## ROWKSHEET6

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## 2023-12-13

```
#1. Create a data frame for the table below. Show your solution.
studentsData <- data.frame (</pre>
  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")
## Installing package into '/cloud/lib/x86_64-pc-linux-gnu-library/4.3'
## (as 'lib' is unspecified)
library(Hmisc)
## Attaching package: 'Hmisc'
## The following objects are masked from 'package:base':
##
##
       format.pval, units
install.packages("pastecs")
## Installing package into '/cloud/lib/x86_64-pc-linux-gnu-library/4.3'
## (as 'lib' is unspecified)
library(pastecs)
stats_hmisc<-describe(studentsData)</pre>
stats_pastics <- stat.desc(studentsData)</pre>
#2.
#a. Write the codes and describe the result.
Fertilize <- c(10,10,10, 20,20,50,10,20,10,50,20,50,20,10)
ordered(Fertilize)
```

```
## [1] 10 10 10 20 20 50 10 20 10 50 20 50 20 10
## Levels: 10 < 20 < 50
# the Fertilize result shows the level as an ordered factor.
#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?
exercise <- c("l", "n", "n", "i", "l", "l", "n", "n", "i", "l")
exercisefactor <- factor(exercise, levels = c("n", "l", "i"), labels = c("none", "light", "intense"))</pre>
exercisefactor
## [1] light
                                intense light
              none
                        none
                                                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
state <- c("tas", "sa", "qld", "nsw", "nsw", "nt", "wa", "qld",</pre>
           "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") )</pre>
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
#the factor_with_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 \leftarrow 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():
incmeans <- tapply(incomes, factorlevel, mean)</pre>
incmeans
        act
                           nt
                                   qld
                                                     tas
                                                               vic
                 nsw
                                             sa
## 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
#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 \leftarrow function(x) \ sqrt(var(x)/length(x)) \ Note: After this assignment, the standard errors are called
#a. What is the standard error? Write the codes.
standardError <- function(x) sqrt(var(x)/length(x))</pre>
incster <- tapply(incomes, factorlevel, standardError)</pre>
incster
##
        act
                nsw
                          nt
                                  qld
                                                    tas
                                                             vic
## 1.500000 4.310195 4.500000 4.106093 2.738613 0.500000 5.244044 2.657536
#b. Interpret the result.
#ANSWER: It displays the computed standard errors for the means of state incomes. A smaller standard er
#7. Use the titanic dataset.
#a. subset the titatic dataset of those who survived and not survived. Show the codes and its result.
library(datasets)
data(Titanic)
Titanic<-as.data.frame(Titanic)</pre>
survivor<-subset(Titanic, Survived=="Yes")</pre>
survivor
##
     Class
              Sex
                    Age Survived Freq
## 17
             Male Child
                             Yes
       1st
## 18
       2nd Male Child
                             Yes
                                   11
## 19
             Male Child
                             Yes
                                   13
       3rd
## 20 Crew
             Male Child
                             Yes
                                    0
## 21 1st Female Child
                             Yes
                                   1
      2nd Female Child
## 22
                             Yes 13
## 23
       3rd Female Child
                             Yes
                                   14
## 24 Crew Female Child
                             Yes
                                   0
      1st Male Adult
## 25
                             Yes 57
## 26
       2nd Male Adult
                             Yes 14
## 27
       3rd Male Adult
                             Yes
                                  75
## 28 Crew Male Adult
                             Yes 192
                             Yes 140
## 29
      1st Female Adult
## 30
       2nd Female Adult
                             Yes 80
## 31
       3rd Female Adult
                             Yes
                                   76
## 32 Crew Female Adult
                             Yes
                                   20
notsurvivor <- subset(Titanic, Survived == "No")</pre>
notsurvivor
##
     Class
              Sex
                    Age Survived Freq
## 1
       1st Male Child
```

```
## 2
       2nd Male Child
                            No
## 3
       3rd Male Child
                            No
                                 35
## 4 Crew Male Child
                            No
                                 0
## 5
      1st Female Child
                                 0
                            No
       2nd Female Child
## 6
                            No
                                 0
## 7
      3rd Female Child
                            No
                                17
## 8 Crew Female Child
                            No
                                 0
            Male Adult
## 9
      1st
                            No 118
## 10
       2nd Male Adult
                            No 154
## 11
       3rd Male Adult
                            No 387
## 12 Crew Male Adult
                            No 670
## 13
      1st Female Adult
                                4
                            No
## 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</pre>
```

##		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
##		1054593	10	5	5	3
##	23	1056784	3	1	1	1
	24	1057013	8	4	5	1
##		1059552	1	1	1	1
##	26	1065726	5	2	3	4
##	27	1066373	3	2	1	1
##		1066979	5	1	1	1
##		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
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##	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	1170413	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
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	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
ii TT	102	1211202	,	0	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
		456282				
	279		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
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	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		2			
		684955		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
π#	JJ+	000001	2	,	10	10

