Life history in songbirds — Martin 2015

Martin (2015) studied songbirds in temperate and tropical environments. He showed (Figure 2A) that peak growth rate is higher in species suffering higher nest predation risk, and is lower in tropical species with the same level of risk as temperate species. In the same Figure (2B) he reported that nestling period covaries with growth rate, with tropical species having a shorter nestling periods (for the same growth rate) than temperate species. The file Martin2015_figure2.pdf contains a figure generated with ggplot2 similar to Figure 2 of the original paper. Reproduce the figure using the file Martin2015_data.csv you can find in the ggplot2/data folder.

As always, we need to read the data:

```
m2015 <- read.csv("../data/Martin2015_data.csv",sep = '\t', stringsAsFactors = FALSE)
dim(m2015)</pre>
```

[1] 72 15

```
head(m2015)
```

```
##
                    species
                                 nstldpr
                                             nstl
                                                      krate
                                                                 kwing
## 1 Empidonax occidentalis 0.030200000 15.04500 0.4273536 0.2817678
## 2
             Vireo_plumbeus 0.042200000 13.40000 0.4755502 0.3078776
## 3
               Vireo_gilvus 0.021600000 13.43700 0.4901149 0.2729885
## 4
              Parus_gambeli 0.007380891 21.35714 0.3596292 0.2301780
## 5
         Turdus_migratorius 0.030100000 14.66700 0.4999061 0.3161703
## 6
          Catharus_guttatus 0.047400000 12.62500 0.5266504 0.3275454
##
     PropWCfldg Propmassfldg trips tripsnstl
                                                 CS
                                                        armort
                                                                   lmas aerial
      0.6903986
## 1
                   1.0529779 19.947
                                      6.136000 3.88 0.4126082 1.037426
## 2
                          NA 7.104
                                     2.921000 3.26
                                                            NA 1.225309
                                                                             0
             NΑ
## 3
     0.6521742
                                                                             0
                   0.9856032 10.903
                                     3.502992 3.66 0.5009575 1.099076
## 4
      0.8500607
                   1.0591968 19.818
                                      3.768000 6.58 0.5178348 1.062556
                                                                             0
## 5
      0.6042932
                   0.7856921
                              6.229
                                      1.884000 3.34 0.5029359 1.888179
                                                                             0
      0.6440784
                   0.8528176 5.934 1.896000 3.78 0.4786893 1.489958
                                                                             0
## 6
##
     regurg site
## 1
          0
## 2
## 3
          0
               1
## 4
          0
               1
## 5
          0
               1
## 6
               1
```

For panel A, we want to plot the peak growth rate (krate) against the nestling predation rate (nstldpr), coloring the points according to site.

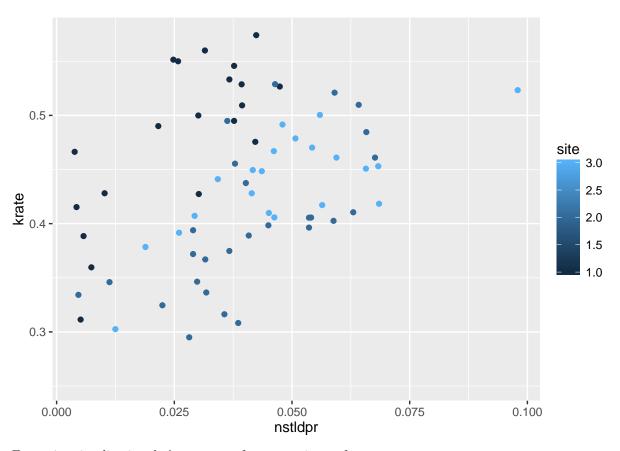
Load the packages:

```
require(ggplot2)
require(dplyr)
require(gridExtra)
```

And start plotting:

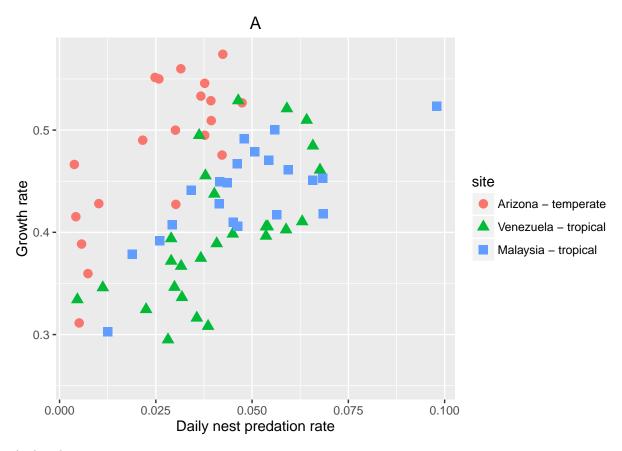
```
plA <- ggplot(data = m2015, aes(x = nstldpr, y = krate, colour = site)) + geom_point()
plA</pre>
```

Warning: Removed 4 rows containing missing values (geom_point).



For easier visualization, let's map transform site into a factor:

Warning: Removed 4 rows containing missing values (geom_point).

