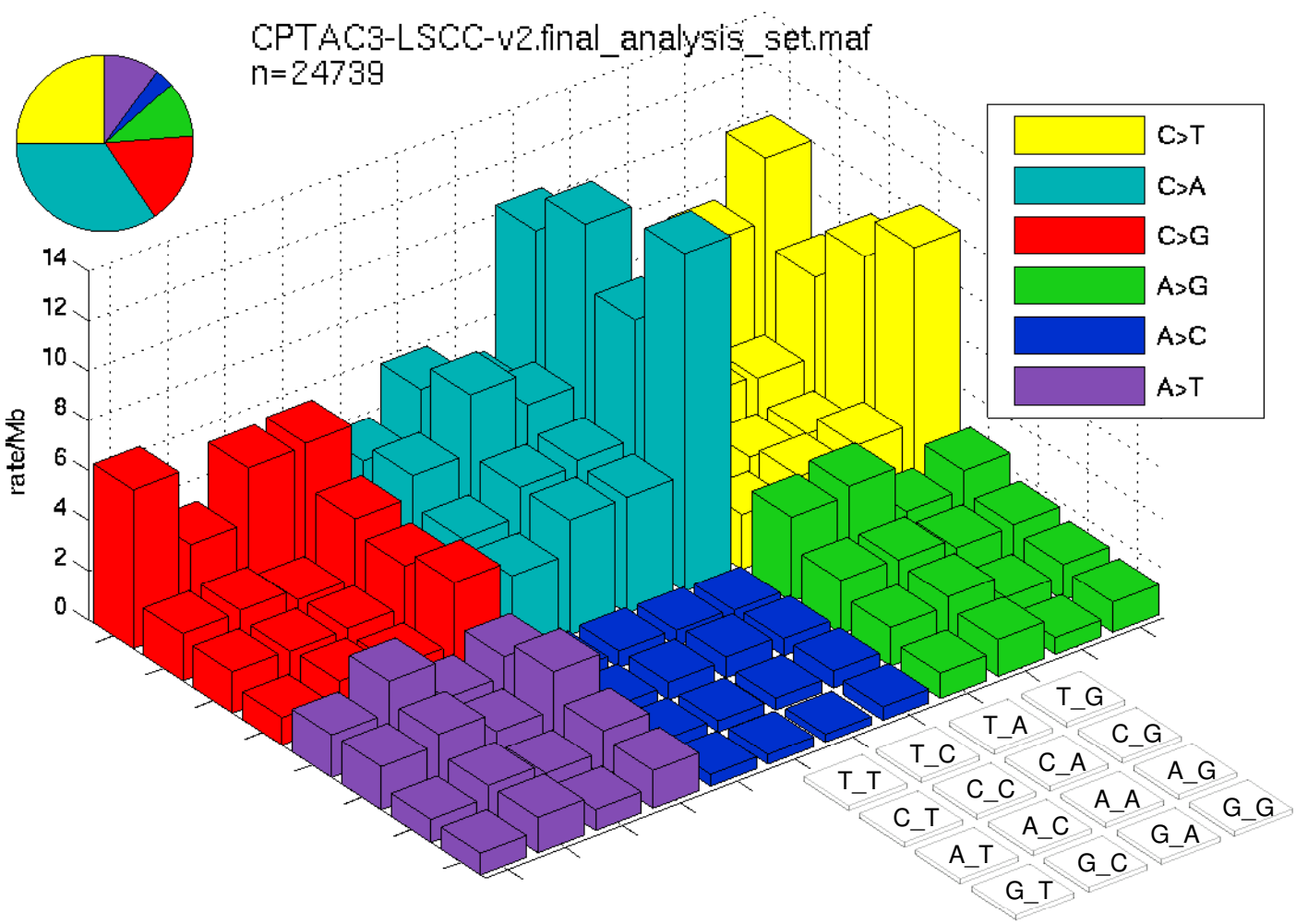
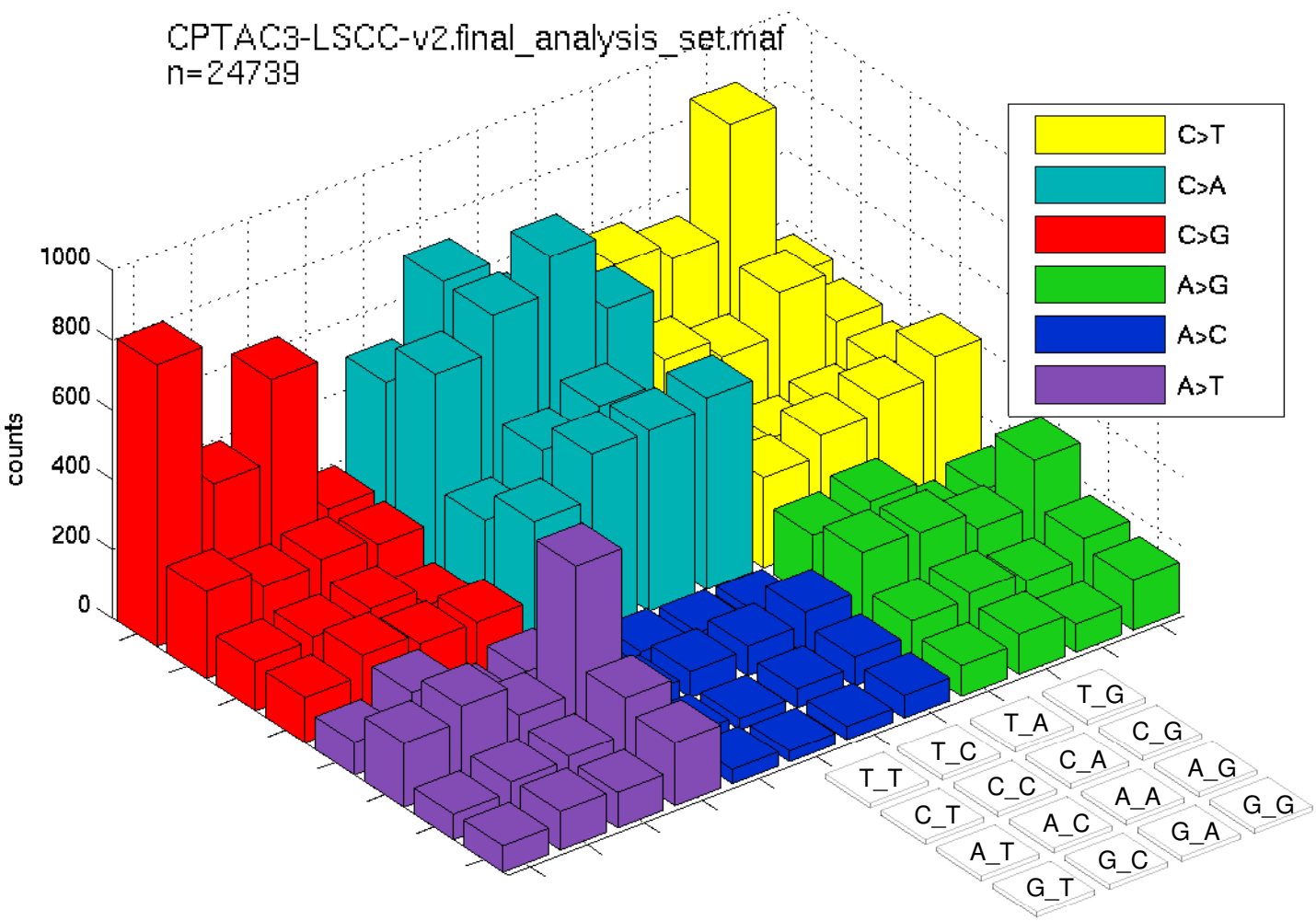
