

# Midterm 1 W26

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2026-01-29

## Instructions

Answer the following questions and complete the exercises in RMarkdown. Please embed all of your code and push your final work to your repository. Your code must be organized, clean, and run free from errors. Remember, you must remove the `#` for any included code chunks to run. Be sure to add your name to the author header above.

Your code must knit in order to be considered. If you are stuck and cannot answer a question, then comment out your code and knit the document. You may use your notes, labs, and homework to help you complete this exam. Do not use any other resources- including AI assistance or other students' work.

Don't forget to answer any questions that are asked in the prompt! Each question must be coded; it cannot be answered by a sort in a spreadsheet or a written response only.

For all plots you create, a title and clearly labeled axes must be provided. We also expect pipes `%>%` to be used wherever possible.

Be sure to push your completed midterm to your repository and upload the document to Gradescope. This exam is worth 50 points.

Please load the following libraries.

```
library(tidyverse)
library(janitor)
```

## Part 1: Repository

**Question 1. (3 points) Before you start analyzing data, please put a link to your GitHub repository below. Your repository should have a clear README and be well-organized. Add `jmledford3115` and `bryshal` as collaborators to your repository if you haven't already done so.**

Link to repository: <https://github.com/wanghouyushenglhj-web/Bis-015L.git>  
(<https://github.com/wanghouyushenglhj-web/Bis-015L.git>)

## Part 2: Data and Analysis

In the midterm 1 folder there is a second folder called `data`. Inside the `data` folder, there is a `.csv` file called `anolis_dat.csv`. These data came from D. Luke Mahler, Liam J. Revell, Richard E. Glor, Jonathan B. Losos, ECOLOGICAL OPPORTUNITY AND THE RATE OF MORPHOLOGICAL EVOLUTION IN THE DIVERSIFICATION OF

GREATER ANTILLEAN ANOLES, Evolution, Volume 64, Issue 9, 1 September 2010, Pages 2731–2745 (<https://academic.oup.com/evolut/article/64/9/2731/6854302?login=true>). The original research article is included in the data folder.

*Anolis* is a genus of lizards commonly known as anoles. Anoles are found throughout the Americas, but are especially diverse in the Caribbean. The data include morphological measurements for *Anolis* lizards from the islands of the Greater Antilles. These data can be used to study patterns of morphological evolution and adaptation in *Anolis* lizards.

The variables include:

- species : Species name of the anole lizard.
- habitat : Habitat type where the lizard was found.
- hindlimb\_length\_mm : Length of the lizard's hindlimbs (in millimeters).
- tail\_length\_mm : Length of the lizard's tail (in millimeters).
- body\_length\_mm : Length of the lizard's body (in millimeters).
- toepad\_lamellae\_count : Count of lamellae on the lizard's toepads.
- island : Island where the lizard was found.

**Question 2. (2 points) Load the data and store it as an object called `anolis` .**

```
anolis <- read.csv("data/anolis_dat.csv")
```

```
names(anolis)
```

```
## [1] "Species"          "Habitat"
## [3] "Hindlimb.length..mm." "Tail.length..mm."
## [5] "Body.length..mm."   "Toepad.lamellae..count."
## [7] "Island"
```

**Question 3. (2 points) Use a summary function of your choice to get an idea of the structure of the data.**

```
str(anolis)
```

```
## 'data.frame':   52 obs. of  7 variables:
## $ Species      : chr  "A. ahli" "A. alayoni" "A. alfaroi" "A. aliniger"
## ...
## $ Habitat      : chr  "Trunk-ground" "Twig" "Grass-bush" "Trunk-crown" ...
## $ Hindlimb.length..mm. : num  50.5 25.5 26.2 36.8 50.4 ...
## $ Tail.length..mm.   : num  82 54.8 79 84.9 154.4 ...
## $ Body.length..mm.   : num  51.7 41.3 30.9 51.5 72.3 ...
## $ Toepad.lamellae..count.: int  27 31 24 36 41 28 29 28 28 31 ...
## $ Island        : chr  "Cuba" "Cuba" "Cuba" "Hispaniola" ...
```

```
glimpse(anolis)
```

```
## Rows: 52
## Columns: 7
## $ Species      <chr> "A. ahli", "A. alayoni", "A. alfaroi", "A. ali...
## $ Habitat      <chr> "Trunk-ground", "Twig", "Grass-bush", "Trunk-c...
## $ Hindlimb.length..mm. <dbl> 50.46, 25.50, 26.17, 36.80, 50.39, 49.37, 29.9...
## $ Tail.length..mm. <dbl> 81.99, 54.75, 79.00, 84.88, 154.45, 91.01, 106...
## $ Body.length..mm. <dbl> 51.67, 41.32, 30.95, 51.53, 72.32, 51.72, 32.9...
## $ Toepad.lamellae..count. <int> 27, 31, 24, 36, 41, 28, 29, 28, 28, 31, 32, 32...
## $ Island       <chr> "Cuba", "Cuba", "Cuba", "Hispaniola", "Cuba", ...
```

```
summary(anolis)
```

##	Species	Habitat	Hindlimb.length..mm.	Tail.length..mm.
##	Length:52	Length:52	Min. :20.00	Min. : 39.79
##	Class :character	Class :character	1st Qu.:31.70	1st Qu.: 82.19
##	Mode :character	Mode :character	Median :40.01	Median :101.86
##			Mean :40.51	Mean :102.81
##			3rd Qu.:49.27	3rd Qu.:121.92
##			Max. :65.48	Max. :154.45
##	Body.length..mm.	Toepad.lamellae..count.	Island	
##	Min. :29.01	Min. :23.00	Length:52	
##	1st Qu.:40.07	1st Qu.:28.00	Class :character	
##	Median :47.05	Median :29.00	Mode :character	
##	Mean :47.65	Mean :30.54		
##	3rd Qu.:54.53	3rd Qu.:33.00		
##	Max. :72.32	Max. :44.00		

**Question 4. (2 points) Clean the variable names so they are all lowercase and without special characters or spaces. Be sure to use the cleaned data for all subsequent analyses.**

```
anolis <- anolis%>%
  clean_names() %>%
  mutate(across(where(is.character), tolower))
```

```
anolis
```

##	species	habitat	hindlimb_length_mm	tail_length_mm
## 1	a. ahli	trunk-ground	50.46	81.99
## 2	a. alayoni	twig	25.50	54.75
## 3	a. alfaroi	grass-bush	26.17	79.00
## 4	a. aliniger	trunk-crown	36.80	84.88
## 5	a. allisoni	trunk-crown	50.39	154.45
## 6	a. allogus	trunk-ground	49.37	91.01
## 7	a. alumina	grass-bush	29.97	106.40
## 8	a. alutaceus	grass-bush	27.39	94.62
## 9	a. angusticeps	twig	24.37	65.10
## 10	a. armouri	trunk-ground	51.81	101.37
## 11	a. bahorucoensis	grass-bush	38.71	115.82
## 12	a. bremeri	trunk-ground	47.44	114.01
## 13	a. breslini	trunk-ground	49.84	122.85
## 14	a. chlorocyanus	trunk-crown	48.55	142.12
## 15	a. clivicola	grass-bush	34.79	98.75
## 16	a. confusus	trunk-ground	41.63	78.31
## 17	a. cooki	trunk-ground	48.50	134.57
## 18	a. cupeyalensis	grass-bush	22.88	93.90
## 19	a. cyanopleurus	grass-bush	31.38	111.68
## 20	a. dolichocephalus	grass-bush	39.49	131.27
## 21	a. grahami	trunk-crown	48.71	119.78
## 22	a. guafe	trunk-ground	39.09	69.45
## 23	a. guazuma	twig	20.46	39.79
## 24	a. gundlachi	trunk-ground	58.32	136.13
## 25	a. haetianus	trunk-ground	65.48	154.41
## 26	a. hendersoni	grass-bush	38.53	128.81
## 27	a. homolechis	trunk-ground	44.97	90.92
## 28	a. imias	trunk-ground	51.67	104.67
## 29	a. inexpectatus	grass-bush	26.80	101.12
## 30	a. insolitus	twig	26.13	50.77
## 31	a. jubar	trunk-ground	41.08	82.15
## 32	a. krugi	grass-bush	40.53	130.76
## 33	a. lineatopus	trunk-ground	52.71	119.97
## 34	a. longitibialis	trunk-ground	61.92	140.36
## 35	a. macilentus	grass-bush	31.80	97.35
## 36	a. marcanoi	trunk-ground	49.24	123.95
## 37	a. mestrei	trunk-ground	44.93	92.11
## 38	a. opalinus	trunk-crown	35.67	82.15
## 39	a. ophiolepis	grass-bush	26.88	81.13
## 40	a. placidus	twig	20.00	47.96
## 41	a. poncensis	grass-bush	33.39	118.20
## 42	a. porcatus	trunk-crown	44.58	140.26
## 43	a. quadriocellifer	trunk-ground	39.05	82.20
## 44	a. rubribarbus	trunk-ground	50.43	93.94
## 45	a. sagrei	trunk-ground	45.47	111.90
## 46	a. semilineatus	grass-bush	33.99	121.61
## 47	a. shrevei	trunk-ground	43.61	102.53
## 48	a. singularis	trunk-crown	38.42	102.34
## 49	a. strahmi	trunk-ground	65.24	138.87
## 50	a. stratulus	trunk-crown	34.38	77.93
## 51	a. vanidicus	grass-bush	26.45	90.60

