix(
120
              waiting_room=waiting_room, num_barbers=num_barbers,
121
122
         pi = get_steady_state_vector(Q)
123
         return pi[-1]
124
```

We can now confirm the previous probability calculated probability of the shop being full:

```
Python input

print(round(get_probability_of_full_shop(), 6))
```

which gives:

```
Python output

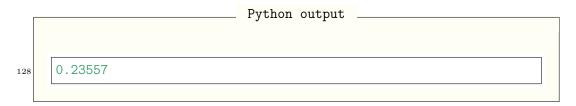
0.261756
```

If we were too have 2 extra space in the waiting room:

```
Python input

print(round(get_probability_of_full_shop(waiting_room=6), 6))
```

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This is a slight improvement however, increasing the number of barbers has a substantial effect:

```
Python input

print(round(get_probability_of_full_shop(num_barbers=3), 6))
```

```
Python output

0.078636
```

2.4 SOLVING WITH R

which gives:

The first step we will take is write a function to obtain the transition rates between two given states:

```
R input
      #' Return the transition rate for two given states.
131
      # '
132
      #' @param in_state an integer
133
      #' @param out_state an integer
134
      #' @param waiting_room an integer (default: 4)
135
      #' @param num_barbers an integer (default: 2)
136
137
      #' @return A real
138
      get_transition_rate <- function(in state,</pre>
139
                                          out_state,
140
                                          waiting_room = 4,
141
                                          num_barbers = 2){
142
        arrival_rate <- 10
143
        service_rate <- 4
144
145
        capacity <- waiting_room + num_barbers</pre>
146
        delta <- out_state - in_state</pre>
147
148
        if (delta == 1) {
149
          if (in state < capacity) {</pre>
150
            return(arrival_rate)
151
          }
152
        }
153
154
        if (delta == -1) {
155
          return(min(in state, num barbers) * service rate)
157
        return(0)
158
159
```

We will not actually use this function but a vectorized version of this:

```
vectorized_get_transition_rate <- Vectorize(
get_transition_rate,
vectorize.args = c("in_state", "out_state")
)
```

This function can now take a vector of inputs for the in_state and out_state variables which will allow us to simplify the following code that creates the matrices:

```
R input
         Return the transition rate matrix Q
164
165
      #' @param waiting_room an integer (default: 4)
166
      #' @param num_barbers an integer (default: 2)
167
168
      #' @return A matrix
169
      get_transition_rate_matrix <- function(waiting_room = 4,</pre>
170
                                                  num_barbers = 2){
171
        max_state <- waiting_room + num_barbers</pre>
172
173
        Q <- outer(0:max_state,</pre>
174
          0:max_state,
175
          vectorized_get_transition_rate,
176
          waiting_room = waiting_room,
177
          num_barbers = num_barbers
178
179
        row sums <- rowSums(Q)</pre>
180
181
        diag(Q) <- -row sums
182
183
      }
184
```

Using this we can obtain the matrix Q for our default system:

```
R input

Q <- get_transition_rate_matrix()
print(Q)
```

which gives:

```
R output
                   [,2] [,3] [,4] [,5]
                                              [,6]
187
       [1,]
              -10
                      10
                              0
                                     0
                                            0
                                                  0
188
       [2,]
                 4
                     -14
                             10
                                     0
                                           0
                                                  0
                                                         0
189
       [3,]
                        8
                            -18
                                   10
                 0
                                           0
                                                  0
                                                         0
190
       [4,]
                              8
                                  -18
                 0
                        0
                                          10
                                                  0
                                                         0
191
       [5,]
                 0
                        0
                              0
                                     8
                                         -18
                                                 10
                                                         0
192
       [6,]
                        0
                              0
                                     0
                                           8
                                                -18
                 0
                                                       10
193
       [7,]
                                           0
                                                       -8
194
                        0
                              0
                                     0
                                                  8
```

One immediate thing we can do with this matrix is take the matrix exponential discussed above. To do this, we need to use an R library call expm.

To be able to make use of the nice %>% "pipe" operator we are also going to load the magrittr library. Now if we wanted to see what would happen after .5 time units we obtain:

```
library(expm, warn.conflicts = FALSE, quietly = TRUE)
library(magrittr, warn.conflicts = FALSE, quietly = TRUE)
print( (Q * .5) %>% expm %>% round(5))
```

which gives:

```
R output
              [,1]
                      [,2]
                               [,3]
                                       [,4]
                                                [,5]
199
     [1,] 0.10492 0.21254 0.20377 0.17142 0.13021 0.09564 0.08150
200
     [2,] 0.08501 0.18292 0.18666 0.17080 0.14377 0.11890 0.11194
201
     [3,] 0.06521 0.14933 0.16338 0.16478 0.15633 0.14751 0.15346
202
     [4,] 0.04388 0.10931 0.13183 0.15181 0.16777 0.18398 0.21142
203
     [5,] 0.02667 0.07361 0.10005 0.13422 0.17393 0.21890 0.27262
204
     [6,] 0.01567 0.04870 0.07552 0.11775 0.17512 0.24484 0.32239
205
     [7,] 0.01068 0.03668 0.06286 0.10824 0.17448 0.25791 0.34914
206
```

After 500 time units we obtain:

