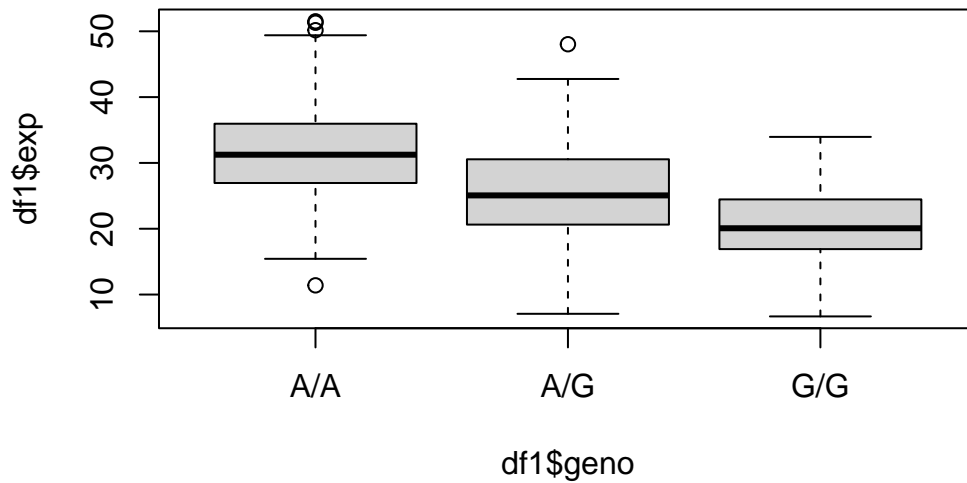


class12hw

sylvia ho a18482382

q13 smaple siz: a/a: 108, a/g: 233, g/g/:121 median: A/A 31.24847
A/G 25.06486
G/G 20.07363

```
df1<-read.table("df.txt")  
bx1<-boxplot(df1$exp ~ df1$geno)
```



```
table(df1$geno)
```

```
A/A A/G G/G  
108 233 121
```

```
library(dplyr)
```

Attaching package: 'dplyr'

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

filter, lag

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

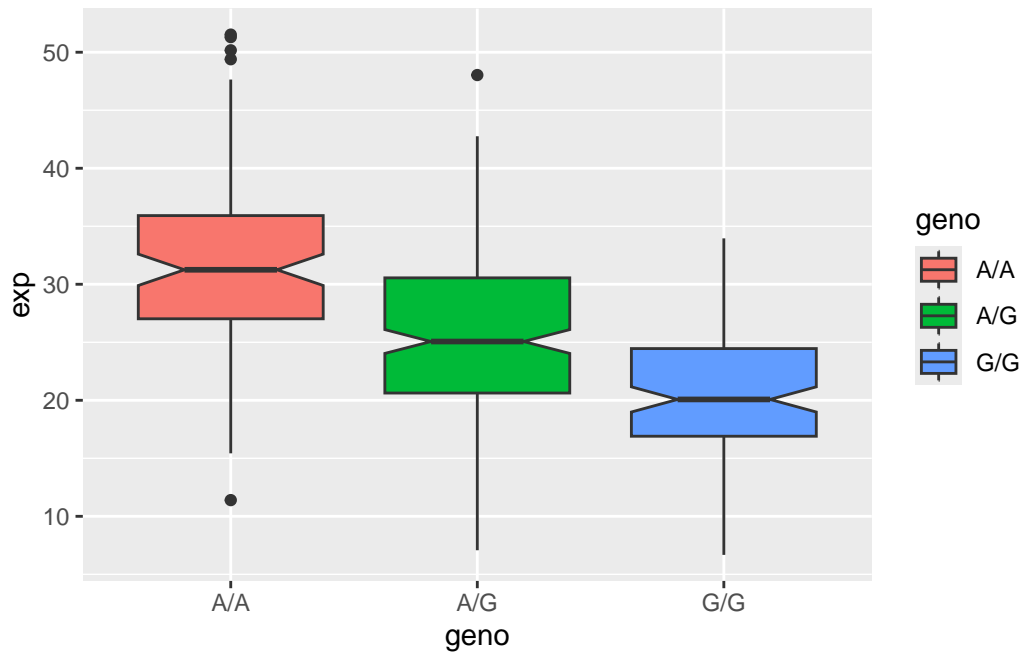
intersect, setdiff, setequal, union

```
df1 %>%  
  group_by(geno) %>%  
  summarize(median_value = median(exp, na.rm = TRUE))
```

```
# A tibble: 3 x 2  
  geno median_value  
  <chr>         <dbl>  
1 A/A          31.2  
2 A/G          25.1  
3 G/G          20.1
```