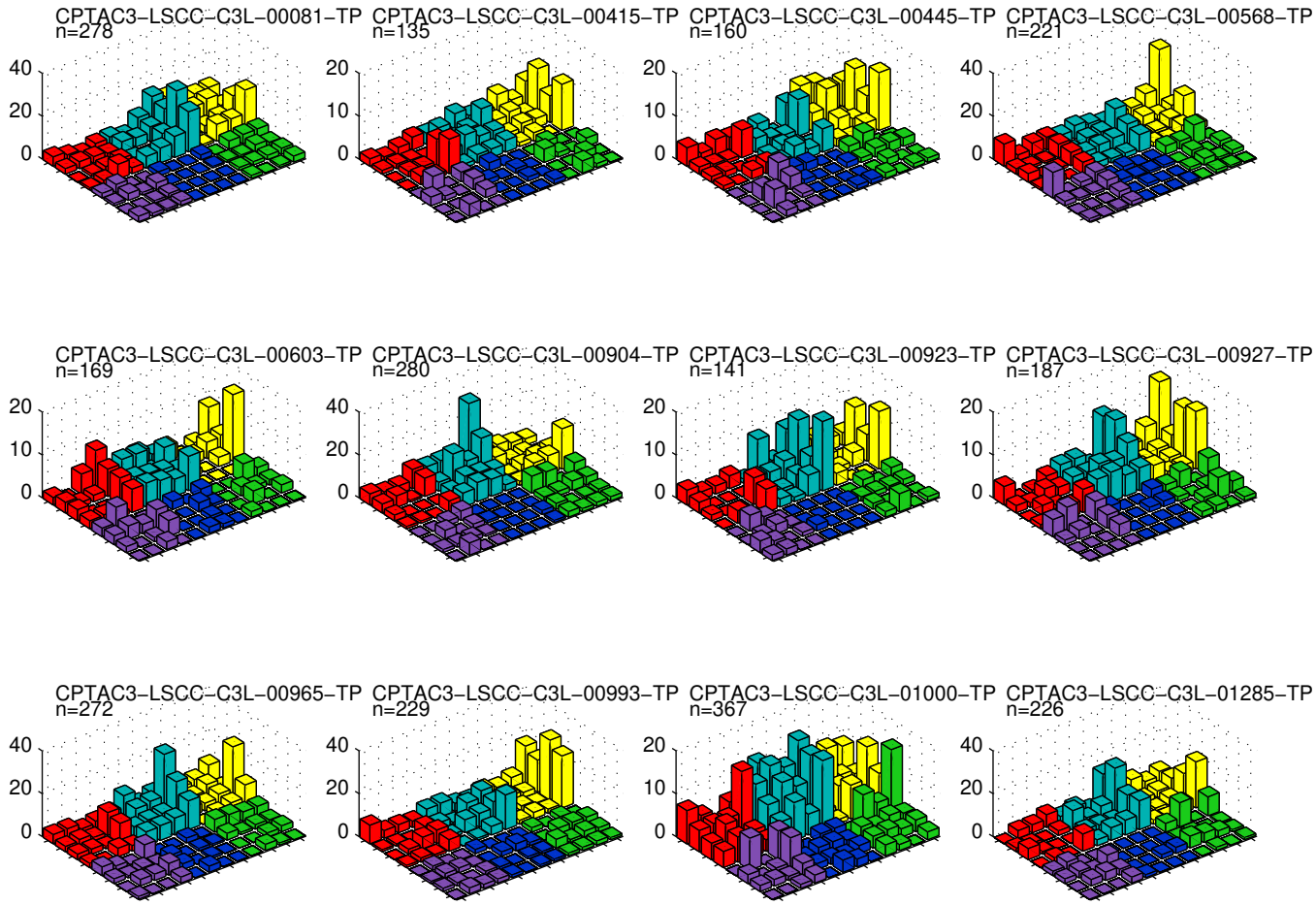
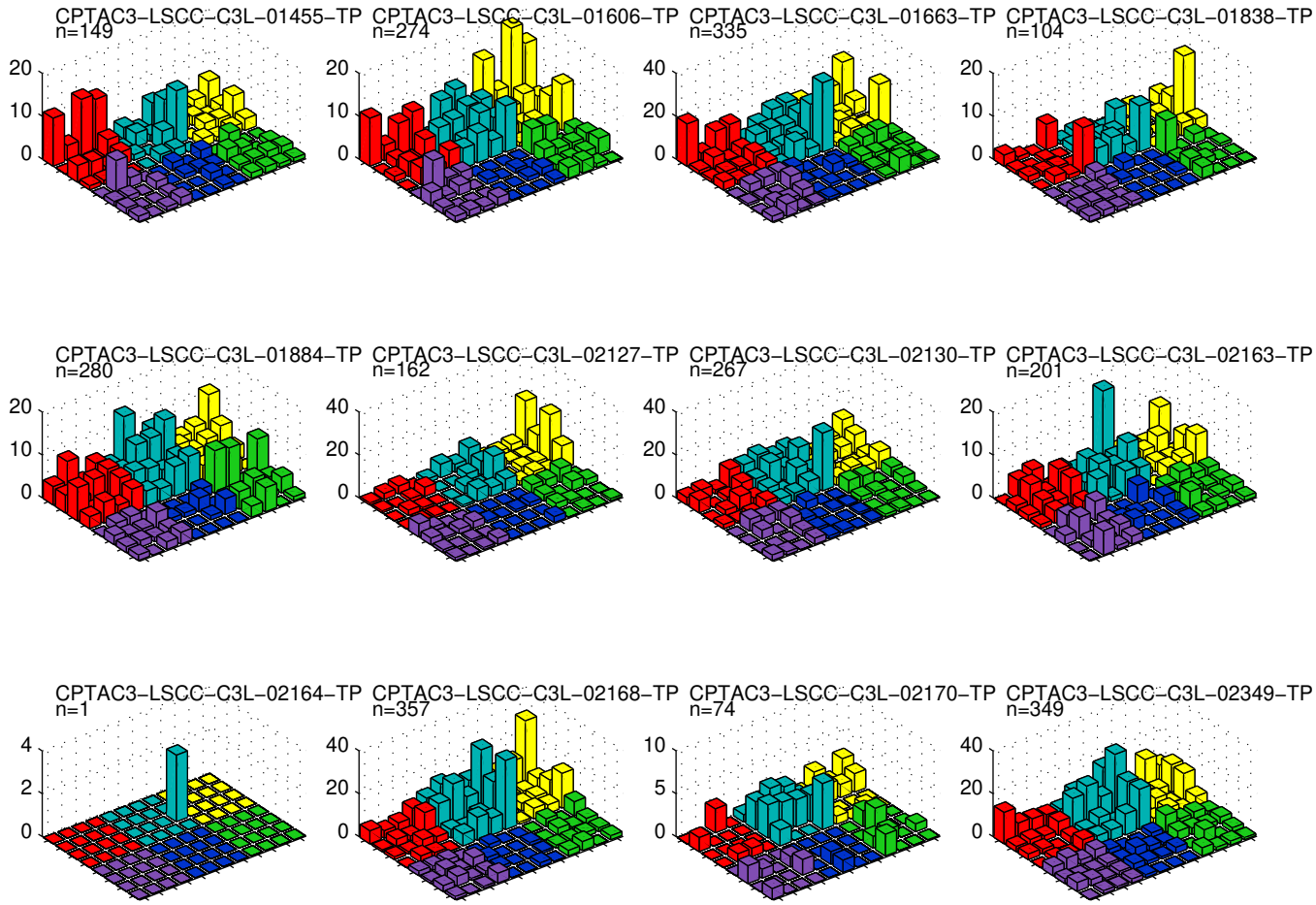