```
R input

print( (Q * 500) %>% expm %>% round(5))
```

which gives:

```
R output
              [,1]
                               [,3]
                      [,2]
                                       [,4]
                                                [,5]
                                                       [,6]
                                                                [,7]
208
     [1,] 0.03431 0.08577 0.10722 0.13402 0.16752 0.2094 0.26176
209
     [2,] 0.03431 0.08577 0.10722 0.13402 0.16752 0.2094 0.26176
210
     [3,] 0.03431 0.08577 0.10722 0.13402 0.16752 0.2094 0.26176
211
     [4,] 0.03431 0.08577 0.10722 0.13402 0.16752 0.2094 0.26176
212
     [5,] 0.03431 0.08577 0.10722 0.13402 0.16752 0.2094 0.26176
213
     [6,] 0.03431 0.08577 0.10722 0.13402 0.16752 0.2094 0.26176
214
     [7,] 0.03431 0.08577 0.10722 0.13402 0.16752 0.2094 0.26176
215
```

We see that no matter what state (row) we are in, after 500 time units the probabilities are all the same. We could in fact stop our analysis here, however our choice of 500 time units was arbitrary and might not be the correct amount for all possible scenarios, as such we will continue to aim to solve the underlying equation 2.4 directly.

To be able to do this, we will make use of the versatile pracma package which includes a number of numerical analysis functions for efficient computations.

```
R input
      library(pracma, warn.conflicts = FALSE, quietly = TRUE)
216
217
      #' Return the steady state vector of any given continuous time
218
      #' transition rate matrix
219
220
      #' @param Q a transition rate matrix
221
222
      #' @return A vector
223
      get steady state vector <- function(Q){</pre>
224
        state_space_size <- dim(Q)[1]</pre>
225
        A \leftarrow rbind(t(Q), 1)
226
        b <- c(integer(state_space_size), 1)</pre>
        mldivide(A, b)
228
229
```

This is making use of pracma's mldivide function which chooses the best numerical algorithm to find the solution to a given matrix equation Ax = b.

So if we now see the steady state vector for our default system:

```
R input

print(get_steady_state_vector(Q))
```

we get:

```
R output
                  [,1]
231
      [1,] 0.03430888
232
      [2,] 0.08577220
233
      [3,] 0.10721525
234
      [4,] 0.13401906
235
      [5,] 0.16752383
236
      [6,] 0.20940479
237
      [7,] 0.26175598
238
```

We can see that the shop is expected to be empty approximately 3.4% of the time and full 26.2% of the time.

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The final piece of this puzzle is to create a single function that uses all of the above to just return the probability of the shop being full.

```
__ R input _
      #' Return the probability of the barber shop being full
239
240
      #' @param waiting_room (default: 4)
241
      #' @param num barbers (default: 2)
242
243
      #' @return A real
244
      get_probability_of_full_shop <- function(waiting_room = 4,</pre>
245
                                                   num barbers = 2){
246
        arrival_rate <- 10
247
        service rate <- 4
248
        pi <- get_transition_rate_matrix(</pre>
249
          waiting_room = waiting_room,
250
          num_barbers = num_barbers
251
          ) %>%
252
          get_steady_state_vector()
253
254
        capacity <- waiting room + num barbers</pre>
255
        pi[capacity + 1]
256
257
```

Now we can run this code efficiently with both scenarios:

```
R input

print(get_probability_of_full_shop(waiting_room = 6))
```

which decreases the probability of a full shop to:

```
R output

[1] 0.2355699
```

but adding another barber and chair:

```
R input

print(get_probability_of_full_shop(num_barbers = 3))

gives:

R output

[1] 0.0786359
```

2.5 RESEARCH

TBA

Discrete Event Simulation

OMPLEX situations further compounded by randomness appear throughout our daily lives. For example, data flowing through a computer network, patients being treated at an emergency services, and daily commutes to work. Mathematics can be used to understand these complex situations so as to make predictions which in turn can be used to make improvements. One tool used to do this is to let a computer create a dynamic virtual representation of the scenario in question, the particular type we are going to cover here is called Discrete Event Simulation.

3.1 PROBLEM

Consider the following situation: a bicycle repair shop would like reconfigure their set-up in order to guarantee that all bicycles processed by the repair shop take a maximum of 30 minutes. Their current set-up is as follows:

- Bicycles arrive randomly at the shop at a rate of 15 per hour.
- They wait in line to be seen at an inspection counter, manned by one member of staff who can inspect one bicycle at a time. On average an inspection takes around 3 minutes.
- After inspection it is found that around 20% of bicycles do not need repair, and they are then ready for collection.
- After inspection is is found that around 80% of bicycles go on to be repaired.
 These then wait in line outside the repair workshop, which is manned by two
 members of staff who can each repair one bicycle at a time. On average a repair
 takes around 6 minutes.
- After repair the bicycles are ready for collection.

A diagram of the system is shown in Figure 3.1

We can also assume that there is infinite capacity at the bicycle repair shop for waiting bicycles. The shop will hire and extra member of staff in order to meet their target of a maximum time in the system of 30 minutes. They would like to know if they should work on the inspection counter or in the repair workshop?



Figure 3.1 Diagrammatic representation of the bicycle repair shop as a queuing system.

3.2 THEORY

A number of the events that govern the behaviour of the bicycle shop above are probabilistic. For example the times that bicycles arrive at the shop, the duration of the inspection and repairs, and whether the bicycle would need to go on to be repaired or not. When a number of these probabilistic events are arranged in a complex system such as the bicycle shop, using analytical methods to manipulate these probabilities can become difficult. One method to deal with this is *simulation*.

Consider one probabilistic event, rolling a die. A die has six sides numbered 1 to 6, each side is equally likely to land. Therefore the probability of rolling a 1 is $\frac{1}{6}$, the probability of rolling a 2 is $\frac{1}{6}$, and so on. This means that that if we roll the die a large number of times, we would except $\frac{1}{6}$ of those rolls to be a 1. This is called the law of large numbers.

Now imagine we have an event in which we do not know the analytical probability of it occurring. Consider rolling a weighted die, in this case a die in which the probability of obtaining one number is much greater than the others. How can we estimate the probability of obtaining a 5 on this die?

Rolling the weighted die once does not give us much information. However due to the law of large numbers, we can roll this die a number of times, and find the proportion of those rolls which gave a 5. The more times we roll the die, the closer this proportion approaches the underlying probability of obtaining a 5.

For a complex system such as the bicycle shop, we would like to estimate the proportion of bicycles that take longer than 30 minutes to be processed. As it is a complex system it is difficult to work this out analytically. So, just like the weighted die, we would like to observe this system a number of times and record the overall proportions of bicycles spending longer than 30 minutes in the shop, which will converge to the true underlying proportion. However unlike rolling a weighted die, it it costly to observe this shop over a number of days with identical conditions. In this case it is costly in terms of time, as the repair shop already exists. However some scenarios, for example the scenario where the repair shop hires and additional member of staff, do not yet exist, so observing this this would be costly in terms of money also. We can however build a virtual representation of this complex system on a computer, and observe a virtual day of work much more quickly and much less costly on the computer, similar to a video game.

In order to do this, the computer needs to be able to generate random outcomes of

each of the smaller events that make up the large complex system. Generating random events are essentially doing things to random numbers, that need to be generated.

Computers are deterministic, therefore true randomness is not always possible. They can however generate pseudorandom numbers: sequences of numbers that look like random numbers, but are entirely determined from the previous numbers in the sequence. Most programming languages have methods of doing this.

In order to simulate an event we can again manipulate the law of large numbers. Let $X \sim U(0,1)$, a uniformly pseudorandom variable between 0 and 1. Let D be the outcome of a roll of an unbiased die. Then D can be defined as:

$$D = \begin{cases} 1 & \text{if } 0 \le X < \frac{1}{6} \\ 2 & \text{if } \frac{1}{6} \le X < \frac{2}{6} \\ 3 & \text{if } \frac{2}{6} \le X < \frac{3}{6} \\ 4 & \text{if } \frac{3}{6} \le X < \frac{4}{6} \\ 5 & \text{if } \frac{4}{6} \le X < \frac{5}{6} \\ 6 & \text{if } \frac{5}{6} \le X < 1 \end{cases}$$

$$(3.1)$$

The bicycle repair shop is a system made up of interactions of a number of other simpler random events. This can be thought of as many interactions of random variables, each generated using pseudorandom numbers.

In this case the fundamental random events that need to be generated are:

- the time each bicycle arrives to the repair shop,
- the time each bicycle spends at the inspection counter,
- whether each bicycle needs to go on the the repair workshop,
- the time each those bicycles spends at the repair shop.

As the simulation progresses these events will be generated, and will interact together as described in Section 4.1. The proportion of customers spending longer than 30 minutes in the shop can then be counted. This proportion itself is a random variable, and so just like the weighted die, running this simulation once does not give us much information. But we can run the simulation many times and take an average proportion, to smooth out any variability.

The process outlined above is a particular implementation of Monte Carlo simulation called *discrete event simulation*, which generates pseudorandom numbers and observes their interactions. In practice there are two main approaches to simulating complex probabilistic systems such as this one: the *event scheduling* approach, and *process based* simulation. It just so happens that the main implementations in Python and R use each of these approaches, so you will see both approaches used here.

3.2.1 Event Scheduling Approach

When using the event scheduling approach, we can think of the 'virtual representation' of the system as being the facilities that the bicycles use, shown in Figure 3.1. Then we let entities (the bicycles) interact with these facilities. It is these facilities that determine how the entities behave.

In a simulation that uses an event scheduling approach, a key concept is that events occur that cause further events to occur in the future, either immediately or after a delay, such as after some time in service. In the bicycle shop examples of such events include a bicycle joining a queue, a bicycle beginning service, and a bicycle finishing service. At each event the event list is updated, and the clock then jumps forward to the next event in this updated list.

3.2.2 Process Based Simulation

When using process based simulation, we can think of the 'virtual representation' of the system as being the sequence of actions that each entity (the bicycles) must take, and these sequences of actions might contain delays as a number of entities seize and release a finite amount of resources. It is the sequence of actions that determine how the entities behave.

For the bicycle repair shop an example of one possible sequence of actions would be:

 $arrive \rightarrow seize \ inspection \ counter \rightarrow delay \rightarrow release \ inspection \ counter \rightarrow seize \ repair \ shop \rightarrow delay \rightarrow release \ repair \ shop \rightarrow leave$

The scheduled delays in this sequence of events correspond to the time spend being inspected and the time spend being repaired. Waiting in line for service at these facilities are not included in the sequence of events; that is implicit by the 'seize' and 'release' actions, as an entity will wait for a free resource before seizing one. Therefore in process based simulations, in addition to defining a sequence of events, resource types and their numbers also need to be defined.

3.3 SOLVING WITH PYTHON

In this book we will use the Ciw library in order to conduct discrete event simulation in Python. Ciw uses the event scheduling approach, which means we must define the system's facilities, and then let customers loose to interact with them.

In this case there are two facilities to define: the inspection desk and the repair workshop. Let's order these as so. For each of these we need to define:

- the distribution of times between consecutive bicycles arriving,
- the distribution of times the bicycles spend in service,
- the number of servers available,
- the probability of routing to each of the other facilities after service.

In this case we will assume that the time between consecutive arrivals follow a exponential distribution, and that the service times also follow exponential distributions. These are common assumptions for this sort of queueing system.

In Ciw, these are defined in a Network object, created using the ciw.create network

function. The function below creates a Network object that defines the for a given set of parameters bicycle repair shop:

```
Python input
      import ciw
262
263
264
     def build network object(
265
          num_inspectors=1, num_repairers=2,
266
     ):
267
          """Returns a Network object that defines the repair shop.
268
269
          Args:
270
               num_inspectors: a positive integer (default: 1)
271
              num_repairers: a positive integer (default: 2)
272
273
          Returns:
274
               a Ciw network object
275
          11 11 11
276
          arrival_rate = 15
277
          inspection_rate = 20
278
          repair_rate = 10
279
          prob_need_repair = 0.8
280
          N = ciw.create_network(
281
              arrival_distributions=[
282
                   ciw.dists.Exponential(arrival rate),
283
                   ciw.dists.NoArrivals(),
284
              ],
285
              service distributions=[
286
                   ciw.dists.Exponential(inspection_rate),
287
                   ciw.dists.Exponential(repair_rate),
288
              ],
289
              number_of_servers=[num_inspectors, num_repairers],
290
              routing=[[0.0, prob_need_repair], [0.0, 0.0]],
291
292
          return N
293
```

A Network object is used by Ciw to access system parameters. For example one piece of information it holds is the number of nodes of the system:

```
Python input

N = build_network_object()
print(N.number_of_nodes)
```

which gives:

```
Python output

296 2
```

Now we have defined the system, we need to use this to build the virtual representation of the system: in Ciw this is a Simulation object. Once this is built the simulation can be run, that is observe it for one virtual day. The following function does this:

```
Python input
      def run simulation(network, seed=0):
297
          """Builds a simulation object and runs it for 8 time units.
298
299
          Args:
300
              network: a Ciw network object
301
              seed: a float (default: 0)
302
303
          Returns:
304
              a Ciw simulation object after a run of the simulation
305
306
         max_time = 8
307
          ciw.seed(seed)
308
          Q = ciw.Simulation(network)
309
          Q.simulate_until_max_time(max_time)
310
          return Q
311
```

Notice here a random seed is set. This is because there is some element of randomness when initialising this object, and much randomness in running the simulation, and in order to ensure reproducible results we force the pseudorandom number generator to produce the same sequence of pseudorandom numbers each time. Notice also that the simulation always begins with an empty system, so the first bicycle to arrive will never wait for service. Depending on the situation this may be an unwanted

feature, though not in this case as it is reasonable to assume that the bicycle shop will begin the day with no customers.

Now we wish to count the number of bicycles that have finished service, and to count the number of those whose entire journey through the system lasted longer than 0.5 hours. In order to do so, we'll use the pandas library for efficient manipulation of data frames.

