

uhuru dataset

2022-10-04

1. Describing the data that we are using

We are using the data set from this study

The Data From UHURU Acacia Survey is based on a study of assessing both direct and indirect consequences of the removal of nonrandom species of the environment. The UHURU consists of three wired electric fences to provoke herbivore exclusion treatments this also has a control group which is un-fenced. The units that we are using are for height meters and for weight kg. These three Fenced treatments include “Mega” excludes elephants and giraffes only; “Meso” excludes both megaherbivores and mesoherbivores, ca. 40 kg and larger; and “Total” excludes all herbivores > 5 kg.

2. reading the data table into R

First make sure we are in the correct working directory “getwd()” “/Users/atziri/Bio 195-197/Data Science”

if it is not right set the working directory with ‘setwd()’

how do i test where the computer is ?

```
read.csv(file = "/Users/atziri/Bio 195-197/Data Science/raw-data/ACACIA_DREPANOLOBIUM_SURVEY.txt", sep = ";")
```

##	SURVEY	YEAR	SITE	BLOCK	TREATMENT	PLOT	ID	HEIGHT	AXIS1	AXIS2	CIRC
## 1	1	2012	SOUTH	1	TOTAL	S1TOTAL	581	2.25	2.75	2.15	20.0
## 2	1	2012	SOUTH	1	TOTAL	S1TOTAL	582	2.65	4.10	3.90	28.0
## 3	1	2012	SOUTH	1	TOTAL	S1TOTAL	3111	1.5	1.70	0.85	17.0
## 4	1	2012	SOUTH	1	TOTAL	S1TOTAL	3112	2.01	1.80	1.60	12.0
## 5	1	2012	SOUTH	1	TOTAL	S1TOTAL	3113	1.75	1.84	1.42	13.0
## 6	1	2012	SOUTH	1	TOTAL	S1TOTAL	3114	1.65	1.62	0.85	15.0
## 7	1	2012	SOUTH	1	TOTAL	S1TOTAL	3115	1.2	1.95	0.90	9.0
## 8	1	2012	SOUTH	1	TOTAL	S1TOTAL	3199	1.45	2.00	1.75	12.2
## 9	1	2012	SOUTH	1	MESO	S1MESO	941	1.87	2.15	1.82	13.0
## 10	1	2012	SOUTH	1	MESO	S1MESO	942	2.38	5.55	4.82	35.0
## 11	1	2012	SOUTH	1	MESO	S1MESO	943	2.58	4.90	4.24	24.0
## 12	1	2012	SOUTH	1	MESO	S1MESO	944	2.65	3.75	3.10	27.0
## 13	1	2012	SOUTH	1	MESO	S1MESO	946	2.35	2.34	2.05	20.0
## 14	1	2012	SOUTH	1	MESO	S1MESO	947	1.88	2.10	1.85	28.0
## 15	1	2012	SOUTH	1	MESO	S1MESO	3116	2.32	3.05	2.63	30.0
## 16	1	2012	SOUTH	1	MESO	S1MESO	3117	2.39	2.21	2.10	13.0
## 17	1	2012	SOUTH	1	MESO	S1MESO	3118	2.2	1.80	1.50	10.0
## 18	1	2012	SOUTH	1	MESO	S1MESO	3119	1.05	0.90	0.55	8.0
## 19	1	2012	SOUTH	1	MESO	S1MESO	3120	2	1.25	1.20	10.0
## 20	1	2012	SOUTH	1	MESO	S1MESO	3131	1.28	1.14	1.00	10.0
## 21	1	2012	SOUTH	2	OPEN	S2OPEN	341	dead	NA	NA	NA
## 22	1	2012	SOUTH	2	TOTAL	S2TOTAL	3178	1.4	2.50	2.15	18.0

## 23	1	2012	SOUTH	2	TOTAL S2TOTAL	101	1.9	3.31	2.65	15.0
## 24	1	2012	SOUTH	2	TOTAL S2TOTAL	102	1.75	2.70	2.55	16.0
## 25	1	2012	SOUTH	2	TOTAL S2TOTAL	103	1.8	2.75	2.30	16.0
## 26	1	2012	SOUTH	2	TOTAL S2TOTAL	104	2.7	4.05	4.00	35.2
## 27	1	2012	SOUTH	2	TOTAL S2TOTAL	105	2.02	2.85	1.49	17.0
## 28	1	2012	SOUTH	2	TOTAL S2TOTAL	108	1.9	3.10	2.85	19.0
## 29	1	2012	SOUTH	2	TOTAL S2TOTAL	109	1.85	2.45	1.90	19.0
## 30	1	2012	SOUTH	2	TOTAL S2TOTAL	110	1.65	1.90	1.54	17.0
## 31	1	2012	SOUTH	2	TOTAL S2TOTAL	111	1.4	2.35	1.45	14.0
## 32	1	2012	SOUTH	2	TOTAL S2TOTAL	113	2.5	3.25	2.30	22.0
## 33	1	2012	SOUTH	2	TOTAL S2TOTAL	115	2.05	5.40	4.50	33.0
## 34	1	2012	SOUTH	2	TOTAL S2TOTAL	116	2.26	3.50	3.10	33.0
## 35	1	2012	SOUTH	2	TOTAL S2TOTAL	117	2.13	2.40	2.30	20.0
## 36	1	2012	SOUTH	2	TOTAL S2TOTAL	118	1.8	3.15	2.55	22.0
## 37	1	2012	SOUTH	2	TOTAL S2TOTAL	1211	1.85	2.00	2.27	20.0
## 38	1	2012	SOUTH	2	TOTAL S2TOTAL	1212	1.5	2.15	1.80	15.0
## 39	1	2012	SOUTH	2	TOTAL S2TOTAL	1213	1.87	2.34	2.05	13.0
## 40	1	2012	SOUTH	2	TOTAL S2TOTAL	1214	1.58	1.28	0.75	11.0
## 41	1	2012	SOUTH	2	TOTAL S2TOTAL	1215	2.05	2.10	1.75	17.0
## 42	1	2012	SOUTH	2	TOTAL S2TOTAL	1216	1.75	2.45	3.28	16.0
## 43	1	2012	SOUTH	2	TOTAL S2TOTAL	1217	1.49	1.50	1.45	13.0
## 44	1	2012	SOUTH	2	TOTAL S2TOTAL	1218	1.28	2.00	0.90	10.0
## 45	1	2012	SOUTH	2	TOTAL S2TOTAL	1219	1.49	2.35	1.65	13.0
## 46	1	2012	SOUTH	2	TOTAL S2TOTAL	1220	1.07	1.20	0.95	11.0
## 47	1	2012	SOUTH	2	TOTAL S2TOTAL	1231	1.48	1.25	1.20	9.0
## 48	1	2012	SOUTH	2	TOTAL S2TOTAL	1232	1.25	1.25	0.90	10.0
## 49	1	2012	SOUTH	2	TOTAL S2TOTAL	1233	1.41	1.41	1.40	14.0
## 50	1	2012	SOUTH	2	TOTAL S2TOTAL	1234	1.6	1.60	1.30	13.0
## 51	1	2012	SOUTH	2	TOTAL S2TOTAL	1235	1.2	1.20	1.30	14.0
## 52	1	2012	SOUTH	2	TOTAL S2TOTAL	1236	1.49	1.49	1.20	8.0
## 53	1	2012	SOUTH	2	TOTAL S2TOTAL	1237	1.5	1.50	1.50	14.0
## 54	1	2012	SOUTH	2	TOTAL S2TOTAL	1238	1.65	1.65	2.00	20.0
## 55	1	2012	SOUTH	2	TOTAL S2TOTAL	1239	1.13	1.13	1.20	10.0
## 56	1	2012	SOUTH	2	TOTAL S2TOTAL	1240	1.25	1.25	0.90	10.0
## 57	1	2012	SOUTH	2	TOTAL S2TOTAL	1251	1.1	1.20	1.10	10.0
## 58	1	2012	SOUTH	2	TOTAL S2TOTAL	1252	2.2	2.70	2.40	25.0
## 59	1	2012	SOUTH	2	TOTAL S2TOTAL	1253	1.45	1.65	1.25	10.0
## 60	1	2012	SOUTH	2	TOTAL S2TOTAL	1254	1.6	2.45	2.10	13.0
## 61	1	2012	SOUTH	2	TOTAL S2TOTAL	1255	1.55	2.40	1.80	13.0
## 62	1	2012	SOUTH	2	TOTAL S2TOTAL	1256	1.5	2.40	2.15	13.0
## 63	1	2012	SOUTH	2	TOTAL S2TOTAL	1257	1.03	1.20	1.00	10.0
## 64	1	2012	SOUTH	2	TOTAL S2TOTAL	1258	2.14	1.90	1.70	13.0
## 65	1	2012	SOUTH	2	TOTAL S2TOTAL	1259	1.2	1.90	1.65	12.0
## 66	1	2012	SOUTH	2	TOTAL S2TOTAL	1260	1.05	1.10	1.00	9.0
## 67	1	2012	SOUTH	2	TOTAL S2TOTAL	2131	1.8	2.60	2.40	15.0
## 68	1	2012	SOUTH	2	TOTAL S2TOTAL	2132	1.2	1.00	0.95	7.0
## 69	1	2012	SOUTH	2	TOTAL S2TOTAL	2133	1.75	1.40	1.10	10.0
## 70	1	2012	SOUTH	2	TOTAL S2TOTAL	2134	1.45	3.10	1.80	10.0
## 71	1	2012	SOUTH	2	TOTAL S2TOTAL	2135	1.17	1.20	1.10	5.0
## 72	1	2012	SOUTH	2	TOTAL S2TOTAL	2136	2.15	3.10	2.58	22.0
## 73	1	2012	SOUTH	2	TOTAL S2TOTAL	2137	1.7	1.70	1.40	12.0
## 74	1	2012	SOUTH	2	TOTAL S2TOTAL	3132	1.98	2.85	2.70	12.0
## 75	1	2012	SOUTH	2	TOTAL S2TOTAL	3133	1.26	1.95	1.75	17.0
## 76	1	2012	SOUTH	2	TOTAL S2TOTAL	3134	1.11	1.95	1.50	10.0

## 77	1	2012	SOUTH	2	TOTAL	S2TOTAL	3135	1.14	1.32	1.05	10.0
## 78	1	2012	SOUTH	2	TOTAL	S2TOTAL	3136	1.26	1.60	1.40	10.0
## 79	1	2012	SOUTH	2	TOTAL	S2TOTAL	3137	1.3	1.40	0.80	10.0
## 80	1	2012	SOUTH	2	TOTAL	S2TOTAL	3138	1.29	1.44	1.35	13.0
## 81	1	2012	SOUTH	2	TOTAL	S2TOTAL	3139	1.31	1.35	1.15	7.0
## 82	1	2012	SOUTH	2	TOTAL	S2TOTAL	3140	1.15	1.70	1.28	10.0
## 83	1	2012	SOUTH	2	TOTAL	S2TOTAL	3151	1.87	3.40	1.85	15.0
## 84	1	2012	SOUTH	2	TOTAL	S2TOTAL	3152	1.47	2.10	1.61	8.0
## 85	1	2012	SOUTH	2	TOTAL	S2TOTAL	3153	1.05	1.79	1.50	10.0
## 86	1	2012	SOUTH	2	TOTAL	S2TOTAL	3154	2.1	4.90	3.75	25.0
## 87	1	2012	SOUTH	2	TOTAL	S2TOTAL	3155	1.99	1.80	1.35	13.0
## 88	1	2012	SOUTH	2	TOTAL	S2TOTAL	3156	1.42	1.90	1.80	14.0
## 89	1	2012	SOUTH	2	TOTAL	S2TOTAL	3157	1.5	2.11	1.75	12.0
## 90	1	2012	SOUTH	2	TOTAL	S2TOTAL	3158	1.06	1.05	0.85	4.0
## 91	1	2012	SOUTH	2	TOTAL	S2TOTAL	3159	1.49	1.50	1.15	13.0
## 92	1	2012	SOUTH	2	TOTAL	S2TOTAL	3160	1.8	1.60	1.50	14.0
## 93	1	2012	SOUTH	2	TOTAL	S2TOTAL	3171	1.93	1.74	1.20	14.0
## 94	1	2012	SOUTH	2	TOTAL	S2TOTAL	3172	1.2	1.60	1.30	10.0
## 95	1	2012	SOUTH	2	TOTAL	S2TOTAL	3173	1.65	1.25	1.10	11.0
## 96	1	2012	SOUTH	2	TOTAL	S2TOTAL	3174	1.52	1.49	1.10	12.0
## 97	1	2012	SOUTH	2	TOTAL	S2TOTAL	3175	1.43	2.05	1.54	13.0
## 98	1	2012	SOUTH	2	TOTAL	S2TOTAL	3176	1.25	1.40	1.25	13.0
## 99	1	2012	SOUTH	2	TOTAL	S2TOTAL	3177	1.88	2.65	2.64	20.0
## 100	1	2012	SOUTH	2	TOTAL	S2TOTAL	3179	1.03	1.40	0.60	13.0
## 101	1	2012	SOUTH	2	TOTAL	S2TOTAL	3180	1.1	1.30	1.20	10.0
## 102	1	2012	SOUTH	2	TOTAL	S2TOTAL	3191	1.4	1.05	1.00	10.0
## 103	1	2012	SOUTH	2	TOTAL	S2TOTAL	3192	1.05	1.55	0.90	10.0
## 104	1	2012	SOUTH	2	TOTAL	S2TOTAL	3193	1.18	1.20	1.00	7.0
## 105	1	2012	SOUTH	2	TOTAL	S2TOTAL	3194	1.4	1.30	1.85	13.0
## 106	1	2012	SOUTH	2	TOTAL	S2TOTAL	3195	1.37	2.67	2.19	19.0
## 107	1	2012	SOUTH	2	TOTAL	S2TOTAL	3196	1.32	2.15	1.55	11.0
## 108	1	2012	SOUTH	2	MEGA	S2MEGA	182	1.55	2.20	1.20	20.0
## 109	1	2012	SOUTH	2	MEGA	S2MEGA	183	1.3	1.80	0.90	8.0
## 110	1	2012	SOUTH	2	MEGA	S2MEGA	184	1.24	1.20	1.20	25.0
## 111	1	2012	SOUTH	2	MEGA	S2MEGA	185	1.5	2.10	1.75	16.0
## 112	1	2012	SOUTH	2	MEGA	S2MEGA	186	1.65	2.50	2.20	15.0
## 113	1	2012	SOUTH	2	MEGA	S2MEGA	187	2.17	2.00	1.20	15.0
## 114	1	2012	SOUTH	2	MEGA	S2MEGA	188	1.28	1.60	1.50	10.0
## 115	1	2012	SOUTH	2	MEGA	S2MEGA	189	1.07	1.50	1.50	10.0
## 116	1	2012	SOUTH	2	MEGA	S2MEGA	190	0.67	1.00	0.80	8.0
## 117	1	2012	SOUTH	2	MEGA	S2MEGA	191	0.68	0.70	0.60	4.0
## 118	1	2012	SOUTH	2	MEGA	S2MEGA	192	1.87	1.60	1.40	9.0
## 119	1	2012	SOUTH	2	MEGA	S2MEGA	193	1.35	1.90	1.50	14.0
## 120	1	2012	SOUTH	2	MEGA	S2MEGA	194	1.75	2.10	2.10	15.0
## 121	1	2012	SOUTH	2	MESO	S2MESO	462	1.75	3.30	2.50	23.0
## 122	1	2012	SOUTH	2	MESO	S2MESO	463	1.64	2.30	2.00	14.0
## 123	1	2012	SOUTH	2	MESO	S2MESO	2138	1.42	0.90	0.80	10.0
## 124	1	2012	SOUTH	3	OPEN	S3OPEN	1301	dead	NA	NA	NA
## 125	1	2012	SOUTH	3	OPEN	S3OPEN	1302	0.9	1.30	1.10	11.0
## 126	1	2012	SOUTH	3	TOTAL	S3TOTAL	1061	dead	NA	NA	NA
## 127	1	2012	SOUTH	3	TOTAL	S3TOTAL	1062	1.8	2.60	2.60	15.0
## 128	1	2012	SOUTH	3	TOTAL	S3TOTAL	1063	2.47	3.10	2.20	18.0
## 129	1	2012	SOUTH	3	TOTAL	S3TOTAL	1064	2.15	1.60	1.10	17.0
## 130	1	2012	SOUTH	3	TOTAL	S3TOTAL	1066	1.7	2.50	2.15	15.0

## 131	1 2012 SOUTH	3	TOTAL S3TOTAL 1066	1.9	1.80	1.50	20.0
## 132	1 2012 SOUTH	3	TOTAL S3TOTAL 1067	1.95	2.10	1.90	13.0
## 133	1 2012 SOUTH	3	TOTAL S3TOTAL 1068	1.8	1.70	1.40	13.0
## 134	1 2012 SOUTH	3	TOTAL S3TOTAL 1069	1.4	2.00	1.60	14.0
## 135	1 2012 SOUTH	3	TOTAL S3TOTAL 1070	1	1.30	1.20	7.0
## 136	1 2012 SOUTH	3	TOTAL S3TOTAL 2139	1.75	1.20	1.10	13.0
## 137	1 2012 SOUTH	3	TOTAL S3TOTAL 2140	1.28	1.50	0.95	4.0
## 138	1 2012 SOUTH	3	TOTAL S3TOTAL 2151	1	1.40	1.20	4.0
## 139	1 2012 SOUTH	3	TOTAL S3TOTAL 2152	1.45	1.50	1.30	10.0
## 140	1 2012 SOUTH	3	TOTAL S3TOTAL 2153	1	1.00	0.75	8.0
## 141	1 2012 SOUTH	3	TOTAL S3TOTAL 2154	1.03	1.00	0.90	6.0
## 142	1 2012 SOUTH	3	TOTAL S3TOTAL 2155	1.51	2.00	1.80	12.0
## 143	1 2012 SOUTH	3	TOTAL S3TOTAL 2156	1.17	1.10	0.90	10.0
## 144	1 2012 SOUTH	3	TOTAL S3TOTAL 2157	1.33	1.90	1.85	14.0
## 145	1 2012 SOUTH	3	TOTAL S3TOTAL 2158	1.3	1.10	0.85	8.0
## 146	1 2012 SOUTH	3	TOTAL S3TOTAL 2159	1.13	1.10	0.90	10.0
## 147	1 2012 SOUTH	3	TOTAL S3TOTAL 2160	1.58	1.40	1.40	13.0
## 148	1 2012 SOUTH	3	TOTAL S3TOTAL 2171	1.06	1.40	1.00	5.0
## 149	1 2012 SOUTH	3	TOTAL S3TOTAL 2172	1.05	1.40	0.95	7.0
## 150	1 2012 SOUTH	3	TOTAL S3TOTAL 2173	1.45	1.60	1.10	6.0
## 151	1 2012 SOUTH	3	TOTAL S3TOTAL 2174	1.15	1.10	0.90	5.0
## 152	1 2012 SOUTH	3	TOTAL S3TOTAL 2175	1.42	1.45	1.30	13.0
## 153	1 2012 SOUTH	3	TOTAL S3TOTAL 2176	1.02	1.20	1.00	8.0
## 154	1 2012 SOUTH	3	TOTAL S3TOTAL 2177	1.4	1.20	1.00	9.0
## 155	1 2012 SOUTH	3	TOTAL S3TOTAL 2178	1.45	2.10	2.05	15.0
## 156	1 2012 SOUTH	3	MESO S3MESO 1421	1.95	2.20	1.60	13.0
## 157	1 2012 SOUTH	3	MESO S3MESO 1422	dead	NA	NA	NA
##	FLOWERS	BUDS	FRUITS	ANT			
## 1	0	0	10	CS			
## 2	0	0	150	TP			
## 3	2	1	50	TP			
## 4	0	0	75	CS			
## 5	0	0	20	CS			
## 6	0	0	0	E			
## 7	0	0	0	CS			
## 8	0	0	25	CS			
## 9	0	0	0	TP			
## 10	0	0	50	TP			
## 11	0	0	5	CS			
## 12	0	0	60	TP			
## 13	0	0	60	TP			
## 14	2	0	60	CS			
## 15	2	0	0	CS			
## 16	0	0	0	TP			
## 17	0	0	0	TP			
## 18	0	0	0	CS			
## 19	0	0	0	CM			
## 20	0	0	0	TP			
## 21	NA	NA	NA				
## 22	0	0	5	CS			
## 23	0	0	45	CS			
## 24	40	50	35	CS			
## 25	8	2	65	CS			
## 26	0	0	20	TP			

