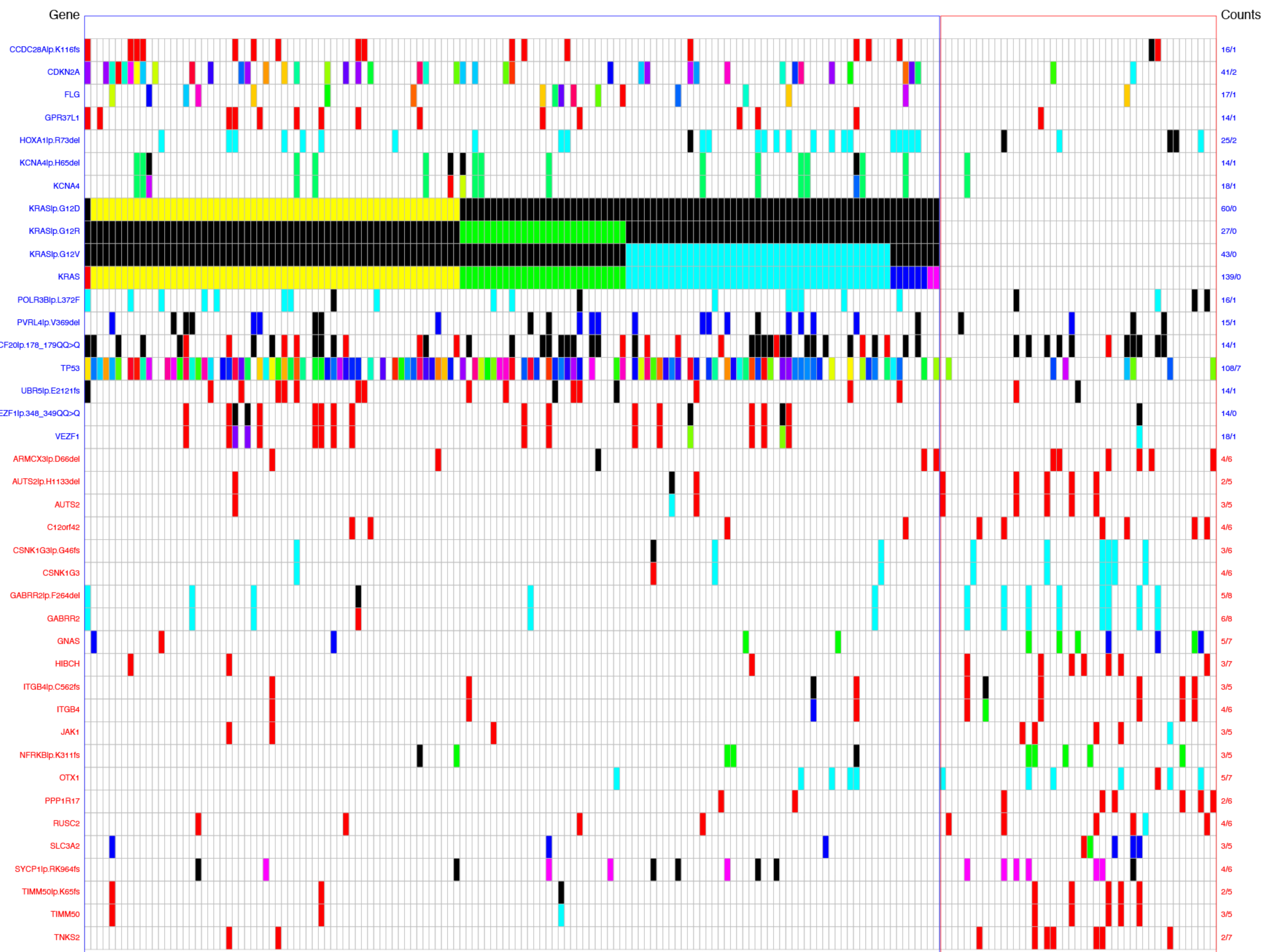
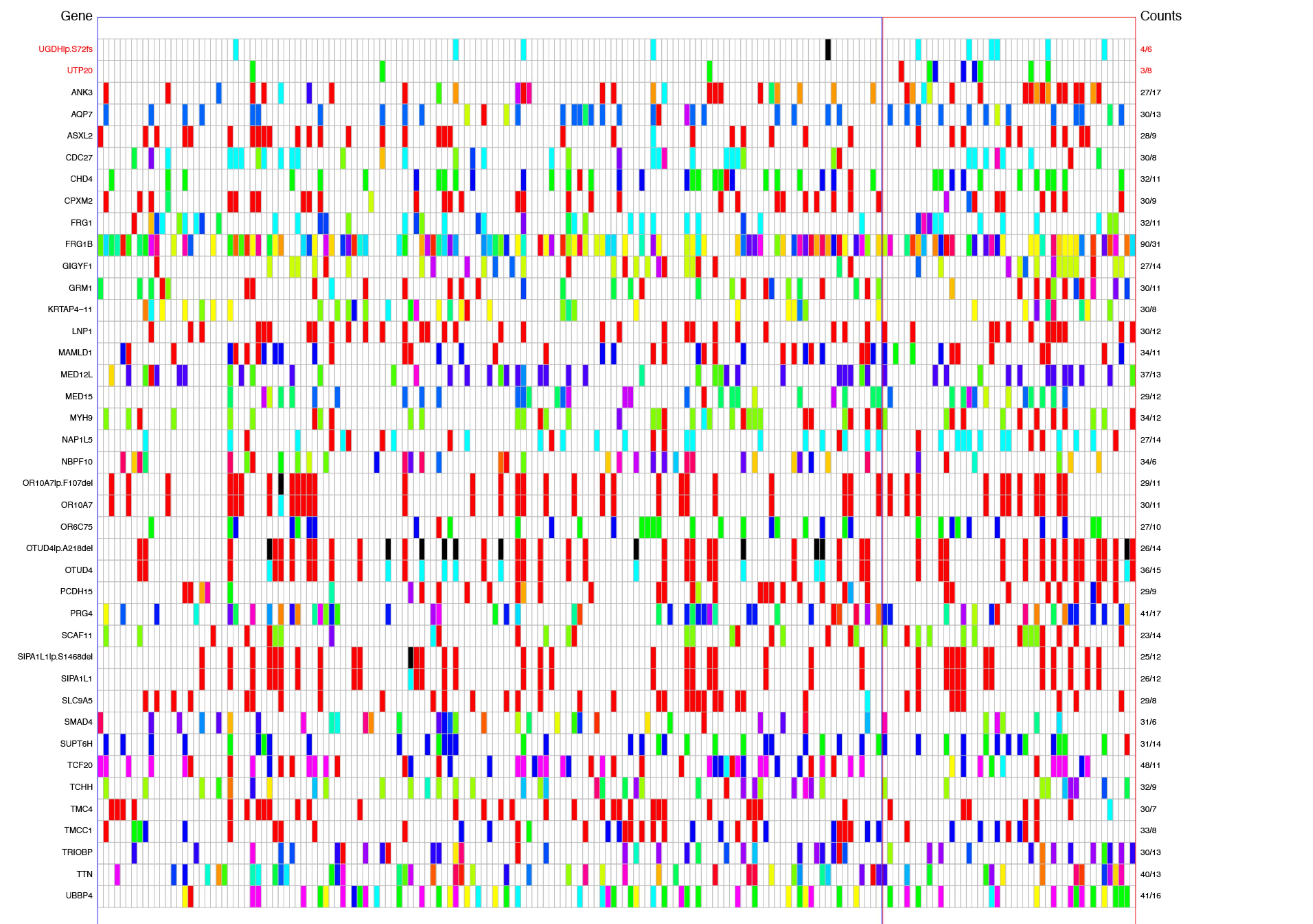