q14

```
library(ggplot2)  
  
ggplot(df1)+aes(geno,exp,fill=geno)+  
  geom_boxplot(notch=TRUE)
```



df1

	sample	geno	exp
1	HG00367	A/G	28.96038
2	NA20768	A/G	20.24449
3	HG00361	A/A	31.32628
4	HG00135	A/A	34.11169
5	NA18870	G/G	18.25141
6	NA11993	A/A	32.89721
7	HG00256	A/G	31.48736
8	NA18498	A/A	47.64556
9	HG00327	G/G	17.67473
10	HG00115	A/G	33.85374
11	NA20806	A/G	16.29854
12	HG00278	A/G	19.73450
13	NA20585	A/A	30.71355
14	NA19137	A/G	13.96175
15	HG00235	A/A	25.44983
16	NA20798	A/A	34.24915
17	NA12546	G/G	18.55622
18	NA19116	A/A	35.15014
19	HG00381	A/G	18.40351
20	NA18488	G/G	23.10383

21	HG00259	A/G	34.21985
22	HG00177	A/G	23.32404
23	NA19214	G/G	30.94554
24	NA19247	A/A	24.54684
25	NA19098	A/G	23.18606
26	NA20589	A/G	18.15997
27	NA19207	A/A	49.39612
28	HG00112	G/G	21.14387
29	NA20518	G/G	18.39547
30	HG00335	A/A	28.20755
31	NA19119	G/G	12.02809
32	HG00247	G/G	17.44761
33	NA12155	A/G	28.03580
34	NA20771	A/G	30.65270
35	NA20758	G/G	29.82254
36	HG00121	A/G	20.51327
37	NA20759	A/A	28.56199
38	NA20816	A/G	29.72309
39	NA20542	A/G	22.50789
40	NA18511	A/G	31.68959
41	NA12249	G/G	23.01983
42	NA11830	A/G	28.76435
43	NA19159	A/G	35.85543
44	NA20778	A/G	37.62403
45	NA18908	A/G	20.54885
46	HG00320	G/G	13.42470
47	NA11843	G/G	22.65437
48	HG00105	A/A	51.51787
49	NA20588	G/G	11.07445
50	NA20510	G/G	28.35841
51	NA12342	A/G	31.04941
52	HG00249	A/G	18.94583
53	NA11894	A/A	38.10956
54	HG00240	A/G	32.29483
55	HG00132	A/A	31.13741
56	HG00118	G/G	28.79371
57	NA18520	G/G	27.08956
58	NA18508	A/G	27.81775
59	HG00353	A/G	19.89903
60	NA20792	A/G	48.03410
61	NA12234	G/G	16.11138
62	HG00377	A/A	39.12999
63	NA19143	A/G	27.90313

64	NA20787	A/G	36.47949
65	NA20513	A/G	20.03116
66	HG00243	A/G	29.65063
67	NA19172	A/A	32.44173
68	NA06994	A/G	34.92257
69	NA18510	A/G	16.71385
70	HG00337	A/G	16.68151
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72	NA19152	G/G	26.61928
73	NA20761	G/G	30.18323
74	NA19235	A/G	11.60808
75	HG00382	A/G	19.30953
76	NA20544	A/A	34.03260
77	NA18923	G/G	19.40790
78	HG00313	A/G	20.49040
79	HG00238	G/G	19.52301
80	NA20754	A/G	22.37224
81	NA11918	A/G	15.20045
82	NA18868	A/A	36.27151
83	NA06986	A/G	20.07459
84	HG00263	A/G	35.42982
85	NA12058	G/G	26.56808
86	NA20507	A/G	19.10884
87	NA12777	A/G	24.81087
88	NA12144	A/G	33.22193
89	HG00129	G/G	17.34076
90	HG00123	A/G	33.40835
91	NA12814	A/G	22.38996
92	HG00183	G/G	10.74263
93	HG00109	G/G	16.66051
94	NA20505	A/G	31.31626
95	NA12273	A/G	9.36055
96	HG00174	A/A	26.10355
97	HG00324	A/A	19.48106
98	HG00365	A/G	23.17937
99	NA20520	A/A	38.77623
100	NA19189	A/G	30.63079
101	HG00155	A/G	19.10420
102	HG00111	A/A	40.82922
103	NA12827	A/G	25.70962
104	NA18517	G/G	29.01720
105	NA20801	G/G	20.69333
106	NA20529	G/G	21.15677