```
Python input
      import pandas as pd
312
313
314
      def get_proportion(Q):
315
          """Returns the proportion of bicycles spending over a given
316
          limit at the repair shop.
317
318
          Args:
319
              Q: a Ciw simulation object after a run of the
320
                  simulation
321
322
323
          Returns:
              a real
324
          n n n
325
          limit = 0.5
326
          inds = Q.nodes[-1].all_individuals
327
          recs = pd.DataFrame(
328
              dr for ind in inds for dr in ind.data records
329
          )
330
          recs["total time"] = (
331
              recs["exit_date"] - recs["arrival_date"]
332
333
          total_times = recs.groupby("id_number")["total_time"].sum()
334
          return (total_times > limit).mean()
335
```

Altogether these functions can define the system, run one day of our system, and then find the proportion of bicycles spending over half an hour in the shop:

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```
Python input

N = build_network_object()
Q = run_simulation(N)
p = get_proportion(Q)
print(round(p, 6))
```

This piece of code gives

```
Python output

0.261261
```

meaning 26.13% of all bicycles spent longer than half an hour at the repair shop. However this particular day may have contained a number of extreme events. For a more accurate proportion this experiment should be repeated, and an average proportion taken. In order to do so, let's write a function that performs the above experiment over a number of trials, then finds an average proportion:

Python input

```
def get_average_proportion(num_inspectors=1, num_repairers=2):
341
          """Returns the average proportion of bicycles spending over
342
          a given limit at the repair shop.
343
344
          Args:
345
              num_inspectors: a positive integer (default: 1)
346
              num_repairers: a positive integer (default: 2)
347
348
          Returns:
349
              a real
350
          .....
351
         num_trials = 100
352
         N = build_network_object(
353
              num inspectors=num inspectors,
354
              num_repairers=num_repairers,
355
356
         proportions = []
357
          for trial in range(num_trials):
358
              Q = run_simulation(N, seed=trial)
359
              proportion = get_proportion(Q=Q)
360
              proportions.append(proportion)
361
          return sum(proportions) / num trials
362
```

This can be used to find the average proportion over 100 trials for the current system of one inspector and two repair people:

```
Python input _
     p = get_average_proportion(num_inspectors=1, num_repairers=2)
363
     print(round(p, 6))
364
```

which gives:

```
_____Python output _____
    0.159354
365
```

that is, on average 15.94% of bicycles will spend longer than 30 minutes at the repair shop.

Now consider the two possible future scenarios we wish top compare: hiring an extra member of staff to serve at the inspection desk, or hiring an extra member of staff at the repair workshop. Which scenario yields a smaller proportion of bicycles spending over 30 minutes at the shop? Let's investigate. First look at the situation where the additional member of staff works at the inspection desk:

```
Python input

p = get_average_proportion(num_inspectors=2, num_repairers=2)
print(round(p, 6))

which gives:

Python output

0.038477
```

that is 3.85% of bicycles.

Now look at the situation where the additional member of staff works at the repair workshop:

```
p = get_average_proportion(num_inspectors=1, num_repairers=3)
print(round(p, 6))

which gives:

Python input

Python input

0.103591
```

that is 10.36% of bicycles.

Therefore an additional member of staff at the inspection desk would be more beneficial than an additional member of staff at the repair workshop.



Figure 3.2 Diagrammatic representation of the forked trajectories a bicycle can take

3.4 SOLVING WITH R

In this book we will use the Simmer package in order to conduct discrete event simulation in R. Simmer uses the process based approach, which means we must define the each bicycle's sequence of actions, and then generate a number of bicycles with these sequences.

In Simmer these sequences of actions are made up of trajectories. The diagram in Figure 3.2 shows the branched trajectories than a bicycle would take at the repair shop:

The function below defines a simmer object that describes these trajectories:

```
R input
     library(simmer)
372
373
      #' Returns a simmer trajectory object outlining the bicycles
374
        path through the repair shop
376
      #' @return A simmer trajectory object
377
      define bicycle trajectories <- function() {</pre>
378
        inspection_rate <- 20</pre>
379
        repair_rate <- 10
380
        prob need repair <- 0.8
381
        bicycle <-
          trajectory("Inspection") %>%
383
          seize("Inspector") %>%
384
          timeout(function() {
385
            rexp(1, inspection_rate)
386
          }) %>%
387
          release("Inspector") %>%
388
          branch(
389
            function() (runif(1) < prob_need_repair),</pre>
390
            continue = c(F),
391
            trajectory("Repair") %>%
392
               seize("Repairer") %>%
393
               timeout(function() {
394
                 rexp(1, repair_rate)
395
              }) %>%
396
              release("Repairer"),
397
            trajectory("Out")
398
          )
399
        return(bicycle)
400
401
```