Broad GDAC **pancreas** adenocarcinomas (N=184; 3 pages)



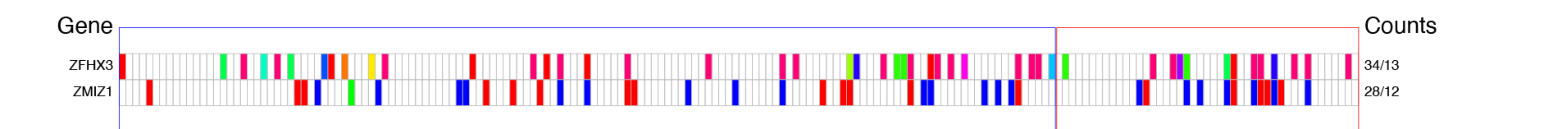
mutants:139

non-mutants:45



mutants:139

non-mutants:45



mutants:139

non-mutants:45

Figure 1A. Mutations in PAAD (pancreas) cancers in the Broad GDAC data.

MSK cBioPortal **pancreas** adenocarcinomas (N=766)

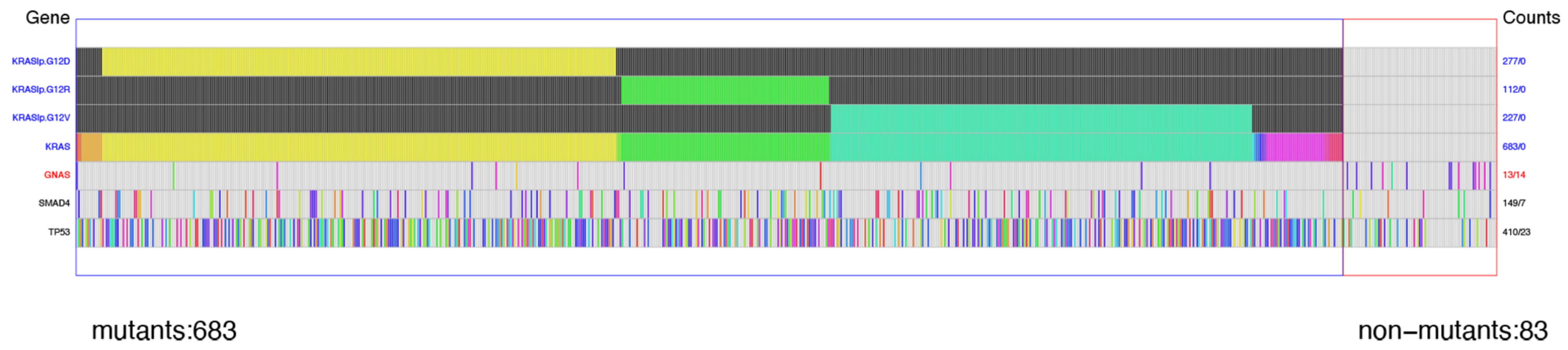
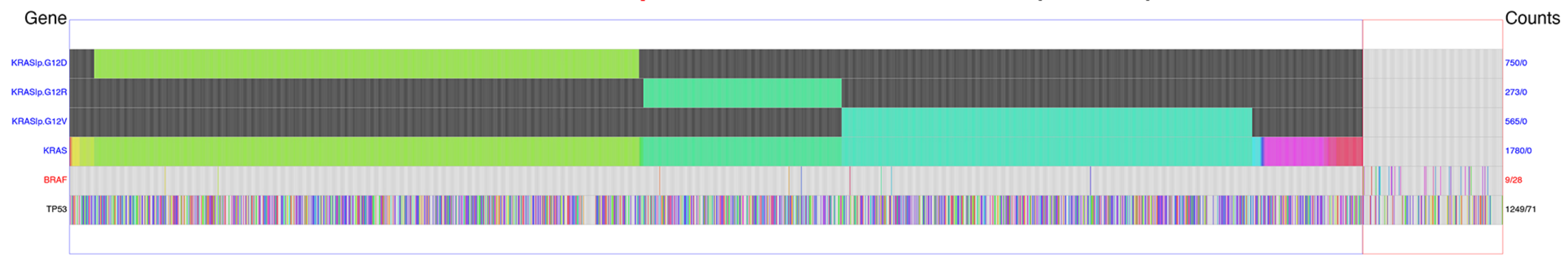


Figure 1B. Mutations in PAAD cancers in the MSK cBioPortal data.

