

dplyr_session2

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Outline

1. `rename()` : rename columns in a data frame
2. `arrange()` : order rows
3. `group_by()` : summary statistics by groups you define, like `ddply(data, grp, transform)`
4. Piping/Chaining `%>%` : connect manipulations into a single call

- <http://www.sharpsightlabs.com/dplyr-intro-data-manipulation-with-r/>
- <https://rpubs.com/justmarkham/dplyr-tutorial>
- <https://rpubs.com/mkapur/ModelingProject>

Future Topics

4. SQL functions
 - Connecting to databases (<http://cran.r-project.org/web/packages/dplyr/vignettes/databases.html>)
<https://github.com/datacarpentry/R-dplyr-ecology/blob/gh-pages/03-data-analysis.Rmd> <https://github.com/justmarkham/dplyr-tutorial/blob/master/dplyr-tutorial.Rmd>
 - Manipulating data from DBs
5. Two-table joins : multi-table data manipulation (<http://cran.r-project.org/web/packages/dplyr/vignettes/two-table.html>)
6. Data Cleaning (<http://stackoverflow.com/questions/23642811/replace-parts-of-a-variable-using-numeric-indices-in-dplyr-rq=1>)

1. `rename()` - rename columns

```
# Let's play with the iris dataset for a bit, RA Fisher first analyzed these data in 1936  
# first print the names of the columns
```

```
data(iris)  
iris
```

##	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
## 1	5.1	3.5	1.4	0.2	setosa
## 2	4.9	3.0	1.4	0.2	setosa
## 3	4.7	3.2	1.3	0.2	setosa
## 4	4.6	3.1	1.5	0.2	setosa
## 5	5.0	3.6	1.4	0.2	setosa
## 6	5.4	3.9	1.7	0.4	setosa
## 7	4.6	3.4	1.4	0.3	setosa
## 8	5.0	3.4	1.5	0.2	setosa
## 9	4.4	2.9	1.4	0.2	setosa

## 10	4.9	3.1	1.5	0.1	setosa
## 11	5.4	3.7	1.5	0.2	setosa
## 12	4.8	3.4	1.6	0.2	setosa
## 13	4.8	3.0	1.4	0.1	setosa
## 14	4.3	3.0	1.1	0.1	setosa
## 15	5.8	4.0	1.2	0.2	setosa
## 16	5.7	4.4	1.5	0.4	setosa
## 17	5.4	3.9	1.3	0.4	setosa
## 18	5.1	3.5	1.4	0.3	setosa
## 19	5.7	3.8	1.7	0.3	setosa
## 20	5.1	3.8	1.5	0.3	setosa
## 21	5.4	3.4	1.7	0.2	setosa
## 22	5.1	3.7	1.5	0.4	setosa
## 23	4.6	3.6	1.0	0.2	setosa
## 24	5.1	3.3	1.7	0.5	setosa
## 25	4.8	3.4	1.9	0.2	setosa
## 26	5.0	3.0	1.6	0.2	setosa
## 27	5.0	3.4	1.6	0.4	setosa
## 28	5.2	3.5	1.5	0.2	setosa
## 29	5.2	3.4	1.4	0.2	setosa
## 30	4.7	3.2	1.6	0.2	setosa
## 31	4.8	3.1	1.6	0.2	setosa
## 32	5.4	3.4	1.5	0.4	setosa
## 33	5.2	4.1	1.5	0.1	setosa
## 34	5.5	4.2	1.4	0.2	setosa
## 35	4.9	3.1	1.5	0.2	setosa
## 36	5.0	3.2	1.2	0.2	setosa
## 37	5.5	3.5	1.3	0.2	setosa
## 38	4.9	3.6	1.4	0.1	setosa
## 39	4.4	3.0	1.3	0.2	setosa
## 40	5.1	3.4	1.5	0.2	setosa
## 41	5.0	3.5	1.3	0.3	setosa
## 42	4.5	2.3	1.3	0.3	setosa
## 43	4.4	3.2	1.3	0.2	setosa
## 44	5.0	3.5	1.6	0.6	setosa
## 45	5.1	3.8	1.9	0.4	setosa
## 46	4.8	3.0	1.4	0.3	setosa
## 47	5.1	3.8	1.6	0.2	setosa
## 48	4.6	3.2	1.4	0.2	setosa
## 49	5.3	3.7	1.5	0.2	setosa
## 50	5.0	3.3	1.4	0.2	setosa
## 51	7.0	3.2	4.7	1.4	versicolor
## 52	6.4	3.2	4.5	1.5	versicolor
## 53	6.9	3.1	4.9	1.5	versicolor
## 54	5.5	2.3	4.0	1.3	versicolor
## 55	6.5	2.8	4.6	1.5	versicolor
## 56	5.7	2.8	4.5	1.3	versicolor
## 57	6.3	3.3	4.7	1.6	versicolor
## 58	4.9	2.4	3.3	1.0	versicolor
## 59	6.6	2.9	4.6	1.3	versicolor
## 60	5.2	2.7	3.9	1.4	versicolor
## 61	5.0	2.0	3.5	1.0	versicolor
## 62	5.9	3.0	4.2	1.5	versicolor
## 63	6.0	2.2	4.0	1.0	versicolor

