

¿Podemos detectar la resistencia a iPARP?: de la preclínica a la clínica

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Prof Asociado Universidad de Valencia



La paradoja del SS John Harvey



Frederijk Guthrie
Describe el gas mostaza
1859



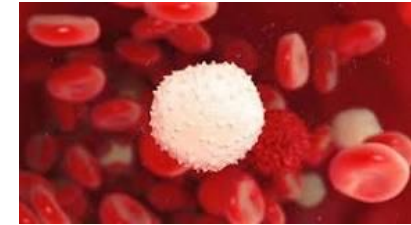
1ª Guerra Mundial → Wilhelm Steinkopf
lo sintetiza a gran escala



2ª Guerra Mundial
Bombardeo sobre Bari 1943
“El 2º Pearl Harbour”



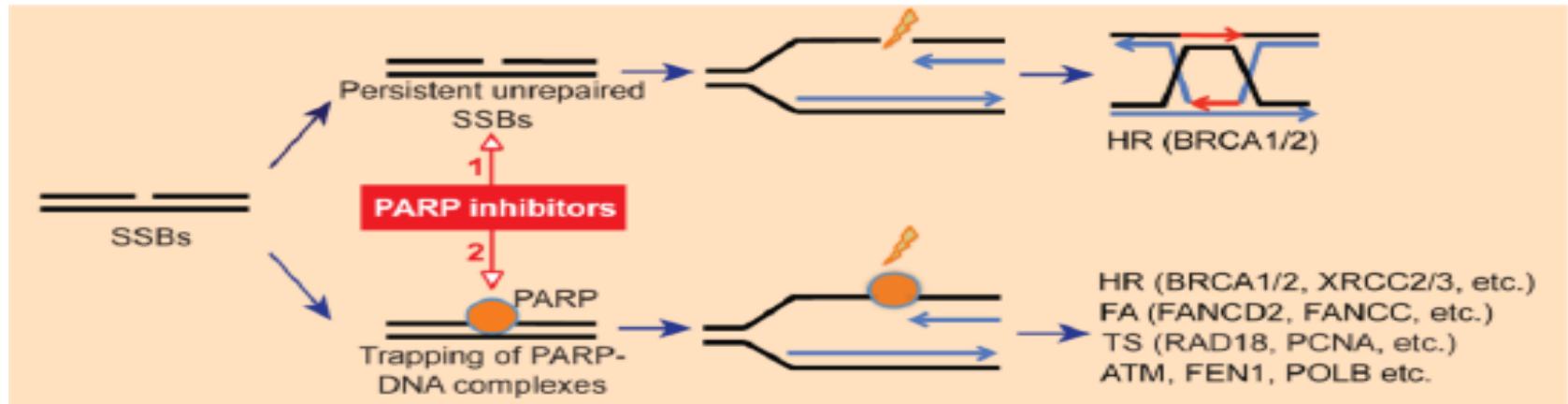
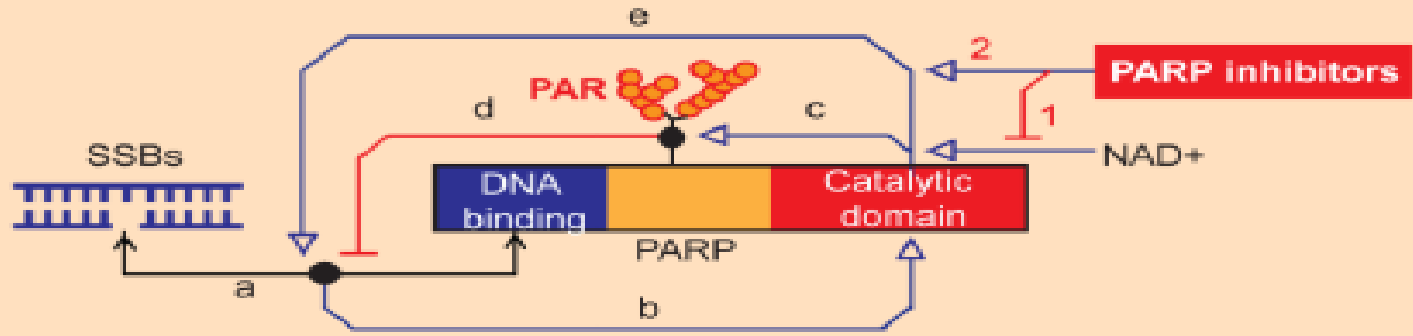
El SS John Harvey se hunde
cargado de gas mostaza



