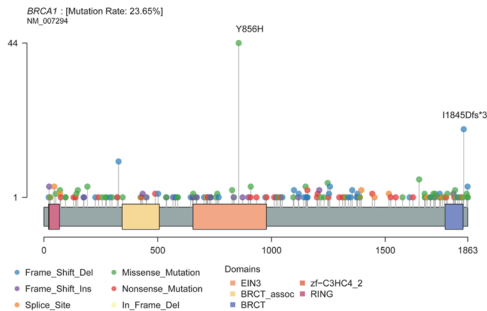


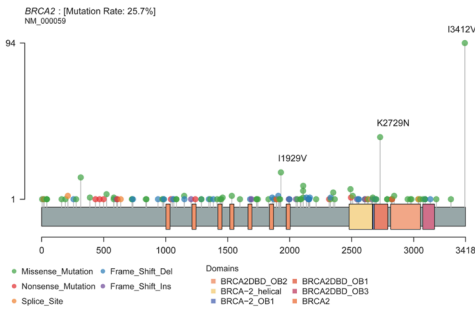
The Chinese ovarian cancer population had characteristic high-frequency mutated genes and hotspots

0.57	0.64	0.45	1.71	ATM
9.50	8.70	17.59	3.42	BRCA1
5.12	5.80	5.57	5.81	BRCA2
1.36	0.92	0.67	3.25	BRIP1
0.00	0.10	0.00	0.00	EPCAM
0.05	0.10	0.00	8.88	MLH1
0.00	0.79	0.45	1.02	MSH2
0.16	0.40	0.00	0.68	MSH6
0.21	0.30	0.00	1.11	PMS2
0.57	0.58	0.00	0.43	RAD51C
0.57	0.48	2.45	2.05	RAD51D
0.00	0.00	0.00	0.00	STK11
Cohort A	Cohort B	Cohort C	Our Cohort	

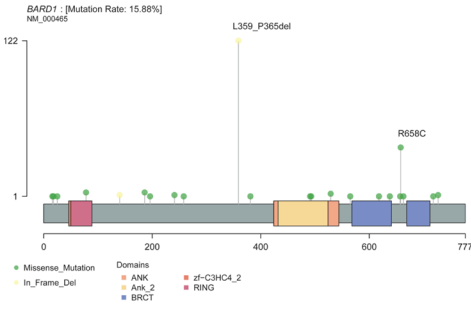
(A)



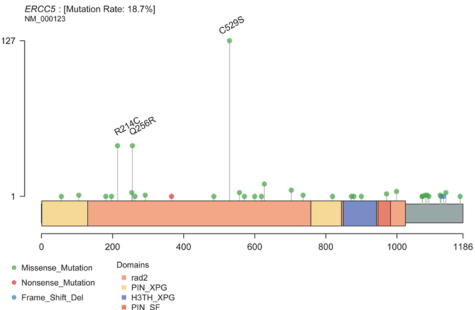
(B)



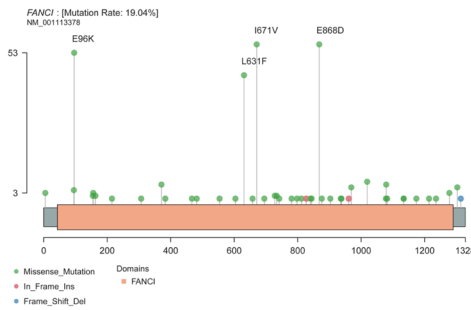
(C)



(D)



(E)



(F)

