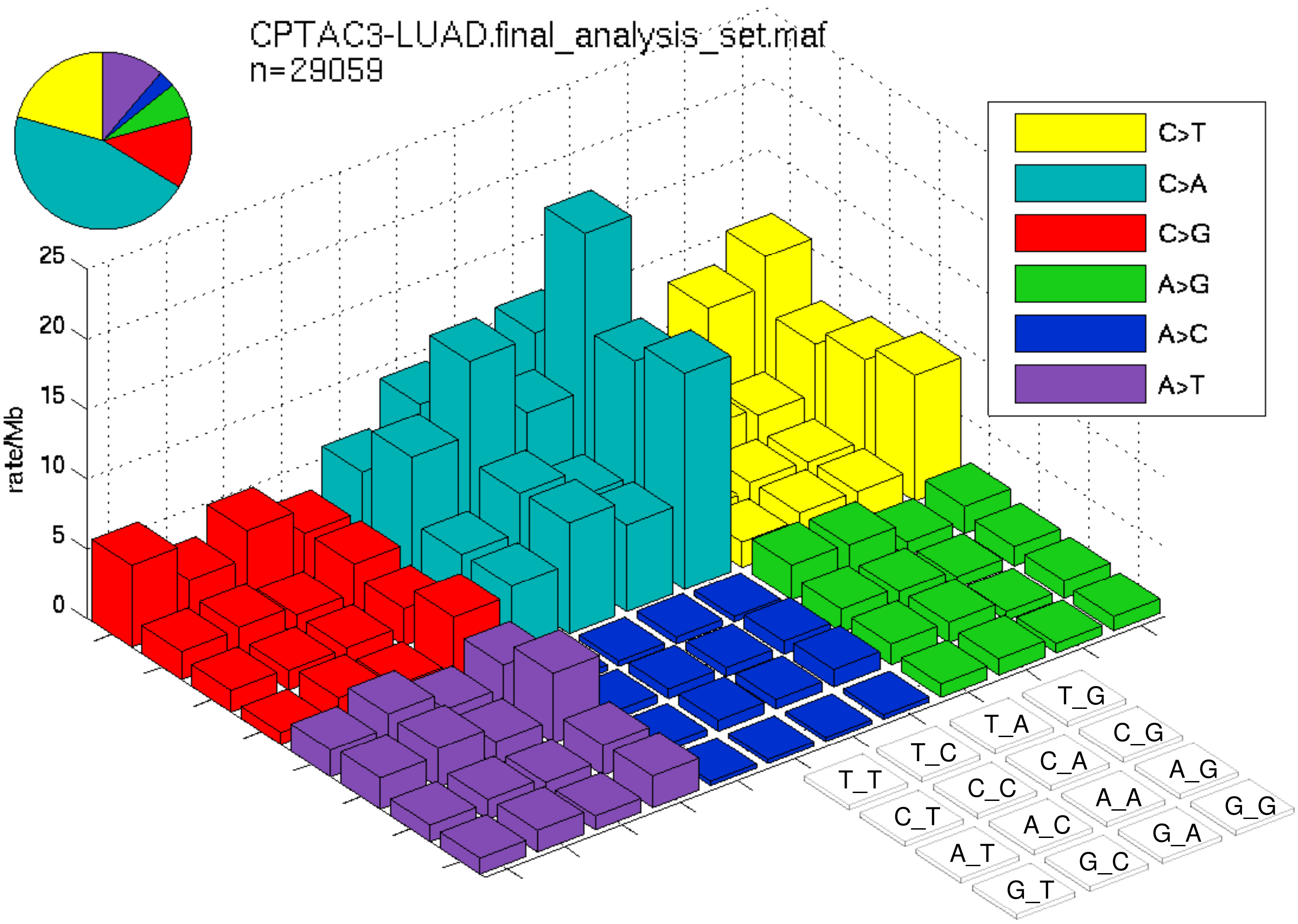
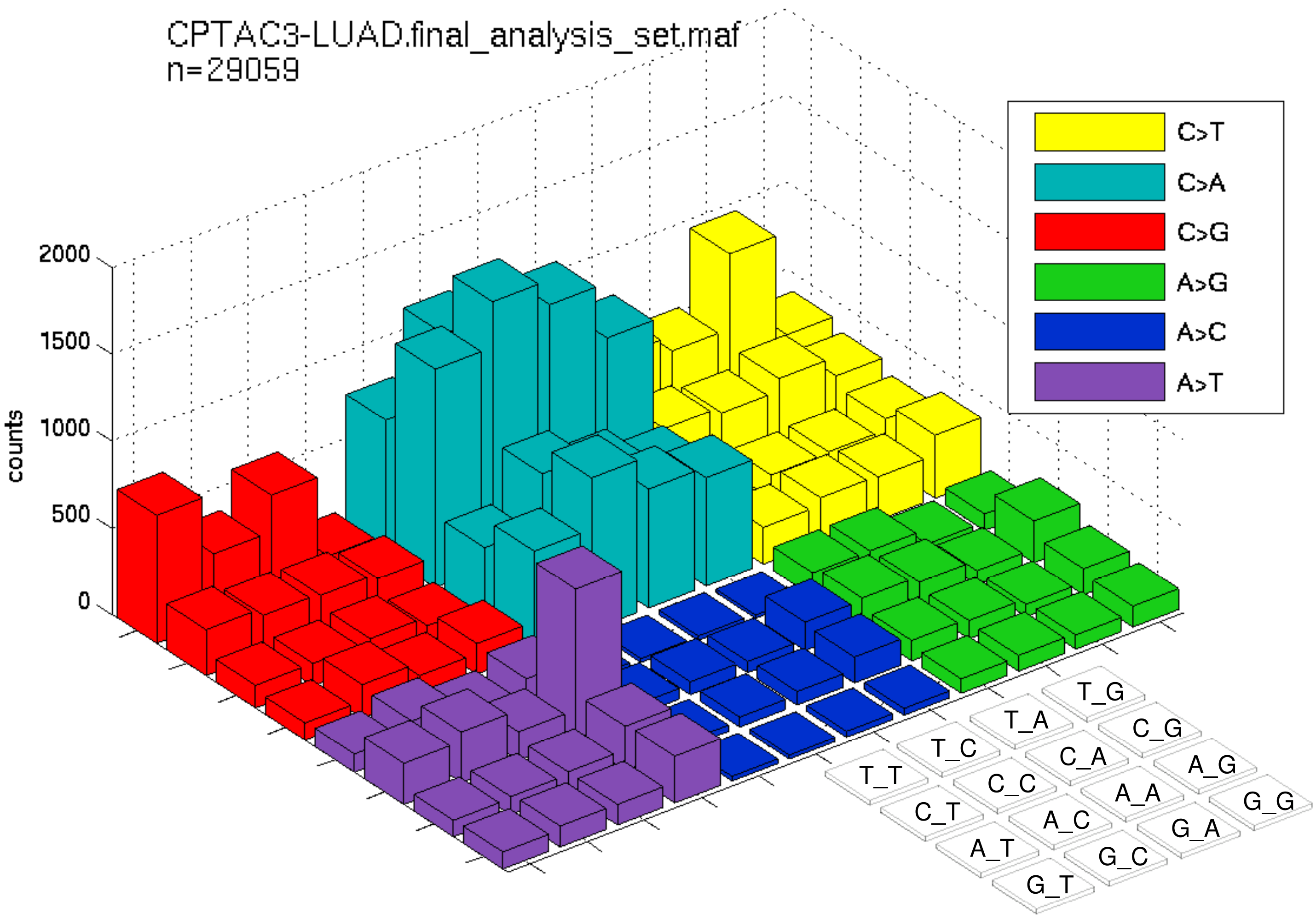
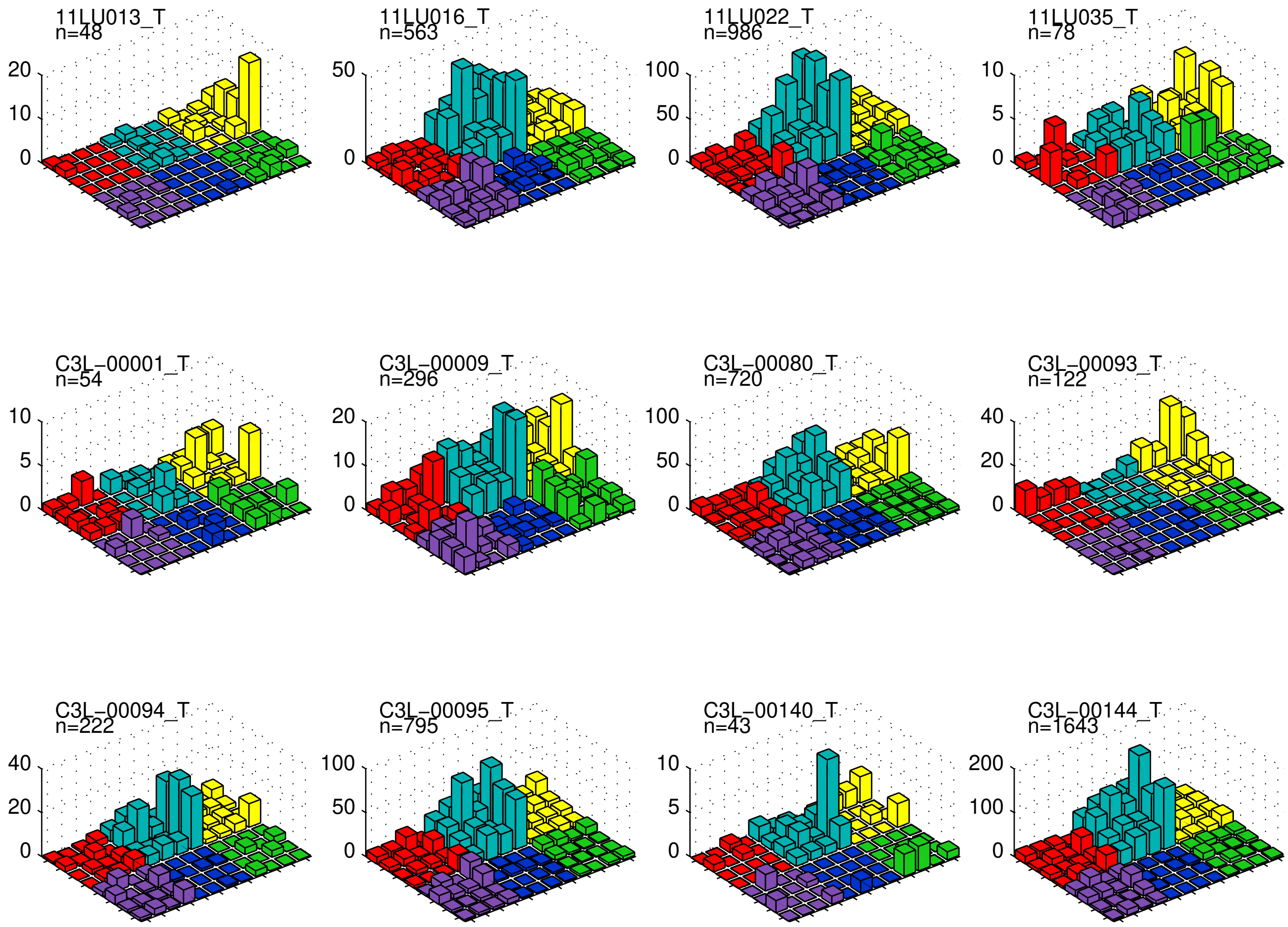
