

# 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/Incidence.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     0     0     1     1
## 2     2     2     6     6     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  Young         0
## 2     2  Young         0
## 3     3  Young         0
## 4     4  Young        23
## 5     5  Young        63
## 6     6  Young        73
## 7     7  Young        66
## 8     8  Young        26
```

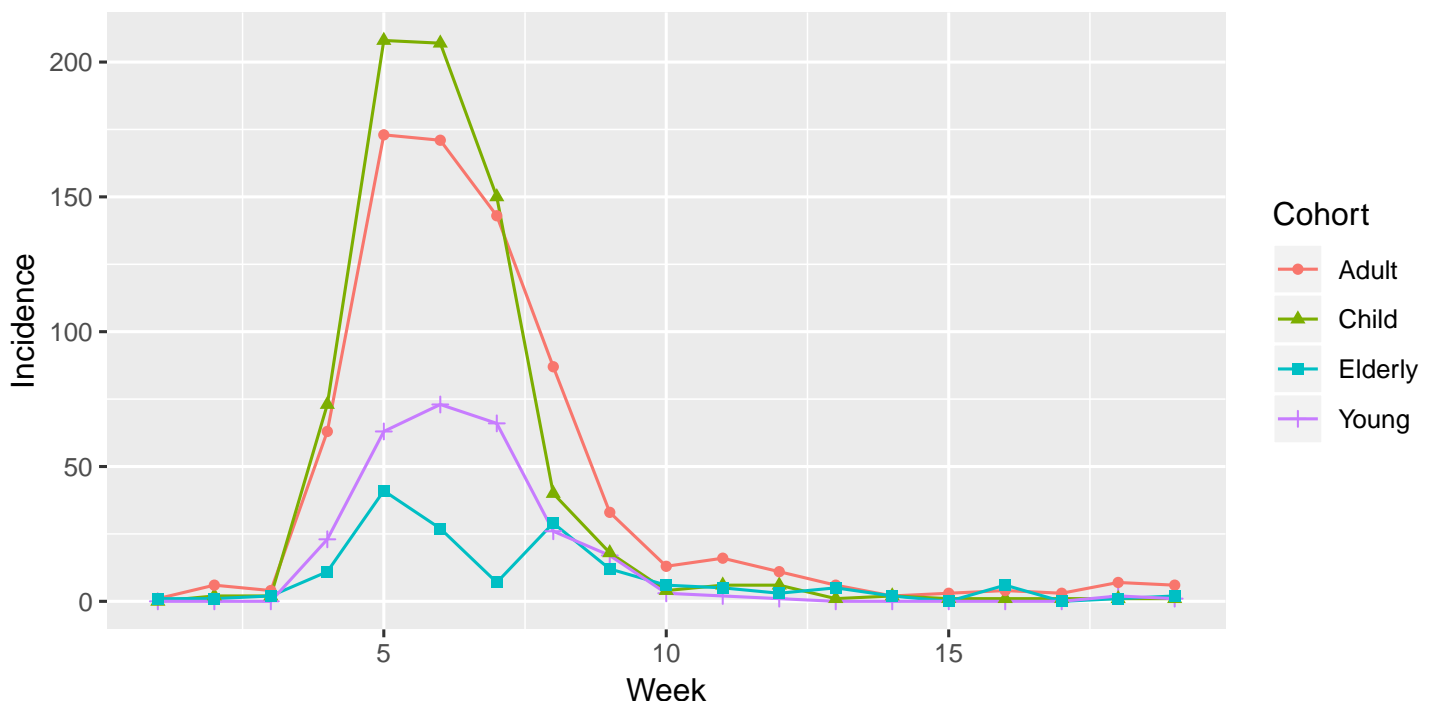
## Summarise Data

