

Supplementary Materials

Impact of ERCC1, XPF and DNA Polymerase β Expression on Platinum Response in Patient-Derived Ovarian Cancer Xenografts

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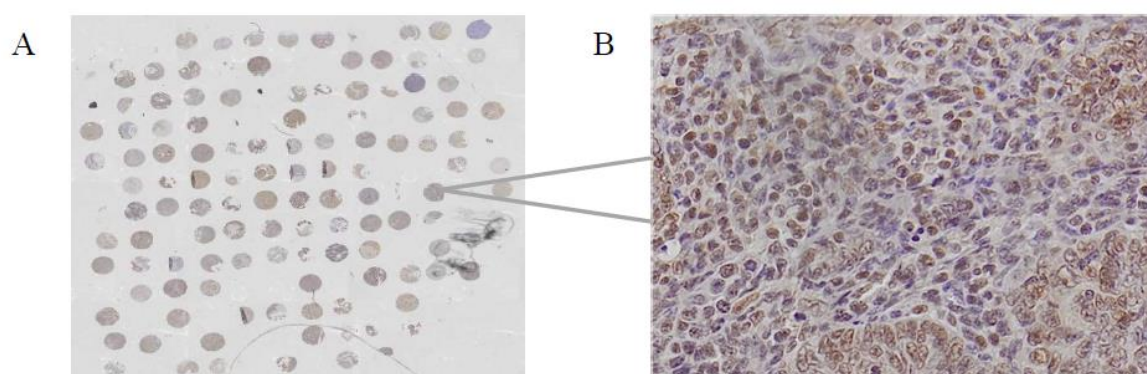


Figure S1. ERCC1 immunohistochemical (IHC) expression on the TMA. (A). Representative image of the FFPE-TMA comprehensive of 52 ovarian cancer PDXs, stained with antibody anti-ERCC1 (Santa Cruz Biotechnology, sc-10785, Dallas, TX, USA) magnification 2 \times . (B). Representative PDX#4, ERCC1-positive (IHC-score = 3.25). ERCC1-dots are present both in the cell nucleus than in the cytoplasm. Magnification 20 \times .

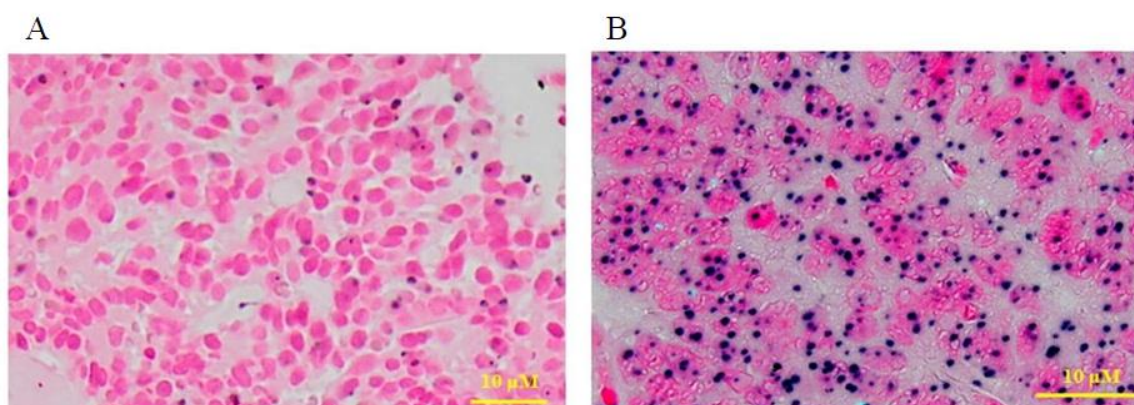


Figure S2. ERCC1/XPF complexes detected by PLA on a FFPE OC-PDX's tumor sample. (A) Negative control. PLA was done without primary antibodies. It is possible to observe nuclei counterstained with Nuclear Fast Red (pink nuclei). (B) The same FFPE OC-PDX sample after PLA technique. ERCC1/XPF foci are visible as violet dots within epithelial cancer cells' nuclei counterstained with Nuclear Fast Red (pink nuclei). Magnification 40 \times , brightfield microscopy.

