

## Worksheet #6

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1. Create a data frame for the table below. Show your solution.

```
students_data <- data.frame (  
  Students = c(1,2,3,4,5,6,7,8,9,10),  
  preTest = c(55,54,47,57,51,61,57,54,63,58),  
  postTest = c(61,60,56,63,56,63,59,56,62,61)  
)
```

students\_data

##	Students	preTest	postTest
## 1	1	55	61
## 2	2	54	60
## 3	3	47	56
## 4	4	57	63
## 5	5	51	56
## 6	6	61	63
## 7	7	57	59
## 8	8	54	56
## 9	9	63	62
## 10	10	58	61

#a. Compute the descriptive statistics using different packages (Hmisc and pastecs). Write the codes and its result

```
library(Hmisc)
```

```
## Warning: package 'Hmisc' was built under R version 4.3.2
```

```
##
```

```
## Attaching package: 'Hmisc'
```

```
## The following objects are masked from 'package:base':
```

```
##
```

```
##      format.pval, units
```

```
library(pastecs)
```

```
## Warning: package 'pastecs' was built under R version 4.3.2
```

```
hmiscSTATS<-describe(students_data)
```

```
pasticsSTATS <- stat.desc(students_data)
```

```
hmiscSTATS
```

```

## students_data
##
## 3 Variables      10 Observations
## -----
##
## Students
##      n missing distinct      Info      Mean      Gmd      .05      .10
##      10      0       10       1      5.5     3.667     1.45     1.90
##      .25     .50     .75     .90     .95
##      3.25     5.50     7.75     9.10     9.55
##
## Value      1  2  3  4  5  6  7  8  9 10
## Frequency  1  1  1  1  1  1  1  1  1  1
## Proportion 0.1 0.1 0.1 0.1 0.1 0.1 0.1 0.1 0.1 0.1
##
## For the frequency table, variable is rounded to the nearest 0
## -----
##
## preTest
##      n missing distinct      Info      Mean      Gmd
##      10      0       8     0.988     55.7     5.444
##
## Value      47 51 54 55 57 58 61 63
## Frequency  1  1  2  1  2  1  1  1
## Proportion 0.1 0.1 0.2 0.1 0.2 0.1 0.1 0.1
##
## For the frequency table, variable is rounded to the nearest 0
## -----
##
## postTest
##      n missing distinct      Info      Mean      Gmd
##      10      0       6     0.964     59.7     3.311
##
## Value      56 59 60 61 62 63
## Frequency  3  1  1  2  1  2
## Proportion 0.3 0.1 0.1 0.2 0.1 0.2
##
## For the frequency table, variable is rounded to the nearest 0
## -----
##
pasticsSTATS
##
##      Students      preTest      postTest
## nbr.val 10.0000000 10.0000000 10.0000000
## nbr.null 0.0000000 0.0000000 0.0000000
## nbr.na  0.0000000 0.0000000 0.0000000
## min     1.0000000 47.0000000 56.0000000
## max     10.0000000 63.0000000 63.0000000
## range    9.0000000 16.0000000 7.0000000

```

```
## sum          55.0000000 557.0000000 597.0000000
## median       5.5000000 56.0000000 60.5000000
## mean         5.5000000 55.7000000 59.7000000
## SE.mean      0.9574271 1.46855938 0.89504811
## CI.mean.0.95 2.1658506 3.32211213 2.02473948
## var          9.1666667 21.5666667 8.01111111
## std.dev      3.0276504 4.64399254 2.83039063
## coef.var     0.5504819 0.08337509 0.04741023
```

.#2. The Department of Agriculture was studying the effects of several levels of a fertilizer on the growth of a plant. For some analyses, it might be useful to convert the fertilizer levels to an ordered factor.

#a. Write the codes and describe the result.

```
fertilizer <- c(10,10,10, 20,20,50,10,20,10,50,20,50,20,10)
ordered(fertilizer)

## [1] 10 10 10 20 20 50 10 20 10 50 20 50 20 10
## Levels: 10 < 20 < 50
```

The data\_fertilize result displays the level as an ordered factor.

#3. Abdul Hassan, president of Floor Coverings Unlimited, has asked you to study the exercise levels undertaken by 10 subjects were "l", "n", "n", "i", "l", "l", "n", "n", "i", "l"; n=none, l=light, i=intense.