## 64	6.1	2.9	4.7	1.4 versicolor
## 65	5.6	2.9	3.6	1.3 versicolor
## 66	6.7	3.1	4.4	1.4 versicolor
## 67	5.6	3.0	4.5	1.5 versicolor
## 68	5.8	2.7	4.1	1.0 versicolor
## 69	6.2	2.2	4.5	1.5 versicolor
## 70	5.6	2.5	3.9	1.1 versicolor
## 71	5.9	3.2	4.8	1.8 versicolor
## 72	6.1	2.8	4.0	1.3 versicolor
## 73	6.3	2.5	4.9	1.5 versicolor
## 74	6.1	2.8	4.7	1.2 versicolor
## 75	6.4	2.9	4.3	1.3 versicolor
## 76	6.6	3.0	4.4	1.4 versicolor
## 77	6.8	2.8	4.8	1.4 versicolor
## 78	6.7	3.0	5.0	1.7 versicolor
## 79	6.0	2.9	4.5	1.5 versicolor
## 80	5.7	2.6	3.5	1.0 versicolor
## 81	5.5	2.4	3.8	1.1 versicolor
## 82	5.5	2.4	3.7	1.0 versicolor
## 83	5.8	2.7	3.9	1.2 versicolor
## 84	6.0	2.7	5.1	1.6 versicolor
## 85	5.4	3.0	4.5	1.5 versicolor
## 86	6.0	3.4	4.5	1.6 versicolor
## 87	6.7	3.1	4.7	1.5 versicolor
## 88	6.3	2.3	4.4	1.3 versicolor
## 89	5.6	3.0	4.1	1.3 versicolor
## 90	5.5	2.5	4.0	1.3 versicolor
## 91	5.5	2.6	4.4	1.2 versicolor
## 92	6.1	3.0	4.6	1.4 versicolor
## 93	5.8	2.6	4.0	1.2 versicolor
## 94	5.0	2.3	3.3	1.0 versicolor
## 95	5.6	2.7	4.2	1.3 versicolor
## 96	5.7	3.0	4.2	1.2 versicolor
## 97	5.7	2.9	4.2	1.3 versicolor
## 98	6.2	2.9	4.3	1.3 versicolor
## 99	5.1	2.5	3.0	1.1 versicolor
## 100	5.7	2.8	4.1	1.3 versicolor
## 101	6.3	3.3	6.0	2.5 virginica
## 102	5.8	2.7	5.1	1.9 virginica
## 103	7.1	3.0	5.9	2.1 virginica
## 104	6.3	2.9	5.6	1.8 virginica
## 105	6.5	3.0	5.8	2.2 virginica
## 106	7.6	3.0	6.6	2.1 virginica
## 107	4.9	2.5	4.5	1.7 virginica
## 108	7.3	2.9	6.3	1.8 virginica
## 109	6.7	2.5	5.8	1.8 virginica
## 110	7.2	3.6	6.1	2.5 virginica
## 111	6.5	3.2	5.1	2.0 virginica
## 112	6.4	2.7	5.3	1.9 virginica
## 113	6.8	3.0	5.5	2.1 virginica
## 114	5.7	2.5	5.0	2.0 virginica
## 115	5.8	2.8	5.1	2.4 virginica
## 116	6.4	3.2	5.3	2.3 virginica
## 117	6.5	3.0	5.5	1.8 virginica

```
## 118      7.7      3.8      6.7      2.2 virginica
## 119      7.7      2.6      6.9      2.3 virginica
## 120      6.0      2.2      5.0      1.5 virginica
## 121      6.9      3.2      5.7      2.3 virginica
## 122      5.6      2.8      4.9      2.0 virginica
## 123      7.7      2.8      6.7      2.0 virginica
## 124      6.3      2.7      4.9      1.8 virginica
## 125      6.7      3.3      5.7      2.1 virginica
## 126      7.2      3.2      6.0      1.8 virginica
## 127      6.2      2.8      4.8      1.8 virginica
## 128      6.1      3.0      4.9      1.8 virginica
## 129      6.4      2.8      5.6      2.1 virginica
## 130      7.2      3.0      5.8      1.6 virginica
## 131      7.4      2.8      6.1      1.9 virginica
## 132      7.9      3.8      6.4      2.0 virginica
## 133      6.4      2.8      5.6      2.2 virginica
## 134      6.3      2.8      5.1      1.5 virginica
## 135      6.1      2.6      5.6      1.4 virginica
## 136      7.7      3.0      6.1      2.3 virginica
## 137      6.3      3.4      5.6      2.4 virginica
## 138      6.4      3.1      5.5      1.8 virginica
## 139      6.0      3.0      4.8      1.8 