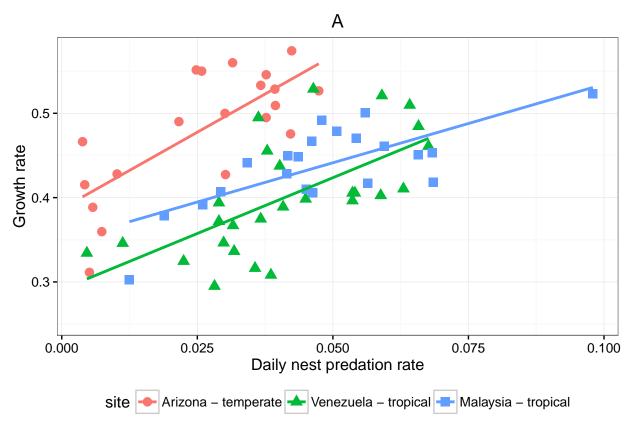


And make it prettier:

```
plA <- plA + theme_bw() + theme(legend.position = "bottom") +
   geom_smooth(method = "glm", se = FALSE)
plA</pre>
```

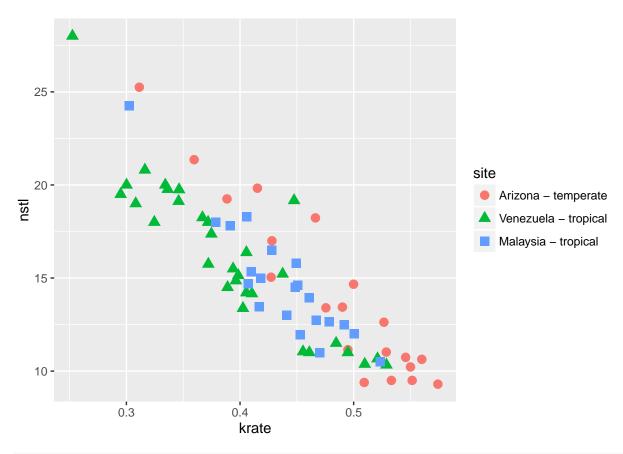
Warning: Removed 4 rows containing non-finite values (stat_smooth).

Warning: Removed 4 rows containing missing values (geom_point).

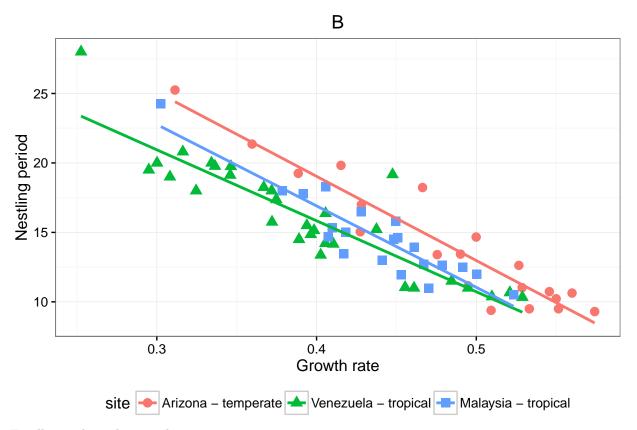


Good. Now let's start working on panel B: we need to plot the nestling period (nstl) against the growth rate (krate). Again, we color and choose shapes according to site.

```
plB <- ggplot(data = m2015, aes(x = krate, y = nstl, colour = site, shape = site)) +
    geom_point(size = 3)
plB</pre>
```



```
# Add labels
plB <- plB + xlab("Growth rate") + ylab("Nestling period") + ggtitle("B")
# Add linear model, and move legend
plB <- plB + theme_bw() + theme(legend.position = "bottom") +
   geom_smooth(method = "glm", se = FALSE)
plB</pre>
```

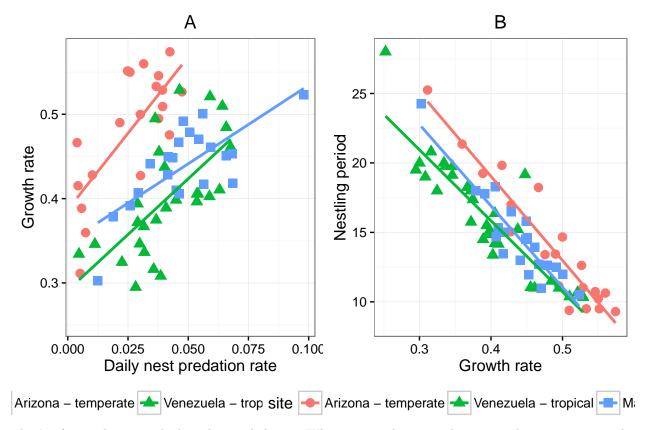


Finally, combine the two plots using gridExtra:

```
grid.arrange(plA, plB, ncol = 2)
```

Warning: Removed 4 rows containing non-finite values (stat_smooth).

Warning: Removed 4 rows containing missing values (geom_point).



That's it! Try playing with the colors and shapes. When you are happy with your results, you can save the graph using the command pdf.

```
pdf(file = "../data/Martin2015_figure2.pdf", width = 12, height = 7)
grid.arrange(plA, plB, ncol = 2)
```

Warning: Removed 4 rows containing non-finite values (stat_smooth).

Warning: Removed 4 rows containing missing values (geom_point).

dev.off()

pdf

2