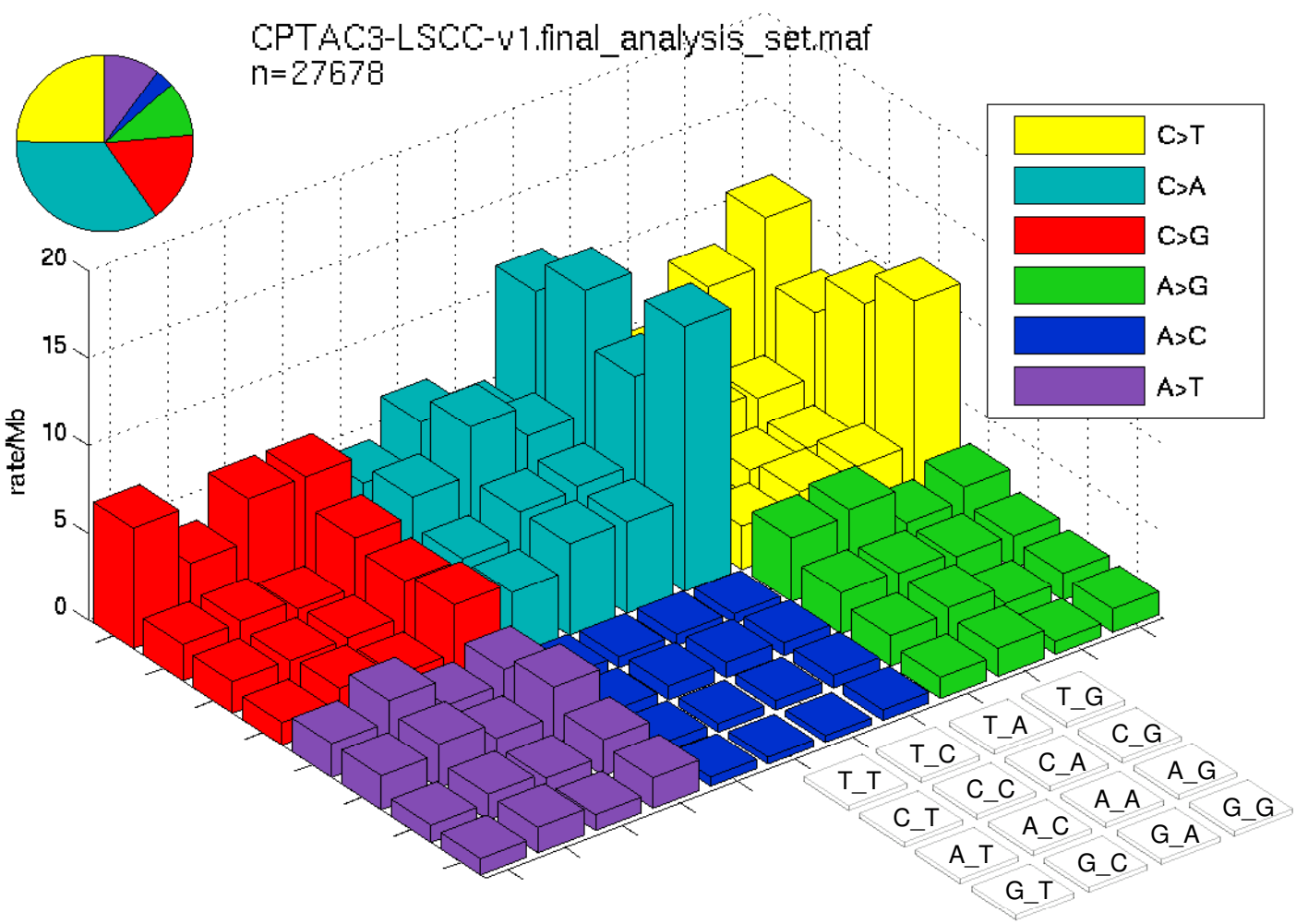
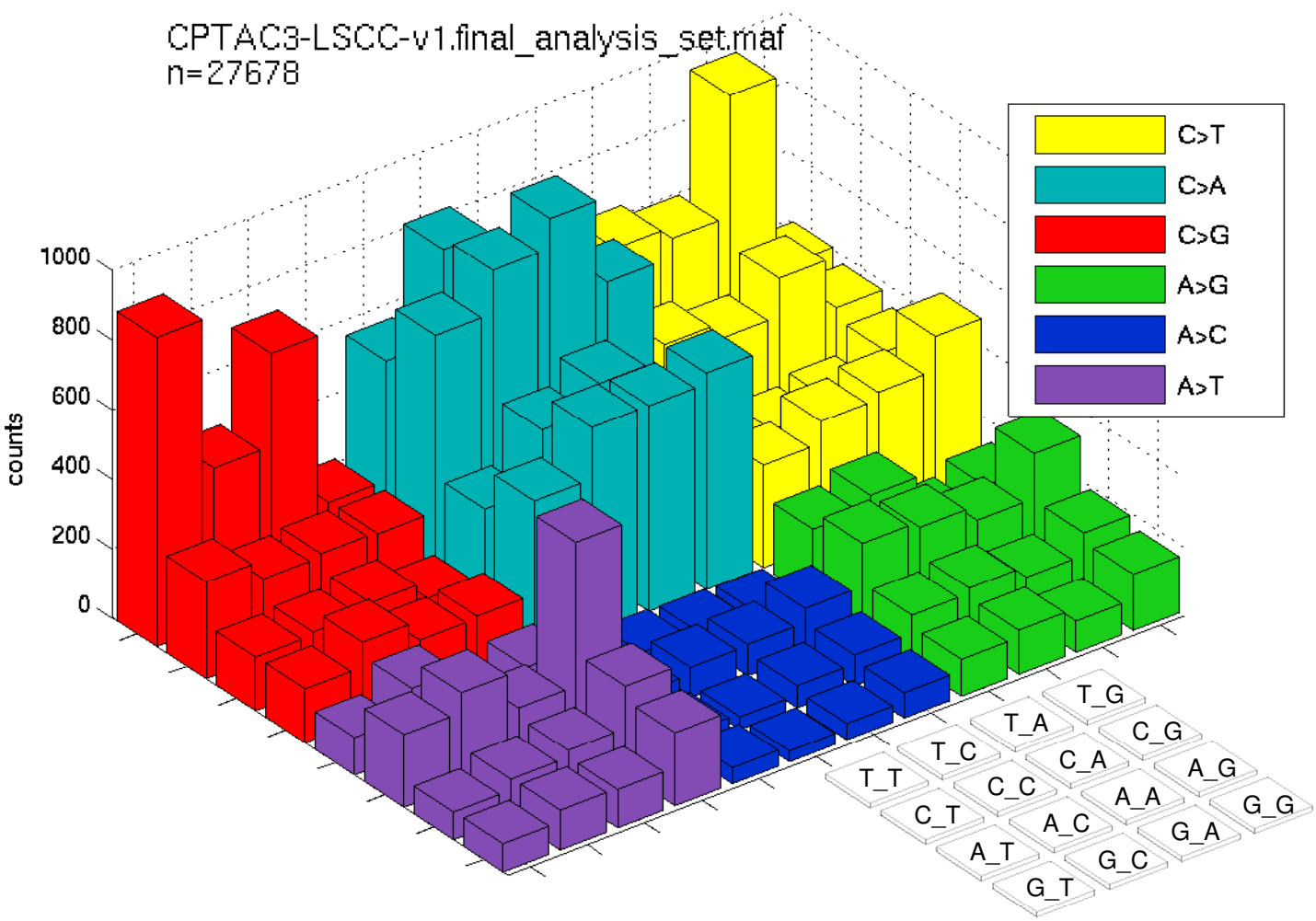