### a. What is the best way to represent this in R?

```
exercise_levels <- c("l", "n", "n", "i", "l", "l", "n", "n", "i", "l")
exercise_factor <- factor(exercise_levels, levels = c("n", "l", "i"), labels = c("none", "light", "intense"))
exercise_factor

## [1] light none none intense light light none none
## [10] light
## Levels: none light intense
```

## 4. Sample of 30 tax accountants from all the states and territories of Australia and their individual state of origin is specified by a character vector of state mnemonics as:

```
state <- c("tas", "sa", "qld", "nsw", "nsw", "nt", "wa", "wa", "qld",
           "vic", "nsw", "vic", "qld", "qld", "sa", "tas", "sa", "nt",
           "wa", "vic", "qld", "nsw", "nsw", "wa", "sa", "act", "nsw",
           "vic", "vic", "act")

factorlevel <- factor(state, levels = c("act", "nsw", "nt", "qld", "sa",
```

```
"tas", "vic", "wa") )
factorlevel

## [1] tas sa qld nsw nsw nt wa wa qld vic nsw vic qld qld sa tas sa
nt wa
## [20] vic qld nsw nsw wa sa act nsw vic vic act
## Levels: act nsw nt qld sa tas vic wa
```

## 5. From #4 - continuation:

- Suppose we have the incomes of the same tax accountants in another vector (in suitably large units of money)

```
income <- c(60, 49, 40, 61, 64, 60, 59, 54,
            62, 69, 70, 42, 56, 61, 61, 61, 58, 51, 48,
            65, 49, 49, 41, 48, 52, 46, 59, 46, 58, 43)
```

- Calculate the sample mean income for each state we can now use the special function `tapply()`:

```
income_means <- tapply(income, factorlevel, mean)

income_means

##      act      nsw      nt      qld      sa      tas      vic      wa
## 44.50000 57.33333 55.50000 53.60000 55.00000 60.50000 56.00000 52.25000
```

- Copy the results and interpret.

#The result has the means of each states that has factor with levels # act nsw nt qld sa tas vic wa #50000 57.33333 55.50000 53.60000 55.00000 60.50000 56.00000 52.25000

#The data suggests that there are variations in mean incomes for tax accountants across different states in Australia, with NSW and Tasmania having higher average incomes compared to other states.

#6. Calculate the standard errors of the state income means (refer again to number 3)  
 #stdError <- function(x) sqrt(var(x)/length(x)) Note: After this assignment, the standard errors are calculated by: incster <- tapply(incomes, statef, stdError)

#a. What is the standard error? Write the codes.

```
stdError <- function(x) sqrt(var(x)/length(x))
incster <- tapply(income, factorlevel, stdError)
incster
```

```
##      act      nsw      nt      qld      sa      tas      vic      wa
## 1.500000 4.310195 4.500000 4.106093 2.738613 0.500000 5.244044 2.657536
```

#b. Interpret the result. #The precision and diversity of the mean income calculations for each state are shown by the standard errors. Greater unpredictability and less precision are typically indicated by higher standard errors, whereas lower standard errors typically indicate more precise estimations. For a thorough analysis of the data, these standard errors must be taken into account in addition to the mean incomes.

#7. Use the titanic dataset.

#a. subset the titanic dataset of those who survived and not survived. Show the codes and its result.

```
library(datasets)
data(Titanic)

titanic <- as.data.frame(Titanic)

survived <- subset(titanic, Survived == "Yes")
print(survived)

##      Class      Sex      Age Survived Freq
## 17    1st    Male  Child      Yes     5
## 18    2nd    Male  Child      Yes    11
## 19    3rd    Male  Child      Yes    13
## 20   Crew    Male  Child      Yes     0
## 21    1st  Female  Child      Yes     1
## 22    2nd  Female  Child      Yes    13
## 23    3rd  Female  Child      Yes    14
## 24   Crew  Female  Child      Yes     0
## 25    1st    Male  Adult      Yes    57
## 26    2nd    Male  Adult      Yes    14
## 27    3rd    Male  Adult      Yes    75
## 28   Crew    Male  Adult      Yes   192
## 29    1st  Female  Adult      Yes   140
## 30    2nd  Female  Adult      Yes    80
## 31    3rd  Female  Adult      Yes    76
## 32   Crew  Female  Adult      Yes    20

did_not_survive <- subset(titanic, Survived == "No")
print(did_not_survive)

##      Class      Sex      Age Survived Freq
## 1     1st    Male  Child      No     0
## 2     2nd    Male  Child      No     0
## 3     3rd    Male  Child      No    35
## 4     Crew    Male  Child      No     0
## 5     1st  Female  Child      No     0
## 6     2nd  Female  Child      No     0
## 7     3rd  Female  Child      No    17
```

```
## 8   Crew Female Child      No    0
## 9    1st   Male Adult      No   118
## 10   2nd   Male Adult      No   154
## 11   3rd   Male Adult      No   387
## 12   Crew   Male Adult      No  670
## 13   1st  Female Adult      No    4
## 14   2nd  Female Adult      No   13
## 15   3rd  Female Adult      No   89
## 16   Crew  Female Adult      No    3
```