```
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

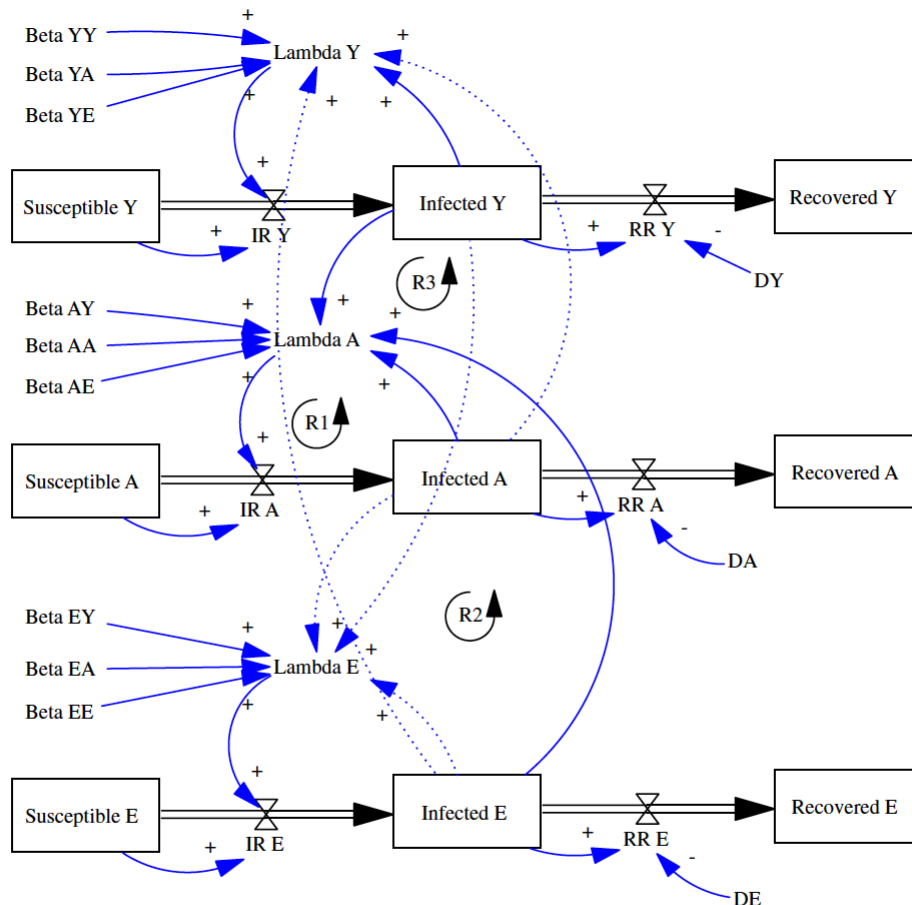
```
t_coh <- t_inc %>%  
  group_by(Cohort) %>%  
  summarise(TotalInfected=sum(Incidence),  
    PeakValue=max(Incidence),  
    PeakWeek=Week[which(Incidence==max(Incidence))],  
    AvrValue=mean(Incidence),  
    SD=sd(Incidence))
```

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, 0.00781, 0.02500,
```

## 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
```

