

Figure S1. Mutational analysis of *BRCA1* and *BRCA2* in SHIN3 cells. (A) Sample *BRCA1* sequence (3,630 bp) was compared with Reference NM_007294 in the NCBI gene database (49) and a single variant of A→G at codon 349 was detected. This variant causes a single amino acid change at position 116 from Gln to Arg, which is currently not known to have functional outcome or be associated with breast or ovarian cancer, therefore it was determined to be variants of unknown significance. Sample Sequence for *BRCA2* (5,018 bp) was compared with Reference NM_000059 in the NCBI gene database (50), and two variants were detected at (B) codon 2660 A→G and (C) codon 4560 G→C. These variants do not cause amino acid changes.



Figure S2. Combination treatment of pharmacological ascorbate and olaparib enhanced DNA double stranded breaks in OVCAR5 cells. Cells were treated with ascorbate, olaparib or a combination of the two at different concentrations for 2 h. Where indicated, cells were pretreated with catalase. Representative western blotting images from two independent experiments are presented. β -actin was used as a loading control. Asc, ascorbate.

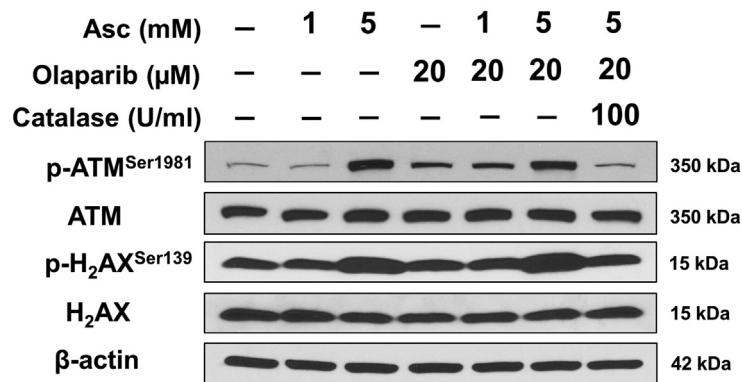


Table SI. Primer sequences used in the present study.

Exon 11 of <i>BRCA1</i> gene	
Nest PCR primers (5'→3')	F1: AGAACCAAGTTTCTGATGGCC R1: GGGACACCTGGATTGCTTT F2: TAGCCAGTTGGTTGATTCCACC R2: GCACCTTAGGAGGAACATGTTTC F3: GGAATAAGCAGAAACTGCCATG F4: ATGAGAATAAAACAAAAGGTGATTG F5: TTGTCAATCCTAGCCTCCAAG F6: TGCCCACCTGGGTCCITAA F7: GAAGTGGGCTCCAGTATTATGA F8: GGATGAAGAGCTTCCCTGCT F9: TCAAGAAGAGCAAAGCATGG
Sequencing primers (5'→3')	
Exon 11 of <i>BRCA2</i> gene	
Nest PCR primers (5'→3')	F1: AAGCAGTCCTGCACCTCAC R1: CCATACTCCCCAAACTGACTA F2: CTGCCTCAGCCTCCAAAAGT R2: CCACTAAGATAAGGGCTCTCC F3: GAAGGACAGTGTGAAAATGATCCAA F4: CAATTCAAAAATAACTGTCAATCC F5: GTGTTGAAATTGTAAATACCTTGGC F6: TGCTGGCCTGTTGAAAAATG F7: TATCTGGCCAGTTTATGAAGGAGG F8: ACAACCGAACGTGATGAAA F9: AAAAACTTCTGTGAGTCAGACTTCA F10: GGTAGGGCCACCTGCATTTA F11: ACAGCAAGTGGAAAATCTGTCC F12: GTTGATAAGAGAAACCCAGAGCAC F13: TTCTAAACTGCCAAGTCATGCC
Sequencing primers (5'→3')	

F, Forward; R, reverse.

Table SII. IC₅₀ values of ascorbate, olaparib and veliparib in the ovarian cancer cell lines.

Cell line	<i>BRCA1</i> mutation status	<i>BRCA2</i> mutation status	Ascorbate IC ₅₀ (mM) ^a	Olaparib IC ₅₀ (μ M) ^a	Veliparib IC ₅₀ (μ M) ^a
A2780	Wild-type	Wild-type	0.15±0.02	-	-
OVCAR10	Wild-type	Wild-type	0.30±0.01	-	-
OVCAR3	Wild-type	Wild-type	0.36±0.03	-	-
OVCAR5	Wild-type	Wild-type	1.35±0.04	219.85±77.65	64.06±10.04
SKOV3	Wild-type	Wild-type	1.49±0.04	-	-
OVCAR8	Wild-type	Wild-type	1.67±0.02	-	-
SHIN3	Wild-type	Wild-type	2.80±0.08	270.95±69.75	95.28±24.82
HIO-80 ^b	-	-	NA ^c	-	-

^aMean ± standard error of mean; ^bnon-tumorigenic human ovarian cell line; ^cNA, not applicable.