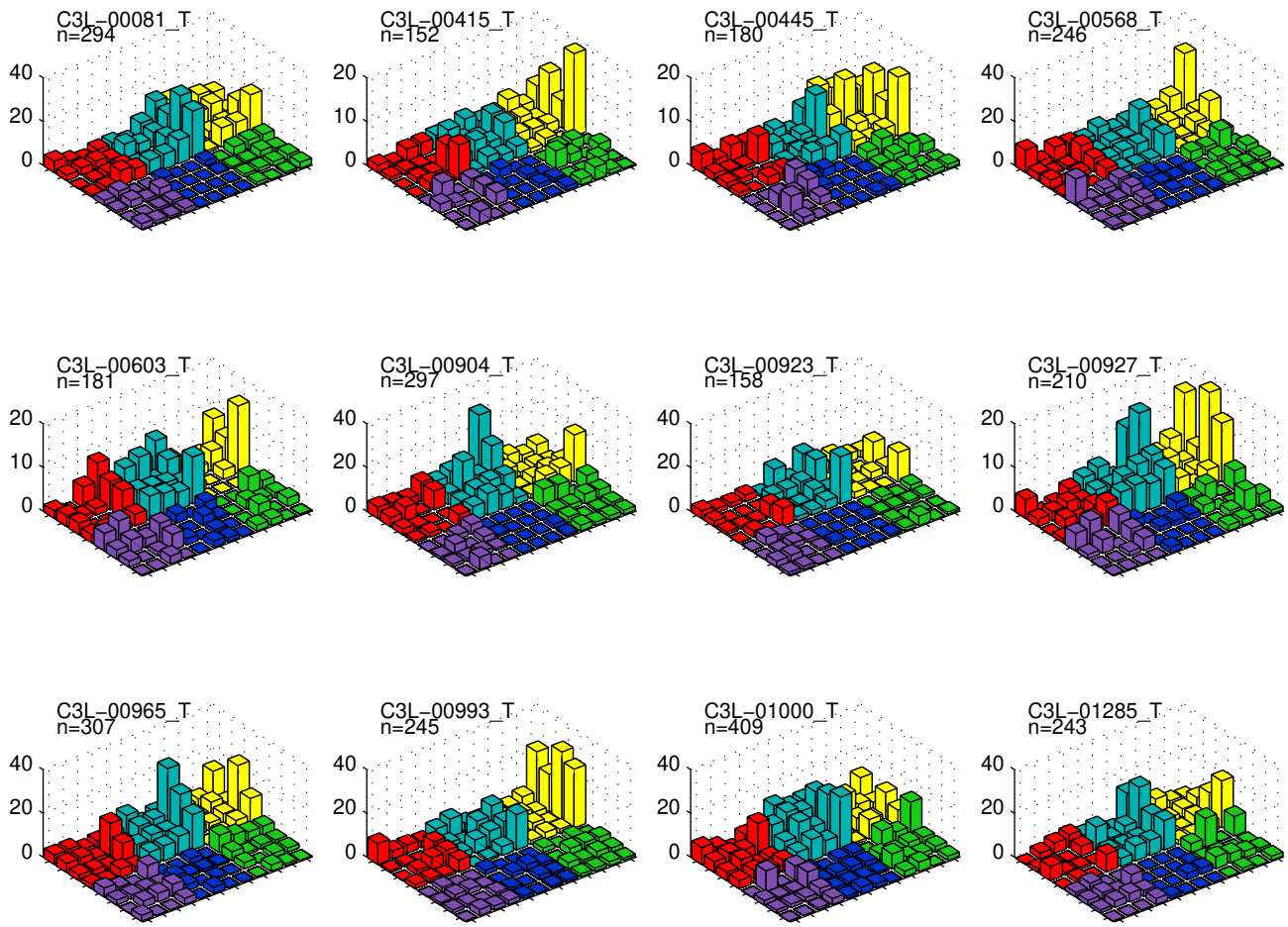
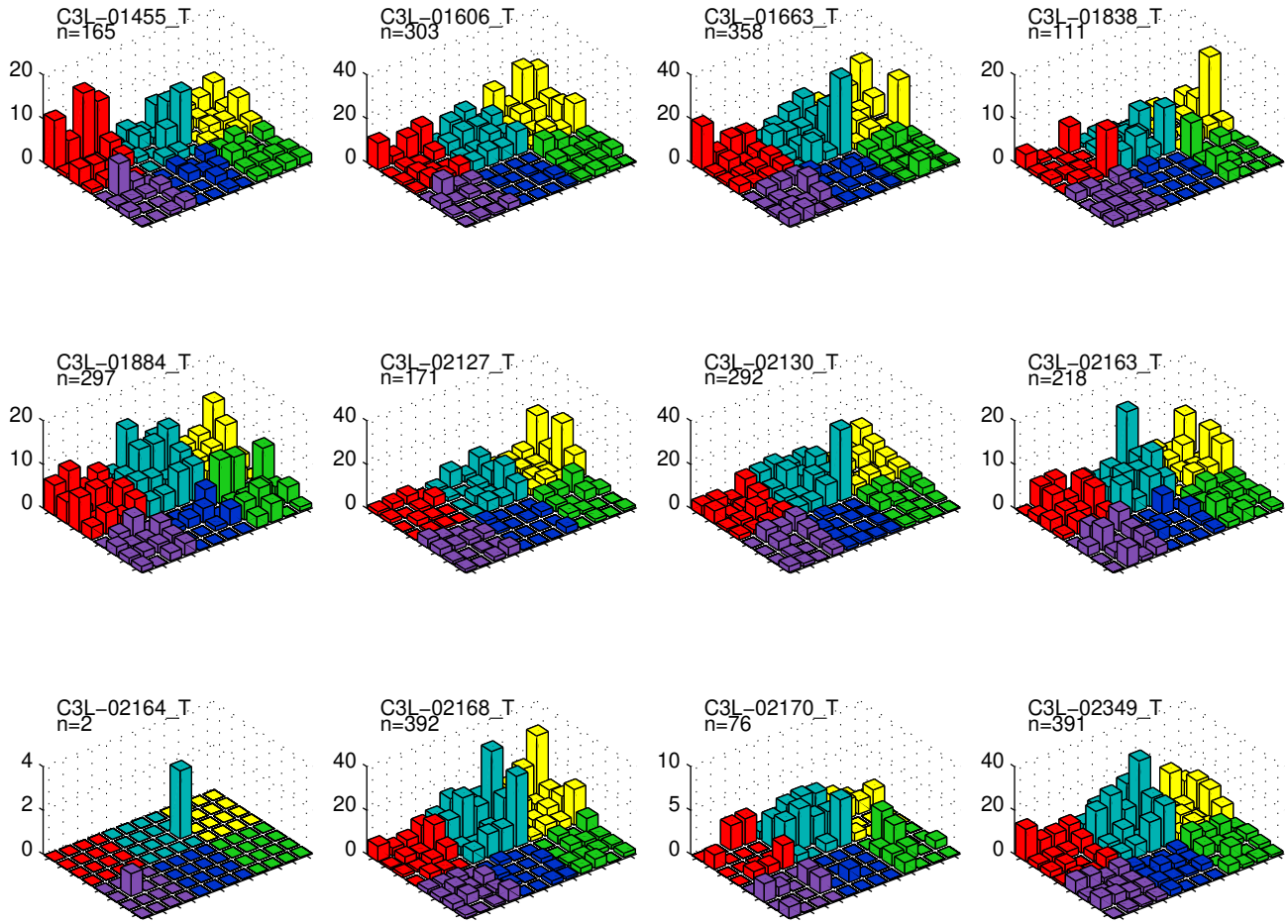