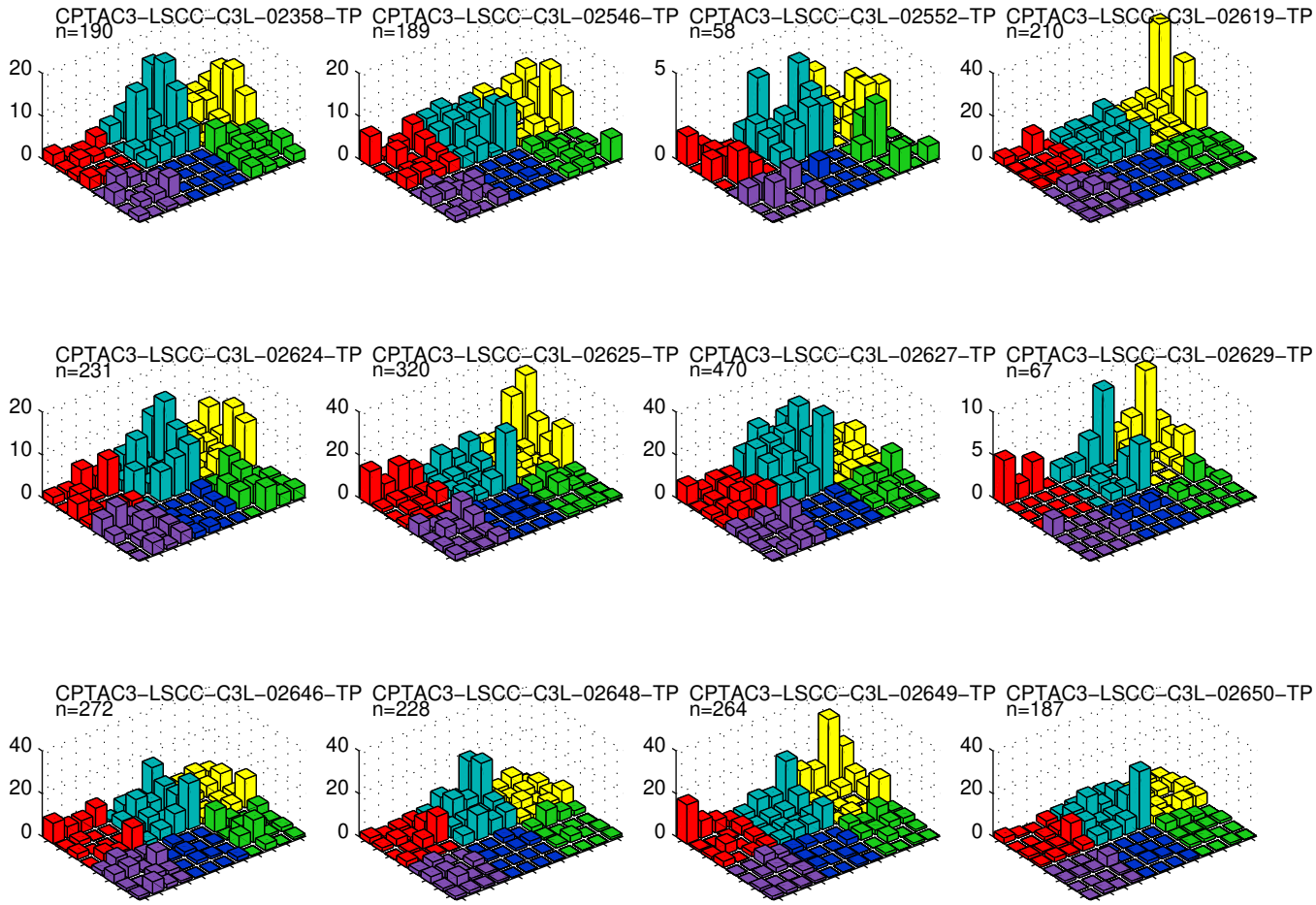
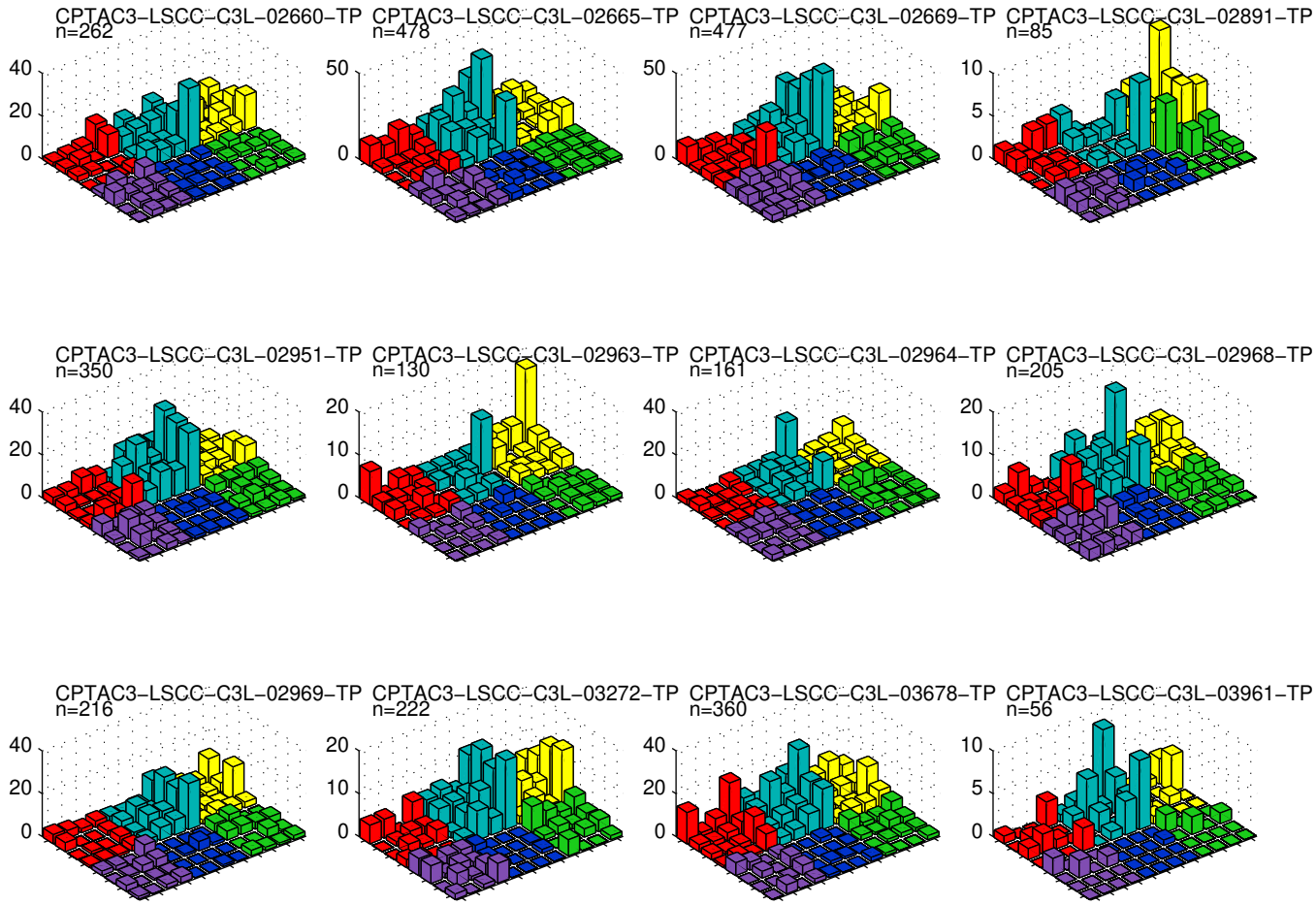