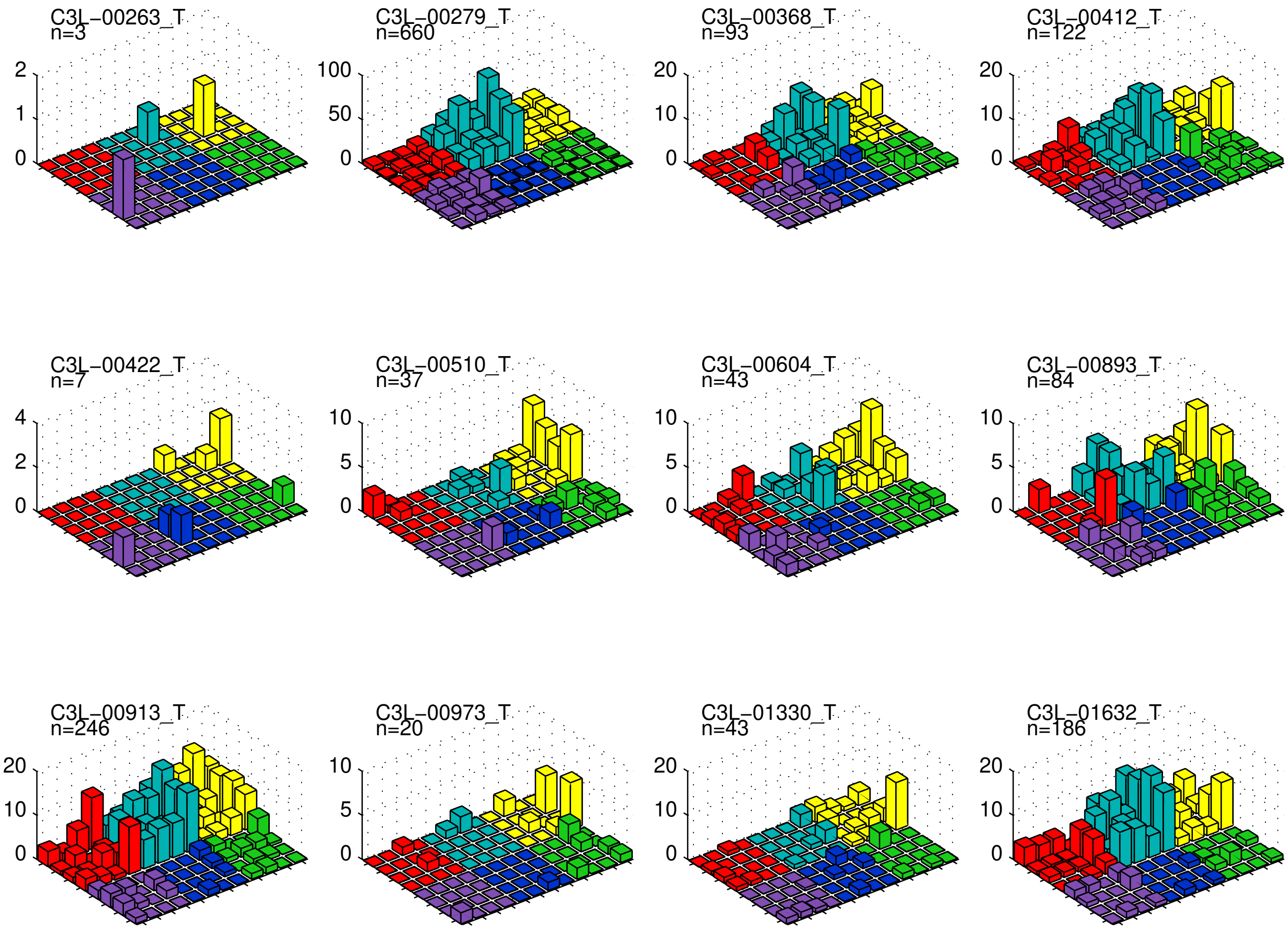
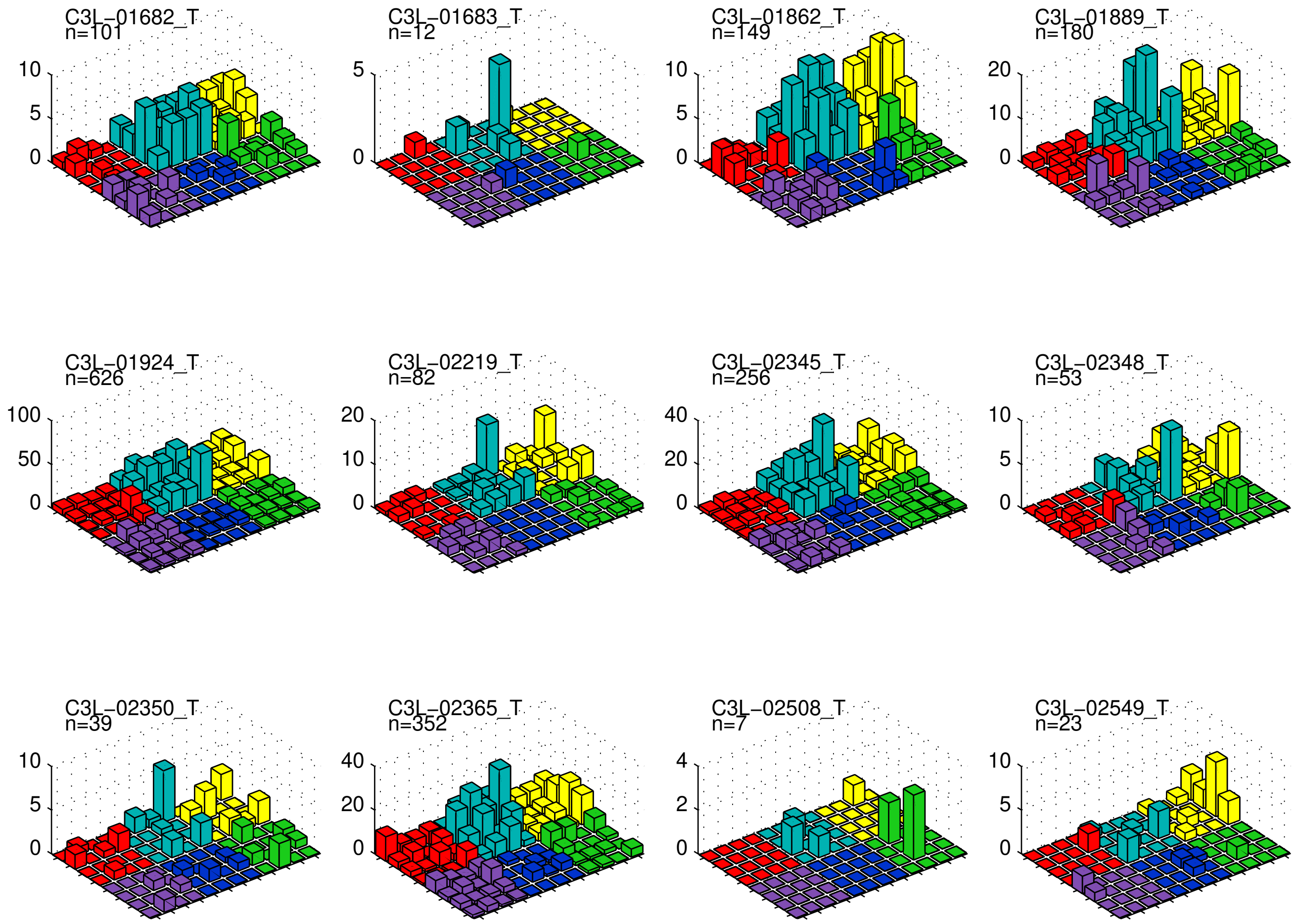


