

TRANSFORMING OUTLIERS

OVERVIEW

- **outliers** = extreme values that tend to exert undue influence on result of analysis
- Know your options for dealing with outliers (so can get results that mean what you think they mean)

SUMMARY

- Log linear transformation brings it closer to the normal distribution - which is the assumption behind so many statistical procedures

PSEUDOCODE

INSTALL AND LOAD PACKAGES

LOAD AND PREPARE DATA

Check existence of outliers - with histogram:

Best way to check for outliers is with boxplot: (bc it marks outliers)

LOOK AT SOME OPTIONS FOR DEALING WITH OUTLIERS

PART1: REMOVE OUTLIERS

See which ones are outliers:

Filter out continents

Create histogram & boxplot

PART2: BRING IN OUTLIERS

Sort observations in descending order:

IF value is greater than 840 THEN change it to 840:

Graph results:

PART3: CREATE NEW GROUPS

Sort in descending order:

Create new variable - called "landmass" > IF value is < 1000 THEN call it an island:

Boxplot of area of continents:

Boxplot of islands:

PART4: TRANSFORMING DATA

Strong positive skew:

Log is natural logarithm with base e (base 10 is log10) > brings in extreme positive values:

INSTALL AND LOAD PACKAGES

```
pacman::p_load(datasets, pacman, tidyverse)
```

LOAD AND PREPARE DATA

from **datasets** package - **islands** dataset - has strong positive skew

```
?islands
```

- area of world's major land masses
- areas on 1000s of square mile - of landmasses that exceed 10,000 sq miles
- named vector - with 48 observations

```
islands
```

- results:

-alphabetical order - with area (from Africa with 11506, down to end of list with Victoria with 82 (thousands of square miles))

```
> islands
```

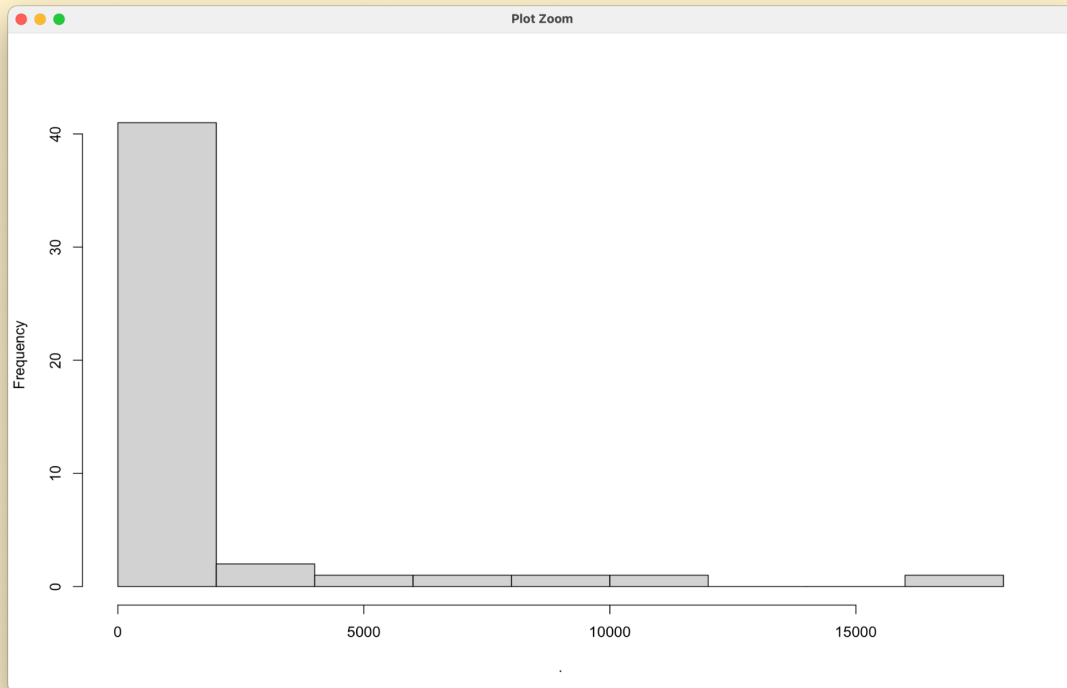
Africa	Antarctica	Asia	Australia
11506	5500	16988	2968
Axel Heiberg	Baffin	Banks	Borneo
16	184	23	280
Britain	Celebes	Celon	Cuba
84	73	25	43
Devon	Ellesmere	Europe	Greenland
21	82	3745	840
Hainan	Hispaniola	Hokkaido	Honshu
13	30	30	89
Iceland	Ireland	Java	Kyushu
40	33	49	14
Luzon	Madagascar	Melville	Mindanao
42	227	16	36
Moluccas	New Britain	New Guinea	New Zealand (N)
29	15	306	44
New Zealand (S)	Newfoundland	North America	Novaya Zemlya
58	43	9390	32
Prince of Wales	Sakhalin	South America	Southampton
13	29	6795	16
Spitsbergen	Sumatra	Taiwan	Tasmania
15	183	14	26
Tierra del Fuego	Timor	Vancouver	Victoria
19	13	12	82