## 27	0	0	70	CS
## 28	0	0	125	CM
## 29	0	0	200	CM
## 30	0	0	10	CS
## 31	0	0	0	CS
## 32	0	0	35	TP
## 33	0	0	300	CM
## 34	2	2	100	CS
## 35	0	0	30	CM
## 36	0	0	50	TP
## 37	0	0	10	CM
## 38	0	0	25	CS
## 39	0	0	15	TP
## 40	0	0	0	TP
## 41	0	0	15	TP
## 42	0	0	0	TP
## 43	0	0	40	TP
## 44	0	0	0	TP
## 45	0	0	15	CM
## 46	0	0	0	CM
## 47	0	0	0	TP
## 48	0	0	0	TP
## 49	0	0	1	TP
## 50	0	0	20	TP
## 51	0	0	0	TP
## 52	0	0	0	TP
## 53	0	0	20	TP
## 54	0	0	0	TP
## 55	0	0	0	CN
## 56	0	0	0	CN
## 57	0	0	0	TP
## 58	0	0	5	TP
## 59	0	0	0	TP
## 60	0	0	25	TP
## 61	0	0	25	TP
## 62	0	0	20	TP
## 63	0	0	0	TP
## 64	0	0	10	CS
## 65	1	0	25	CS
## 66	0	0	0	TP
## 67	0	0	10	TP
## 68	0	0	0	TP
## 69	0	0	0	TP
## 70	0	0	0	TP
## 71	0	0	0	TP
## 72	0	0	0	CS
## 73	0	0	0	CS
## 74	0	0	25	AB_TP
## 75	0	0	0	TP
## 76	0	0	0	TP
## 77	0	0	0	TP
## 78	0	0	0	CS
## 79	0	0	0	CS
## 80	0	0	0	CS

## 81	0	0	0	CS
## 82	0	0	5	CS
## 83	6	0	0	CS
## 84	0	0	0	CS
## 85	0	0	1	CS
## 86	0	0	25	CS
## 87	0	0	0	CS
## 88	0	0	0	CS
## 89	0	0	10	CS
## 90	0	0	0	CS
## 91	0	0	35	CS
## 92	0	0	0	CS
## 93	0	0	0	CS
## 94	0	0	0	CS
## 95	0	0	0	CS
## 96	0	0	20	CS
## 97	0	0	0	CS
## 98	0	0	0	CM
## 99	0	0	100	CM
## 100	0	0	0	CS
## 101	0	0	0	CS
## 102	0	0	0	CS
## 103	0	0	0	CM
## 104	0	0	0	TP
## 105	0	0	30	CS
## 106	0	0	50	TP
## 107	0	0	10	CS
## 108	0	0	0	CS
## 109	0	0	15	CS
## 110	0	0	10	CS
## 111	5	0	200	CS
## 112	0	0	80	CS
## 113	0	0	150	TP
## 114	0	0	40	TP
## 115	0	0	60	TP
## 116	0	0	0	CS
## 117	0	0	0	TP
## 118	0	0	40	CS
## 119	0	0	20	CS
## 120	0	0	75	TP
## 121	0	0	20	CM
## 122	0	0	0	TP
## 123	0	0	0	E
## 124	NA	NA	NA	
## 125	0	0	0	TP
## 126	NA	NA	NA	
## 127	0	0	50	TP
## 128	0	0	0	TP
## 129	0	0	0	TP
## 130	0	0	2	TP
## 131	0	0	25	TP
## 132	0	0	0	TP
## 133	0	0	0	TP
## 134	0	0	0	TP

```

## 135      0      0      0      TP
## 136      0      0      0      TP
## 137      0      0      0      TP
## 138      0      0      0      TP
## 139      0      0      0      TP
## 140      0      0      0      TP
## 141      0      0      0      TP
## 142      0      0      0      TP
## 143      0      0      0      TP
## 144      0      0      0      TP
## 145      0      0      0      TP
## 146      0      0      0      TP
## 147      0      0      0      TP
## 148      0      0      8      TP
## 149      0      0      0      TP
## 150      0      0      0      TP
## 151      0      0      0      TP
## 152      0      0      0      TP
## 153      0      0      0      TP
## 154      0      0      0      TP
## 155      0      0     20      TP
## 156      0      0      2      CS
## 157     NA     NA     NA

```

```
read.csv(file = "../raw-data/ACACIA_DREPANOLOBIUM_SURVEY.txt", sep = "\t")
```

```

##      SURVEY YEAR  SITE BLOCK TREATMENT  PLOT  ID HEIGHT AXIS1 AXIS2 CIRC
## 1      1 2012 SOUTH      1      TOTAL S1TOTAL  581   2.25  2.75  2.15 20.0
## 2      1 2012 SOUTH      1      TOTAL S1TOTAL  582   2.65  4.10  3.90 28.0
## 3      1 2012 SOUTH      1      TOTAL S1TOTAL 3111    1.5  1.70  0.85 17.0
## 4      1 2012 SOUTH      1      TOTAL S1TOTAL 3112    2.01  1.80  1.60 12.0
## 5      1 2012 SOUTH      1      TOTAL S1TOTAL 3113    1.75  1.84  1.42 13.0
## 6      1 2012 SOUTH      1      TOTAL S1TOTAL 3114    1.65  1.62  0.85 15.0
## 7      1 2012 SOUTH      1      TOTAL S1TOTAL 3115    1.2  1.95  0.90  9.0
## 8      1 2012 SOUTH      1      TOTAL S1TOTAL 3199    1.45  2.00  1.75 12.2
## 9      1 2012 SOUTH      1      MESO  S1MESO  941    1.87  2.15  1.82 13.0
## 10     1 2012 SOUTH      1      MESO  S1MESO  942    2.38  5.55  4.82 35.0
## 11     1 2012 SOUTH      1      MESO  S1MESO  943    2.58  4.90  4.24 24.0
## 12     1 2012 SOUTH      1      MESO  S1MESO  944    2.65  3.75  3.10 27.0
## 13     1 2012 SOUTH      1      MESO  S1MESO  946    2.35  2.34  2.05 20.0
## 14     1 2012 SOUTH      1      MESO  S1MESO  947    1.88  2.10  1.85 28.0
## 15     1 2012 SOUTH      1      MESO  S1MESO 3116    2.32  3.05  2.63 30.0
## 16     1 2012 SOUTH      1      MESO  S1MESO 3117    2.39  2.21  2.10 13.0
## 17     1 2012 SOUTH      1      MESO  S1MESO 3118    2.2  1.80  1.50 10.0
## 18     1 2012 SOUTH      1      MESO  S1MESO 3119    1.05  0.90  0.55  8.0
## 19     1 2012 SOUTH      1      MESO  S1MESO 3120    2    1.25  1.20 10.0
## 20     1 2012 SOUTH      1      MESO  S1MESO 3131    1.28  1.14  1.00 10.0
## 21     1 2012 SOUTH      2      OPEN  S2OPEN  341   dead    NA    NA    NA
## 22     1 2012 SOUTH      2      TOTAL S2TOTAL 3178    1.4  2.50  2.15 18.0
## 23     1 2012 SOUTH      2      TOTAL S2TOTAL  101    1.9  3.31  2.65 15.0
## 24     1 2012 SOUTH      2      TOTAL S2TOTAL  102    1.75  2.70  2.55 16.0
## 25     1 2012 SOUTH      2      TOTAL S2TOTAL  103    1.8  2.75  2.30 16.0
## 26     1 2012 SOUTH      2      TOTAL S2TOTAL  104    2.7  4.05  4.00 35.2
## 27     1 2012 SOUTH      2      TOTAL S2TOTAL  105    2.02  2.85  1.49 17.0

```

## 28	1	2012	SOUTH	2	TOTAL S2TOTAL	108	1.9	3.10	2.85	19.0
## 29	1	2012	SOUTH	2	TOTAL S2TOTAL	109	1.85	2.45	1.90	19.0
## 30	1	2012	SOUTH	2	TOTAL S2TOTAL	110	1.65	1.90	1.54	17.0
## 31	1	2012	SOUTH	2	TOTAL S2TOTAL	111	1.4	2.35	1.45	14.0
## 32	1	2012	SOUTH	2	TOTAL S2TOTAL	113	2.5	3.25	2.30	22.0
## 33	1	2012	SOUTH	2	TOTAL S2TOTAL	115	2.05	5.40	4.50	33.0
## 34	1	2012	SOUTH	2	TOTAL S2TOTAL	116	2.26	3.50	3.10	33.0
## 35	1	2012	SOUTH	2	TOTAL S2TOTAL	117	2.13	2.40	2.30	20.0
## 36	1	2012	SOUTH	2	TOTAL S2TOTAL	118	1.8	3.15	2.55	22.0
## 37	1	2012	SOUTH	2	TOTAL S2TOTAL	1211	1.85	2.00	2.27	20.0
## 38	1	2012	SOUTH	2	TOTAL S2TOTAL	1212	1.5	2.15	1.80	15.0
## 39	1	2012	SOUTH	2	TOTAL S2TOTAL	1213	1.87	2.34	2.05	13.0
## 40	1	2012	SOUTH	2	TOTAL S2TOTAL	1214	1.58	1.28	0.75	11.0
## 41	1	2012	SOUTH	2	TOTAL S2TOTAL	1215	2.05	2.10	1.75	17.0
## 42	1	2012	SOUTH	2	TOTAL S2TOTAL	1216	1.75	2.45	3.28	16.0
## 43	1	2012	SOUTH	2	TOTAL S2TOTAL	1217	1.49	1.50	1.45	13.0
## 44	1	2012	SOUTH	2	TOTAL S2TOTAL	1218	1.28	2.00	0.90	10.0
## 45	1	2012	SOUTH	2	TOTAL S2TOTAL	1219	1.49	2.35	1.65	13.0
## 46	1	2012	SOUTH	2	TOTAL S2TOTAL	1220	1.07	1.20	0.95	11.0
## 47	1	2012	SOUTH	2	TOTAL S2TOTAL	1231	1.48	1.25	1.20	9.0
## 48	1	2012	SOUTH	2	TOTAL S2TOTAL	1232	1.25	1.25	0.90	10.0
## 49	1	2012	SOUTH	2	TOTAL S2TOTAL	1233	1.41	1.41	1.40	14.0
## 50	1	2012	SOUTH	2	TOTAL S2TOTAL	1234	1.6	1.60	1.30	13.0
## 51	1	2012	SOUTH	2	TOTAL S2TOTAL	1235	1.2	1.20	1.30	14.0
## 52	1	2012	SOUTH	2	TOTAL S2TOTAL	1236	1.49	1.49	1.20	8.0
## 53	1	2012	SOUTH	2	TOTAL S2TOTAL	1237	1.5	1.50	1.50	14.0
## 54	1	2012	SOUTH	2	TOTAL S2TOTAL	1238	1.65	1.65	2.00	20.0
## 55	1	2012	SOUTH	2	TOTAL S2TOTAL	1239	1.13	1.13	1.20	10.0
## 56	1	2012	SOUTH	2	TOTAL S2TOTAL	1240	1.25	1.25	0.90	10.0
## 57	1	2012	SOUTH	2	TOTAL S2TOTAL	1251	1.1	1.20	1.10	10.0
## 58	1	2012	SOUTH	2	TOTAL S2TOTAL	1252	2.2	2.70	2.40	25.0
## 59	1	2012	SOUTH	2	TOTAL S2TOTAL	1253	1.45	1.65	1.25	10.0
## 60	1	2012	SOUTH	2	TOTAL S2TOTAL	1254	1.6	2.45	2.10	13.0
## 61	1	2012	SOUTH	2	TOTAL S2TOTAL	1255	1.55	2.40	1.80	13.0
## 62	1	2012	SOUTH	2	TOTAL S2TOTAL	1256	1.5	2.40	2.15	13.0
## 63	1	2012	SOUTH	2	TOTAL S2TOTAL	1257	1.03	1.20	1.00	10.0
## 64	1	2012	SOUTH	2	TOTAL S2TOTAL	1258	2.14	1.90	1.70	13.0
## 65	1	2012	SOUTH	2	TOTAL S2TOTAL	1259	1.2	1.90	1.65	12.0
## 66	1	2012	SOUTH	2	TOTAL S2TOTAL	1260	1.05	1.10	1.00	9.0
## 67	1	2012	SOUTH	2	TOTAL S2TOTAL	2131	1.8	2.60	2.40	15.0
## 68	1	2012	SOUTH	2	TOTAL S2TOTAL	2132	1.2	1.00	0.95	7.0
## 69	1	2012	SOUTH	2	TOTAL S2TOTAL	2133	1.75	1.40	1.10	10.0
## 70	1	2012	SOUTH	2	TOTAL S2TOTAL	2134	1.45	3.10	1.80	10.0
## 71	1	2012	SOUTH	2	TOTAL S2TOTAL	2135	1.17	1.20	1.10	5.0
## 72	1	2012	SOUTH	2	TOTAL S2TOTAL	2136	2.15	3.10	2.58	22.0
## 73	1	2012	SOUTH	2	TOTAL S2TOTAL	2137	1.7	1.70	1.40	12.0
## 74	1	2012	SOUTH	2	TOTAL S2TOTAL	3132	1.98	2.85	2.70	12.0
## 75	1	2012	SOUTH	2	TOTAL S2TOTAL	3133	1.26	1.95	1.75	17.0
## 76	1	2012	SOUTH	2	TOTAL S2TOTAL	3134	1.11	1.95	1.50	10.0
## 77	1	2012	SOUTH	2	TOTAL S2TOTAL	3135	1.14	1.32	1.05	10.0
## 78	1	2012	SOUTH	2	TOTAL S2TOTAL	3136	1.26	1.60	1.40	10.0
## 79	1	2012	SOUTH	2	TOTAL S2TOTAL	3137	1.3	1.40	0.80	10.0
## 80	1	2012	SOUTH	2	TOTAL S2TOTAL	3138	1.29	1.44	1.35	13.0
## 81	1	2012	SOUTH	2	TOTAL S2TOTAL	3139	1.31	1.35	1.15	7.0

## 82	1	2012	SOUTH	2	TOTAL S2TOTAL	3140	1.15	1.70	1.28	10.0
## 83	1	2012	SOUTH	2	TOTAL S2TOTAL	3151	1.87	3.40	1.85	15.0
## 84	1	2012	SOUTH	2	TOTAL S2TOTAL	3152	1.47	2.10	1.61	8.0
## 85	1	2012	SOUTH	2	TOTAL S2TOTAL	3153	1.05	1.79	1.50	10.0
## 86	1	2012	SOUTH	2	TOTAL S2TOTAL	3154	2.1	4.90	3.75	25.0
## 87	1	2012	SOUTH	2	TOTAL S2TOTAL	3155	1.99	1.80	1.35	13.0
## 88	1	2012	SOUTH	2	TOTAL S2TOTAL	3156	1.42	1.90	1.80	14.0
## 89	1	2012	SOUTH	2	TOTAL S2TOTAL	3157	1.5	2.11	1.75	12.0
## 90	1	2012	SOUTH	2	TOTAL S2TOTAL	3158	1.06	1.05	0.85	4.0
## 91	1	2012	SOUTH	2	TOTAL S2TOTAL	3159	1.49	1.50	1.15	13.0
## 92	1	2012	SOUTH	2	TOTAL S2TOTAL	3160	1.8	1.60	1.50	14.0
## 93	1	2012	SOUTH	2	TOTAL S2TOTAL	3171	1.93	1.74	1.20	14.0
## 94	1	2012	SOUTH	2	TOTAL S2TOTAL	3172	1.2	1.60	1.30	10.0
## 95	1	2012	SOUTH	2	TOTAL S2TOTAL	3173	1.65	1.25	1.10	11.0
## 96	1	2012	SOUTH	2	TOTAL S2TOTAL	3174	1.52	1.49	1.10	12.0
## 97	1	2012	SOUTH	2	TOTAL S2TOTAL	3175	1.43	2.05	1.54	13.0
## 98	1	2012	SOUTH	2	TOTAL S2TOTAL	3176	1.25	1.40	1.25	13.0
## 99	1	2012	SOUTH	2	TOTAL S2TOTAL	3177	1.88	2.65	2.64	20.0
## 100	1	2012	SOUTH	2	TOTAL S2TOTAL	3179	1.03	1.40	0.60	13.0
## 101	1	2012	SOUTH	2	TOTAL S2TOTAL	3180	1.1	1.30	1.20	10.0
## 102	1	2012	SOUTH	2	TOTAL S2TOTAL	3191	1.4	1.05	1.00	10.0
## 103	1	2012	SOUTH	2	TOTAL S2TOTAL	3192	1.05	1.55	0.90	10.0
## 104	1	2012	SOUTH	2	TOTAL S2TOTAL	3193	1.18	1.20	1.00	7.0
## 105	1	2012	SOUTH	2	TOTAL S2TOTAL	3194	1.4	1.30	1.85	13.0
## 106	1	2012	SOUTH	2	TOTAL S2TOTAL	3195	1.37	2.67	2.19	19.0
## 107	1	2012	SOUTH	2	TOTAL S2TOTAL	3196	1.32	2.15	1.55	11.0
## 108	1	2012	SOUTH	2	MEGA S2MEGA	182	1.55	2.20	1.20	20.0
## 109	1	2012	SOUTH	2	MEGA S2MEGA	183	1.3	1.80	0.90	8.0
## 110	1	2012	SOUTH	2	MEGA S2MEGA	184	1.24	1.20	1.20	25.0
## 111	1	2012	SOUTH	2	MEGA S2MEGA	185	1.5	2.10	1.75	16.0
## 112	1	2012	SOUTH	2	MEGA S2MEGA	186	1.65	2.50	2.20	15.0
## 113	1	2012	SOUTH	2	MEGA S2MEGA	187	2.17	2.00	1.20	15.0
## 114	1	2012	SOUTH	2	MEGA S2MEGA	188	1.28	1.60	1.50	10.0
## 115	1	2012	SOUTH	2	MEGA S2MEGA	189	1.07	1.50	1.50	10.0
## 116	1	2012	SOUTH	2	MEGA S2MEGA	190	0.67	1.00	0.80	8.0
## 117	1	2012	SOUTH	2	MEGA S2MEGA	191	0.68	0.70	0.60	4.0
## 118	1	2012	SOUTH	2	MEGA S2MEGA	192	1.87	1.60	1.40	9.0
## 119	1	2012	SOUTH	2	MEGA S2MEGA	193	1.35	1.90	1.50	14.0
## 120	1	2012	SOUTH	2	MEGA S2MEGA	194	1.75	2.10	2.10	15.0
## 121	1	2012	SOUTH	2	MESO S2MESO	462	1.75	3.30	2.50	23.0
## 122	1	2012	SOUTH	2	MESO S2MESO	463	1.64	2.30	2.00	14.0
## 123	1	2012	SOUTH	2	MESO S2MESO	2138	1.42	0.90	0.80	10.0
## 124	1	2012	SOUTH	3	OPEN S3OPEN	1301	dead	NA	NA	NA
## 125	1	2012	SOUTH	3	OPEN S3OPEN	1302	0.9	1.30	1.10	11.0
## 126	1	2012	SOUTH	3	TOTAL S3TOTAL	1061	dead	NA	NA	NA
## 127	1	2012	SOUTH	3	TOTAL S3TOTAL	1062	1.8	2.60	2.60	15.0
## 128	1	2012	SOUTH	3	TOTAL S3TOTAL	1063	2.47	3.10	2.20	18.0
## 129	1	2012	SOUTH	3	TOTAL S3TOTAL	1064	2.15	1.60	1.10	17.0
## 130	1	2012	SOUTH	3	TOTAL S3TOTAL	1066	1.7	2.50	2.15	15.0
## 131	1	2012	SOUTH	3	TOTAL S3TOTAL	1066	1.9	1.80	1.50	20.0
## 132	1	2012	SOUTH	3	TOTAL S3TOTAL	1067	1.95	2.10	1.90	13.0
## 133	1	2012	SOUTH	3	TOTAL S3TOTAL	1068	1.8	1.70	1.40	13.0
## 134	1	2012	SOUTH	3	TOTAL S3TOTAL	1069	1.4	2.00	1.60	14.0
## 135	1	2012	SOUTH	3	TOTAL S3TOTAL	1070	1	1.30	1.20	7.0