CPTAC3-LSCC-v2.final_analysis_set.maf
n=24739
rate/Mb
0
2
4
6
8
10
12
14
C>T
C>A
C>G
A>G
A>C
A>T
T_G
T_A
C_G
T_C
C_A
A_G
T_T
C_C
A_A
G_G
C_T
A_C
G_A
A_T
G_C
G_T

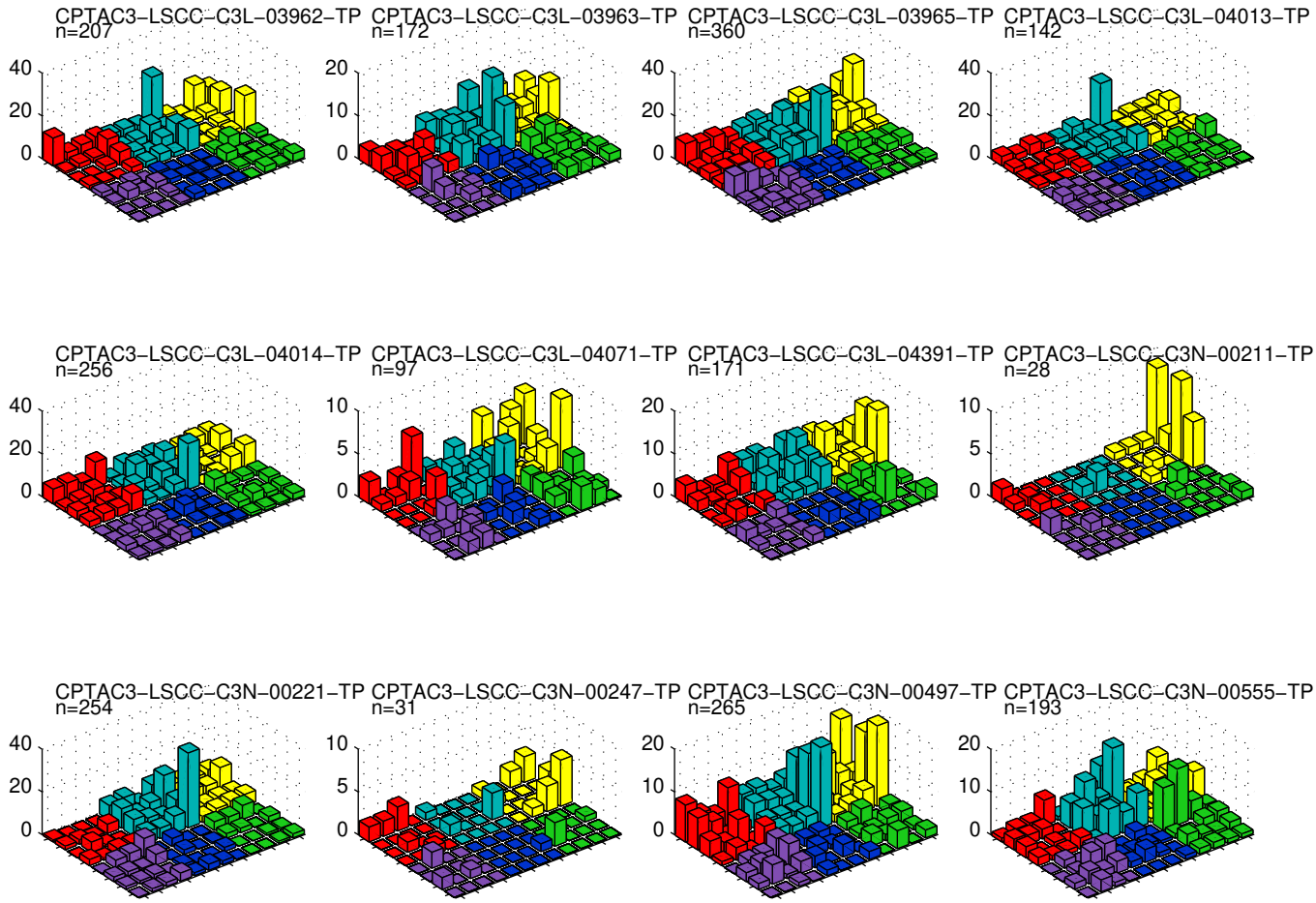

CPTAC3-LSCC-v2.final_analysis_set.maf
n=24739
counts
1000
800
600
400
200
0
C>T
C>A
C>G
A>G
A>C
A>T
T_G
T_A
C_G
T_C
C_A
A_G
T_T
C_C
A_A
G_G
C_T
A_C
G_A
A_T
G_C
G_T

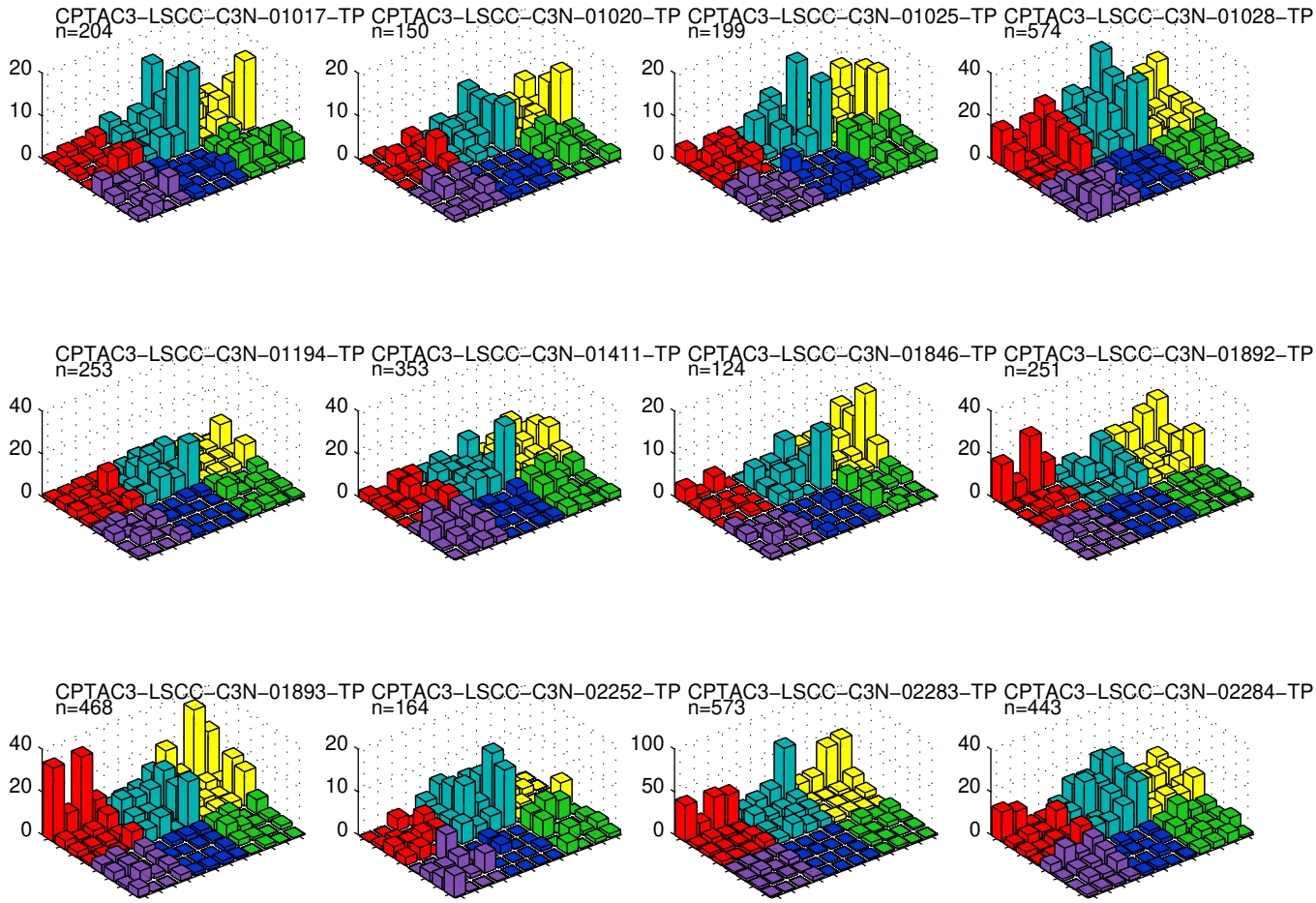

CPTAC3−LSCC−C3L−00081−TP
n=278
CPTAC3−LSCC−C3L−00415−TP
n=135
CPTAC3−LSCC−C3L−00445−TP
n=160
CPTAC3−LSCC−C3L−00568−TP
n=221
CPTAC3−LSCC−C3L−00603−TP
n=169
CPTAC3−LSCC−C3L−00904−TP
n=280
CPTAC3−LSCC−C3L−00923−TP
n=141
CPTAC3−LSCC−C3L−00927−TP
n=187
CPTAC3−LSCC−C3L−00965−TP
n=272
CPTAC3−LSCC−C3L−00993−TP
n=229
CPTAC3−LSCC−C3L−01000−TP
n=367
CPTAC3−LSCC−C3L−01285−TP
n=226