107	NA18909	A/G	38.34531
108	HG00173	A/G	19.03976
109	HG00349	G/G	18.58691
110	HG00234	G/G	19.04962
111	NA19248	G/G	22.81974
112	NA20810	A/A	46.50527
113	HG00255	A/G	28.81770
114	NA12813	G/G	32.01142
115	NA20537	G/G	21.12823
116	NA18912	A/G	42.75662
117	HG00332	G/G	18.61268
118	HG00152	G/G	19.37093
119	NA20783	G/G	31.42162
120	NA12154	A/G	25.61662
121	HG00236	A/A	33.07320
122	NA19146	A/A	25.47283
123	HG00312	A/G	26.48467
124	HG00148	A/G	28.02486
125	HG00364	A/G	24.23377
126	HG00311	A/G	21.03717
127	NA11881	A/A	29.50655
128	HG00185	G/G	16.67764
129	NA20807	A/G	33.51752
130	NA19184	A/G	20.73493
131	HG00133	A/G	33.55650
132	NA20531	G/G	19.08659
133	NA19138	A/A	27.48438
134	NA19206	A/G	36.62034
135	HG00277	G/G	21.55001
136	NA18858	A/G	40.06318
137	HG00375	A/G	33.92744
138	HG00127	A/G	21.02084
139	NA19099	A/G	29.95687
140	HG00336	G/G	8.29591
141	HG00097	A/G	25.80393
142	HG00267	A/G	21.49924
143	NA20581	G/G	12.58869
144	NA12286	A/G	34.79575
145	NA20797	A/G	34.57705
146	NA12872	A/G	30.03549
147	HG00360	A/G	16.59638
148	NA20530	A/G	27.22300
149	NA12348	A/G	24.35621

150	NA20538	G/G	17.34109
151	NA12760	A/G	22.86793
152	NA12763	A/G	23.19511
153	NA20814	G/G	28.23642
154	NA19222	A/A	35.69719
155	NA06989	A/A	32.42236
156	NA19171	G/G	19.99979
157	NA11829	A/G	33.74015
158	NA11992	A/G	24.08401
159	HG00141	G/G	25.55413
160	NA19150	A/G	26.39419
161	NA20828	A/G	32.33359
162	NA12749	A/A	28.91526
163	NA19190	G/G	24.45672
164	NA06985	A/G	11.36287
165	HG00178	A/G	21.16515
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171	NA12272	G/G	14.66862
172	NA19096	G/G	33.95602
173	NA20800	A/G	22.73049
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175	NA19236	G/G	18.26466
176	HG00264	A/G	25.57669
177	NA20521	A/A	27.87464
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180	HG00329	A/A	16.86780
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188	NA20804	A/A	36.51922
189	NA11994	A/G	30.83577
190	HG00156	G/G	17.32504
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192	HG00180	A/G	19.66773

193	HG00282	G/G	19.14766
194	HG00343	G/G	12.57599
195	HG00139	G/G	22.28749
196	HG01789	A/G	24.64870
197	HG00321	A/G	17.03159
198	HG00306	A/A	27.43637
199	HG00232	G/G	17.29261
200	NA20528	A/G	22.27101
201	HG00122	G/G	24.18141
202	NA07037	A/A	35.63983
203	NA07056	A/G	15.92557
204	HG00151	A/G	32.54150
205	NA19129	A/A	38.85161
206	NA20517	A/G	22.40203
207	NA19149	G/G	16.07627
208	HG00341	A/G	27.41638
209	HG00274	A/G	31.99645
210	HG00106	A/G	30.05415
211	HG00189	G/G	14.80495
212	HG00252	A/G	20.01602
213	NA11832	A/G	34.47373
214	HG00323	A/A	22.44576
215	NA18916	A/A	37.06379
216	NA18867	A/G	28.75978
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218	HG00126	G/G	23.46573
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220	NA20504	A/G	15.71646
221	NA20532	A/G	21.76610
222	NA12812	A/G	9.62656
223	HG00244	A/G	28.53965
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225	HG00378	G/G	27.78837
226	NA20790	A/A	50.16704
227	NA20512	A/A	37.94544
228	HG00268	A/A	29.15536
229	HG00380	A/A	28.85309
230	NA12761	A/A	38.57101
231	HG00384	A/G	29.49417
232	NA20796	G/G	23.92355
233	NA12399	G/G	9.55902
234	HG00310	A/G	29.55520
235	HG00096	A/A	30.89365