## 136	1	2012	SOUTH	3	TOTAL	S3TOTAL	2139	1.75	1.20	1.10	13.0
## 137	1	2012	SOUTH	3	TOTAL	S3TOTAL	2140	1.28	1.50	0.95	4.0
## 138	1	2012	SOUTH	3	TOTAL	S3TOTAL	2151	1	1.40	1.20	4.0
## 139	1	2012	SOUTH	3	TOTAL	S3TOTAL	2152	1.45	1.50	1.30	10.0
## 140	1	2012	SOUTH	3	TOTAL	S3TOTAL	2153	1	1.00	0.75	8.0
## 141	1	2012	SOUTH	3	TOTAL	S3TOTAL	2154	1.03	1.00	0.90	6.0
## 142	1	2012	SOUTH	3	TOTAL	S3TOTAL	2155	1.51	2.00	1.80	12.0
## 143	1	2012	SOUTH	3	TOTAL	S3TOTAL	2156	1.17	1.10	0.90	10.0
## 144	1	2012	SOUTH	3	TOTAL	S3TOTAL	2157	1.33	1.90	1.85	14.0
## 145	1	2012	SOUTH	3	TOTAL	S3TOTAL	2158	1.3	1.10	0.85	8.0
## 146	1	2012	SOUTH	3	TOTAL	S3TOTAL	2159	1.13	1.10	0.90	10.0
## 147	1	2012	SOUTH	3	TOTAL	S3TOTAL	2160	1.58	1.40	1.40	13.0
## 148	1	2012	SOUTH	3	TOTAL	S3TOTAL	2171	1.06	1.40	1.00	5.0
## 149	1	2012	SOUTH	3	TOTAL	S3TOTAL	2172	1.05	1.40	0.95	7.0
## 150	1	2012	SOUTH	3	TOTAL	S3TOTAL	2173	1.45	1.60	1.10	6.0
## 151	1	2012	SOUTH	3	TOTAL	S3TOTAL	2174	1.15	1.10	0.90	5.0
## 152	1	2012	SOUTH	3	TOTAL	S3TOTAL	2175	1.42	1.45	1.30	13.0
## 153	1	2012	SOUTH	3	TOTAL	S3TOTAL	2176	1.02	1.20	1.00	8.0
## 154	1	2012	SOUTH	3	TOTAL	S3TOTAL	2177	1.4	1.20	1.00	9.0
## 155	1	2012	SOUTH	3	TOTAL	S3TOTAL	2178	1.45	2.10	2.05	15.0
## 156	1	2012	SOUTH	3	MESO	S3MESO	1421	1.95	2.20	1.60	13.0
## 157	1	2012	SOUTH	3	MESO	S3MESO	1422	dead	NA	NA	NA
##	FLOWERS	BUDS	FRUITS	ANT							
## 1	0	0	10	CS							
## 2	0	0	150	TP							
## 3	2	1	50	TP							
## 4	0	0	75	CS							
## 5	0	0	20	CS							
## 6	0	0	0	E							
## 7	0	0	0	CS							
## 8	0	0	25	CS							
## 9	0	0	0	TP							
## 10	0	0	50	TP							
## 11	0	0	5	CS							
## 12	0	0	60	TP							
## 13	0	0	60	TP							
## 14	2	0	60	CS							
## 15	2	0	0	CS							
## 16	0	0	0	TP							
## 17	0	0	0	TP							
## 18	0	0	0	CS							
## 19	0	0	0	CM							
## 20	0	0	0	TP							
## 21	NA	NA	NA								
## 22	0	0	5	CS							
## 23	0	0	45	CS							
## 24	40	50	35	CS							
## 25	8	2	65	CS							
## 26	0	0	20	TP							
## 27	0	0	70	CS							
## 28	0	0	125	CM							
## 29	0	0	200	CM							
## 30	0	0	10	CS							
## 31	0	0	0	CS							

## 32	0	0	35	TP
## 33	0	0	300	CM
## 34	2	2	100	CS
## 35	0	0	30	CM
## 36	0	0	50	TP
## 37	0	0	10	CM
## 38	0	0	25	CS
## 39	0	0	15	TP
## 40	0	0	0	TP
## 41	0	0	15	TP
## 42	0	0	0	TP
## 43	0	0	40	TP
## 44	0	0	0	TP
## 45	0	0	15	CM
## 46	0	0	0	CM
## 47	0	0	0	TP
## 48	0	0	0	TP
## 49	0	0	1	TP
## 50	0	0	20	TP
## 51	0	0	0	TP
## 52	0	0	0	TP
## 53	0	0	20	TP
## 54	0	0	0	TP
## 55	0	0	0	CN
## 56	0	0	0	CN
## 57	0	0	0	TP
## 58	0	0	5	TP
## 59	0	0	0	TP
## 60	0	0	25	TP
## 61	0	0	25	TP
## 62	0	0	20	TP
## 63	0	0	0	TP
## 64	0	0	10	CS
## 65	1	0	25	CS
## 66	0	0	0	TP
## 67	0	0	10	TP
## 68	0	0	0	TP
## 69	0	0	0	TP
## 70	0	0	0	TP
## 71	0	0	0	TP
## 72	0	0	0	CS
## 73	0	0	0	CS
## 74	0	0	25	AB_TP
## 75	0	0	0	TP
## 76	0	0	0	TP
## 77	0	0	0	TP
## 78	0	0	0	CS
## 79	0	0	0	CS
## 80	0	0	0	CS
## 81	0	0	0	CS
## 82	0	0	5	CS
## 83	6	0	0	CS
## 84	0	0	0	CS
## 85	0	0	1	CS

## 86	0	0	25	CS
## 87	0	0	0	CS
## 88	0	0	0	CS
## 89	0	0	10	CS
## 90	0	0	0	CS
## 91	0	0	35	CS
## 92	0	0	0	CS
## 93	0	0	0	CS
## 94	0	0	0	CS
## 95	0	0	0	CS
## 96	0	0	20	CS
## 97	0	0	0	CS
## 98	0	0	0	CM
## 99	0	0	100	CM
## 100	0	0	0	CS
## 101	0	0	0	CS
## 102	0	0	0	CS
## 103	0	0	0	CM
## 104	0	0	0	TP
## 105	0	0	30	CS
## 106	0	0	50	TP
## 107	0	0	10	CS
## 108	0	0	0	CS
## 109	0	0	15	CS
## 110	0	0	10	CS
## 111	5	0	200	CS
## 112	0	0	80	CS
## 113	0	0	150	TP
## 114	0	0	40	TP
## 115	0	0	60	TP
## 116	0	0	0	CS
## 117	0	0	0	TP
## 118	0	0	40	CS
## 119	0	0	20	CS
## 120	0	0	75	TP
## 121	0	0	20	CM
## 122	0	0	0	TP
## 123	0	0	0	E
## 124	NA	NA	NA	
## 125	0	0	0	TP
## 126	NA	NA	NA	
## 127	0	0	50	TP
## 128	0	0	0	TP
## 129	0	0	0	TP
## 130	0	0	2	TP
## 131	0	0	25	TP
## 132	0	0	0	TP
## 133	0	0	0	TP
## 134	0	0	0	TP
## 135	0	0	0	TP
## 136	0	0	0	TP
## 137	0	0	0	TP
## 138	0	0	0	TP
## 139	0	0	0	TP

```
## 140      0      0      0      TP
## 141      0      0      0      TP
## 142      0      0      0      TP
## 143      0      0      0      TP
## 144      0      0      0      TP
## 145      0      0      0      TP
## 146      0      0      0      TP
## 147      0      0      0      TP
## 148      0      0      8      TP
## 149      0      0      0      TP
## 150      0      0      0      TP
## 151      0      0      0      TP
## 152      0      0      0      TP
## 153      0      0      0      TP
## 154      0      0      0      TP
## 155      0      0     20      TP
## 156      0      0      2      CS
## 157     NA     NA     NA
```

```
r_proj_wd <- "/Users/atziri/Bio 195-197/Data Science"
r_chunk_wd <- getwd()
r_proj_wd == r_chunk_wd
```

```
## [1] FALSE
```

```
r_chunk_wd
```

```
## [1] "/Users/atziri/Bio 195-197/Data Science/documents"
```

```
acacia <- read.csv(file = "../raw-data/ACACIA_DREPANOLOBIUM_SURVEY.txt", sep = "\t")
```

#The two dots represents what we need to do to go one up to the folder nedded

3. explore our data set

'head()' gives us the first six rows

```
head(acacia)
```

```
##  SURVEY YEAR  SITE BLOCK TREATMENT  PLOT  ID HEIGHT AXIS1 AXIS2 CIRC
## 1      1 2012 SOUTH      1      TOTAL S1TOTAL 581   2.25  2.75  2.15  20
## 2      1 2012 SOUTH      1      TOTAL S1TOTAL 582   2.65  4.10  3.90  28
## 3      1 2012 SOUTH      1      TOTAL S1TOTAL 3111   1.5   1.70  0.85  17
## 4      1 2012 SOUTH      1      TOTAL S1TOTAL 3112   2.01  1.80  1.60  12
## 5      1 2012 SOUTH      1      TOTAL S1TOTAL 3113   1.75  1.84  1.42  13
## 6      1 2012 SOUTH      1      TOTAL S1TOTAL 3114   1.65  1.62  0.85  15
##  FLOWERS BUDS FRUITS ANT
## 1      0      0      10 CS
## 2      0      0     150 TP
## 3      2      1      50 TP
```

```
## 4      0      0      75 CS
## 5      0      0      20 CS
## 6      0      0       0 E
```

```
summary(acacia)
```

```
##      SURVEY      YEAR      SITE      BLOCK
## Min.   :1   Min.   :2012   Length:157   Min.   :1.000
## 1st Qu.:1   1st Qu.:2012   Class :character 1st Qu.:2.000
## Median :1   Median :2012   Mode  :character Median :2.000
## Mean   :1   Mean   :2012                   Mean  :2.089
## 3rd Qu.:1   3rd Qu.:2012                   3rd Qu.:2.000
## Max.   :1   Max.   :2012                   Max.   :3.000
##
##      TREATMENT      PLOT      ID      HEIGHT
## Length:157      Length:157   Min.   : 101   Length:157
## Class :character Class :character 1st Qu.:1062   Class :character
## Mode  :character Mode  :character Median :1301   Mode  :character
##                                     Mean  :1743
##                                     3rd Qu.:3118
##                                     Max.   :3199
##
##      AXIS1      AXIS2      CIRC      FLOWERS
## Min.   :0.700   Min.   :0.550   Min.   : 4.00   Min.   : 0.0000
## 1st Qu.:1.400   1st Qu.:1.100   1st Qu.:10.00   1st Qu.: 0.0000
## Median :1.800   Median :1.490   Median :13.00   Median : 0.0000
## Mean   :1.972   Mean   :1.636   Mean   :13.76   Mean   : 0.4444
## 3rd Qu.:2.350   3rd Qu.:2.000   3rd Qu.:16.00   3rd Qu.: 0.0000
## Max.   :5.550   Max.   :4.820   Max.   :35.20   Max.   :40.0000
## NA's   :4      NA's   :4      NA's   :4      NA's   :4
##
##      BUDS      FRUITS      ANT
## Min.   : 0.0000   Min.   : 0.00   Length:157
## 1st Qu.: 0.0000   1st Qu.: 0.00   Class :character
## Median : 0.0000   Median : 0.00   Mode  :character
## Mean   : 0.3595   Mean   : 20.03
## 3rd Qu.: 0.0000   3rd Qu.: 25.00
## Max.   :50.0000   Max.   :300.00
## NA's   :4      NA's   :4
```

```
colnames(acacia)
```

```
## [1] "SURVEY" "YEAR" "SITE" "BLOCK" "TREATMENT" "PLOT"
## [7] "ID" "HEIGHT" "AXIS1" "AXIS2" "CIRC" "FLOWERS"
## [13] "BUDS" "FRUITS" "ANT"
```

```
row.names(acacia)
```

```
## [1] "1" "2" "3" "4" "5" "6" "7" "8" "9" "10" "11" "12"
## [13] "13" "14" "15" "16" "17" "18" "19" "20" "21" "22" "23" "24"
## [25] "25" "26" "27" "28" "29" "30" "31" "32" "33" "34" "35" "36"
## [37] "37" "38" "39" "40" "41" "42" "43" "44" "45" "46" "47" "48"
## [49] "49" "50" "51" "52" "53" "54" "55" "56" "57" "58" "59" "60"
```

```
## [61] "61" "62" "63" "64" "65" "66" "67" "68" "69" "70" "71" "72"
## [73] "73" "74" "75" "76" "77" "78" "79" "80" "81" "82" "83" "84"
## [85] "85" "86" "87" "88" "89" "90" "91" "92" "93" "94" "95" "96"
## [97] "97" "98" "99" "100" "101" "102" "103" "104" "105" "106" "107" "108"
## [109] "109" "110" "111" "112" "113" "114" "115" "116" "117" "118" "119" "120"
## [121] "121" "122" "123" "124" "125" "126" "127" "128" "129" "130" "131" "132"
## [133] "133" "134" "135" "136" "137" "138" "139" "140" "141" "142" "143" "144"
## [145] "145" "146" "147" "148" "149" "150" "151" "152" "153" "154" "155" "156"
## [157] "157"
```

```
class(acacia$SURVEY)
```

```
## [1] "integer"
```

```
#the sapply function all owes to apply a function to a list of objects
#a data frame is a list of vectors of the same length
sapply(acacia, class)
```

```
##      SURVEY      YEAR      SITE      BLOCK  TREATMENT      PLOT
## "integer" "integer" "character" "integer" "character" "character"
##      ID      HEIGHT      AXIS1      AXIS2      CIRC      FLOWERS
## "integer" "character" "numeric" "numeric" "numeric" "integer"
##      BUDS      FRUITS      ANT
## "integer" "integer" "character"
```

make sure that everything that should be a numeric value is a number one way to check is the 'summary()' command

another way is using the type function

```
typeof(acacia[, "HEIGHT"])
```

```
## [1] "character"
```

```
acacia$HEIGHT
```

```
## [1] "2.25" "2.65" "1.5" "2.01" "1.75" "1.65" "1.2" "1.45" "1.87" "2.38"
## [11] "2.58" "2.65" "2.35" "1.88" "2.32" "2.39" "2.2" "1.05" "2" "1.28"
## [21] "dead" "1.4" "1.9" "1.75" "1.8" "2.7" "2.02" "1.9" "1.85" "1.65"
## [31] "1.4" "2.5" "2.05" "2.26" "2.13" "1.8" "1.85" "1.5" "1.87" "1.58"
## [41] "2.05" "1.75" "1.49" "1.28" "1.49" "1.07" "1.48" "1.25" "1.41" "1.6"
## [51] "1.2" "1.49" "1.5" "1.65" "1.13" "1.25" "1.1" "2.2" "1.45" "1.6"
## [61] "1.55" "1.5" "1.03" "2.14" "1.2" "1.05" "1.8" "1.2" "1.75" "1.45"
## [71] "1.17" "2.15" "1.7" "1.98" "1.26" "1.11" "1.14" "1.26" "1.3" "1.29"
## [81] "1.31" "1.15" "1.87" "1.47" "1.05" "2.1" "1.99" "1.42" "1.5" "1.06"
## [91] "1.49" "1.8" "1.93" "1.2" "1.65" "1.52" "1.43" "1.25" "1.88" "1.03"
## [101] "1.1" "1.4" "1.05" "1.18" "1.4" "1.37" "1.32" "1.55" "1.3" "1.24"
## [111] "1.5" "1.65" "2.17" "1.28" "1.07" "0.67" "0.68" "1.87" "1.35" "1.75"
## [121] "1.75" "1.64" "1.42" "dead" "0.9" "dead" "1.8" "2.47" "2.15" "1.7"
## [131] "1.9" "1.95" "1.8" "1.4" "1" "1.75" "1.28" "1" "1.45" "1"
## [141] "1.03" "1.51" "1.17" "1.33" "1.3" "1.13" "1.58" "1.06" "1.05" "1.45"
## [151] "1.15" "1.42" "1.02" "1.4" "1.45" "1.95" "dead"
```

we identified a column that has problematic data we need to fix this

Cleaning our raw data

3.2 assign 'NA' lable to missing

We are going to read the data table again, but we are going to assign 'NA' to the "dead value" that we don't want in our arguments are always plain text

```
acacia <- read.csv(file = "/Users/atziri/Bio 195-197/Data Science/raw-data/ACACIA_DREPANOLOBIUM_SURVEY.")
```

4. Visualize our data

For this we are using the 'ggplot' package. let's install and load it

```
# install.packages("ggplot2")
library(ggplot2)
```

Now We are going to create our first plotting layer with the function 'ggplot'.

```
colnames(acacia)
```

```
## [1] "SURVEY" "YEAR" "SITE" "BLOCK" "TREATMENT" "PLOT"
## [7] "ID" "HEIGHT" "AXIS1" "AXIS2" "CIRC" "FLOWERS"
## [13] "BUDS" "FRUITS" "ANT"
```

```
acacia$CIRC
```

```
## [1] 20.0 28.0 17.0 12.0 13.0 15.0 9.0 12.2 13.0 35.0 24.0 27.0 20.0 28.0 30.0
## [16] 13.0 10.0 8.0 10.0 10.0 NA 18.0 15.0 16.0 16.0 35.2 17.0 19.0 19.0 17.0
## [31] 14.0 22.0 33.0 33.0 20.0 22.0 20.0 15.0 13.0 11.0 17.0 16.0 13.0 10.0 13.0
## [46] 11.0 9.0 10.0 14.0 13.0 14.0 8.0 14.0 20.0 10.0 10.0 10.0 25.0 10.0 13.0
## [61] 13.0 13.0 10.0 13.0 12.0 9.0 15.0 7.0 10.0 10.0 5.0 22.0 12.0 12.0 17.0
## [76] 10.0 10.0 10.0 10.0 13.0 7.0 10.0 15.0 8.0 10.0 25.0 13.0 14.0 12.0 4.0
## [91] 13.0 14.0 14.0 10.0 11.0 12.0 13.0 13.0 20.0 13.0 10.0 10.0 10.0 7.0 13.0
## [106] 19.0 11.0 20.0 8.0 25.0 16.0 15.0 15.0 10.0 10.0 8.0 4.0 9.0 14.0 15.0
## [121] 23.0 14.0 10.0 NA 11.0 NA 15.0 18.0 17.0 15.0 20.0 13.0 13.0 14.0 7.0
## [136] 13.0 4.0 4.0 10.0 8.0 6.0 12.0 10.0 14.0 8.0 10.0 13.0 5.0 7.0 6.0
## [151] 5.0 13.0 8.0 9.0 15.0 13.0 NA
```

The ggplot function creates a blank canvas. the canvas contains our data and the variables that we plot.