virginica
## 140      6.9      3.1      5.4      2.1 virginica
## 141      6.7      3.1      5.6      2.4 virginica
## 142      6.9      3.1      5.1      2.3 virginica
## 143      5.8      2.7      5.1      1.9 virginica
## 144      6.8      3.2      5.9      2.3 virginica
## 145      6.7      3.3      5.7      2.5 virginica
## 146      6.7      3.0      5.2      2.3 virginica
## 147      6.3      2.5      5.0      1.9 virginica
## 148      6.5      3.0      5.2      2.0 virginica
## 149      6.2      3.4      5.4      2.3 virginica
## 150      5.9      3.0      5.1      1.8 virginica
```

```
# list the column names first, so you know what you're working with
names(iris)
```

```
## [1] "Sepal.Length" "Sepal.Width"  "Petal.Length" "Petal.Width"
## [5] "Species"
```

```
# The periods in each column name are kind of annoying, let's fix that
rename(iris, petLength = Petal.Length,
  petWidth = Petal.Width,
  sepWidth = Sepal.Width,
  sepLength = Sepal.Length) -> iris2 # turn the assignment around, remember this for piping later
```

```
# rename and select have special arguments for finding columns
select(iris, matches("sepal", ignore.case = FALSE)) %>% head # case sensitive
```

```
## data frame with 0 columns and 6 rows
```

```
select(iris, matches("sepal")) %>% head # not case sensitive
```

```
##   Sepal.Length Sepal.Width
## 1         5.1         3.5
## 2         4.9         3.0
## 3         4.7         3.2
## 4         4.6         3.1
## 5         5.0         3.6
## 6         5.4         3.9
```

```
# another example - notice that it is not case sensitive
```

```
select(iris, contains("l.l")) %>% head
```

```
##   Sepal.Length Petal.Length
## 1         5.1         1.4
## 2         4.9         1.4
## 3         4.7         1.3
## 4         4.6         1.5
## 5         5.0         1.4
## 6         5.4         1.7
```

```
# similar to a regex function i.e. stringr::sub_str()
```

```
select(iris, ends_with("th")) %>% head # the regex starts at the end of a string; if end of string does
```

```
##   Sepal.Length Sepal.Width Petal.Length Petal.Width
## 1         5.1         3.5         1.4         0.2
## 2         4.9         3.0         1.4         0.2
## 3         4.7         3.2         1.3         0.2
## 4         4.6         3.1         1.5         0.2
## 5         5.0         3.6         1.4         0.2
## 6         5.4         3.9         1.7         0.4
```

```
# what if you have lots of columns, named/numbered something?
```

```
# create a silly example
```

```
df1 = data.frame(x05 = rnorm(5), x06= runif(5), x07 = rpois(5, 2), type = sample(letters, 5))
```

```
# now select a few columns
```

```
select(df1, num_range("x", 5:6, width=2)) # the width argument specifies that the string is at least 2
```

```
##           x05           x06
## 1 0.06081869 0.2043535
## 2 0.14850527 0.1432178
## 3 -0.94231104 0.8971633
## 4 1.02005376 0.5503973
## 5 -0.84575074 0.2712184
```