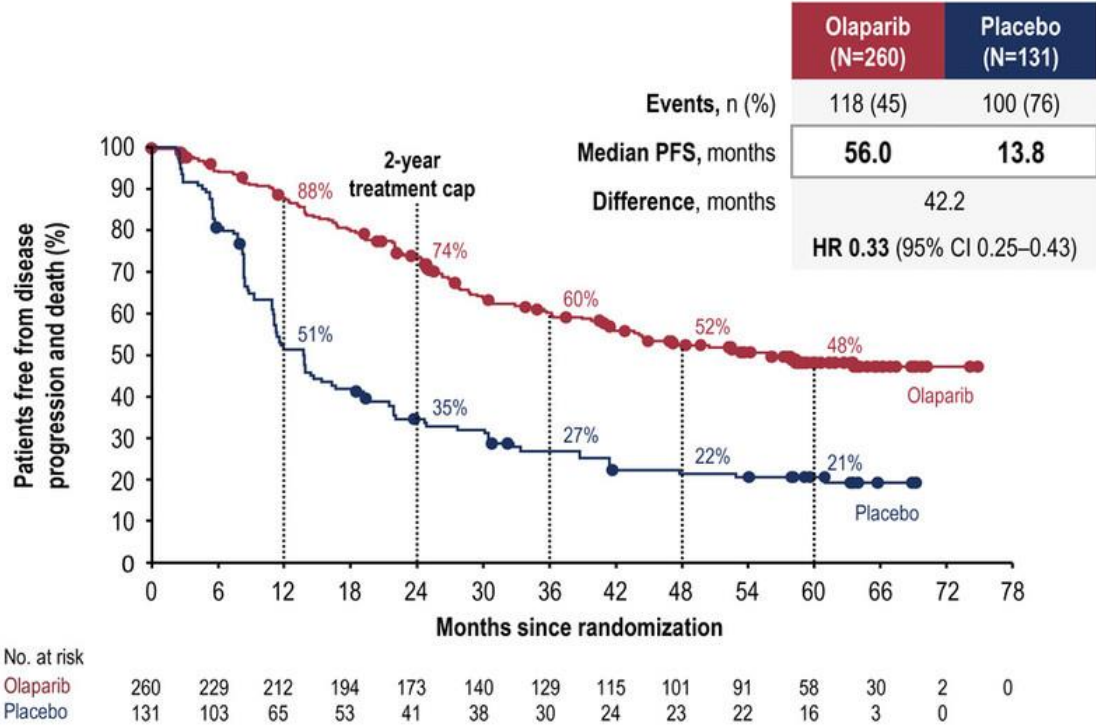
Cientos de habitantes de Bari
fallecen con leucopenia