## 52	a. whitemani trunk-ground	51.06	115.26
##	body_length_mm toepad_lamellae_count	island	
## 1	51.67	27	cuba
## 2	41.32	31	cuba
## 3	30.95	24	cuba
## 4	51.53	36	hispaniola
## 5	72.32	41	cuba
## 6	51.72	28	cuba
## 7	32.94	29	hispaniola
## 8	31.84	28	cuba
## 9	40.22	28	cuba
## 10	56.11	31	hispaniola
## 11	41.81	32	hispaniola
## 12	55.65	32	cuba
## 13	52.29	29	hispaniola
## 14	65.44	42	hispaniola
## 15	39.03	26	cuba
## 16	46.72	32	cuba
## 17	54.45	33	puerto rico
## 18	29.01	24	cuba
## 19	34.32	26	cuba
## 20	45.34	36	hispaniola
## 21	57.97	38	jamacia
## 22	43.95	29	cuba
## 23	39.24	28	cuba
## 24	59.97	28	puerto rico
## 25	68.19	33	hispaniola
## 26	43.19	34	hispaniola
## 27	51.34	32	cuba
## 28	54.89	28	cuba
## 29	31.28	27	cuba
## 30	40.70	25	hispaniola
## 31	47.38	29	cuba
## 32	44.35	33	puerto rico
## 33	56.50	29	jamacia
## 34	63.29	28	hispaniola
## 35	37.39	28	cuba
## 36	53.79	33	hispaniola
## 37	49.05	30	cuba
## 38	42.27	35	jamacia
## 39	34.59	24	cuba
## 40	39.63	26	hispaniola
## 41	41.52	29	puerto rico
## 42	64.37	44	cuba
## 43	45.03	28	cuba
## 44	53.74	29	cuba
## 45	53.14	33	cuba
## 46	36.68	30	hispaniola
## 47	48.85	28	hispaniola
## 48	52.65	38	hispaniola
## 49	65.36	31	hispaniola
## 50	43.62	34	puerto rico

```
## 51      34.19      23      cuba
## 52      54.77      29  hispaniola
```

**Question 5. (4 points) Convert the `habitat` and `island` variables to factors.**

```
names(anolis)
```

```
## [1] "species"      "habitat"      "hindlimb_length_mm"
## [4] "tail_length_mm" "body_length_mm" "toepad_lamellae_count"
## [7] "island"
```

```
anolis %>%
  mutate(across(c(habitat, island), as.factor))
```

##	species	habitat	hindlimb_length_mm	tail_length_mm
## 1	a. ahli	trunk-ground	50.46	81.99
## 2	a. alayoni	twig	25.50	54.75
## 3	a. alfaroi	grass-bush	26.17	79.00
## 4	a. aliniger	trunk-crown	36.80	84.88
## 5	a. allisoni	trunk-crown	50.39	154.45
## 6	a. allogus	trunk-ground	49.37	91.01
## 7	a. alumina	grass-bush	29.97	106.40
## 8	a. alutaceus	grass-bush	27.39	94.62
## 9	a. angusticeps	twig	24.37	65.10
## 10	a. armouri	trunk-ground	51.81	101.37
## 11	a. bahorucoensis	grass-bush	38.71	115.82
## 12	a. bremeri	trunk-ground	47.44	114.01
## 13	a. breslini	trunk-ground	49.84	122.85
## 14	a. chlorocyanus	trunk-crown	48.55	142.12
## 15	a. clivicola	grass-bush	34.79	98.75
## 16	a. confusus	trunk-ground	41.63	78.31
## 17	a. cooki	trunk-ground	48.50	134.57
## 18	a. cupeyalensis	grass-bush	22.88	93.90
## 19	a. cyanopleurus	grass-bush	31.38	111.68
## 20	a. dolichocephalus	grass-bush	39.49	131.27
## 21	a. grahami	trunk-crown	48.71	119.78
## 22	a. guafe	trunk-ground	39.09	69.45
## 23	a. guazuma	twig	20.46	39.79
## 24	a. gundlachi	trunk-ground	58.32	136.13
## 25	a. haetianus	trunk-ground	65.48	154.41
## 26	a. hendersoni	grass-bush	38.53	128.81
## 27	a. homolechis	trunk-ground	44.97	90.92
## 28	a. imias	trunk-ground	51.67	104.67
## 29	a. inexpectatus	grass-bush	26.80	101.12
## 30	a. insolitus	twig	26.13	50.77
## 31	a. jubar	trunk-ground	41.08	82.15
## 32	a. krugi	grass-bush	40.53	130.76
## 33	a. lineatopus	trunk-ground	52.71	119.97
## 34	a. longitibialis	trunk-ground	61.92	140.36
## 35	a. macilentus	grass-bush	31.80	97.35
## 36	a. marcanoi	trunk-ground	49.24	123.95
## 37	a. mestrei	trunk-ground	44.93	92.11
## 38	a. opalinus	trunk-crown	35.67	82.15
## 39	a. ophiolepis	grass-bush	26.88	81.13
## 40	a. placidus	twig	20.00	47.96
## 41	a. poncensis	grass-bush	33.39	118.20
## 42	a. porcatus	trunk-crown	44.58	140.26
## 43	a. quadriocellifer	trunk-ground	39.05	82.20
## 44	a. rubribarbus	trunk-ground	50.43	93.94
## 45	a. sagrei	trunk-ground	45.47	111.90
## 46	a. semilineatus	grass-bush	33.99	121.61
## 47	a. shrevei	trunk-ground	43.61	102.53
## 48	a. singularis	trunk-crown	38.42	102.34
## 49	a. strahmi	trunk-ground	65.24	138.87
## 50	a. stratulus	trunk-crown	34.38	77.93
## 51	a. vanidicus	grass-bush	26.45	90.60

## 52	a. whitemani trunk-ground	51.06	115.26
##	body_length_mm toepad_lamellae_count	island	
## 1	51.67	27	cuba
## 2	41.32	31	cuba
## 3	30.95	24	cuba
## 4	51.53	36	hispaniola
## 5	72.32	41	cuba
## 6	51.72	28	cuba
## 7	32.94	29	hispaniola
## 8	31.84	28	cuba
## 9	40.22	28	cuba
## 10	56.11	31	hispaniola
## 11	41.81	32	hispaniola
## 12	55.65	32	cuba
## 13	52.29	29	hispaniola
## 14	65.44	42	hispaniola
## 15	39.03	26	cuba
## 16	46.72	32	cuba
## 17	54.45	33	puerto rico
## 18	29.01	24	cuba
## 19	34.32	26	cuba
## 20	45.34	36	hispaniola
## 21	57.97	38	jamacia
## 22	43.95	29	cuba
## 23	39.24	28	cuba
## 24	59.97	28	puerto rico
## 25	68.19	33	hispaniola
## 26	43.19	34	hispaniola
## 27	51.34	32	cuba
## 28	54.89	28	cuba
## 29	31.28	27	cuba
## 30	40.70	25	hispaniola
## 31	47.38	29	cuba
## 32	44.35	33	puerto rico
## 33	56.50	29	jamacia
## 34	63.29	28	hispaniola
## 35	37.39	28	cuba
## 36	53.79	33	hispaniola
## 37	49.05	30	cuba
## 38	42.27	35	jamacia
## 39	34.59	24	cuba
## 40	39.63	26	hispaniola
## 41	41.52	29	puerto rico
## 42	64.37	44	cuba
## 43	45.03	28	cuba
## 44	53.74	29	cuba
## 45	53.14	33	cuba
## 46	36.68	30	hispaniola
## 47	48.85	28	hispaniola
## 48	52.65	38	hispaniola
## 49	65.36	31	hispaniola
## 50	43.62	34	puerto rico



```
## 51          34.19          23          cuba
## 52          54.77          29  hispaniola
```

