## 5-Homework Using dplyr and ggplot2 BIOL 5000

You will need these libraries:

```
library(ggplot2)
library(dplyr)
```

You will need this data:

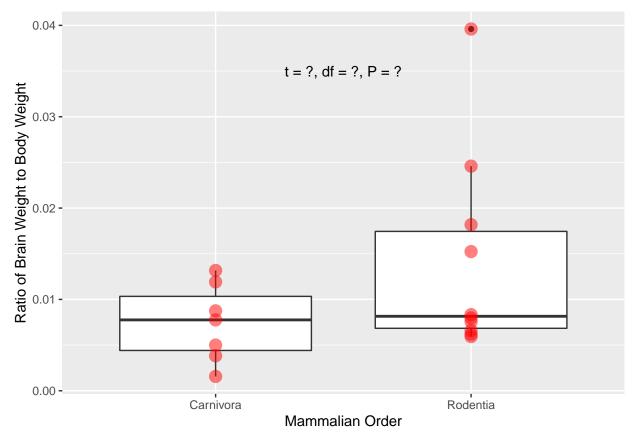
data(msleep)

Question 1. Use dplyr to recreate the summary table below using the msleep dataset (built into ggplot). You will need to use several of the dplyr verbs (functions) we have learned in class.

## # A tibble: 17 x 5						
##		name	order	brainwt	bodywt	ratio
##		<chr></chr>	<chr></chr>	<dbl></dbl>	<dbl></dbl>	<dbl></dbl>
##	1	Jaguar	${\tt Carnivora}$	0.157	100	0.00157
##	2	Gray seal	${\tt Carnivora}$	0.325	85	0.00382
##	3	Dog	${\tt Carnivora}$	0.07	14	0.005
##	4	Red fox	${\tt Carnivora}$	0.0504	4.23	0.0119
##	5	Arctic fox	${\tt Carnivora}$	0.0445	3.38	0.0132
##	6	Domestic cat	${\tt Carnivora}$	0.0256	3.3	0.00776
##	7	Genet	${\tt Carnivora}$	0.0175	2	0.00875
##	8	African giant pouched rat	Rodentia	0.0066	1	0.0066
##	9	Arctic ground squirrel	Rodentia	0.0057	0.92	0.00620
##	10	Guinea pig	Rodentia	0.0055	0.728	0.00755
##	11	Chinchilla	Rodentia	0.0064	0.42	0.0152
##	12	Laboratory rat	Rodentia	0.0019	0.32	0.00594
##	13	Cotton rat	Rodentia	0.00118	0.148	0.00797
##	14	Mole rat	Rodentia	0.003	0.122	0.0246
##	15	Golden hamster	Rodentia	0.001	0.12	0.00833
##	16	Thirteen-lined ground squirrel	Rodentia	0.004	0.101	0.0396
##	17	House mouse	Rodentia	0.0004	0.022	0.0182

**Question 2**. Use a t-test to compare the two mammal orders, Rodentia and Carnivora, in their ratio of brain weight to body weight.

Question 3. Using ggplot2, replicate the following figure. Fill in the values from your t-test (Question 2).



Question 4. Save the plot to your Rproject folder for this exercise as a 6" X 6" .jpg file.

Homework (folder containing .rproj file and all associated input/output files) is due to the D2L Assignments dropbox **before class on November 2nd**. As before, make each of your 'answers' an object. (e.g.) Answer $_3 \leftarrow ggplot(data\_summary, aes(x = ....... and 'print' the name of the object below the specified code (e.g. Answer<math>_3$ ). Each object is then displayed automatically in the console (if a model or table) or plot viewer (if its a graph) without having to click on anything.