#8. The data sets are about the breast cancer Wisconsin. The samples arrive periodically as Dr. Wolberg reports his clinical cases. The database therefore reflects this chronology [https://drive.google.com/file/d/16MFL0ehCgx2MJuNSAuB2CsBy6eDIru/view?usp=drive\\_link](https://drive.google.com/file/d/16MFL0ehCgx2MJuNSAuB2CsBy6eDIru/view?usp=drive_link)

```
library(readr)
csv.file<-"breastcancer_wisconsin.csv"
breastcancer<-read.csv("breastcancer_wisconsin.csv")
breastcancer

##           id clump_thickness size_uniformity shape_uniformity
marginal_adhesion
## 1      1000025              5              1              1
1
## 2      1002945              5              4              4
5
## 3      1015425              3              1              1
1
## 4      1016277              6              8              8
1
## 5      1017023              4              1              1
3
## 6      1017122              8             10             10
8
## 7      1018099              1              1              1
1
## 8      1018561              2              1              2
1
## 9      1033078              2              1              1
1
## 10     1033078              4              2              1
1
## 11     1035283              1              1              1
1
## 12     1036172              2              1              1
1
## 13     1041801              5              3              3
3
## 14     1043999              1              1              1
1
```

## 15	1044572	8	7	5
10				
## 16	1047630	7	4	6
4				
## 17	1048672	4	1	1
1				
## 18	1049815	4	1	1
1				
## 19	1050670	10	7	7
6				
## 20	1050718	6	1	1
1				
## 21	1054590	7	3	2
10				
## 22	1054593	10	5	5
3				
## 23	1056784	3	1	1
1				
## 24	1057013	8	4	5
1				
## 25	1059552	1	1	1
1				
## 26	1065726	5	2	3
4				
## 27	1066373	3	2	1
1				
## 28	1066979	5	1	1
1				
## 29	1067444	2	1	1
1				
## 30	1070935	1	1	3
1				
## 31	1070935	3	1	1
1				
## 32	1071760	2	1	1
1				
## 33	1072179	10	7	7
3				
## 34	1074610	2	1	1
2				
## 35	1075123	3	1	2
1				
## 36	1079304	2	1	1
1				
## 37	1080185	10	10	10
8				
## 38	1081791	6	2	1
1				
## 39	1084584	5	4	4
9				

## 40	1091262	2	5	3
3				
## 41	1096800	6	6	6
9				
## 42	1099510	10	4	3
1				
## 43	1100524	6	10	10
2				
## 44	1102573	5	6	5
6				
## 45	1103608	10	10	10
4				
## 46	1103722	1	1	1
1				
## 47	1105257	3	7	7
4				
## 48	1105524	1	1	1
1				
## 49	1106095	4	1	1
3				
## 50	1106829	7	8	7
2				
## 51	1108370	9	5	8
1				
## 52	1108449	5	3	3
4				
## 53	1110102	10	3	6
2				
## 54	1110503	5	5	5
8				
## 55	1110524	10	5	5
6				
## 56	1111249	10	6	6
3				
## 57	1112209	8	10	10
1				
## 58	1113038	8	2	4
1				
## 59	1113483	5	2	3
1				
## 60	1113906	9	5	5
2				
## 61	1115282	5	3	5
5				
## 62	1115293	1	1	1
1				
## 63	1116116	9	10	10
1				
## 64	1116132	6	3	4
1				