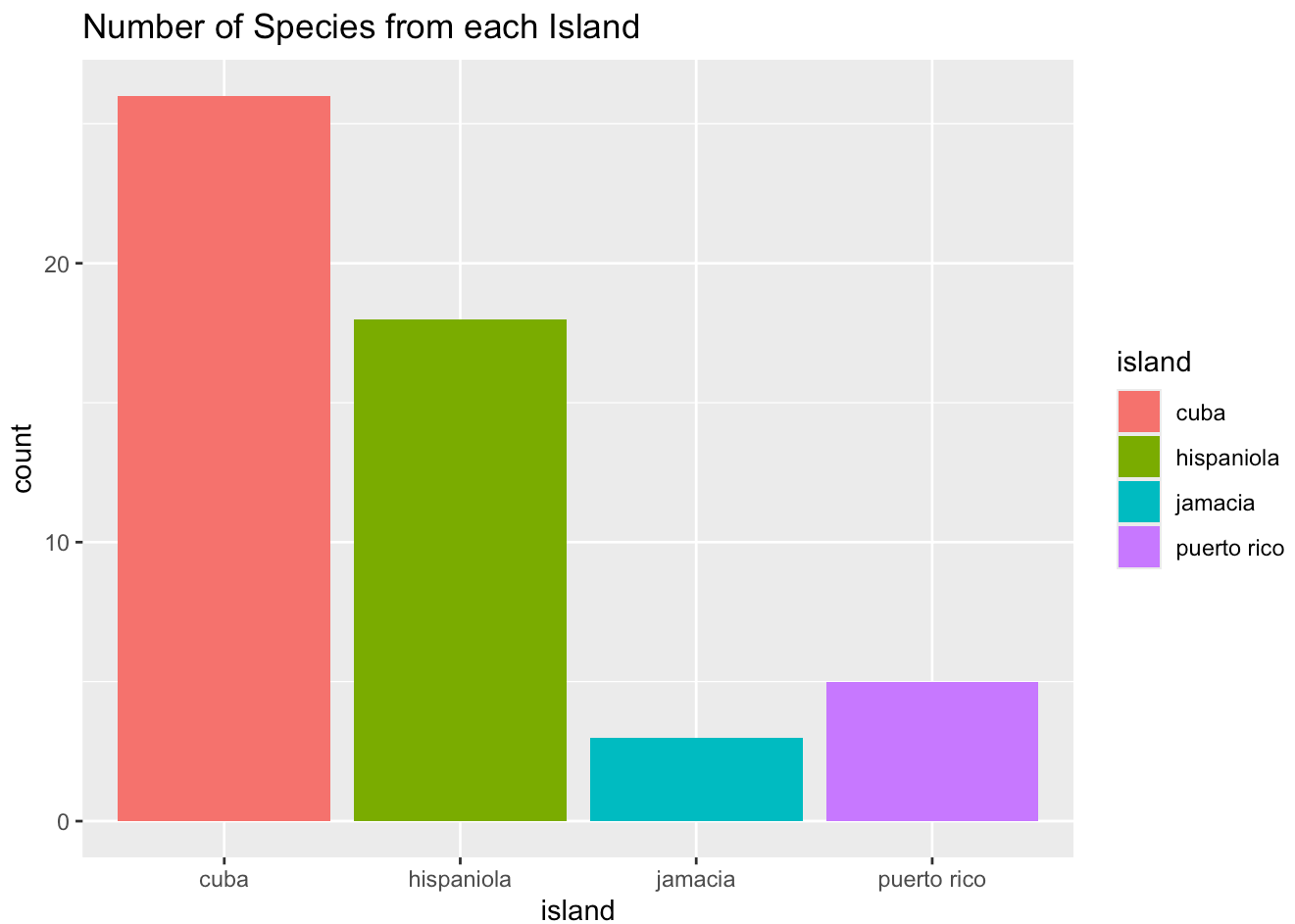
**Question 6. (2 points)** Anole species were sampled from multiple islands. Which islands are represented in the data? Display the island names.

```
anolis %>%
  count(island)#Cuba is represented island in the data.
```

```
##      island  n
## 1      cuba 26
## 2  hispaniola 18
## 3    jamacia  3
## 4  puerto rico  5
```

**Question 7. (4 points)** Is sampling equal across islands? Create a plot to visualize the number of anole species sampled from each island. Be sure to label your axes and add a title.

```
anolis %>%
  ggplot(mapping=aes(x=island))+
  geom_bar(mapping = aes(fill=island))+
  labs(title="Number of Species from each Island")
```



**Question 8. (2 points)** Which habitat types are represented in the data? Display the names of the habitat types.

```
anolis %>%
  count(habitat) %>%
  arrange(desc(n))#trunk_ground represented in the data for habitat.
```

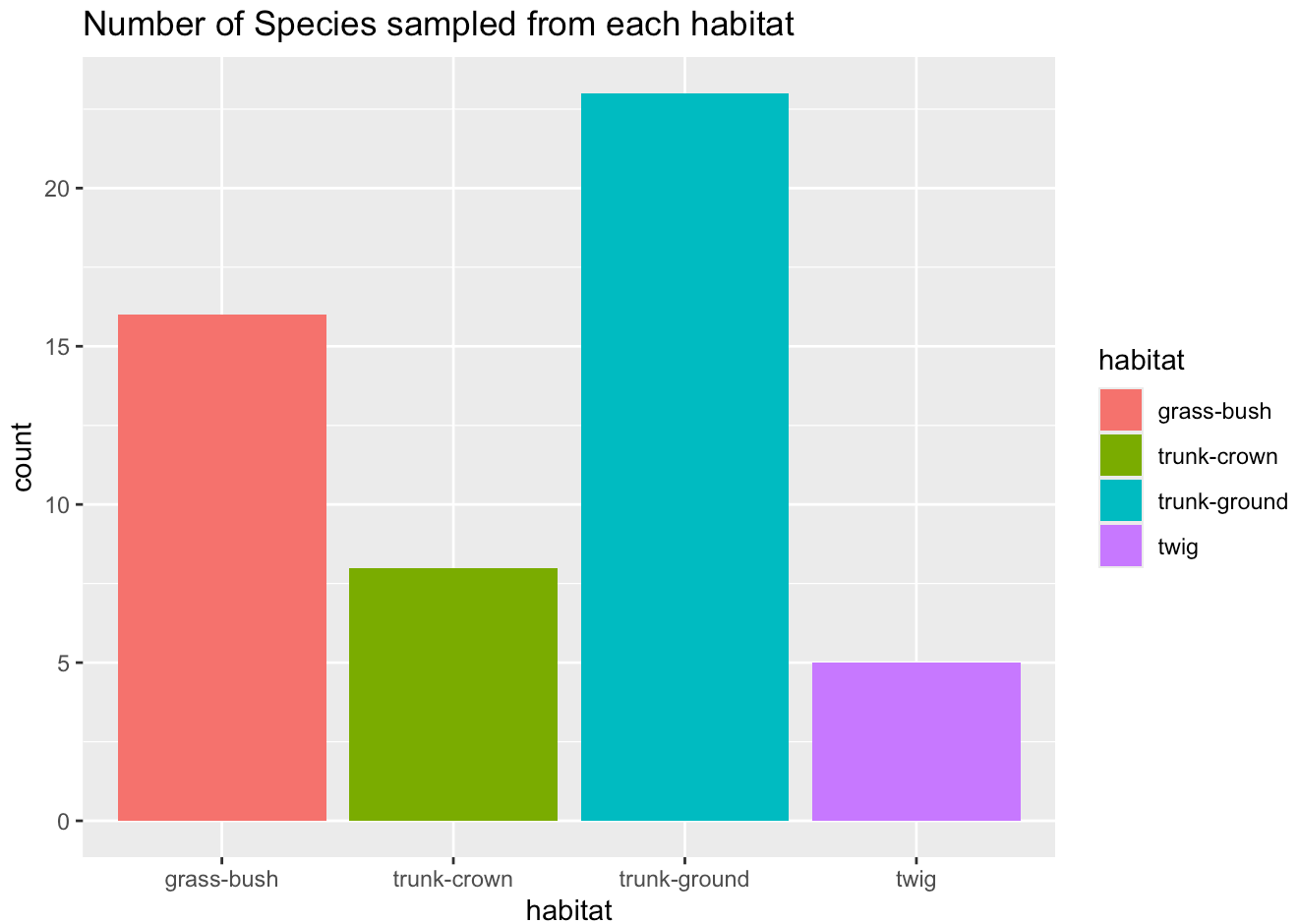
```
##      habitat  n
## 1 trunk-ground 23
## 2  grass-bush 16
## 3 trunk-crown  8
## 4      twig   5
```