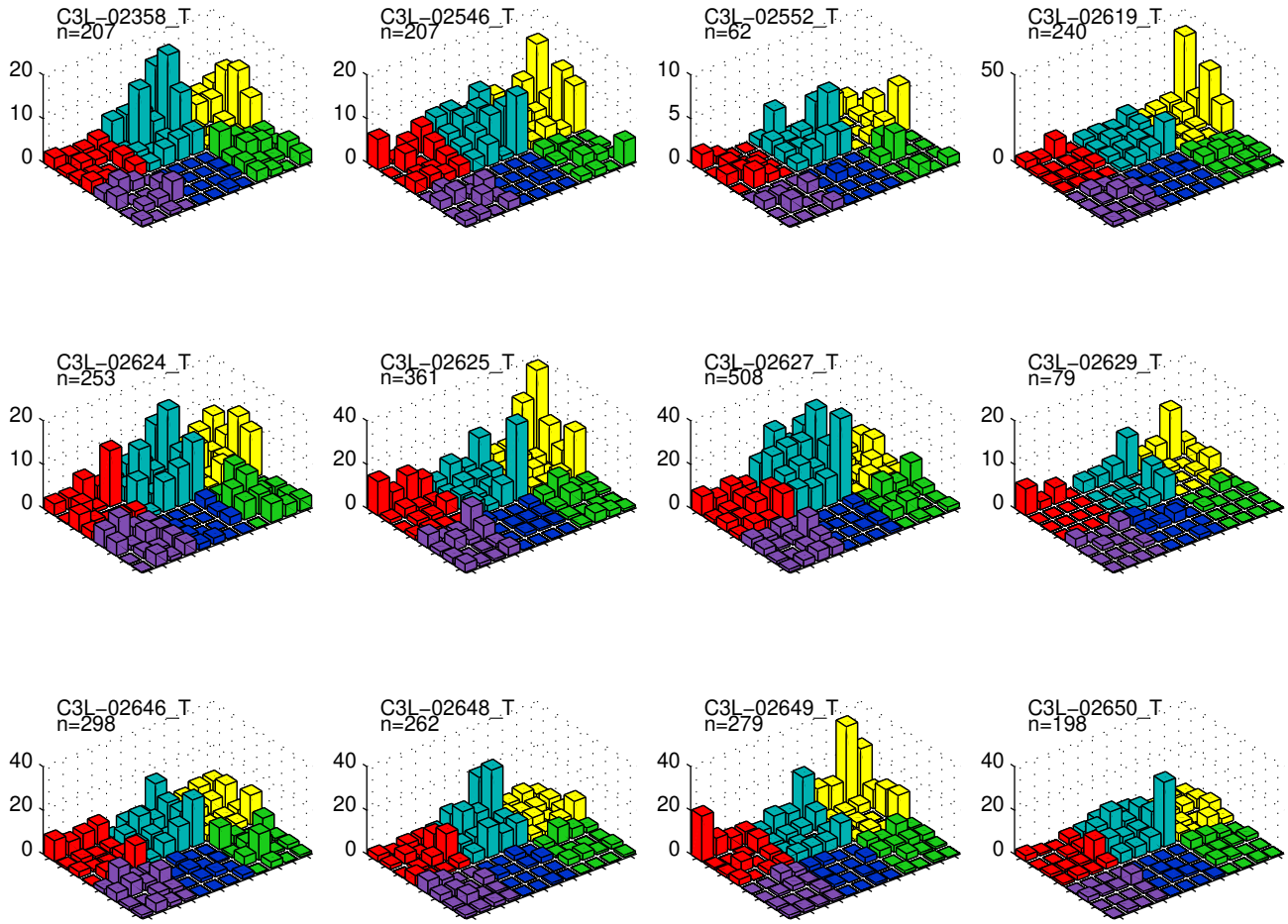
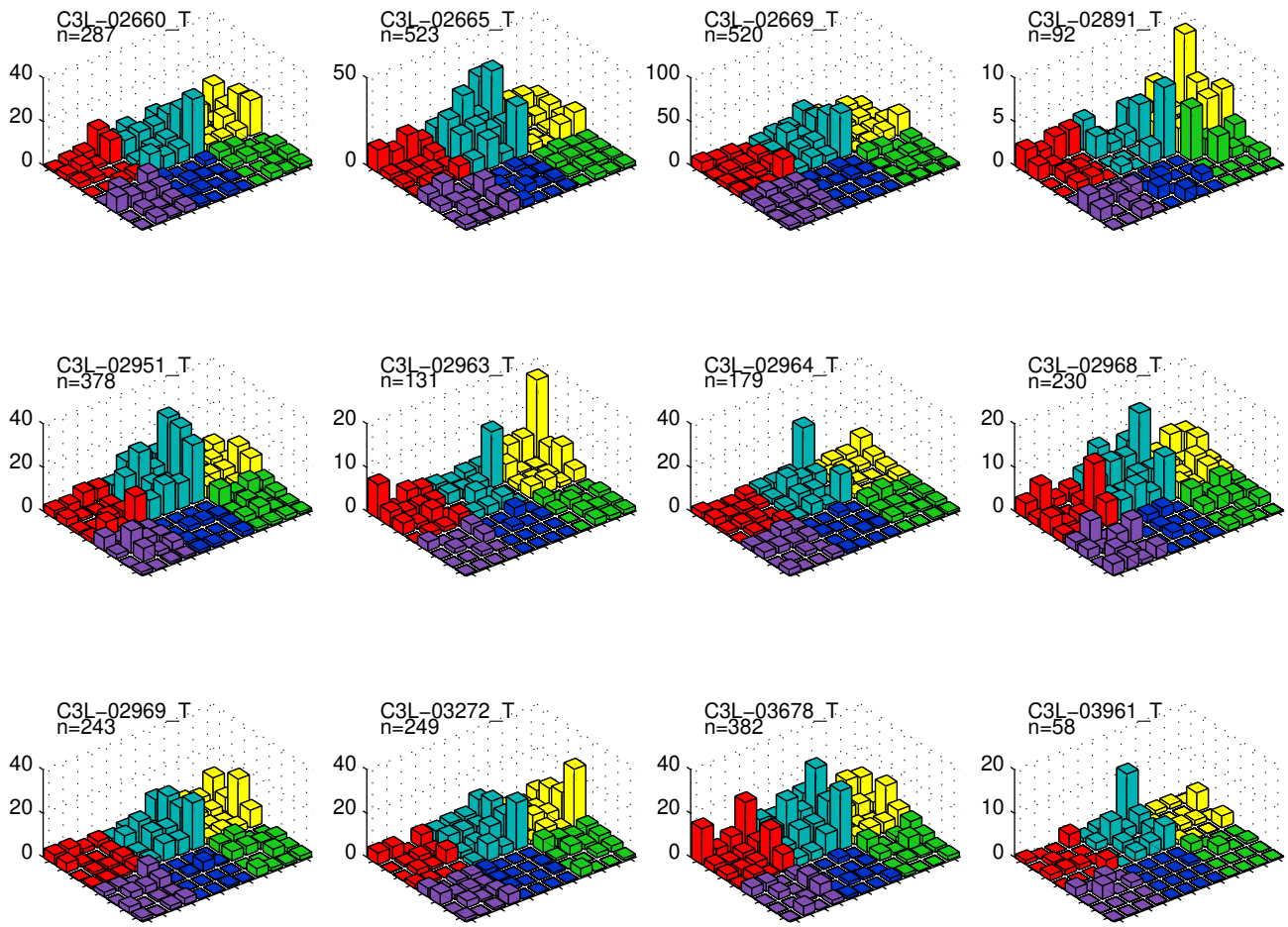