236	NA19147	A/G	19.44178
237	NA20752	A/G	21.43751
238	NA19107	A/G	30.40382
239	HG00099	G/G	12.35836
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241	NA19114	G/G	22.53910
242	HG00376	A/A	31.43743
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244	HG00130	A/G	28.50982
245	HG00158	A/A	22.37043
246	HG00269	A/A	28.46943
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248	HG00258	A/A	30.15636
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250	HG00276	G/G	16.40569
251	HG00331	A/G	31.10134
252	NA12751	A/G	35.99067
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255	NA11920	A/G	26.42877
256	HG00326	A/G	26.28329
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258	NA12716	A/G	20.72639
259	HG00142	G/G	19.42882
260	HG00309	A/G	21.09140
261	HG00315	G/G	26.56993
262	HG00338	A/G	23.79292
263	NA11995	A/A	32.59723
264	NA19209	A/A	36.02549
265	NA20540	A/A	23.86454
266	NA12890	A/A	28.38114
267	HG00250	G/G	13.34557
268	NA20769	G/G	16.60507
269	HG00138	A/A	25.14243
270	NA19200	A/A	51.30170
271	NA19144	G/G	24.85165
272	NA12815	G/G	21.56943
273	NA12043	A/G	18.79569
274	HG00350	A/G	29.54042
275	NA12383	A/A	28.14811
276	NA19201	A/G	18.78700
277	HG00187	A/G	21.41071
278	NA06984	A/A	29.18390

279	NA20508	A/G	21.29782
280	NA19175	G/G	23.95528
281	NA20815	A/G	33.91853
282	NA12044	A/G	27.20808
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287	HG00260	G/G	26.04123
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289	NA07347	A/G	37.73840
290	NA07357	A/A	27.09760
291	NA20543	A/G	34.14567
292	HG00261	G/G	20.07363
293	HG00273	G/G	19.76527
294	NA12341	A/G	15.36874
295	HG00245	A/G	29.50350
296	NA19198	A/G	25.70400
297	NA20757	A/G	20.07219
298	NA11930	A/A	33.89656
299	HG00358	G/G	18.50772
300	NA18933	A/G	24.53928
301	HG00242	A/G	17.84487
302	NA20773	A/G	23.35766
303	NA12282	A/G	15.71243
304	NA19131	A/A	33.48253
305	NA18499	A/A	15.43178
306	HG00117	A/A	29.45277
307	NA19121	G/G	20.14146
308	NA20515	G/G	18.07151
309	HG00355	A/G	19.89034
310	NA12775	A/G	25.37234
311	NA12005	A/G	16.12745
312	NA11893	A/G	24.18529
313	NA20808	A/G	21.97051
314	NA10847	G/G	6.94390
315	NA19102	A/G	13.08172
316	NA12400	G/G	22.14277
317	NA18487	A/G	32.00764
318	NA19093	A/G	30.59653
319	HG00342	G/G	14.23742
320	NA19160	A/G	29.74443
321	NA19095	A/G	27.88354

322	HG00160	A/A	26.80283
323	NA20766	A/G	11.12451
324	NA12717	A/G	7.07505
325	HG00125	A/G	23.13726
326	HG00171	A/G	21.09331
327	NA12873	A/G	8.20002
328	NA20525	A/G	20.62572
329	NA20826	A/G	18.24345
330	HG00136	G/G	19.85388
331	HG00272	A/G	11.13478
332	NA12340	A/A	43.51943
333	HG00251	A/G	24.43943
334	HG00369	A/G	22.24289
335	NA20803	A/G	24.67325
336	NA12842	A/G	41.03924
337	HG00146	A/A	45.80808
338	HG01790	A/G	33.31795
339	NA20809	A/G	27.98844
340	NA20765	G/G	27.73467
341	HG00362	A/A	26.55972
342	HG00114	A/G	31.57994
343	NA18917	A/A	24.87330
344	NA18502	G/G	19.02064
345	HG00150	A/G	36.73337
346	NA20527	A/A	29.99549
347	HG00179	A/G	18.45322
348	NA20805	A/A	26.68589
349	NA19117	A/G	23.60431
350	HG00285	A/G	24.33489
351	NA20772	G/G	14.49816
352	NA19213	A/G	35.74662
353	HG00344	A/G	22.75684
354	NA12156	A/A	39.37193
355	HG00257	G/G	26.78940
356	NA18486	G/G	20.84709
357	HG00188	G/G	10.77316
358	HG00366	A/G	34.42403
359	HG00157	A/A	38.39523
360	HG00262	A/A	41.23635
361	HG00280	G/G	12.82128
362	HG00308	G/G	16.90256
363	NA11831	A/G	25.34866
364	NA18910	G/G	29.60045