These trajectories are not very useful alone, we are yet to define the resources used, or a way to generate bicycles with these trajectories. This is done in the function below, which begins by defining a repair_shop with one resource labelled "Inspector", and two resources labelled "Repairer". Once this is built the simulation can be run, that is observe it for one virtual day. The following function does all this:

R input

```
Runs one trial of the simulation.
402
     # '
403
     #' Oparam bicycle a simmer trajectory object
404
     #' Oparam num_inspectors positive integer (default: 1)
405
     #' Oparam num_repairers positive integer (default: 2)
406
     #' @param seed a float (default: 0)
407
408
     #' @return A simmer simulation object after one run of
409
                  the simulation
410
     run simulation <- function(bicycle,</pre>
411
                                   num_inspectors = 1,
                                   num_repairers = 2,
413
                                   seed = 0) {
414
       arrival rate <- 15
415
       max_time <- 8
416
       repair_shop <-
417
          simmer("Repair Shop") %>%
418
          add_resource("Inspector", num_inspectors) %>%
419
          add_resource("Repairer", num_repairers) %>%
420
          add generator("Bicycle", bicycle, function() {
421
            rexp(1, arrival rate)
422
          })
423
424
       set.seed(seed)
425
       repair_shop %>% run(until = 8)
426
       return(repair_shop)
427
428
```

Notice here a random seed is set. This is because there are elements of randomness when running the simulation, and in order to ensure reproducible results we force the pseudorandom number generator to produce the same sequence of pseudorandom numbers each time. Notice also that the simulation always begins with an empty system, so the first bicycle to arrive will never wait for service. Depending on the situation this may be an unwanted feature, though not in this case as it is reasonable to assume that the bicycle shop will begin the day with no customers.

Now we wish to count the number of bicycles that have finished service, and to count the number of those whose entire journey through the system lasted longer than 0.5 hours. Using Simmer's get mon arrivals() function we can get a data frame of records to manipulate.

```
R input
        Returns the proportion of bicycles spending over 30
429
        minutes in the repair shop
430
431
        Oparam repair_shop a simmer simulation object
432
433
     #' @return a float between 0 and 1
434
     get proportion <- function(repair shop) {</pre>
435
       limit <- 0.5
436
       recs <- repair shop %>% get mon arrivals()
437
       total times <- recs$end time - recs$start time
438
       return(mean(total times > 0.5))
439
440
```

Altogether these functions can define the system, run one day of our system, and then find the proportion of bicycles spending over half an hour in the shop:

```
bicycle <- define_bicycle_trajectories()
repair_shop <- run_simulation(bicycle = bicycle)
print(get_proportion(repair_shop = repair_shop))
```

This piece of code gives

```
R output

[1] 0.04032258
```

meaning 4.03% of all bicycles spent longer than half an hour at the repair shop. However this particular day may have contained a number of extreme events. For a more accurate proportion this experiment should be repeated, and an average proportion taken. In order to do so, let's write a function that performs the above experiment over a number of trials, then finds an average proportion:

```
R input
      #' Returns the average proportion of bicycles spending over
445
      #' a given limit at the repair shop.
446
447
      #' Oparam num_inspectors positive integer (default: 1)
448
      #' Oparam num_repairers positive integer (default: 2)
449
450
      #' @return a float between 0 and 1
451
      get_average_proportion <- function(num_inspectors = 1,</pre>
452
                                             num_repairers = 2) {
453
        num_trials <- 100</pre>
454
        bicycle <- define bicycle trajectories()</pre>
455
        proportions <- c()</pre>
456
        for (trial in 1:num trials) {
457
          repair shop <- run simulation(</pre>
458
            bicycle = bicycle,
459
            num_inspectors = num_inspectors,
460
            num_repairers = num_repairers,
461
            seed = trial
462
463
          proportion <- get proportion(</pre>
464
            repair_shop = repair_shop
465
466
          proportions[trial] <- proportion</pre>
467
468
        return(mean(proportions))
469
470
```

This can be used to find the average proportion over 100 trials:

```
R input
     print(
471
       get average proportion(
472
         num_inspectors = 1,
473
         num_repairers = 2)
474
475
```

which gives:

```
R output

[1] 0.1551579
```

that is, on average 15.52% of bicycles will spend longer than 30 minutes at the repair shop.

Now consider the two possible future scenarios we wish top compare: hiring an extra member of staff to serve at the inspection desk, or hiring an extra member of staff at the repair workshop. Which scenario yields a smaller proportion of bicycles spending over 30 minutes at the shop? Let's investigate. First look at the situation where the additional member of staff works at the inspection desk:

```
print(
    get_average_proportion(
    num_inspectors = 2,
    num_repairers = 2)
)
```

which gives:

```
R output
[1] 0.04115338
```

that is 4.12% of bicycles.