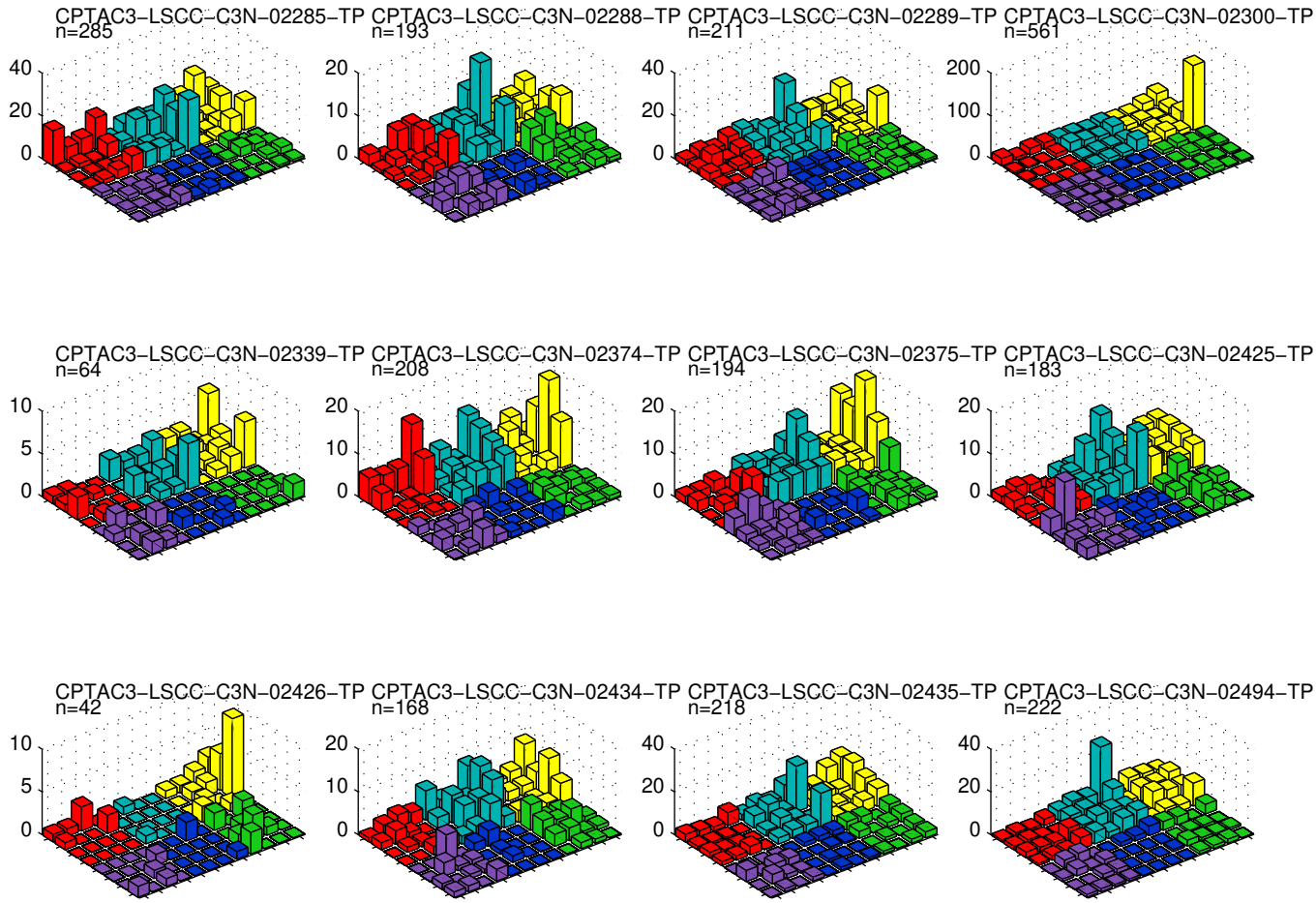

CPTAC3−LSCC−C3L−01455−TP
n=149
CPTAC3−LSCC−C3L−01606−TP
n=274
CPTAC3−LSCC−C3L−01663−TP
n=335
CPTAC3−LSCC−C3L−01838−TP
n=104
CPTAC3−LSCC−C3L−01884−TP
n=280
CPTAC3−LSCC−C3L−02127−TP
n=162
CPTAC3−LSCC−C3L−02130−TP
n=267
CPTAC3−LSCC−C3L−02163−TP
n=201
CPTAC3−LSCC−C3L−02164−TP
n=1
CPTAC3−LSCC−C3L−02168−TP
n=357
CPTAC3−LSCC−C3L−02170−TP
n=74
CPTAC3−LSCC−C3L−02349−TP
n=349

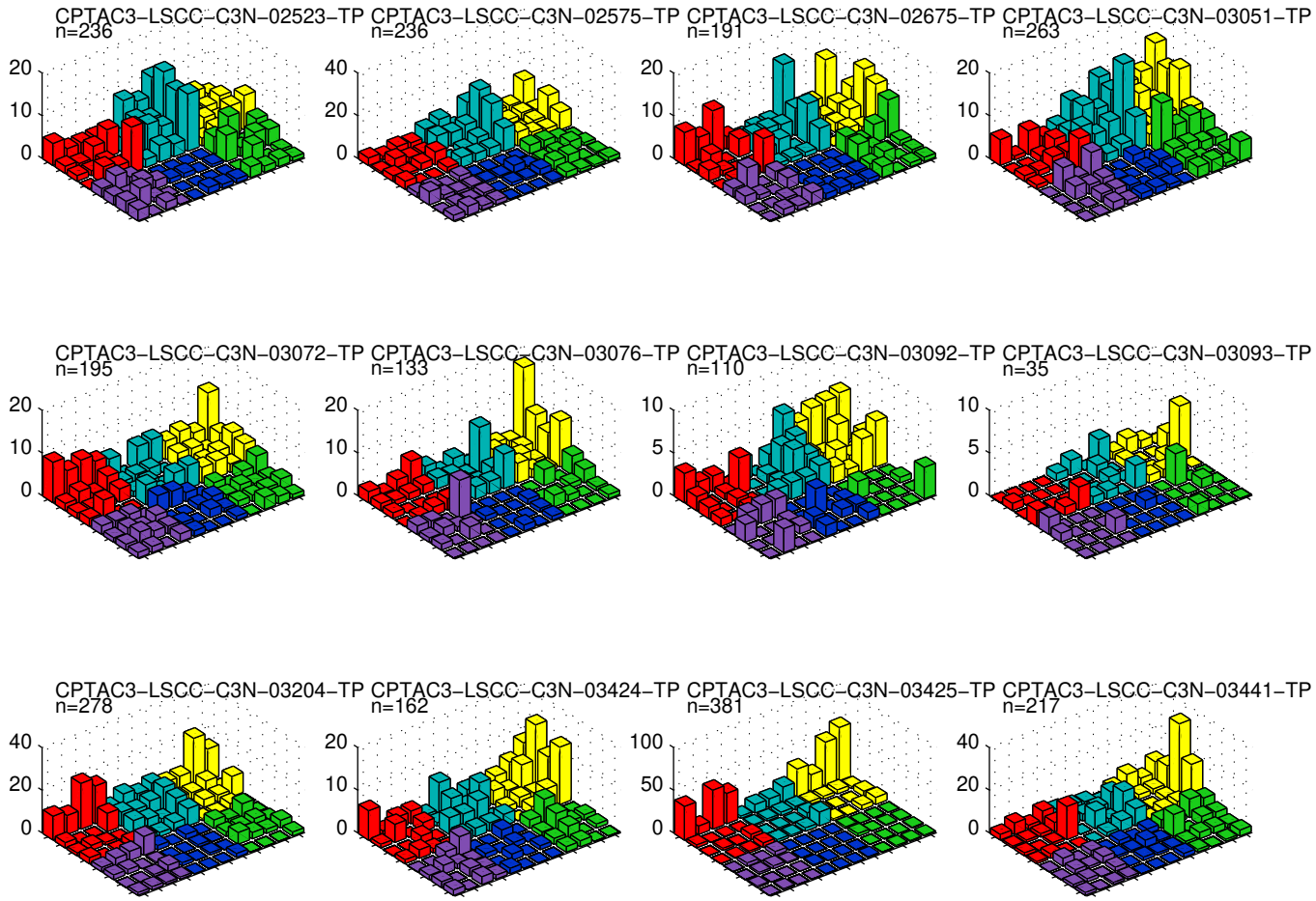

CPTAC3−LSCC−C3L−02358−TP
n=190
CPTAC3−LSCC−C3L−02546−TP
n=189
CPTAC3−LSCC−C3L−02552−TP
n=58
CPTAC3−LSCC−C3L−02619−TP
n=210
CPTAC3−LSCC−C3L−02624−TP
n=231
CPTAC3−LSCC−C3L−02625−TP
n=320
CPTAC3−LSCC−C3L−02627−TP
n=470
CPTAC3−LSCC−C3L−02629−TP
n=67
CPTAC3−LSCC−C3L−02646−TP
n=272
CPTAC3−LSCC−C3L−02648−TP
n=228
CPTAC3−LSCC−C3L−02649−TP
n=264
CPTAC3−LSCC−C3L−02650−TP
n=187