AACR Genie **pancreas** adenocarcinomas (N=1973)

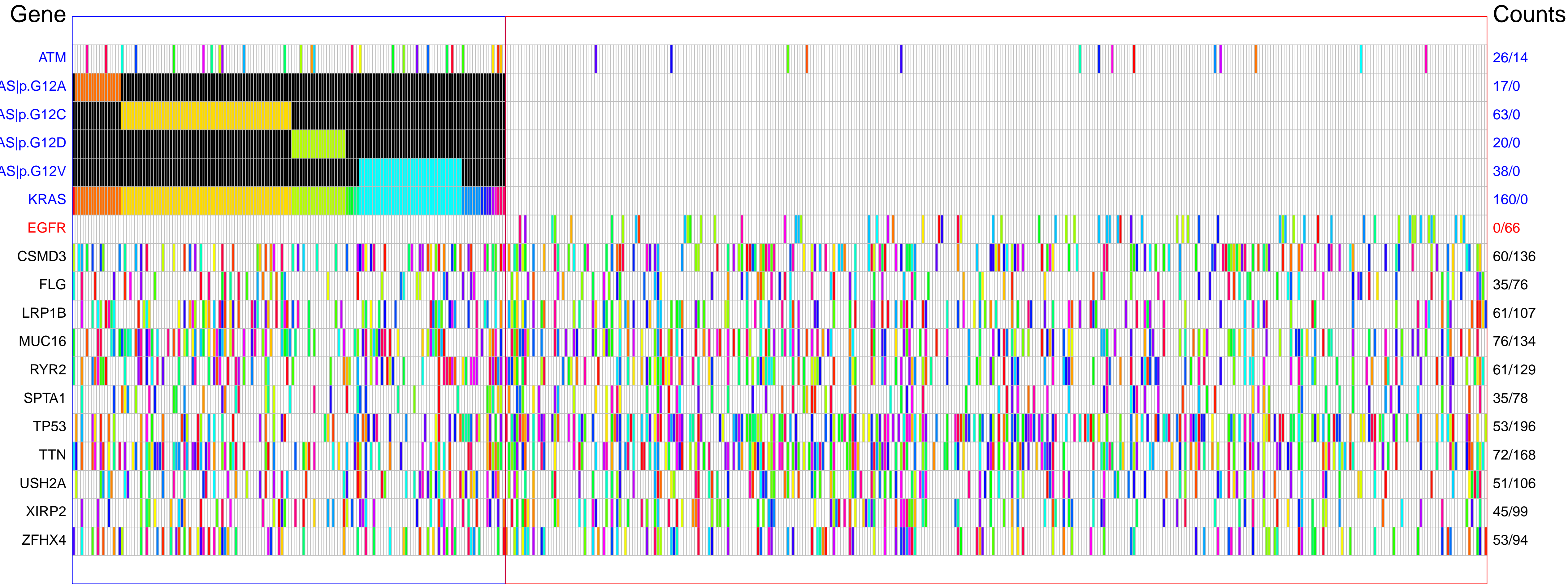


mutants:1780

non-mutants:193

Figure 1C. Mutations in PAAD cancers in the AACR Genie data.

Broad GDAC lung adenocarcinomas (corrected; N=523)

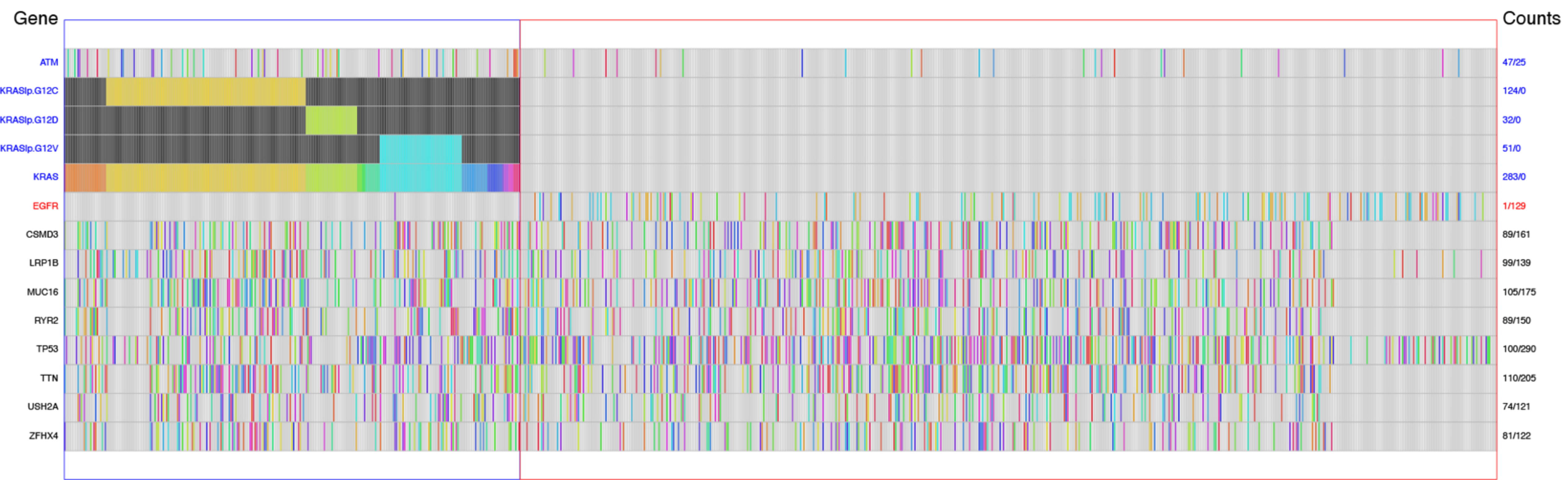


mutants:160

non-mutants:363

Corrected Figure 2A. Mutations in LUAD (lung) cancers in the Broad GDAC data.