Check existence of outliers - with histogram:

```
islands %>% hist(main = NULL)
```

- result:

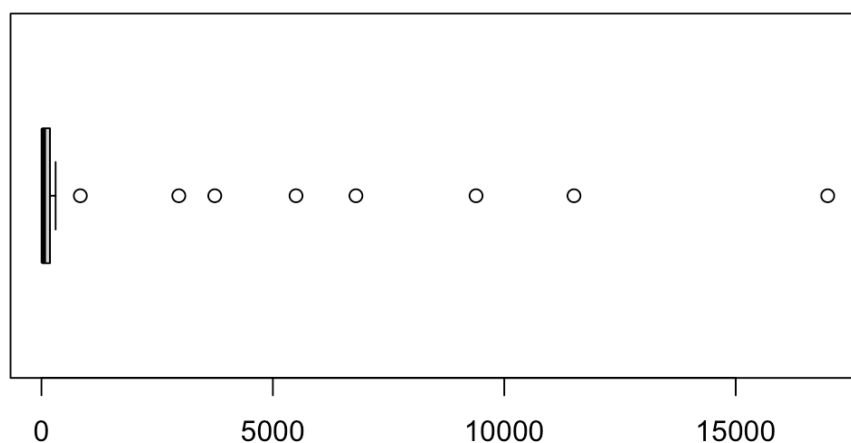
-most in smallest size bin - and few way out past rest of plot >> NOT normal distribution & big outliers (throws off analysis)

**Best way to check for outliers is with boxplot: (bc it marks outliers)**

```
islands %>% boxplot(horizontal = T)
```

- result:

-range of 50% scores is super compressed and at low end
-even the highest non-outlying data point is compressed
-8 outliers - one extreme (asia)



LOOK AT SOME OPTIONS FOR LEADING WITH OUTLIERS

there are sophisticated algorithms that deal with outliers:

- when using the non-parametric approach
 - decision trees dn get thrown off by outliers
 - neural networks - are more flexible wrt outliers

IF standard analysis (scatterplots, means) THEN want to deal with outliers:

PART1: REMOVE OUTLIERS

- draconian - cut off outliers (throw them away)
- appropriate - as long as
 - 1) only care about non-outlier scores
 - 2) are specific, clear that you did that (and that focusing only on the major ones)

See which ones are outliers:

- Sort observation in descending value:

```
islands2 <- islands %>%
  enframe() %>%           # convert vector to tibble
  arrange(desc(value)) %>% # sort by descending values
  print()
```

- result:

-Asia 16K Africa 11K .. (top masses are continents - of course they are going to be huge)

```
> islands2 <- islands %>%
+   enframe() %>%           # Convert vector to tibble
+   arrange(desc(value)) %>% # Sort by descending values
+   print()
# A tibble: 48 × 2
   name      value
  <chr>    <dbl>
1 Asia      16988
2 Africa    11506
3 North America 9390
4 South America 6795
5 Antarctica   5500
6 Europe      3745
7 Australia   2968
8 Greenland   840
9 New Guinea  306
10 Borneo     280
# i 38 more rows
# i Use `print(n = ...)` to see more rows
```

Data

islands2 48 obs. of 2 variables

- → can say - we don't want to focus on continents > want to focus on islands (a way of defining your sample, that helps you deal with some of these outliers)

Filter out continents

```
islands2 %>%
  filter(value < 1000) %>%
  print()
```

- **result:**

-filters out everything to Australia, and keep greenland and below

```
> islands2 %<>%
+   filter(value < 1000) %>%
+   print()
# A tibble: 41 × 2
   name      value
  <chr>    <dbl>
1 Greenland  840
2 New Guinea 306
3 Borneo    280
4 Madagascar 227
5 Baffin    184
6 Sumatra   183
7 Honshu     89
8 Britain    84
9 Ellesmere  82
10 Victoria  82
# i 31 more rows
# i Use `print(n = ...)` to see more rows
```

Data

islands2	41 obs. of 2 variables
----------	------------------------

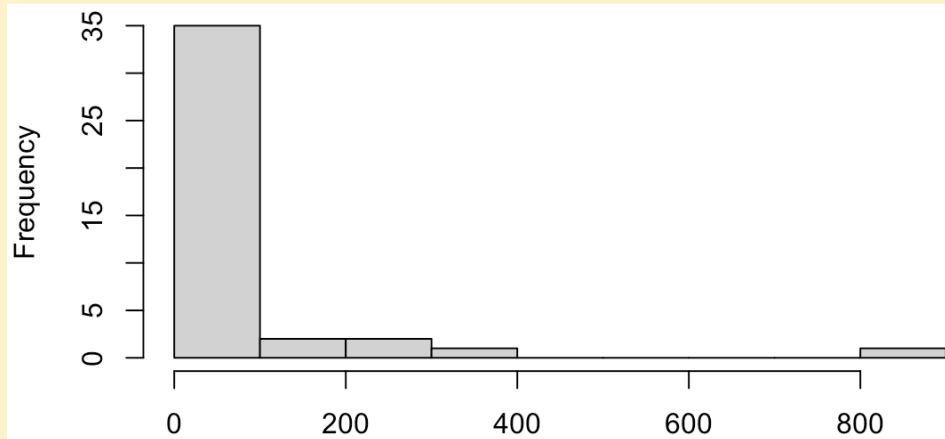
Create histogram & boxplot

- **pull()** creates a VECTOR (like `islands$value`)
- **select()** create a DATAFRAME
- (dift functions need dift data formats)

```
islands2 %>% pull(value) %>% histogram(main = NULL)
```

- result:

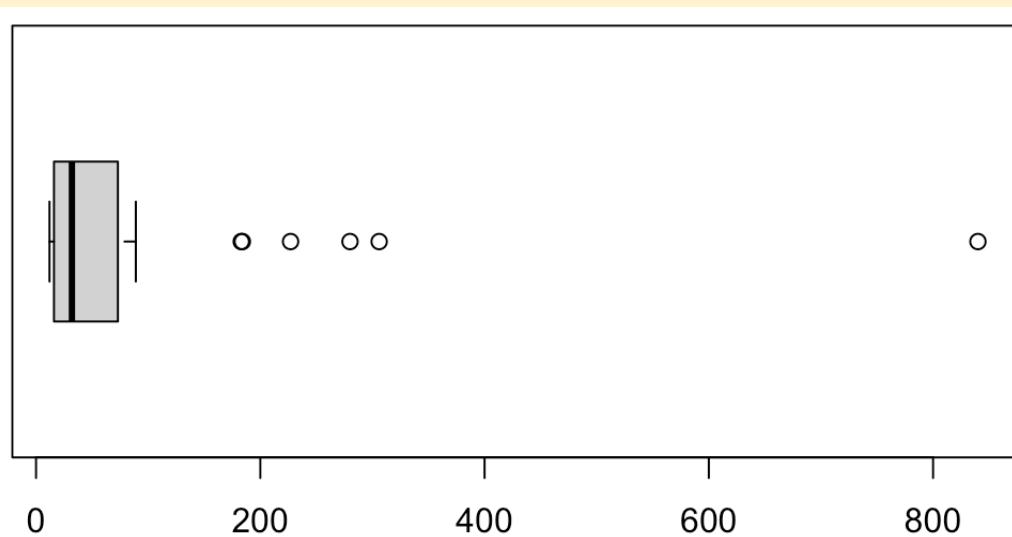
-still have outliers (greenland)



```
islands2 %>% select(value) %>% boxplot(horizontal = T)
```

- result:

-not quite as pathological as before (bd redefining group that interested in)



PART2: BRING IN OUTLIERS

- **winsorizing** - option - that is not used very often
 - ex: times on races / time to graduation / financial data
- take extreme values -> change them to highest non-outlier value
 - ex: time to graduation - we are going up to 8 years, and anything after 8 we are going to code as 8

Sort observations in descending order:

```
islands3 <- islands %>%
  enframe() %>%           # convert vector to tibble
  arrange(desc(value)) %>% # sort by descending values
  print()
```

- results:

The largest non-continent is Greenland, with an area of about 840,000 square miles (coded as 840 in the dataset)

```
> islands3 <- islands %>%
+   enframe() %>%           # Convert vector to tibble
+   arrange(desc(value)) %>% # Sort by descending values
+   print()
# A tibble: 48 x 2
   name      value
  <chr>    <dbl>
1 Asia      16988
2 Africa    11506
3 North America 9390
4 South America 6795
5 Antarctica 5500
6 Europe     3745
7 Australia  2968
8 Greenland   840
9 New Guinea  306
10 Borneo     280
# i 38 more rows
# i Use `print(n = ...)` to see more rows
```

▶ islands3 48 obs. of 2 variables

IF value is greater than 840 THEN change it to 840:

```
islands3 <- islands3 %>%
  mutate(
    value = ifelse(
      value > 840,      # test if value is greater than 840
      840,              # if true, replace value with 840
      value             # if false, keep existing value
    )
  ) %>%
  print()
```

- results:

Now many 840s (for all the continents)

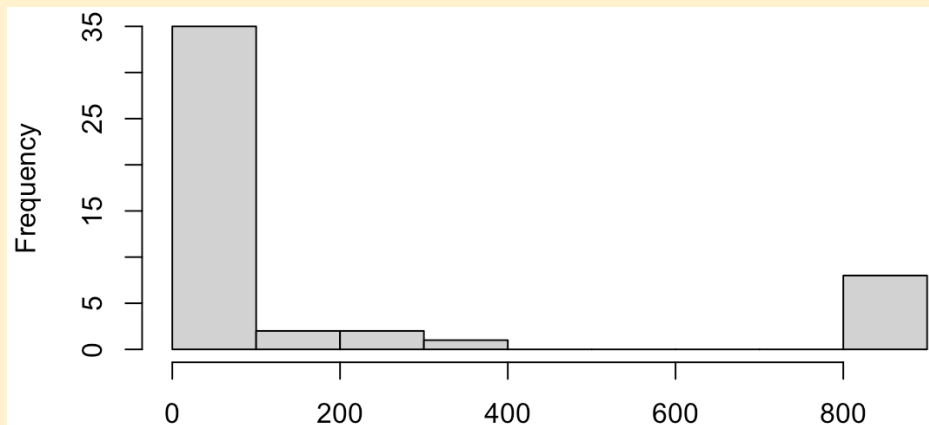
```
> islands3 <- islands3 %>%
+   mutate(
+     value = ifelse(
+       value > 840, # Test if value is greater than 840
+       840,        # If true, replace value with 840
+       value       # If false, keep existing value
+     )
+   ) %>%
+   print()
# A tibble: 48 × 2
  name      value
  <chr>    <dbl>
1 Asia      840
2 Africa    840
3 North America 840
4 South America 840
5 Antarctica 840
6 Europe     840
7 Australia  840
8 Greenland  840
9 New Guinea 306
10 Borneo    280
# i 38 more rows
# i Use `print(n = ...)` to see more rows
```


