

# RWorksheet\_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)  
)
```

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

```
#install.packages("Hmisc")  
library(Hmisc)
```

```
##  
## Attaching package: 'Hmisc'  
## The following objects are masked from 'package:base':  
##  
##      format.pval, units
```

```
#install.packages("pastecs")  
library(pastecs)
```

```
stats_hmisc<-describe(students_data)  
stats_pastics <- stat.desc(students_data)
```

*#2. The Department of Agriculture was studying the effects of several levels of a fertilizer on the grow*

*#a. Write the codes and describe the result.*

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

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

*# The result of the 'ordered' function applied to the 'data\_fertilize' vector indicates that the fertil*

*#3. Abdul Hassan, president of Floor Coverings Unlimited, has asked you to study the ex-ercise levels u*

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

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

```
exercise_factor
```

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

*# 4. Sample of 30 tax accountants from all the states and territories of Australia and their individual*

```
tax_accountants <- 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")
```

```
factor_level <- factor(tax_accountants, levels = c("act", "nsw", "nt", "qld", "sa", "tas", "vic", "wa"))
factor_level
```

```
## [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
```

*#the factor\_level variable result is factor with level.*

*# 5. From #4 - continuation:*

*# • Suppose we have the incomes of the same tax accountants in another vector (in suitably large units)*

```
incomes <- 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)
```

*# a. Calculate the sample mean income for each state we can now use the special function tapply():*

```
mean_income <- tapply(incomes, factor_level, mean)
```

```
mean_income
```

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

*# b. 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
```

*#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 ca*

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

```
stdError <- function(x) sqrt(var(x)/length(x))
incster <- tapply(incomes, factor_level, 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.*

*#answer. The outcome presents the computed standard errors associated with the mean incomes of differen*

*#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")
```

```
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