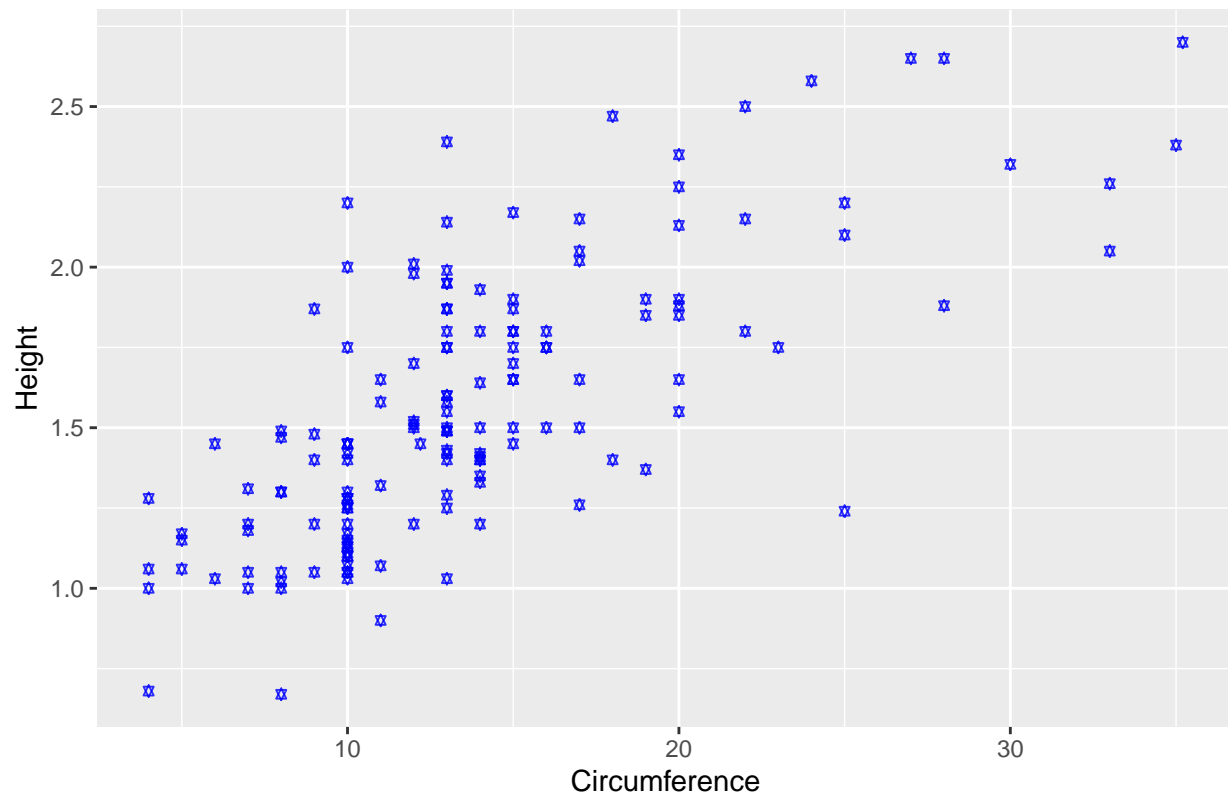
The aes function allows for the plugging of all lines colors and data we want to read

A scatter plot can be created with the function (geom_point) it can have a blank () however if you want to specify color size and transparency you add it

```
ggplot(data = acacia, mapping = aes(x = CIRC, y = HEIGHT )) + geom_point(size = 1, color = "blue", alpha = 0.5)
labs(x = "Circumference", y = "Height", title = "Data From UHURU Acacia Survey ")
```

```
## Warning: Removed 4 rows containing missing values (geom_point).
```

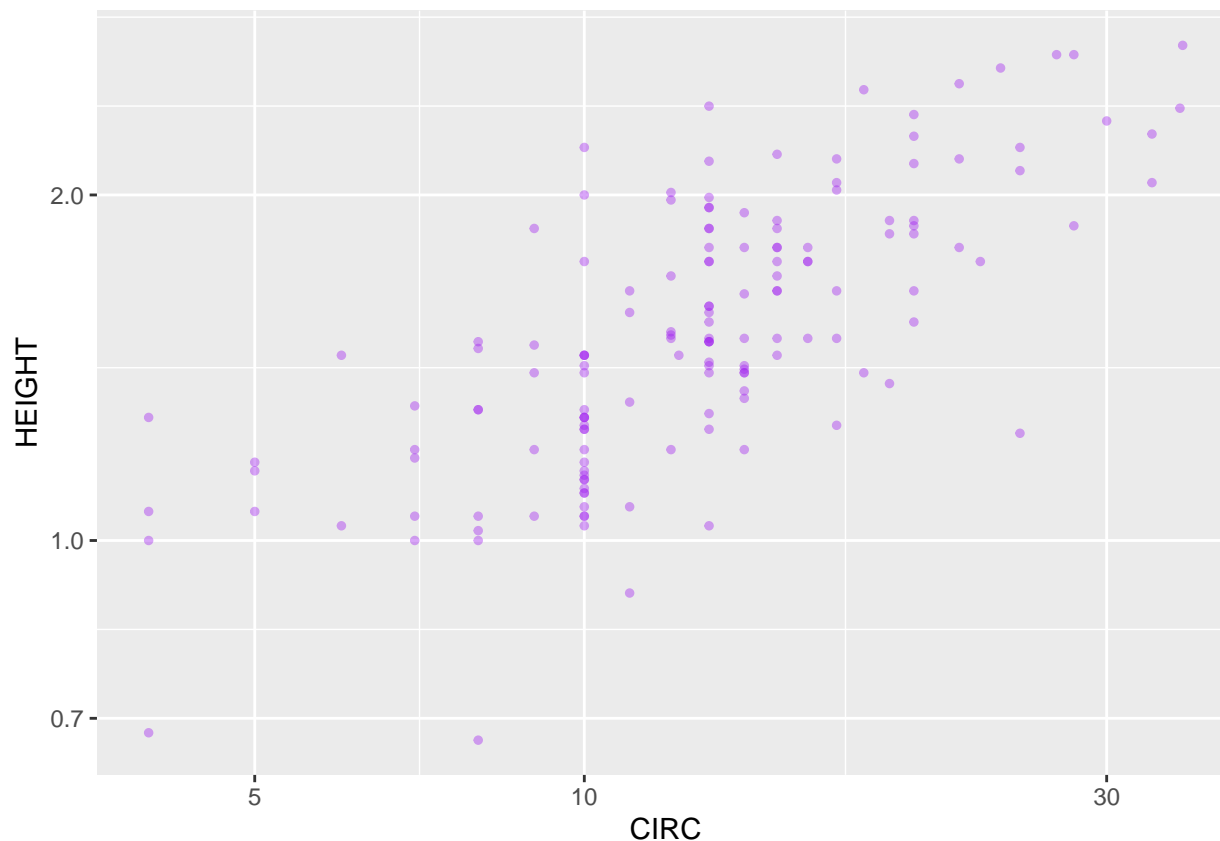

Data From UHURU Acacia Survey



to rescale the plotting of the axis to log scale we use the function 'scale_y_log10()'

```
ggplot(data = acacia, mapping = aes(x = CIRC, y = HEIGHT )) +  
  geom_point(size = 1, color = "purple", alpha = 0.4) +  
  scale_x_log10() +  
  scale_y_log10()
```

```
## Warning: Removed 4 rows containing missing values (geom_point).
```

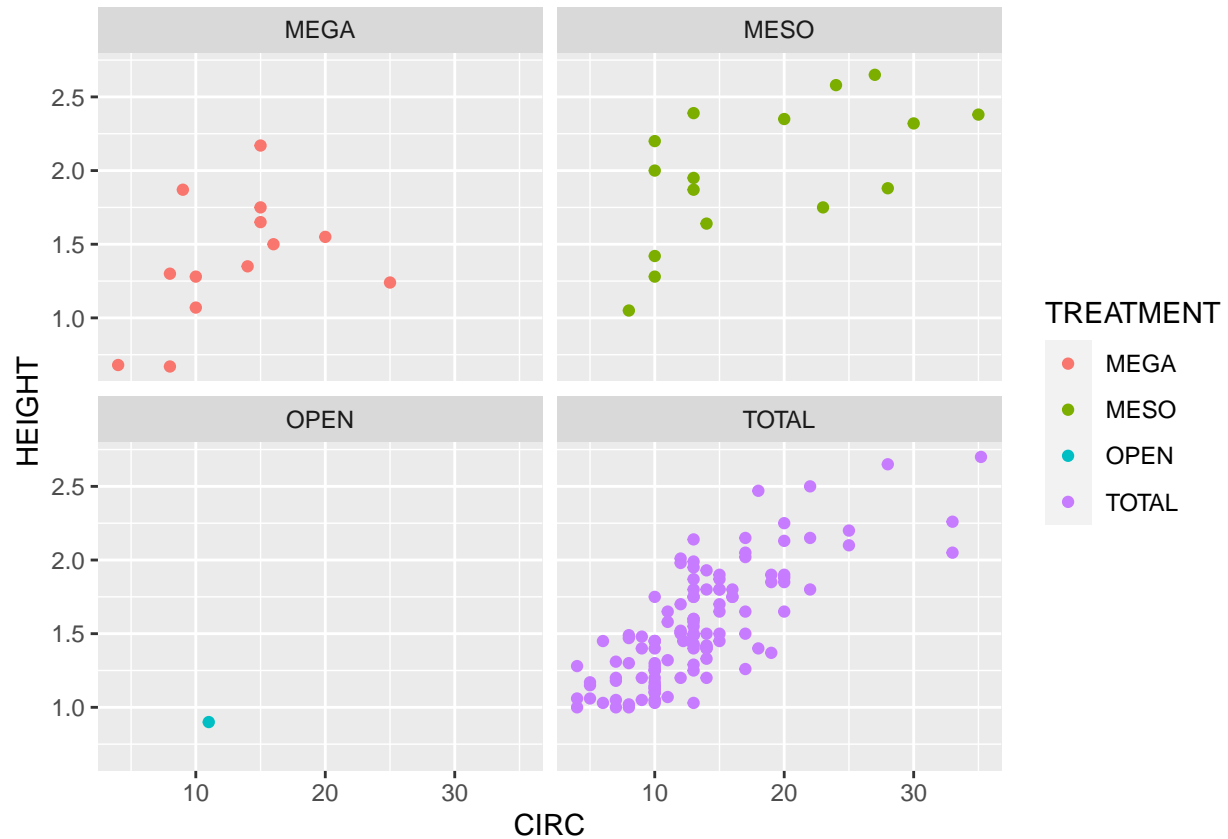


```
acacia$TREATMENT
```

```
## [1] "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "MESO"
## [10] "MESO" "MESO" "MESO" "MESO" "MESO" "MESO" "MESO" "MESO" "MESO"
## [19] "MESO" "MESO" "OPEN" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL"
## [28] "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL"
## [37] "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL"
## [46] "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL"
## [55] "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL"
## [64] "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL"
## [73] "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL"
## [82] "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL"
## [91] "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL"
## [100] "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL"
## [109] "MEGA" "MEGA" "MEGA" "MEGA" "MEGA" "MEGA" "MEGA" "MEGA" "MEGA"
## [118] "MEGA" "MEGA" "MEGA" "MESO" "MESO" "MESO" "OPEN" "OPEN" "TOTAL"
## [127] "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL"
## [136] "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL"
## [145] "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL" "TOTAL"
## [154] "TOTAL" "TOTAL" "MESO" "MESO"
```

```
ggplot(data = acacia, mapping = aes (x = CIRC, y = HEIGHT, color = TREATMENT)) +
  geom_point() +
  facet_wrap(~TREATMENT)
```

```
## Warning: Removed 4 rows containing missing values (geom_point).
```



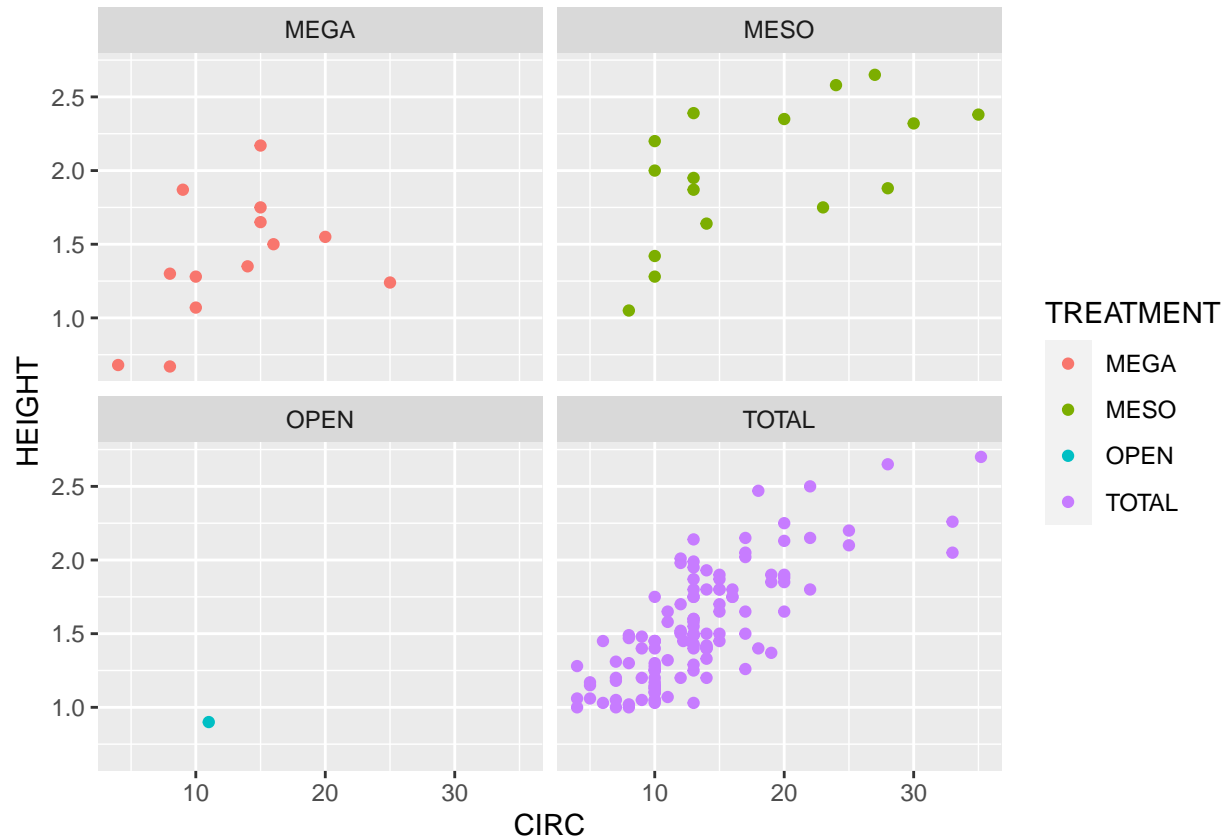
4.2 Visualize a statistical analysis of correlation

Subplots or facets

The function to create subplots by a third variable is called 'facet_wrap()'

```
ggplot(data = acacia, mapping = aes(x = CIRC, y = HEIGHT, color = TREATMENT)) + geom_point() +
  facet_wrap(~TREATMENT)
```

```
## Warning: Removed 4 rows containing missing values (geom_point).
```



The total that excludes all the herbivores has more trees than the fenced spaces with select herbivores meso has taller trees with larger circumference mega has shorter trees with smaller circumference