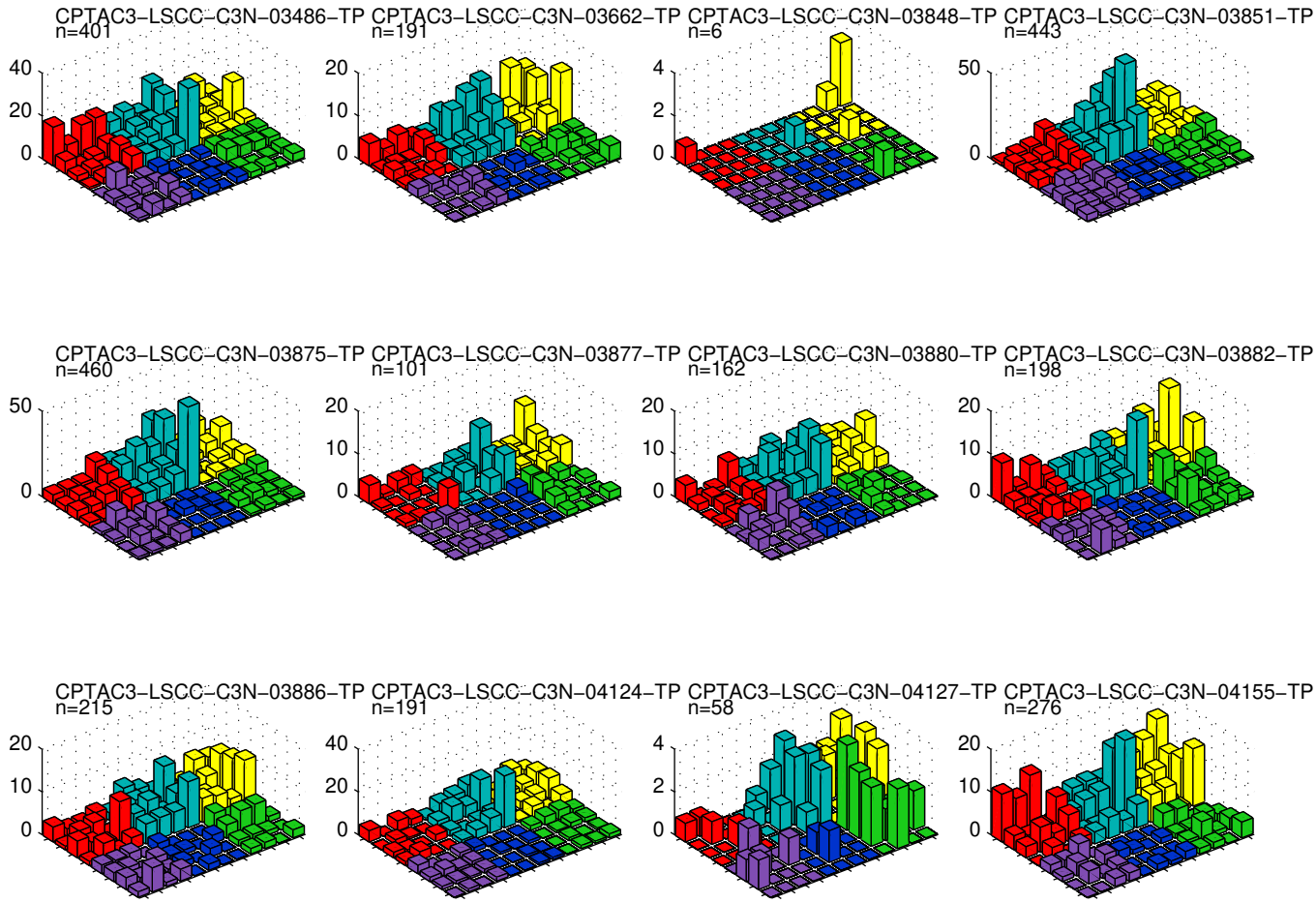

CPTAC3−LSCC−C3L−02660−TP
n=262
40
20
0
CPTAC3−LSCC−C3L−02665−TP
n=478
50
0
CPTAC3−LSCC−C3L−02669−TP
n=477
50
0
CPTAC3−LSCC−C3L−02891−TP
n=85
10
5
0
CPTAC3−LSCC−C3L−02951−TP
n=350
40
20
0
CPTAC3−LSCC−C3L−02963−TP
n=130
20
10
0
CPTAC3−LSCC−C3L−02964−TP
n=161
40
20
0
CPTAC3−LSCC−C3L−02968−TP
n=205
20
10
0
CPTAC3−LSCC−C3L−02969−TP
n=216
40
20
0
CPTAC3−LSCC−C3L−03272−TP
n=222
20
10
0
CPTAC3−LSCC−C3L−03678−TP
n=360
40
20
0
CPTAC3−LSCC−C3L−03961−TP
n=56
10
5
0

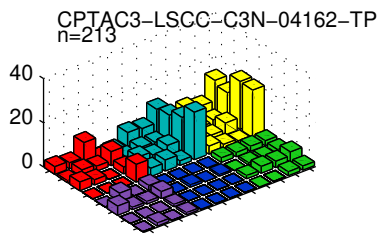

CPTAC3−LSCC−C3L−03962−TP
n=207
CPTAC3−LSCC−C3L−03963−TP
n=172
CPTAC3−LSCC−C3L−03965−TP
n=360
CPTAC3−LSCC−C3L−04013−TP
n=142
CPTAC3−LSCC−C3L−04014−TP
n=256
CPTAC3−LSCC−C3L−04071−TP
n=97
CPTAC3−LSCC−C3L−04391−TP
n=171
CPTAC3−LSCC−C3N−00211−TP
n=28
CPTAC3−LSCC−C3N−00221−TP
n=254
CPTAC3−LSCC−C3N−00247−TP
n=31
CPTAC3−LSCC−C3N−00497−TP
n=265
CPTAC3−LSCC−C3N−00555−TP
n=193

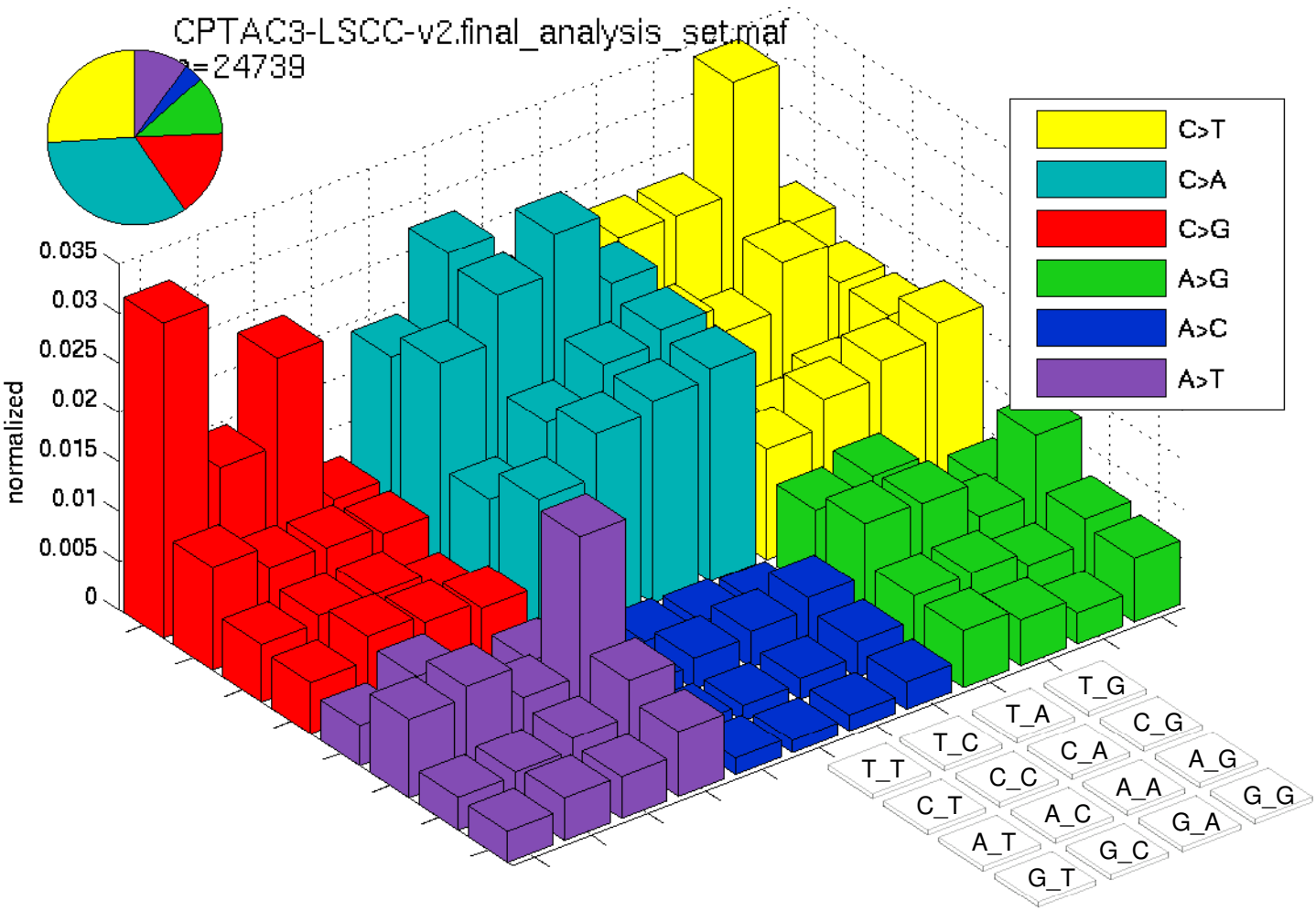

CPTAC3−LSCC−C3N−01017−TP
n=204
CPTAC3−LSCC−C3N−01020−TP
n=150
CPTAC3−LSCC−C3N−01025−TP
n=199
CPTAC3−LSCC−C3N−01028−TP
n=574
CPTAC3−LSCC−C3N−01194−TP
n=253
CPTAC3−LSCC−C3N−01411−TP
n=353
CPTAC3−LSCC−C3N−01846−TP
n=124
CPTAC3−LSCC−C3N−01892−TP
n=251
CPTAC3−LSCC−C3N−01893−TP
n=468
CPTAC3−LSCC−C3N−02252−TP
n=164
CPTAC3−LSCC−C3N−02283−TP
n=573
CPTAC3−LSCC−C3N−02284−TP
n=443