## 255	055504	4	1	1	1
## 355		1	1	1	1
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##	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	1	2	1	1	2
	666	2	1	1	1	1	2
##	667	2	1	1	1	2	2
##	668	2	1	3	1	1	2
##	669	6	1	7	10	3	4
##	670	5	5	7	10	1	4
	671	5	8	7	4	1	4
	672	2	1	3	1	1	2
	673	2	1	3	1	1	2
	674	3	1	1	1	1	2
	675	2	1	2	1	1	2
	676	2	1	1	1	1	2
	677	2		2	1		2
		2	1			1	
	678		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

## summary(breastcancer\_wisconsin)

standarderrorclumpthickness

```
##
                      clump_thickness size_uniformity shape_uniformity
         id
              61634
                      Min.
                           : 1.000
                                      Min. : 1.000
                                                       Min. : 1.000
## Min.
  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
                                                              :10.000
                                                       Max.
## marginal_adhesion epithelial_size
                                     bare_nucleoli
                                                        bland_chromatin
## Min.
         : 1.000
                          : 1.000
                                      Length:699
                     Min.
                                                        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
         : 2.807
                           : 3.216
                                                              : 3.438
## Mean
                     Mean
                                                        Mean
## 3rd Qu.: 4.000
                                                        3rd Qu.: 5.000
                     3rd Qu.: 4.000
## Max.
          :10.000
                     Max.
                            :10.000
                                                        Max.
                                                               :10.000
## normal nucleoli
                       mitoses
                                         class
          : 1.000
## Min.
                    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
                          : 1.589
## Mean
         : 2.867
                                    Mean :2.69
                    Mean
## 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 'breastcancer_wisconsin' dataset comprises clinical records detailing various cases. These
#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_wisconsin$clump_thickness
standarderrorclumpthickness <- standardError(clump_thickness)</pre>
```

```
## [1] 0.1065011
```

```
#0.1065011
#d.2 Coefficient of variability for Marginal Adhesion.
#Using mean and standard deviation to get the Coefficient of Variation.
marginalAdhesion <- breastcancer_wisconsin$marginal_adhesion</pre>
mean <- mean(marginalAdhesion)</pre>
sd <- sd(marginalAdhesion)</pre>
cv <- sd / mean
CV
## [1] 1.017283
cv<-cv*100 #Getting the percentage
## [1] 101.7283
#d.3 Number of null values of Bare Nuclei.
bareNuclei <- breastcancer_wisconsin$bare_nucleoli</pre>
num_null <- sum(is.na(bareNuclei))</pre>
num_null
## [1] 15
#d.4 Mean and standard deviation for Bland Chromatin
\#Using\ mean\ and\ standard\ deviation
blandchromatinData <- breastcancer_wisconsin$bland_chromatin</pre>
mean_bland_chromatin <- mean(blandchromatinData)</pre>
sdblandchromatin <- sd(blandchromatinData)</pre>
mean_bland_chromatin
## [1] 3.437768
sdblandchromatin
## [1] 2.438364
#d.5 Confidence interval of the mean for Uniformity of Cell Shape
#Using t.test function
uniformityCellShape <- breastcancer_wisconsin$shape_uniformity</pre>
confidenceInterval <- t.test(uniformityCellShape, na.rm = TRUE)$conf.int</pre>
print(confidenceInterval)
## [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"
                                                 "normal_nucleoli"
  [7] "bare_nucleoli"
                             "bland_chromatin"
## [10] "mitoses"
                            "class"
#e. Find the percentage of respondents who are malignant. Interpret the results
malignant <- sum(breastcancer_wisconsin$class == 4) / nrow(breastcancer_wisconsin) * 100</pre>
malignant
## [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
       М
                 0.455
                          0.365 0.095
                                             0.5140
                                                           0.2245
                                                                          0.1010
## 2
        М
                 0.350
                          0.265 0.090
                                             0.2255
                                                           0.0995
                                                                          0.0485
        F
## 3
                                                           0.2565
                 0.530
                          0.420 0.135
                                             0.6770
                                                                          0.1415
## 4
        М
                 0.440
                          0.365 0.125
                                             0.5160
                                                           0.2155
                                                                          0.1140
## 5
        Ι
                 0.330
                          0.255 0.080
                                             0.2050
                                                           0.0895
                                                                          0.0395
## 6
                 0.425
                          0.300 0.095
                                             0.3515
                                                           0.1410
                                                                          0.0775
        Ι
##
     ShellWeight Rings
           0.150
## 1
## 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
```

Min.

:0.0000

1st Qu.:0.1150

Min.

:0.0020

1st Qu.:0.4415

:0.0550

1st Qu.:0.3500

## F:1307

## I:1342

:0.075

1st Qu.:0.450

Min.

```
## 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
                 VisceraWeight
                               ShellWeight
                                              Rings
## ShuckedWeight
## 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
```