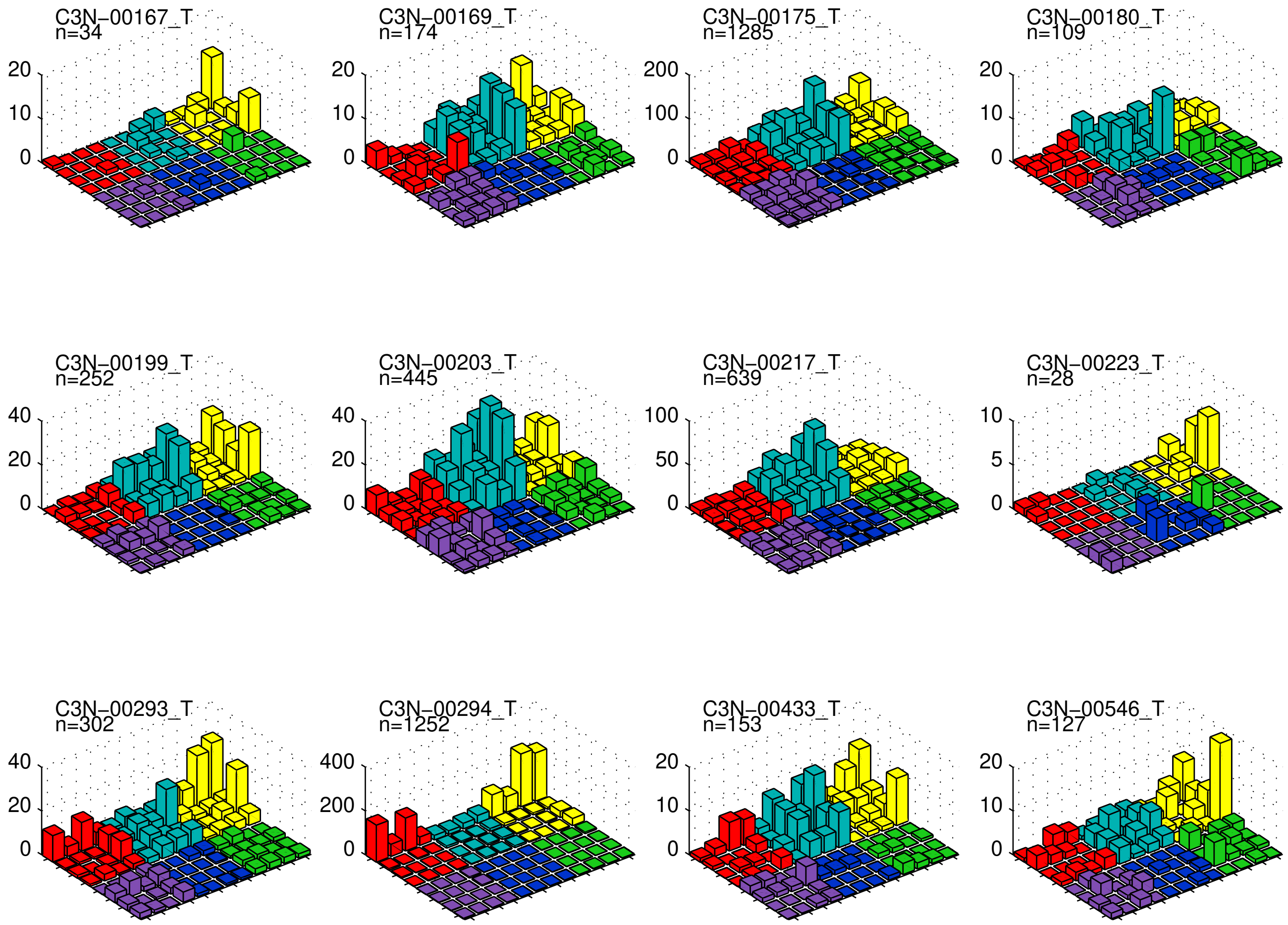
CPTAC3-LUAD.final_analysis_set.maf
n=29059
rate/Mb
25
20
15
10
5
0
C>T
C>A
C>G
A>G
A>C
A>T
T_G
T_A
C_G
T_C
C_A
A_G
T_T
C_C
A_A
G_G
C_T
A_C
G_A
A_T
G_C
G_T

