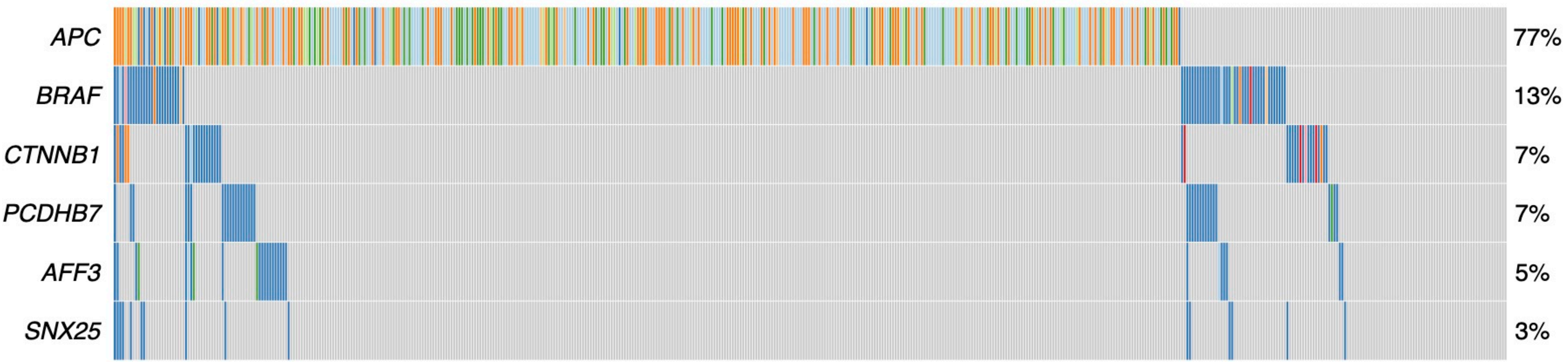
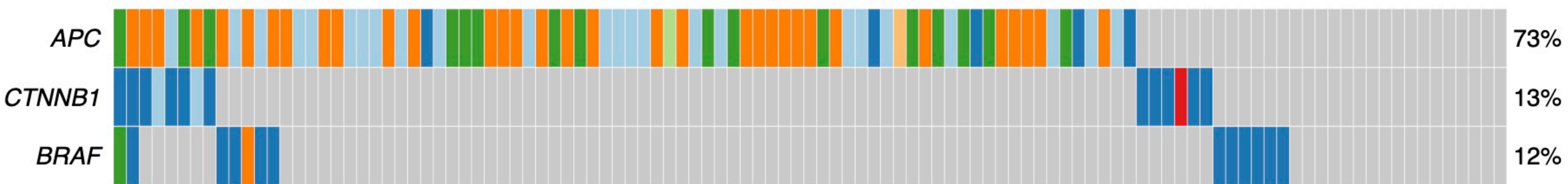


A

Altered in 469 (88.49%) of 530 samples (TCGA cohort)

**B**

Altered in 92 (84.4%) of 109 samples (MSKCC cohort)



- Missense_Mutation
- Nonsense_Mutation
- Frame_Shift_Del
- Splice_Site
- In_Frame_Del
- Multi_Hit