MSK cBioPortal lung adenocarcinomas (N=890)

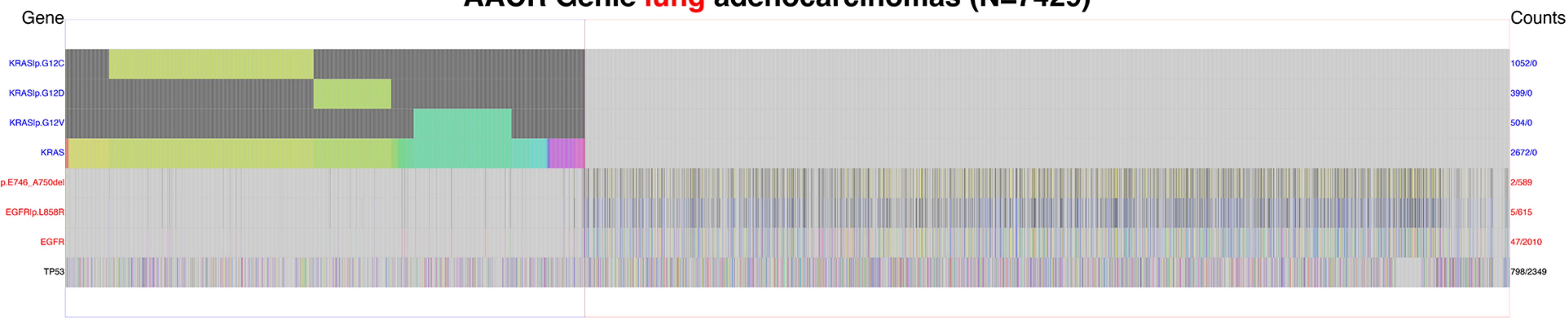


mutants:283

non-mutants:607

Figure 2B. Mutations in LUAD cancers in the MSK cBioPortal data.

AACR Genie lung adenocarcinomas (N=7429)

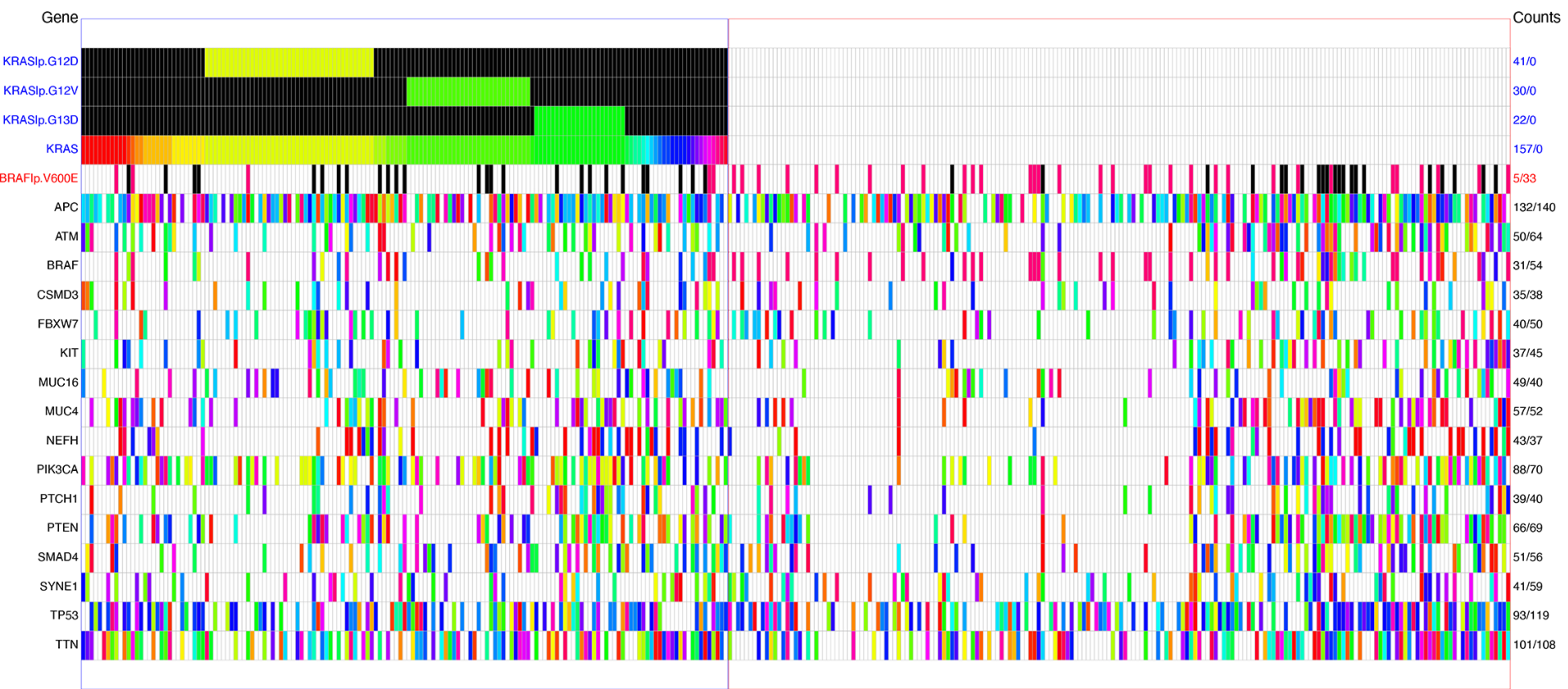


mutants:2672

non-mutants:4757

Figure 2C. Mutations in LUAD cancers in the AACR Genie data.

Broad GDAC **colon** adenocarcinomas (N=347)



mutants:157

non-mutants:190

Figure 3A. Mutations in COAD (colon) cancers is the Broad GDAC data.