CPTAC3-LSCC-v1.final_analysis_set.maf
n=27678
rate/Mb
20
15
10
5
0
C>T
C>A
C>G
A>G
A>C
A>T
T_G
T_A
C_G
T_C
C_A
A_G
T_T
C_C
A_A
G_G
C_T
A_C
G_A
A_T
G_C
G_T

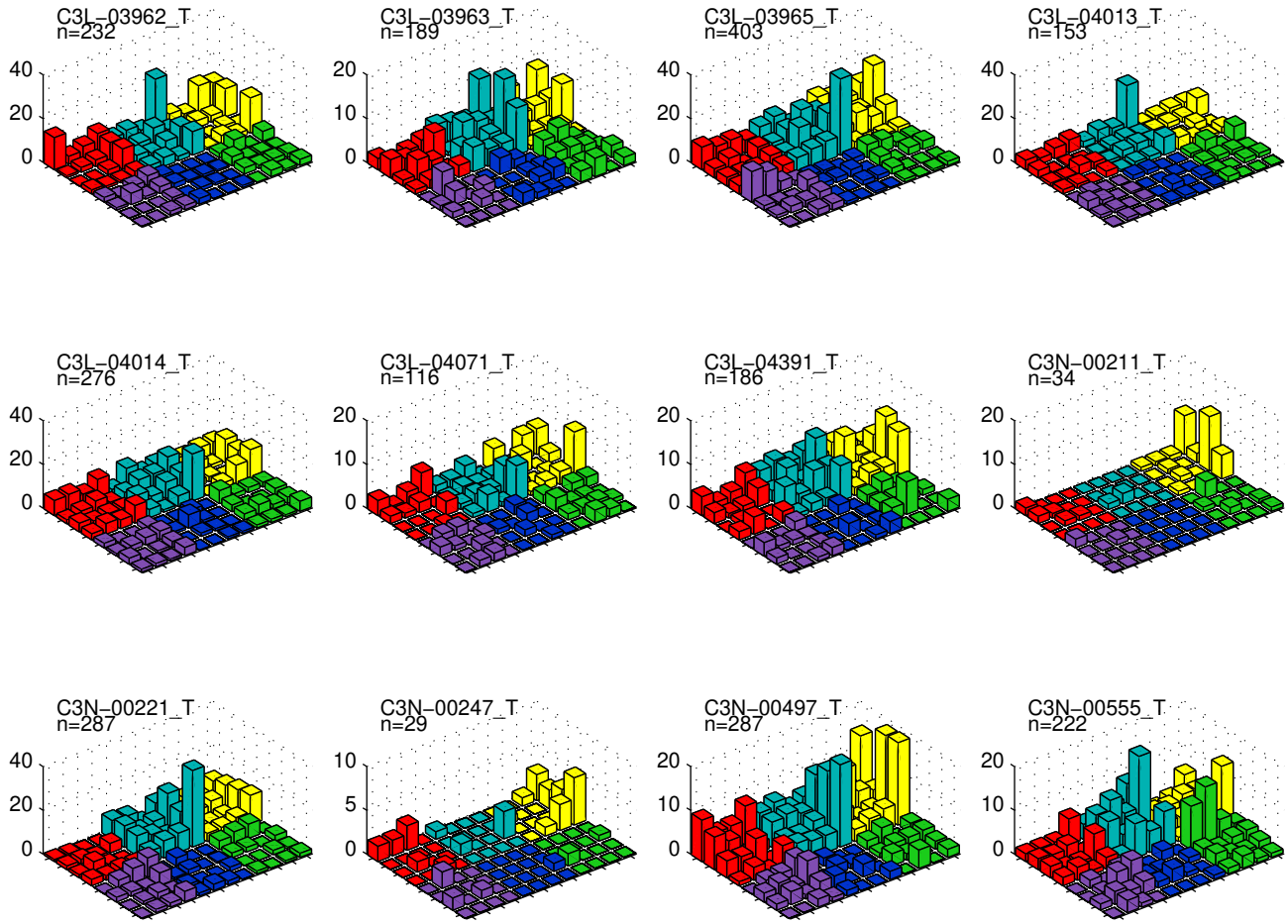

CPTAC3-LSCC-v1.final_analysis_set.maf
n=27678
counts
1000
800
600
400
200
0
C>T
C>A
C>G
A>G
A>C
A>T
T_G
T_A
C_G
T_C
C_A
A_G
T_T
C_C
A_A
G_G
C_T
A_C
G_A
A_T
G_C
G_T

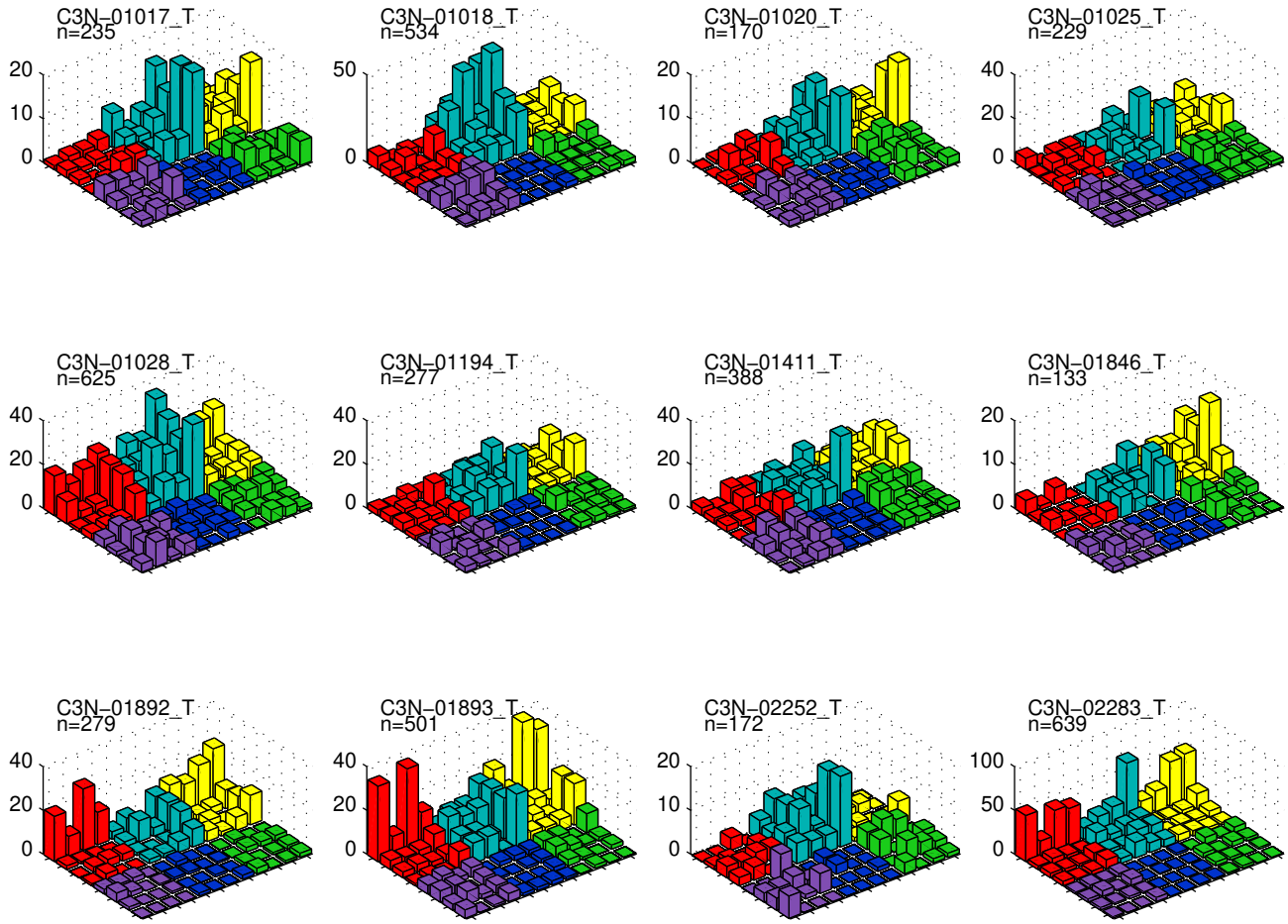

C3L−00081_T
n=294
C3L−00415_T
n=152
C3L−00445_T
n=180
C3L−00568_T
n=246
C3L−00603_T
n=181
C3L−00904_T
n=297
C3L−00923_T
n=158
C3L−00927_T
n=210
C3L−00965_T
n=307
C3L−00993_T
n=245
C3L−01000_T
n=409
C3L−01285_T
n=243