2. arrange() - re-order rows by column

```
# arrange(dataframe, column1, column2, ...)
```

```
head(arrange(iris2, desc(Species), sepLength))
```

```
##      sepLength sepWidth petLength petWidth  Species
## 1      4.9      2.5      4.5      1.7 virginica
## 2      5.6      2.8      4.9      2.0 virginica
## 3      5.7      2.5      5.0      2.0 virginica
## 4      5.8      2.7      5.1      1.9 virginica
## 5      5.8      2.8      5.1      2.4 virginica
## 6      5.8      2.7      5.1      1.9 virginica
```

```
# the piping way:
arrange(iris2, desc(Species), sepLength) %>% head
```

```
##      sepLength sepWidth petLength petWidth  Species
## 1      4.9      2.5      4.5      1.7 virginica
## 2      5.6      2.8      4.9      2.0 virginica
## 3      5.7      2.5      5.0      2.0 virginica
## 4      5.8      2.7      5.1      1.9 virginica
## 5      5.8      2.8      5.1      2.4 virginica
## 6      5.8      2.7      5.1      1.9 virginica
```

3. group_by() - equivalent to plyr::ddply()

```
library(plyr)
```

```
## -----
## You have loaded plyr after dplyr - this is likely to cause problems.
## If you need functions from both plyr and dplyr, please load plyr first, then dplyr:
## library(plyr); library(dplyr)
## -----
##
## Attaching package: 'plyr'
##
## The following objects are masked from 'package:dplyr':
##
##      arrange, count, desc, failwith, id, mutate, rename, summarise,
##      summarize
```

```
# mutate iris2, replace iris2 with mutated data
iris2 %>% mutate(sepalRatio = sepWidth/sepLength, # a simple ratio
                avgSepLen = mean(sepLength)) -> iris2 # can use mutate to do column-wise operations to

# transmute only outputs new columns
transmute(iris2, sepalRatio = sepWidth/sepLength)
```

```
##      sepalRatio
## 1      0.6862745
## 2      0.6122449
## 3      0.6808511
## 4      0.6739130
## 5      0.7200000
## 6      0.7222222
```

## 7	0.7391304
## 8	0.6800000
## 9	0.6590909
## 10	0.6326531
## 11	0.6851852
## 12	0.7083333
## 13	0.6250000
## 14	0.6976744
## 15	0.6896552
## 16	0.7719298
## 17	0.7222222
## 18	0.6862745
## 19	0.6666667
## 20	0.7450980
## 21	0.6296296
## 22	0.7254902
## 23	0.7826087
## 24	0.6470588
## 25	0.7083333
## 26	0.6000000
## 27	0.6800000
## 28	0.6730769
## 29	0.6538462
## 30	0.6808511
## 31	0.6458333
## 32	0.6296296
## 33	0.7884615
## 34	0.7636364
## 35	0.6326531
## 36	0.6400000
## 37	0.6363636
## 38	0.7346939
## 39	0.6818182
## 40	0.6666667
## 41	0.7000000
## 42	0.5111111
## 43	0.7272727
## 44	0.7000000
## 45	0.7450980
## 46	0.6250000
## 47	0.7450980
## 48	0.6956522
## 49	0.6981132
## 50	0.6600000
## 51	0.4571429
## 52	0.5000000
## 53	0.4492754
## 54	0.4181818
## 55	0.4307692
## 56	0.4912281
## 57	0.5238095
## 58	0.4897959
## 59	0.4393939
## 60	0.5192308

