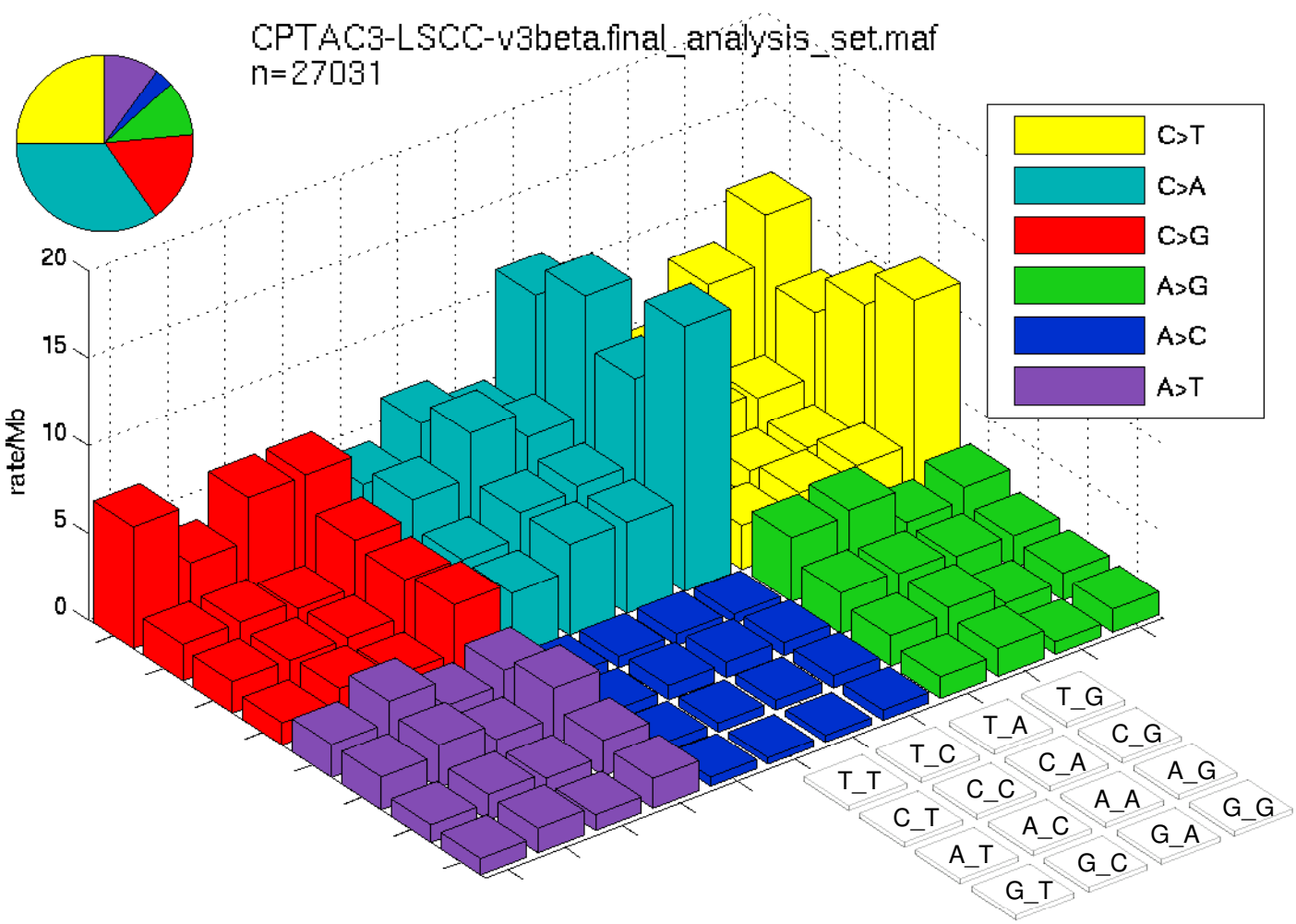
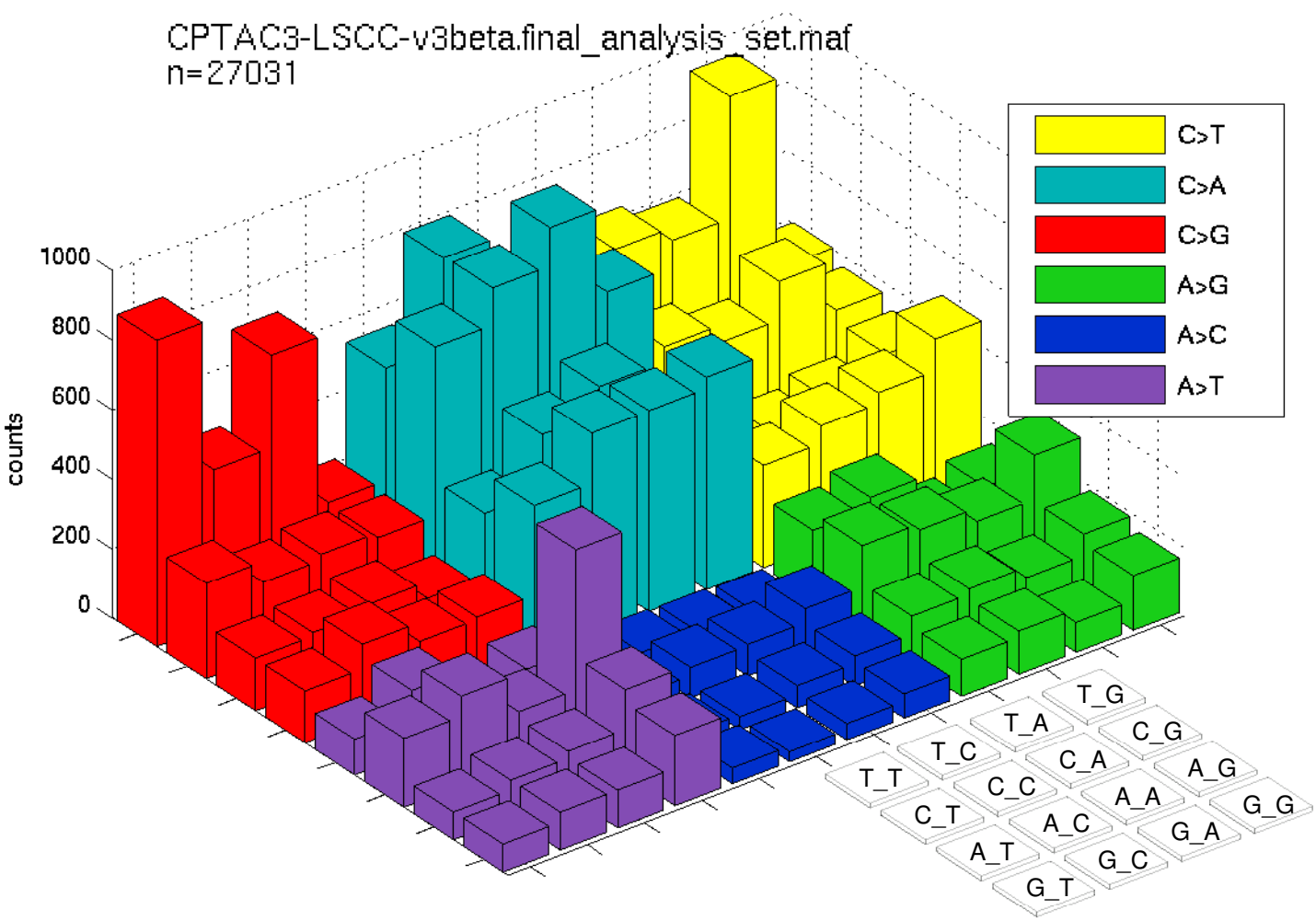

CPTAC3-LSCC-v3beta.final_analysis_set.maf
n=27031
rate/Mb
20
15
10
5
0
C>T
C>A
C>G
A>G
A>C
A>T
T_G
T_A
C_G
T_C
C_A
A_G
T_T
C_C
A_A
G_G
C_T
A_C
G_A
A_T
G_C
G_T