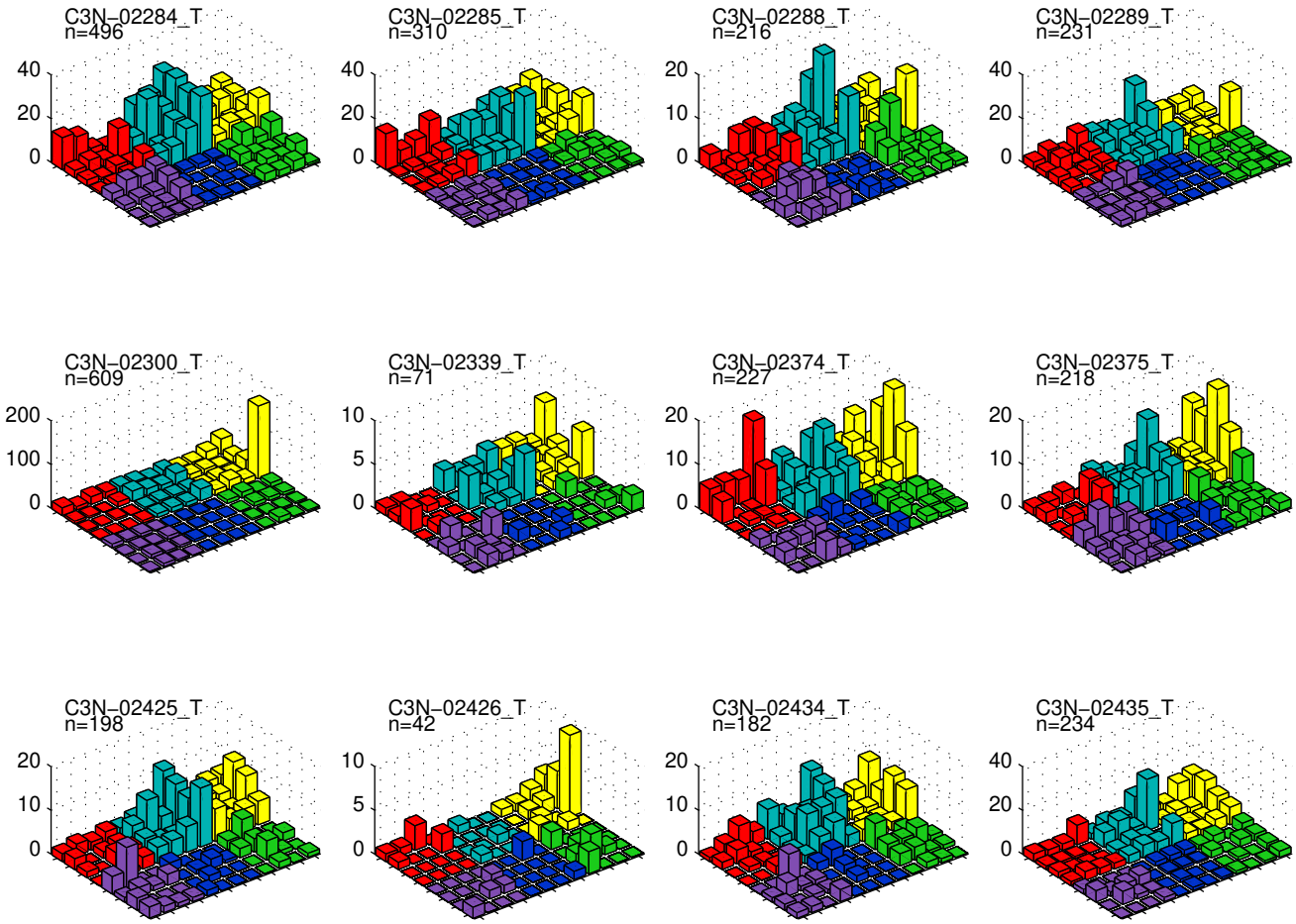

C3L−01455_T
n=165
20
10
0
C3L−01606_T
n=303
40
20
0
C3L−01663_T
n=358
40
20
0
C3L−01838_T
n=111
20
10
0
C3L−01884_T
n=297
20
10
0
C3L−02127_T
n=171
40
20
0
C3L−02130_T
n=292
40
20
0
C3L−02163_T
n=218
20
10
0
C3L−02164_T
n=2
4
2
0
C3L−02168_T
n=392
40
20
0
C3L−02170_T
n=76
10
5
0
C3L−02349_T
n=391
40
20
0

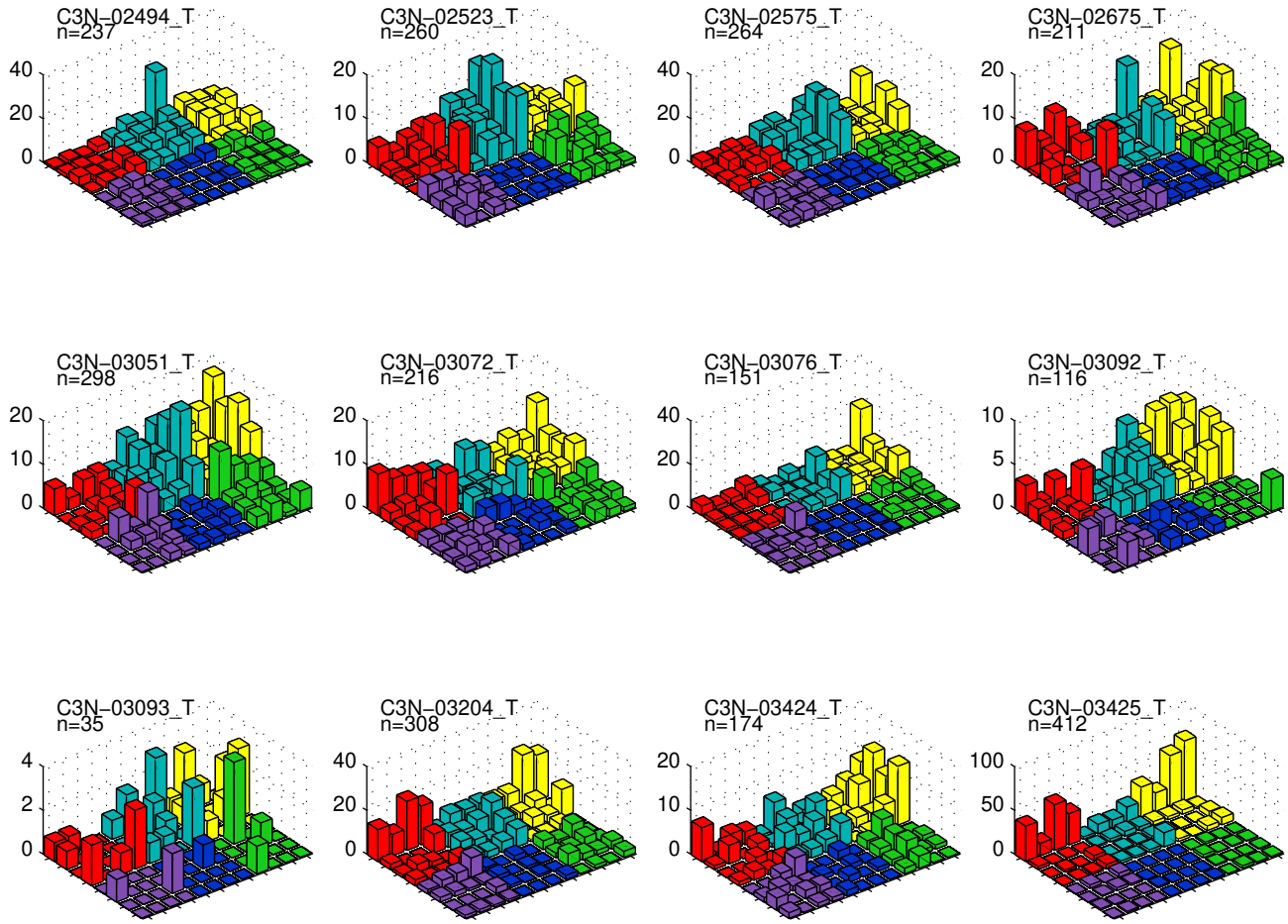

C3L−02358_T
n=207
20
10
0
C3L−02546_T
n=207
20
10
0
C3L−02552_T
n=62
10
5
0
C3L−02619_T
n=240
50
0
C3L−02624_T
n=253
20
10
0
C3L−02625_T
n=361
40
20
0
C3L−02627_T
n=508
40
20
0
C3L−02629_T
n=79
20
10
0
C3L−02646_T
n=298
40
20
0
C3L−02648_T
n=262
40
20
0
C3L−02649_T
n=279
40
20
0
C3L−02650_T
n=198
40
20
0