## 65	1116192	1	1	1
1				
## 66	1116998	10	4	2
1				
## 67	1117152	4	1	1
1				
## 68	1118039	5	3	4
1				
## 69	1120559	8	3	8
3				
## 70	1121732	1	1	1
1				
## 71	1121919	5	1	3
1				
## 72	1123061	6	10	2
8				
## 73	1124651	1	3	3
2				
## 74	1125035	9	4	5
10				
## 75	1126417	10	6	4
1				
## 76	1131294	1	1	2
1				
## 77	1132347	1	1	4
1				
## 78	1133041	5	3	1
2				
## 79	1133136	3	1	1
1				
## 80	1136142	2	1	1
1				
## 81	1137156	2	2	2
1				
## 82	1143978	4	1	1
2				
## 83	1143978	5	2	1
1				
## 84	1147044	3	1	1
1				
## 85	1147699	3	5	7
8				
## 86	1147748	5	10	6
1				
## 87	1148278	3	3	6
4				
## 88	1148873	3	6	6
6				
## 89	1152331	4	1	1
1				

## 90	1155546	2	1	1
2				
## 91	1156272	1	1	1
1				
## 92	1156948	3	1	1
2				
## 93	1157734	4	1	1
1				
## 94	1158247	1	1	1
1				
## 95	1160476	2	1	1
1				
## 96	1164066	1	1	1
1				
## 97	1165297	2	1	1
2				
## 98	1165790	5	1	1
1				
## 99	1165926	9	6	9
2				
## 100	1166630	7	5	6
10				
## 101	1166654	10	3	5
1				
## 102	1167439	2	3	4
4				
## 103	1167471	4	1	2
1				
## 104	1168359	8	2	3
1				
## 105	1168736	10	10	10
10				
## 106	1169049	7	3	4
4				
## 107	1170419	10	10	10
8				
## 108	1170420	1	6	8
10				
## 109	1171710	1	1	1
1				
## 110	1171710	6	5	4
4				
## 111	1171795	1	3	1
2				
## 112	1171845	8	6	4
3				
## 113	1172152	10	3	3
10				
## 114	1173216	10	10	10
3				

## 115	1173235	3	3	2
1				
## 116	1173347	1	1	1
1				
## 117	1173347	8	3	3
1				
## 118	1173509	4	5	5
10				
## 119	1173514	1	1	1
1				
## 120	1173681	3	2	1
1				
## 121	1174057	1	1	2
2				
## 122	1174057	4	2	1
1				
## 123	1174131	10	10	10
2				
## 124	1174428	5	3	5
1				
## 125	1175937	5	4	6
7				
## 126	1176406	1	1	1
1				
## 127	1176881	7	5	3
7				
## 128	1177027	3	1	1
1				
## 129	1177399	8	3	5
4				
## 130	1177512	1	1	1
1				
## 131	1178580	5	1	3
1				
## 132	1179818	2	1	1
1				
## 133	1180194	5	10	8
10				
## 134	1180523	3	1	1
1				
## 135	1180831	3	1	1
1				
## 136	1181356	5	1	1
1				
## 137	1182404	4	1	1
1				
## 138	1182410	3	1	1
1				
## 139	1183240	4	1	2
1				

## 140	1183246	1	1	1
1				
## 141	1183516	3	1	1
1				
## 142	1183911	2	1	1
1				
## 143	1183983	9	5	5
4				
## 144	1184184	1	1	1
1				
## 145	1184241	2	1	1
1				
## 146	1184840	1	1	3
1				
## 147	1185609	3	4	5
2				
## 148	1185610	1	1	1
1				
## 149	1187457	3	1	1
3				
## 150	1187805	8	8	7
4				
## 151	1188472	1	1	1
1				
## 152	1189266	7	2	4
1				
## 153	1189286	10	10	8
6				
## 154	1190394	4	1	1
1				
## 155	1190485	1	1	1
1				
## 156	1192325	5	5	5
6				
## 157	1193091	1	2	2
1				
## 158	1193210	2	1	1
1				
## 159	1193683	1	1	2
1				
## 160	1196295	9	9	10
3				
## 161	1196915	10	7	7
4				
## 162	1197080	4	1	1
1				
## 163	1197270	3	1	1
1				
## 164	1197440	1	1	1
2				

## 165	1197510	5	1	1
1				
## 166	1197979	4	1	1
1				
## 167	1197993	5	6	7
8				
## 168	1198128	10	8	10
10				
## 169	1198641	3	1	1
1				
## 170	1199219	1	1	1
2				
## 171	1199731	3	1	1
1				
## 172	1199983	1	1	1
1				
## 173	1200772	1	1	1
1				
## 174	1200847	6	10	10
10				
## 175	1200892	8	6	5
4				
## 176	1200952	5	8	7
7				
## 177	1201834	2	1	1
1				
## 178	1201936	5	10	10
3				
## 179	1202125	4	1	1
1				
## 180	1202812	5	3	3
3				
## 181	1203096	1	1	1
1				
## 182	1204242	1	1	1
1				
## 183	1204898	6	1	1
1				
## 184	1205138	5	8	8
8				
## 185	1205579	8	7	6
4				
## 186	1206089	2	1	1
1				
## 187	1206695	1	5	8
6				
## 188	1206841	10	5	6
10				
## 189	1207986	5	8	4
10				