Se crea la primera
Quimioterapia
La mostaza nitrogenada

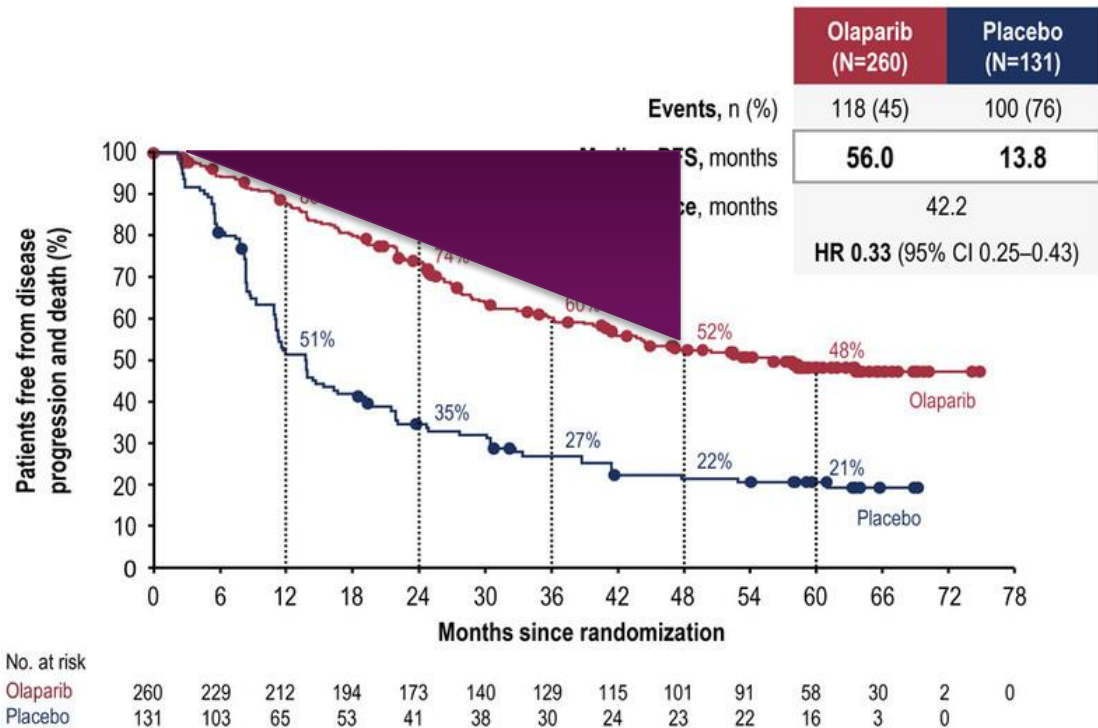
La paradoja de los iPARP



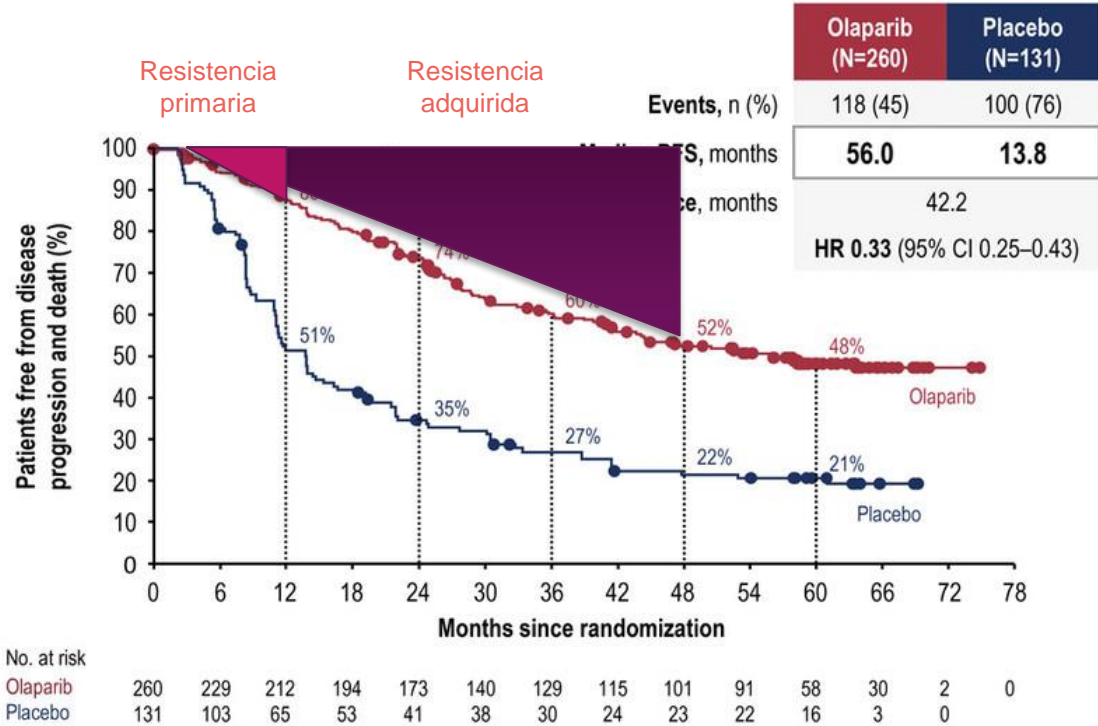
SOLO-1




SOLO-1



SOLO-1






**¿Cuáles son los
mecanismos de
resistencia a iPARP?**

**¿Cuáles son las bases
biológicas para