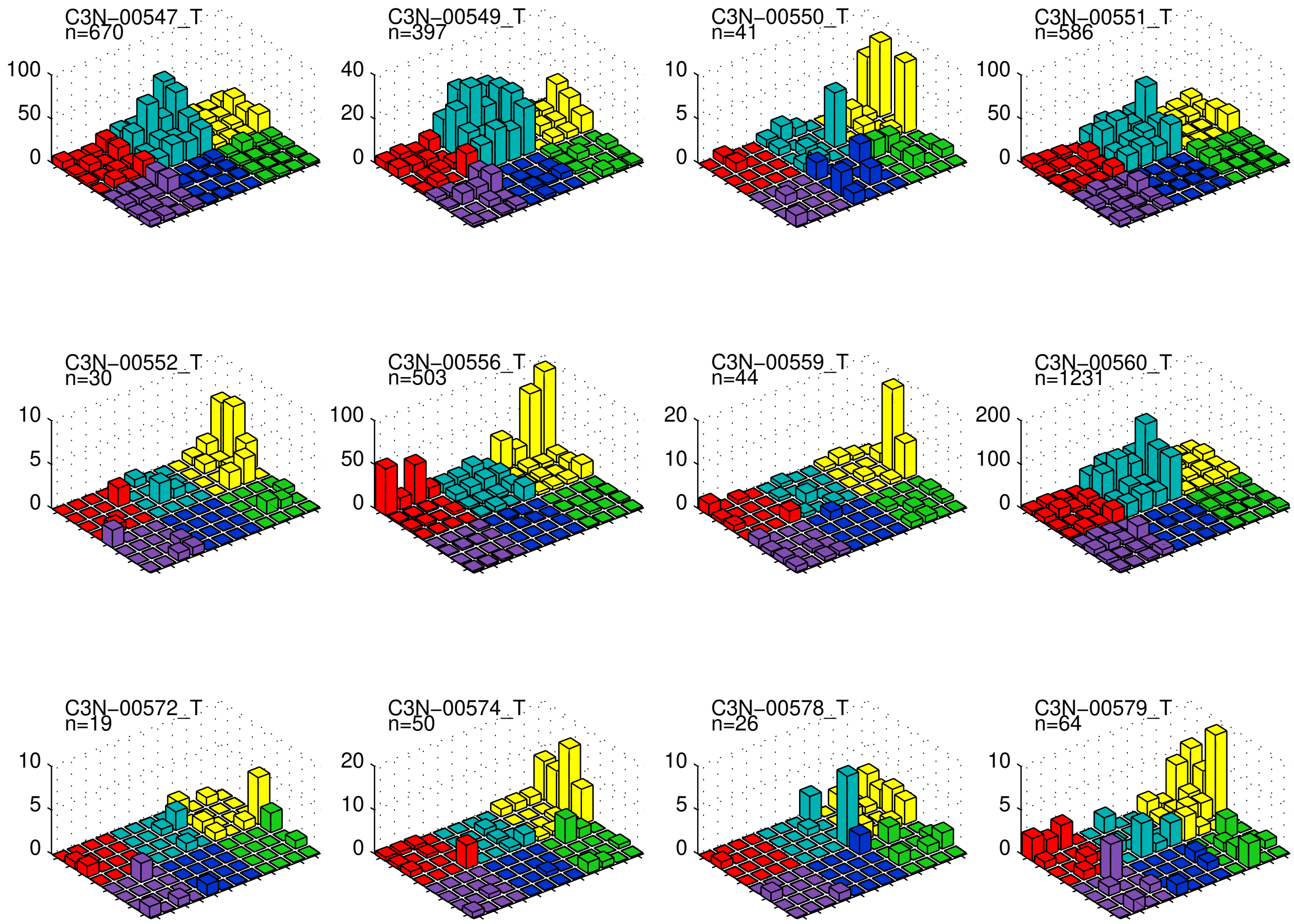

CPTAC3-LUAD.final_analysis_set.maf
n=29059
counts
2000
1500
1000
500
0
C>T
C>A
C>G
A>G
A>C
A>T
T_G
T_A
C_G
T_C
C_A
A_G
T_T
C_C
A_A
G_G
C_T
A_C
G_A
A_T
G_C
G_T

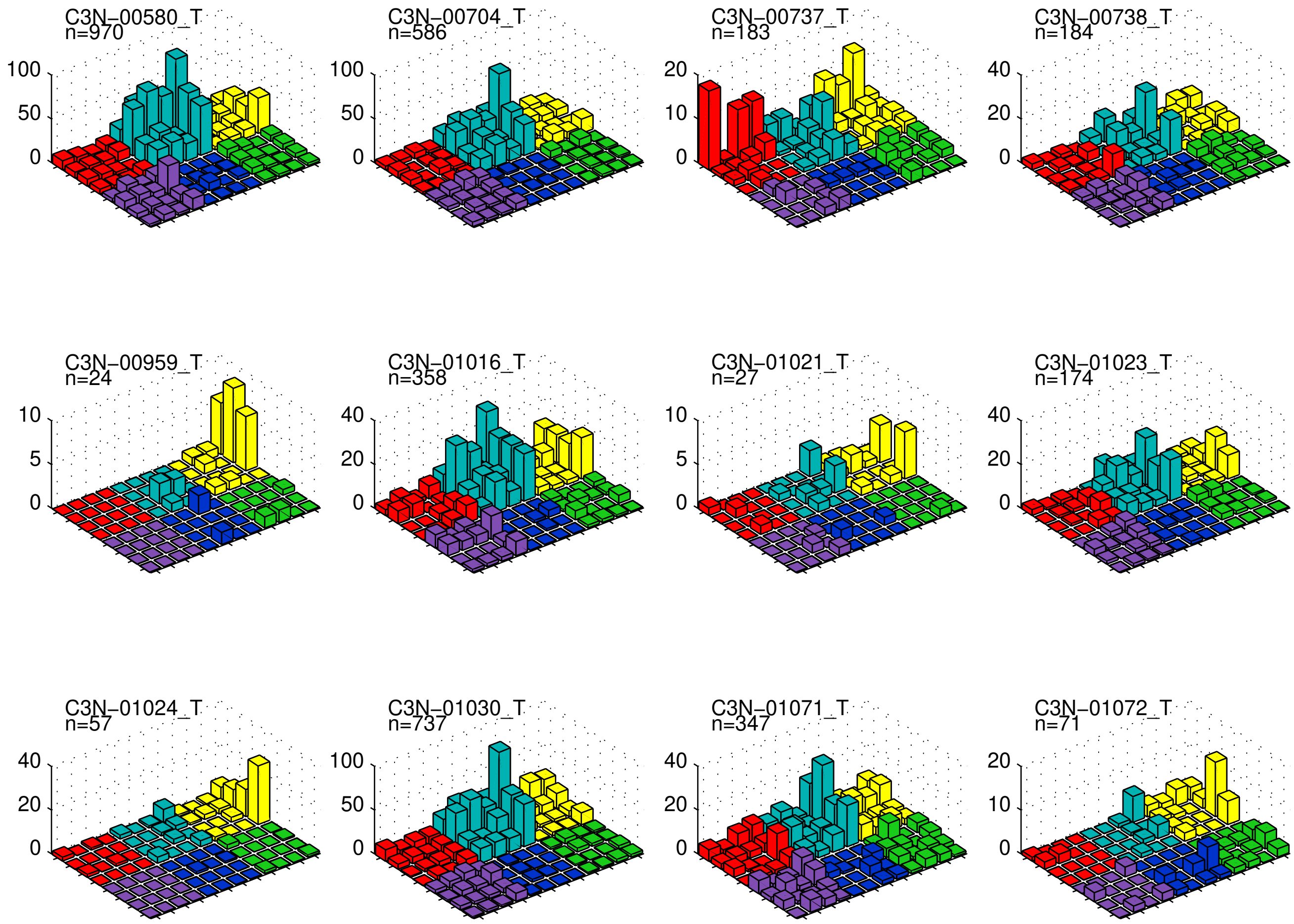

11LU013_T
n=48
11LU016_T
n=563
11LU022_T
n=986
11LU035_T
n=78
C3L−00001_T
n=54
C3L−00009_T
n=296
C3L−00080_T
n=720
C3L−00093_T
n=122
C3L−00094_T
n=222
C3L−00095_T
n=795
C3L−00140_T
n=43
C3L−00144_T
n=1643