```
not_survived <- subset(Titanic, Survived == "No")
```

```
not_survived
```

##	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*

```
library(readr)
```

```
csv.file<-"breastcancer_wisconsin.csv"
```

```
breastcancer_wisconsin<-read.csv("breastcancer_wisconsin.csv")
```

```
breastcancer_wisconsin
```

##	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
## 271	390840	8	4	7	1
## 272	411453	5	1	1	1
## 273	320675	3	3	5	2
## 274	428903	7	2	4	1
## 275	431495	3	1	1	1
## 276	432809	3	1	3	1
## 277	434518	3	1	1	1
## 278	452264	1	1	1	1
## 279	456282	1	1	1	1
## 280	476903	10	5	7	3
## 281	486283	3	1	1	1
## 282	486662	2	1	1	2
## 283	488173	1	4	3	10
## 284	492268	10	4	6	1
## 285	508234	7	4	5	10
## 286	527363	8	10	10	10
## 287	529329	10	10	10	10
## 288	535331	3	1	1	1
## 289	543558	6	1	3	1
## 290	555977	5	6	6	8
## 291	560680	1	1	1	1
## 292	561477	1	1	1	1
## 293	563649	8	8	8	1
## 294	601265	10	4	4	6
## 295	606140	1	1	1	1
## 296	606722	5	5	7	8
## 297	616240	5	3	4	3
## 298	61634	5	4	3	1
## 299	625201	8	2	1	1
## 300	63375	9	1	2	6
## 301	635844	8	4	10	5
## 302	636130	1	1	1	1
## 303	640744	10	10	10	7
## 304	646904	1	1	1	1
## 305	653777	8	3	4	9
## 306	659642	10	8	4	4
## 307	666090	1	1	1	1
## 308	666942	1	1	1	1
## 309	667204	7	8	7	6
## 310	673637	3	1	1	1
## 311	684955	2	1	1	1
## 312	688033	1	1	1	1
## 313	691628	8	6	4	10
## 314	693702	1	1	1	1
## 315	704097	1	1	1	1
## 316	704168	4	6	5	6
## 317	706426	5	5	5	2
## 318	709287	6	8	7	8
## 319	718641	1	1	1	1
## 320	721482	4	4	4	4
## 321	730881	7	6	3	2
## 322	733639	3	1	1	1
## 323	733639	3	1	1	1

## 324	733823	5	4	6	10
## 325	740492	1	1	1	1
## 326	743348	3	2	2	1
## 327	752904	10	1	1	1
## 328	756136	1	1	1	1
## 329	760001	8	10	3	2
## 330	760239	10	4	6	4
## 331	76389	10	4	7	2
## 332	764974	5	1	1	1
## 333	770066	5	2	2	2
## 334	785208	5	4	6	6
## 335	785615	8	6	7	3
## 336	792744	1	1	1	1
## 337	797327	6	5	5	8
## 338	798429	1	1	1	1
## 339	704097	1	1	1	1
## 340	806423	8	5	5	5
## 341	809912	10	3	3	1
## 342	810104	1	1	1	1
## 343	814265	2	1	1	1
## 344	814911	1	1	1	1
## 345	822829	7	6	4	8
## 346	826923	1	1	1	1
## 347	830690	5	2	2	2
## 348	831268	1	1	1	1
## 349	832226	3	4	4	10
## 350	832567	4	2	3	5
## 351	836433	5	1	1	3
## 352	837082	2	1	1	1
## 353	846832	3	4	5	3
## 354	850831	2	7	10	10
## 355	855524	1	1	1	1
## 356	857774	4	1	1	1
## 357	859164	5	3	3	1
## 358	859350	8	10	10	7
## 359	866325	8	10	5	3
## 360	873549	10	3	5	4
## 361	877291	6	10	10	10
## 362	877943	3	10	3	10
## 363	888169	3	2	2	1
## 364	888523	4	4	4	2
## 365	896404	2	1	1	1
## 366	897172	2	1	1	1
## 367	95719	6	10	10	10
## 368	160296	5	8	8	10
## 369	342245	1	1	3	1
## 370	428598	1	1	3	1
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## 674	3	1	1	1	1	2
## 675	2	1	2	1	1	2
## 676	2	1	1	1	1	2
## 677	2	1	2	1	1	2
## 678	2	1	1	1	1	2
## 679	2	1	1	1	1	2
## 680	2	1	1	1	1	2
## 681	5	10	10	10	7	4
## 682	4	10	5	6	3	4
## 683	2	1	3	2	1	2
## 684	2	1	1	1	1	2
## 685	2	1	1	1	1	2
## 686	2	1	1	1	1	2
## 687	2	1	1	1	1	2
## 688	2	1	2	3	1	2
## 689	2	1	1	1	1	2
## 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

*#a. describe what is the dataset all about.*

*#answer.The 'breastcancer\_wisconsin' dataset comprises clinical records of cases, providing identificat*

*#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
clumpthcknss_data <- breastcancer_wisconsin$clump_thickness
std_error_clump_thickness <- stdError(clumpthcknss_data)
std_error_clump_thickness

## [1] 0.1065011
#0.1065011

#d.2 Coefficient of variability for Marginal Adhesion.
#Using mean and standard deviation to get the Coefficient of Variation.
marginal_adhesion <- breastcancer_wisconsin$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 <- breastcancer_wisconsin$bare_nucleoli
num_null <- sum(is.na(bare_nuclei))
num_null

## [1] 15
#d.4 Mean and standard deviation for Bland Chromatin
#Using mean and standard deviation
bland_chromatin_data <- breastcancer_wisconsin$bland_chromatin
mean_bland_chromatin <- mean(bland_chromatin_data)
sd_bland_chromatin <- sd(bland_chromatin_data)
mean_bland_chromatin

## [1] 3.437768
sd_bland_chromatin

## [1] 2.438364
#d.5 Confidence interval of the mean for Uniformity of Cell Shape
#Using t.test function
uniformity_cellShape_data <- breastcancer_wisconsin$shape_uniformity
confidence_interval <- t.test(uniformity_cellShape_data, na.rm = TRUE)$conf.int
print(confidence_interval)

## [1] 2.986741 3.428138
## attr(,"conf.level")
## [1] 0.95
#d. How many attributes?
length(breastcancer_wisconsin)

## [1] 11

```

```
names(breastcancer_wisconsin)
```

```
## [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*

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

```
## [1] 34.47783
```

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

```
install.packages("AppliedPredictiveModeling")
```

```
## Installing package into '/cloud/lib/x86_64-pc-linux-gnu-library/4.3'
## (as 'lib' is unspecified)
```

```
library("AppliedPredictiveModeling")
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
```

```
getwd()

## [1] "/cloud/project/RWorksheet_6"

Abalone_excel<-"/cloud/project/RWorksheet_6/AbaloneData.xlsx"
#install.packages("writexl")
library(writexl)

write_xlsx(abalone, Abalone_excel)
```