MSK cBioPortal colon adenocarcinomas (N=132; 3 pages)

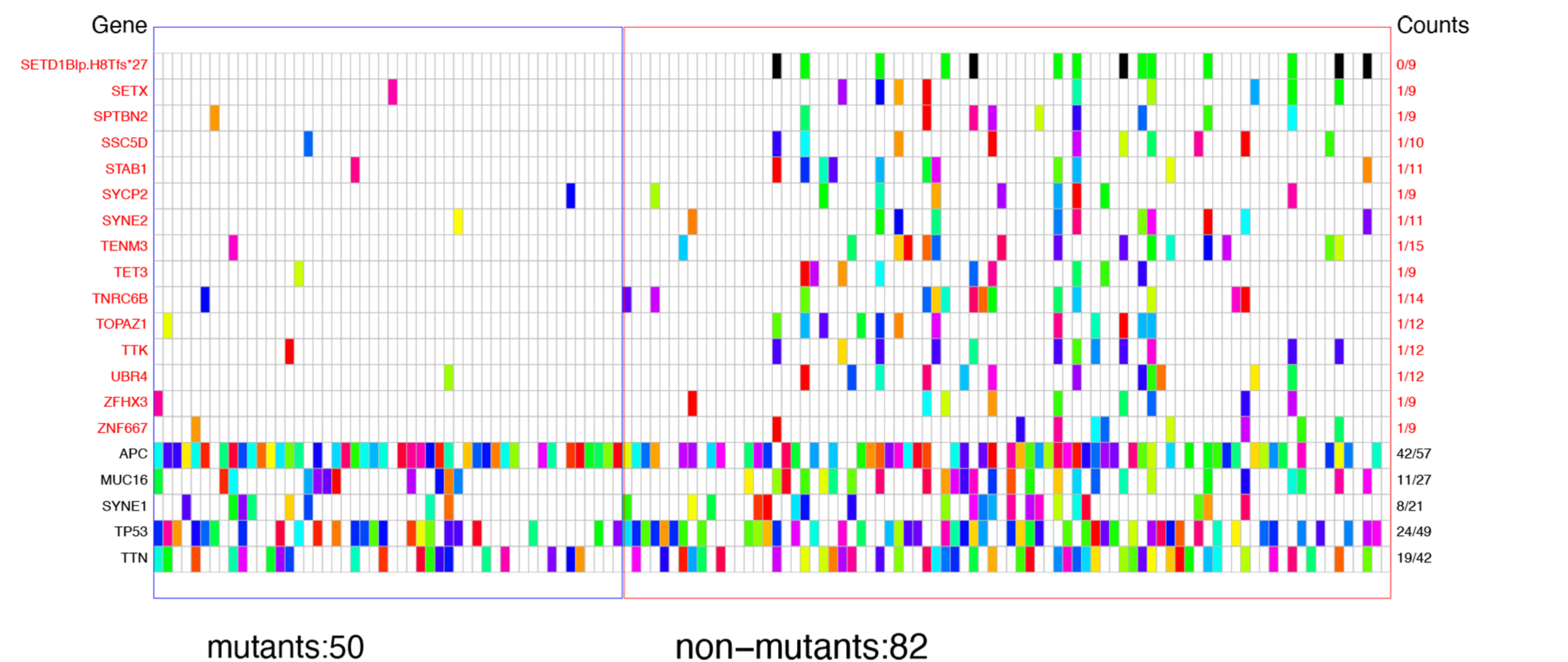