CPTAC3−LSCC−C3N−02285−TP
n=285
CPTAC3−LSCC−C3N−02288−TP
n=193
CPTAC3−LSCC−C3N−02289−TP
n=211
CPTAC3−LSCC−C3N−02300−TP
n=561
CPTAC3−LSCC−C3N−02339−TP
n=64
CPTAC3−LSCC−C3N−02374−TP
n=208
CPTAC3−LSCC−C3N−02375−TP
n=194
CPTAC3−LSCC−C3N−02425−TP
n=183
CPTAC3−LSCC−C3N−02426−TP
n=42
CPTAC3−LSCC−C3N−02434−TP
n=168
CPTAC3−LSCC−C3N−02435−TP
n=218
CPTAC3−LSCC−C3N−02494−TP
n=222


CPTAC3−LSCC−C3N−02523−TP
n=236
CPTAC3−LSCC−C3N−02575−TP
n=236
CPTAC3−LSCC−C3N−02675−TP
n=191
CPTAC3−LSCC−C3N−03051−TP
n=263
CPTAC3−LSCC−C3N−03072−TP
n=195
CPTAC3−LSCC−C3N−03076−TP
n=133
CPTAC3−LSCC−C3N−03092−TP
n=110
CPTAC3−LSCC−C3N−03093−TP
n=35
CPTAC3−LSCC−C3N−03204−TP
n=278
CPTAC3−LSCC−C3N−03424−TP
n=162
CPTAC3−LSCC−C3N−03425−TP
n=381
CPTAC3−LSCC−C3N−03441−TP
n=217

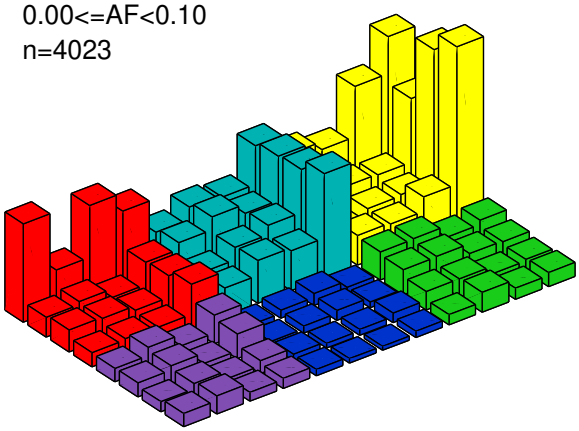

CPTAC3−LSCC−C3N−03486−TP
n=401
40
20
0
CPTAC3−LSCC−C3N−03662−TP
n=191
20
10
0
CPTAC3−LSCC−C3N−03848−TP
n=6
4
2
0
CPTAC3−LSCC−C3N−03851−TP
n=443
50
0
CPTAC3−LSCC−C3N−03875−TP
n=460
50
0
CPTAC3−LSCC−C3N−03877−TP
n=101
20
10
0
CPTAC3−LSCC−C3N−03880−TP
n=162
20
10
0
CPTAC3−LSCC−C3N−03882−TP
n=198
20
10
0
CPTAC3−LSCC−C3N−03886−TP
n=215
20
10
0
CPTAC3−LSCC−C3N−04124−TP
n=191
40
20
0
CPTAC3−LSCC−C3N−04127−TP
n=58
4
2
0
CPTAC3−LSCC−C3N−04155−TP
n=276
20
10
0


CPTAC3−LSCC−C3N−04162−TP
n=213
40
20
0

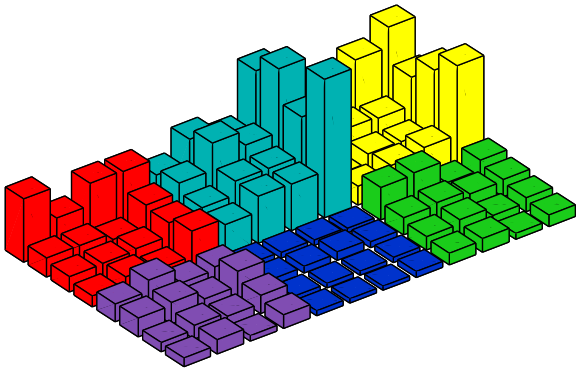

CPTAC3-LSCC-v2.final_analysis_set.maf
=24739
normalized
0.035
0.03
0.025
0.02
0.015
0.01
0.005
0
C>T
C>A
C>G
A>G
A>C
A>T
T_G
T_A
C_G
T_C
C_A
A_G
T_T
C_C
A_A
G_G
C_T
A_C
G_A
A_T
G_C
G_T