estrategias que
reviertan las
resistencias a PARPi?**

**¿Existe una resistencia
cruzada PARPi-platino?**

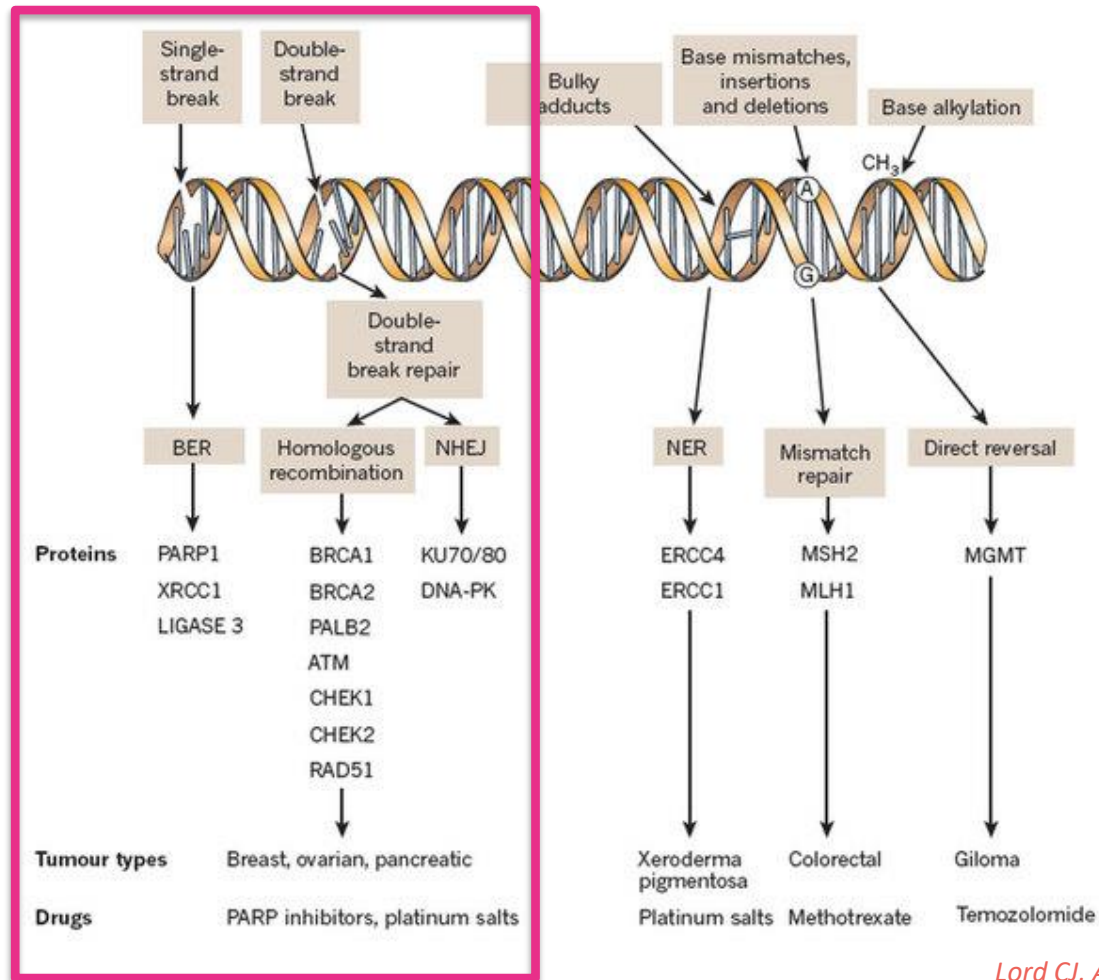


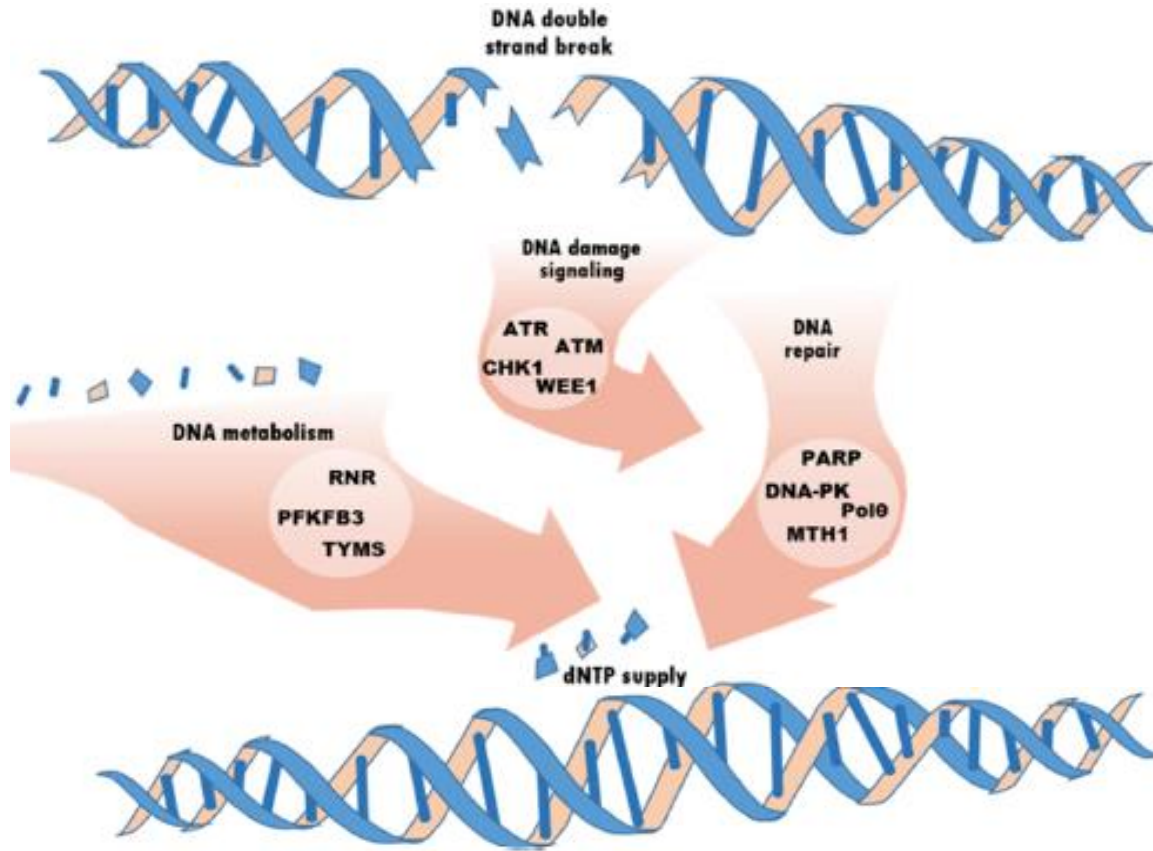
**¿Cuáles son los
mecanismos de
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**¿Cuáles son las bases