CPTAC3-LSCC-v3beta.final_analysis_set.maf
n=27031
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
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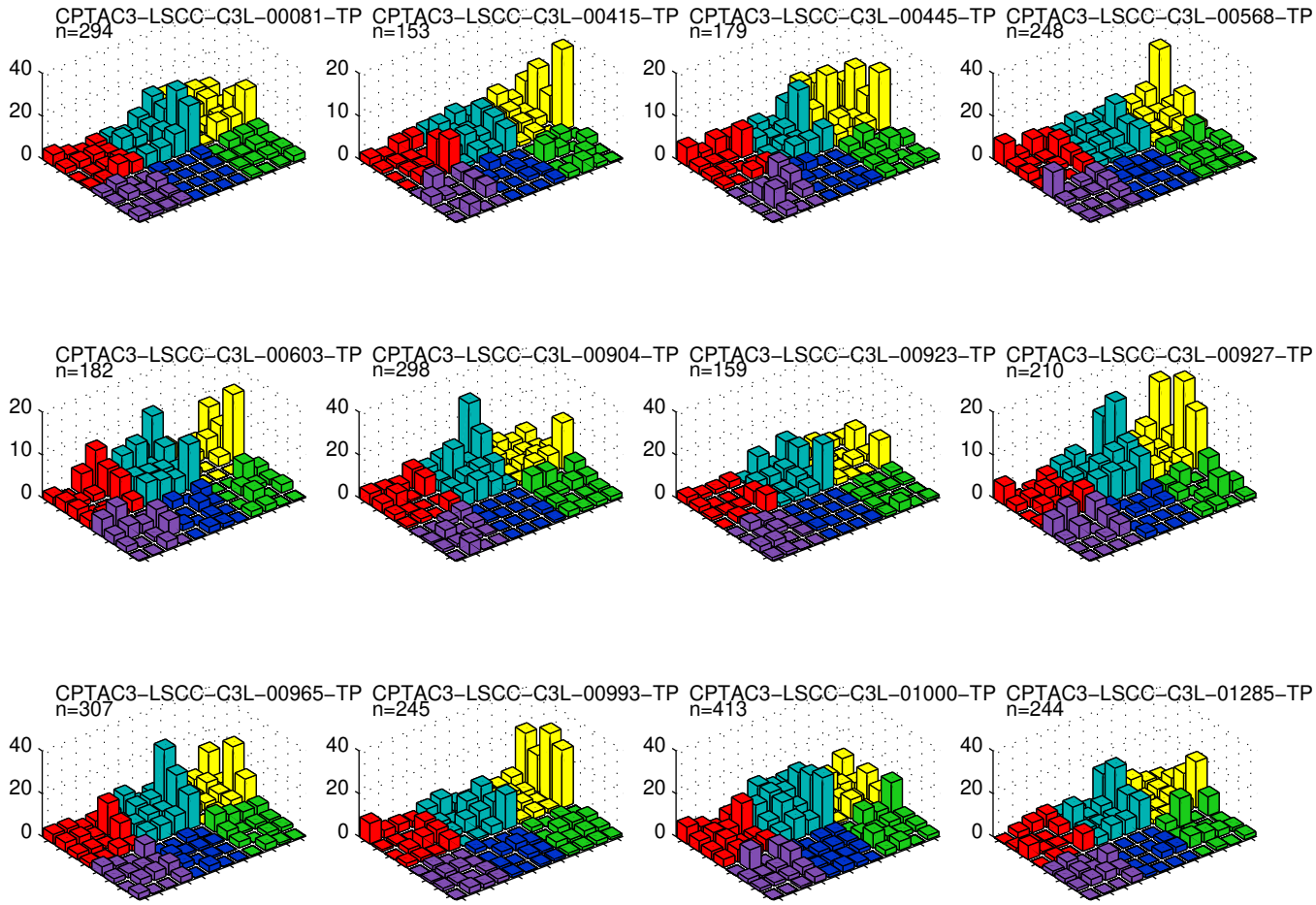