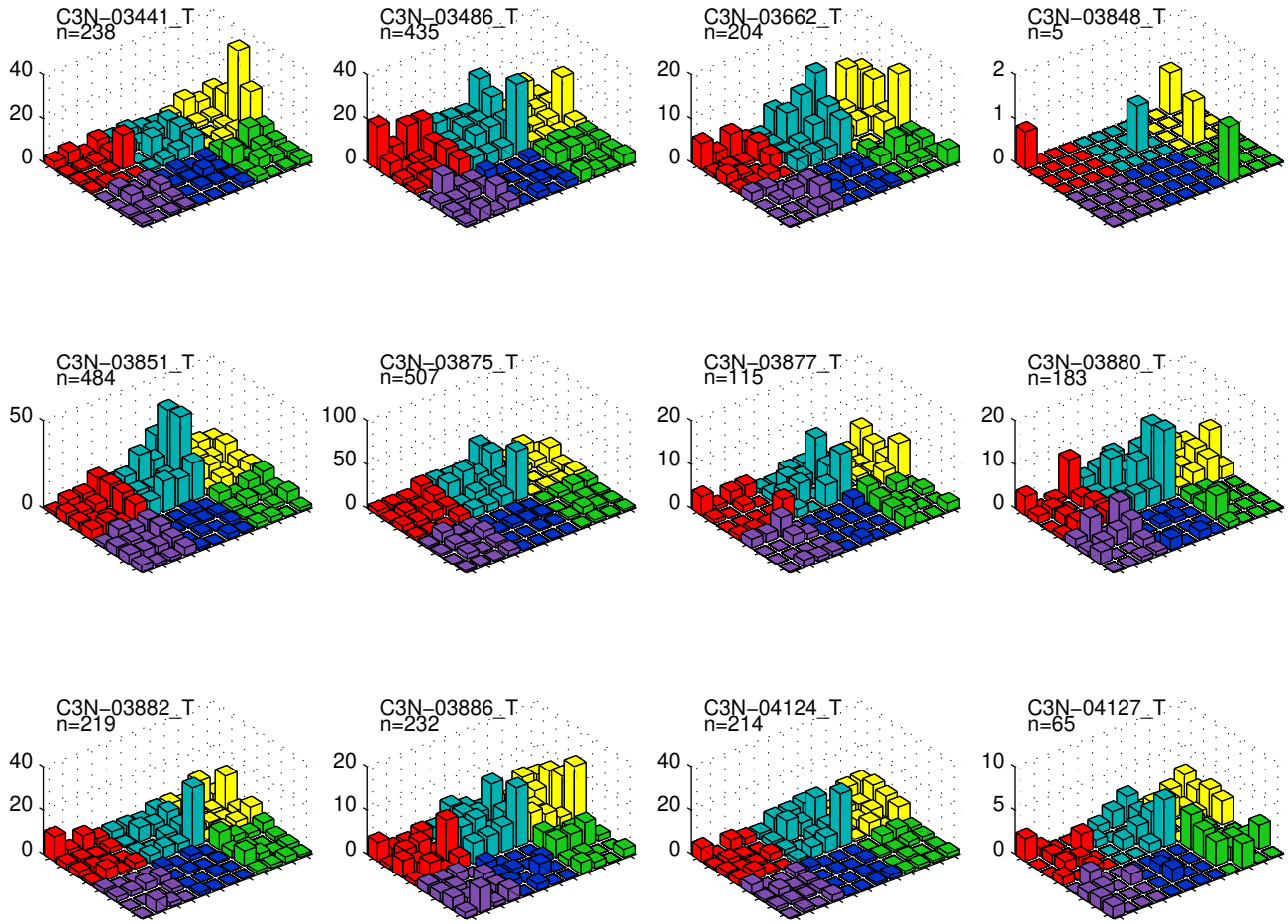

C3L−02660_T
n=287
40
20
0
C3L−02665_T
n=523
50
0
C3L−02669_T
n=520
100
50
0
C3L−02891_T
n=92
10
5
0
C3L−02951_T
n=378
40
20
0
C3L−02963_T
n=131
20
10
0
C3L−02964_T
n=179
40
20
0
C3L−02968_T
n=230
20
10
0
C3L−02969_T
n=243
40
20
0
C3L−03272_T
n=249
40
20
0
C3L−03678_T
n=382
40
20
0
C3L−03961_T
n=58
20
10
0

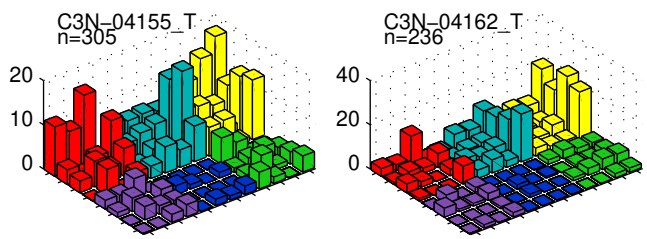

C3L−03962_T
n=232
40
20
0
C3L−03963_T
n=189
20
10
0
C3L−03965_T
n=403
40
20
0
C3L−04013_T
n=153
40
20
0
C3L−04014_T
n=276
40
20
0
C3L−04071_T
n=116
20
10
0
C3L−04391_T
n=186
20
10
0
C3N−00211_T
n=34
20
10
0
C3N−00221_T
n=287
40
20
0
C3N−00247_T
n=29
10
5
0
C3N−00497_T
n=287
20
10
0
C3N−00555_T
n=222
20
10
0

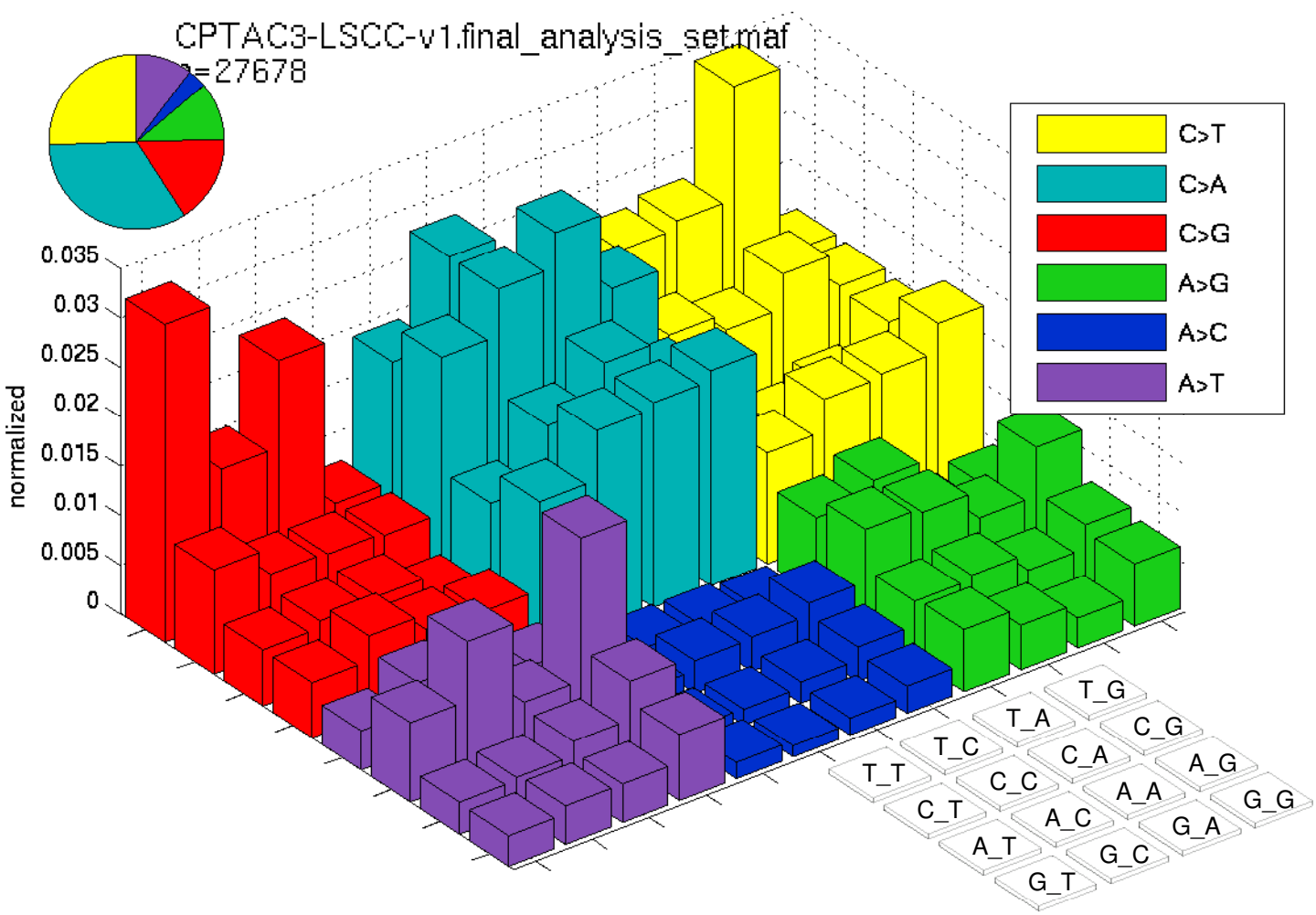

C3N-01017_T
n=235
20
10
0
C3N-01018_T
n=534
50
0
C3N-01020_T
n=170
20
10
0
C3N-01025_T
n=229
40
20
0
C3N-01028_T
n=625
40
20
0
C3N-01194_T
n=277
40
20
0
C3N-01411_T
n=388
40
20
0
C3N-01846_T
n=133
20
10
0
C3N-01892_T
n=279
40
20
0
C3N-01893_T
n=501
40
20
0
C3N-02252_T
n=172
20
10
0
C3N-02283_T
n=639
100
50
0