## 190	1208301	1	2	3
1				
## 191	1210963	10	10	10
8				
## 192	1211202	7	5	10
10				
## 193	1212232	5	1	1
1				
## 194	1212251	1	1	1
1				
## 195	1212422	3	1	1
1				
## 196	1212422	4	1	1
1				
## 197	1213375	8	4	4
5				
## 198	1213383	5	1	1
4				
## 199	1214092	1	1	1
1				
## 200	1214556	3	1	1
1				
## 201	1214966	9	7	7
5				
## 202	1216694	10	8	8
4				
## 203	1216947	1	1	1
1				
## 204	1217051	5	1	1
1				
## 205	1217264	1	1	1
1				
## 206	1218105	5	10	10
9				
## 207	1218741	10	10	9
3				
## 208	1218860	1	1	1
1				
## 209	1218860	1	1	1
1				
## 210	1219406	5	1	1
1				
## 211	1219525	8	10	10
10				
## 212	1219859	8	10	8
8				
## 213	1220330	1	1	1
1				
## 214	1221863	10	10	10
10				

## 215	1222047	10	10	10
10				
## 216	1222936	8	7	8
7				
## 217	1223282	1	1	1
1				
## 218	1223426	1	1	1
1				
## 219	1223793	6	10	7
7				
## 220	1223967	6	1	3
1				
## 221	1224329	1	1	1
2				
## 222	1225799	10	6	4
3				
## 223	1226012	4	1	1
3				
## 224	1226612	7	5	6
3				
## 225	1227210	10	5	5
6				
## 226	1227244	1	1	1
1				
## 227	1227481	10	5	7
4				
## 228	1228152	8	9	9
5				
## 229	1228311	1	1	1
1				
## 230	1230175	10	10	10
3				
## 231	1230688	7	4	7
4				
## 232	1231387	6	8	7
5				
## 233	1231706	8	4	6
3				
## 234	1232225	10	4	5
5				
## 235	1236043	3	3	2
1				
## 236	1241232	3	1	4
1				
## 237	1241559	10	8	8
2				
## 238	1241679	9	8	8
5				
## 239	1242364	8	10	10
8				

## 240	1243256	10	4	3
2				
## 241	1270479	5	1	3
3				
## 242	1276091	3	1	1
3				
## 243	1277018	2	1	1
1				
## 244	128059	1	1	1
1				
## 245	1285531	1	1	1
1				
## 246	1287775	5	1	1
2				
## 247	144888	8	10	10
8				
## 248	145447	8	4	4
1				
## 249	167528	4	1	1
1				
## 250	169356	3	1	1
1				
## 251	183913	1	2	2
1				
## 252	191250	10	4	4
10				
## 253	1017023	6	3	3
5				
## 254	1100524	6	10	10
2				
## 255	1116116	9	10	10
1				
## 256	1168736	5	6	6
2				
## 257	1182404	3	1	1
1				
## 258	1182404	3	1	1
1				
## 259	1198641	3	1	1
1				
## 260	242970	5	7	7
1				
## 261	255644	10	5	8
10				
## 262	263538	5	10	10
6				
## 263	274137	8	8	9
4				
## 264	303213	10	4	4
10				



## 265	314428	7	9	4
10				
## 266	1182404	5	1	4
1				
## 267	1198641	10	10	6
3				
## 268	320675	3	3	5
2				
## 269	324427	10	8	8
2				
## 270	385103	1	1	1
1				
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## 560	2	1	2	1	1
2					
## 561	2	1	3	1	1
2					
## 562	2	1	3	1	1
2					
## 563	2	1	3	1	1
2					
## 564	2	1	2	1	1
2					



