

S2 Table. Differences in number of HR-related gene aberrations between the *gBRCA1/2p* and *gBRCA1/2w* groups

	HR-related gene aberration		Non- <i>BRCA1/2</i> HR-related gene aberration	
	<i>gBRCA1/2p</i>	<i>gBRCA1/2w</i>	<i>gBRCA1/2p</i>	<i>gBRCA1/2w</i>
Mean (SD)	2.00 (1.23)	1.65 (1.43)	0.54 (0.71)	1.26 (1.26)
Median (IQR)	2 (1-2)	1 (1-2)	0 (0-1)	1 (0-2)
p-value	0.156 ^{a)}		0.008 ^{a)}	

HR, homologous recombination; IQR, interquartile range; SD, standard deviation. ^{a)}By Mann-Whitney U test.