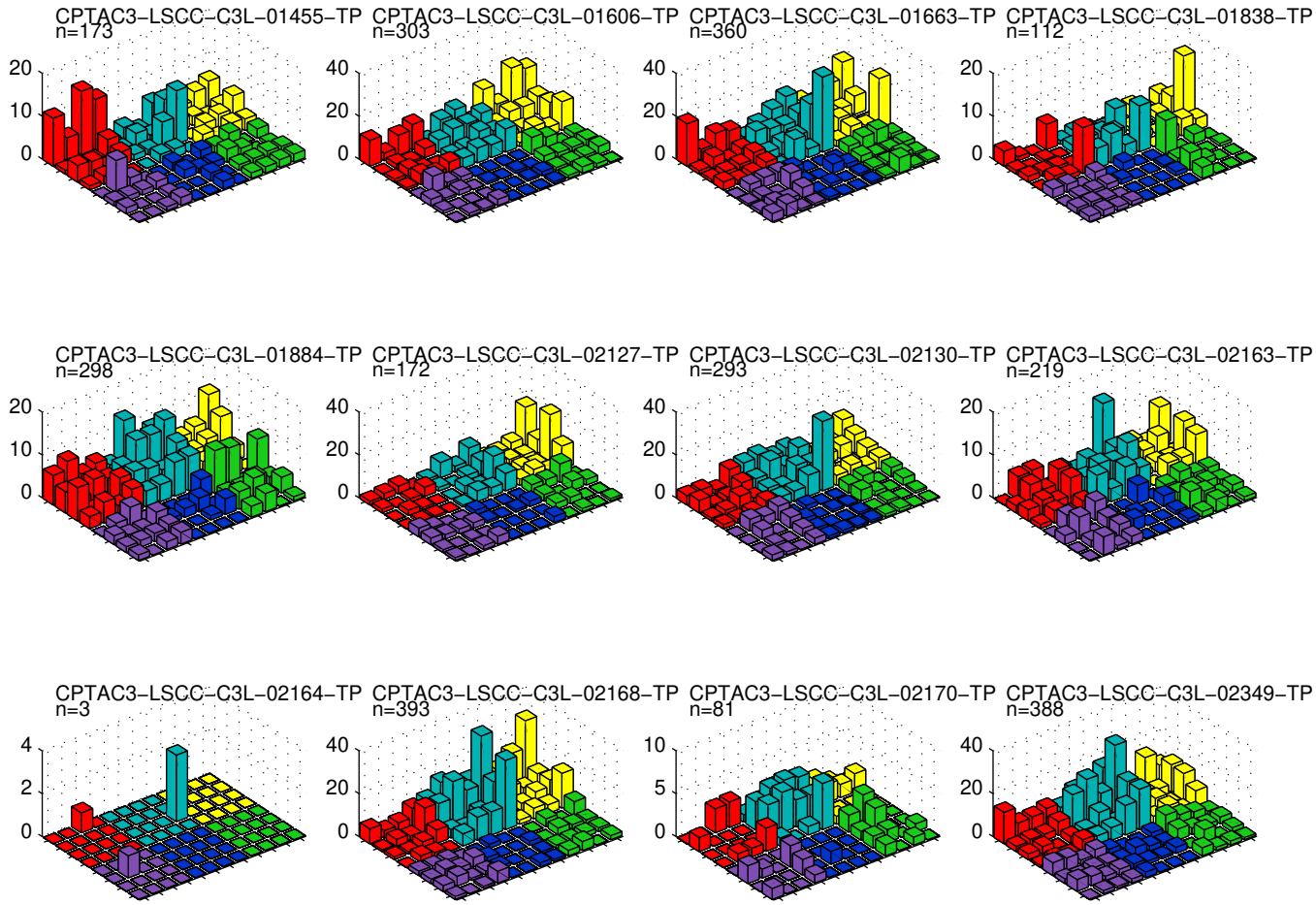
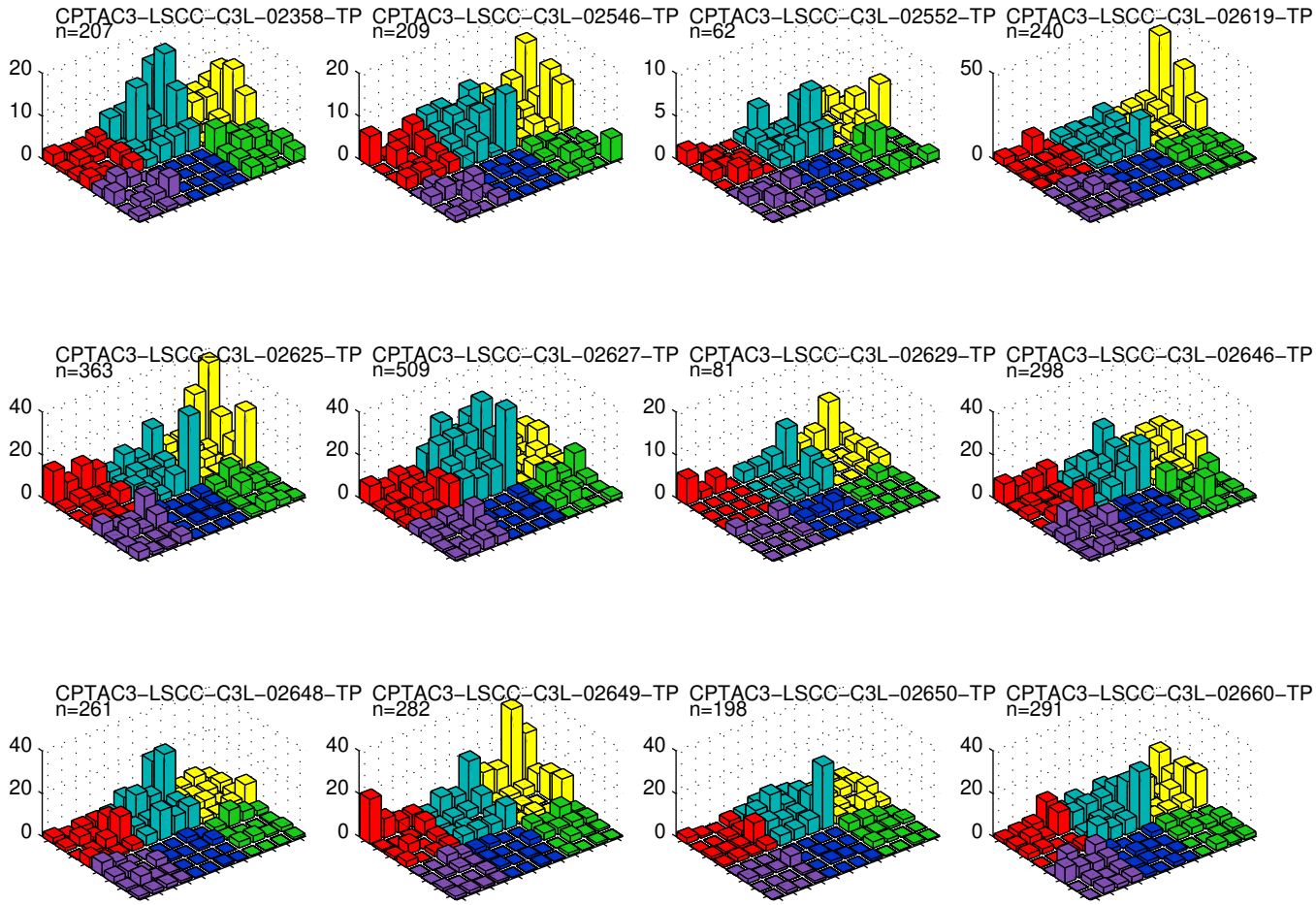
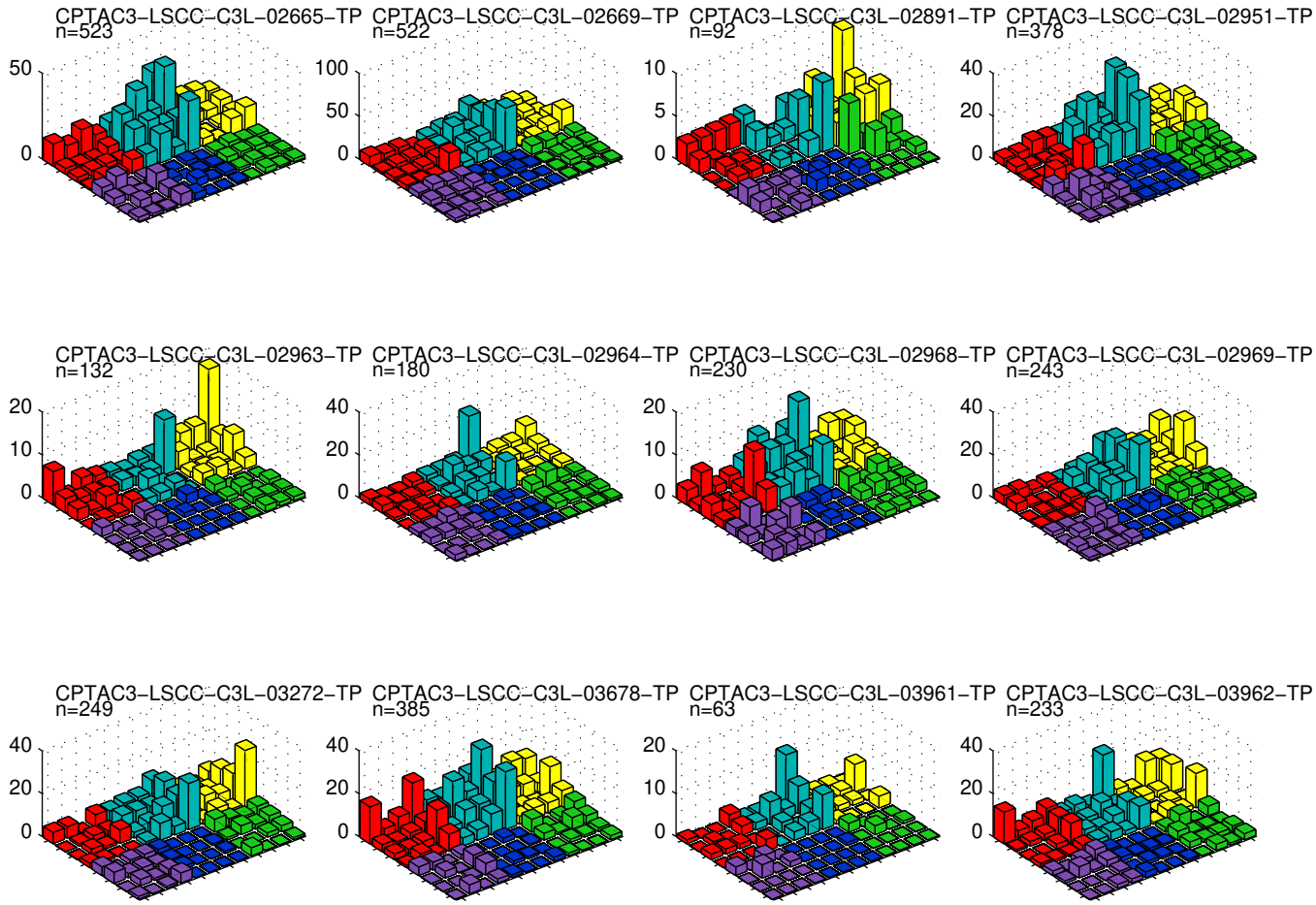
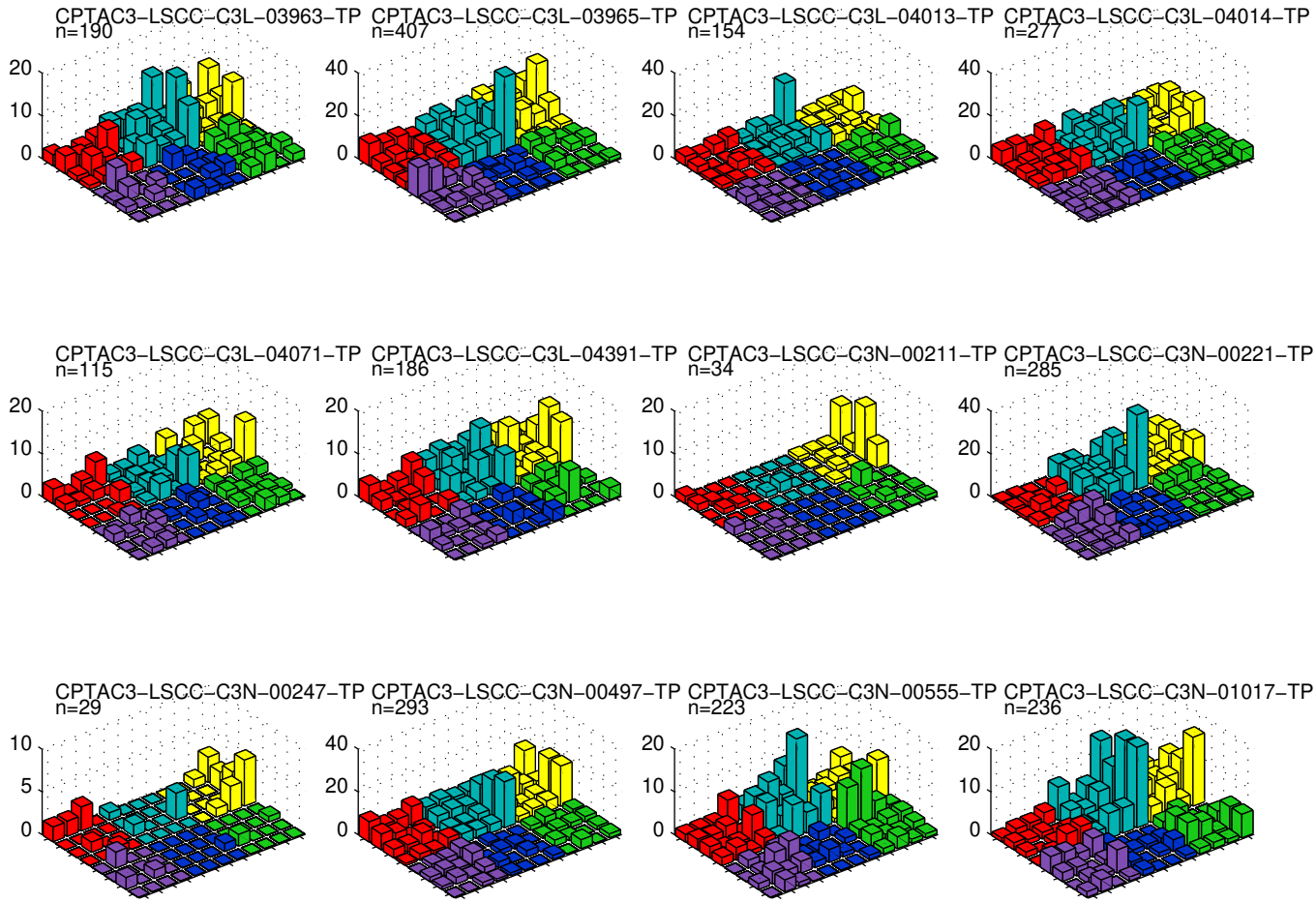
A_T
G_C
G_T