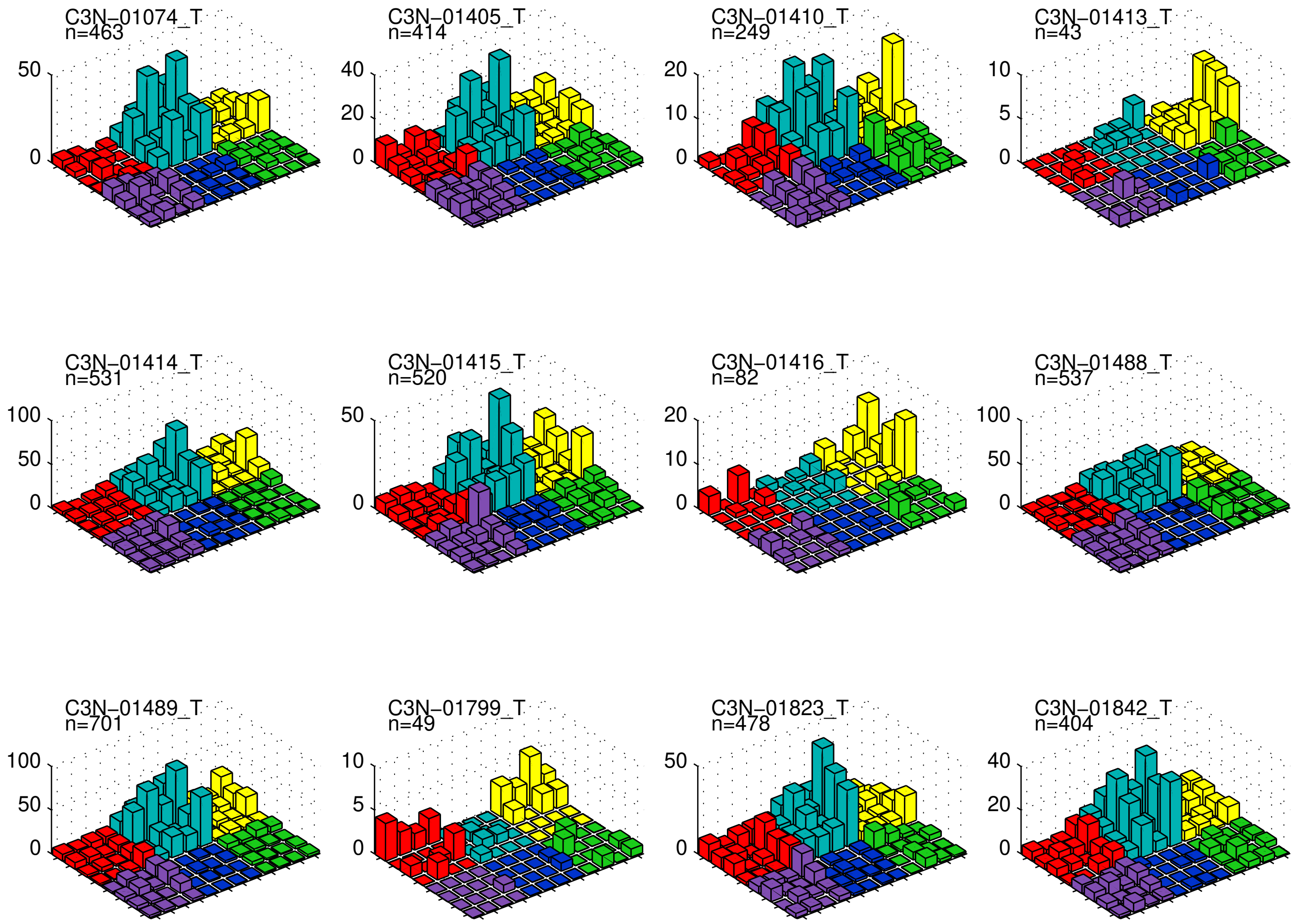

C3L−00263_T
n=3
C3L−00279_T
n=660
C3L−00368_T
n=93
C3L−00412_T
n=122
C3L−00422_T
n=7
C3L−00510_T
n=37
C3L−00604_T
n=43
C3L−00893_T
n=84
C3L−00913_T
n=246
C3L−00973_T
n=20
C3L−01330_T
n=43
C3L−01632_T
n=186


C3L−01682_T
n=101
C3L−01683_T
n=12
C3L−01862_T
n=149
C3L−01889_T
n=180
C3L−01924_T
n=626
C3L−02219_T
n=82
C3L−02345_T
n=256
C3L−02348_T
n=53
C3L−02350_T
n=39
C3L−02365_T
n=352
C3L−02508_T
n=7
C3L−02549_T
n=23

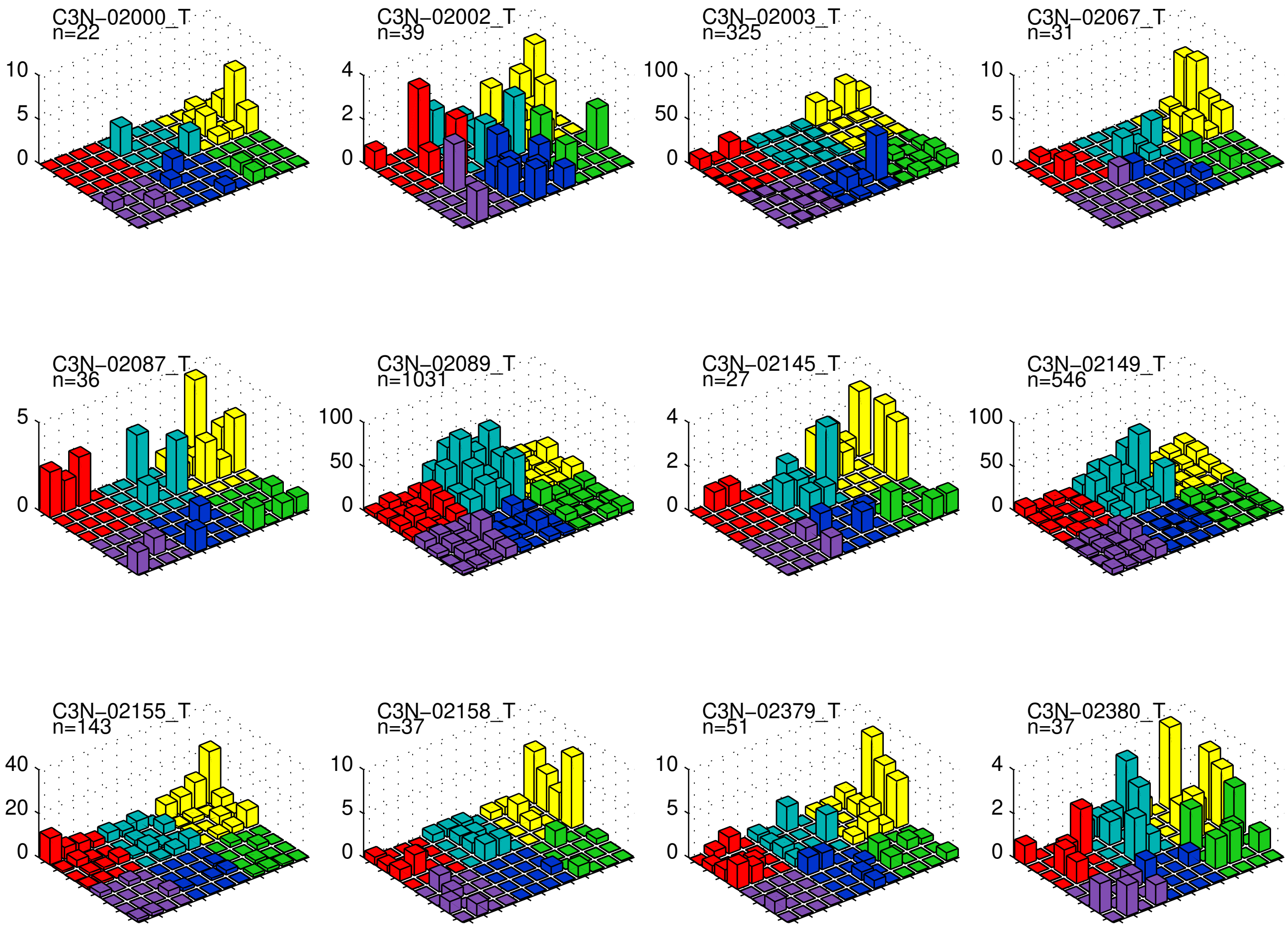

C3N–00167_T
n=34
C3N–00169_T
n=174
C3N–00175_T
n=1285
C3N–00180_T
n=109
C3N–00199_T
n=252
C3N–00203_T
n=445
C3N–00217_T
n=639
C3N–00223_T
n=28
C3N–00293_T
n=302
C3N–00294_T
n=1252
C3N–00433_T
n=153
C3N–00546_T
n=127