**Question 9. (4 points)** Is sampling equal across habitat types? Create a plot to visualize the number of anole species sampled from each habitat type. Be sure to label your axes and add a title.

```
names(anolis)
```

```
## [1] "species"          "habitat"          "hindlimb_length_mm"
## [4] "tail_length_mm"   "body_length_mm"   "toepad_lamellae_count"
## [7] "island"
```

```
anolis %>%
  ggplot(mapping=aes(x=habitat))+
  geom_bar(mapping = aes(fill=habitat))+
  labs(title="Number of Species sampled from each habitat")
```

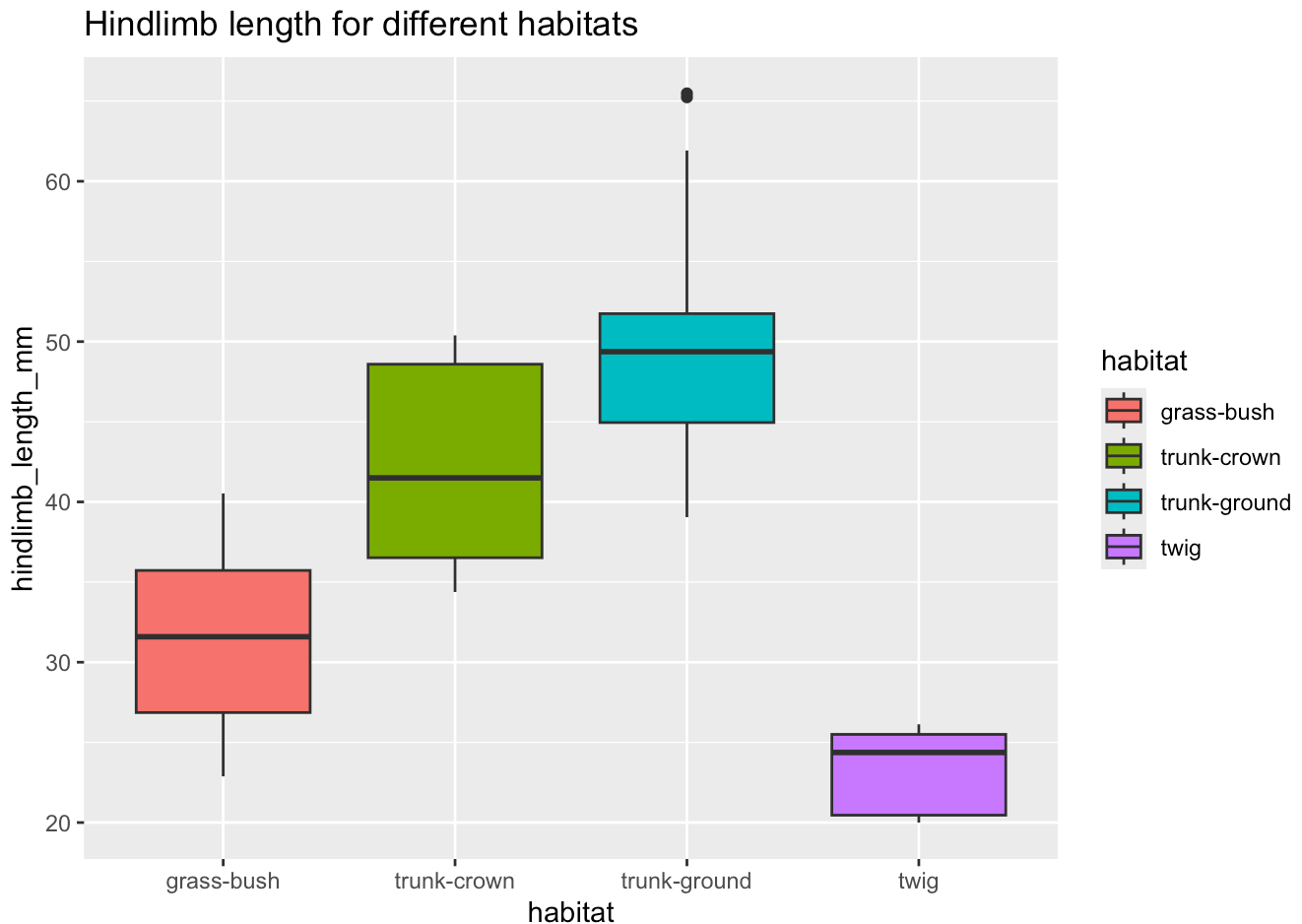


**Question 10. (4 points)** The morphology of anoles varies based on their habitat. How does the range of hindlimb length compare among different habitats? Create a plot to visualize the distribution of hindlimb lengths across habitat types. Be sure to label your axes and add a title.

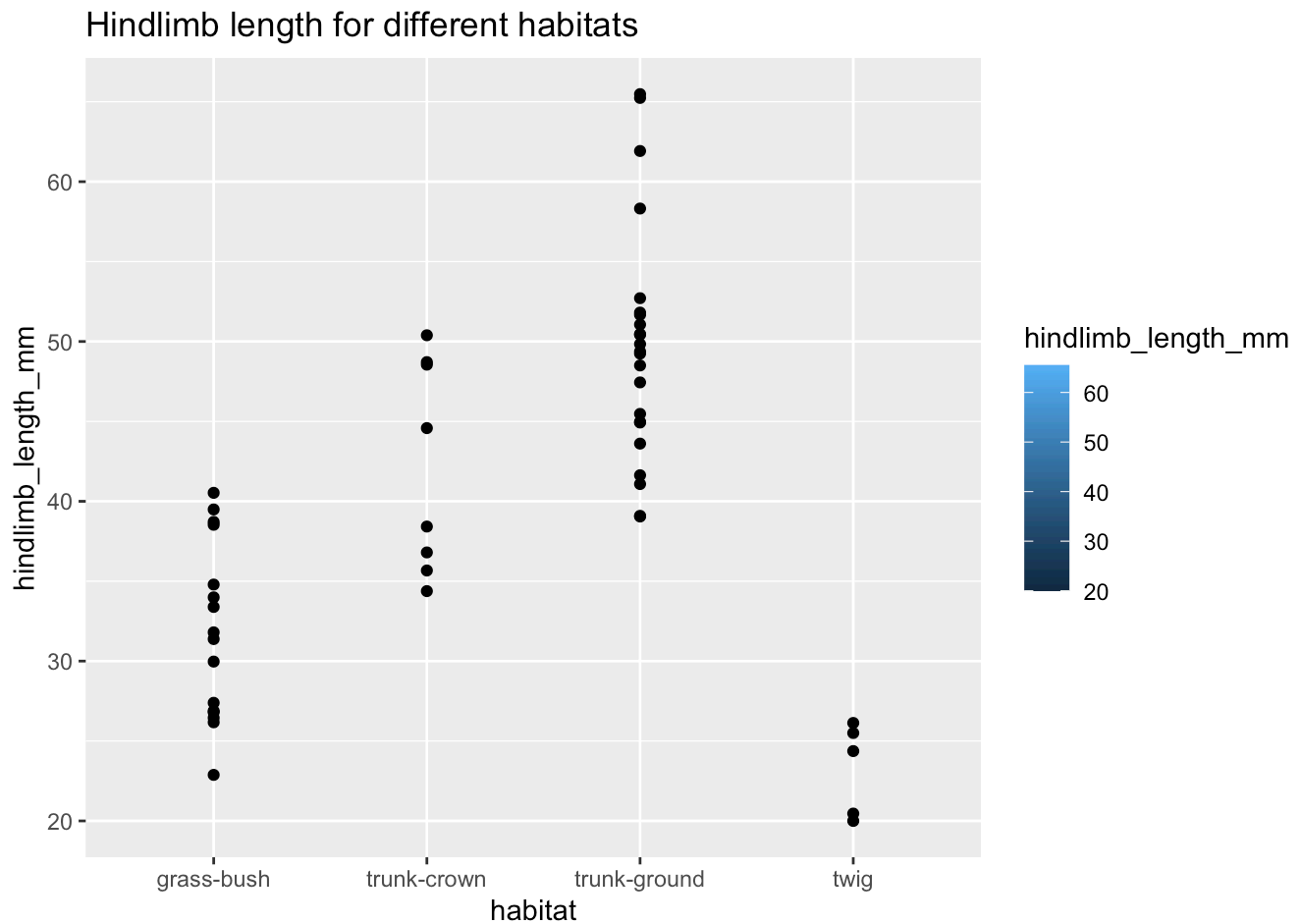
```
names(anolis)
```

```
## [1] "species"          "habitat"          "hindlimb_length_mm"
## [4] "tail_length_mm"   "body_length_mm"   "toepad_lamellae_count"
## [7] "island"
```

```
anolis %>%  
  ggplot(mapping=aes(x=habitat,y=hindlimb_length_mm))+  
  geom_boxplot(mapping = aes(fill=habitat))+  
  labs(title="Hindlimb length for different habitats")
```



```
anolis %>%  
  ggplot(mapping=aes(x=habitat,y=hindlimb_length_mm))+  
  geom_point(mapping = aes(fill=hindlimb_length_mm))+  
  labs(title="Hindlimb length for different habitats")
```