A little interpretation of what is going on here

How to test a Hypothesis

Model Fitting functions

The `geom_smooth()` function

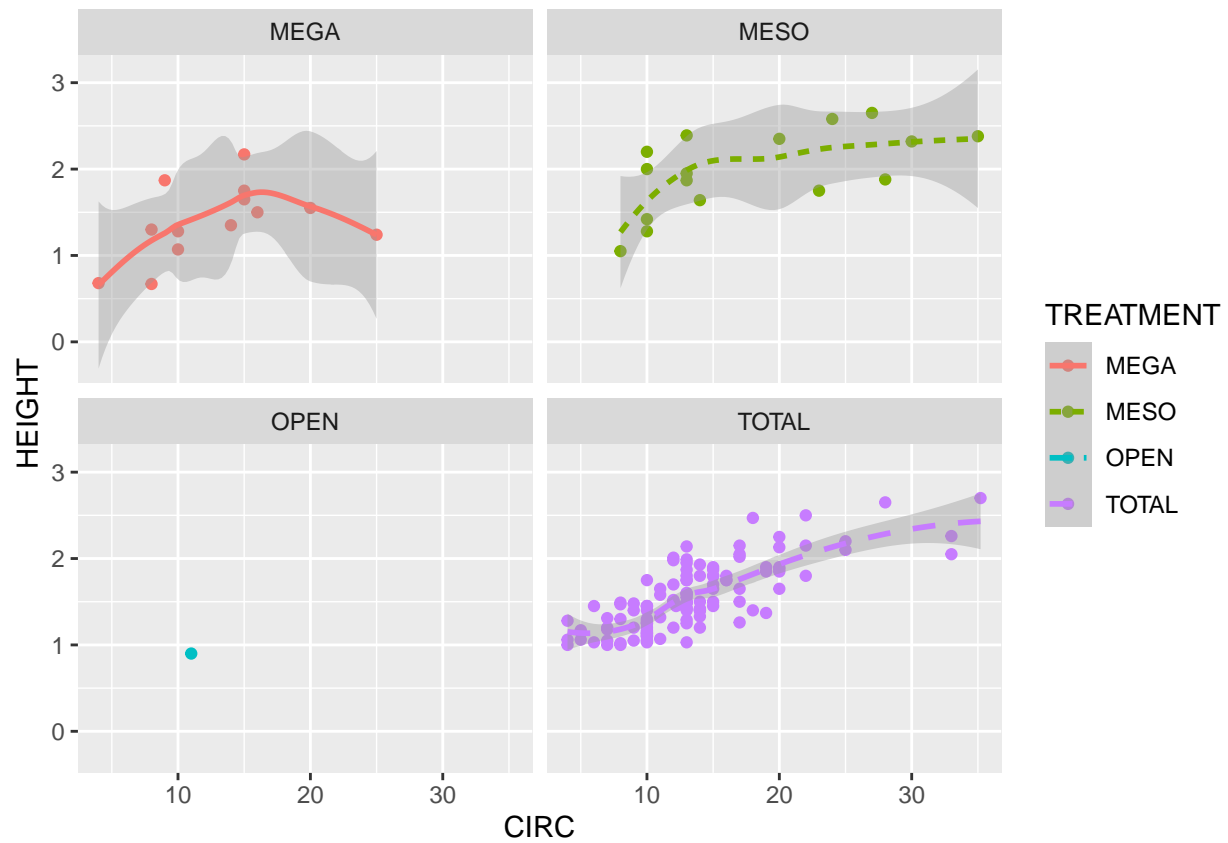
```
ggplot(data = acacia, mapping = aes(x = CIRC, y = HEIGHT, color = TREATMENT, linetype = TREATMENT)) +
  geom_smooth(method = "loess") +
  facet_wrap(~TREATMENT)
```

```
## Warning: Ignoring unknown parameters: method
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 4 rows containing non-finite values (stat_smooth).
```

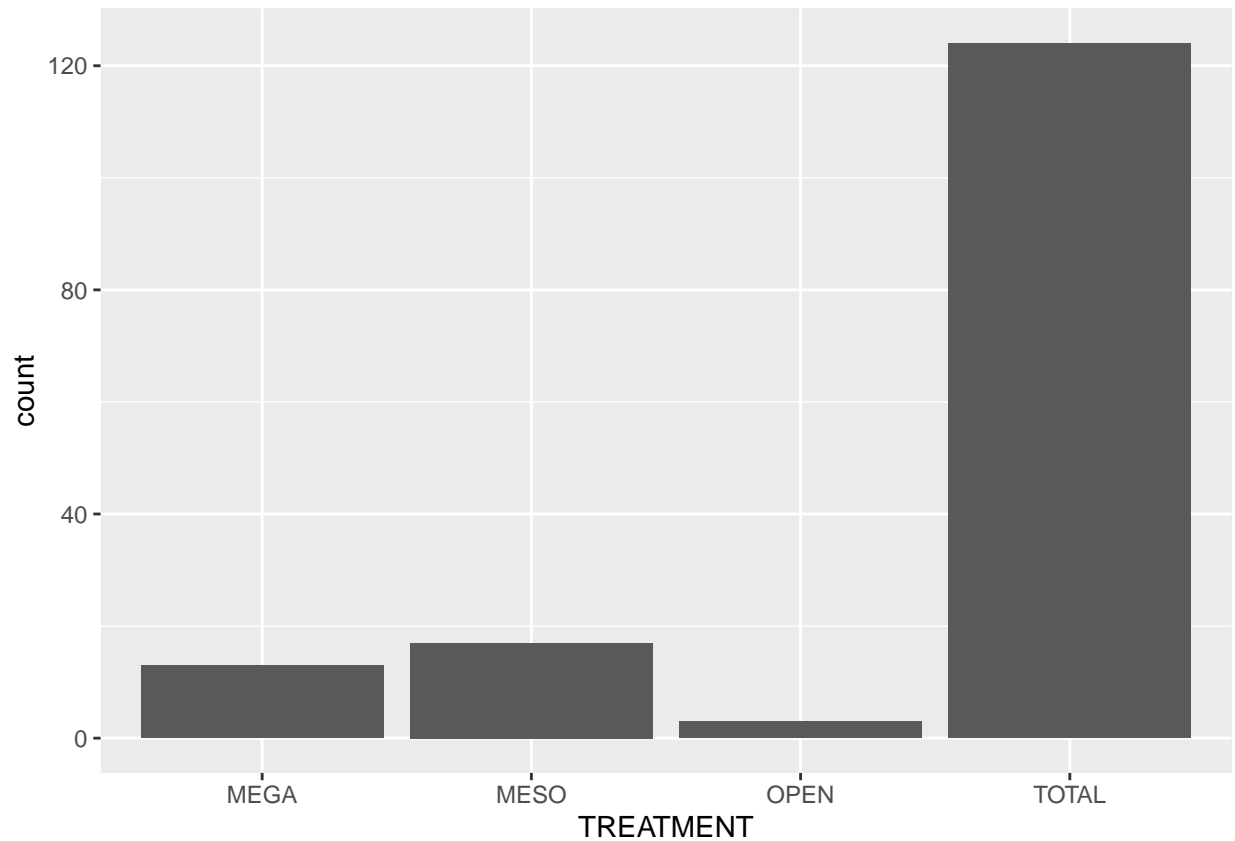
```
## Warning: Removed 4 rows containing missing values (geom_point).
```



Histograms and barplots

For bar plots use the `geom_bar()` function:

```
ggplot(data = acacia, aes(x = TREATMENT)) +  
  geom_bar()
```

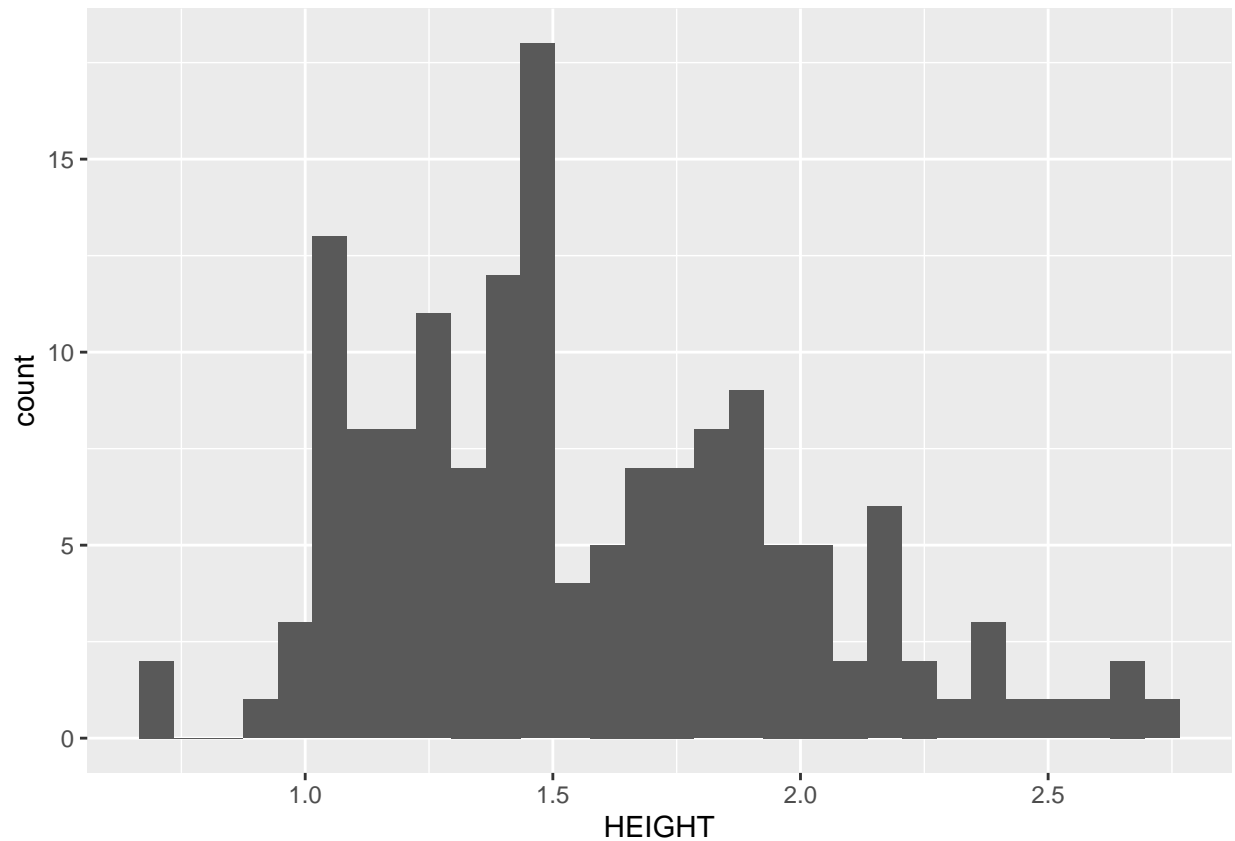


If you want to see the distribution of a continuous variable we use the `geom_histogram()` function:

```
ggplot(data = acacia, mapping = aes(x = HEIGHT)) +  
  geom_histogram()
```

```
## 'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.
```

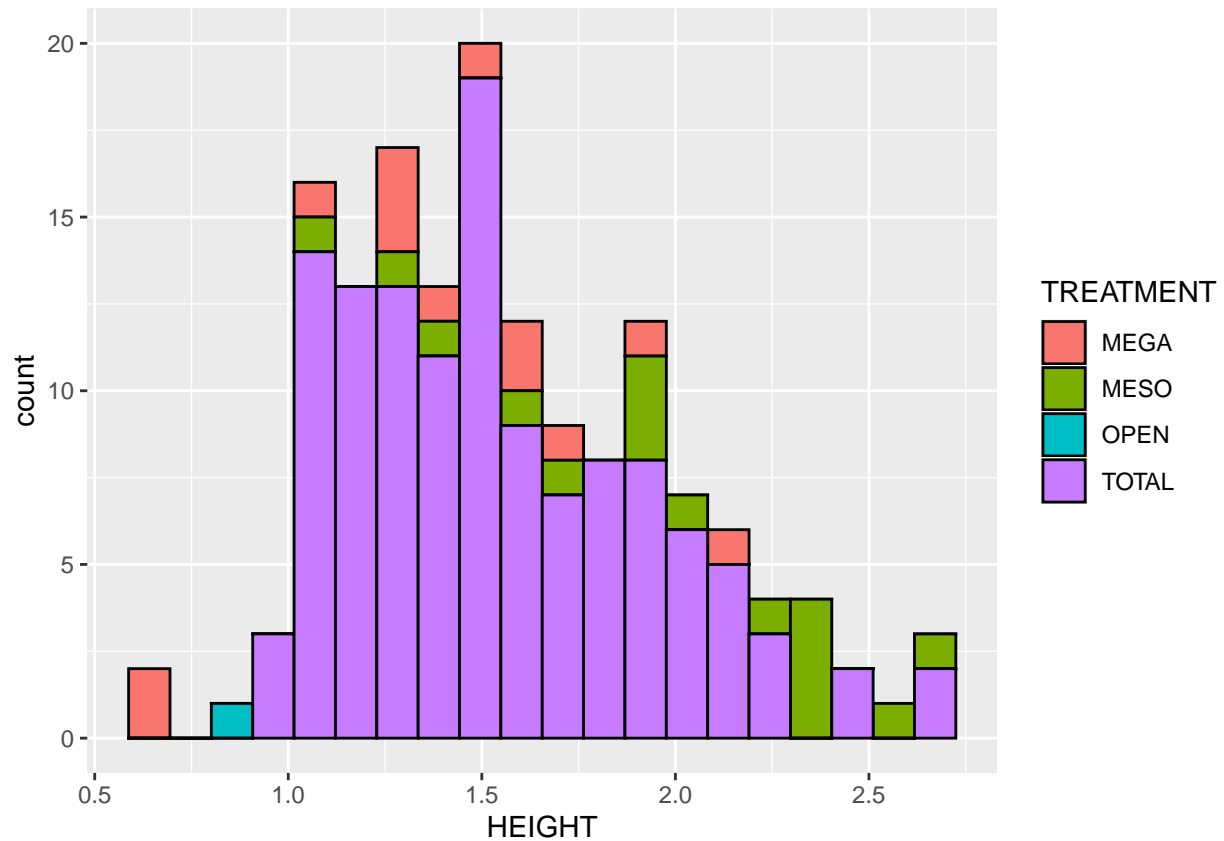
```
## Warning: Removed 4 rows containing non-finite values (stat_bin).
```



The 'fill =' command gives the inside of the bar color and the 'color = ""' command gives the outline color of the bars

```
ggplot(data = acacia, mapping = aes(x = HEIGHT, fill = TREATMENT)) +  
  geom_histogram(bins = 20, color = "black")
```

```
## Warning: Removed 4 rows containing non-finite values (stat_bin).
```



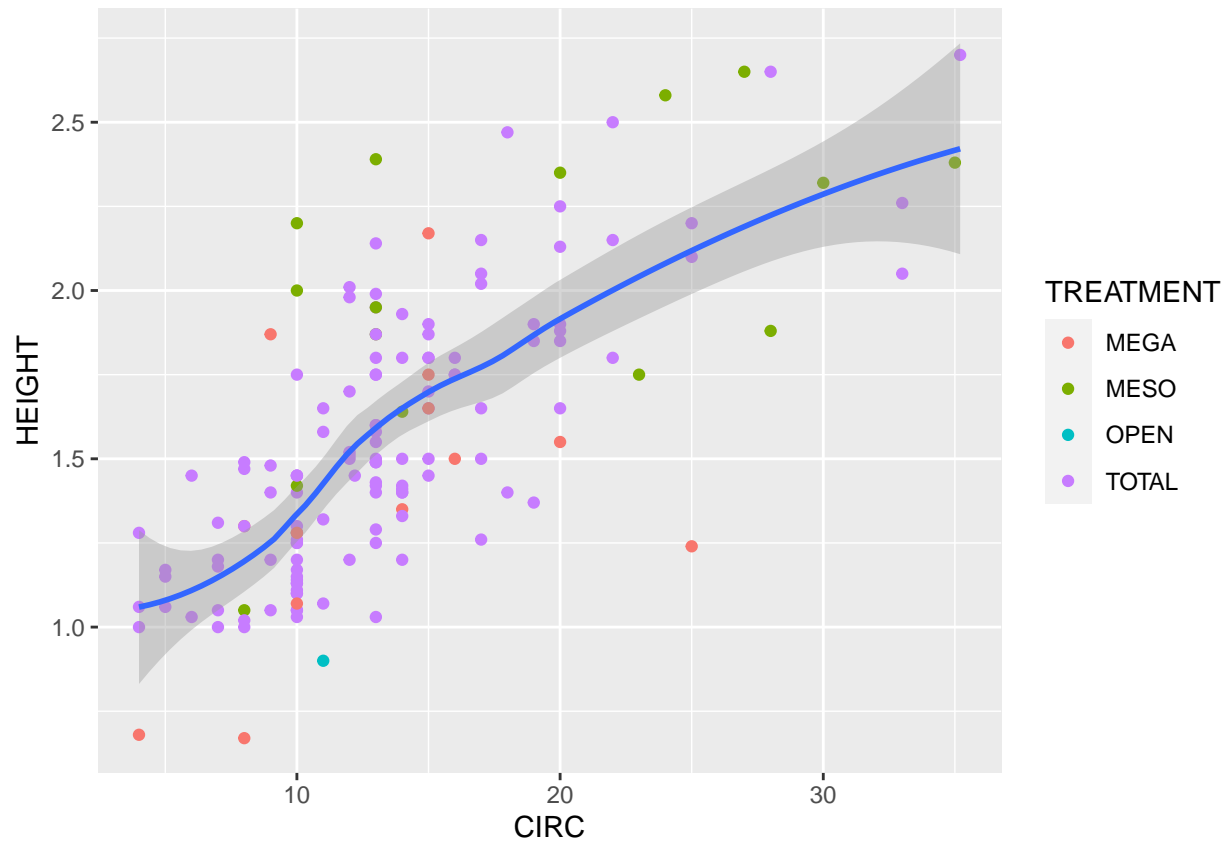
layer multiple data from the same or different data sets

```
ggplot() +
  geom_point(data = acacia,
            mapping = aes(x = CIRC, y = HEIGHT,
                          color = TREATMENT)) +
  geom_smooth(data = acacia,
            mapping = aes(x = CIRC, y = HEIGHT))
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 4 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 4 rows containing missing values (geom_point).
```

Save images as files

```
ggsave(filename = "acacia_by_treatment.pdf")

## Saving 6.5 x 4.5 in image

## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'

## Warning: Removed 4 rows containing non-finite values (stat_smooth).

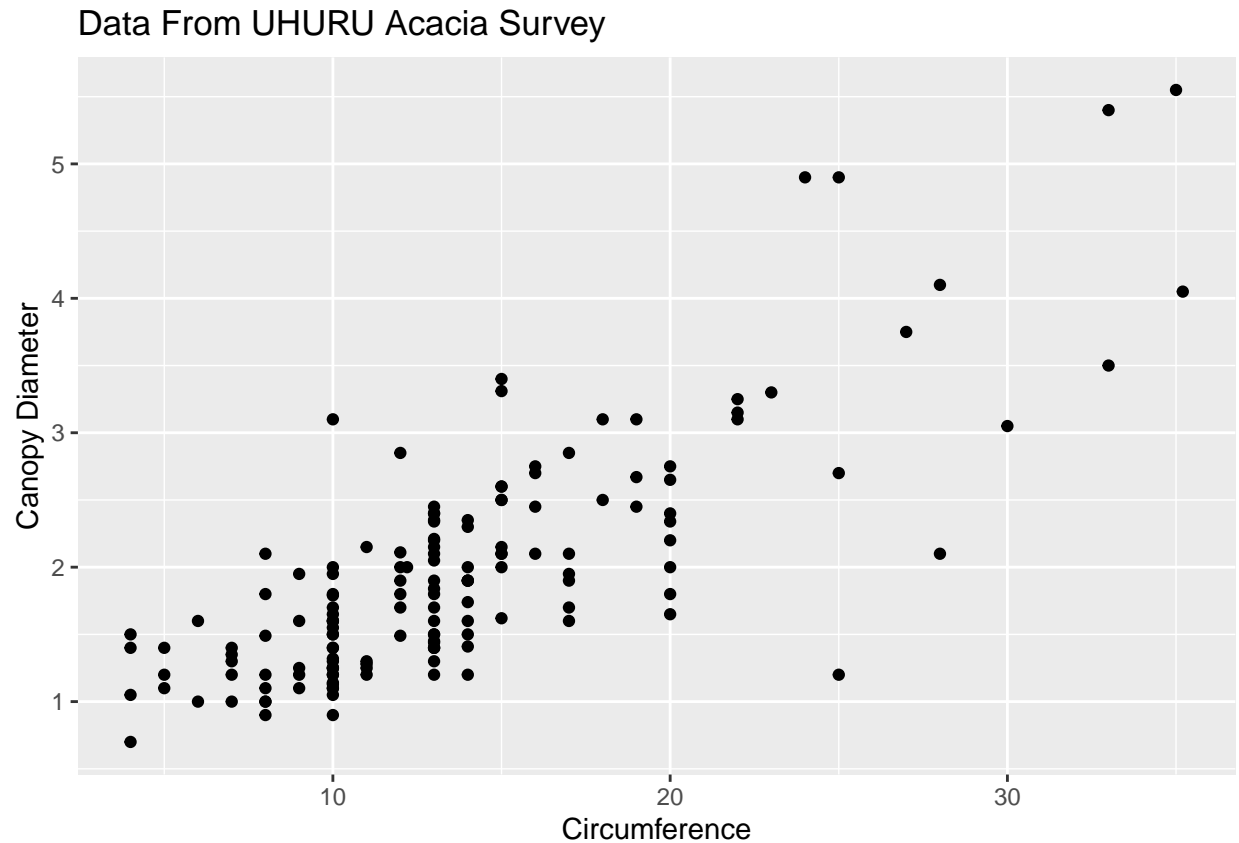
## Warning: Removed 4 rows containing missing values (geom_point).
```

In class exercise 1.