biológicas para
estrategias que
reviertan las
resistencias a PARPi?**

Tipos de daño en DNA

Mecanismos de reparación





Mecanismos de resistencia a iPARP



Sobreexpresión
ABCB1



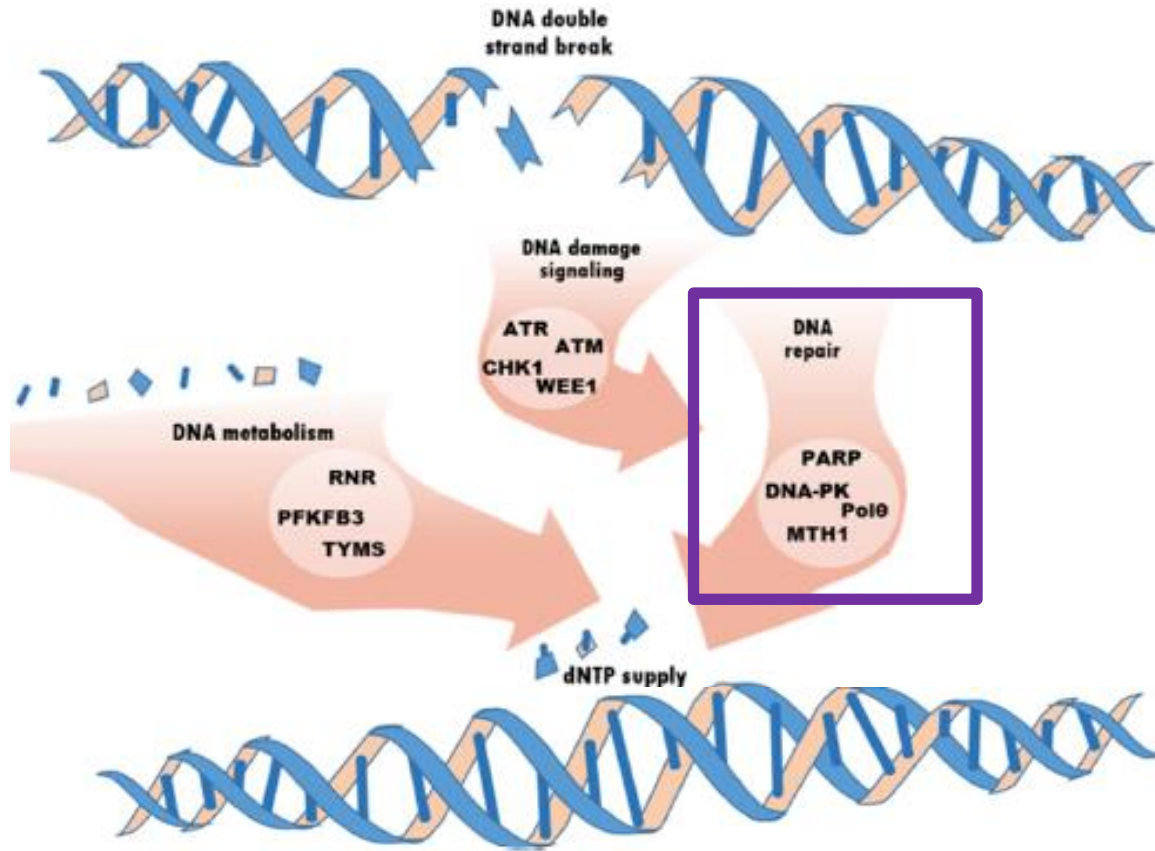
Estabilizan el replicación fork
– anulan el efecto trapping