## 565 2	2	1	3	2	1
## 566 4	5	10	10	10	1
## 567 2	2	1	3	1	1
## 568 2	2	3	2	1	1
## 569 4	6	10	2	5	2
## 570 4	6	5	10	3	1
## 571 4	8	10	8	2	1
## 572 4	3	10	9	10	2
## 573 2	2	1	2	1	1
## 574 2	2	1	2	1	1
## 575 4	4	2	7	7	1
## 576 2	2	1	3	1	1
## 577 2	2	1	2	1	1
## 578 2	2	1	2	1	1
## 579 2	2	1	2	1	1
## 580 2	2	1	3	1	1
## 581 2	2	1	2	1	1
## 582 4	5	10	7	5	1
## 583 4	4	10	6	10	1
## 584 2	2	1	1	1	1
## 585 2	3	1	1	1	1
## 586 2	2	1	1	1	1
## 587 4	6	10	10	10	1
## 588 2	2	1	2	2	1
## 589 4	6	3	4	1	1

## 590 2	2	1	1	1	1
## 591 4	4	1	10	1	1
## 592 4	4	10	7	6	1
## 593 4	3	10	4	1	1
## 594 2	2	1	1	1	1
## 595 4	4	10	7	1	1
## 596 2	2	1	2	1	1
## 597 2	2	1	2	1	1
## 598 2	2	1	3	1	1
## 599 2	2	1	2	1	1
## 600 2	1	1	1	1	1
## 601 2	2	1	2	1	1
## 602 2	1	1	2	1	1
## 603 2	2	1	2	1	1
## 604 4	4	1	8	10	1
## 605 4	5	10	8	1	2
## 606 4	5	8	7	8	3
## 607 2	2	1	1	1	1
## 608 2	2	1	1	1	1
## 609 4	10	10	10	1	1
## 610 2	2	1	1	1	1
## 611 4	3	10	7	1	2
## 612 4	5	2	8	5	1
## 613 4	6	10	10	10	10
## 614 2	2	1	2	1	1

## 615 2	1	1	2	1	1
## 616 2	2	1	2	1	1
## 617 2	2	1	2	1	1
## 618 2	1	?	1	1	1
## 619 2	2	1	2	1	1
## 620 2	2	1	2	1	1
## 621 2	2	1	2	1	1
## 622 2	3	2	6	1	1
## 623 2	2	1	2	1	1
## 624 2	2	1	1	1	1
## 625 2	1	1	2	1	1
## 626 2	3	4	1	1	1
## 627 4	7	6	7	7	3
## 628 2	2	5	1	1	1
## 629 2	2	1	1	1	1
## 630 2	2	1	1	1	1
## 631 2	2	1	1	1	1
## 632 2	2	1	2	1	1
## 633 2	2	1	1	1	1
## 634 4	5	3	5	10	1
## 635 2	2	1	1	1	1
## 636 2	2	1	1	1	1
## 637 4	7	1	10	10	3
## 638 2	2	2	2	1	1
## 639 2	2	1	1	1	1

## 640	2	1	1	1	1
2					
## 641	2	1	1	1	1
2					
## 642	2	1	2	1	1
2					
## 643	2	1	2	1	1
2					
## 644	2	1	1	1	1
2					
## 645	2	1	1	1	1
2					
## 646	2	1	2	1	1
2					
## 647	2	1	1	1	1
2					
## 648	2	1	1	1	1
2					
## 649	10	2	10	10	10
4					
## 650	2	1	2	1	1
2					
## 651	3	4	1	1	1
2					
## 652	2	1	2	1	1
2					
## 653	2	1	2	2	1
2					
## 654	2	1	2	1	1
2					
## 655	2	1	3	1	1
2					
## 656	2	1	2	1	1
2					
## 657	2	1	2	1	1
2					
## 658	8	1	3	6	1
2					
## 659	3	10	7	2	3
4					
## 660	2	1	1	1	1
2					
## 661	2	1	2	1	1
2					
## 662	2	1	3	1	1
2					
## 663	2	1	2	1	1
2					
## 664	2	1	2	1	1
2					

