

# 11. Using R to Support Simulation


Data Science for OR - J. Duggan

# Using the tidyverse to support simulation

- Tidying input data
- Analysing simulation output
- Running sensitivity analysis

## NOTES AND INSIGHTS

### **Input and output data analysis for system dynamics modelling using the tidyverse libraries of R**

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# (1) Tidying input data

Table 2. Time series influenza data from the 1957 pandemic (U.K. data)

| Week | Young | Child | Adult | Elderly |
|------|-------|-------|-------|---------|
| 1    | 0     | 0     | 1     | 1       |
| 2    | 0     | 2     | 6     | 1       |
| 3    | 0     | 2     | 4     | 2       |
| 4    | 23    | 73    | 63    | 11      |
| 5    | 63    | 208   | 173   | 41      |
| 6    | 73    | 207   | 171   | 27      |
| 7    | 66    | 150   | 143   | 7       |
| 8    | 26    | 40    | 87    | 29      |
| 9    | 17    | 18    | 33    | 12      |
| 10   | 3     | 4     | 13    | 6       |
| 11   | 2     | 6     | 16    | 5       |
| 12   | 1     | 6     | 11    | 3       |
| 13   | 0     | 1     | 6     | 5       |
| 14   | 0     | 2     | 2     | 2       |
| 15   | 0     | 1     | 3     | 0       |
| 16   | 0     | 1     | 4     | 6       |
| 17   | 0     | 1     | 3     | 0       |
| 18   | 2     | 1     | 7     | 1       |
| 19   | 1     | 1     | 6     | 2       |

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## Using readr to access data

```
inc <- read_csv("../..//11 simulation/code/sdr_paper1/data/Inc.csv")
```

```
## Parsed with column specification:
## cols(
##   Week = col_double(),
##   Young = col_double(),
##   Child = col_double(),
##   Adult = col_double(),
##   Elderly = col_double()
## )
```

```
slice(inc,1:2)
```

```
## # A tibble: 2 x 5
##   Week Young Child Adult Elderly
##   <dbl> <dbl> <dbl> <dbl>   <dbl>
## 1     1     1     0     0     1
## 2     1     0     0     1     1
```

# Convert to Tidy Data

```
t_inc <- gather(inc, Cohort, Incidence, Young:Elderly)
slice(t_inc, 1:8)
```

```
## # A tibble: 8 x 3
##   Week Cohort Incidence
##   <dbl> <chr>      <dbl>
## 1     1  1 Young          0
## 2     2  2 Young          0
## 3     3  3 Young          0
## 4     4  4 Young         23
## 5     5  5 Young         63
## 6     6  6 Young         73
## 7     7  7 Young         66
## 8     8  8 Young         26
```

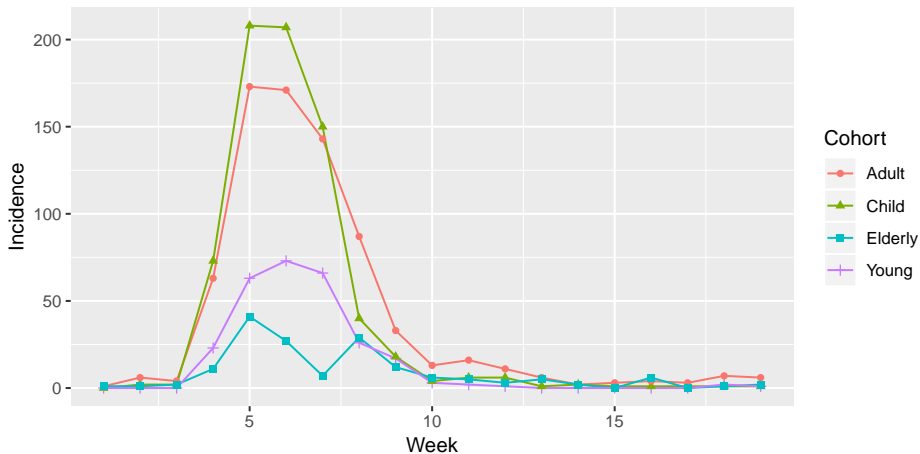
# Summarise Data

```
wk_tot <- t_inc %>% group_by(Week) %>%  
  summarise(Incidence=sum(Incidence)) %>%  
  arrange(desc(Incidence))  
slice(wk_tot,1:6)
```

```
## # A tibble: 6 x 2  
##   Week Incidence  
##   <dbl>     <dbl>  
## 1     5       485  
## 2     6       478  
## 3     7       366  
## 4     8       182  
## 5     4       170  
## 6     9        80
```