Inicio del NHEJ → pérdida de
53BP1

Pérdida de fenotipo HRD

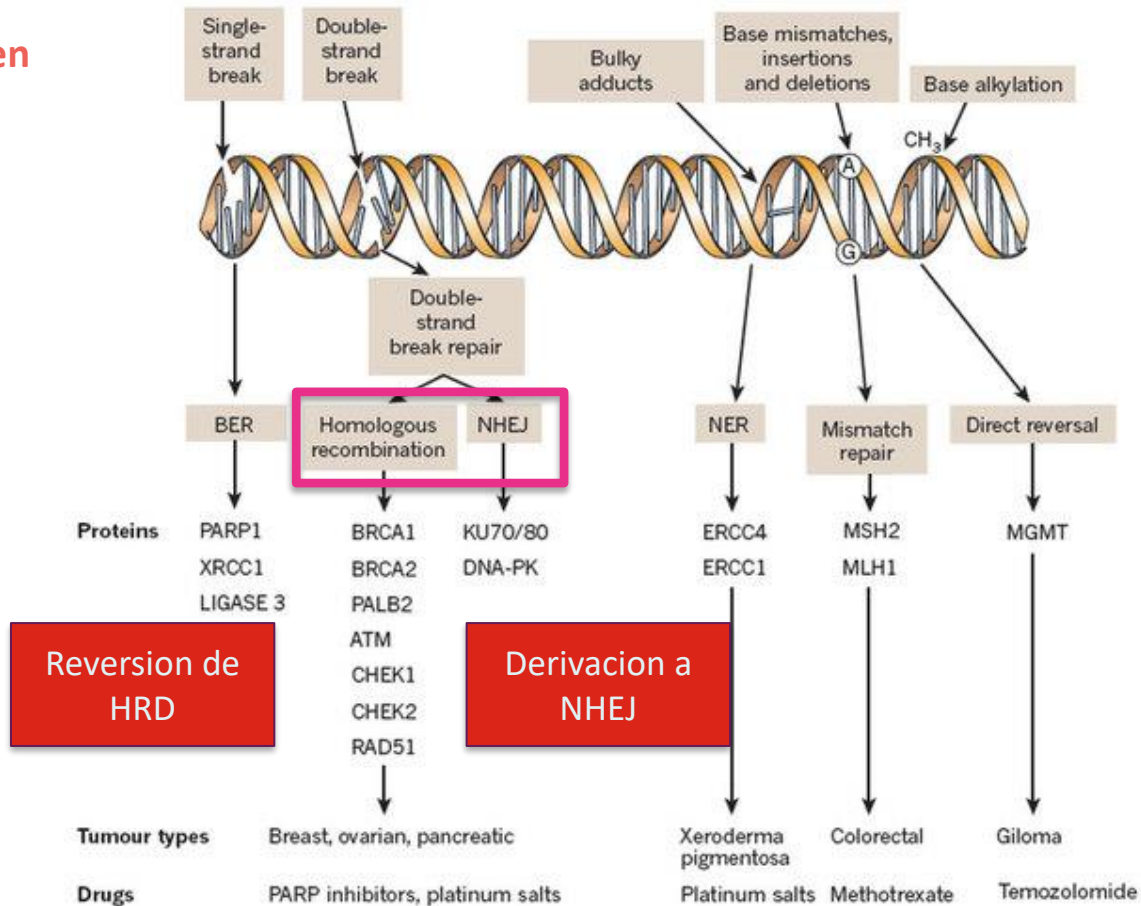


Modifican checkpoint
intrafase S
Amplificación CCNE1
Pérdida de SLFN11



Tipos de daño en DNA

Mecanismos de reparación



RING domain-deficient BRCA1 promotes PARP inhibitor and platinum resistance

Therapeutics, Targets, and Chemical Biology

Cancer
Research

The BRCA1- Δ 11q Alternative Splice Isoform Bypasses Germline Mutations and Promotes Therapeutic Resistance to PARP Inhibition and Cisplatin

