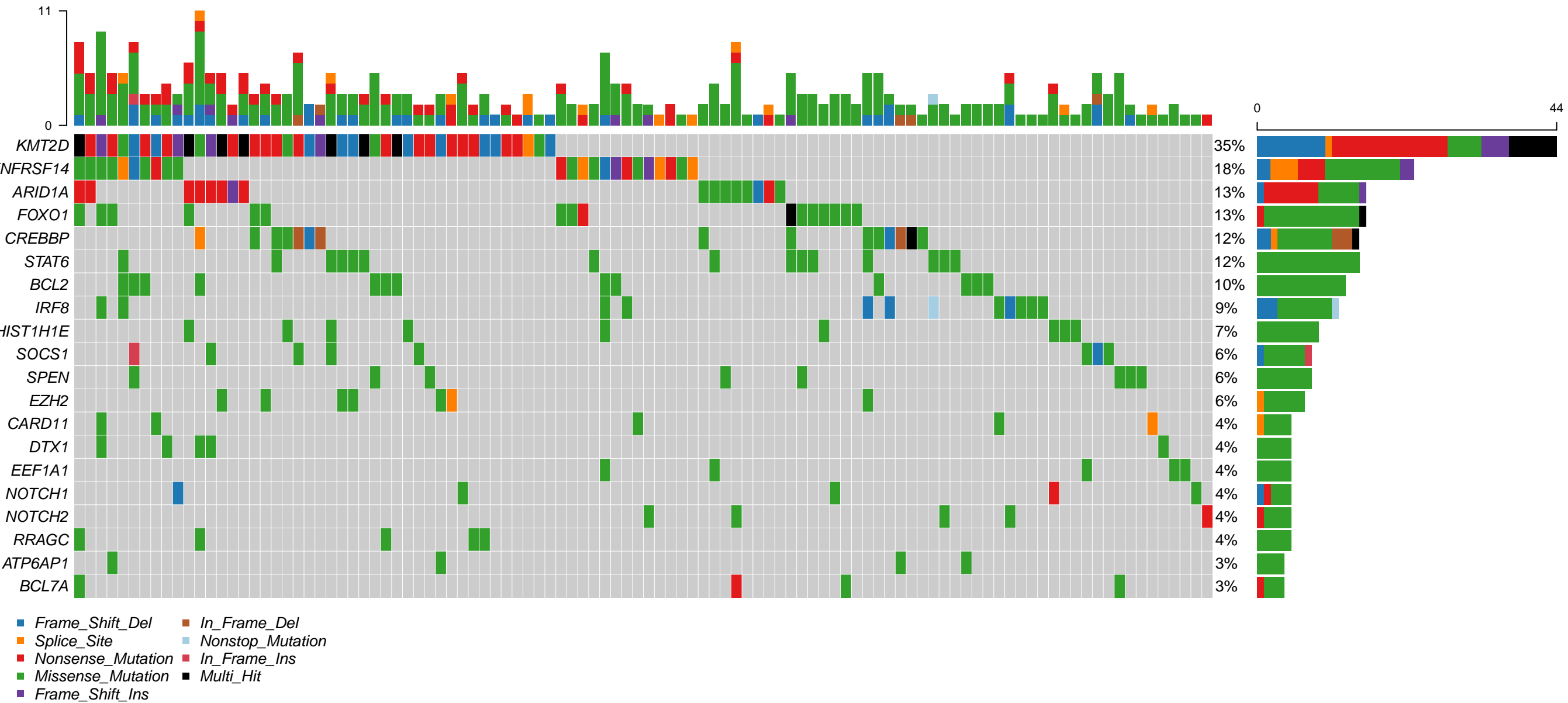
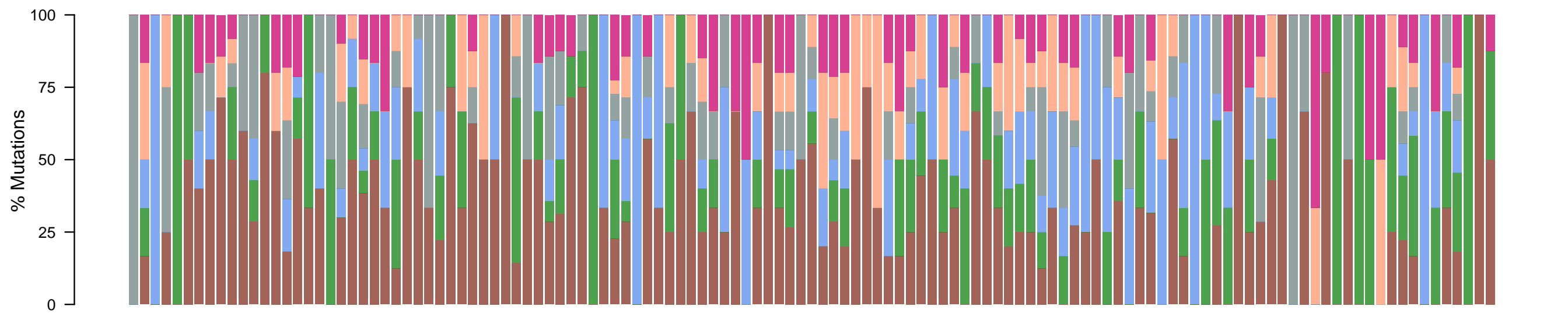