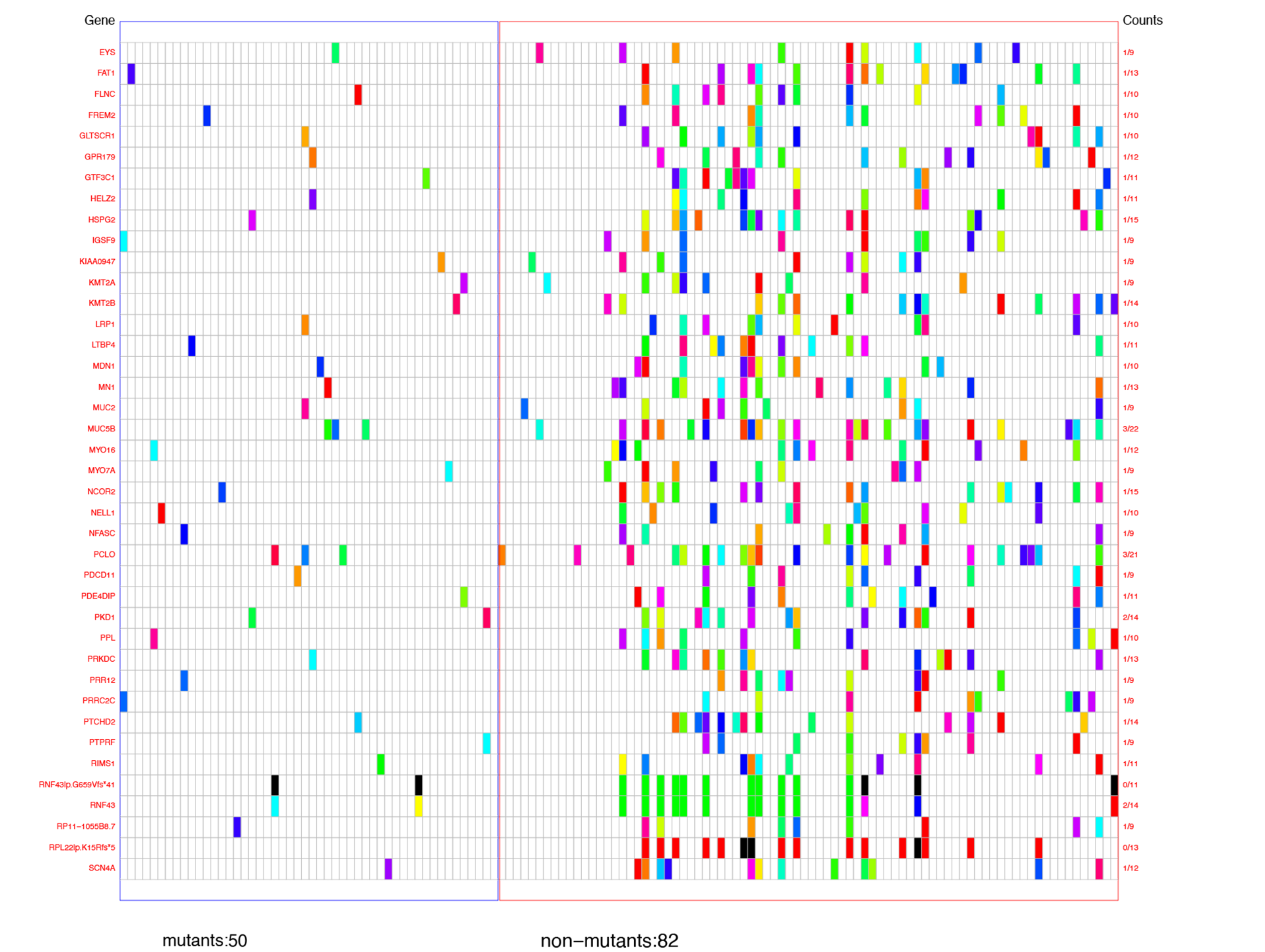
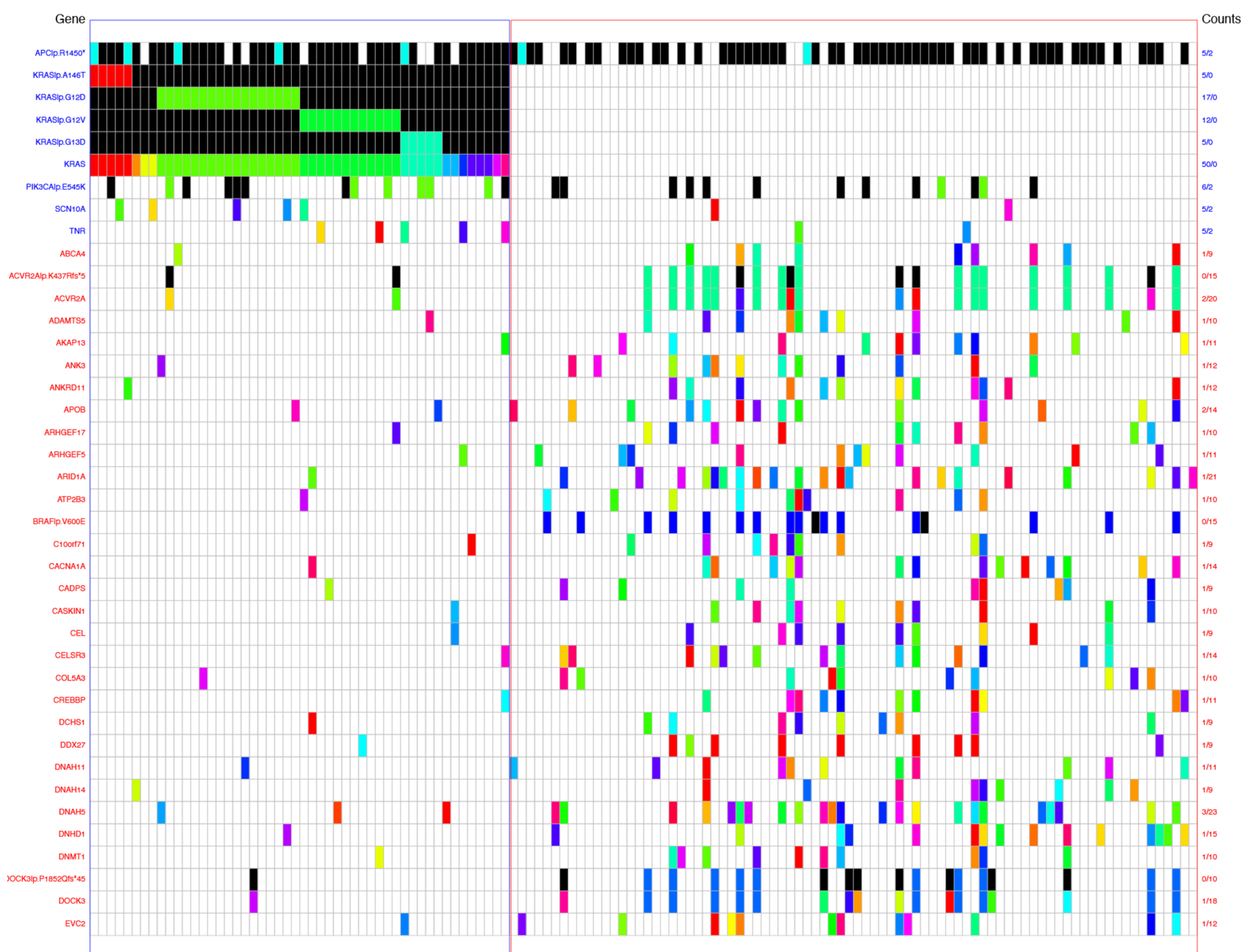
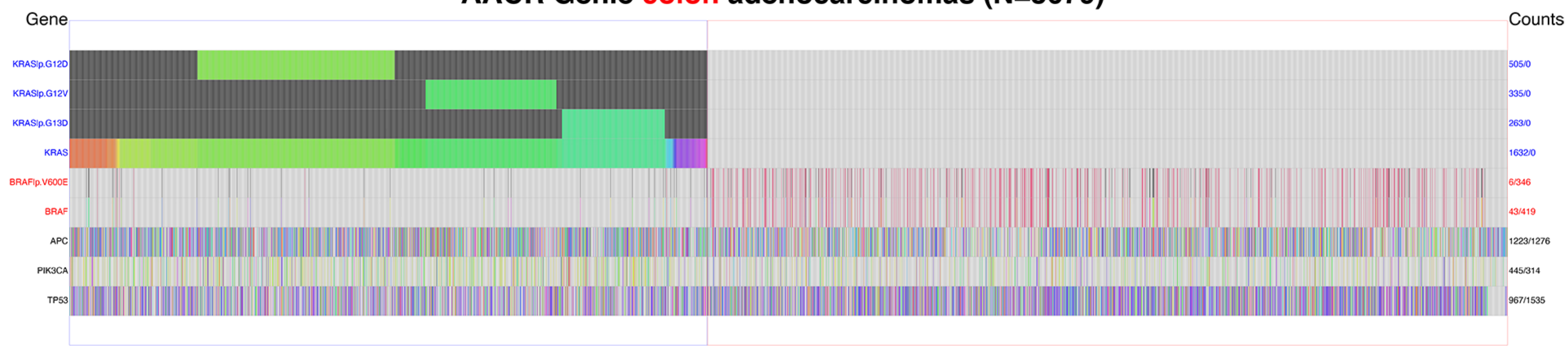