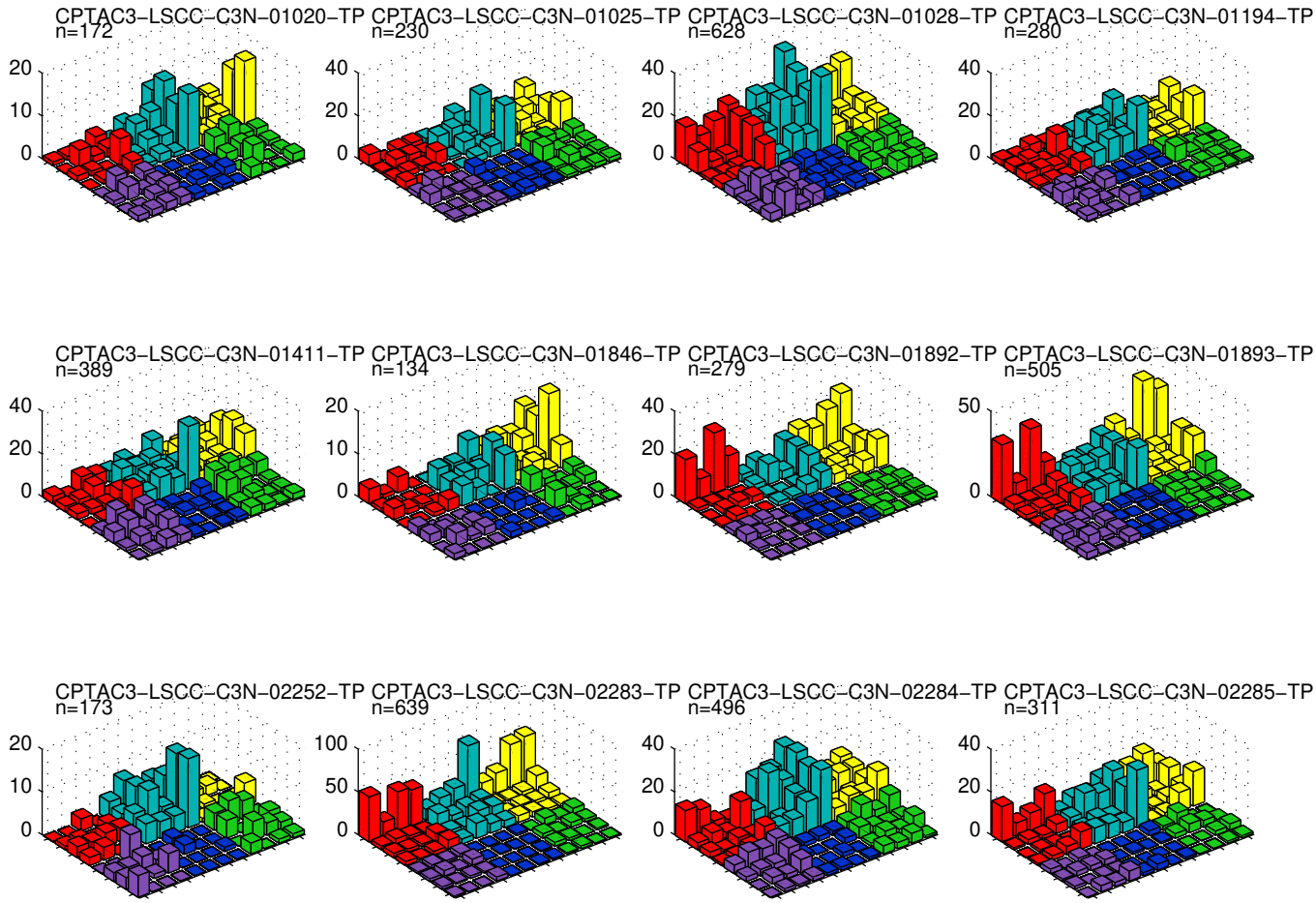

CPTAC3−LSCC−C3L−00081−TP
n=294
CPTAC3−LSCC−C3L−00415−TP
n=153
CPTAC3−LSCC−C3L−00445−TP
n=179
CPTAC3−LSCC−C3L−00568−TP
n=248
CPTAC3−LSCC−C3L−00603−TP
n=182
CPTAC3−LSCC−C3L−00904−TP
n=298
CPTAC3−LSCC−C3L−00923−TP
n=159
CPTAC3−LSCC−C3L−00927−TP
n=210
CPTAC3−LSCC−C3L−00965−TP
n=307
CPTAC3−LSCC−C3L−00993−TP
n=245
CPTAC3−LSCC−C3L−01000−TP
n=413
CPTAC3−LSCC−C3L−01285−TP
n=244