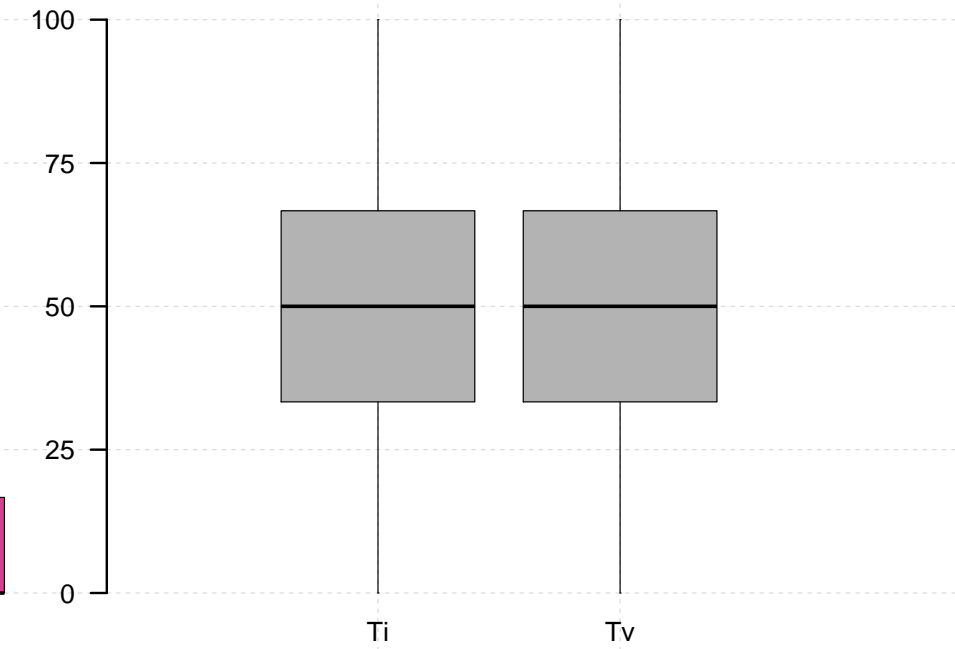
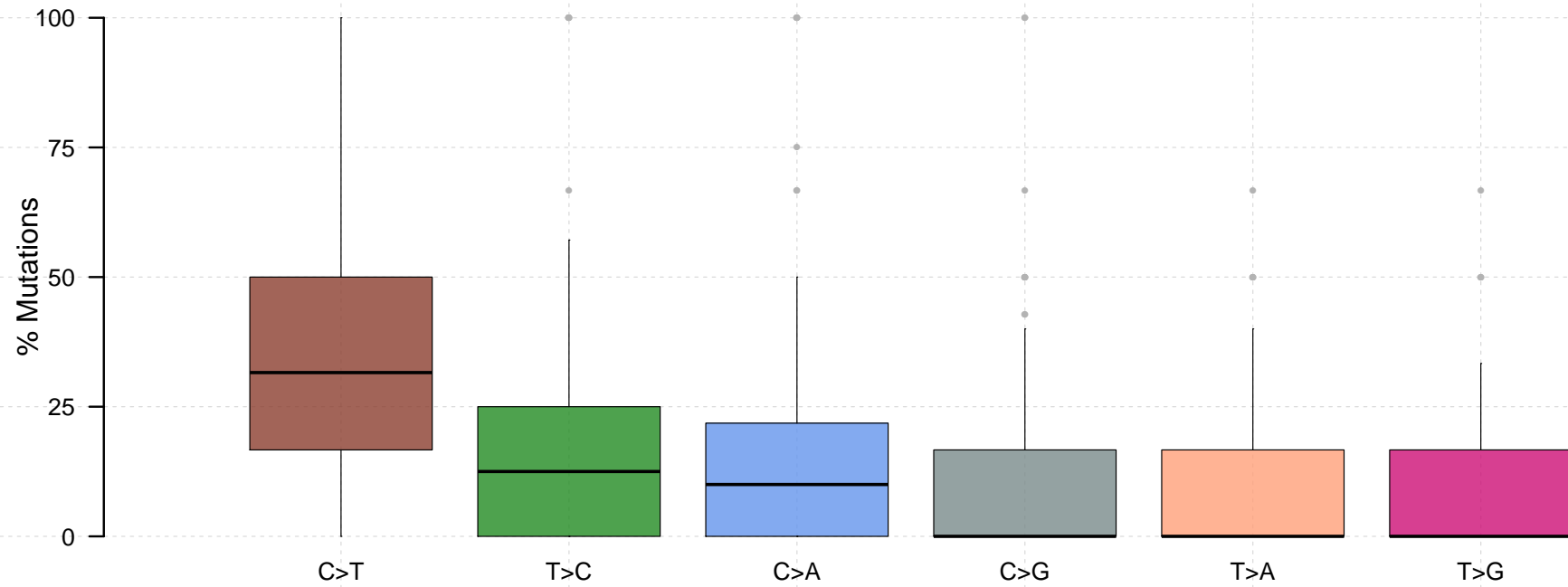
