

S5 Table. TMB values according to number of non-*BRCA1/2* HR-related gene aberrations

Non- <i>BRCA1/2</i> HR-related gene aberrations	0	1	≥ 2	p-value ^{a)}
Mean	11.15	11.70	14.96	0.076
SD	5.06	4.88	6.77	

HR, homologous recombination; TMB, tumor mutation burden. ^{a)}p-value by ANOVA.