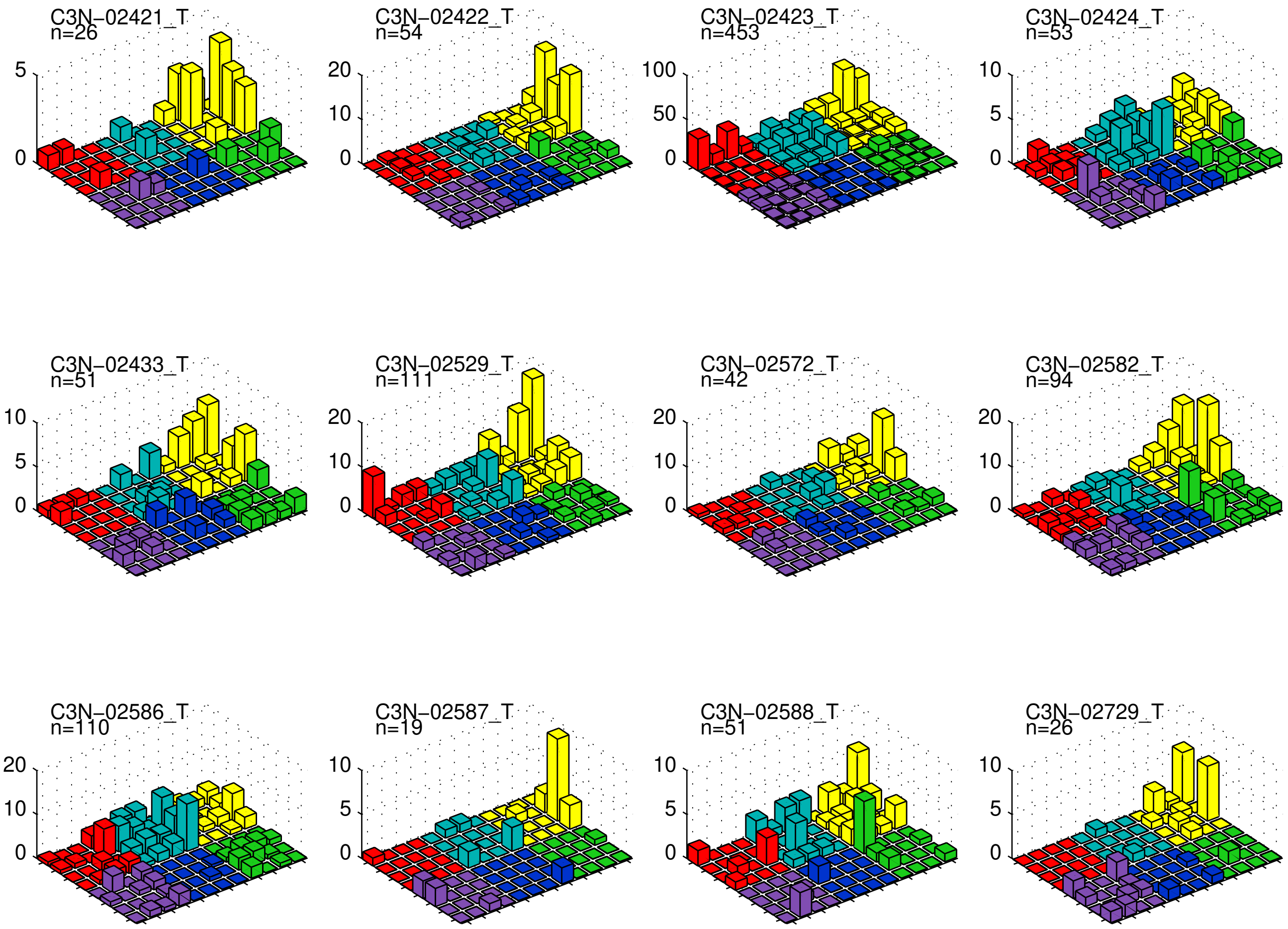

C3N–00547_T
n=670
100
50
0
C3N–00549_T
n=397
40
20
0
C3N–00550_T
n=41
10
5
0
C3N–00551_T
n=586
100
50
0
C3N–00552_T
n=30
10
5
0
C3N–00556_T
n=503
100
50
0
C3N–00559_T
n=44
20
10
0
C3N–00560_T
n=1231
200
100
0
C3N–00572_T
n=19
10
5
0
C3N–00574_T
n=50
20
10
0
C3N–00578_T
n=26
10
5
0
C3N–00579_T
n=64
10
5
0

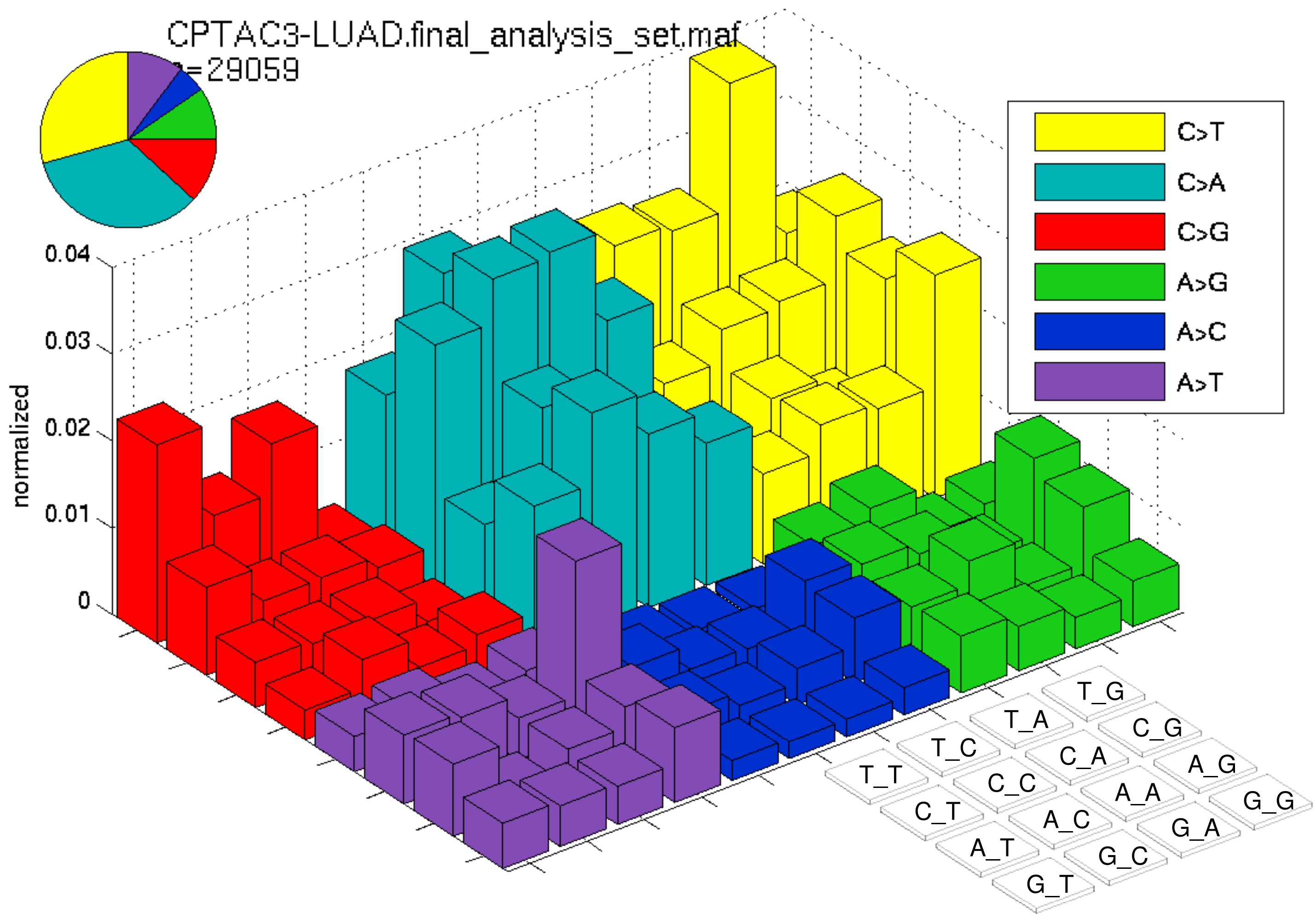

C3N-00580_T
n=970
100
50
0
C3N-00704_T
n=586
100
50
0
C3N-00737_T
n=183
20
10
0
C3N-00738_T
n=184
40
20
0
C3N-00959_T
n=24
10
5
0
C3N-01016_T
n=358
40
20
0
C3N-01021_T
n=27
10
5
0
C3N-01023_T
n=174
40
20
0
C3N-01024_T
n=57
40
20
0
C3N-01030_T
n=737
100
50
0
C3N-01071_T
n=347
40
20
0
C3N-01072_T
n=71
20
10
0