## 61	0.4000000
## 62	0.5084746
## 63	0.3666667
## 64	0.4754098
## 65	0.5178571
## 66	0.4626866
## 67	0.5357143
## 68	0.4655172
## 69	0.3548387
## 70	0.4464286
## 71	0.5423729
## 72	0.4590164
## 73	0.3968254
## 74	0.4590164
## 75	0.4531250
## 76	0.4545455
## 77	0.4117647
## 78	0.4477612
## 79	0.4833333
## 80	0.4561404
## 81	0.4363636
## 82	0.4363636
## 83	0.4655172
## 84	0.4500000
## 85	0.5555556
## 86	0.5666667
## 87	0.4626866
## 88	0.3650794
## 89	0.5357143
## 90	0.4545455
## 91	0.4727273
## 92	0.4918033
## 93	0.4482759
## 94	0.4600000
## 95	0.4821429
## 96	0.5263158
## 97	0.5087719
## 98	0.4677419
## 99	0.4901961
## 100	0.4912281
## 101	0.5238095
## 102	0.4655172
## 103	0.4225352
## 104	0.4603175
## 105	0.4615385
## 106	0.3947368
## 107	0.5102041
## 108	0.3972603
## 109	0.3731343
## 110	0.5000000
## 111	0.4923077
## 112	0.4218750
## 113	0.4411765
## 114	0.4385965


```
## 115 0.4827586
## 116 0.5000000
## 117 0.4615385
## 118 0.4935065
## 119 0.3376623
## 120 0.3666667
## 121 0.4637681
## 122 0.5000000
## 123 0.3636364
## 124 0.4285714
## 125 0.4925373
## 126 0.4444444
## 127 0.4516129
## 128 0.4918033
## 129 0.4375000
## 130 0.4166667
## 131 0.3783784
## 132 0.4810127
## 133 0.4375000
## 134 0.4444444
## 135 0.4262295
## 136 0.3896104
## 137 0.5396825
## 138 0.4843750
## 139 0.5000000
## 140 0.4492754
## 141 0.4626866
## 142 0.4492754
## 143 0.4655172
## 144 0.4705882
## 145 0.4925373
## 146 0.4477612
## 147 0.3968254
## 148 0.4615385
## 149 0.5483871
## 150 0.5084746
```

```
# group the data by column "Species" - summarise by mean sepal and petal lengths
iris2 %>% group_by(Species) %>% summarise(mean(sepLength), mean(petLength))
```

```
##      mean(sepLength) mean(petLength)
## 1          5.843333          3.758
```

```
# how you would do this in plyr; I think it makes more sense to use a single function call
# syntax: ddply(dataframe, groups, special function, function calls)
head(ddply(iris2, .(Species), summarize,
           avgSepLen = mean(sepLength)))
```

```
##      Species avgSepLen
## 1      setosa      5.006
## 2 versicolor      5.936
## 3  virginica      6.588
```

```
# transform adds column(s) all at once
head(ddply(iris2, .(Species), transform,
  avgSepLen = mean(sepLength)))
```