Graph results:

```
islands3 %>% pull(value) %>% hist(main = NULL)
```

- results

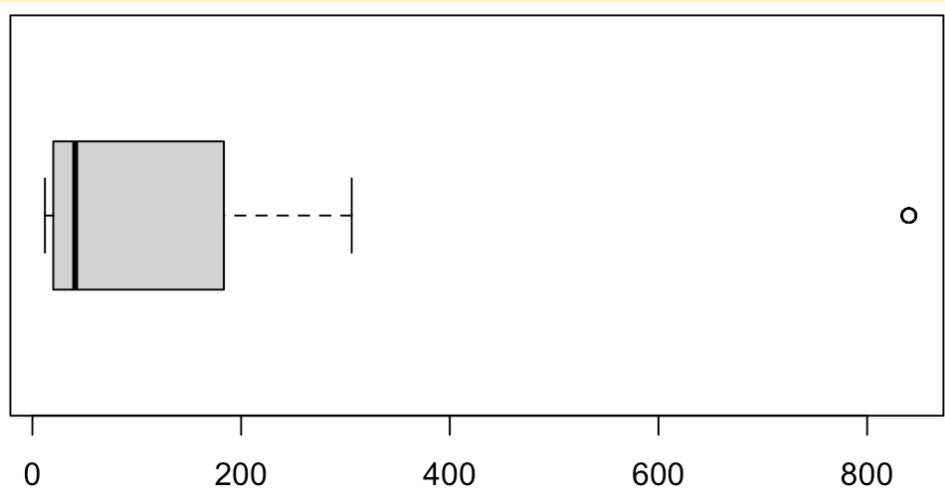
Big bump at high end - bc many observations at 840



```
islands3 %>% select(value) %>% boxplot(horizontal = T)
```

- results

Several observations at high end stacked into of each other



PART3: CREATE NEW GROUPS

Split into 2 groups: continents, islands > will treat then separately

Sort in descending order:

```
islands4 <- islands %>%
  enframe() %>%           # convert vector to tibble
  arrange(desc(value)) %>% # sort in descending order
  print()
```

- **results:**

```
> islands4 <- islands %>%
+   enframe() %>%           # Convert vector to tibble
+   arrange(desc(value)) %>% # Sort by descending values
+   print()
# A tibble: 48 × 2
   name      value
  <chr>    <dbl>
1 Asia      16988
2 Africa    11506
3 North America 9390
4 South America 6795
5 Antarctica   5500
6 Europe       3745
7 Australia    2968
8 Greenland     840
9 New Guinea    306
10 Borneo       280
# i 38 more rows
# i Use `print(n = ...)` to see more rows
```

▶ islands4 48 obs. of 2 variables

Create new variable - called "landmass" > IF value is < 1000 THEN call it an island:

```
island4 %>%
  mutate(
    landmass = ifelse
      value < 1000,      # test if value is less than 1k sq mi
      "island",         # if true, it's an island
      "continent"       # if false, it's a continent
  )
) %>%
print()
```