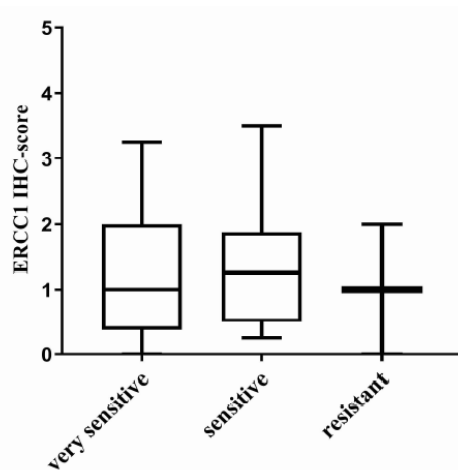


Figure S3. ERCC1-IHC score in the high grade serous/endometrioid PDXs. Distribution of ERCC1 protein expression in PDX very sensitive ($n = 10$), sensitive ($n = 12$) and resistant ($n = 2$) to DDP (see Material and Methods). Data are expressed as mean \pm standard deviation.

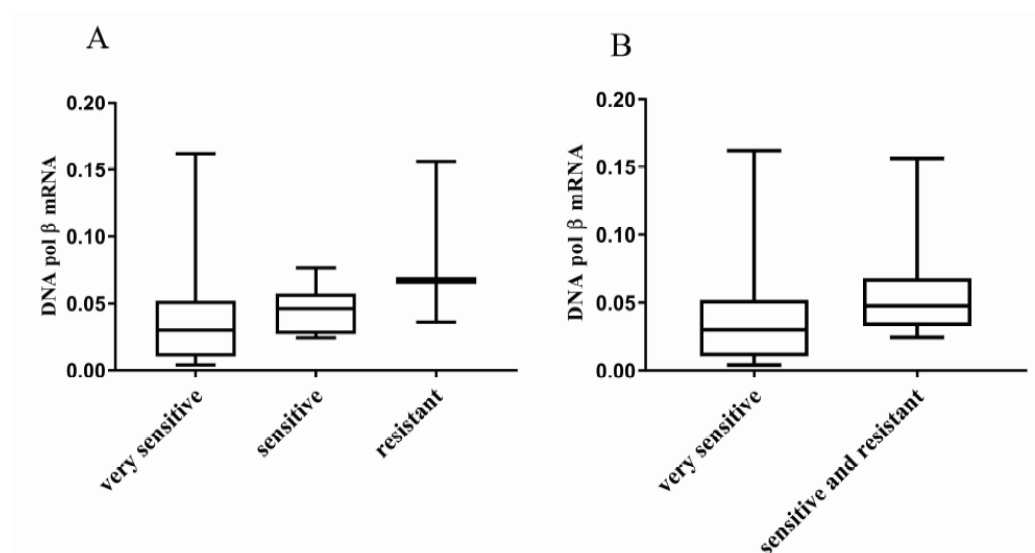


Figure S4. DNA pol β mRNA in high grade serous and endometrioid PDXs. (A). Distribution of DNA pol β mRNA in high grade PDXs very sensitive ($n = 9$), sensitive ($n = 11$) and resistant ($n = 3$) (see Material and Methods) to DDP. (B). Distribution of DNA pol β mRNA in very sensitive PDXs and sensitive/resistant to DDP. Data are expressed as mean \pm standard deviation.

Table S1. Correlations found in all the OC-PDXs.