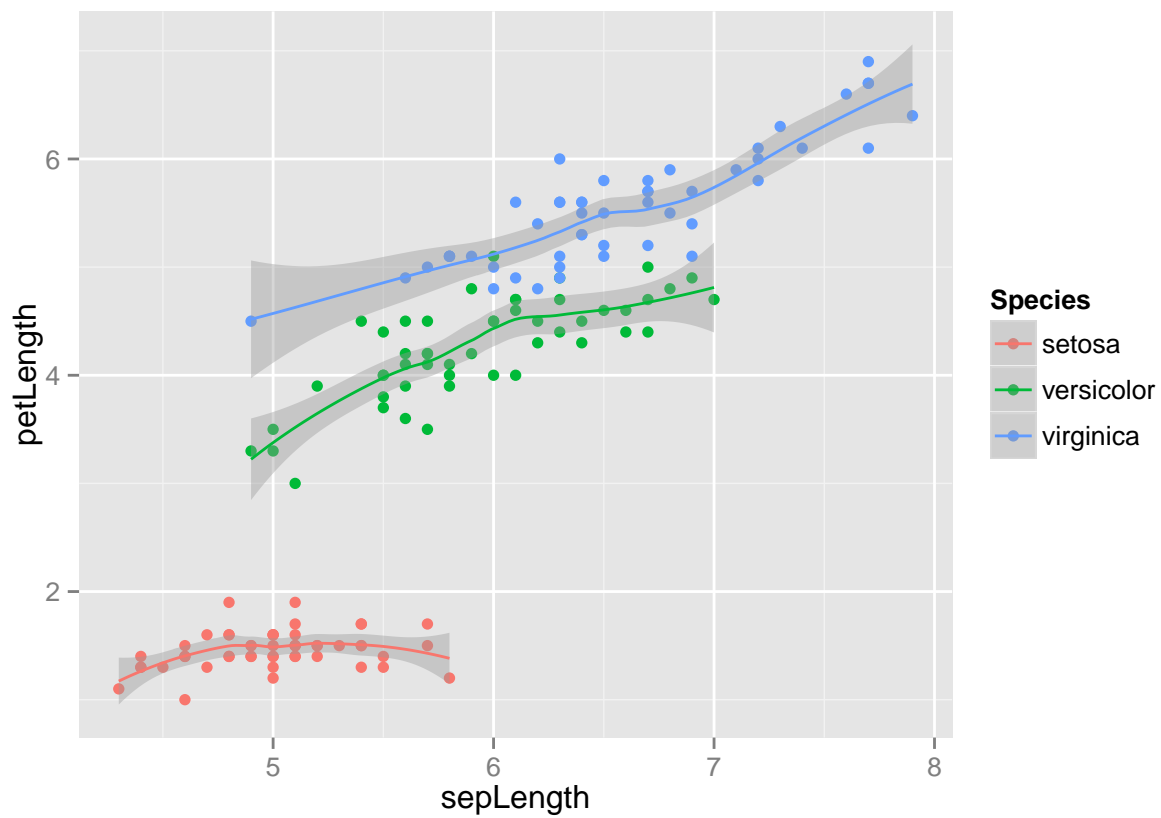
```
##   sepLength sepWidth petLength petWidth Species sepalRatio avgSepLen
## 1      5.1      3.5      1.4      0.2   setosa  0.6862745      5.006
## 2      4.9      3.0      1.4      0.2   setosa  0.6122449      5.006
## 3      4.7      3.2      1.3      0.2   setosa  0.6808511      5.006
## 4      4.6      3.1      1.5      0.2   setosa  0.6739130      5.006
## 5      5.0      3.6      1.4      0.2   setosa  0.7200000      5.006
## 6      5.4      3.9      1.7      0.4   setosa  0.7222222      5.006
```

```
# mutate lets you adds column(s) sequentially, so they can be referenced internally within the function
head(ddply(iris2, .(Species), mutate,
  avgSepLen = mean(sepLength), ratko = avgSepLen*10)) # ratko references the previous column: avgSepLen
```

```
##   sepLength sepWidth petLength petWidth Species sepalRatio avgSepLen ratko
## 1      5.1      3.5      1.4      0.2   setosa  0.6862745      5.006 50.06
## 2      4.9      3.0      1.4      0.2   setosa  0.6122449      5.006 50.06
## 3      4.7      3.2      1.3      0.2   setosa  0.6808511      5.006 50.06
## 4      4.6      3.1      1.5      0.2   setosa  0.6739130      5.006 50.06
## 5      5.0      3.6      1.4      0.2   setosa  0.7200000      5.006 50.06
## 6      5.4      3.9      1.7      0.4   setosa  0.7222222      5.006 50.06
```

```
# can we use a pipe to plot? Si se puede, a few examples below:
# plot the groups on the same panel, using color to identify groups
iris2 %>% group_by(Species) %>% ggplot(aes(sepLength,petLength, color = Species)) + geom_point() + geom_smooth()
```

```
## geom_smooth: method="auto" and size of largest group is <1000, so using loess. Use 'method = x' to control
```