1.

```
ggplot() +
  geom_point(data = acacia,
             mapping = aes(x = CIRC, y = AXIS1, )) +
  labs(x = "Circumference", y = "Canopy Diameter", title = "Data From UHURU Acacia Survey ")
```

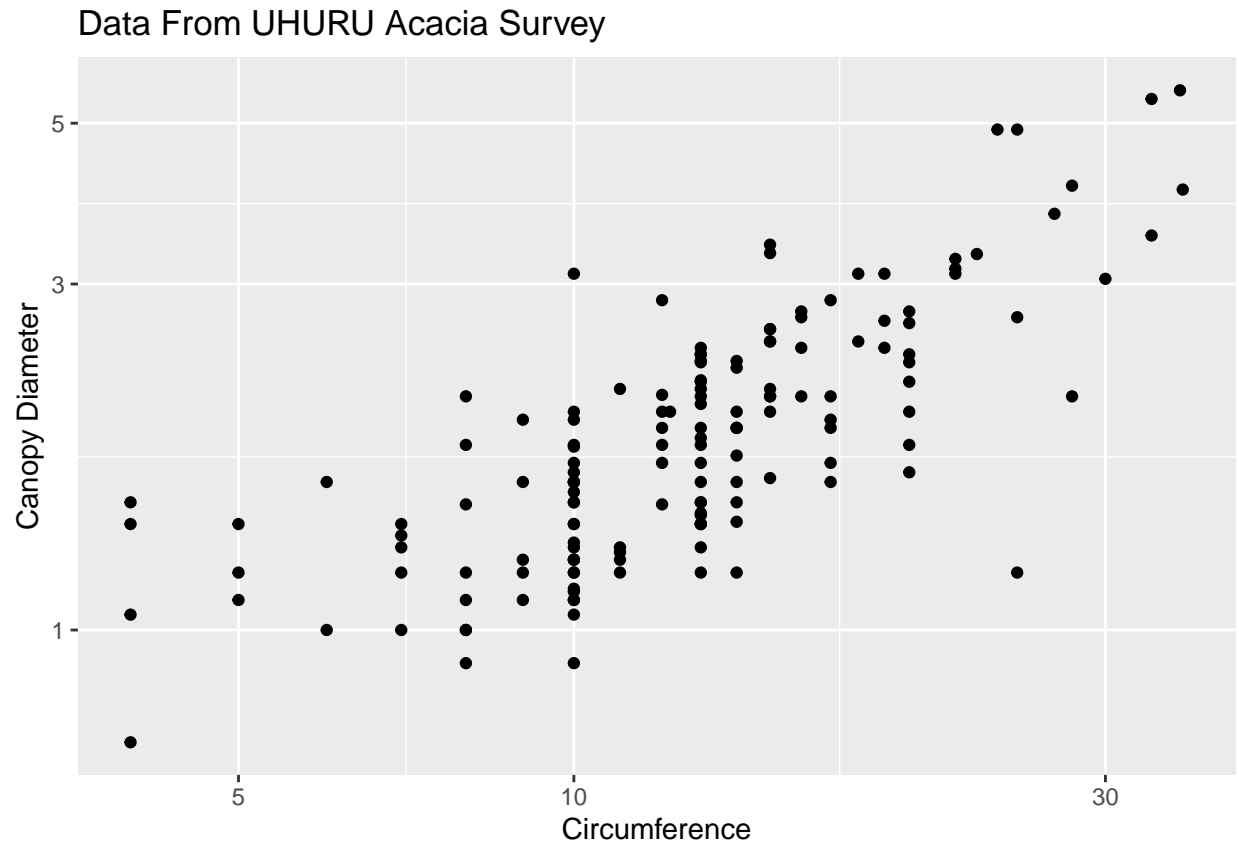
```
## Warning: Removed 4 rows containing missing values (geom_point).
```



```
# 2.
```

```
ggplot() +  
  geom_point(data = acacia,  
             mapping = aes(x = CIRC, y = AXIS1, )) +  
  labs(x = "Circumference", y = "Canopy Diameter", title = "Data From UHURU Acacia Survey ") +  
  scale_x_log10() +  
  scale_y_log10()
```

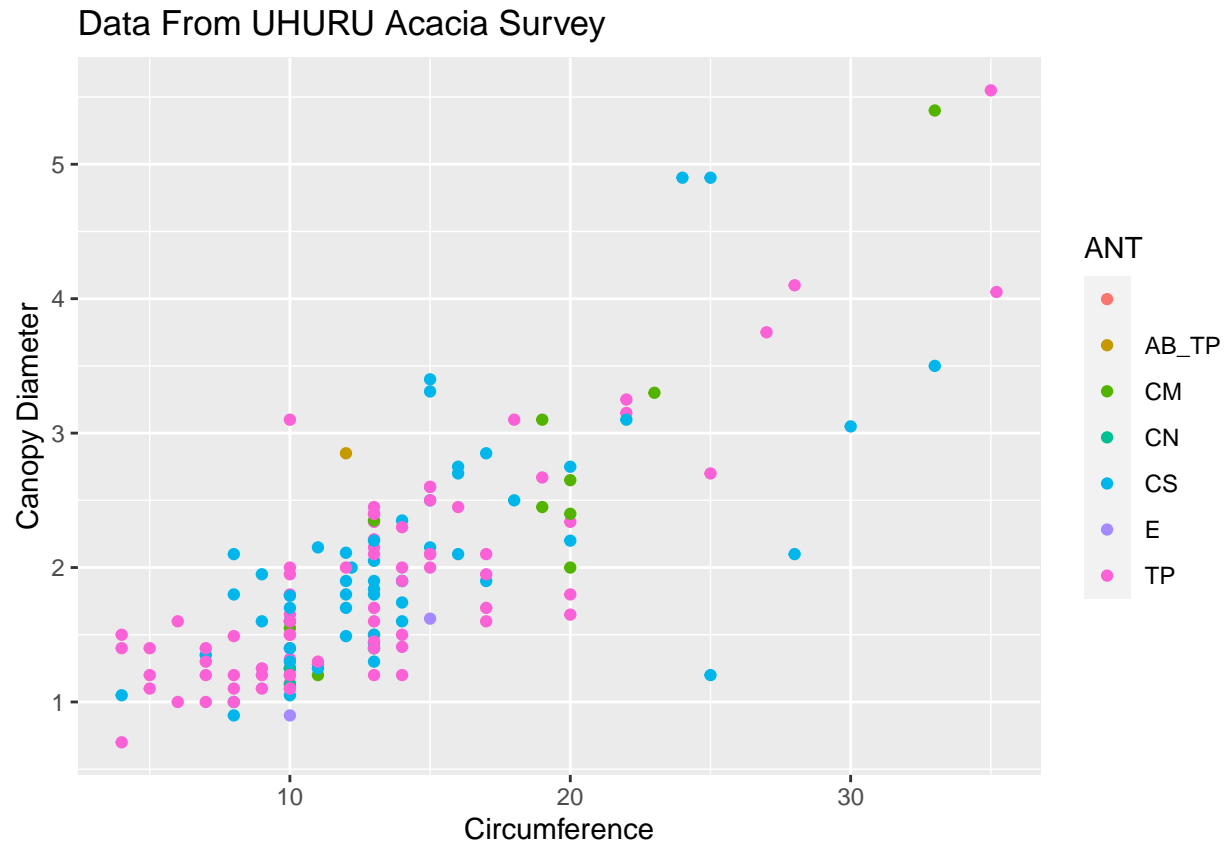
```
## Warning: Removed 4 rows containing missing values (geom_point).
```



3.

```
ggplot() +  
  geom_point(data = acacia,  
            mapping = aes(x = CIRC, y = AXIS1,  
                          color = ANT)) +  
  labs(x = "Circumference", y = "Canopy Diameter", title = "Data From UHURU Acacia Survey ")
```

Warning: Removed 4 rows containing missing values (geom_point).

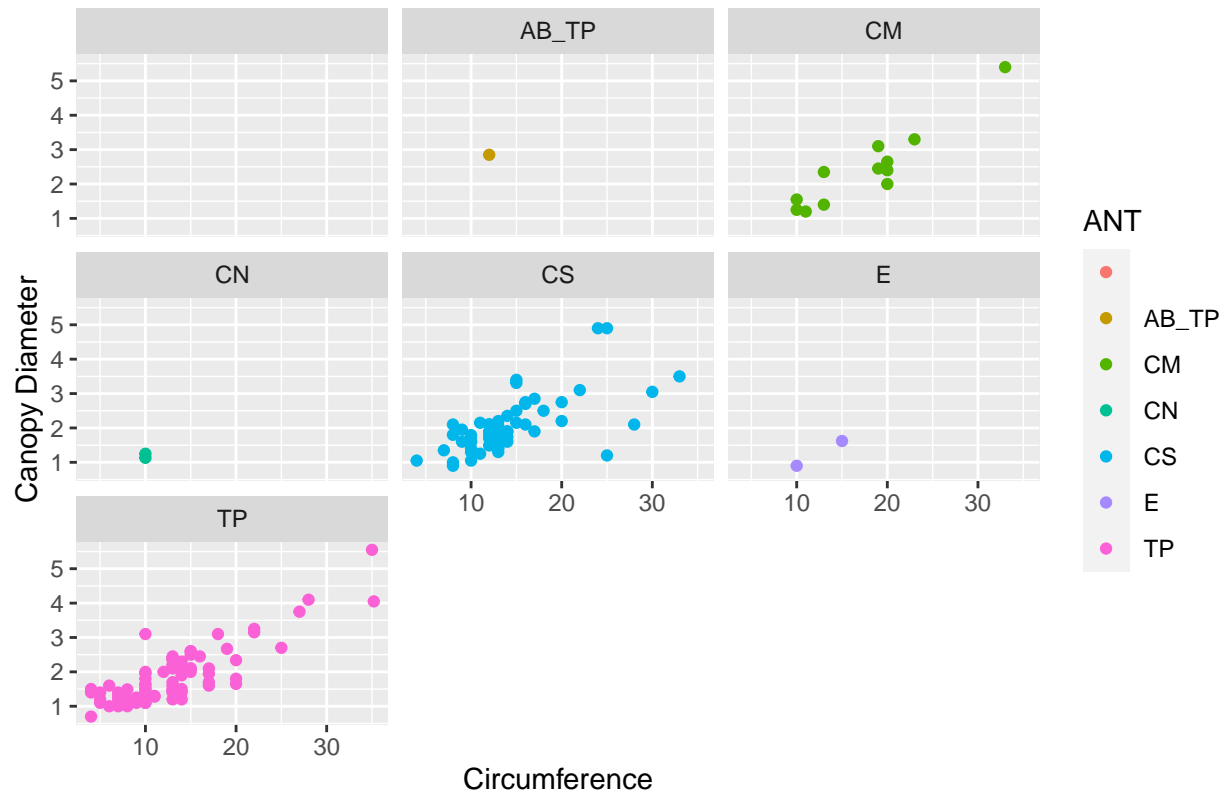


4.

```
ggplot() +
  geom_point(data = acacia,
             mapping = aes(x = CIRC, y = AXIS1,
                           color = ANT)) +
  labs(x = "Circumference", y = "Canopy Diameter", title = "Data From UHURU Acacia Survey ") +
  geom_point() +
  facet_wrap(~ANT)
```

Warning: Removed 4 rows containing missing values (geom_point).

Data From UHURU Acacia Survey



5.

```
ggplot(data = acacia,
       mapping = aes(x = CIRC, y = AXIS1,
                     color = ANT)) +
  labs(x = "Circumference", y = "Canopy Diameter", title = "Data From UHURU Acacia Survey ") +
  geom_point() +
  geom_smooth()+
  facet_wrap(~ANT)
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 4 rows containing non-finite values (stat_smooth).
```

```
## Warning in simpleLoess(y, x, w, span, degree = degree, parametric =
## parametric, : span too small. fewer data values than degrees of freedom.
```

```
## Warning in simpleLoess(y, x, w, span, degree = degree, parametric =
## parametric, : at 9.975
```

```
## Warning in simpleLoess(y, x, w, span, degree = degree, parametric =
## parametric, : radius 0.000625
```

```
## Warning in simpleLoess(y, x, w, span, degree = degree, parametric =
## parametric, : all data on boundary of neighborhood. make span bigger
```

```

## Warning in simpleLoess(y, x, w, span, degree = degree, parametric =
## parametric, : pseudoinverse used at 9.975

## Warning in simpleLoess(y, x, w, span, degree = degree, parametric =
## parametric, : neighborhood radius 0.025

## Warning in simpleLoess(y, x, w, span, degree = degree, parametric =
## parametric, : reciprocal condition number 1

## Warning in simpleLoess(y, x, w, span, degree = degree, parametric =
## parametric, : at 15.025

## Warning in simpleLoess(y, x, w, span, degree = degree, parametric =
## parametric, : radius 0.000625

## Warning in simpleLoess(y, x, w, span, degree = degree, parametric =
## parametric, : all data on boundary of neighborhood. make span bigger

## Warning in simpleLoess(y, x, w, span, degree = degree, parametric =
## parametric, : There are other near singularities as well. 0.000625

## Warning in simpleLoess(y, x, w, span, degree = degree, parametric =
## parametric, : zero-width neighborhood. make span bigger

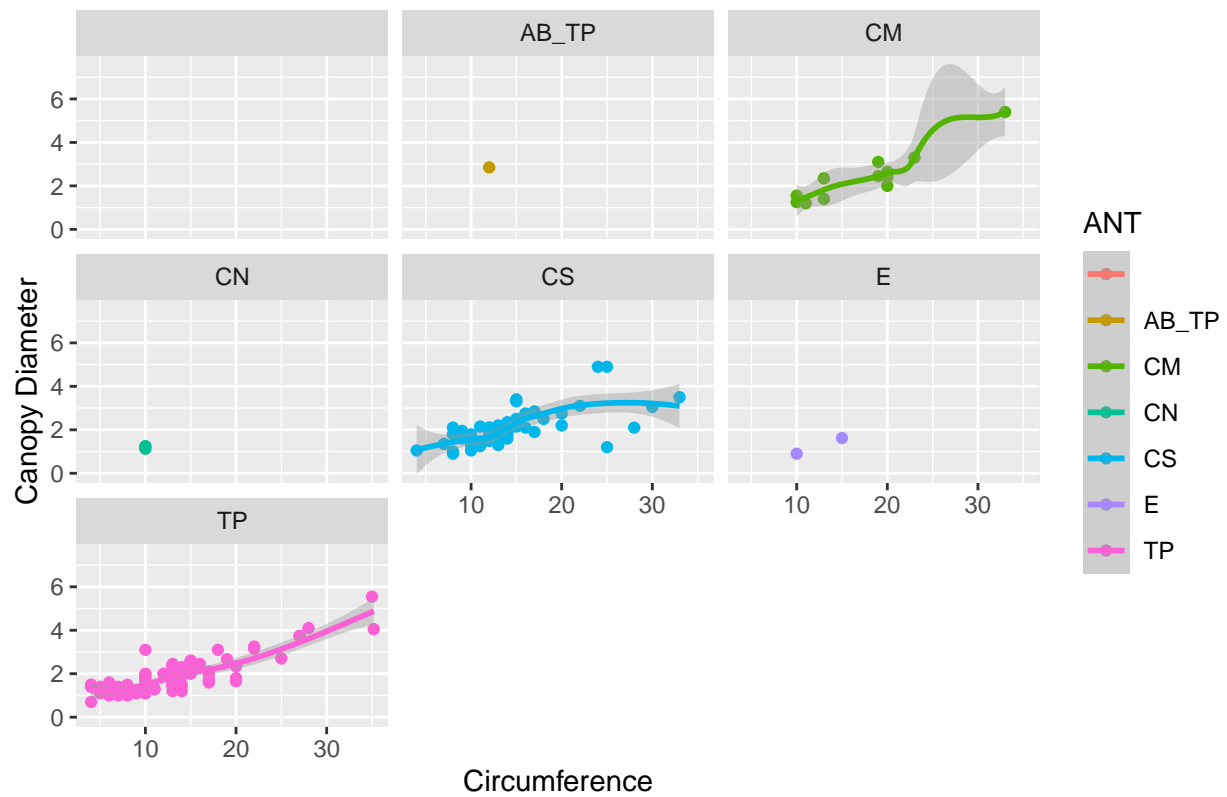
## Warning in simpleLoess(y, x, w, span, degree = degree, parametric =
## parametric, : zero-width neighborhood. make span bigger

## Warning: Computation failed in 'stat_smooth()':
## NA/NaN/Inf in foreign function call (arg 5)

## Warning: Removed 4 rows containing missing values (geom_point).

