# Expected answers

#### 2023-10-11

### Part 1: body mass and metabolic rate

#### Ex.2)

```
Expected output:
```

```
## Rows: 25 Columns: 3
## -- Column specification ------
## Delimiter: ","
## chr (1): family
## dbl (2): body_mass, metabolic_rate
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.
## # A tibble: 6 x 3
    body_mass metabolic_rate family
##
        <dbl>
                    <dbl> <chr>
## 1
        32000
                      50.0 Antilocapridae
## 2
       37800
                      52.0 Antilocapridae
## 3
       347000
                     307. Bovidae
                      10.1 Bovidae
## 4
        4200
## 5
       196500
                      230. Bovidae
                      149. Bovidae
## 6
       100000
Ex.5)
## spc_tbl_ [25 x 3] (S3: spec_tbl_df/tbl_df/tbl/data.frame)
                 : num [1:25] 32000 37800 347000 4200 196500 ...
## $ body mass
## $ metabolic_rate: num [1:25] 50 52 306.8 10.1 230.1 ...
                  : chr [1:25] "Antilocapridae" "Antilocapridae" "Bovidae" "Bovidae" ...
   - attr(*, "spec")=
##
    .. cols(
         body_mass = col_double(),
##
         metabolic_rate = col_double(),
##
         family = col_character()
   . .
## - attr(*, "problems")=<externalptr>
Ex.6)
Expected output:
## [1] "Antilocapridae" "Bovidae"
                                       "Camelidae"
                                                       "Canidae"
## [5] "Cervidae"
                      "Suidae"
                                       "Tayassuidae"
                                                       "Tragulidae"
```

```
## [1] "Antilocapridae" "Bovidae" "Camelidae" "Canidae"
## [5] "Cervidae" "Suidae" "Tayassuidae" "Tragulidae"
```

### Ex.9)

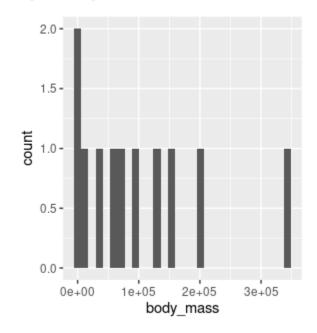
Expected output:

```
## # A tibble: 1 x 1
## range
## <dbl>
## 1 405387

## # A tibble: 1 x 1
## range
## <dbl>
## 1 342800
```

# Ex.10)

Expected output:

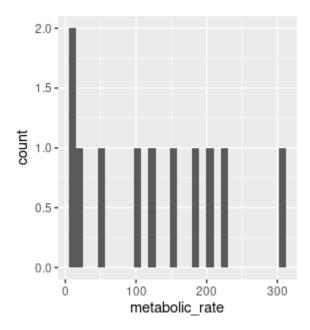


# Ex.11)

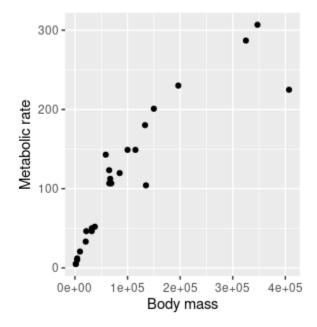
Expected outputs:

```
## # A tibble: 1 x 1
## range
## <dbl>
## # A tibble: 1 x 1
## range
## <dbl>
## 1 297.
```

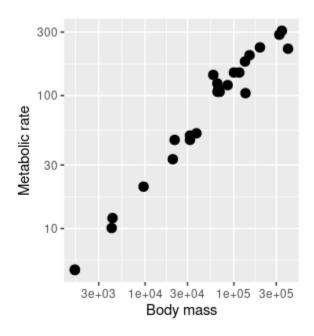
Ex.12)
Expected output:



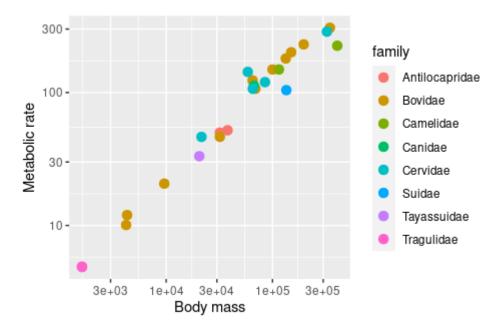
Ex.13)
Expected output:



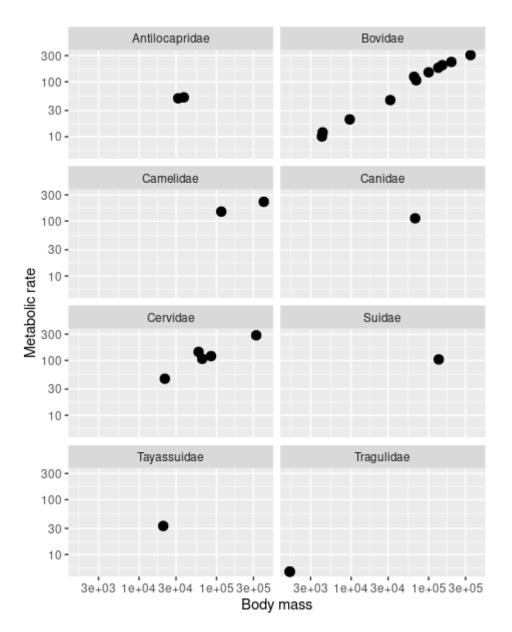
Ex.14)
Expected output:



Ex.15)
Expected output:



Ex.16)
Expected output:



# Ex.17)

Expected output:

##	#	A tibble: 8 x 3	3	
##		family	N	Range_body_mass
##		<fct></fct>	<int></int>	<dbl></dbl>
##	1	Antilocapridae	2	5800
##	2	Bovidae	11	342800
##	3	Camelidae	2	292000
##	4	Canidae	1	0
##	5	Cervidae	5	303500
##	6	Suidae	1	0
##	7	Tayassuidae	1	0
##	8	Tragulidae	2	5

### Part 2: Carbon storage in shrubs

#### Ex.18)

Expected output:

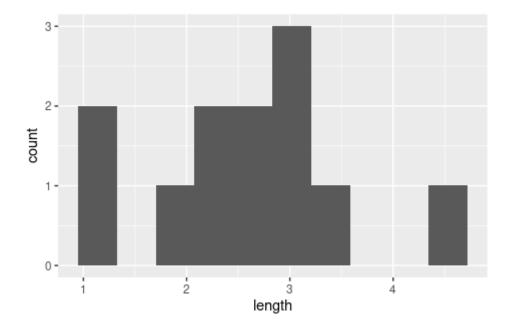
```
## Rows: 12 Columns: 5
## Delimiter: ","
## dbl (5): site, experiment, length, width, height
##
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.
## # A tibble: 6 x 5
##
     site experiment length width height
##
    <dbl>
            <dbl> <dbl> <dbl>
## 1
        1
                  1
                       2.2
                            1.3
                                   9.6
## 2
        1
                  2
                       2.1
                            2.2
                                   7.6
## 3
        1
                  3
                       2.7
                            1.5
                                   2.2
        2
                       3
## 4
                  1
                            4.5
                                   1.5
        2
                  2
## 5
                       3.1
                            3.1
                                   4
## 6
                            2.8
                       2.5
                                   3
## [1] 12 5
## spc_tbl_ [12 x 5] (S3: spec_tbl_df/tbl_df/tbl/data.frame)
           : num [1:12] 1 1 1 2 2 2 3 3 3 4 ...
## $ experiment: num [1:12] 1 2 3 1 2 3 1 2 3 1 ...
## $ length
            : num [1:12] 2.2 2.1 2.7 3 3.1 2.5 1.9 1.1 3.5 2.9 ...
              : num [1:12] 1.3 2.2 1.5 4.5 3.1 2.8 1.8 0.5 2 2.7 ...
## $ width
             : num [1:12] 9.6 7.6 2.2 1.5 4 3 4.5 2.3 7.5 3.2 ...
##
   $ height
   - attr(*, "spec")=
##
##
    .. cols(
##
         site = col_double(),
##
         experiment = col_double(),
##
        length = col_double(),
    . .
##
        width = col double(),
##
        height = col_double()
    . .
##
    ..)
   - attr(*, "problems")=<externalptr>
##
        site
                   experiment
                                 length
                                                width
                                                               height
## Min. :1.00
                 Min. :1
                             Min. :1.100
                                            Min. :0.500
                                                           Min. :1.50
  1st Qu.:1.75
                 1st Qu.:1
                             1st Qu.:2.050
                                            1st Qu.:1.725
                                                           1st Qu.:2.60
## Median :2.50
                 Median :2
                             Median :2.600
                                            Median :2.100
                                                           Median:3.60
## Mean
         :2.50
                 Mean :2
                             Mean :2.558
                                            Mean
                                                 :2.417
                                                           Mean
                                                                 :4.55
                                            3rd Qu.:2.875
## 3rd Qu.:3.25
                 3rd Qu.:3
                             3rd Qu.:3.025
                                                           3rd Qu.:6.75
## Max. :4.00
                                                   :4.800
                 Max.
                        :3
                             Max. :4.500
                                            Max.
                                                           Max. :9.60
## site experiment
                       length
                                      width
                                                     height
## 1:3
        1:4
                         :1.100
                                  Min.
                                         :0.500
                   Min.
                                                 Min.
                                                      :1.50
## 2:3
                   1st Qu.:2.050
                                  1st Qu.:1.725
       2:4
                                                 1st Qu.:2.60
```

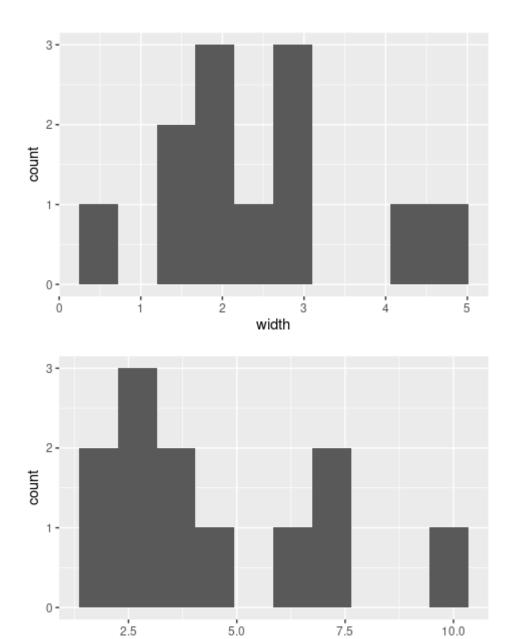
##	3:3	3:4	Median	:2.600	Median	:2.100	Median	:3.60
##	4:3		Mean	:2.558	Mean	:2.417	Mean	:4.55
##			3rd Qu.	:3.025	3rd Qu.	:2.875	3rd Qu.	:6.75
##			Max.	:4.500	Max.	:4.800	Max.	:9.60

# Ex.19)

Expected outputs:

##	# 1	A tibbl	le: 12 x 5			
##		site	experiment	length	${\tt width}$	height
##		<fct></fct>	<fct></fct>	<dbl></dbl>	<dbl></dbl>	<dbl></dbl>
##	1	1	1	2.2	1.3	9.6
##	2	1	2	2.1	2.2	7.6
##	3	1	3	2.7	1.5	2.2
##	4	2	1	3	4.5	1.5
##	5	2	2	3.1	3.1	4
##	6	2	3	2.5	2.8	3
##	7	3	1	1.9	1.8	4.5
##	8	3	2	1.1	0.5	2.3
##	9	3	3	3.5	2	7.5
##	10	4	1	2.9	2.7	3.2
##	11	4	2	4.5	4.8	6.5
##	12	4	3	1.2	1.8	2.7





height

Ex.20)