## 665 2	2	1	2	1	1
## 666 2	2	1	1	1	1
## 667 2	2	1	1	1	2
## 668 2	2	1	3	1	1
## 669 4	6	1	7	10	3
## 670 4	5	5	7	10	1
## 671 4	5	8	7	4	1
## 672 2	2	1	3	1	1
## 673 2	2	1	3	1	1
## 674 2	3	1	1	1	1
## 675 2	2	1	2	1	1
## 676 2	2	1	1	1	1
## 677 2	2	1	2	1	1
## 678 2	2	1	1	1	1
## 679 2	2	1	1	1	1
## 680 2	2	1	1	1	1
## 681 4	5	10	10	10	7
## 682 4	4	10	5	6	3
## 683 2	2	1	3	2	1
## 684 2	2	1	1	1	1
## 685 2	2	1	1	1	1
## 686 2	2	1	1	1	1
## 687 2	2	1	1	1	1
## 688 2	2	1	2	3	1
## 689 2	2	1	1	1	1

```
## 690      2      1      1      1      8
2
## 691      2      1      1      1      1
2
## 692      4      5      4      4      1
4
## 693      2      1      1      1      1
2
## 694      2      1      2      1      2
2
## 695      3      2      1      1      1
2
## 696      2      1      1      1      1
2
## 697      7      3      8     10      2
4
## 698      3      4     10      6      1
4
## 699      4      5     10      4      1
4
```

```
summary(breastcancer)
```

```
##      id      clump_thickness  size_uniformity  shape_uniformity
## Min.   : 61634  Min.   : 1.000  Min.   : 1.000  Min.   : 1.000
## 1st Qu.: 870688 1st Qu.: 2.000  1st Qu.: 1.000  1st Qu.: 1.000
## Median : 1171710 Median : 4.000  Median : 1.000  Median : 1.000
## Mean   : 1071704 Mean   : 4.418  Mean   : 3.134  Mean   : 3.207
## 3rd Qu.: 1238298 3rd Qu.: 6.000  3rd Qu.: 5.000  3rd Qu.: 5.000
## Max.   :13454352 Max.   :10.000  Max.   :10.000  Max.   :10.000
## marginal_adhesion epithelial_size bare_nucleoli  bland_chromatin
## Min.   : 1.000  Min.   : 1.000  Length:699  Min.   : 1.000
## 1st Qu.: 1.000  1st Qu.: 2.000  Class :character  1st Qu.: 2.000
## Median : 1.000  Median : 2.000  Mode  :character  Median : 3.000
## Mean   : 2.807  Mean   : 3.216  Mean   : 3.438
## 3rd Qu.: 4.000  3rd Qu.: 4.000  3rd Qu.: 5.000
## Max.   :10.000  Max.   :10.000  Max.   :10.000
## normal_nucleoli  mitoses      class
## Min.   : 1.000  Min.   : 1.000  Min.   :2.00
## 1st Qu.: 1.000  1st Qu.: 1.000  1st Qu.:2.00
## Median : 1.000  Median : 1.000  Median :2.00
## Mean   : 2.867  Mean   : 1.589  Mean   :2.69
## 3rd Qu.: 4.000  3rd Qu.: 1.000  3rd Qu.:4.00
## Max.   :10.000  Max.   :10.000  Max.   :4.00
```

#a. describe what is the dataset all about. #ANSWER: The dataset 'breastcancer\_wisconsin' is a database of clinical reports of breastcancer cases. The dataset contains various features or characteristics of breast cancer tumors, including information about their size, shape, adhesion properties, cell characteristics, and mitotic activity. The goal might be to analyze

these features to predict or understand the nature of the tumors, particularly whether they are benign or malignant.

#d. Compute the descriptive statistics using different packages. Find the values of: #d.1 Standard error of the mean for clump thickness. #Using stdError function

```
clump_thickness <- breastcancer$clump_thickness
stderror_clump_thickness <- stdError(clump_thickness)
stderror_clump_thickness

## [1] 0.1065011
```

#d.2 Coefficient of variability for Marginal Adhesion. #Using mean and standard deviation to get the Coefficient of Variation.

```
marginal_adhesion <- breastcancer$marginal_adhesion
mean <- mean(marginal_adhesion)
sd <- sd(marginal_adhesion)
cv <- sd / mean
cv

## [1] 1.017283

cv<-cv*100
cv

## [1] 101.7283
```

#d.3 Number of null values of Bare Nuclei.

```
bare_nuclei_data <- breastcancer$bare_nucleoli
null_values <- sum(is.na(bare_nuclei_data))
null_values

## [1] 15
```

#d.4 Mean and standard deviation for Bland Chromatin #Using mean and standard deviation

```
bland_chromatin <- breastcancer$bland_chromatin
bland_chromatin_mean <- mean(bland_chromatin)
bland_chromatin_sd <- sd(bland_chromatin)
bland_chromatin_mean

## [1] 3.437768

bland_chromatin_sd

## [1] 2.438364
```

#d.5 Confidence interval of the mean for Uniformity of Cell Shape #Using t.test function

```

uniformity_cell_shape <- breastcancer$shape_uniformity
confidence_interval <- t.test(uniformity_cell_shape, na.rm = TRUE)$conf.int
print(confidence_interval)

## [1] 2.986741 3.428138
## attr(,"conf.level")
## [1] 0.95