Yifan Wang¹, Andrea J. Bernhardt¹, Cristina Cruz^{2,3}, John J. Krais¹, Joseph Nacson¹,

Downloaded from <http://ajci.jci.org/>

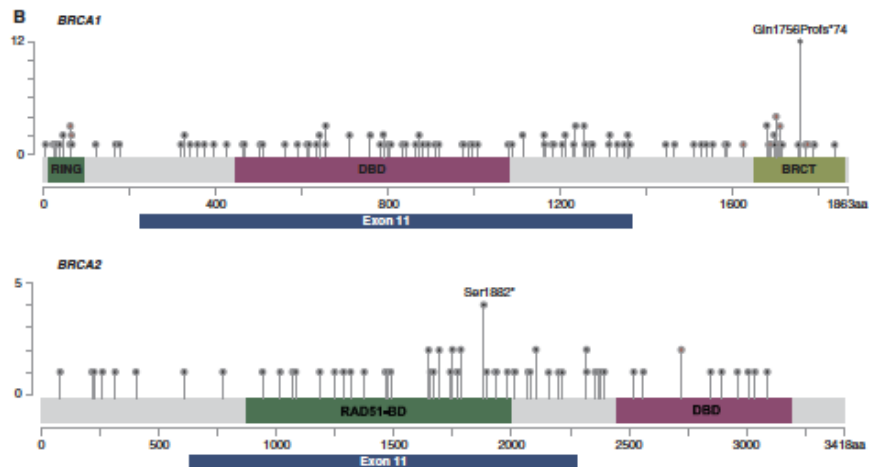
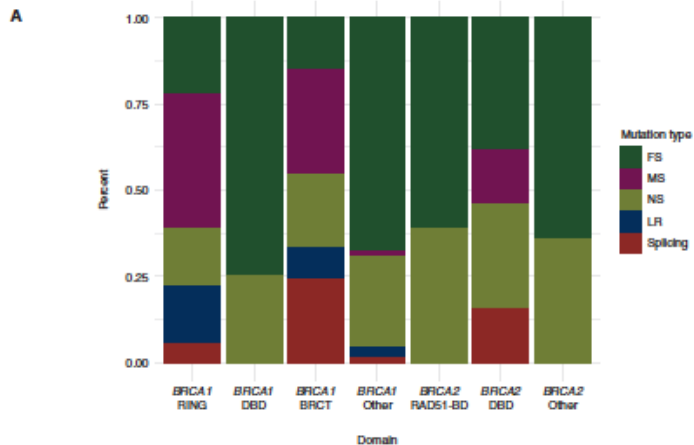
BRCA1^{185delAG} tumors may acquire therapy resistance through expression of RING-less BRCA1

Rinske Drost,¹ Kiranjit K. Dhillon,² Hanneke van der Gulden,¹ Ingrid van der Heijden,¹ Inger Brandsma,³ Cristina Cruz,^{4,5}

ORIGINAL ARTICLE

Association of location of *BRCA1* and *BRCA2* mutations with benefit from olaparib and bevacizumab maintenance in high-grade ovarian cancer: phase III PAOLA-1/ENGOT-ov25 trial subgroup exploratory analysis

S. I. Labidi-Galy^{1,2,7†}, M. Rodrigues^{2,4,5†}, J. L. Sandoval^{1,7}, J. E. Kurtz^{2,6}, F. Heitz^{7,8,9}, A. M. Mosconi^{11,11}, I. Romero^{12,13}, U. Denison^{14,15}, S. Nagao^{14,17}, I. Vergote^{18,19}, G. Parma^{20,21}, T. J. Nøttrup^{22,23}, E. Rouleau²⁴, G. Garnier^{25,26}, A. B-Balat²⁷, C. Zamagni²⁸, C. Martín-Lorente^{12,10}, E. Pujade-Lauraine^{1,6}, A. Fiévet²⁴ & I. L. Ray-Coquard^{2,6,11,12}



Association of location of *BRCA1* and *BRCA2* mutations with benefit from olaparib and bevacizumab maintenance in high-grade ovarian cancer: phase III PAOLA-1/ENGOT-ov25 trial subgroup exploratory analysis

