AUDPC Analysis

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Background

Rmarkdown is great for combining code and notes into coesive reports on analyses. In this example, I will be using data found on this tutorial to explore the effects of wheat cultivar and fugicide treatment on the incidence of disease caused by *Puccinia striiformis* (stripe rust). I will do this by comparing the **AUDPC** (area under disease progress curve) for the three cultivars and two fungicide treatments.

Analysis

To accomplish this, I will do the following:

- 1. Download and parse the file using read.csv
- 2. Tidy up the data using reshape2 and dplyr functions
- 3. Graph the data using ggplot2

Download and parse data set

data <- read.csv("http://www.apsnet.org/edcenter/advanced/topics/EcologyAndEpidemiologyInR/DiseaseProgr

Tidy up data set

More information on "Tidy" data can be found here.

```
library('dplyr')
library('reshape2')
long_data <- data %>%
  melt(id.vars = "Julian.Date", value.name = "disease") %>%  # long form
  mutate(fungicide = grepl(pattern = "\\.[Tt]rt$", variable)) %>%  # add fungicide column
  transform(fungicide = ifelse(fungicide, "treated", "untreated")) %>%  # make it sensible
  transform(variable = gsub("\\.[Tt]rt$", "", variable))  # clean up the variable
```

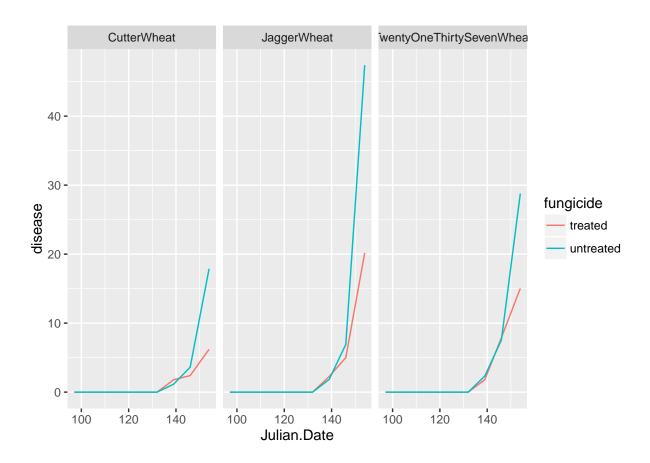
Lets see what the data set looks like after reformatting:

head(long_data)

```
Julian.Date
                                  variable disease fungicide
##
## 1
             97 TwentyOneThirtySevenWheat
                                                  0 untreated
## 2
             104 TwentyOneThirtySevenWheat
                                                  0 untreated
## 3
             111 TwentyOneThirtySevenWheat
                                                  0 untreated
             118 TwentyOneThirtySevenWheat
## 4
                                                  0 untreated
             125 TwentyOneThirtySevenWheat
## 5
                                                  0 untreated
             132 TwentyOneThirtySevenWheat
## 6
                                                  0 untreated
```

Graph AUDPC

```
library('ggplot2')
ggplot(long_data) +
  geom_line(aes(x = Julian.Date, y = disease, color = fungicide)) +
  facet_wrap(~variable)
```



Conclusions

In general, the AUDPC only changes between the fungicide treatments during the last 10 days of the experiment. The "Cutter" seemed to be most resistant to the disease followed by "2137" and "Jagger".