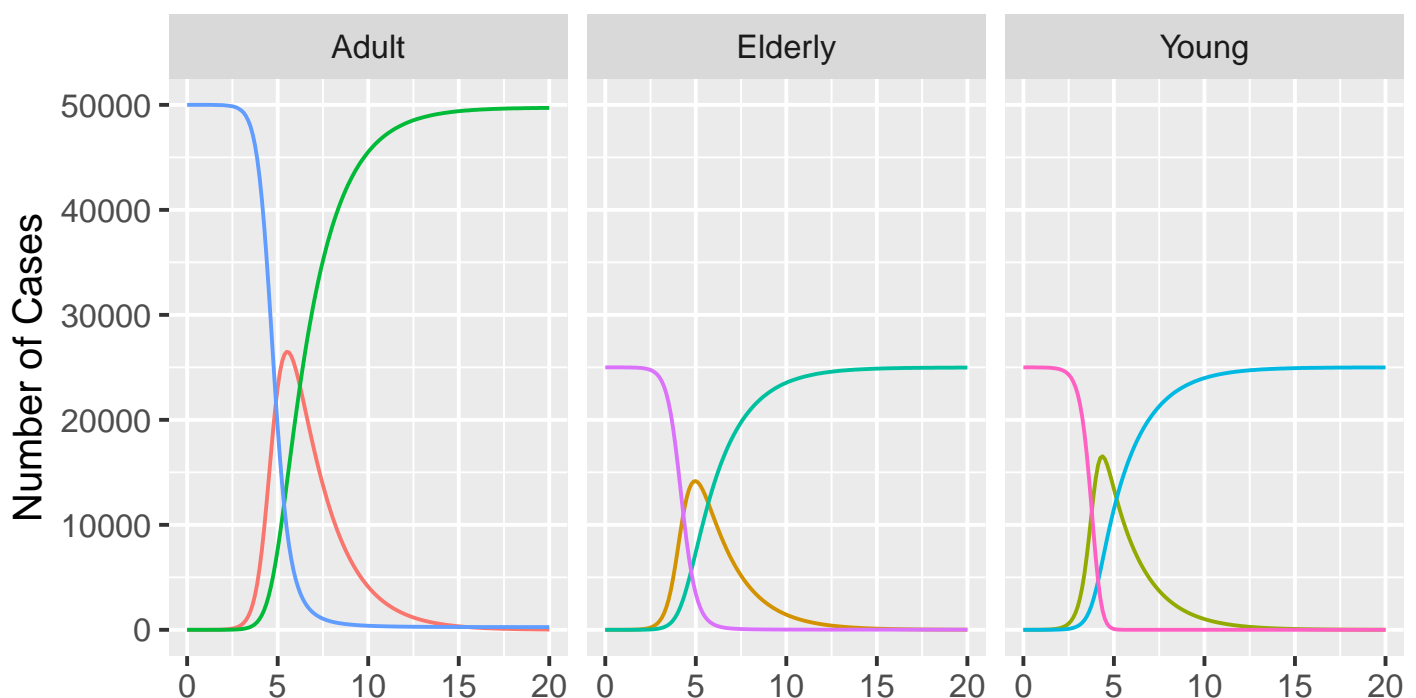
## Add cohort and stock information

```
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/Sensit
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)
```

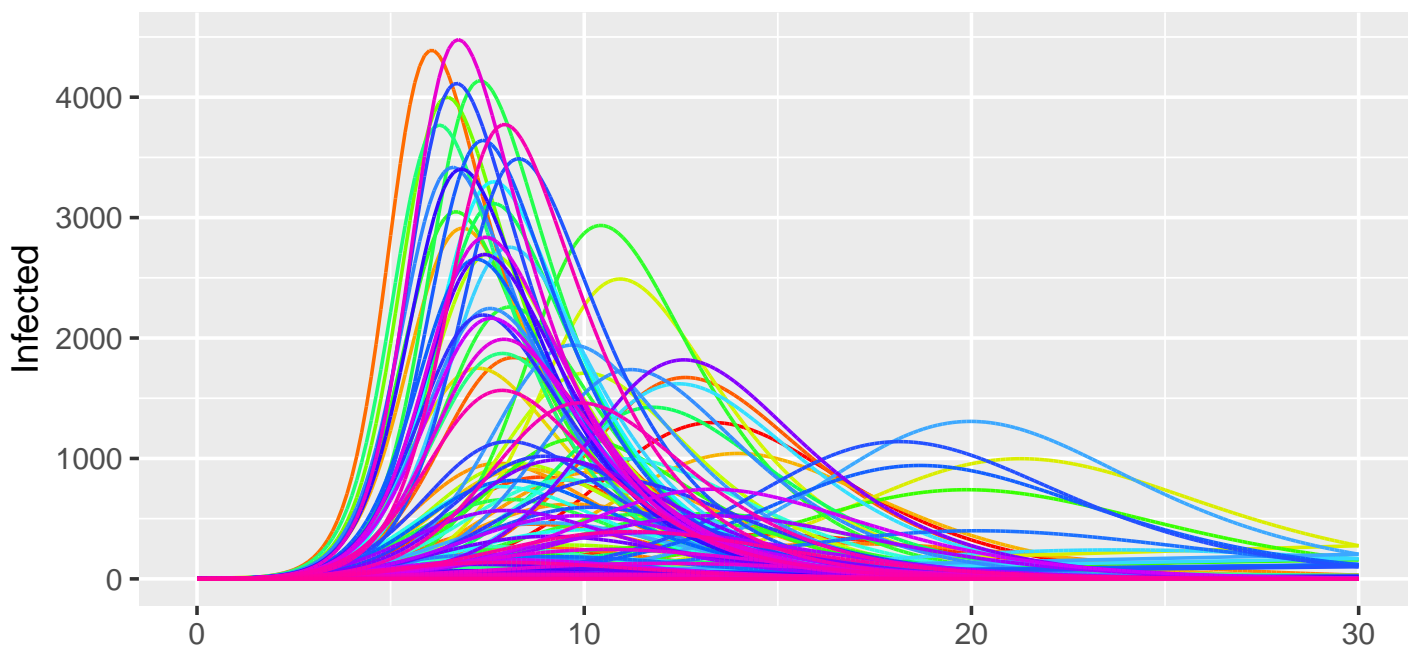
```
## # A tibble: 2 x 6
```

```
##   Simulation SimTime    R0      VF Variable Value
##   <dbl>    <dbl> <dbl> <dbl> <chr>    <dbl>
## 1         1      0  2.76 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)
```