Now look at the situation where the additional member of staff works at the repair workshop:

```
R input

print(
get_average_proportion(
num_inspectors = 1,
num_repairers = 3)
)
```

which gives:

that is 10.01% of bicycles.

Therefore an additional member of staff at the inspection desk would be more beneficial than an additional member of staff at the repair workshop.

3.5 RESEARCH

TBA

		_

		_

Modelling with Differential Equations

Stems often change in a way that depends on their current state. For example, the speed at which a cup of coffee cools down depends on its current temperature. These types of systems are called dynamical systems and are modelled mathematically using differential equations. In this chapter we will consider a direct solution approach using symbolic mathematics.

4.1 PROBLEM

Consider the following situation: the entire population of a small rural town has caught a cold. All 100 individuals will recover at an average rate of 2 per day. The town leadership have noticed that being ill costs approximately £10 per day, this is due to general lack of productivity, poorer mood and other intangible aspects. They need to decide whether or not to order cold medicine which would **double** the recover rate. The cost of of the cold medicine is a one off cost of £5 per person.

4.2 THEORY

In the case of this town, the overall rate at which people get better is dependent on the number of people in how are ill. This can be represented mathematically using a differential equation which is a way of relating the rate of change of a system to the state of the system itself.

In general if we are interested in some variable x over time t the differential function equation will be of the form:

$$\frac{dx}{dt} = f(x) \tag{4.1}$$

For some function f. In our case, if we denote the number of infected individuals as I where we implicitly mean that I is a function of time: I = I(t) and the rate at which individuals recover by α then the differential equation that describes the above situation is:

$$\frac{dI}{dt} = -\alpha I \tag{4.2}$$

Finding a solution to this differential equation means finding an expression for I that when differentiated gives -alphaI.

In this particular case, one such function is:

$$I(t) = e^{-\alpha t} \tag{4.3}$$

However, I(0) = 1 whereas for our problem we know that at time t = 0 there are 100 infected individuals. Indeed a differential equation defines a family of solutions and we need to know some sort of initial (also referred to as boundary) condition to have the exact solution. Which in this case would be:

$$I(t) = 100e^{-\alpha t} \tag{4.4}$$

To evaluate the cost we then need to know the sum of the values of that function over time. Integration gives us exactly this, so the cost would be:

$$K \int_0^\infty I(t)dt \tag{4.5}$$

where K is the cost per person per unit time.

In the upcoming sections we will confirm and use code to carry out the above efficiently so as to answer the original question.

4.3 SOLVING WITH PYTHON

The first step we will take is to write a function to obtain the differential equation. Note that here we will be using the Python library sympy which allows us to carry out symbolic calculations.

Python input _ import sympy as sym 489 490 t = sym.Symbol("t") 491 alpha = sym.Symbol("alpha") 492 $I_0 = sym.Symbol("I_0")$ 493 I = sym.Function("I") 494 495 496 def get_equation(alpha=alpha): 497 """Return the differential equation. 498 499 Args: 500 alpha: a float (default: symbolic alpha) 501 502 Returns: 503 A symbolic equation 504 505 return sym.Eq(sym.Derivative(I(t), t), -alpha * I(t)) 506

Using this we can get the equation that defines the population change over time:

```
____ Python input
     eq = get_equation()
507
     print(eq)
508
```

which gives:

```
Python output
    Eq(Derivative(I(t), t), -alpha*I(t))
509
```

Note that if you are using Jupyter then your output will actually be a well rendered mathematical equation:

$$\frac{d}{dt}I(t) = -\alpha I(t)$$

Note that we can pass a value to α if we want to:

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```
Python input

eq = get_equation(alpha=1)
print(eq)
```

```
Python output

Eq(Derivative(I(t), t), -I(t))
```

We will now write a function to obtain the solution to this differential

```
Python input
     def get_solution(I_0=I_0, alpha=alpha):
513
          """Return the solution to the differential equation.
514
515
          Args:
516
              I\_0: a float (default: symbolic I\_0)
517
              alpha: a float (default: symbolic alpha)
518
519
          Returns:
520
              A symbolic equation
521
522
          eq = get_equation(alpha=alpha)
523
         return sym.dsolve(eq, I(t), ics={I(0): I 0})
524
```

We can verify the solution discussed previously:

```
Python input

sol = get_solution()
print(sol)
```

which gives:

```
Python output

Eq(I(t), I_0*exp(-alpha*t))
```

$$I(t) = I_0 e^{-\alpha t}$$

We can use sympy itself to verify the result, by taking the derivative of the right hand side of our solution.