```

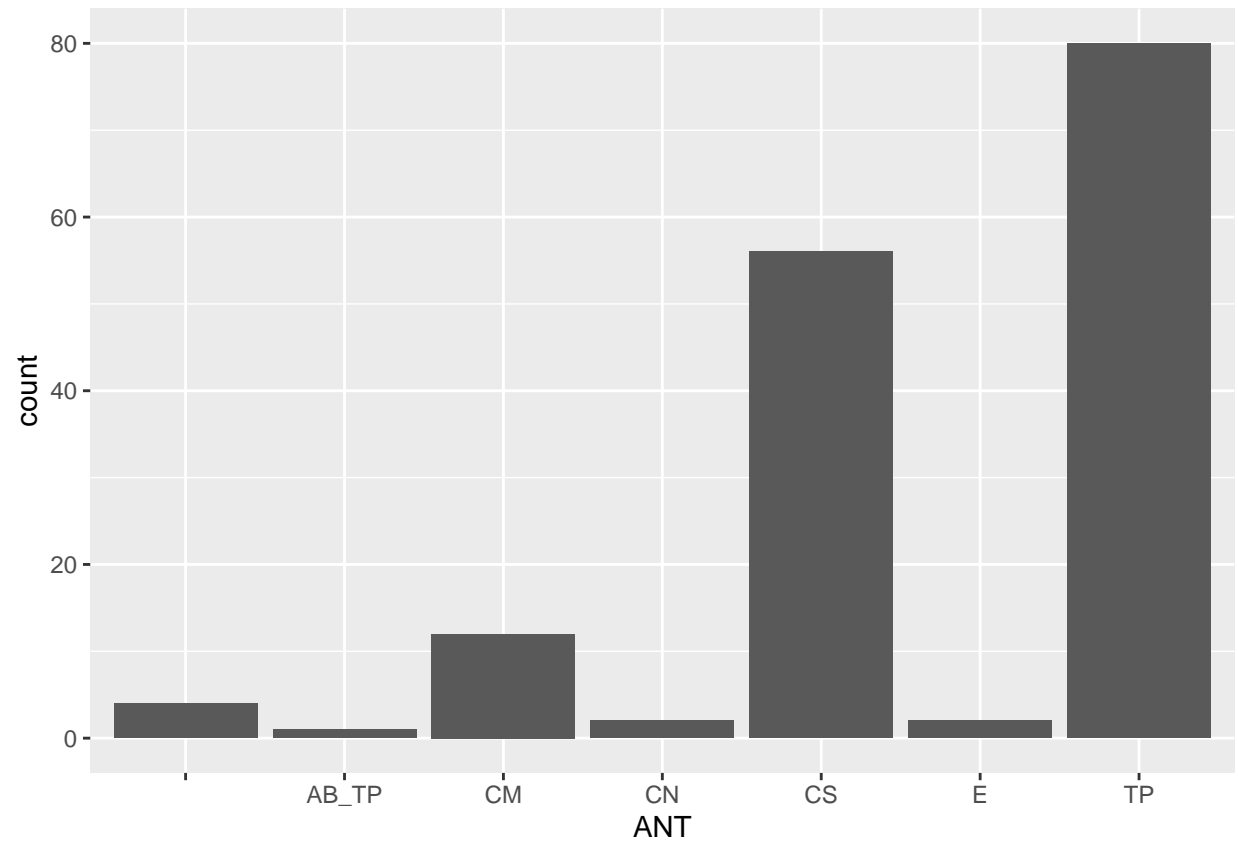
Data From UHURU Acacia Survey



#Exercise 2. Histograms

1.

```
ggplot(data = acacia, aes(x = ANT)) +
  geom_bar()
```

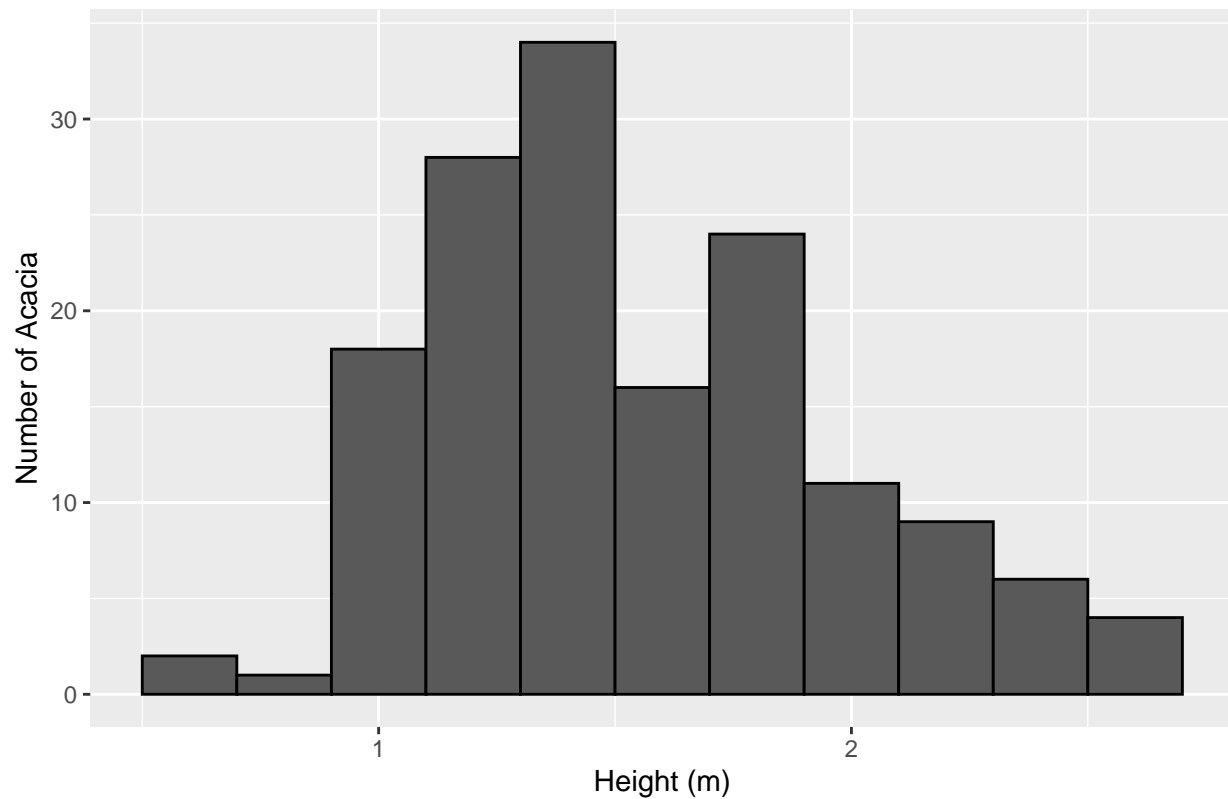


2.

```
ggplot(data = acacia, mapping = aes(x = HEIGHT)) +
  labs(x = "Height (m)", y = "Number of Acacia", title = "Data From UHURU Acacia Survey ") +
  geom_histogram(binwidth = .20 , color = "black")
```

```
## Warning: Removed 4 rows containing non-finite values (stat_bin).
```


Data From UHURU Acacia Survey



3.

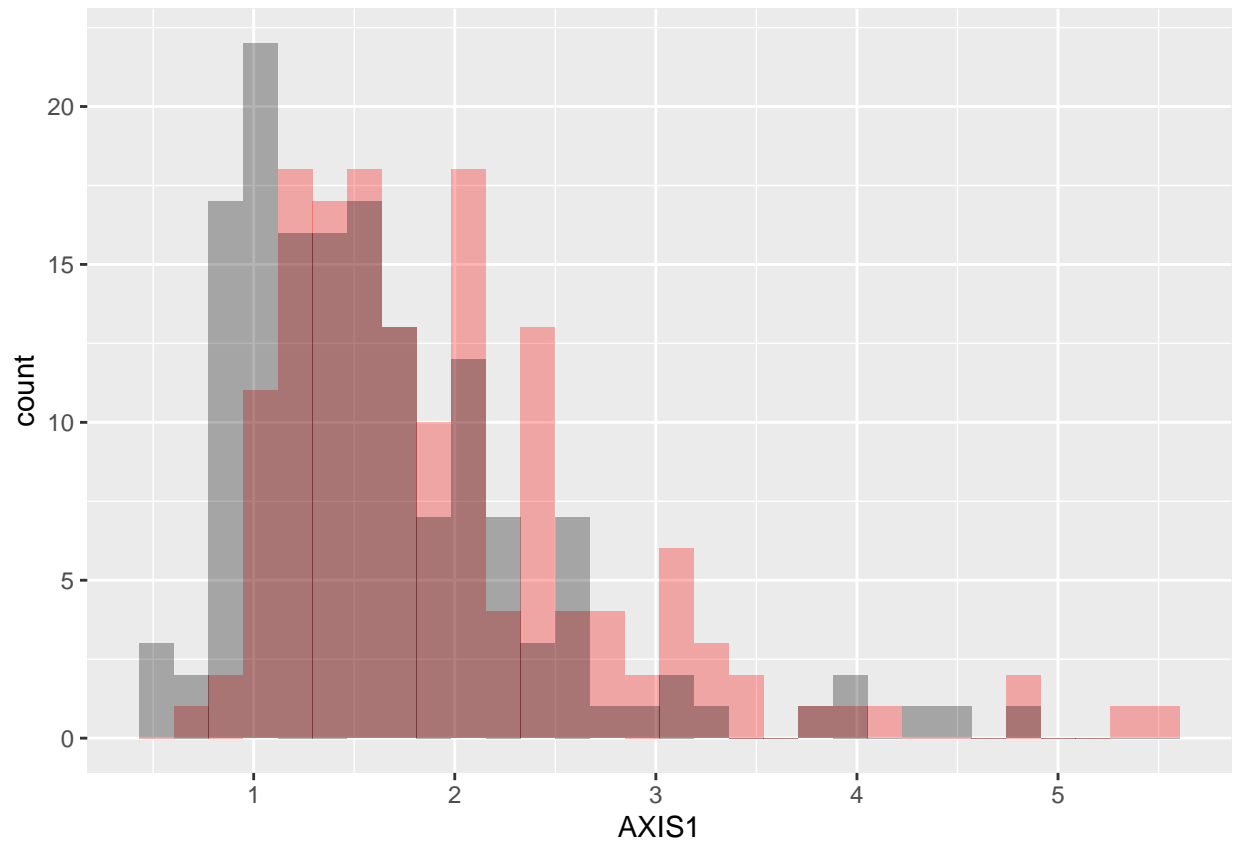
```
ggplot() +
  geom_histogram(data = acacia,
                 mapping = aes(x = AXIS1) , alpha = 0.3, fill = "red") +
  geom_histogram(data = acacia,
                 mapping = aes(x = AXIS2), alpha = 0.3, fill = "black")

## 'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.

## Warning: Removed 4 rows containing non-finite values (stat_bin).

## 'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.

## Warning: Removed 4 rows containing non-finite values (stat_bin).
```



4.

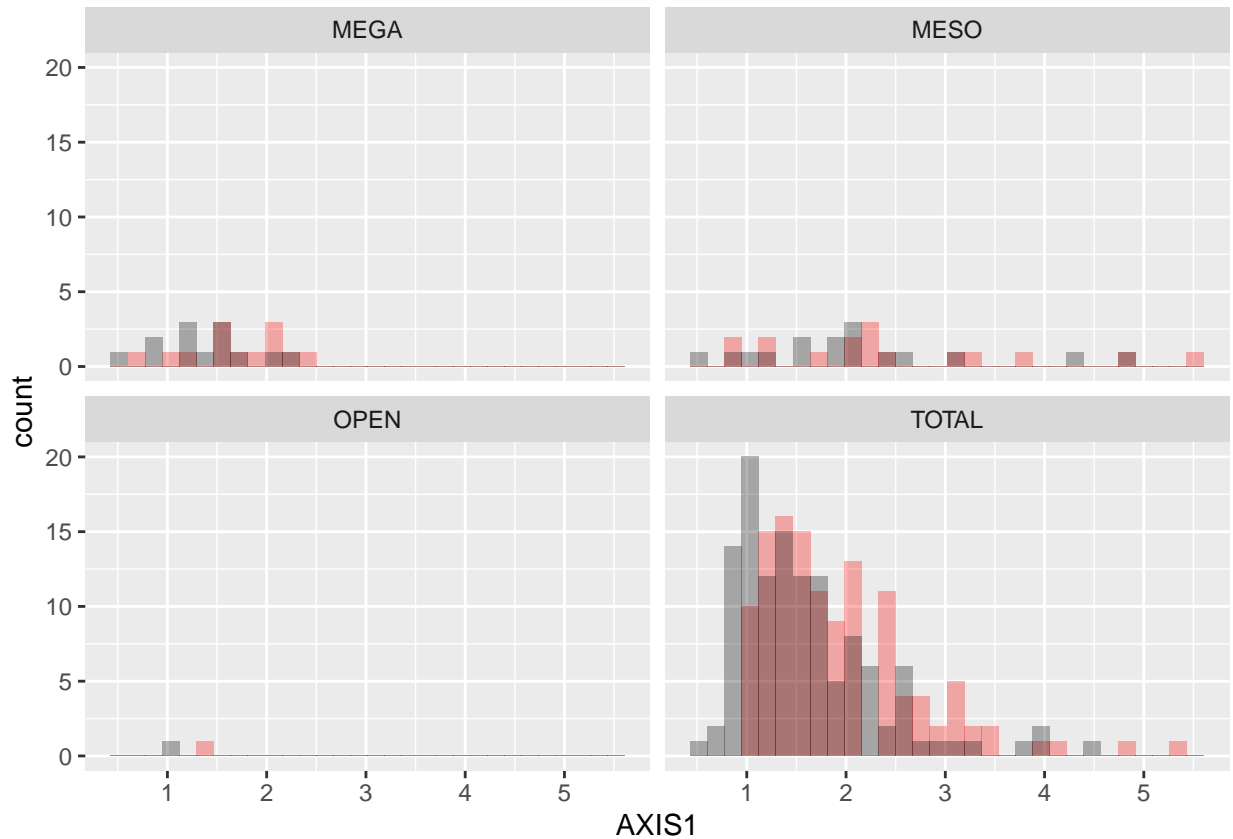
```
ggplot() +
  geom_histogram(data = acacia,
    mapping = aes(x = AXIS1) , alpha = 0.3, fill = "red") +
  geom_histogram(data = acacia,
    mapping = aes(x = AXIS2), alpha = 0.3, fill = "black") +
  facet_wrap(~ TREATMENT)
```

```
## 'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.
```

```
## Warning: Removed 4 rows containing non-finite values (stat_bin).
```

```
## 'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.
```

```
## Warning: Removed 4 rows containing non-finite values (stat_bin).
```



Home exercises

Exercised 3.

#1 To installed a package use the function 'install.packages(" ")'

```
#install.packages("readr")
```

function library allows me to open the directory where the package i just downloaded is stored

```
library(readr)
```

getting and setting my working directory

```
setwd("~/Bio 195-197/Data Science/raw-data")
getwd()
```

```
## [1] "/Users/atziri/Bio 195-197/Data Science/raw-data"
```

```
read_tsv(file = "/Users/atziri/Bio 195-197/Data Science/raw-data/TREE_SURVEYS.txt")
```

```
## Rows: 7508 Columns: 16-- Column specification -----
```

```
## Delimiter: "\t"
## chr (9): SITE, TREATMENT, PLOT, SPECIES, DEAD, HEIGHT, AXIS_2, MEASUREMENT, ...
## dbl (7): SURVEY, YEAR, BLOCK, ORIGINAL_TAG, NEW_TAG, AXIS_1, CIRC
## 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: 7,508 x 16
##   SURVEY YEAR SITE TREATMENT BLOCK PLOT SPECIES ORIGI~1 NEW_TAG DEAD HEIGHT
##   <dbl> <dbl> <chr> <chr>    <dbl> <chr> <chr>    <dbl>    <dbl> <chr> <chr>
## 1     1     2009 SOUTH TOTAL      2 S2T0~ Acacia~      1      NA N    3.4
## 2     2     2010 SOUTH TOTAL      2 S2T0~ Acacia~      1      NA N    3.32
## 3     3     2011 SOUTH TOTAL      2 S2T0~ Acacia~      1      NA N    3.65
## 4     4     2012 SOUTH TOTAL      2 S2T0~ Acacia~      1      NA N    3.74
## 5     5     2013 SOUTH TOTAL      2 S2T0~ Acacia~      1      NA N    3.59
## 6     1     2009 SOUTH TOTAL      2 S2T0~ Acacia~      2      NA N    2.3
## 7     2     2010 SOUTH TOTAL      2 S2T0~ Acacia~      2      NA N    2.32
## 8     3     2011 SOUTH TOTAL      2 S2T0~ Acacia~      2      NA N    2.75
## 9     4     2012 SOUTH TOTAL      2 S2T0~ Acacia~      2      NA Y    <NA>
## 10    5     2013 SOUTH TOTAL      2 S2T0~ Acacia~      2      NA N    2.86
## # ... with 7,498 more rows, 5 more variables: AXIS_1 <dbl>, AXIS_2 <chr>,
## #   CIRC <dbl>, MEASUREMENT <chr>, STEMS <chr>, and abbreviated variable name
## #   1: ORIGINAL_TAG
```

```
read_tsv(file = "../raw-data/TREE_SURVEYS.txt")
```

```
## Rows: 7508 Columns: 16-- Column specification -----
## Delimiter: "\t"
## chr (9): SITE, TREATMENT, PLOT, SPECIES, DEAD, HEIGHT, AXIS_2, MEASUREMENT, ...
## dbl (7): SURVEY, YEAR, BLOCK, ORIGINAL_TAG, NEW_TAG, AXIS_1, CIRC
## 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: 7,508 x 16
##   SURVEY YEAR SITE TREATMENT BLOCK PLOT SPECIES ORIGI~1 NEW_TAG DEAD HEIGHT
##   <dbl> <dbl> <chr> <chr>    <dbl> <chr> <chr>    <dbl>    <dbl> <chr> <chr>
## 1     1     2009 SOUTH TOTAL      2 S2T0~ Acacia~      1      NA N    3.4
## 2     2     2010 SOUTH TOTAL      2 S2T0~ Acacia~      1      NA N    3.32
## 3     3     2011 SOUTH TOTAL      2 S2T0~ Acacia~      1      NA N    3.65
## 4     4     2012 SOUTH TOTAL      2 S2T0~ Acacia~      1      NA N    3.74
## 5     5     2013 SOUTH TOTAL      2 S2T0~ Acacia~      1      NA N    3.59
## 6     1     2009 SOUTH TOTAL      2 S2T0~ Acacia~      2      NA N    2.3
## 7     2     2010 SOUTH TOTAL      2 S2T0~ Acacia~      2      NA N    2.32
## 8     3     2011 SOUTH TOTAL      2 S2T0~ Acacia~      2      NA N    2.75
## 9     4     2012 SOUTH TOTAL      2 S2T0~ Acacia~      2      NA Y    <NA>
## 10    5     2013 SOUTH TOTAL      2 S2T0~ Acacia~      2      NA N    2.86
## # ... with 7,498 more rows, 5 more variables: AXIS_1 <dbl>, AXIS_2 <chr>,
## #   CIRC <dbl>, MEASUREMENT <chr>, STEMS <chr>, and abbreviated variable name
## #   1: ORIGINAL_TAG
```

```
#2 reading the file of the data set with function read_tsv and then assign it to the trees object
```

```
trees <- read_tsv(file = "../raw-data/TREE_SURVEYS.txt",
  col_types = list(HEIGHT = col_double(),
    AXIS_2 = col_double()))
```

```
## Warning: One or more parsing issues, call 'problems()' on your data frame for details,
## e.g.:
##   dat <- vroom(...)
##   problems(dat)
```