## Plot tidy data

```
ggplot(t_inc, aes(x=Week, y=Incidence, color=Cohort,  
  shape=Cohort)) + geom_line() + geom_point()
```



# Descriptive Statistics

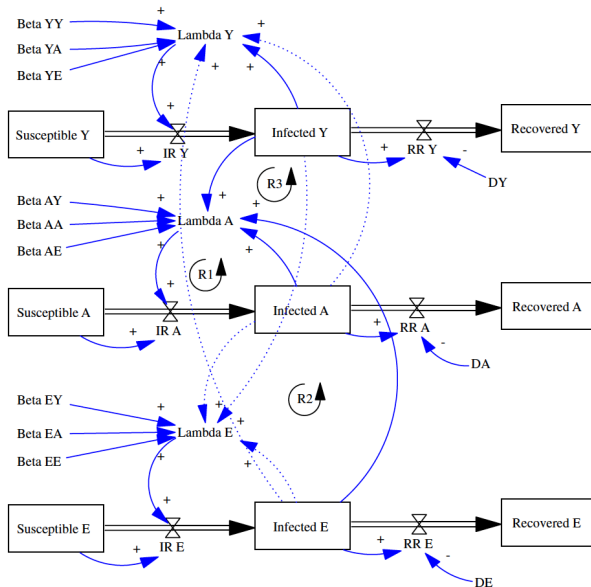
```
t_coh <- t_inc %>%  
  group_by(Cohort) %>%  
  summarise(TotalInfected=sum(Incidence),  
            PeakValue=max(Incidence),  
            PeakWeek=Week[which(Incidence==max(Incidence))],  
            AvrValue=mean(Incidence),  
            SD=sd(Incidence))  
  
t_coh
```

```
## # A tibble: 4 x 6
```

```
## Cohort TotalInfected PeakValue PeakWeek AvrValue SD  
## <chr> <dbl> <dbl> <dbl> <dbl> <dbl>  
## 1 Adult 752 173 5 39.6 59.3  
## 2 Child 724 208 5 38.1 70.1  
## 3 Elderly 161 41 5 8.47 11.4  
## 4 Young 277 73 6 14.6 24.9
```



## (2) Analysing Simulation Output



## Simulation results - many columns

|    |      |                   |                    |                   |
|----|------|-------------------|--------------------|-------------------|
| ## | [1]  | "Time"            | "Beta AA"          | "Beta AE"         |
| ## | [4]  | "Beta AY"         | "Beta EA"          | "Beta EE"         |
| ## | [7]  | "Beta EY"         | "Beta YA"          | "Beta YE"         |
| ## | [10] | "Beta YY"         | "CE AA"            | "CE AE"           |
| ## | [13] | "CE AY"           | "CE EA"            | "CE EE"           |
| ## | [16] | "CE EY"           | "CE YA"            | "CE YE"           |
| ## | [19] | "CE YY"           | "DA"               | "DE"              |
| ## | [22] | "DY"              | "Infected A"       | "Infected E"      |
| ## | [25] | "Infected Y"      | "IR A"             | "IR E"            |
| ## | [28] | "IR Y"            | "Lambda A"         | "Lambda E"        |
| ## | [31] | "Lambda Y"        | "Pop A"            | "Pop E"           |
| ## | [34] | "Pop Y"           | "Prop A Infected"  | "Prop E Infected" |
| ## | [37] | "Prop Y Infected" | "Recovered A"      | "Recovered E"     |
| ## | [40] | "Recovered Y"     | "RR A"             | "RR E"            |
| ## | [43] | "RR Y"            | "Susceptible A"    | "Susceptible E"   |
| ## | [46] | "Susceptible Y"   | "Total Population" |                   |

## Selecting the stocks

```
out <- res %>%  
  select(Time, starts_with("Susceptible"),  
         starts_with("Infected"),  
         starts_with("Recovered"))  
glimpse(out)
```

```
## Observations: 161
```

```
## Variables: 10
```

```
## $ Time          <dbl> 0.000, 0.125, 0.250, 0.375, 0.500,  
## $ `Susceptible A` <dbl> 50000, 50000, 50000, 50000, 50000,  
## $ `Susceptible E` <dbl> 25000, 25000, 25000, 25000, 25000,  
## $ `Susceptible Y` <dbl> 25000, 25000, 25000, 25000, 25000,  
## $ `Infected A`    <dbl> 0.00000, 0.00000, 0.01562, 0.05469,  
## $ `Infected E`    <dbl> 0.0000, 0.1250, 0.2891, 0.5083, 0.8  
## $ `Infected Y`    <dbl> 1.000, 1.312, 1.738, 2.321, 3.124,  
## $ `Recovered A`   <dbl> 0.00000, 0.00000, 0.00000, 0.00097,  
## $ `Recovered E`   <dbl> 0.00000, 0.00000, 0.00000, 0.00704, 0.02500,  
## $ `Recovered Y`   <dbl> 0.00000, 0.00000, 0.00000, 0.00000, 0.00000
```