Expected output:

```
## # A tibble: 3 x 4
                 N Range_length Range_width
     experiment
##
##
     <fct>
                <int>
                             <dbl>
                                         <dbl>
## 1 1
                                           3.2
                    4
                               1.1
## 2 2
                    4
                               3.4
                                           4.3
## 3 3
                               2.3
                                           1.3
```

# Ex.21)

Expected output:

```
## # A tibble: 4 x 6
     site experiment length width height area
     <fct> <fct>
                       <dbl> <dbl> <dbl> <dbl> <
## 1 1
           2
                         2.1
                               2.2
                                      7.6 4.62
## 2 1
                                      9.6 2.86
                         2.2
                               1.3
           1
## 3 3
           3
                         3.5
                               2
                                      7.5 7
## 4 4
                                      6.5 21.6
           2
                         4.5
                               4.8
```

#### Ex.22)

Expected output:

## # A tibble: 2 x 5
## site experiment length width height
## <fct> <fct> <fct> <dbl> <dbl> <dbl> <dbl> 
## 1 1 2 2.1 2.2 7.6
## 2 4 2 4.5 4.8 6.5

#### Ex.23)

Expected output:

## # A tibble: 8 x 5 ## site experiment length width height <dbl> <dbl> <dbl> <fct> <fct> ## 1 1 2.2 1.3 9.6 1 ## 2 1 2 2.1 2.2 7.6 ## 3 2 3 4.5 1.5 1 ## 4 2 2 3.1 3.1 ## 5 3 1.9 1.8 1 4.5 ## 6 3 0.5 2.3 2 1.1 ## 7 4 1 2.9 2.7 3.2 ## 8 4 2 4.5 4.8 6.5

### Ex.24

 ${\bf Expected\ output:}$ 

## # A tibble: 12 x 5 ## site experiment length width height ## <dbl> <dbl> <dbl> <dbl> <dbl> < ## 1 2.2 1.3 9.6 1 1 ## 2 2 2.1 2.2 7.6 1 ## 3 1 3 2.7 1.5 2.2 ## 4 2 3 4.5 1 1.5 ## 5 2 2 3.1 3.1 4 ## 6 2.5 2 3 2.8 3 ## 7 1.9 3 1 1.8 4.5 ## 8 3 2 1.1 0.5 2.3 ## 9 3 3 3.5 2 7.5 ## 10 4 1 2.9 2.7 3.2 ## 11 2 4.5 4.8 6.5 ## 12 4 3 1.2 1.8 2.7

Ex. 25)
Expected output:

##	# A	tibb	le: 12 x 7					
##		site	experiment	length	${\tt width}$	height	area	volume
##		<dbl></dbl>	<dbl></dbl>	<dbl></dbl>	<dbl></dbl>	<dbl></dbl>	<dbl></dbl>	<dbl></dbl>
##	1	1	1	2.2	1.3	9.6	2.86	27.5
##	2	1	2	2.1	2.2	7.6	4.62	35.1
##	3	1	3	2.7	1.5	2.2	4.05	8.91
##	4	2	1	3	4.5	1.5	13.5	20.2
##	5	2	2	3.1	3.1	4	9.61	38.4
##	6	2	3	2.5	2.8	3	7	21
##	7	3	1	1.9	1.8	4.5	3.42	15.4
##	8	3	2	1.1	0.5	2.3	0.55	1.26
##	9	3	3	3.5	2	7.5	7	52.5
##	10	4	1	2.9	2.7	3.2	7.83	25.1
##	11	4	2	4.5	4.8	6.5	21.6	140.
##	12	4	3	1.2	1.8	2.7	2.16	5.83