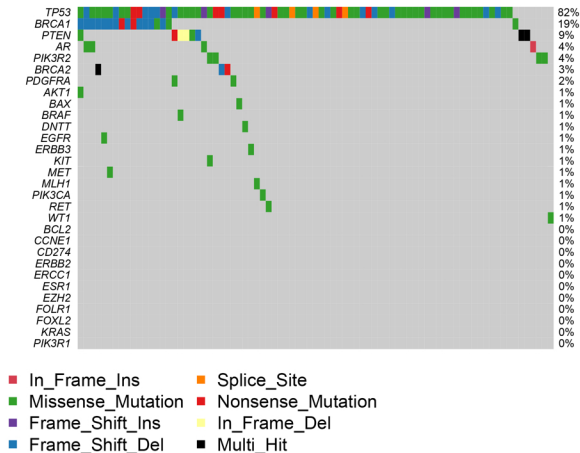
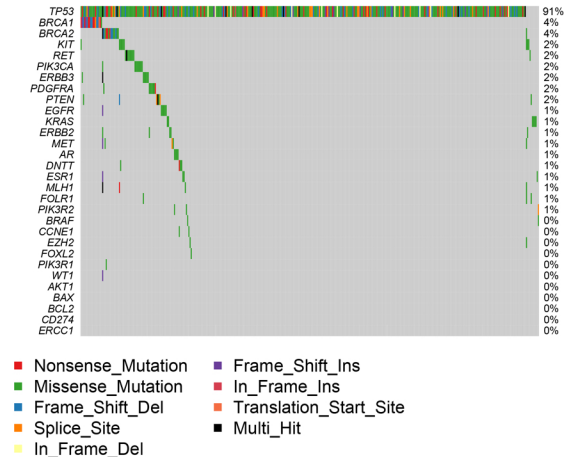


Altered in 81 (90%) of 90 samples.



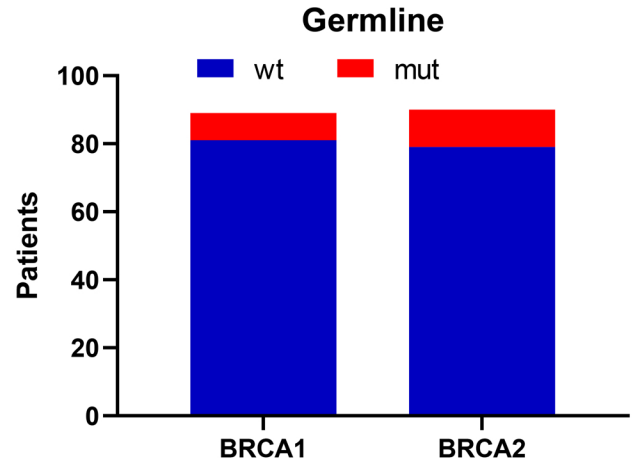
Altered in 383 (93.19%) of 411 samples.



C



D



S-1 A. HRD-related gene mutations in PUTH data. B. HRD-related gene mutations in TCGA data. C. Mutation word cloud generated from PUTH data. D. Proportions of germline mutations in BRCA1 and BRCA2 in PUTH.