

**S1 Table.** Differences in tumor tissue NGS results of the *gBRCA1/2p* and *gBRCA1/2w* groups

	TMB		SNV		CNV		Fusion	
	<i>gBRCA1/2p</i>	<i>gBRCA1/2w</i>	<i>gBRCA1/2p</i>	<i>gBRCA1/2w</i>	<i>gBRCA1/2p</i>	<i>gBRCA1/2w</i>	<i>gBRCA1/2p</i>	<i>gBRCA1/2w</i>
Mean (SD)	10.21 (4.61)	13.47 (5.78)	14.81 (4.95)	18.86 (18.44)	5.77 (4.61)	5.47 (6.87)	0.27 (0.47)	0.23 (0.43)
Median (IQR)	10.45 (6.89-13.17)	12.68 (9.75-16.37)	14 (12-17.75)	16 (12-19.5)	5 (1.5-9.75)	3 (1-7.5)	0 (0-0.75)	0 (0-0)
p-value	0.017 <sup>a)</sup>		0.278 <sup>a)</sup>		0.343 <sup>b)</sup>		0.740 <sup>b)</sup>	

CNV, copy number variant; IQR, interquartile range; NGS, next-generation sequencing; SD, standard deviation; SNV, single-nucleotide variant; TMB, tumor mutation burden. <sup>a)</sup>By Student's t test, <sup>b)</sup>By Mann-Whitney U test.