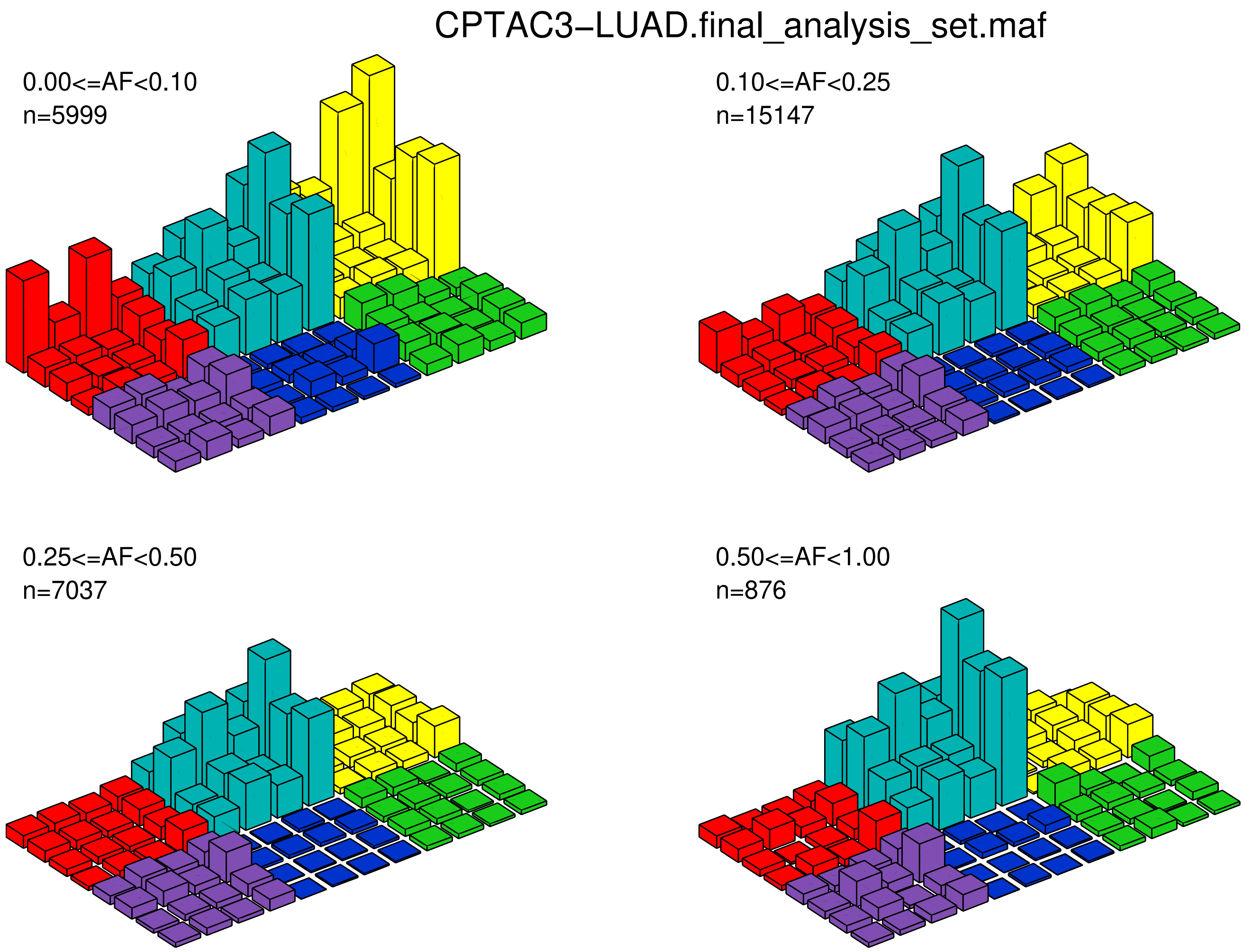
C3N−01074_T
n=463
50
0
C3N−01405_T
n=414
40
20
0
C3N−01410_T
n=249
20
10
0
C3N−01413_T
n=43
10
5
0
C3N−01414_T
n=531
100
50
0
C3N−01415_T
n=520
50
0
C3N−01416_T
n=82
20
10
0
C3N−01488_T
n=537
100
50
0
C3N−01489_T
n=701
100
50
0
C3N−01799_T
n=49
10
5
0
C3N−01823_T
n=478
50
0
C3N−01842_T
n=404
40
20
0

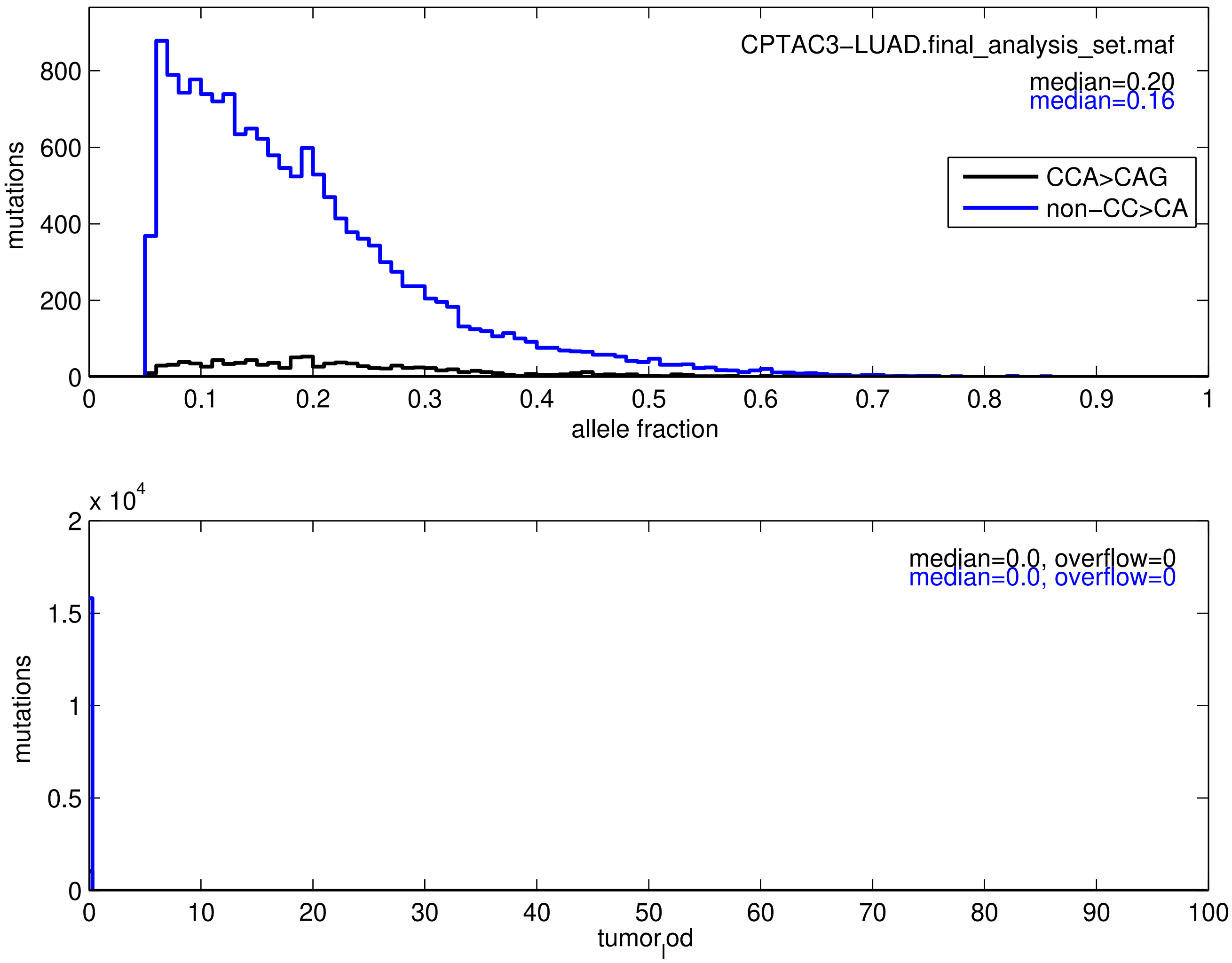

C3N−02000_T
n=22
C3N−02002_T
n=39
C3N−02003_T
n=325
C3N−02067_T
n=31
C3N−02087_T
n=36
C3N−02089_T
n=1031
C3N−02145_T
n=27
C3N−02149_T
n=546
C3N−02155_T
n=143
C3N−02158_T
n=37
C3N−02379_T
n=51
C3N−02380_T
n=37