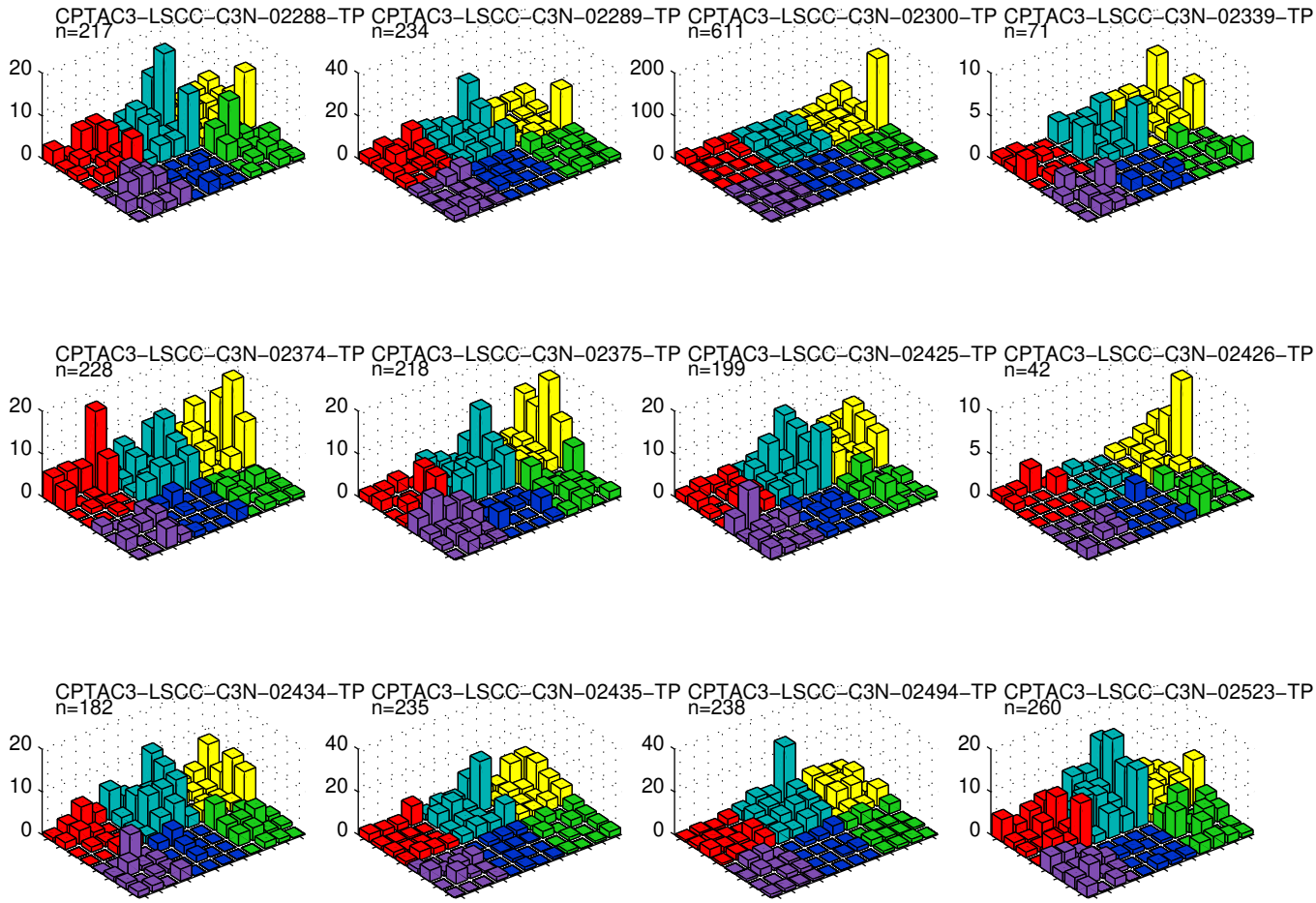

CPTAC3−LSCC−C3L−01455−TP
n=173
20
10
0
CPTAC3−LSCC−C3L−01606−TP
n=303
40
20
0
CPTAC3−LSCC−C3L−01663−TP
n=360
40
20
0
CPTAC3−LSCC−C3L−01838−TP
n=112
20
10
0
CPTAC3−LSCC−C3L−01884−TP
n=298
20
10
0
CPTAC3−LSCC−C3L−02127−TP
n=172
40
20
0
CPTAC3−LSCC−C3L−02130−TP
n=293
40
20
0
CPTAC3−LSCC−C3L−02163−TP
n=219
20
10
0
CPTAC3−LSCC−C3L−02164−TP
n=3
4
2
0
CPTAC3−LSCC−C3L−02168−TP
n=393
40
20
0
CPTAC3−LSCC−C3L−02170−TP
n=81
10
5
0
CPTAC3−LSCC−C3L−02349−TP
n=388
40
20
0

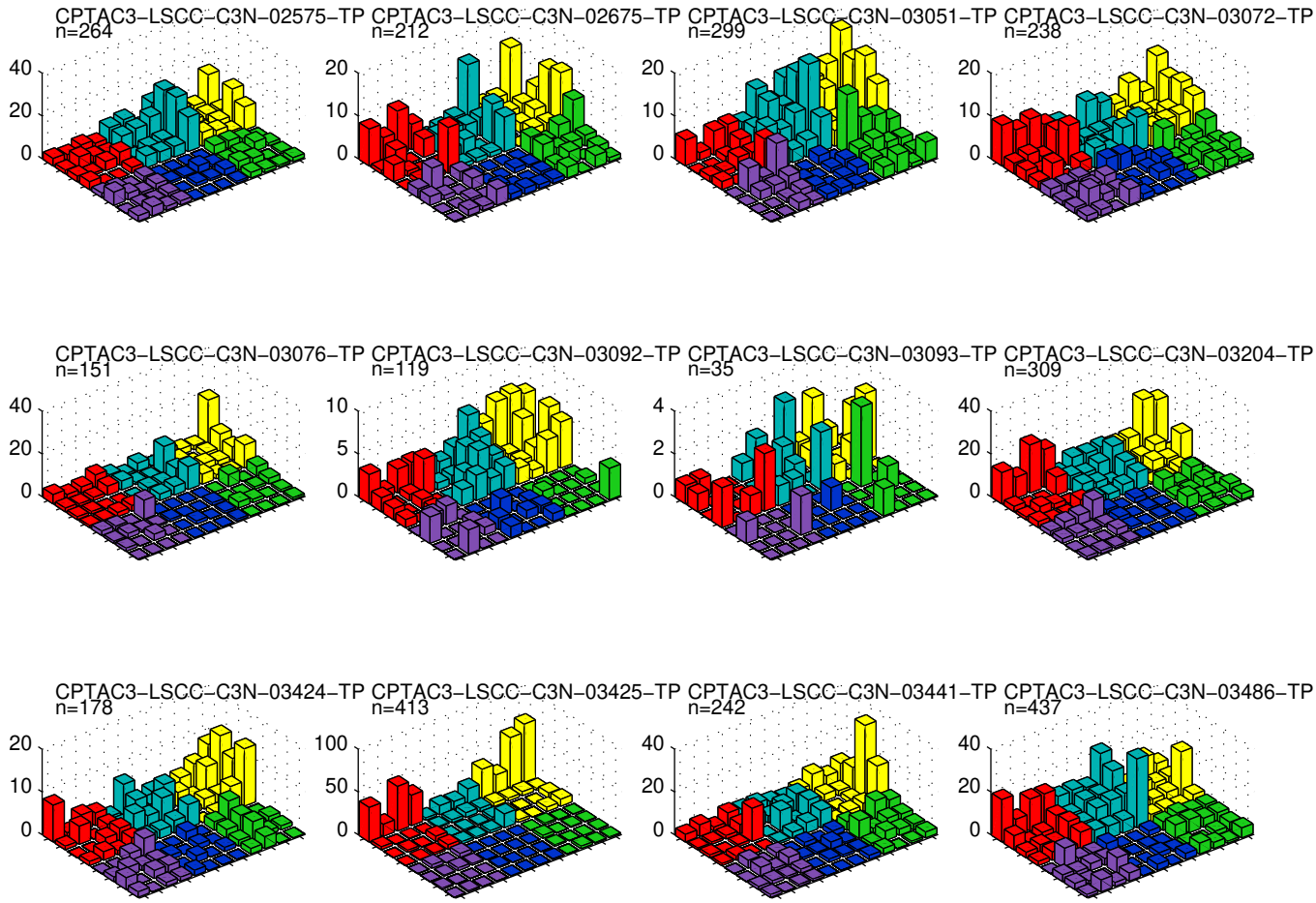

CPTAC3−LSCC−C3L−02358−TP
n=207
20
10
0
CPTAC3−LSCC−C3L−02546−TP
n=209
20
10
0
CPTAC3−LSCC−C3L−02552−TP
n=62
10
5
0
CPTAC3−LSCC−C3L−02619−TP
n=240
50
0
CPTAC3−LSCC−C3L−02625−TP
n=363
40
20
0
CPTAC3−LSCC−C3L−02627−TP
n=509
40
20
0
CPTAC3−LSCC−C3L−02629−TP
n=81
20
10
0
CPTAC3−LSCC−C3L−02646−TP
n=298
40
20
0
CPTAC3−LSCC−C3L−02648−TP
n=261
40
20
0
CPTAC3−LSCC−C3L−02649−TP
n=282
40
20
0
CPTAC3−LSCC−C3L−02650−TP
n=198
40
20
0
CPTAC3−LSCC−C3L−02660−TP
n=291
40
20
0