CPTAC3−LSCC−v2.final_analysis_set.maf

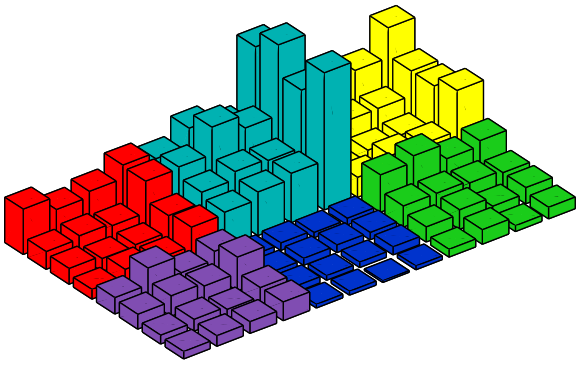
0.00<=AF<0.10
n=4023



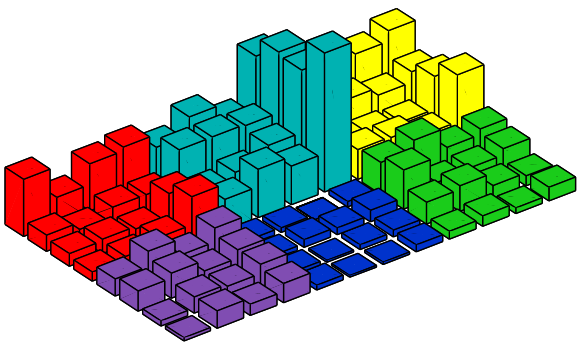
0.10<=AF<0.25
n=11819

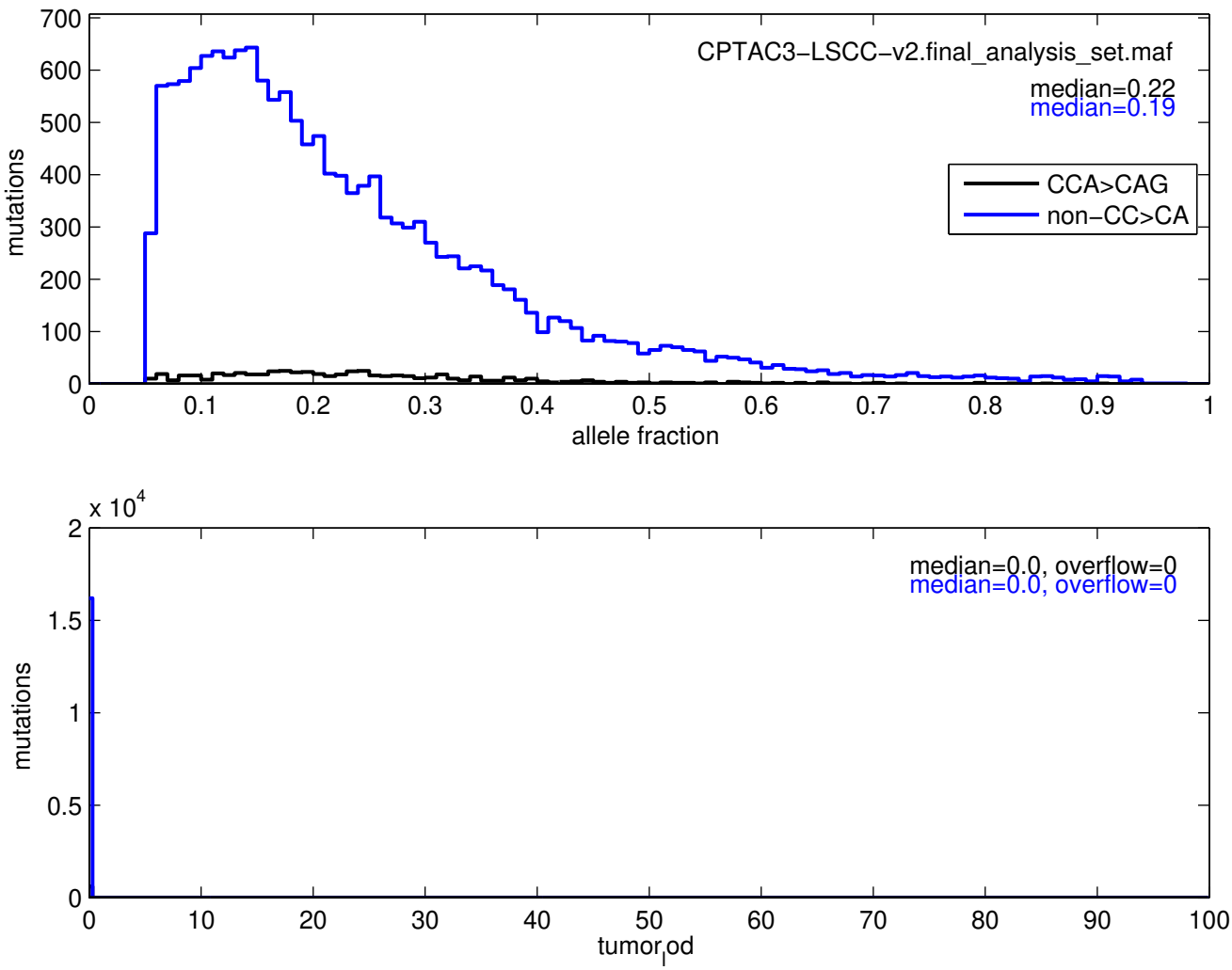


0.25<=AF<0.50
n=7234

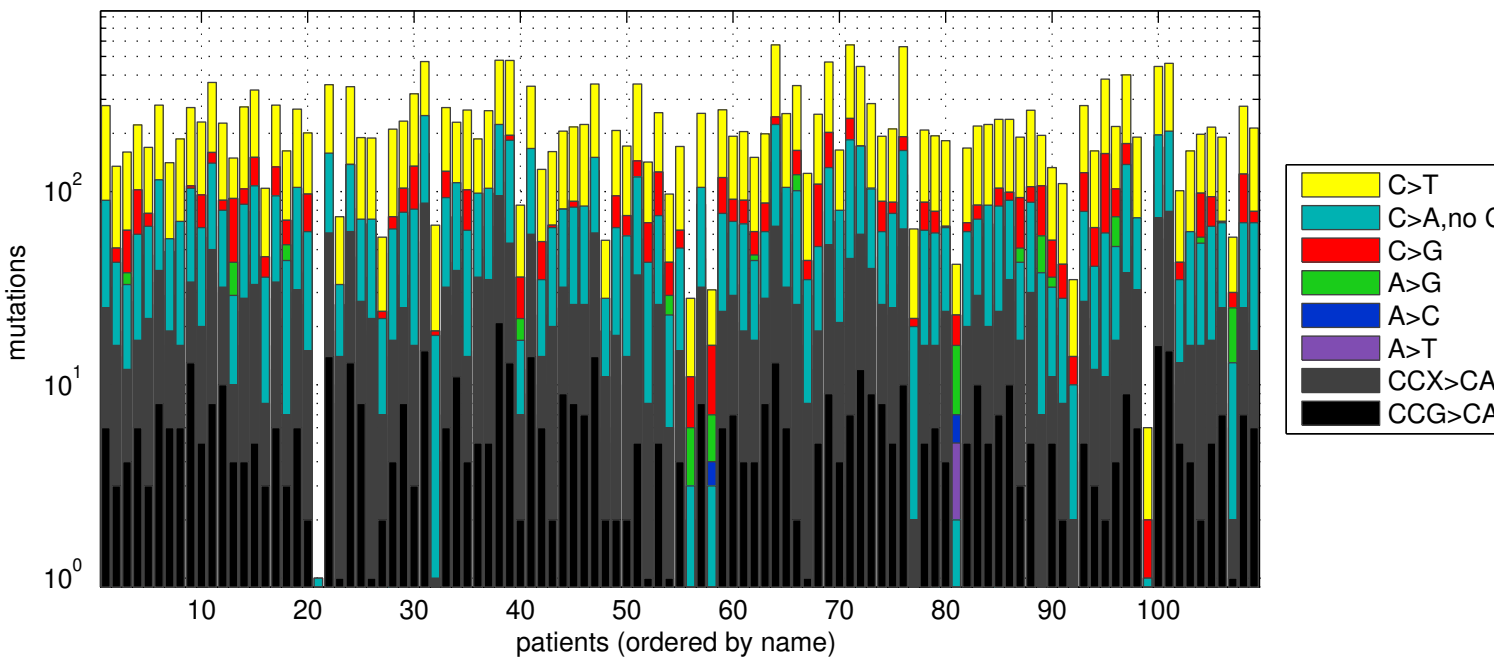


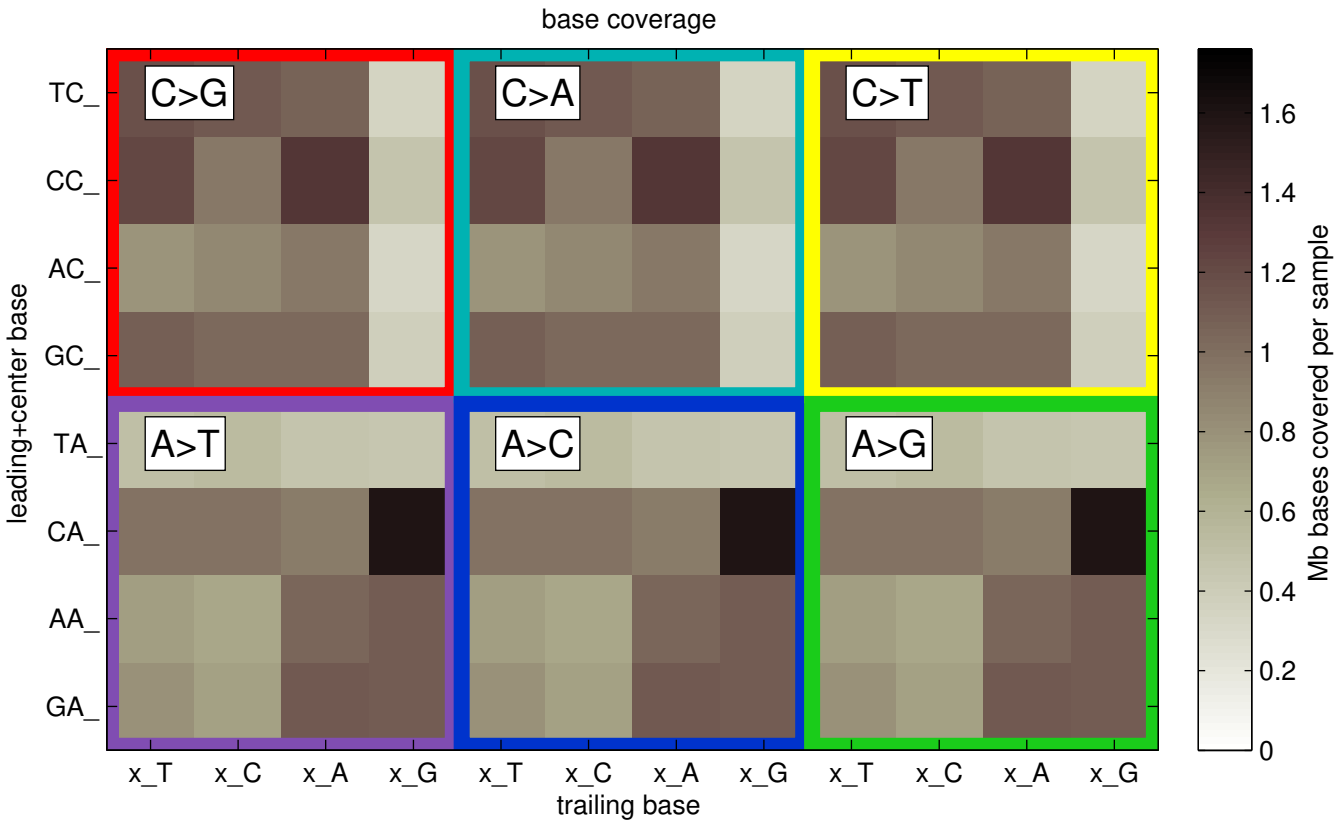
0.50<=AF<1.00
n=1662




CPTAC3−LSCC−v2.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
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