365	NA20795	A/G	25.06486
366	HG00231	A/G	36.78028
367	NA19197	A/G	30.67131
368	HG00101	A/A	27.13936
369	HG00281	G/G	14.81945
370	NA20760	A/A	36.55643
371	HG00176	A/A	28.34688
372	NA18489	A/G	37.82860
373	NA12275	G/G	17.46326
374	NA20514	A/A	15.42908
375	HG00351	G/G	23.26922
376	HG00186	G/G	21.39806
377	NA20586	A/G	25.44086
378	HG00275	G/G	18.06320
379	HG00325	G/G	15.91528
380	NA19118	G/G	24.80823
381	HG00124	G/G	26.04514
382	NA20785	A/A	47.50579
383	HG02215	G/G	18.28089
384	HG00253	A/A	30.15754
385	HG00134	G/G	23.24907
386	HG00339	A/A	34.88439
387	NA20519	A/G	29.49548
388	NA12778	A/G	23.27255
389	NA18861	A/A	29.29955
390	NA20539	A/A	32.87767
391	NA11931	G/G	17.91118
392	NA20812	A/G	28.69506
393	HG00120	G/G	21.09502
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395	HG00328	A/G	27.49975
396	NA20774	A/G	24.66196
397	NA18873	A/G	25.81562
398	NA20502	A/G	22.49429
399	HG00143	A/G	26.88264
400	HG00145	A/A	43.43665
401	NA19225	A/A	26.56050
402	NA12829	A/G	28.98200
403	HG00137	A/G	34.31875
404	NA20524	A/G	26.40231
405	HG00379	A/A	21.87746
406	NA18505	A/G	21.67621
407	HG01334	A/G	27.56805

408	NA18907	A/A	33.42582
409	NA19204	A/A	25.38406
410	NA12874	A/G	16.16277
411	NA20506	A/G	18.28963
412	NA20770	A/A	18.20442
413	NA12776	A/G	30.55183
414	NA18934	A/G	20.70871
415	NA19153	A/G	17.66476
416	HG00356	A/G	22.79543
417	NA12283	A/G	24.03419
418	HG00284	A/G	18.02351
419	NA12489	A/G	21.63102
420	HG00104	A/A	21.62336
421	NA20582	G/G	24.74366
422	NA11840	A/G	27.54976
423	HG00383	A/G	14.79717
424	NA20786	A/A	35.80093
425	NA20802	A/G	25.34921
426	NA20756	A/A	32.26844
427	NA19113	A/G	21.34916
428	NA12889	G/G	27.40521
429	NA12718	A/G	21.20080
430	HG00266	A/G	28.36006
431	NA12287	A/G	22.43773
432	HG00319	A/G	25.56306
433	NA12762	A/A	34.40756
434	HG00334	A/G	19.50634
435	NA12006	G/G	24.85772
436	NA19108	G/G	23.08482
437	NA19185	A/G	28.93651
438	HG00246	A/G	31.79897
439	NA12045	A/G	30.80067
440	NA19257	A/G	33.95134
441	NA12413	A/G	39.43243
442	HG00159	A/A	23.99631
443	NA20811	A/A	11.39643
444	HG00149	A/G	23.91465
445	NA19223	A/G	20.97560
446	NA07346	G/G	16.56929
447	NA20536	A/G	20.02507
448	HG01791	A/A	35.24632
449	HG00271	A/G	33.44170
450	HG00373	A/G	17.32813

451	HG00182	A/A	23.38376
452	HG00110	A/G	32.61856
453	NA20819	A/G	36.77906
454	HG00154	G/G	16.69044
455	HG00330	A/G	16.84776
456	NA12750	A/A	34.94395
457	HG00233	G/G	25.08880
458	HG00131	G/G	32.78519
459	HG00108	A/A	31.92036
460	HG00119	A/G	31.53069
461	NA19130	A/A	44.27738
462	HG00239	A/G	23.18250