## Convert to tidy format

```
out_td <- out %>%  
  gather(key=Variable,value = Amount,  
         `Susceptible A`:`Recovered Y`)  
slice(out_td,1:5)
```

```
## # A tibble: 5 x 3  
##   Time Variable      Amount  
##   <dbl> <chr>         <dbl>  
## 1 0      Susceptible A  50000  
## 2 0.125 Susceptible A  50000  
## 3 0.25  Susceptible A  50000  
## 4 0.375 Susceptible A  50000  
## 5 0.5   Susceptible A  50000
```

## Add cohort and stock information

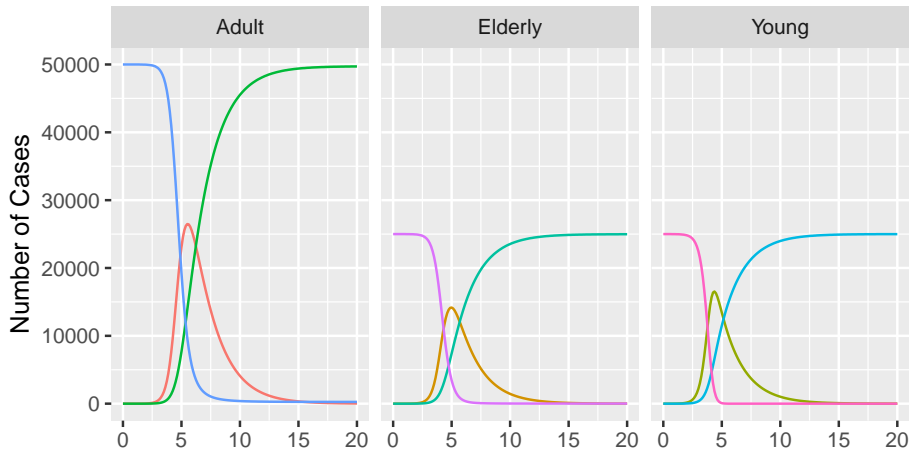
```
new_td <- out_td %>%  
  mutate(Cohort=case_when(  
    grepl("A$",Variable) ~ "Adult",  
    grepl("E$",Variable) ~ "Elderly",  
    grepl("Y$",Variable) ~ "Young"),  
  Class=case_when(  
    grepl("^S",Variable) ~ "Susceptible",  
    grepl("^I",Variable) ~ "Infected",  
    grepl("^R",Variable) ~ "Recovered"))  
slice(new_td,1:2)
```

```
## # A tibble: 2 x 5
```

```
##   Time Variable      Amount Cohort Class  
##   <dbl> <chr>          <dbl> <chr> <chr>  
## 1 0      Susceptible A  50000 Adult  Susceptible  
## 2 0.125 Susceptible A  50000 Adult  Susceptible
```

# Display chart

```
ggplot(new_td) + geom_path(aes(x=Time,y=Amount,  
  colour=Variable))+ylab("Number of Cases")+  
  facet_wrap(~Cohort)+guides(colour=F)
```



### (3) Exploring Sensitivity Data

```
d <- read_tsv("../..//11 simulation/code/sdr_paper1/data/Sensiti
dim(d)
```

```
## [1] 200 244
```

```
d[1:3,1:5]
```

```
## # A tibble: 3 x 5
```

```
##   Simulation      R0      VF `T1 Infected` `T2 Infected`
##   <dbl> <dbl> <dbl>          <dbl>          <dbl>
## 1         1  2.76 0.0263             1             1.11
## 2         2  2.66 0.0739             1             1.10
## 3         3  4.06 0.159              1             1.19
```

## Convert to Tidy Data

```
START_TIME <- 0
DT <- 0.125

td <- gather(d, TimeVariable, Value, -(Simulation:VF)) %>%
  mutate(TSeq=parse_integer(
    str_extract(TimeVariable, "\\d+"))) %>%
  mutate(SimTime=START_TIME+(TSeq-1)*DT) %>%
  separate(TimeVariable, into = c("T", "Variable")) %>%
  select(Simulation, SimTime, R0, VF, Variable, Value) %>%
  arrange(Simulation, SimTime)

slice(td, 1:2)
```