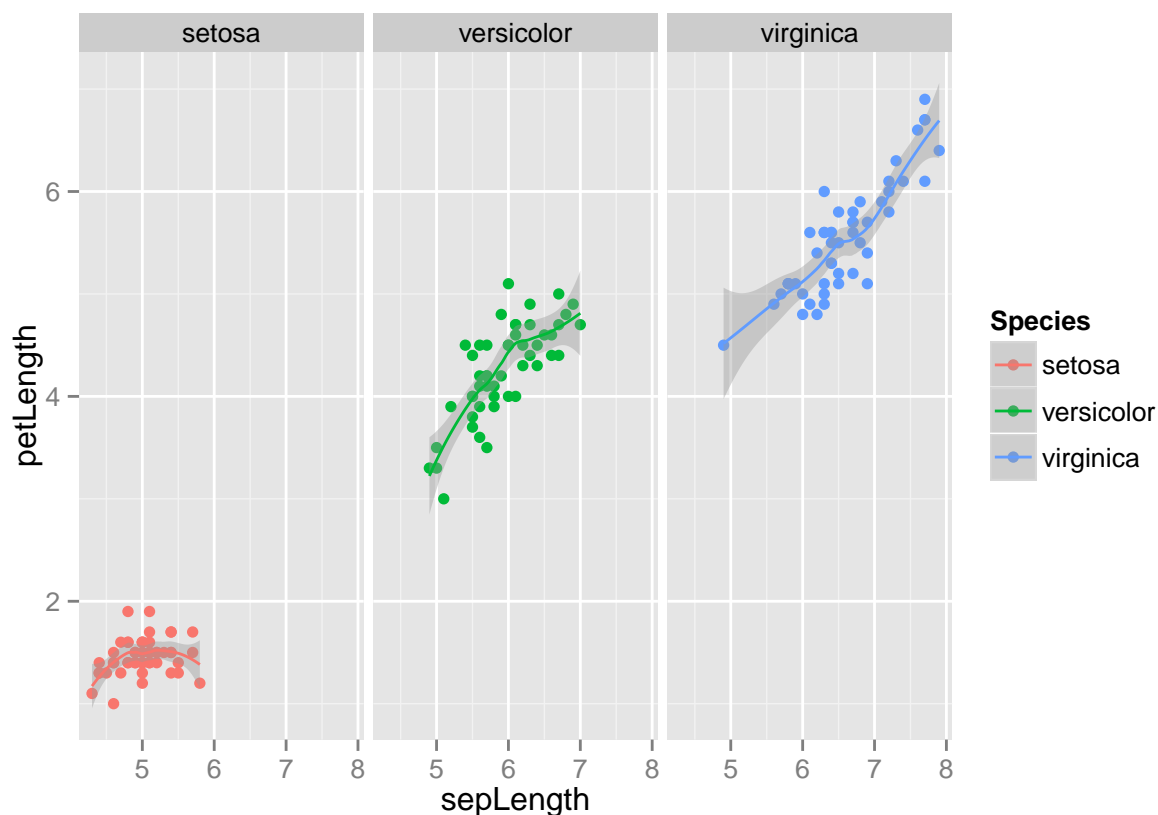
```
# facet_wrap the different groups onto different panels
```

```
iris2 %>% group_by(Species) %>% ggplot(aes(sepLength,petLength, color = Species)) + geom_point() + geom_smooth()
```

```
## geom_smooth: method="auto" and size of largest group is <1000, so using loess. Use 'method = x' to change this
```

```
## geom_smooth: method="auto" and size of largest group is <1000, so using loess. Use 'method = x' to change this
```

```
## geom_smooth: method="auto" and size of largest group is <1000, so using loess. Use 'method = x' to change this
```



```
# sfsmisc as an alternative for plotting?
```

```
# what about modelling; Can we fit a linear model to each group, then examine the summary data? - not s  
iris2 %>% group_by(Species) %>% do(model = lm(sepLength ~ petLength, data = .)) -> iris.mods
```

```
# look at the structure - HOW DO I GET TO YOU?
```

```
str(iris.mods[1,2])
```

```
## Classes 'tbl_df' and 'data.frame':  1 obs. of  1 variable:  
## $ model:List of 1  
## ..$ :List of 12  
## .. ..$ coefficients : Named num  4.213 0.542  
## .. ..- attr(*, "names")= chr  "(Intercept)" "petLength"  
## .. ..$ residuals : Named num  0.1276 -0.0724 -0.2181 -0.4266 0.0276 ...  
## .. ..- attr(*, "names")= chr  "1" "2" "3" "4" ...  
## .. ..$ effects : Named num  -35.3978 0.6592 -0.2232 -0.4456 0.0156 ...  
## .. ..- attr(*, "names")= chr  "(Intercept)" "petLength" "" "" ...  
## .. ..$ rank : int 2  
## .. ..$ fitted.values: Named num  4.97 4.97 4.92 5.03 4.97 ...  
## .. ..- attr(*, "names")= chr  "1" "2" "3" "4" ...  
## .. ..$ assign : int 0 1  
## .. ..$ qr :List of 5  
## .. .. ..$ qr : num [1:50, 1:2] -7.071 0.141 0.141 0.141 0.141 ...  
## .. .. ..- attr(*, "dimnames")=List of 2  
## .. .. .. ..$ : chr  "1" "2" "3" "4" ...  
## .. .. .. ..$ : chr  "(Intercept)" "petLength"
```

```