- results:

console: name <chr> / value <dbl> / landmass <chr>

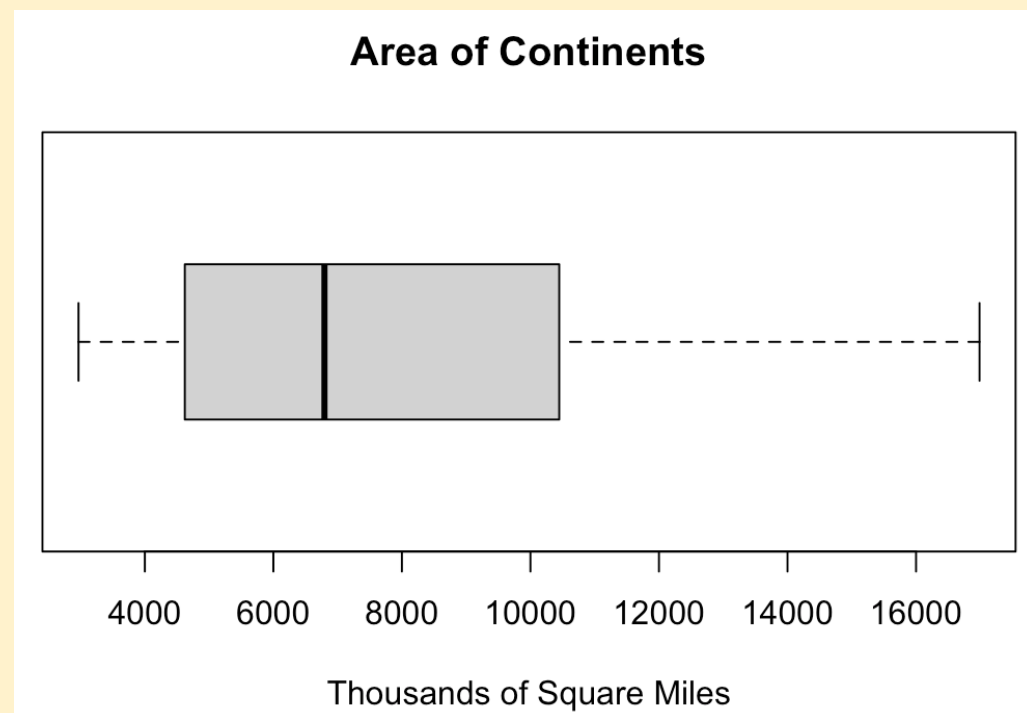
```
> islands4 %<>%
+ mutate(
+   landmass = ifelse(
+     value < 1000, # Test if value is less than 1m sq mi
+     "island",    # If true, it's an island
+     "continent"  # If false, it's a continent
+   )
+ ) %>%
+ print()
# A tibble: 48 × 3
  name      value landmass
  <chr>    <dbl> <chr>
1 Asia      16988 continent
2 Africa    11506 continent
3 North America 9390 continent
4 South America 6795 continent
5 Antarctica   5500 continent
6 Europe      3745 continent
7 Australia   2968 continent
8 Greenland    840 island
9 New Guinea   306 island
10 Borneo      280 island
# i 38 more rows
# i Use `print(n = ...)` to see more rows
```

Boxplot of area of continents:

```
islands4 %>%  
  filter(landmass == "continent") %>%  
  select(value) %>%  
  boxplot(  
    horizontal = T,  
    main = "Area of Continents",  
    xlab = "Thousands of Square Miles"  
  )
```

- results:

No outliers

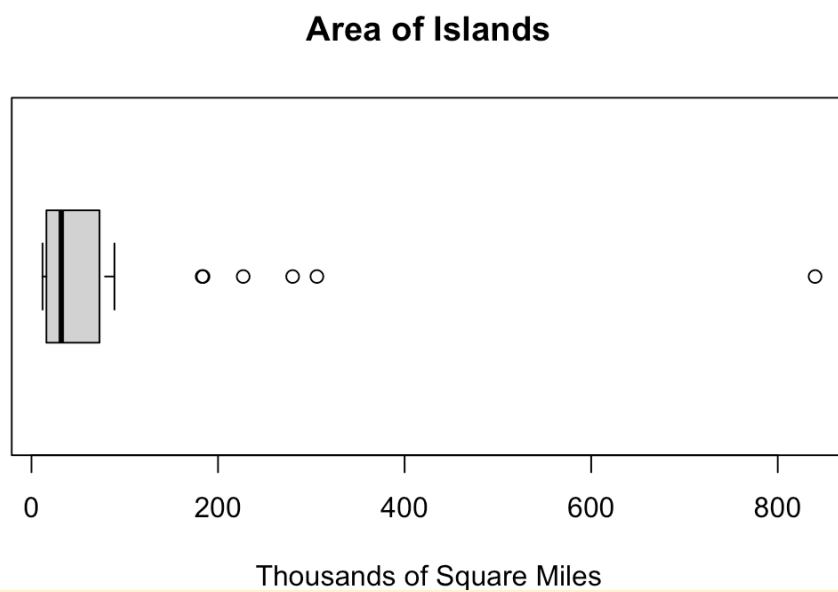


Boxplot of islands:

```
islands4 %>%  
  filter(landmass == "island") %>%  
  select(value) %>%  
  boxplot(  
    horizontal = T,  
    main = "Area of Islands",  
    xlab = "Thousands of Square Miles"  
  )
```

- results:

Greenland is still outlier - but at least you can see what the box looks like now



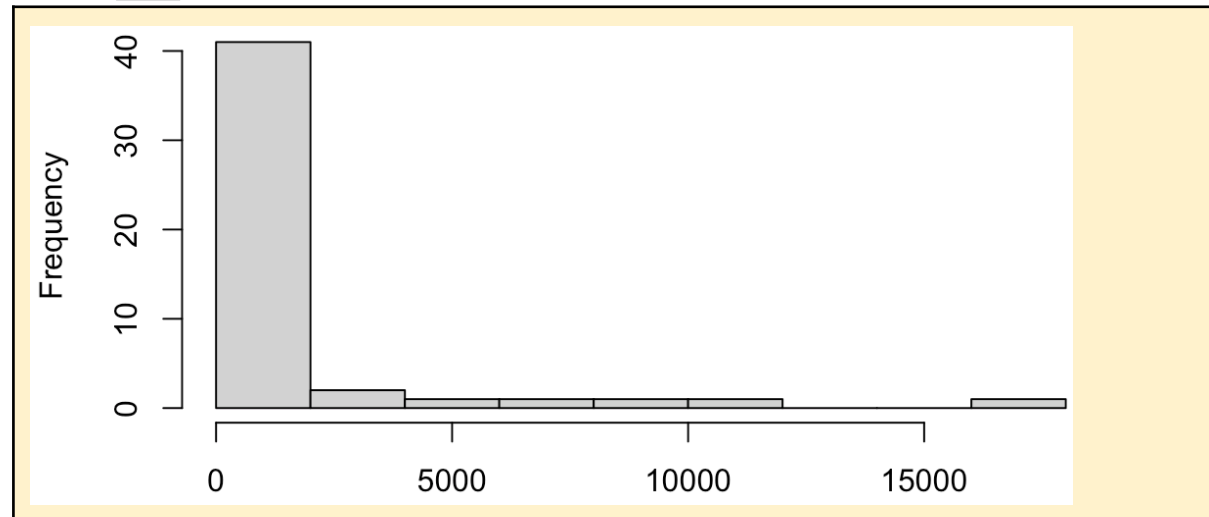
PART4: TRANSFORMING DATA

- linear transformation - do transformation to all the data
- IF positively skewed data - and all values at least 1 THEN logarithm

Strong positive skew:

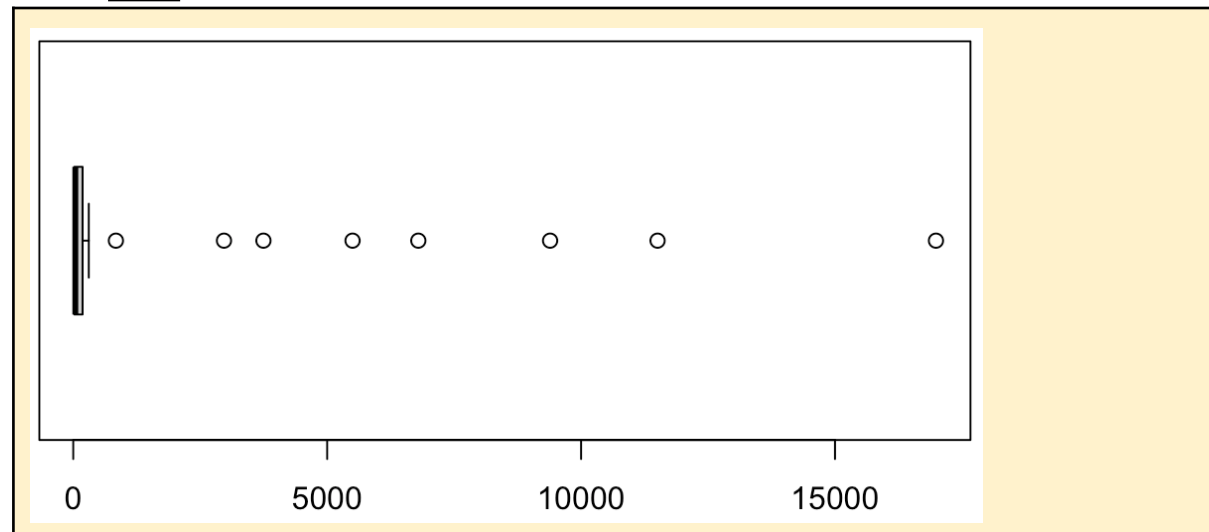
```
islands %>% histogram(main = NULL)
```

- result:



```
islands %>% boxplot(horizontal = T)
```

- result:

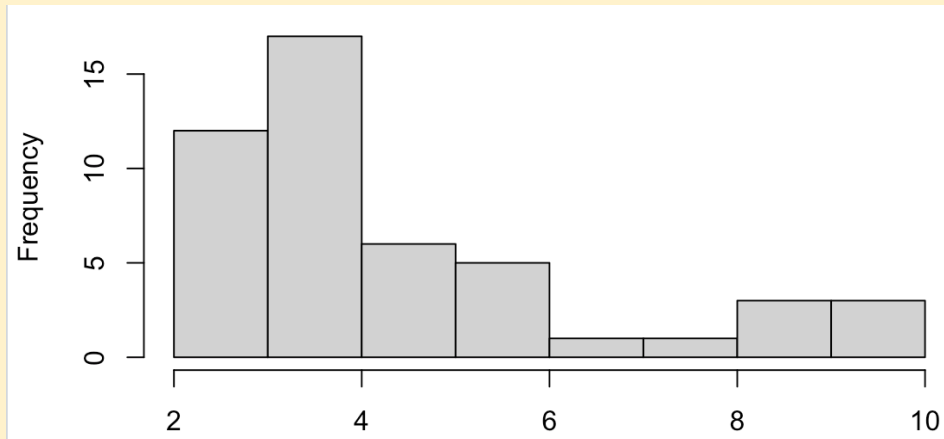


Log is natural logarithm with base e (base 10 is log10) > brings in extreme positive values:

```
islands %>% log() %>% histogram(main = NULL)
```

- result:

Still outliers , but not miles away from others



```
islands %>% log() %>% boxplot(horizontal = T)
```

- result:

Some outliers, but box is predominant