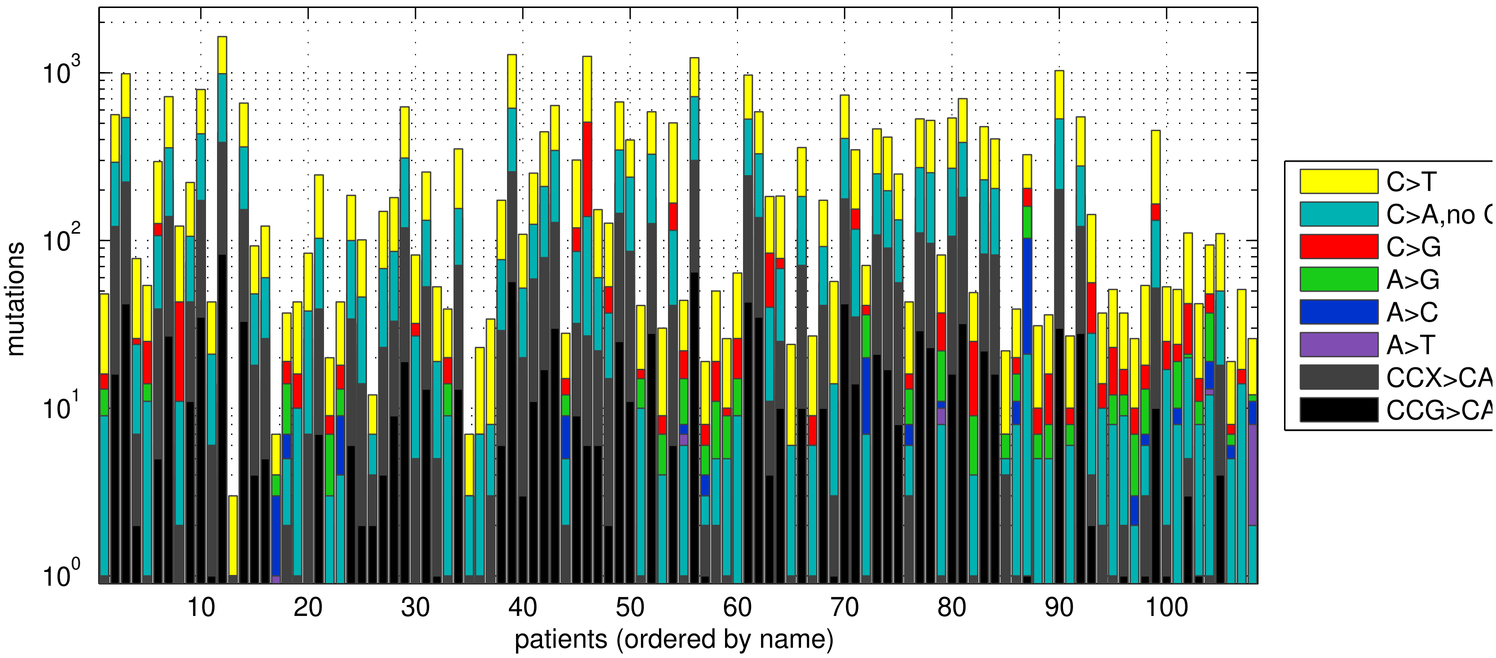

C3N−02421_T
n=26
C3N−02422_T
n=54
C3N−02423_T
n=453
C3N−02424_T
n=53
C3N−02433_T
n=51
C3N−02529_T
n=111
C3N−02572_T
n=42
C3N−02582_T
n=94
C3N−02586_T
n=110
C3N−02587_T
n=19
C3N−02588_T
n=51
C3N−02729_T
n=26



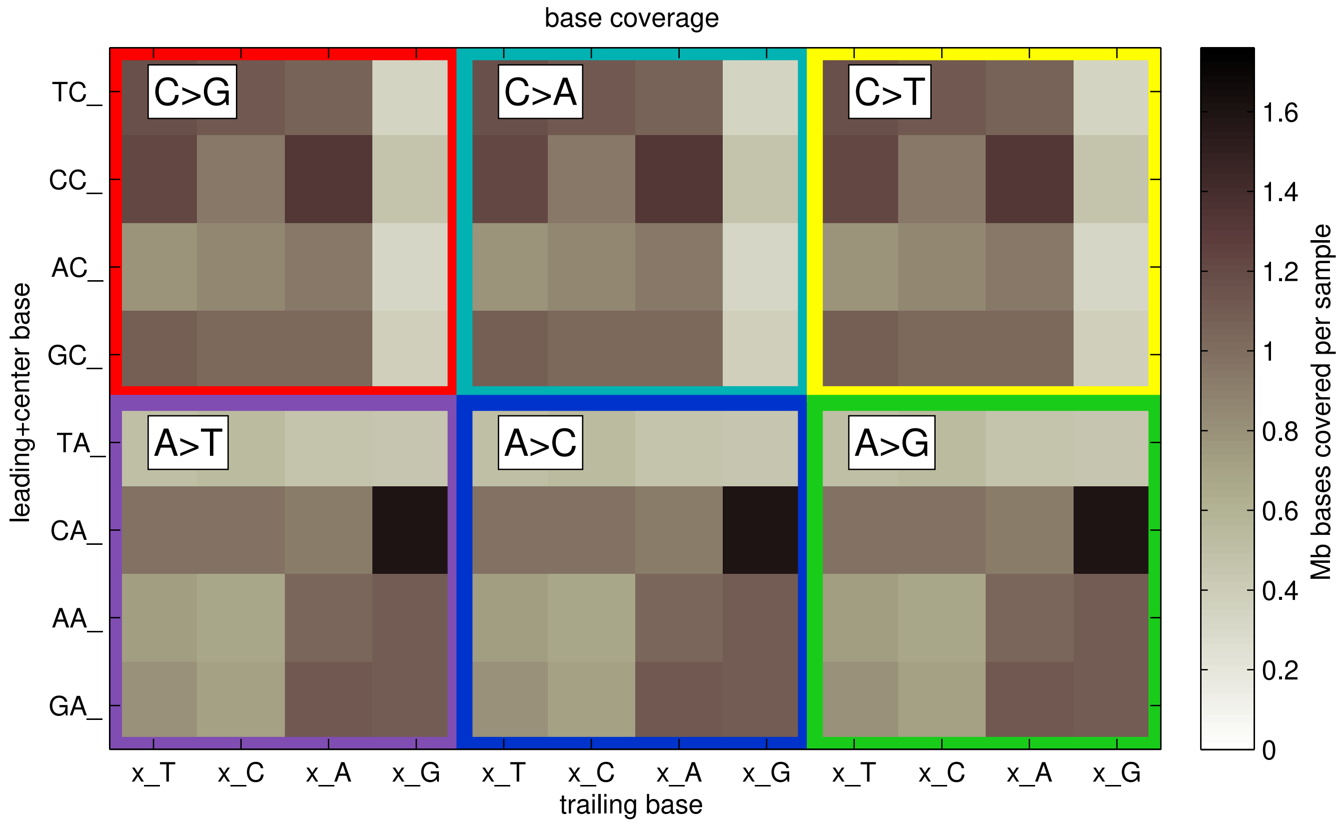
CPTAC3-LUAD.final_analysis_set.maf
n=29059
0.04
0.03
0.02
0.01
0
normalized
C>T
C>A
C>G
A>G
A>C
A>T
T_G
T_A
C_G
T_C
C_A
A_G
T_T
C_C
A_A
G_G
C_T
A_C
G_A
A_T
G_C
G_T


CPTAC3−LUAD.final_analysis_set.maf
0.00<=AF<0.10
n=5999
0.10<=AF<0.25
n=15147
0.25<=AF<0.50
n=7037
0.50<=AF<1.00
n=876


CPTAC3−LUAD.final_analysis_set.maf
median=0.20
median=0.16
CCA>CAG
non−CC>CA
mutations
allele fraction
x 10^4
median=0.0, overflow=0
median=0.0, overflow=0
mutations
tumor_lod



mutations
$10^3$
$10^2$
$10^1$
$10^0$
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
0
0.2
0.4
0.6
0.8
1
1.2
1.4
1.6
Mb bases covered per sample