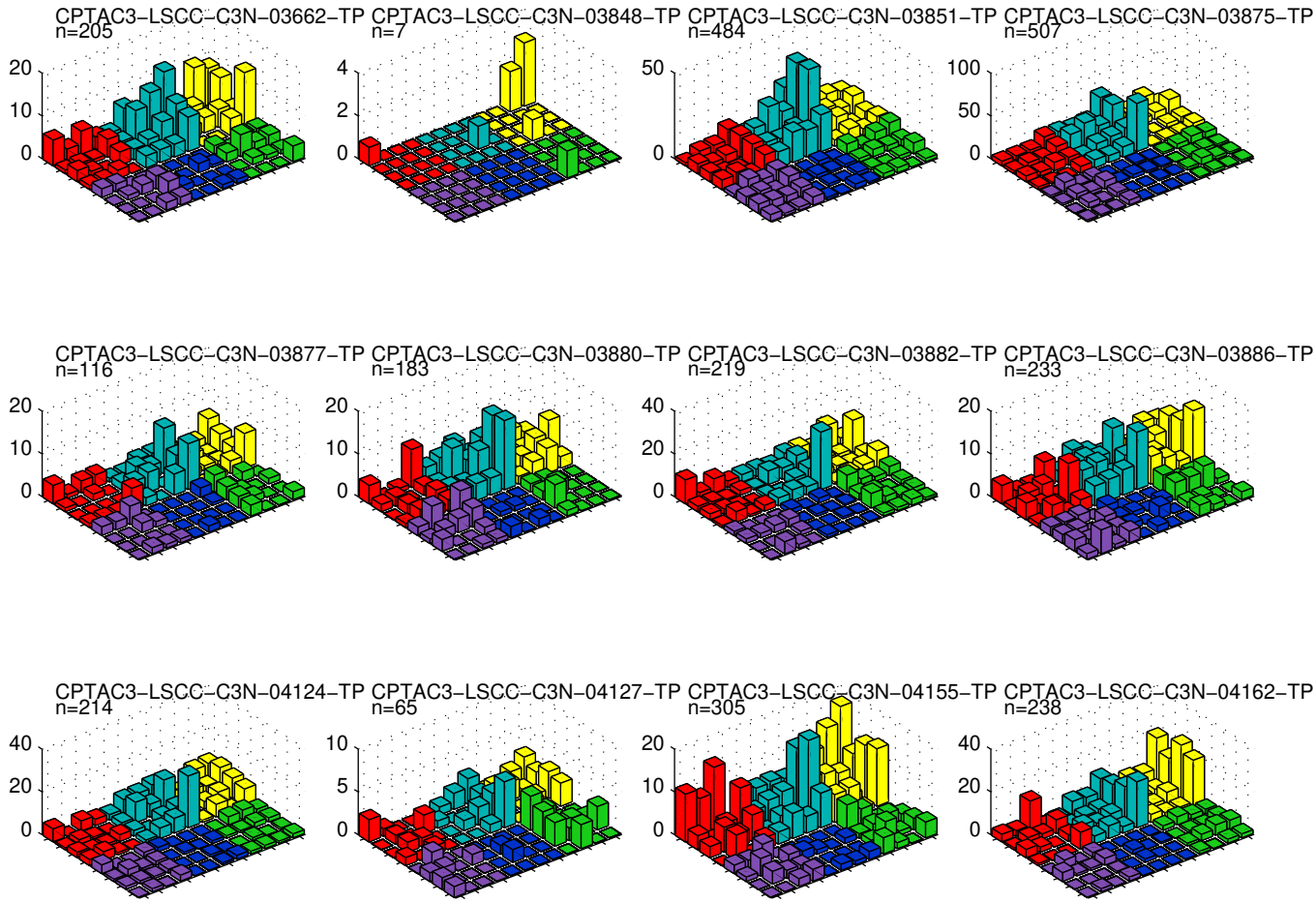

CPTAC3−LSCC−C3L−02665−TP
n=523
50
0
CPTAC3−LSCC−C3L−02669−TP
n=522
100
50
0
CPTAC3−LSCC−C3L−02891−TP
n=92
10
5
0
CPTAC3−LSCC−C3L−02951−TP
n=378
40
20
0
CPTAC3−LSCC−C3L−02963−TP
n=132
20
10
0
CPTAC3−LSCC−C3L−02964−TP
n=180
40
20
0
CPTAC3−LSCC−C3L−02968−TP
n=230
20
10
0
CPTAC3−LSCC−C3L−02969−TP
n=243
40
20
0
CPTAC3−LSCC−C3L−03272−TP
n=249
40
20
0
CPTAC3−LSCC−C3L−03678−TP
n=385
40
20
0
CPTAC3−LSCC−C3L−03961−TP
n=63
20
10
0
CPTAC3−LSCC−C3L−03962−TP
n=233
40
20
0

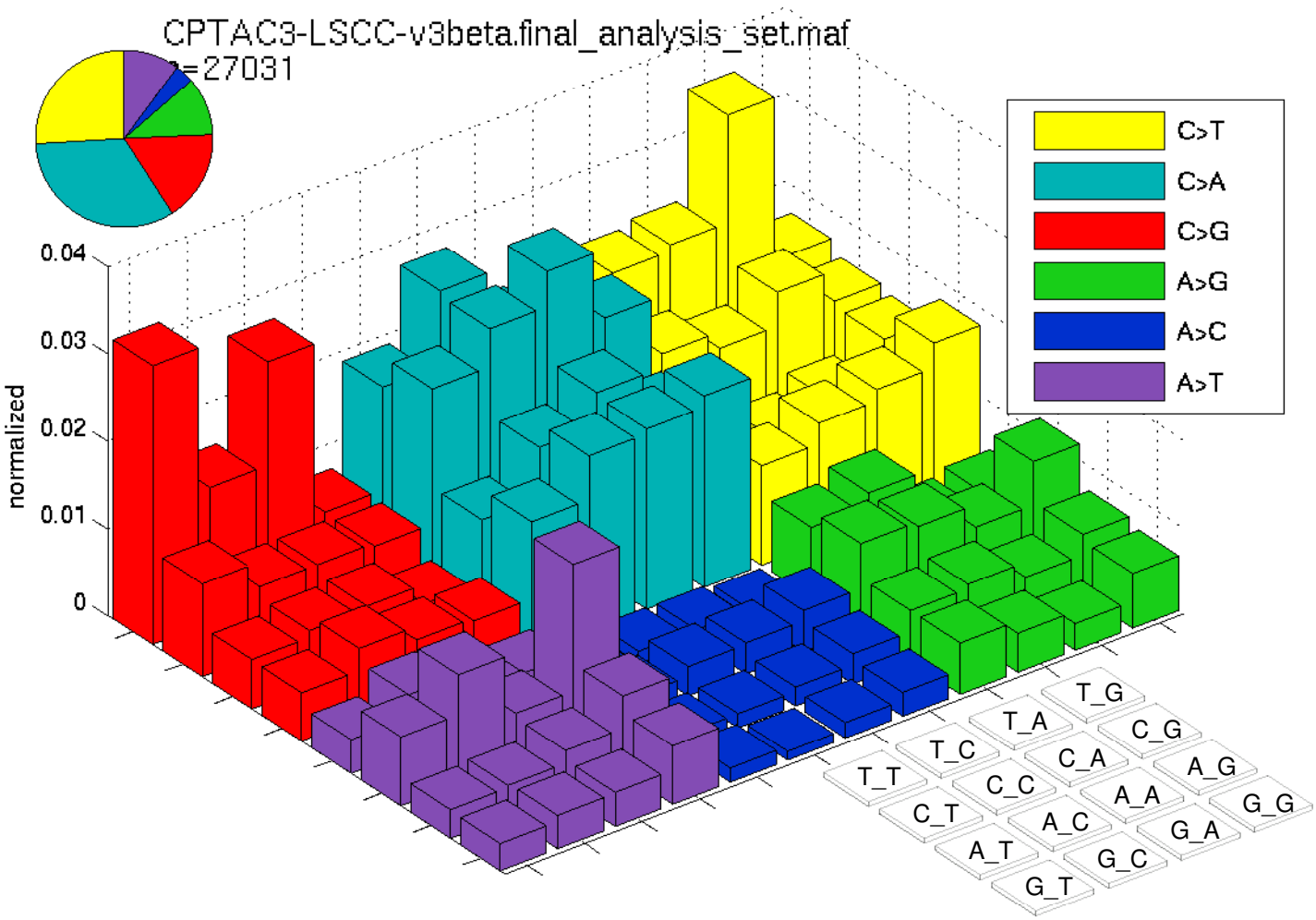

CPTAC3−LSCC−C3L−03963−TP
n=190
CPTAC3−LSCC−C3L−03965−TP
n=407
CPTAC3−LSCC−C3L−04013−TP
n=154
CPTAC3−LSCC−C3L−04014−TP
n=277
CPTAC3−LSCC−C3L−04071−TP
n=115
CPTAC3−LSCC−C3L−04391−TP
n=186
CPTAC3−LSCC−C3N−00211−TP
n=34
CPTAC3−LSCC−C3N−00221−TP
n=285
CPTAC3−LSCC−C3N−00247−TP
n=29
CPTAC3−LSCC−C3N−00497−TP
n=293
CPTAC3−LSCC−C3N−00555−TP
n=223
CPTAC3−LSCC−C3N−01017−TP
n=236


CPTAC3−LSCC−C3N−01020−TP
n=172
CPTAC3−LSCC−C3N−01025−TP
n=230
CPTAC3−LSCC−C3N−01028−TP
n=628
CPTAC3−LSCC−C3N−01194−TP
n=280
CPTAC3−LSCC−C3N−01411−TP
n=389
CPTAC3−LSCC−C3N−01846−TP
n=134
CPTAC3−LSCC−C3N−01892−TP
n=279
CPTAC3−LSCC−C3N−01893−TP
n=505
CPTAC3−LSCC−C3N−02252−TP
n=173
CPTAC3−LSCC−C3N−02283−TP
n=639
CPTAC3−LSCC−C3N−02284−TP
n=496
CPTAC3−LSCC−C3N−02285−TP
n=311