function gives me a summary of the object trees which is now the dataset needed for the assignment

```
summary(trees)
```

```
##      SURVEY      YEAR      SITE      TREATMENT
## Min.   :1   Min.   :2009 Length:7508 Length:7508
## 1st Qu.:2   1st Qu.:2010 Class :character Class :character
## Median :3   Median :2011 Mode  :character Mode  :character
## Mean   :3   Mean   :2011
## 3rd Qu.:4   3rd Qu.:2012
## Max.   :5   Max.   :2013
## NA's   :7   NA's   :7
##      BLOCK      PLOT      SPECIES      ORIGINAL_TAG
## Min.   :1.000 Length:7508 Length:7508 Min.   : 1
## 1st Qu.:1.000 Class :character Class :character 1st Qu.:1003
## Median :2.000 Mode  :character Mode  :character Median :1890
## Mean   :2.031
## 3rd Qu.:3.000
## Max.   :3.000
## NA's   :7
##      NEW_TAG      DEAD      HEIGHT      AXIS_1
## Min.   :1674 Length:7508 Min.   : 0.230 Min.   : 0.200
## 1st Qu.:3549 Class :character 1st Qu.: 1.850 1st Qu.: 1.900
## Median :3582 Mode  :character Median : 2.390 Median : 2.800
## Mean   :3732
## 3rd Qu.:4504
## Max.   :4600
## NA's   :7458
##      HEIGHT      AXIS_1
## Min.   : 0.230 Min.   : 0.200
## 1st Qu.: 1.850 1st Qu.: 1.900
## Median : 2.390 Median : 2.800
## Mean   : 2.515 Mean   : 3.083
## 3rd Qu.: 3.000 3rd Qu.: 4.000
## Max.   :10.000 Max.   :47.000
## NA's   :207 NA's   :210
##      AXIS_2      CIRC      MEASUREMENT      STEMS
## Min.   : 0.000 Min.   : 0.3 Length:7508 Length:7508
## 1st Qu.: 1.800 1st Qu.: 7.0 Class :character Class :character
## Median : 2.610 Median :11.5 Mode  :character Mode  :character
## Mean   : 2.948 Mean   :16.0
## 3rd Qu.: 3.800 3rd Qu.:20.4
## Max.   :44.000 Max.   :136.0
## NA's   :215 NA's   :414
```

#3 add a new column with the command 'nameofdataframehere\$newcolnamehere' then give it the command for the value we need the estimated canopy area calculated as the value in the AXIS_1 column times the value in the AXIS_2 column.

```
trees$canopy_area <- trees$AXIS_1 * trees$AXIS_2
```

#4 Subset the trees data frame with just the SURVEY, YEAR, SITE, and canopy_area columns.

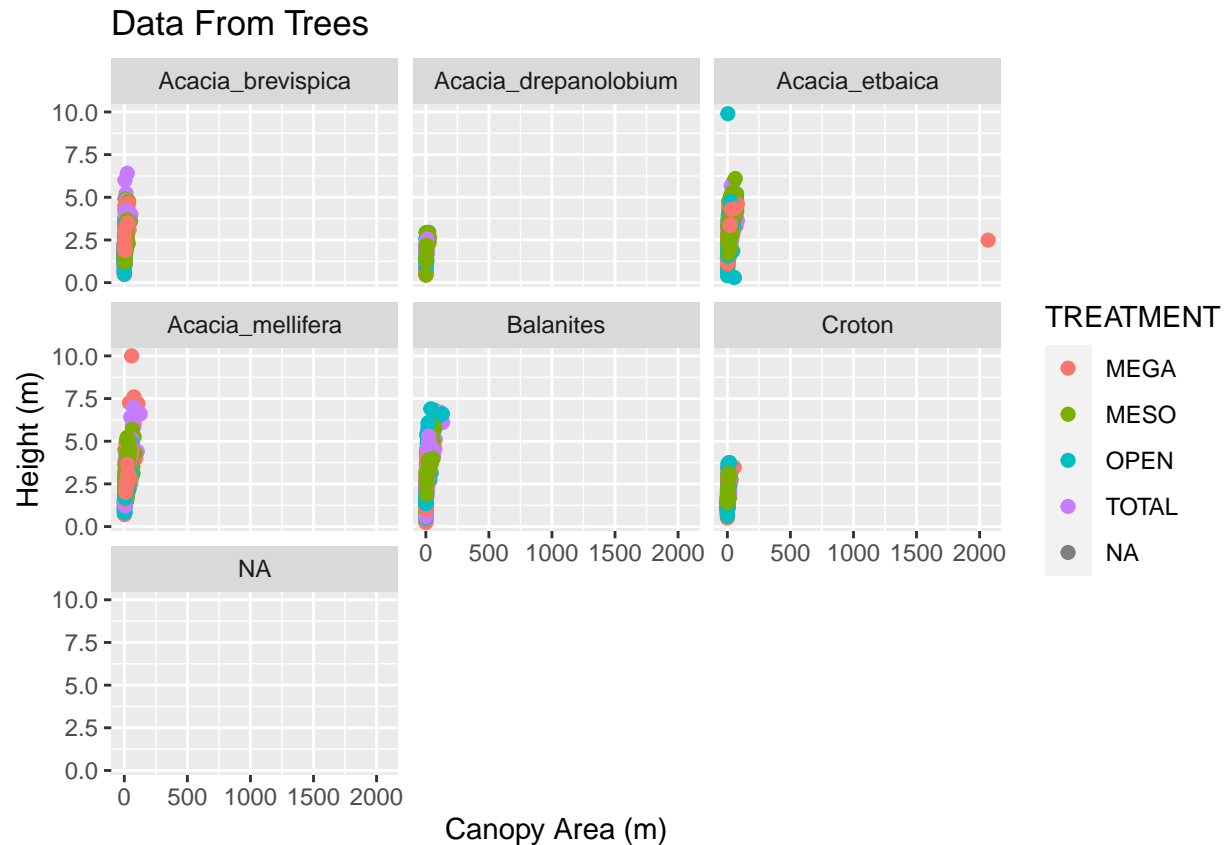
```
subset(trees, select = c('SURVEY' , 'YEAR' , 'SITE' , 'canopy_area'))
```

```
## # A tibble: 7,508 x 4
##   SURVEY  YEAR SITE  canopy_area
##   <dbl> <dbl> <chr>      <dbl>
## 1      1      1 2009 SOUTH        30.5
## 2      2      2 2010 SOUTH        69.7
## 3      3      3 2011 SOUTH        79.6
## 4      4      4 2012 SOUTH        39.0
## 5      5      5 2013 SOUTH        40.8
## 6      1      1 2009 SOUTH         6.16
## 7      2      2 2010 SOUTH         7.29
## 8      3      3 2011 SOUTH        12.5
## 9      4      4 2012 SOUTH         NA
## 10     5      5 2013 SOUTH         9.62
## # ... with 7,498 more rows
```

#5 creating a scatter plot using the ggplot() function. using the mapping function to add the aesthetics of the plot such as the data used for the x and y axis using the 'geom_point()'function to with the command size to add a size to the plots using the facet_wrap(~SPECIES) function to create a subplot for each species. I used the function labs to add the labeling of each of the axis and the title of the graph.

```
ggplot(data = trees,
       mapping = aes(x = canopy_area, y = HEIGHT, color = TREATMENT )) +
  geom_point( size = 2) +
  facet_wrap(~SPECIES ) +
  labs(x = "Canopy Area (m)", y = "Height (m)", title = "Data From Trees ")
```

```
## Warning: Removed 215 rows containing missing values (geom_point).
```



#6 logical vector that evaluates all rows that are in AXIS_1 and AXIS_2 that are below 20 this wa to eliminate outliers all values above 20.

```
trees_2 <- trees$AXIS_1 <= 20 | trees$AXIS_2 <= 20
summary(trees_2)
```

```
##      Mode   FALSE    TRUE   NA's
## logical      1    7297    210
```

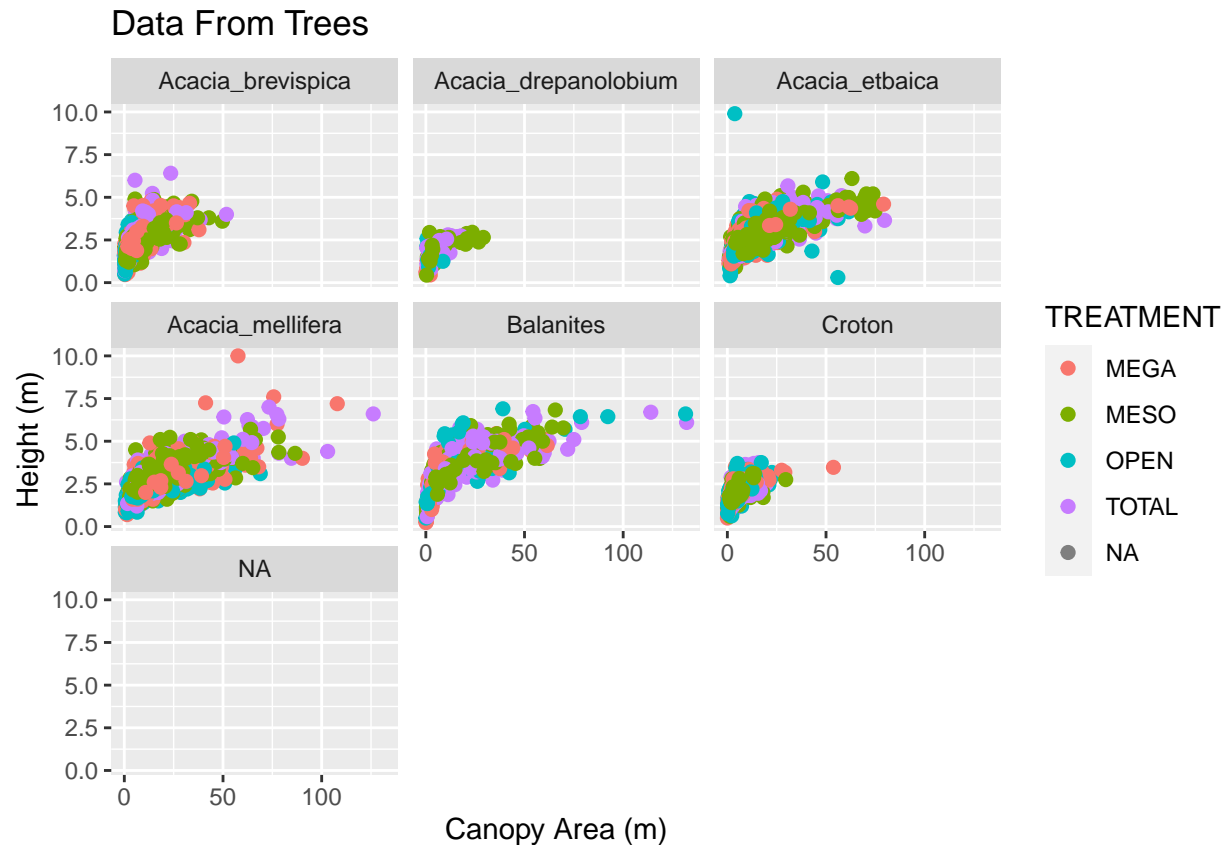
```
nrow(trees)
```

```
## [1] 7508
```

```
trees <- trees[trees_2, ]
```

```
ggplot(data = trees,
       mapping = aes(x = canopy_area, y = HEIGHT, color = TREATMENT )) +
  geom_point( size = 2) +
  facet_wrap(~SPECIES ) +
  labs(x = "Canopy Area (m)", y = "Height (m)", title = "Data From Trees ")
```

```
## Warning: Removed 215 rows containing missing values (geom_point).
```



#7 used the pipe symbol %>% which takes the output of the function and into another function as an argument this linked a sequence of analysis steps.

Assignment needed to be used to the right rather than the left

```
library(dplyr)
```

```
##
```

```
## Attaching package: 'dplyr'
```

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

```
##
```

```
## filter, lag
```

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

```
##
```

```
## intersect, setdiff, setequal, union
```

```
trees %>%
```

```
group_by(YEAR, SPECIES) %>%
```

```
  summarize(species_abundance=n()) -> abundance
```

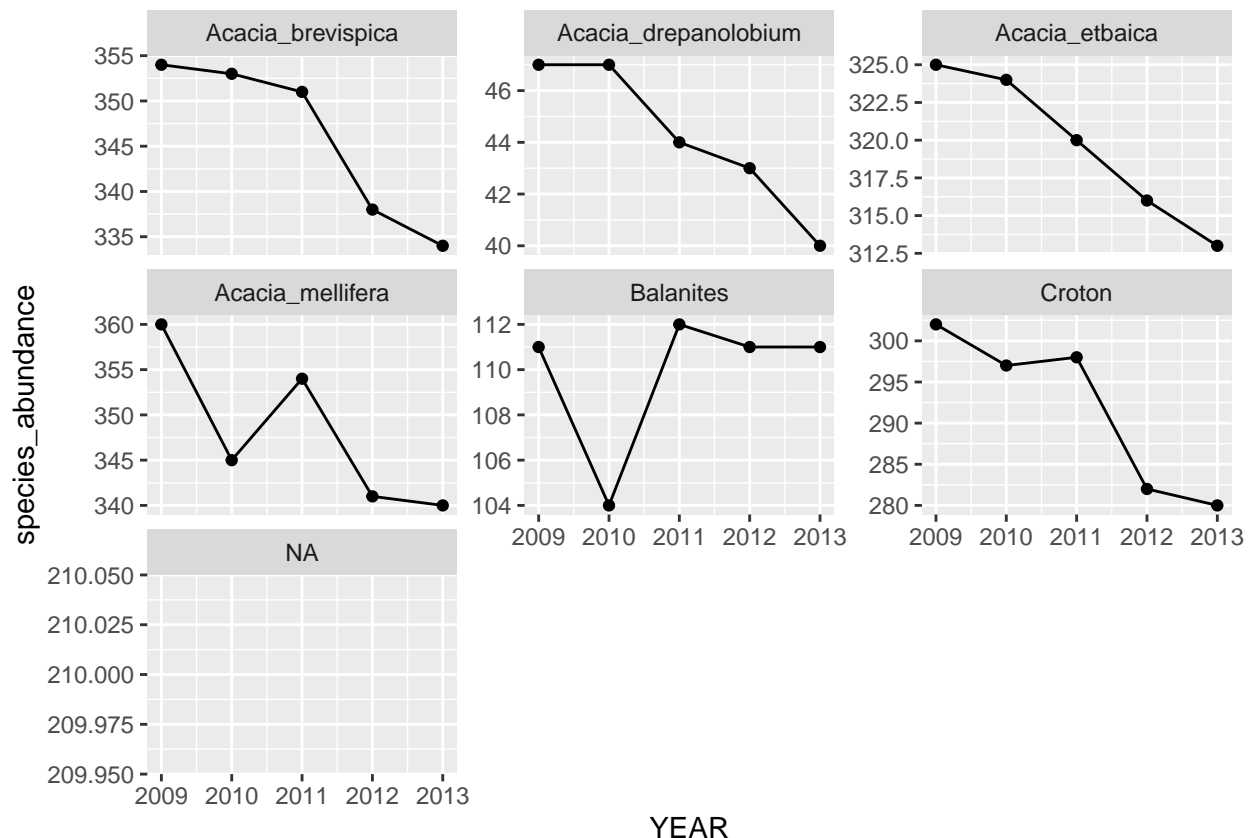
```
## 'summarise()' has grouped output by 'YEAR'. You can override using the '.groups'
## argument.
```

#8 Here I made a line plot with points by using the geom_line() and the geom_point function used the facet_wrap(~SPECIES, scales = "free_y") command to see each trend clearly


```
ggplot(data = abundance,
       mapping = aes(x = YEAR, y = species_abundance)) +
  geom_line()+
  geom_point()+
  facet_wrap(~SPECIES, scales = "free_y")
```

```
## Warning: Removed 1 row(s) containing missing values (geom_path).
```

```
## Warning: Removed 1 rows containing missing values (geom_point).
```



Exercise 4 Comparing the circumference to height relationship in acacia and to the same relationship for trees in the region. This compares two data sets in a scatter plot I used the `geom_points()` function to specify the information we need from each of both data sets first are the commands of the trees data set- what we want on the back goes first the `geom smooth` command gives the smooth line to each one of the layers in the scatter plot the `scale_x_log10()` and `scale_y_log10()` functions allow us to scale the data in the scatter plot we had the issue in office hrs to get the trend lines to be straight rather than curved. used the `labs()` functions to add labels to the scatter plot.

```
ggplot() +
  geom_point(data = trees,
            mapping = aes(x = CIRC , y = HEIGHT), color = "gray") +
  geom_smooth(data = trees,
            mapping = aes(x = CIRC , y = HEIGHT), color = "black") +
  scale_x_log10() +
  scale_y_log10()+
```

```
geom_point(data = acacia,
           mapping = aes(x = CIRC , y = HEIGHT ), color= "red") +
geom_smooth(data = acacia,
           mapping = aes(x = CIRC , y = HEIGHT ), color= "red") +
  scale_x_log10() +
  scale_y_log10()+
  labs(x = "Circumference (cm)", y = "Height (m)", title = "Data From Acacia VS Trees ")
```

```
## Scale for 'x' is already present. Adding another scale for 'x', which will
## replace the existing scale.
```

```
## Scale for 'y' is already present. Adding another scale for 'y', which will
## replace the existing scale.
```

```
## 'geom_smooth()' using method = 'gam' and formula 'y ~ s(x, bs = "cs")'
```

```
## Warning: Removed 416 rows containing non-finite values (stat_smooth).
```

```
## 'geom_smooth()' using method = 'loess' and formula 'y ~ x'
```

```
## Warning: Removed 4 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 416 rows containing missing values (geom_point).
```

```
## Warning: Removed 4 rows containing missing values (geom_point).
```