**Question 11. (4 points)** The plot above is compelling, but don't we expect larger lizards to have longer limbs? What about tail length? Shouldn't longer lizards have longer tails? To correct for this, make two new columns: 1. `ratio_of_hindlimb_to_body`, and 2. `ratio_of_tail_to_body`. Don't forget to add these columns to the `anolis` data frame.

```
names(anolis)
```

```
## [1] "species"          "habitat"           "hindlimb_length_mm"
## [4] "tail_length_mm"   "body_length_mm"    "toepad_lamellae_count"
## [7] "island"
```

```
anolis <- anolis %>%
  mutate(ratio_of_hindlimb_to_body=hindlimb_length_mm/body_length_mm,
         ratio_of_tail_to_body=tail_length_mm/body_length_mm)
```

```
anolis
```

##	species	habitat	hindlimb_length_mm	tail_length_mm
## 1	a. ahli	trunk-ground	50.46	81.99
## 2	a. alayoni	twig	25.50	54.75
## 3	a. alfaroi	grass-bush	26.17	79.00
## 4	a. aliniger	trunk-crown	36.80	84.88
## 5	a. allisoni	trunk-crown	50.39	154.45
## 6	a. allogus	trunk-ground	49.37	91.01
## 7	a. alumina	grass-bush	29.97	106.40
## 8	a. alutaceus	grass-bush	27.39	94.62
## 9	a. angusticeps	twig	24.37	65.10
## 10	a. armouri	trunk-ground	51.81	101.37
## 11	a. bahorucoensis	grass-bush	38.71	115.82
## 12	a. bremeri	trunk-ground	47.44	114.01
## 13	a. breslini	trunk-ground	49.84	122.85
## 14	a. chlorocyanus	trunk-crown	48.55	142.12
## 15	a. clivicola	grass-bush	34.79	98.75
## 16	a. confusus	trunk-ground	41.63	78.31
## 17	a. cooki	trunk-ground	48.50	134.57
## 18	a. cupeyalensis	grass-bush	22.88	93.90
## 19	a. cyanopleurus	grass-bush	31.38	111.68
## 20	a. dolichocephalus	grass-bush	39.49	131.27
## 21	a. grahami	trunk-crown	48.71	119.78
## 22	a. guafe	trunk-ground	39.09	69.45
## 23	a. guazuma	twig	20.46	39.79
## 24	a. gundlachi	trunk-ground	58.32	136.13
## 25	a. haetianus	trunk-ground	65.48	154.41
## 26	a. hendersoni	grass-bush	38.53	128.81
## 27	a. homolechis	trunk-ground	44.97	90.92
## 28	a. imias	trunk-ground	51.67	104.67
## 29	a. inexpectatus	grass-bush	26.80	101.12
## 30	a. insolitus	twig	26.13	50.77
## 31	a. jubar	trunk-ground	41.08	82.15
## 32	a. krugi	grass-bush	40.53	130.76
## 33	a. lineatopus	trunk-ground	52.71	119.97
## 34	a. longitibialis	trunk-ground	61.92	140.36
## 35	a. macilentus	grass-bush	31.80	97.35
## 36	a. marcanoi	trunk-ground	49.24	123.95
## 37	a. mestrei	trunk-ground	44.93	92.11
## 38	a. opalinus	trunk-crown	35.67	82.15
## 39	a. ophiolepis	grass-bush	26.88	81.13
## 40	a. placidus	twig	20.00	47.96
## 41	a. poncensis	grass-bush	33.39	118.20
## 42	a. porcatus	trunk-crown	44.58	140.26
## 43	a. quadriocellifer	trunk-ground	39.05	82.20
## 44	a. rubribarbus	trunk-ground	50.43	93.94
## 45	a. sagrei	trunk-ground	45.47	111.90
## 46	a. semilineatus	grass-bush	33.99	121.61
## 47	a. shrevei	trunk-ground	43.61	102.53
## 48	a. singularis	trunk-crown	38.42	102.34
## 49	a. strahmi	trunk-ground	65.24	138.87
## 50	a. stratulus	trunk-crown	34.38	77.93
## 51	a. vanidicus	grass-bush	26.45	90.60

##	52	a. whitemani trunk-ground	51.06	115.26
##		body_length_mm toepad_lamellae_count	island	ratio_of_hindlimb_to_body
##	1	51.67 27	cuba	0.9765822
##	2	41.32 31	cuba	0.6171346
##	3	30.95 24	cuba	0.8455574
##	4	51.53 36	hispaniola	0.7141471
##	5	72.32 41	cuba	0.6967644
##	6	51.72 28	cuba	0.9545630
##	7	32.94 29	hispaniola	0.9098361
##	8	31.84 28	cuba	0.8602387
##	9	40.22 28	cuba	0.6059175
##	10	56.11 31	hispaniola	0.9233648
##	11	41.81 32	hispaniola	0.9258551
##	12	55.65 32	cuba	0.8524708
##	13	52.29 29	hispaniola	0.9531459
##	14	65.44 42	hispaniola	0.7419010
##	15	39.03 26	cuba	0.8913656
##	16	46.72 32	cuba	0.8910531
##	17	54.45 33	puerto rico	0.8907254
##	18	29.01 24	cuba	0.7886936
##	19	34.32 26	cuba	0.9143357
##	20	45.34 36	hispaniola	0.8709749
##	21	57.97 38	jamacia	0.8402622
##	22	43.95 29	cuba	0.8894198
##	23	39.24 28	cuba	0.5214067
##	24	59.97 28	puerto rico	0.9724862
##	25	68.19 33	hispaniola	0.9602581
##	26	43.19 34	hispaniola	0.8921047
##	27	51.34 32	cuba	0.8759252
##	28	54.89 28	cuba	0.9413372
##	29	31.28 27	cuba	0.8567775
##	30	40.70 25	hispaniola	0.6420147
##	31	47.38 29	cuba	0.8670325
##	32	44.35 33	puerto rico	0.9138670
##	33	56.50 29	jamacia	0.9329204
##	34	63.29 28	hispaniola	0.9783536
##	35	37.39 28	cuba	0.8504948
##	36	53.79 33	hispaniola	0.9154118
##	37	49.05 30	cuba	0.9160041
##	38	42.27 35	jamacia	0.8438609
##	39	34.59 24	cuba	0.7771032
##	40	39.63 26	hispaniola	0.5046682
##	41	41.52 29	puerto rico	0.8041908
##	42	64.37 44	cuba	0.6925586
##	43	45.03 28	cuba	0.8671996
##	44	53.74 29	cuba	0.9384071
##	45	53.14 33	cuba	0.8556643
##	46	36.68 30	hispaniola	0.9266630
##	47	48.85 28	hispaniola	0.8927329
##	48	52.65 38	hispaniola	0.7297246
##	49	65.36 31	hispaniola	0.9981640
##	50	43.62 34	puerto rico	0.7881706