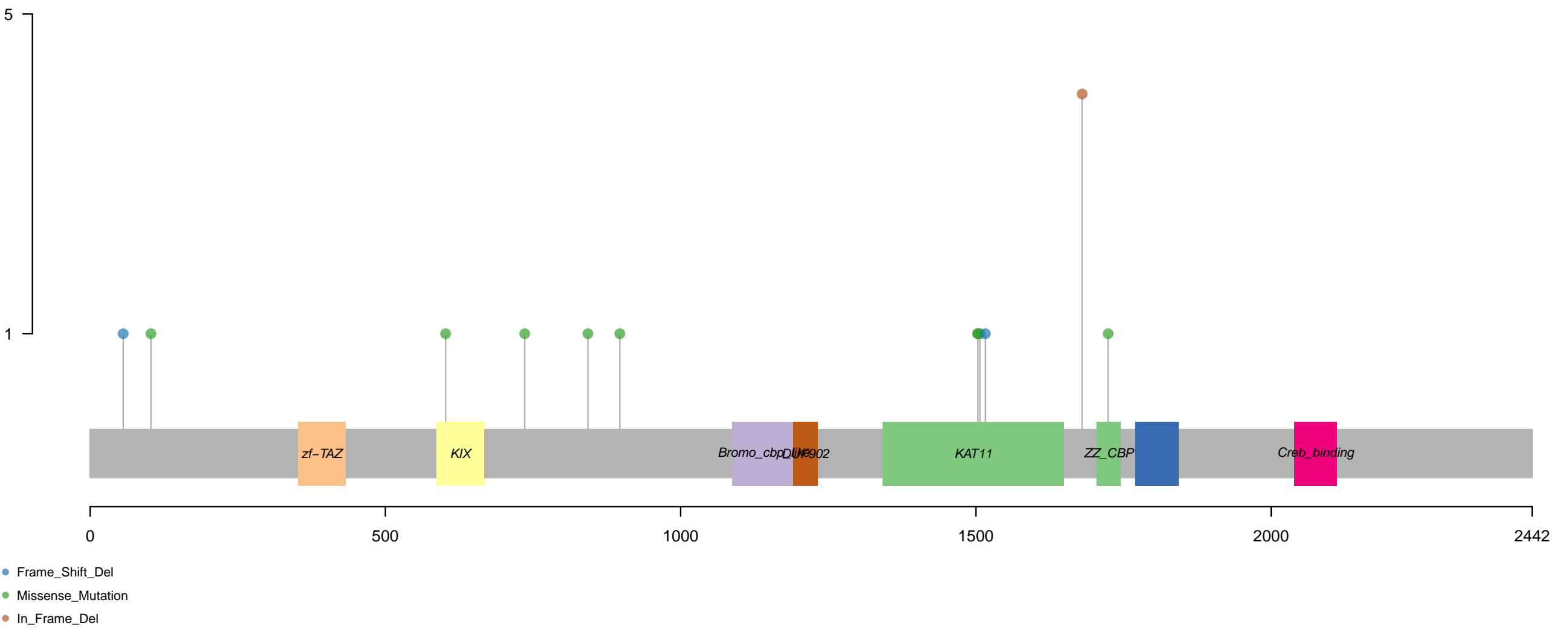
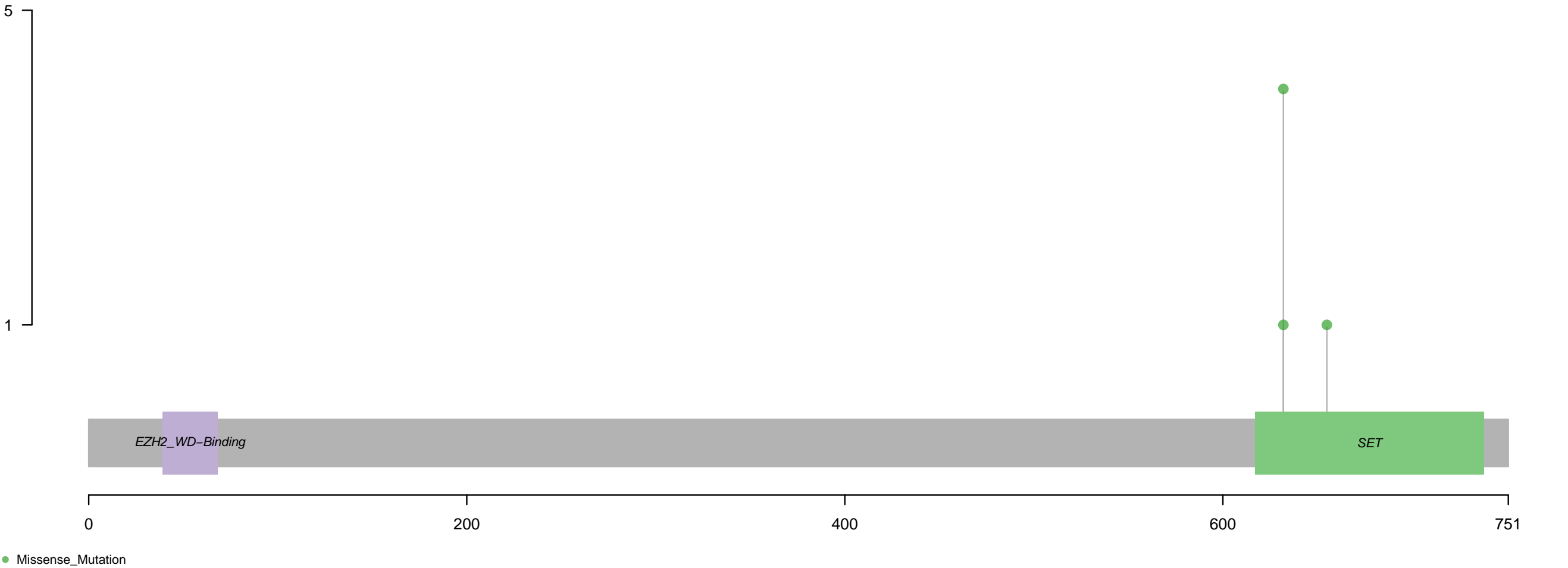


Altered in 104 (81.89%) of 127 samples.









KMT2D : [Somatic Mutation Rate: 34.65%]

NM_003482