C3N–02284_T
n=496
40
20
0
C3N–02285_T
n=310
40
20
0
C3N–02288_T
n=216
20
10
0
C3N–02289_T
n=231
40
20
0
C3N–02300_T
n=609
200
100
0
C3N–02339_T
n=71
10
5
0
C3N–02374_T
n=227
20
10
0
C3N–02375_T
n=218
20
10
0
C3N–02425_T
n=198
20
10
0
C3N–02426_T
n=42
10
5
0
C3N–02434_T
n=182
20
10
0
C3N–02435_T
n=234
40
20
0


C3N–02494_T
n=237
40
20
0
C3N–02523_T
n=260
20
10
0
C3N–02575_T
n=264
40
20
0
C3N–02675_T
n=211
20
10
0
C3N–03051_T
n=298
20
10
0
C3N–03072_T
n=216
20
10
0
C3N–03076_T
n=151
40
20
0
C3N–03092_T
n=116
10
5
0
C3N–03093_T
n=35
4
2
0
C3N–03204_T
n=308
40
20
0
C3N–03424_T
n=174
20
10
0
C3N–03425_T
n=412
100
50
0

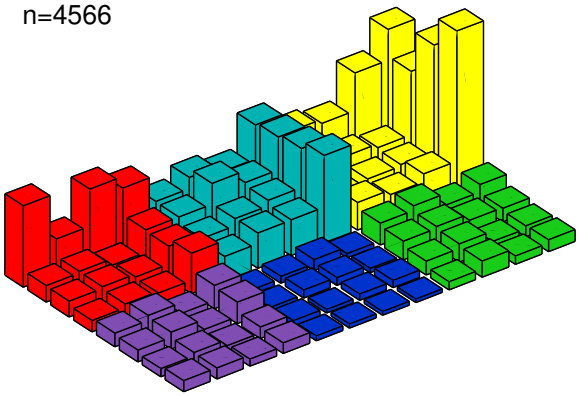

C3N−03441_T
n=238
40
20
0
C3N−03486_T
n=435
40
20
0
C3N−03662_T
n=204
20
10
0
C3N−03848_T
n=5
2
1
0
C3N−03851_T
n=484
50
0
C3N−03875_T
n=507
100
50
0
C3N−03877_T
n=115
20
10
0
C3N−03880_T
n=183
20
10
0
C3N−03882_T
n=219
40
20
0
C3N−03886_T
n=232
20
10
0
C3N−04124_T
n=214
40
20
0
C3N−04127_T
n=65
10
5
0


C3N-04155_T
n=305
20
10
0
C3N-04162_T
n=236
40
20
0

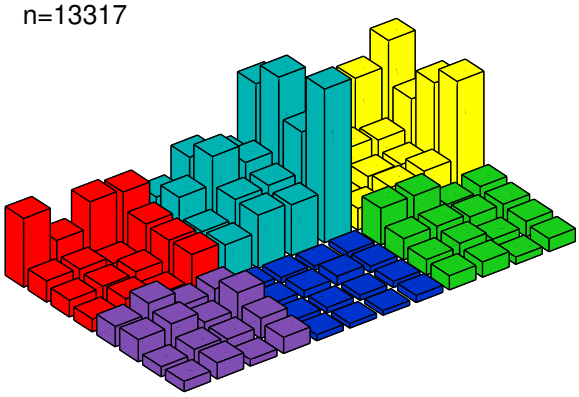

CPTAC3-LSCC-v1.final_analysis_set.maf
=27678
normalized
0.035
0.03
0.025
0.02
0.015
0.01
0.005
0
C>T
C>A
C>G
A>G
A>C
A>T
T_G
T_A
C_G
T_C
C_A
A_G
T_T
C_C
A_A
G_G
C_T
A_C
G_A
A_T
G_C
G_T