Figure 3B. Mutations in COAD cancers in the MSK cBioPortal data.

AACR Genie colon adenocarcinomas (N=3679)



mutants:1632

non-mutants:2047

Figure 3C. Mutations in COAD cancers in the AACR Genie data.

Broad GDAC **rectal** adenocarcinomas (N=121)

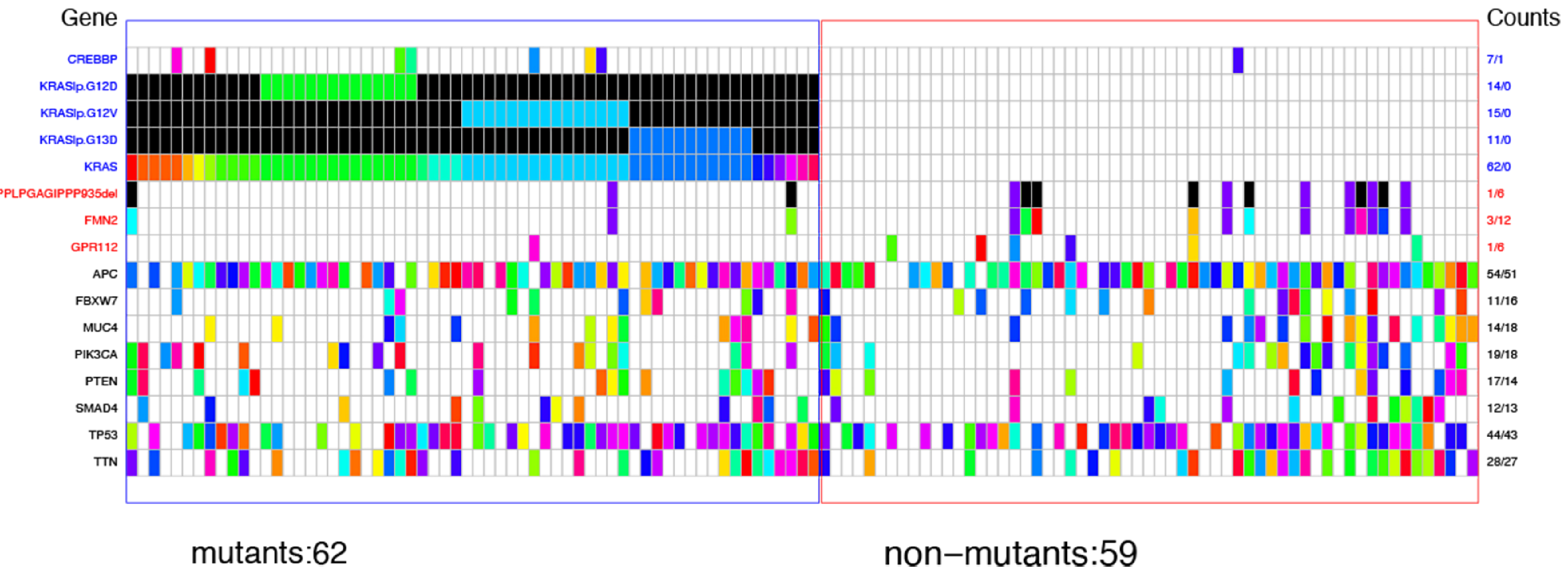


Figure 4A. Mutations in READ (rectal) cancers in the Broad GDAC data.

MSK cBioPortal **rectal** adenocarcinomas (N=327)