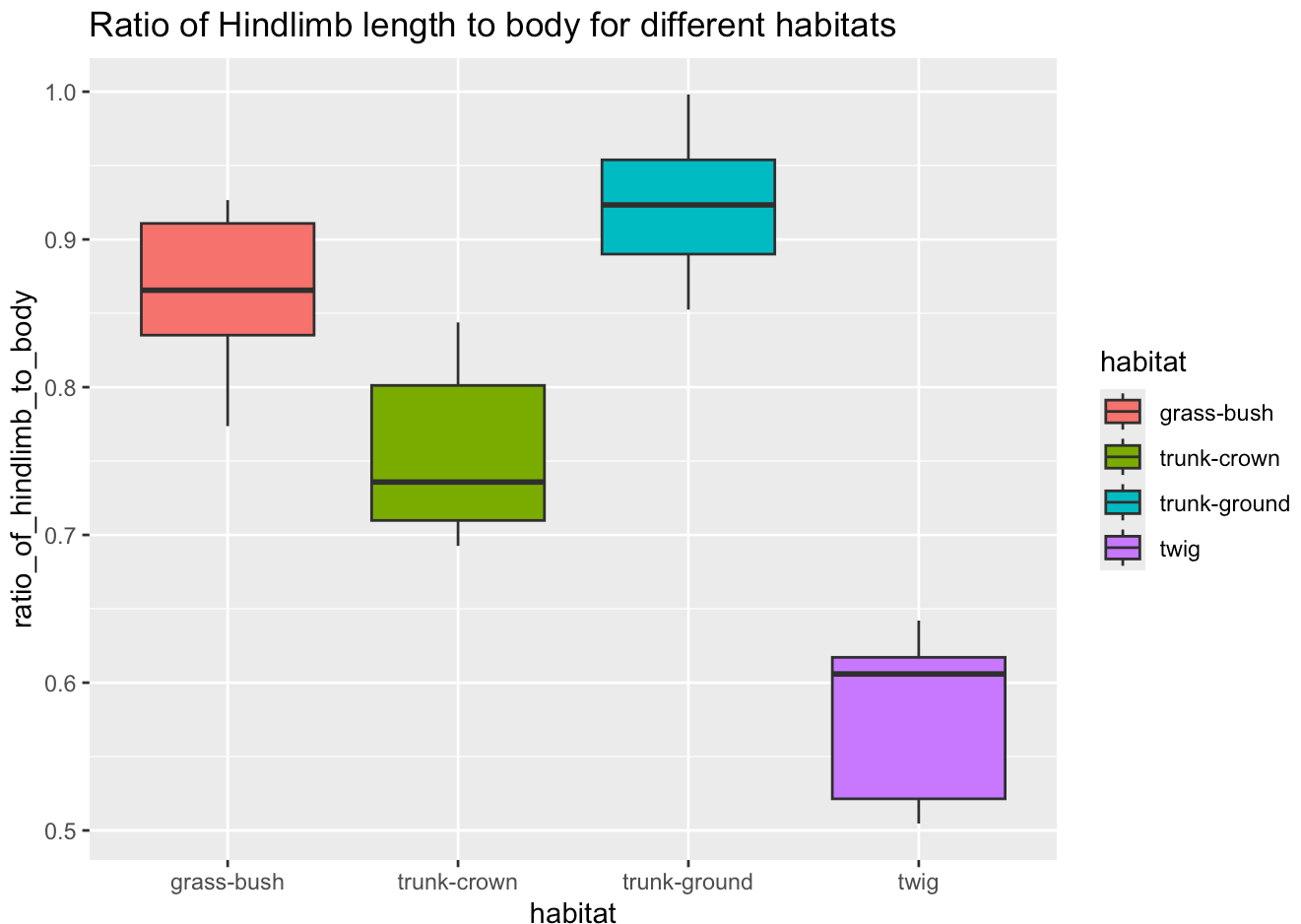
## 51	34.19	23	cuba	0.7736180
## 52	54.77	29	hispaniola	0.9322622
##	ratio_of_tail_to_body			
## 1	1.586801			
## 2	1.325024			
## 3	2.552504			
## 4	1.647196			
## 5	2.135647			
## 6	1.759667			
## 7	3.230115			
## 8	2.971734			
## 9	1.618598			
## 10	1.806630			
## 11	2.770151			
## 12	2.048697			
## 13	2.349398			
## 14	2.171760			
## 15	2.530105			
## 16	1.676156			
## 17	2.471442			
## 18	3.236815			
## 19	3.254079			
## 20	2.895236			
## 21	2.066241			
## 22	1.580205			
## 23	1.014016			
## 24	2.269968			
## 25	2.264408			
## 26	2.982403			
## 27	1.770939			
## 28	1.906905			
## 29	3.232737			
## 30	1.247420			
## 31	1.733854			
## 32	2.948365			
## 33	2.123363			
## 34	2.217728			
## 35	2.603637			
## 36	2.304332			
## 37	1.877880			
## 38	1.943459			
## 39	2.345476			
## 40	1.210194			
## 41	2.846821			
## 42	2.178965			
## 43	1.825450			
## 44	1.748046			
## 45	2.105758			
## 46	3.315431			
## 47	2.098874			
## 48	1.943780			
## 49	2.124694			



```
## 50      1.786566
## 51      2.649898
## 52      2.104437
```

**Question 12. (4 points)** Create a new plot that examines the distribution of `ratio_of_hindlimb_to_body` across habitat types. How does this plot differ from the one you made in Problem 10? Be sure to label your axes and add a title.

```
anolis %>%
  ggplot(mapping=aes(x=habitat,y=ratio_of_hindlimb_to_body))+
  geom_boxplot(mapping = aes(fill=habitat))+
  labs(title="Ratio of Hindlimb length to body for different habitats")#The trunk_ground
not only have the greatest average hindlimb length but also greatest ratio of hindlimb t
o body length.The grass_bush doesn't have large hindlimbs but the ratio of hindlimb to b
ody is kind higher.
```



**Problem 13. (4 points)** A longer tail provides better balance and agility. Create a plot that examines the relationship between body length and tail length. Color the points by habitat type and add a line of best fit. What does this plot suggest about the relationship between body length and tail length? What do you

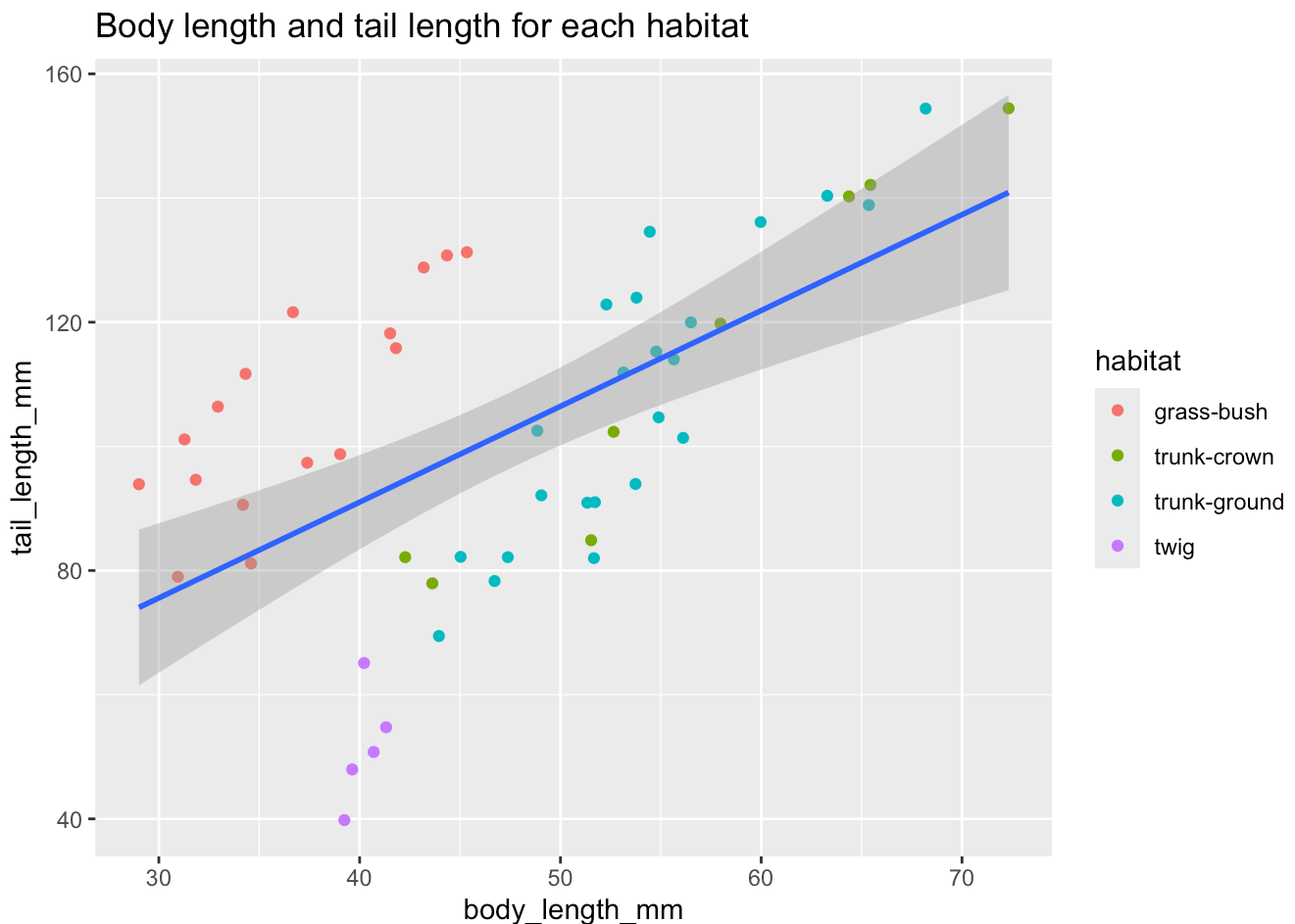
notice about lizards in the **Grass-bush** habitat? Be sure to label your axes and add a title.

```
names(anolis)
```

```
## [1] "species"           "habitat"
## [3] "hindlimb_length_mm" "tail_length_mm"
## [5] "body_length_mm"    "toepad_lamellae_count"
## [7] "island"            "ratio_of_hindlimb_to_body"
## [9] "ratio_of_tail_to_body"
```

```
analis %>%
  ggplot(mapping = aes(x=body_length_mm,y=tail_length_mm))+
  geom_point(mapping = aes(color=habitat))+
  geom_smooth(method = "lm")+
  labs(title="Body length and tail length for each habitat")#1.For most of the data it s
hows a relationship of larger body length have a longer tail length.2.But fir grass_bush
habitat with low body length there are some individuals have a long tail length.
```

```
## `geom_smooth()` using formula = 'y ~ x'
```