```
Python input

[print(sym.diff(sol.rhs, t) == -alpha * sol.rhs)
```

which gives:

```
Python output

True
```

All of the above has given us the general solution in terms of $I(0) = I_0$ and α however we have written the code in such a way as we can pass the actual parameters:

```
Python input

sol = get_solution(alpha=2, I_0=100)
print(sol)
```

which gives:

```
Python output

Eq(I(t), 100*exp(-2*t))
```

Now, to calculate the cost we will write a function to integrate our result:

```
Python input
     def get_cost(
533
          I_0=I_0, alpha=alpha, cost_per_person=10, cost_of_cure=0,
534
     ):
535
          """Return the cost.
536
537
          Args:
538
              I_0: a float (default: symbolic I_0)
539
              alpha: a float (default: symbolic alpha)
540
              cost_per_person: a float (default: 10)
541
              cost_of_cure: a float (default: 0)
542
543
          Returns:
544
              A symbolic expression
545
546
          I_sol = get_solution(I_0=I_0, alpha=alpha)
547
          return (
548
              sym.integrate(I_sol.rhs, (t, 0, sym.oo))
549
              * cost_per_person
550
              + cost_of_cure * I_0
551
          )
552
```

We can now obtain the cost without purchasing the cure:

```
Python input

I_0 = 100
alpha = 2
cost_without_cure = get_cost(I_0=I_0, alpha=alpha)
print(cost_without_cure)
```

which gives:

```
Python output

557

500
```

The cost with cure can use the above with a modified α and a non zero cost of the cure itself:

```
Python input
     cost_of_cure = 5
558
     cost_with_cure = get_cost(
559
         I_0=I_0, alpha=2 * alpha, cost_of_cure=cost_of_cure
560
561
     print(cost_with_cure)
562
```

which gives:

```
Python output
     750
563
```

So given the current parameters it is not worth purchasing the cure.

SOLVING WITH R

R does not have a bespoke symbolic mathematics library however it has an interface to use Python's sympy called rSymPy. In general we would recommend to use Python directly when needing to carry out symbolic mathematical calculations however this is not always possible. Furthermore, the development of rSymPy seems to no longer be active and due to the dependency on Java could be problematic to use. One of other the difficulties of using rSymPy is that it works by passing string back and forth to Python's sympy. As such, we will need to have more of a manual approach and so we will not be able to write the modular code we have written throughout the book.

First let us setup and solve the differential equation:

```
R input
      library("rSymPy", warn.conflicts = FALSE, quietly = TRUE)
564
565
      t <- Var("t")
566
      alpha <- Var("alpha")</pre>
567
      I <- Var("I")</pre>
568
      sympy("dsolve(Derivative(I(t), t) - alpha *I(t), I(t))")
569
```

this will give us:

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```
R output

[1] "C1*exp(alpha*t)"
```

We can then solve the equation directly to find the initial condition:

```
R input

sympy("solve(C1*exp(alpha*0) - I_0, C1)")
```

This gives:

```
772 [1] "[I_0]"
```

Note here that this is in fact using a much older version of Sympy so we cannot take advantage of some of the newer commands.

So we now the form of our particular solution, thus we can now write a function that will give us the cost:

```
R input
      #' Return the cost
573
      #'
574
      #' @param I_O a float (default: symbolic I_O)
575
      #' @param alpha a float (default: symbolic alpha)
576
      #' @param cost_per_person a float (default: 10)
577
      #' @param cost_of_cure a float (default: 0)
578
579
      #' @return A symbolic expression
580
      get cost <- function(I 0 = Var("I 0"),</pre>
581
                             alpha = Var("alpha"),
582
                             cost_per_person = 10,
583
                             cost_of_cure = 0) {
584
        t <- Var("t")
585
        solution cmd <- sprintf(</pre>
586
          "I_sol = %s * exp(-%s * t)",
587
          I_0,
588
          alpha
589
590
        sympy(solution_cmd)
591
        cost cmd <- sprintf(</pre>
592
          "integrate(I_{sol}, (t, 0, oo)) * %s + %s * %s",
593
          cost_per_person, cost_of_cure,
594
          I_0
595
596
        return(sympy(cost_cmd))
597
598
```

Using this:

```
R input
     I_0 <- 100
599
     alpha <- 2
600
     cost_without_cure <- get_cost(I_0 = I_0, alpha = alpha)</pre>
601
     print(cost_without_cure)
602
```

which gives:

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The cost with cure can use the above with a modified α and a non zero cost of the cure itself:

```
Cost_of_cure <- 5
cost_with_cure <- get_cost(
    I_0 = I_0, alpha = 2 * alpha, cost_of_cure = cost_of_cure
)
print(cost_with_cure)
```

which gives:

```
R output

750
```

So given the current parameters it is not worth purchasing the cure.

4.5 RESEARCH

TBA

Bibliography

		_