Method	Molecular Target	IHC-Score		PLA	Normalized Gene Expression Levels		
		<i>ERCC1</i>	<i>DNA pol β</i>	<i>ERCC1/XPF</i>	<i>ERCC1</i>	<i>XPF</i>	<i>DNA pol β</i>
IHC-Score	<i>ERCC1</i>	1	0.03	0.29	-0.11	0.65	0.14
			0.85	0.05	0.96	< 0.0001	0.43
		49	48	46	34	34	34
	<i>DNA pol β</i>	0.03	1	0.58	-0.11	0.11	0.17
		0.85		< 0.0001	0.53	0.54	0.33
		48	51	48	35	35	35
PLA	<i>ERCC1/XPF</i>	0.29	0.58	1	0.04	0.32	0.09
		0.05	< 0.0001		0.82	0.06	0.60
		46	48	49	35	35	35
Normalized Gene Expression Levels by RT-PCR	<i>ERCC1</i>	-0.11	-0.11	0.04	1	0.15	0.32
		0.96	0.53	0.82		0.40	0.05
		34	35	35	36	36	36
	<i>XPF</i>	0.65	0.11	0.32	0.15	1	0.40
		< 0.0001	0.54	0.06	0.40		0.02
		34	35	35	36	36	36
	<i>DNA pol β</i>	0.14	0.17	0.09	0.32	0.40	1
		0.43	0.33	0.60	0.05	0.02	
		34	35	35	36	36	36

Legend: in each box in the first line is reported the Spearman's correlation index, in the second line the p-value (in grey significant p-values), in the third line the number of observations.

Table S2. Tumor histotypes, DDP sensitivity and raw data from IHC, PLA and gene expression analyses in the OC-PDX population under study.

#ID PDX	Histotype	DDP Sensitivity	DNA pol β (IHC-score)	ERCC1 (IHC-score)	PLA	Normalized Gene Expression Levels (mRNA copies)		
					Mean (SD)	ERCC1	XPF	DNA pol β
#PDX1	HG Serous	VS	9	2	7.090 (0.82)	3.0696	0.0029	0.0580
#PDX2	HG Serous	VS	8	0.5	6.079 (1.30)	0.5889	0.0009	0.0170
#PDX3	HG Serous	VS	4.5	0	5.752 (0.92)	2.1873	0.0013	0.1619
#PDX4	HG Serous	VS	0	3.25	3.685 (0.65)	0.5486	0.0026	0.0095
#PDX5	HG Serous	VS	1.5	1	3.267 (0.39)	1.5639	0.0011	0.0460
#PDX6	HG Serous	VS	6	2	4.891 (2.16)	-	-	-
#PDX7	HG Serous/Endom.	VS	0	0.5	0.925 (0.10)	0.7970	0.0018	0.0109
#PDX8	HG Endometrioid	VS	5	1	3.383 (0.74)	0.4336	0.0018	0.0040
#PDX9	HG Endometrioid	VS	-	0	1.250 (0.27)	1.4209	0.0010	0.0300
#PDX10	HG Endometrioid	VS	6	1	5.963 (1.04)	0.3397	0.0023	0.0400
#PDX11	HG Serous	S	8	1.5	7.171 (0.90)	0.8954	0.0038	0.0491
#PDX12	HG Serous	S	12	0.25	5.673 (1.06)	0.3985	0.0010	0.0551
#PDX13	HG Serous	S	1.5	3.5	3.173 (2.60)	0.8543	0.0055	0.0573
#PDX14	HG Serous	S	0	1.5	3.774 (0.44)	2.7480	0.0018	0.0462
#PDX15	HG Serous	S	9	1.5	4.152 (0.76)	0.2073	0.0019	0.0375
#PDX16	HG Serous	S	4.5	0.5	5.272 (1.00)	0.7744	0.0022	0.0244
#PDX17	HG Serous	S	10	0.5	7.039 (0.42)	0.7962	0.0013	0.0271
#PDX18	HG Serous	S	10.5	2	-	-	-	-
#PDX19	HG Endometrioid	S	8	1	4.243 (1.07)	0.4143	0.0040	0.0706
#PDX20	HG Endometrioid	S	0	0.5	2.779 (0.40)	1.5158	0.0014	0.0765
#PDX21	HG Endometrioid	S	0	2	-	0.8398	0.0038	0.0272
#PDX22	HG Endometrioid	S	6	0.75	5.408 (-)	1.2359	0.0019	0.0344
#PDX23	HG Serous	R	7.5	2	8.598 (1.95)	1.0627	0.0090	0.0673
#PDX24	HG Serous	R	0	0	4.202 (0.99)	2.9602	0.0030	0.0358
#PDX25	HG Serous	R	10	-	0.920 (0.16)	0.8152	0.0031	0.1563
#PDX26	HG Serous	-	12	3	6.941 (1.60)	3.071	0.0030	0.0336
#PDX27	HG Serous	-	10	2	8.765 (-)	1.5871	0.0033	0.0591
#PDX28	HG Serous	-	0	2	3.724 (0.68)	0.4129	0.0044	0.0590
#PDX29	HG Serous	-	0	-	3.245 (2.25)	-	-	-
#PDX30	HG Serous	-	7	2.5	6.309 (1.05)	-	-	-
#PDX31	HG Serous	-	7	1	1.306 (0.72)	-	-	-
#PDX32	HG Serous	-	4.5	2	6.063 (0.89)	-	-	-
#PDX33	HG Serous	-	9	1	6.572 (-)	-	-	-
#PDX34	HG Serous	-	5.5	2.5	7.377 (1.95)	-	-	-
#PDX35	HG Serous	-	7	0.5	5.928 (0.76)	-	-	-
#PDX36	HG Serous	-	10	2	6.260 (0.81)	-	-	-
#PDX37	HG Serous	-	12	2	8.597 (1.86)	-	-	-