CPTAC3-LSCC-C3N-02288-TP
n=217
CPTAC3-LSCC-C3N-02289-TP
n=234
CPTAC3-LSCC-C3N-02300-TP
n=611
CPTAC3-LSCC-C3N-02339-TP
n=71
CPTAC3-LSCC-C3N-02374-TP
n=228
CPTAC3-LSCC-C3N-02375-TP
n=218
CPTAC3-LSCC-C3N-02425-TP
n=199
CPTAC3-LSCC-C3N-02426-TP
n=42
CPTAC3-LSCC-C3N-02434-TP
n=182
CPTAC3-LSCC-C3N-02435-TP
n=235
CPTAC3-LSCC-C3N-02494-TP
n=238
CPTAC3-LSCC-C3N-02523-TP
n=260

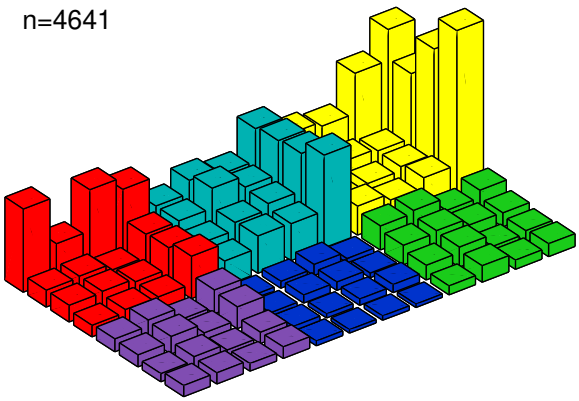

CPTAC3−LSCC−C3N−02575−TP
n=264
CPTAC3−LSCC−C3N−02675−TP
n=212
CPTAC3−LSCC−C3N−03051−TP
n=299
CPTAC3−LSCC−C3N−03072−TP
n=238
CPTAC3−LSCC−C3N−03076−TP
n=151
CPTAC3−LSCC−C3N−03092−TP
n=119
CPTAC3−LSCC−C3N−03093−TP
n=35
CPTAC3−LSCC−C3N−03204−TP
n=309
CPTAC3−LSCC−C3N−03424−TP
n=178
CPTAC3−LSCC−C3N−03425−TP
n=413
CPTAC3−LSCC−C3N−03441−TP
n=242
CPTAC3−LSCC−C3N−03486−TP
n=437


CPTAC3−LSCC−C3N−03662−TP
n=205
20
10
0
CPTAC3−LSCC−C3N−03848−TP
n=7
4
2
0
CPTAC3−LSCC−C3N−03851−TP
n=484
50
0
CPTAC3−LSCC−C3N−03875−TP
n=507
100
50
0
CPTAC3−LSCC−C3N−03877−TP
n=116
20
10
0
CPTAC3−LSCC−C3N−03880−TP
n=183
20
10
0
CPTAC3−LSCC−C3N−03882−TP
n=219
40
20
0
CPTAC3−LSCC−C3N−03886−TP
n=233
20
10
0
CPTAC3−LSCC−C3N−04124−TP
n=214
40
20
0
CPTAC3−LSCC−C3N−04127−TP
n=65
10
5
0
CPTAC3−LSCC−C3N−04155−TP
n=305
20
10
0
CPTAC3−LSCC−C3N−04162−TP
n=238
40
20
0

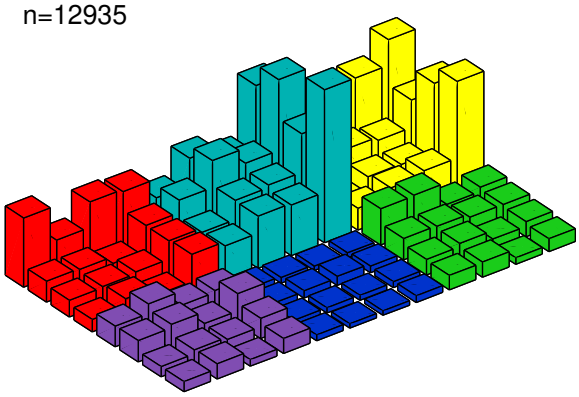

CPTAC3-LSCC-v3beta.final_analysis_set.maf
=27031
normalized
0.04
0.03
0.02
0.01
0
C>T
C>A
C>G
A>G
A>C
A>T
T_G
T_A
C_G
T_C
C_A
A_G
T_T
C_C
A_A
G_G
C_T
A_C
G_A
A_T
G_C
G_T