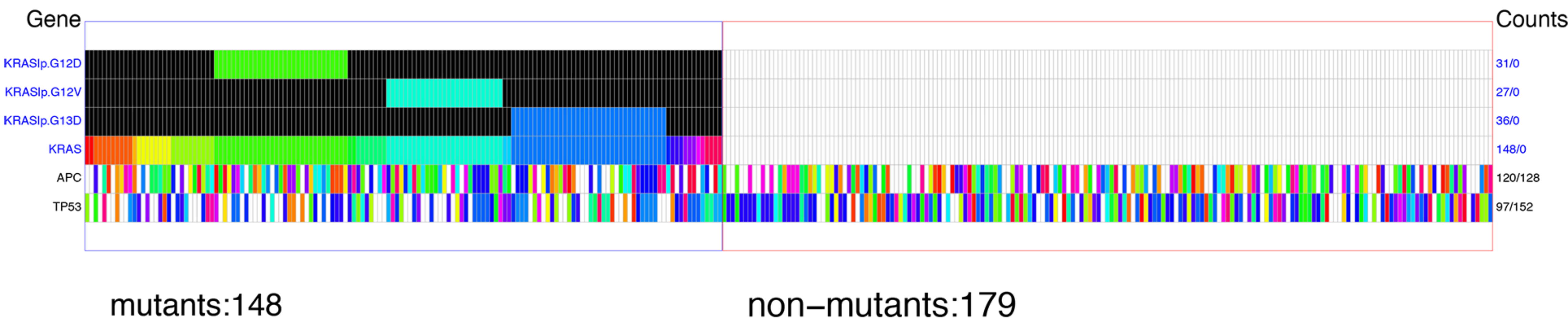
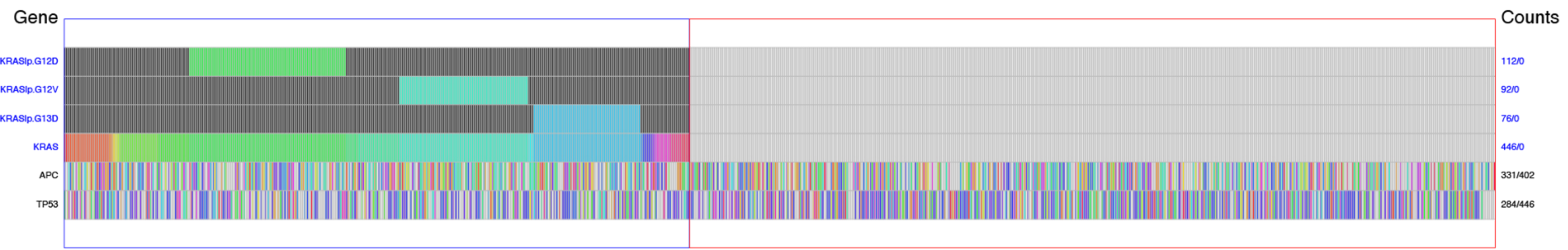


Figure 4B. Mutations in READ cancers in the MSK cBioPortal data.

AACR Genie **rectal** adenocarcinomas (N=1021)

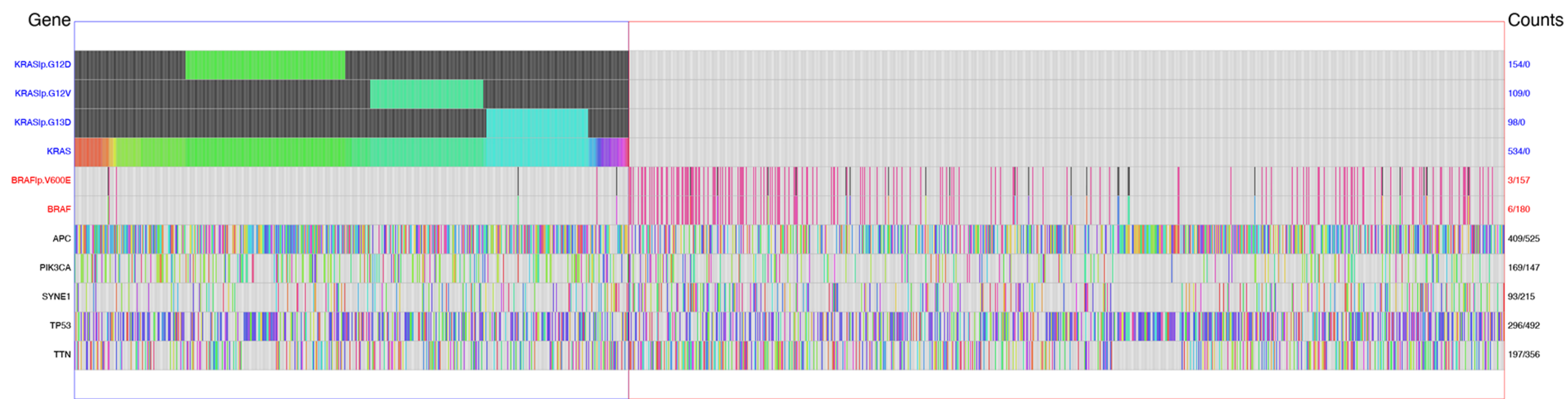


mutants:446

non-mutants:575

Figure 4C. Mutations in READ cancers in the AACR Genie data.

MSK cBioPortal colorectal adenocarcinomas (N=1378)

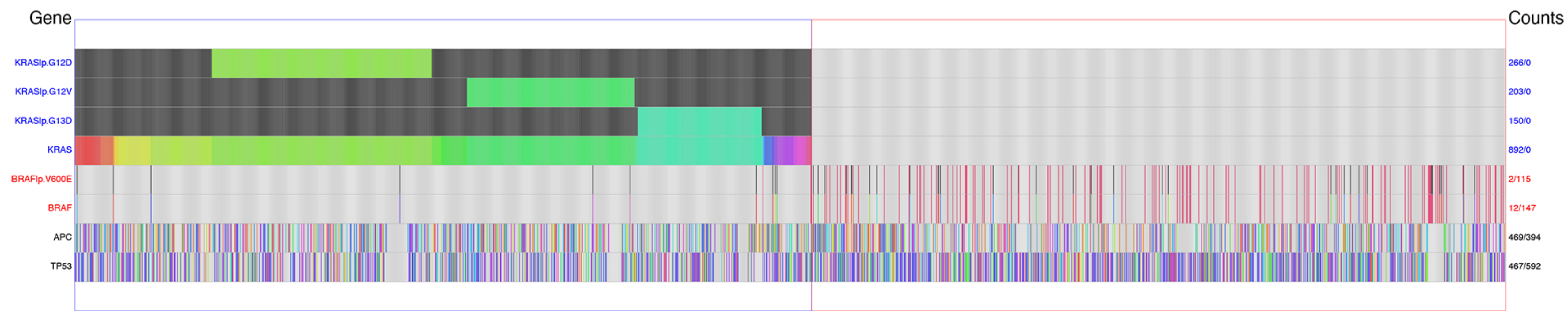


mutants:534

non-mutants:844

Figure 5A. Mutations in COADREAD (colorectal) cancers in the MSK cBioPortal data.

AACR Genie colorectal adenocarcinomas (N=1733)



mutants:892

non-mutants:841

Figure 5B. Mutations in COADREAD cancers in the AACR Genie data.