```

#d. How many attributes?

```

length(breastcancer)

## [1] 11

names(breastcancer)

## [1] "id" "clump_thickness" "size_uniformity"
## [4] "shape_uniformity" "marginal_adhesion" "epithelial_size"
## [7] "bare_nucleoli" "bland_chromatin" "normal_nucleoli"
## [10] "mitoses" "class"

```

#e. Find the percentage of respondents who are malignant. Interpret the results

```

str(breastcancer)

## 'data.frame': 699 obs. of 11 variables:
## $ id : int 1000025 1002945 1015425 1016277 1017023 1017122
1018099 1018561 1033078 1033078 ...
## $ clump_thickness : int 5 5 3 6 4 8 1 2 2 4 ...
## $ size_uniformity : int 1 4 1 8 1 10 1 1 1 2 ...
## $ shape_uniformity : int 1 4 1 8 1 10 1 2 1 1 ...
## $ marginal_adhesion: int 1 5 1 1 3 8 1 1 1 1 ...
## $ epithelial_size : int 2 7 2 3 2 7 2 2 2 2 ...
## $ bare_nucleoli : chr "1" "10" "2" "4" ...
## $ bland_chromatin : int 3 3 3 3 3 9 3 3 1 2 ...
## $ normal_nucleoli : int 1 2 1 7 1 7 1 1 1 1 ...
## $ mitoses : int 1 1 1 1 1 1 1 1 5 1 ...
## $ class : int 2 2 2 2 2 4 2 2 2 2 ...

malignant_percentage <- sum(breastcancer$class == 4) / nrow(breastcancer) *
100
malignant_percentage

## [1] 34.47783

```

#9. Export the data abalone to the Microsoft excel file. Copy the codes.

```

library("AppliedPredictiveModeling")

## Warning: package 'AppliedPredictiveModeling' was built under R version
4.3.2

```



```
data("abalone")
head(abalone)
```

```
##   Type LongestShell Diameter Height WholeWeight ShuckedWeight
VisceraWeight
## 1    M          0.455    0.365  0.095      0.5140      0.2245
0.1010
## 2    M          0.350    0.265  0.090      0.2255      0.0995
0.0485
## 3    F          0.530    0.420  0.135      0.6770      0.2565
0.1415
## 4    M          0.440    0.365  0.125      0.5160      0.2155
0.1140
## 5    I          0.330    0.255  0.080      0.2050      0.0895
0.0395
## 6    I          0.425    0.300  0.095      0.3515      0.1410
0.0775
##   ShellWeight Rings
## 1          0.150   15
## 2          0.070    7
## 3          0.210    9
## 4          0.155   10
## 5          0.055    7
## 6          0.120    8
```

```
summary(abalone)
```

```
##   Type      LongestShell      Diameter      Height      WholeWeight
## F:1307  Min.   :0.075    Min.   :0.0550  Min.   :0.0000  Min.   :0.0020
## I:1342  1st Qu.:0.450    1st Qu.:0.3500  1st Qu.:0.1150  1st Qu.:0.4415
## M:1528  Median :0.545    Median :0.4250  Median :0.1400  Median :0.7995
##          Mean   :0.524    Mean   :0.4079  Mean   :0.1395  Mean   :0.8287
##          3rd Qu.:0.615    3rd Qu.:0.4800  3rd Qu.:0.1650  3rd Qu.:1.1530
##          Max.   :0.815    Max.   :0.6500  Max.   :1.1300  Max.   :2.8255
## ShuckedWeight VisceraWeight ShellWeight Rings
## Min.   :0.0010  Min.   :0.0005  Min.   :0.0015  Min.   : 1.000
## 1st Qu.:0.1860  1st Qu.:0.0935  1st Qu.:0.1300  1st Qu.: 8.000
## Median :0.3360  Median :0.1710  Median :0.2340  Median : 9.000
## Mean   :0.3594  Mean   :0.1806  Mean   :0.2388  Mean   : 9.934
## 3rd Qu.:0.5020  3rd Qu.:0.2530  3rd Qu.:0.3290  3rd Qu.:11.000
## Max.   :1.4880  Max.   :0.7600  Max.   :1.0050  Max.   :29.000
```

```
Abalone_excel <- "~/rPorgrammingCodes/Activities/worksheet
6/Abalone_excel.xlsx"
writexl::write_xlsx(abalone, Abalone_excel, col_names = TRUE)
```