**Problem 14. (4 points)** Toepad lamellae are transverse, plate-like structures found on the ventral surface of the digits. They are a key adaptation that allows anoles to cling to and move efficiently on smooth and vertical surfaces. What is the mean number of toepad lamellae for each habitat type?

```
names(anolis)
```

```
## [1] "species"          "habitat"  
## [3] "hindlimb_length_mm" "tail_length_mm"  
## [5] "body_length_mm"    "toepad_lamellae_count"  
## [7] "island"            "ratio_of_hindlimb_to_body"  
## [9] "ratio_of_tail_to_body"
```

```
anolis %>%  
  filter(habitat=="trunk-ground") %>%  
  mutate(mean_toepad_lamellae_trunk_ground=mean(toepad_lamellae_count))#mean number of t  
oepad lamellae for trunk-ground is 29.95652.
```

##	species	habitat	hindlimb_length_mm	tail_length_mm
## 1	a. ahli	trunk-ground	50.46	81.99
## 2	a. allogus	trunk-ground	49.37	91.01
## 3	a. armouri	trunk-ground	51.81	101.37
## 4	a. bremeri	trunk-ground	47.44	114.01
## 5	a. breslini	trunk-ground	49.84	122.85
## 6	a. confusus	trunk-ground	41.63	78.31
## 7	a. cooki	trunk-ground	48.50	134.57
## 8	a. guafe	trunk-ground	39.09	69.45
## 9	a. gundlachi	trunk-ground	58.32	136.13
## 10	a. haetianus	trunk-ground	65.48	154.41
## 11	a. homolechis	trunk-ground	44.97	90.92
## 12	a. imias	trunk-ground	51.67	104.67
## 13	a. jubar	trunk-ground	41.08	82.15
## 14	a. lineatopus	trunk-ground	52.71	119.97
## 15	a. longitibialis	trunk-ground	61.92	140.36
## 16	a. marcanoi	trunk-ground	49.24	123.95
## 17	a. mestrei	trunk-ground	44.93	92.11
## 18	a. quadriocellifer	trunk-ground	39.05	82.20
## 19	a. rubribarbus	trunk-ground	50.43	93.94
## 20	a. sagrei	trunk-ground	45.47	111.90
## 21	a. shrevei	trunk-ground	43.61	102.53
## 22	a. strahmi	trunk-ground	65.24	138.87
## 23	a. whitemani	trunk-ground	51.06	115.26
##	body_length_mm	toepad_lamellae_count	island	ratio_of_hindlimb_to_body
## 1	51.67	27	cuba	0.9765822
## 2	51.72	28	cuba	0.9545630
## 3	56.11	31	hispaniola	0.9233648
## 4	55.65	32	cuba	0.8524708
## 5	52.29	29	hispaniola	0.9531459
## 6	46.72	32	cuba	0.8910531
## 7	54.45	33	puerto rico	0.8907254
## 8	43.95	29	cuba	0.8894198
## 9	59.97	28	puerto rico	0.9724862
## 10	68.19	33	hispaniola	0.9602581
## 11	51.34	32	cuba	0.8759252
## 12	54.89	28	cuba	0.9413372
## 13	47.38	29	cuba	0.8670325
## 14	56.50	29	jamacia	0.9329204
## 15	63.29	28	hispaniola	0.9783536
## 16	53.79	33	hispaniola	0.9154118
## 17	49.05	30	cuba	0.9160041
## 18	45.03	28	cuba	0.8671996
## 19	53.74	29	cuba	0.9384071
## 20	53.14	33	cuba	0.8556643
## 21	48.85	28	hispaniola	0.8927329
## 22	65.36	31	hispaniola	0.9981640
## 23	54.77	29	hispaniola	0.9322622
##	ratio_of_tail_to_body	mean_toepad_lamellae_trunk_ground		
## 1	1.586801	29.95652		
## 2	1.759667	29.95652		
## 3	1.806630	29.95652		

```
## 4      2.048697      29.95652
## 5      2.349398      29.95652
## 6      1.676156      29.95652
## 7      2.471442      29.95652
## 8      1.580205      29.95652
## 9      2.269968      29.95652
## 10     2.264408      29.95652
## 11     1.770939      29.95652
## 12     1.906905      29.95652
## 13     1.733854      29.95652
## 14     2.123363      29.95652
## 15     2.217728      29.95652
## 16     2.304332      29.95652
## 17     1.877880      29.95652
## 18     1.825450      29.95652
## 19     1.748046      29.95652
## 20     2.105758      29.95652
## 21     2.098874      29.95652
## 22     2.124694      29.95652
## 23     2.104437      29.95652
```

```
anolis %>%
  filter(habitat=="twig")%>%
  mutate(mean_toepad_lamellae_twig=mean(toepad_lamellae_count))#mean number of toepad la
mellae for twig is 27.6.
```

```
##      species habitat hindlimb_length_mm tail_length_mm body_length_mm
## 1    a. alayoni   twig           25.50           54.75           41.32
## 2    a. angusticeps twig           24.37           65.10           40.22
## 3    a. guazuma   twig           20.46           39.79           39.24
## 4    a. insolitus twig           26.13           50.77           40.70
## 5    a. placidus  twig           20.00           47.96           39.63
##      toepad_lamellae_count      island ratio_of_hindlimb_to_body
## 1                        31      cuba           0.6171346
## 2                        28      cuba           0.6059175
## 3                        28      cuba           0.5214067
## 4                        25 hispaniola           0.6420147
## 5                        26 hispaniola           0.5046682
##      ratio_of_tail_to_body mean_toepad_lamellae_twig
## 1                1.325024                27.6
## 2                1.618598                27.6
## 3                1.014016                27.6
## 4                1.247420                27.6
## 5                1.210194                27.6
```

```
anolis %>%
  filter(habitat=="grass-bush")%>%
  mutate(mean_toepad_lamellae_grass_bush=mean(toepad_lamellae_count))#mean number of toe
pad lamellae for grass_bush is 28.3125.
```