CPTAC3−LSCC−v3beta.final_analysis_set.maf

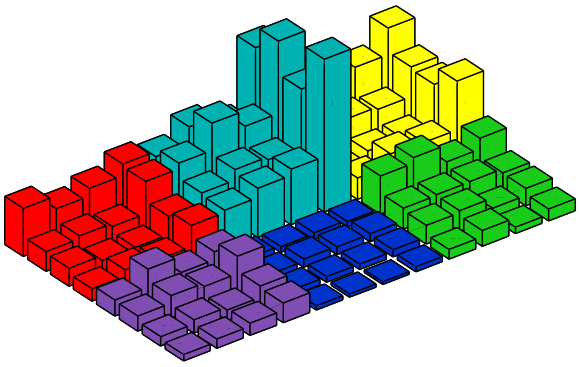
0.00<=AF<0.10
n=4641



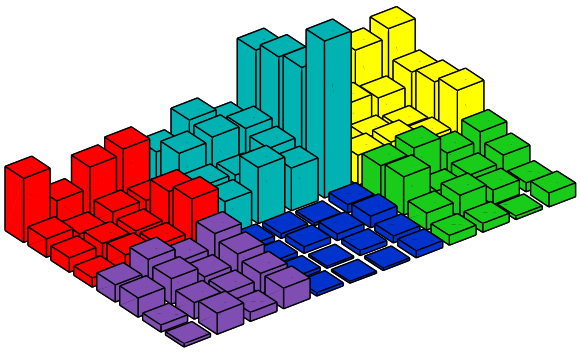
0.10<=AF<0.25
n=12935

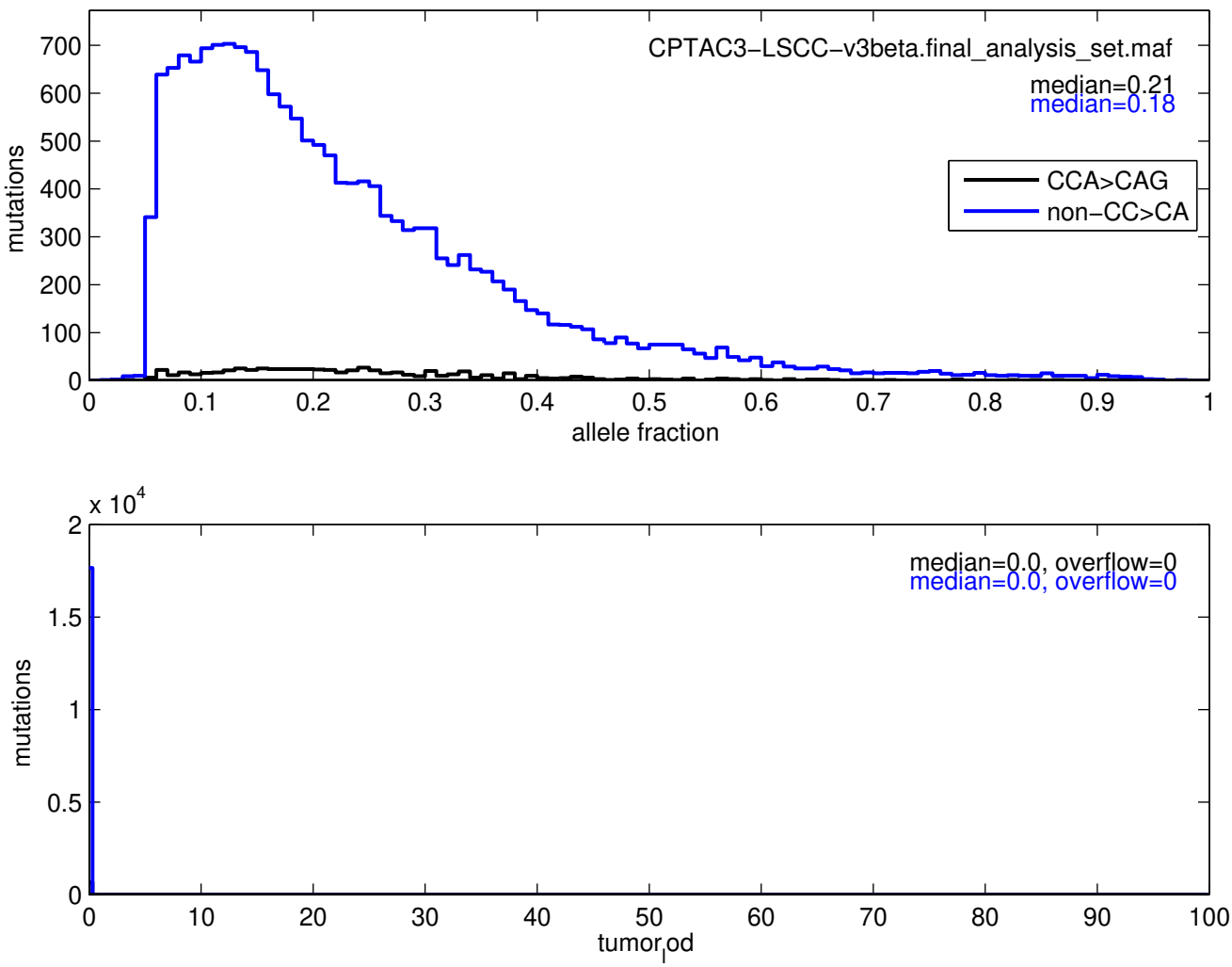


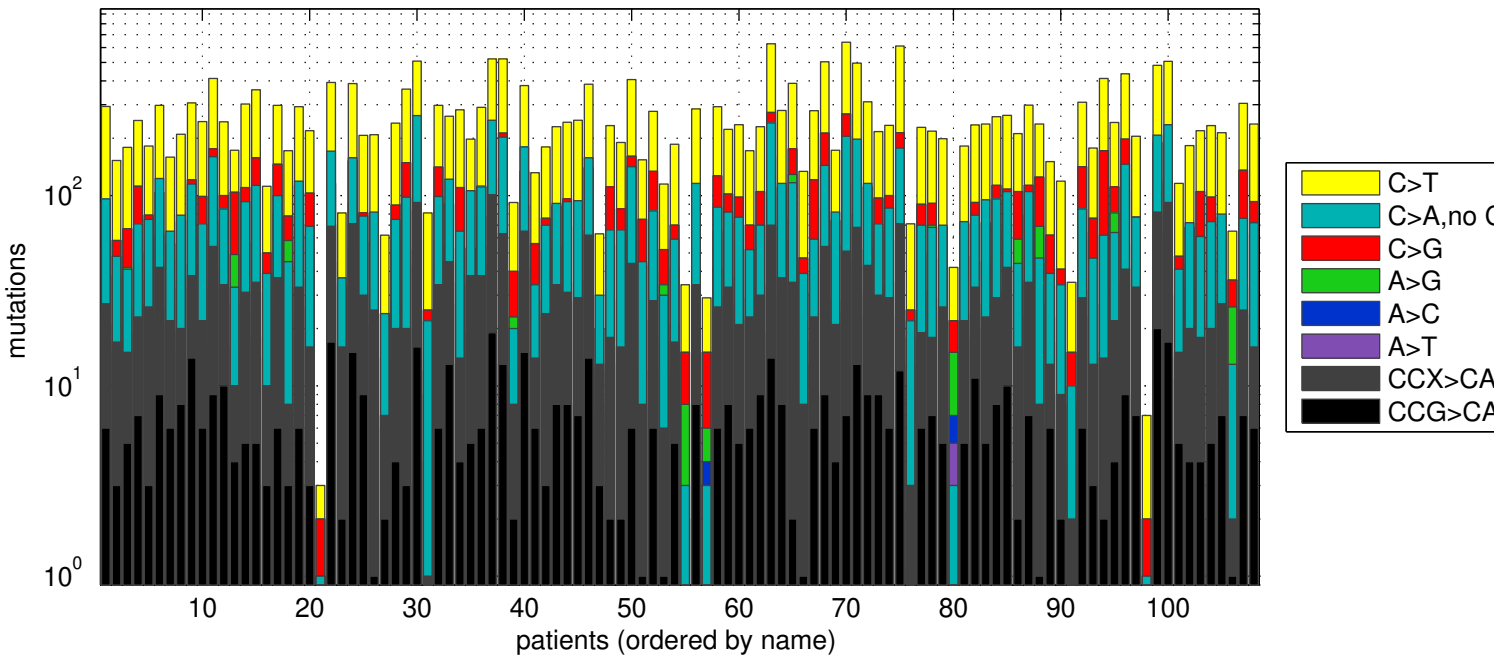
0.25<=AF<0.50
n=7711

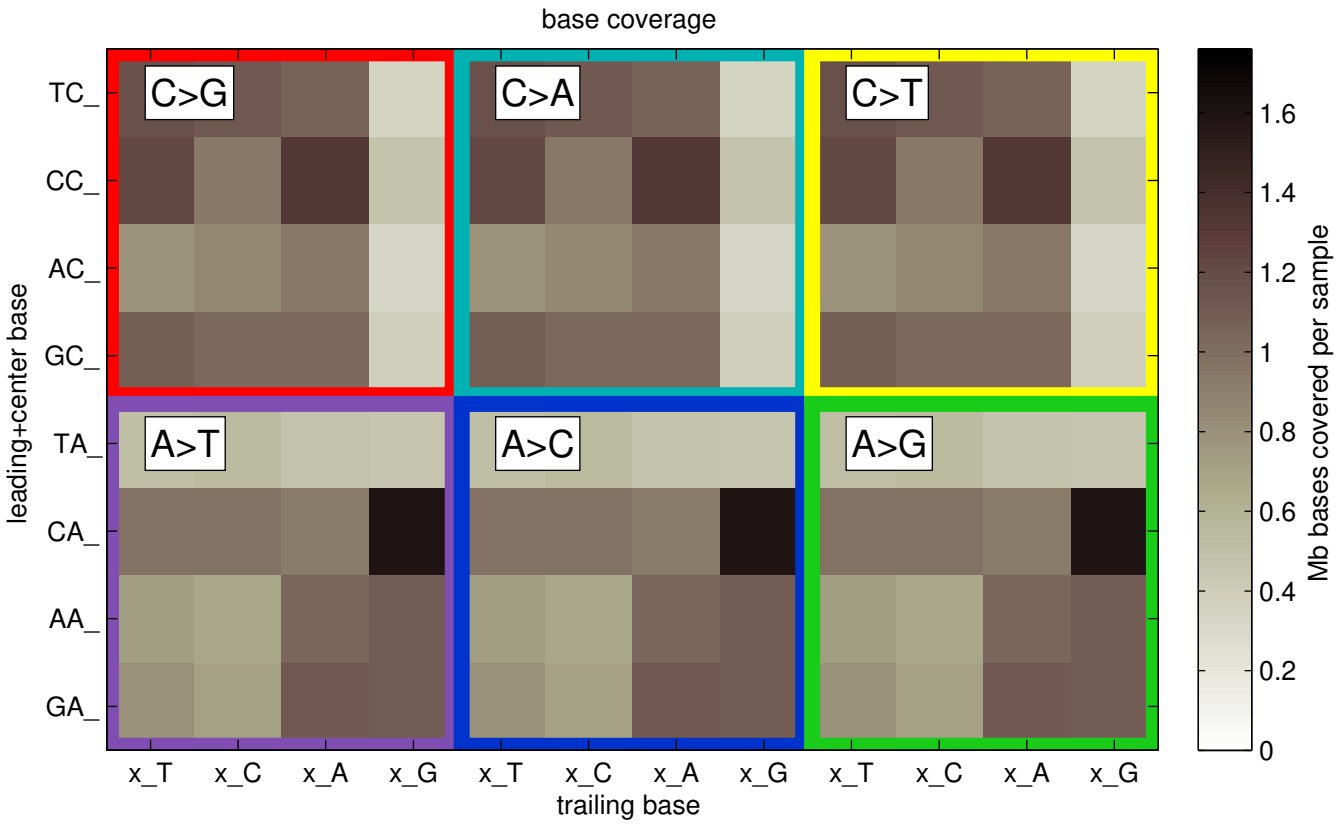


0.50<=AF<1.00
n=1743




CPTAC3−LSCC−v3beta.final_analysis_set.maf
median=0.21
median=0.18
CCA>CAG
non−CC>CA
mutations
allele fraction
x 10^4
median=0.0, overflow=0
median=0.0, overflow=0
mutations
tumor_od


mutations
10^2
10^1
10^0
10
20
30
40
50
60
70
80
90
100
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