CPTAC3−LSCC−v1.final_analysis_set.maf

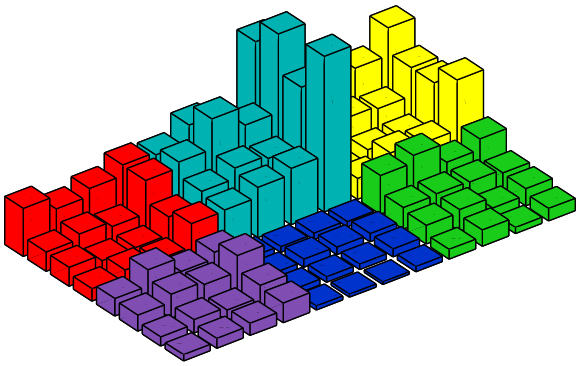
0.00<=AF<0.10
n=4566



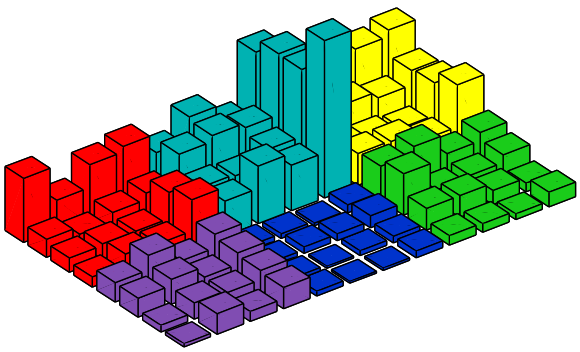
0.10<=AF<0.25
n=13317

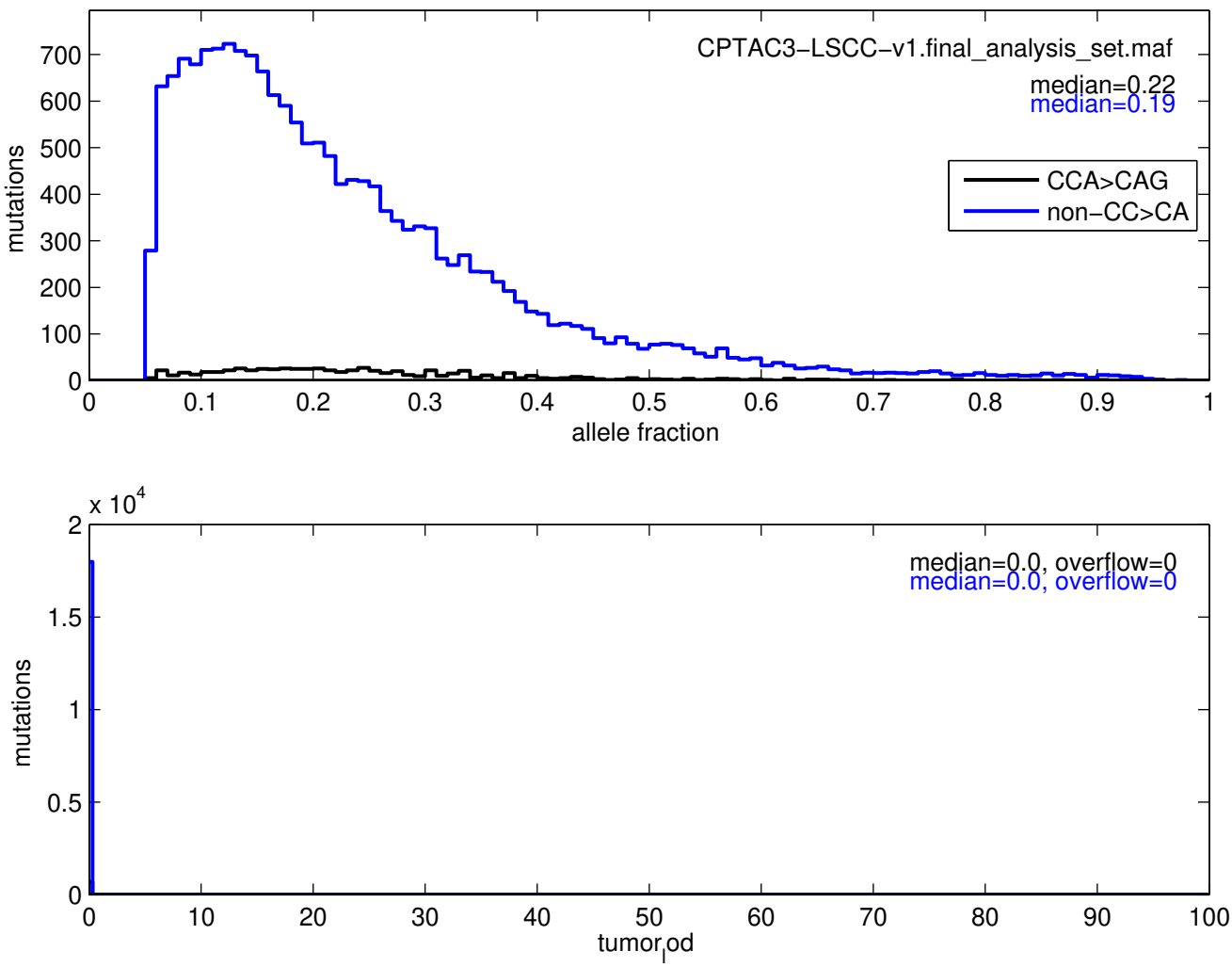


0.25<=AF<0.50
n=7984

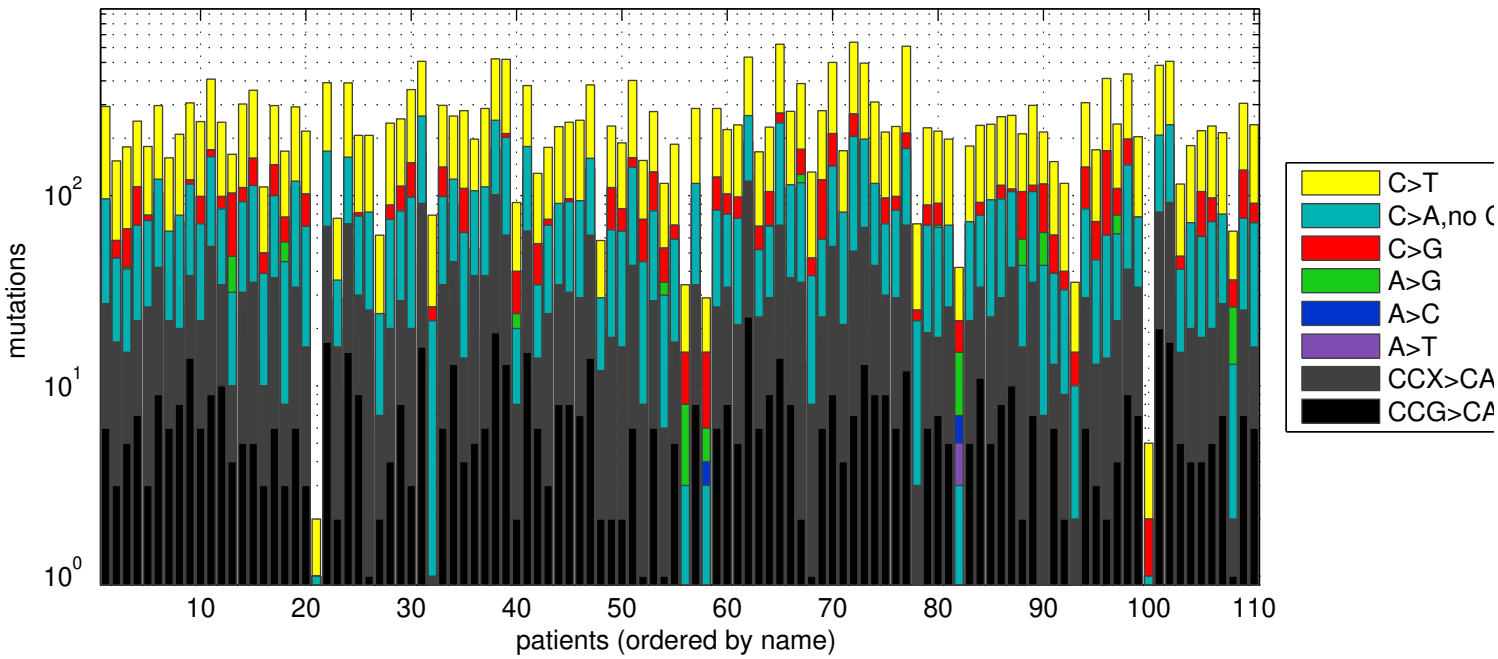


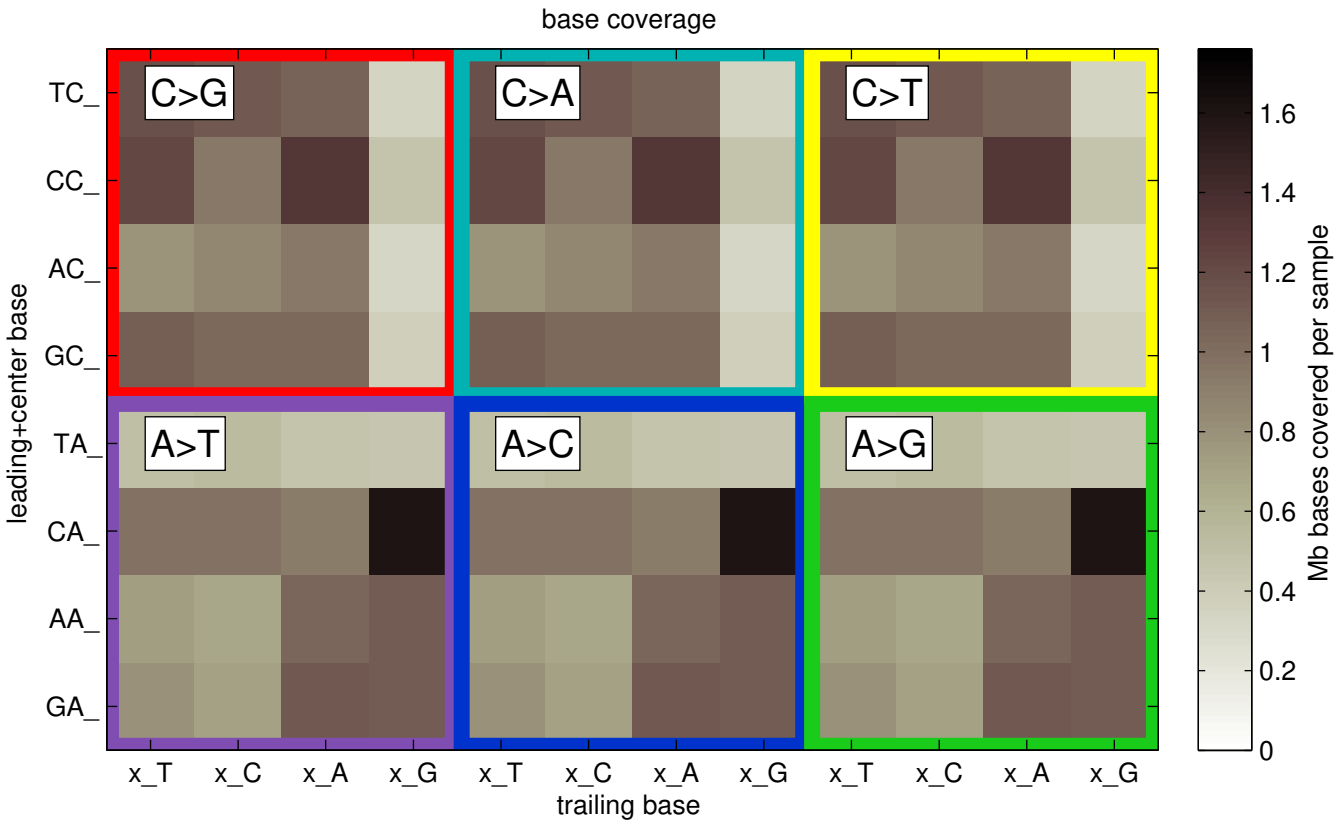
0.50<=AF<1.00
n=1810




CPTAC3−LSCC−v1.final_analysis_set.maf
median=0.22
median=0.19
CCA>CAG
non−CC>CA
mutations
allele fraction
0
100
200
300
400
500
600
700
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
1

x 10^4
median=0.0, overflow=0
median=0.0, overflow=0
mutations
tumor_od
0
0.5
1
1.5
2
10
20
30
40
50
60
70
80
90
100


mutations
10^2
10^1
10^0
10
20
30
40
50
60
70
80
90
100
110
patients (ordered by name)
C>T
C>A,no C
C>G
A>G
A>C
A>T
CCX>CA
CCG>CA


base coverage
C>G
C>A
C>T
A>T
A>C
A>G
TC_
CC_
AC_
GC_
TA_
CA_
AA_
GA_
leading+center base
x_T
x_C
x_A
x_G
trailing base
Mb bases covered per sample
0
0.2
0.4
0.6
0.8
1
1.2
1.4
1.6