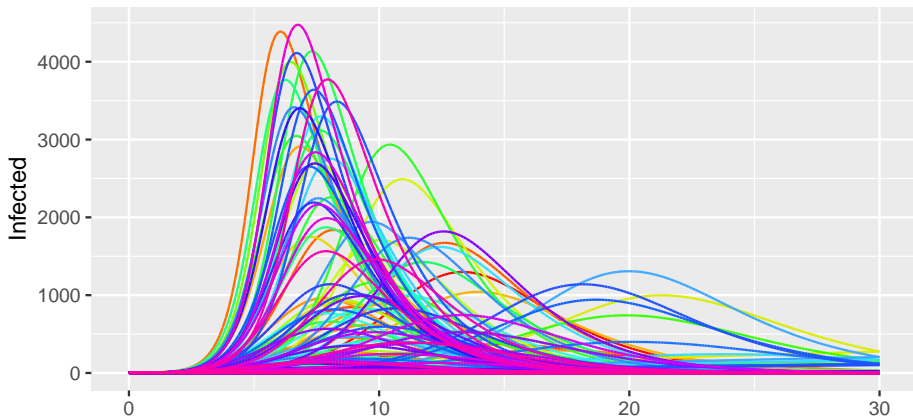
```
## # A tibble: 2 x 6
```

```
##   Simulation SimTime    R0      VF Variable Value
##   <dbl>    <dbl> <dbl> <dbl> <chr>    <dbl>
## 1         1      0.125  0.76  0.0263 Infected 1
```



# Display simulation traces

```
ggplot(td,aes(x=SimTime,y=Value,color=Simulation)) +  
  geom_path() + ylab("Infected") +  
  scale_colour_gradientn(colours=rainbow(10))+  
  xlab("Time (Days)") + guides(color=FALSE)
```



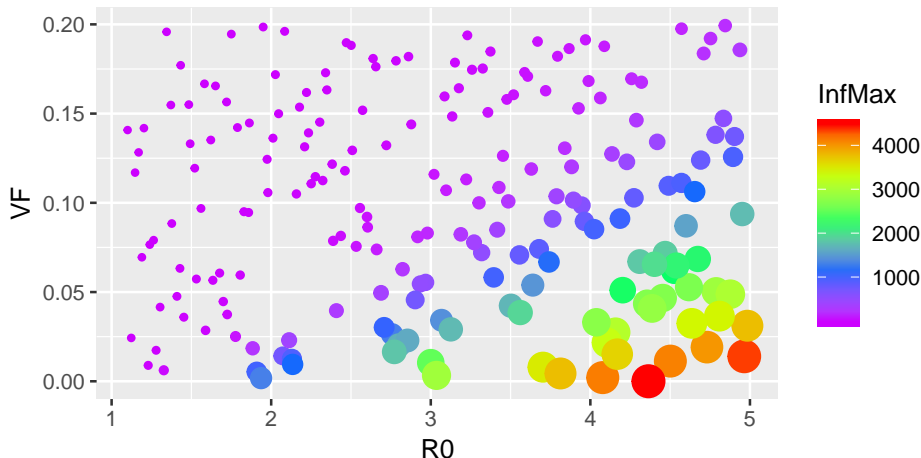
# Calculate Summary Data

```
i_td <- td %>% group_by(Simulation) %>%  
  summarise(InfMax=max(Value),  
            R0=R0[1],  
            VF=VF[1])  
  
slice(i_td,1:5)
```

```
## # A tibble: 5 x 4  
##   Simulation InfMax    R0    VF  
##   <dbl>    <dbl> <dbl> <dbl>  
## 1         1 1298.   2.76 0.0263  
## 2         2   78.9   2.66 0.0739  
## 3         3  127.   4.06 0.159  
## 4         4  367.   2.69 0.0496  
## 5         5   14.0   3.23 0.194
```

# Explore Parameter Space

```
ggplot(data=i_td,aes(x=R0,y=VF,size=InfMax,colour=InfMax)) +  
  geom_point() + guides(size=F) +  
  scale_colour_gradientn(colours=rev(rainbow(5)))
```



## Conclusion

While R is primarily viewed as a toolset to support data scientists, innovative new libraries such as the `tidyverse` can be leveraged to support the system dynamics model-building process. This paper has shown how time series data can be accessed and manipulated, and how the entire model output from a simulation run can be processed for informative summaries and for data visualisation. A further application of the `tidyverse` is to support the process of analysing large datasets produced through sensitivity analysis of system dynamics models.