#PDX38	HG Serous	-	12	0.75	-	-	-	-
#PDX39	HG Serous	-	2	1.25	4.535 (0.93)	-	-	-
#PDX40	HG Serous	-	12	0.75	8.732 (-)	-	-	-
#PDX41	HG Serous/Endom.	-	7.5	2	4.585 (1.03)	-	-	-
#PDX42	LG Serous	VS	8	0.75	3.964 (0.65)	1.0772	0.0029	0.0347
#PDX43	LG Endometrioid	S	0	0.25	3.165 (1.22)	0.6881	0.0005	0.0263
#PDX44	Mixed mullerian	S	1.5	0.5	3.146 (0.58)	0.9031	0.0011	0.0362
#PDX45	Carcinosarcoma	-	0	2.75	3.389 (1.10)	1.9701	0.0025	0.0653
#PDX46	Undifferentiated	-	3.5	2	5.899 (0.84)	-	-	-
#PDX47	Undifferentiated	S	3	2	4.453 (1.70)	0.8242	0.0019	0.0421
#PDX48	Mucinous	S	3	-	3.665 (0.81)	0.6693	0.0008	0.0089
#PDX49	Mucinous	R	12	2	5.069 (1.74)	0.7853	0.0030	0.0342
#PDX50	Clear cell	S	1.5	2.5	7.922 (-)	0.2010	0.0025	0.0252
#PDX51	Clear cell	S	0	1	5.318 (0.43)	1.8312	0.0028	0.0198
#PDX52	Clear cell	R	3.5	0.75	3.617 (1.85)	0.5060	0.0013	0.0250

-: data not available; VS: very sensitive (light grey box) to DDP; S: sensitive to DDP (grey); R: resistant to DDP (dark gray) as explained in Materials and Methods.

Table S3. List of the primers used in real time-PCR for the selected genes.

Function	Gene	Primer Forward	Primer Reverse
Base Excision repair (BER)	<i>DNA pol β</i>	TGCCTGGAGTAGGAACAAAA	GGAAATTGATGGATGAACTCG
Nucleotide excision repair (NER)	<i>ERCC1</i>	CCAACAGCATCATTGTGAGC	TCTTGGCCCAGCACATAGTC
Nucleotide excision repair (NER)	<i>XPF</i>	TTGTGAGGAACTGTATCTGTGG	AGCAAGCATGGTAGGTGTCA
Housekeeping	<i>ACTβ</i>	TCACCCACACTGTGCCATCTACGA	CAGCGGAACCGCTCATTGCCAATGG
Housekeeping	<i>CYPA</i>	GACCCAACACAAATGGTTCC	TTTCACTTTGCCAAACACCA



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