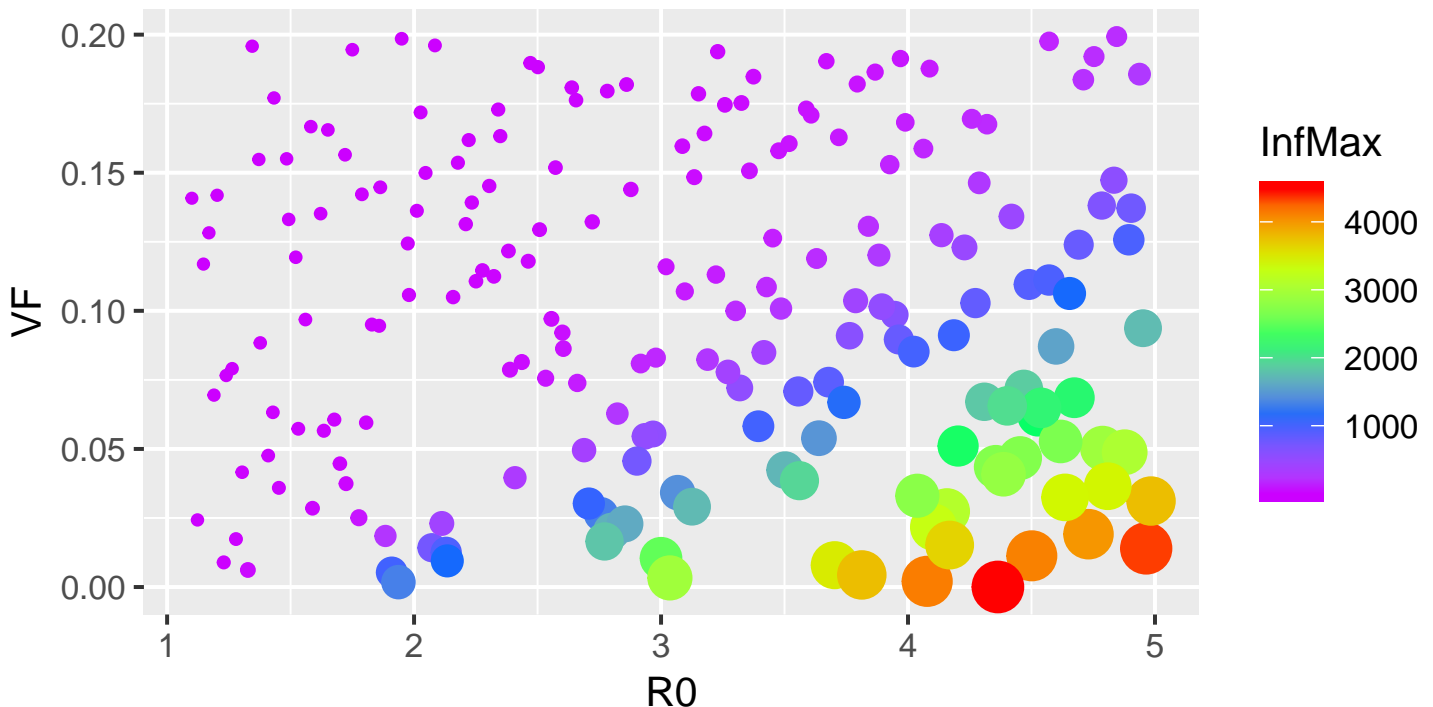
## Calculate Summary Data

```
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)))
```



## Summary

### 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.