## .. - attr(*, "assign")= int 0 1
## ..$ graux: num 1.14 1.04
## ..$ pivot: int 1 2
## ..$ tol : num 1e-07
## ..$ rank : int 2
## .. - attr(*, "class")= chr "qr"
## ..$ df.residual : int 48
## ..$ xlevels : Named list()
## ..$ call : language lm(formula = sepLength ~ petLength, data = .)
## ..$ terms :Classes 'terms', 'formula' length 3 sepLength ~ petLength
## .. - attr(*, "variables")= language list(sepLength, petLength)
## .. - attr(*, "factors")= int [1:2, 1] 0 1
## .. - attr(*, "dimnames")=List of 2
## ..$ : chr [1:2] "sepLength" "petLength"
## ..$ : chr "petLength"
## .. - attr(*, "term.labels")= chr "petLength"
## .. - attr(*, "order")= int 1
## .. - attr(*, "intercept")= int 1
## .. - attr(*, "response")= int 1
## .. - attr(*, ".Environment")=<environment: 0x00000000a0f2ff0>
## .. - attr(*, "predvars")= language list(sepLength, petLength)
## .. - attr(*, "dataClasses")= Named chr [1:2] "numeric" "numeric"
## .. - attr(*, "names")= chr [1:2] "sepLength" "petLength"
## ..$ model :'data.frame': 50 obs. of 2 variables:
## ..$ sepLength: num 5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
## ..$ petLength: num 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 ...
## .. - attr(*, "terms")=Classes 'terms', 'formula' length 3 sepLength ~ petLength
## .. - attr(*, "variables")= language list(sepLength, petLength)
## .. - attr(*, "factors")= int [1:2, 1] 0 1
## .. - attr(*, "dimnames")=List of 2
## ..$ : chr [1:2] "sepLength" "petLength"
## ..$ : chr "petLength"
## .. - attr(*, "term.labels")= chr "petLength"
## .. - attr(*, "order")= int 1
## .. - attr(*, "intercept")= int 1
## .. - attr(*, "response")= int 1
## .. - attr(*, ".Environment")=<environment: 0x00000000a0f2ff0>
## .. - attr(*, "predvars")= language list(sepLength, petLength)
## .. - attr(*, "dataClasses")= Named chr [1:2] "numeric" "numeric"
## .. - attr(*, "names")= chr [1:2] "sepLength" "petLength"
## .. - attr(*, "class")= chr "lm"

```