##	species	habitat	hindlimb_length_mm	tail_length_mm
## 1	a. alfaroi	grass-bush	26.17	79.00
## 2	a. alumina	grass-bush	29.97	106.40
## 3	a. alutaceus	grass-bush	27.39	94.62
## 4	a. bahorucoensis	grass-bush	38.71	115.82
## 5	a. clivicola	grass-bush	34.79	98.75
## 6	a. cupeyalensis	grass-bush	22.88	93.90
## 7	a. cyanopleurus	grass-bush	31.38	111.68
## 8	a. dolichocephalus	grass-bush	39.49	131.27
## 9	a. hendersoni	grass-bush	38.53	128.81
## 10	a. inexpectatus	grass-bush	26.80	101.12
## 11	a. krugi	grass-bush	40.53	130.76
## 12	a. macilentus	grass-bush	31.80	97.35
## 13	a. ophiolepis	grass-bush	26.88	81.13
## 14	a. poncensis	grass-bush	33.39	118.20
## 15	a. semilineatus	grass-bush	33.99	121.61
## 16	a. vanidicus	grass-bush	26.45	90.60
##	body_length_mm	toepad_lamellae_count	island	ratio_of_hindlimb_to_body
## 1	30.95	24	cuba	0.8455574
## 2	32.94	29	hispaniola	0.9098361
## 3	31.84	28	cuba	0.8602387
## 4	41.81	32	hispaniola	0.9258551
## 5	39.03	26	cuba	0.8913656
## 6	29.01	24	cuba	0.7886936
## 7	34.32	26	cuba	0.9143357
## 8	45.34	36	hispaniola	0.8709749
## 9	43.19	34	hispaniola	0.8921047
## 10	31.28	27	cuba	0.8567775
## 11	44.35	33	puerto rico	0.9138670
## 12	37.39	28	cuba	0.8504948
## 13	34.59	24	cuba	0.7771032
## 14	41.52	29	puerto rico	0.8041908
## 15	36.68	30	hispaniola	0.9266630
## 16	34.19	23	cuba	0.7736180
##	ratio_of_tail_to_body	mean_toepad_lamellae_grass_bush		
## 1	2.552504	28.3125		
## 2	3.230115	28.3125		
## 3	2.971734	28.3125		
## 4	2.770151	28.3125		
## 5	2.530105	28.3125		
## 6	3.236815	28.3125		
## 7	3.254079	28.3125		
## 8	2.895236	28.3125		
## 9	2.982403	28.3125		
## 10	3.232737	28.3125		
## 11	2.948365	28.3125		
## 12	2.603637	28.3125		
## 13	2.345476	28.3125		
## 14	2.846821	28.3125		
## 15	3.315431	28.3125		
## 16	2.649898	28.3125		

```

anolis %>%
  filter(habitat=="trunk-crown")%>%
  mutate(mean_toepad_lamellae_trunk_crown=mean(toepad_lamellae_count))#mean number of to
epad lamellae for trunk-crown is 38.5.

```

```

##      species      habitat hindlimb_length_mm tail_length_mm body_length_mm
## 1    a. aliniger trunk-crown          36.80          84.88          51.53
## 2    a. allisoni trunk-crown          50.39          154.45          72.32
## 3 a. chlorocyanus trunk-crown          48.55          142.12          65.44
## 4    a. grahami trunk-crown          48.71          119.78          57.97
## 5    a. opalinus trunk-crown          35.67           82.15          42.27
## 6    a. porcatus trunk-crown          44.58          140.26          64.37
## 7    a. singularis trunk-crown          38.42          102.34          52.65
## 8    a. stratulus trunk-crown          34.38           77.93          43.62
##  toepad_lamellae_count      island ratio_of_hindlimb_to_body
## 1                      36 hispaniola          0.7141471
## 2                      41      cuba          0.6967644
## 3                      42 hispaniola          0.7419010
## 4                      38      jamacia          0.8402622
## 5                      35      jamacia          0.8438609
## 6                      44      cuba          0.6925586
## 7                      38 hispaniola          0.7297246
## 8                      34 puerto rico          0.7881706
##  ratio_of_tail_to_body mean_toepad_lamellae_trunk_crown
## 1              1.647196              38.5
## 2              2.135647              38.5
## 3              2.171760              38.5
## 4              2.066241              38.5
## 5              1.943459              38.5
## 6              2.178965              38.5
## 7              1.943780              38.5
## 8              1.786566              38.5

```

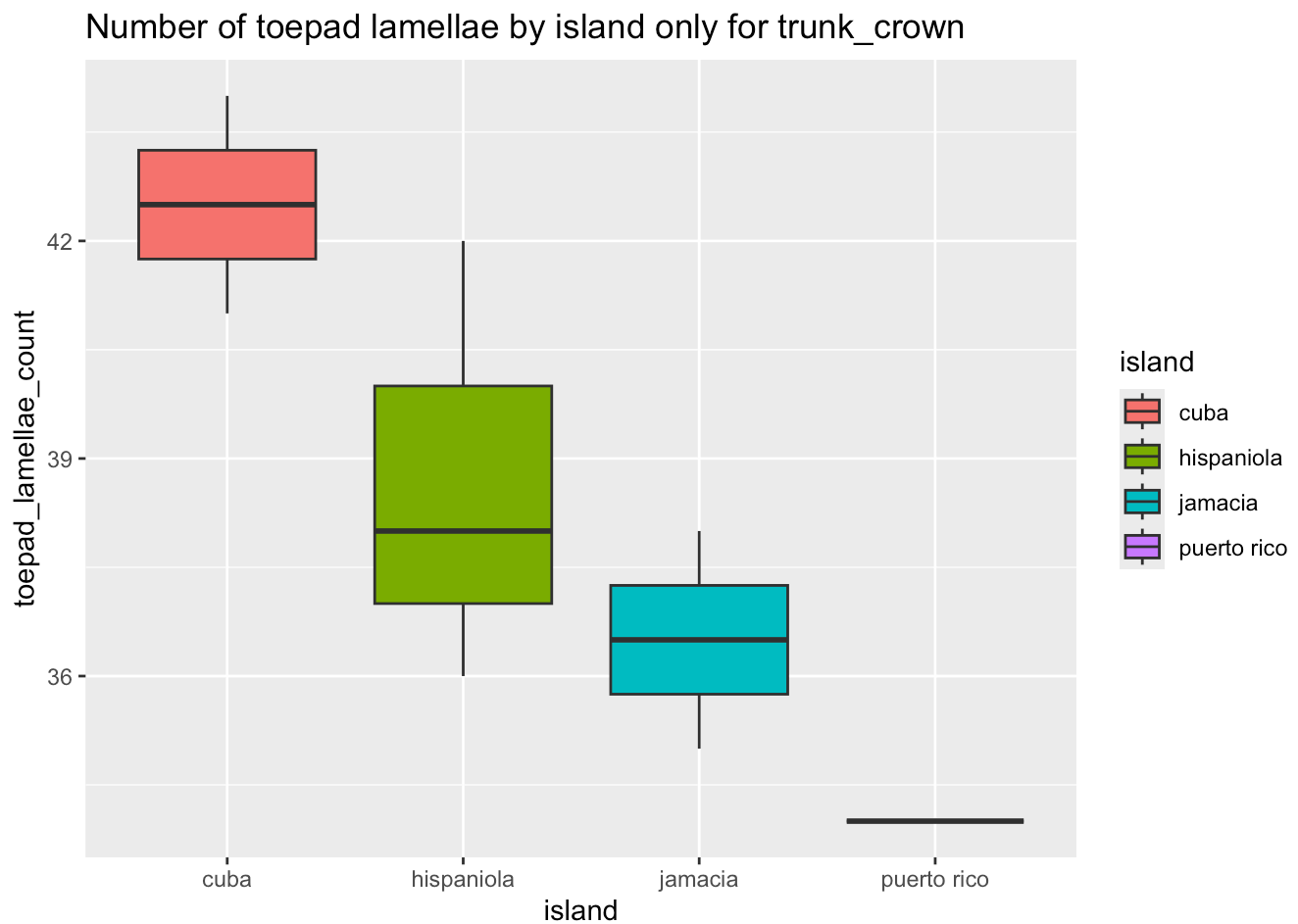
#The mean number of toepad lamellae of trunk\_ground is:29.95652 #The mean number of toepad lamellae of trunk-crown is:38.5 #The mean number of toepad lamellae of grass\_bush is:28.3125 #The mean number of toepad lamellae of twig is:27.6

**Problem 15. (5 points)** The number of toepad lamellae is significantly different for trunk-crown species. But, is this consistent across all islands? Make a plot that shows the range in number of toepad lamellae by island for trunk-crown species only. Be sure to label your axes and add a title.

```

anolis %>%
  filter(habitat=="trunk-crown") %>%
  ggplot(mapping = aes(x=island,y=toepad_lamellae_count))+
  geom_boxplot(mapping = aes(fill=island))+
  labs(title="Number of toepad lamellae by island only for trunk_crown")

```



## Submit the Midterm

1. Save your work and knit the .rmd file.
2. Open the .html file and “print” it to a .pdf file in Google Chrome (not Safari).
3. Go to the class Canvas page and open Gradescope.
4. Submit your .pdf file to the midterm assignment- be sure to assign the pages to the correct questions.
5. Commit and push your work to your repository.