S. I. Labidi-Galy^{1,2,7*}, M. Rodrigues^{3,4,5*}, J. L. Sandoval^{1,7}, J. E. Kurtz^{2,6*}, F. Hellz^{7,8,9*}, A. M. Moconi^{10,11}, I. Romero U. Denison^{12,13}, S. Nagai^{14,15}, L. Vergote^{16,17}, G. Parma^{18,19}, T. J. Nattrop^{20,21}, E. Rouleau²², G. Gamier^{23,24}, A. El-Balat^{1,25,26}, C. Zamagni²⁷, C. Marin-Lorente^{13,28}, E. Pujade-Lauraine¹³, A. Févet²⁴ & L. L. Ray-Coquard^{1,6,13,27}

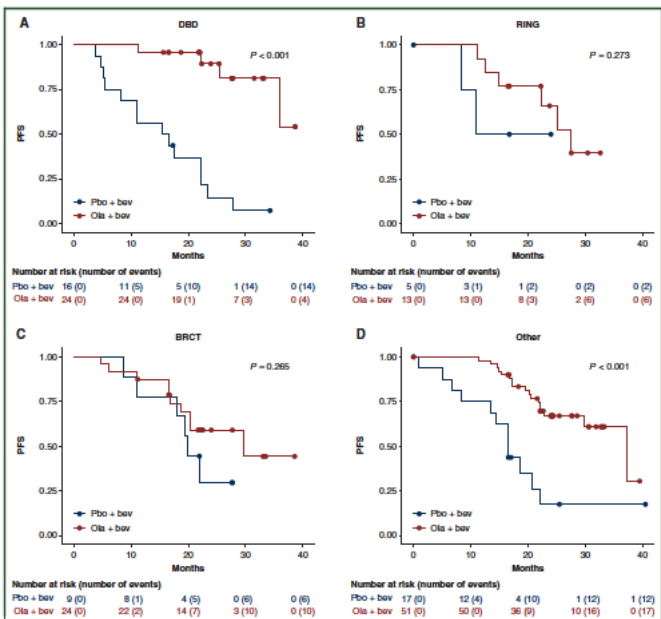
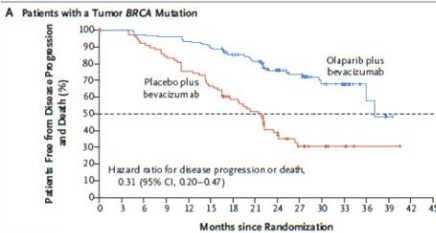


Figure 2. PFS according to the location of mutations in *BRCA1*. (A) DBD. (B) RING. (C) BRCT. (D) Other locations. Bev, bevacizumab; BRCT, C-terminal domain of *BRCA1*; DBD, DNA-binding domain; Ola, olaparib; Pbo, placebo; PFS, progression-free survival; RING, Really Interacting New Gene.

BRCA1



No. at Risk
Olaparib plus bevacizumab
Placebo plus bevacizumab

No parece que el tipo de mutación sean predictor de no actividad de ola-beva

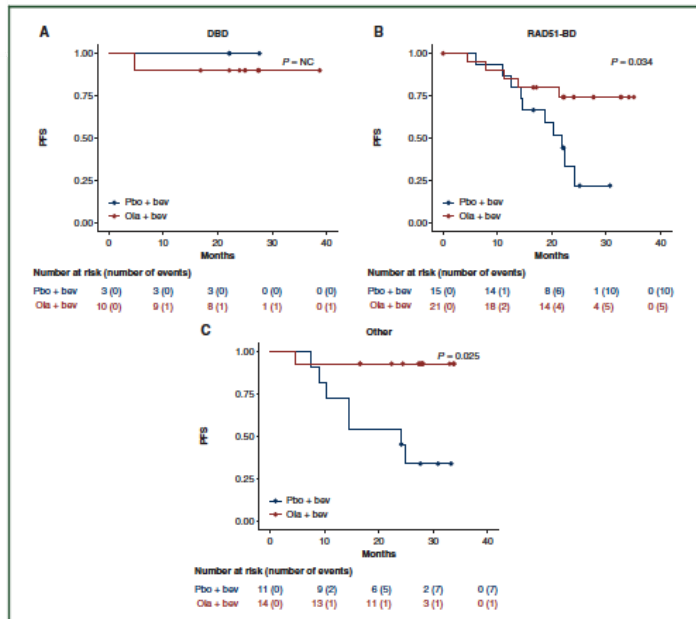


Figure 3. PFS according to the location of mutations in *BRCA2*. (A) DBD. (B) RAD51-BD. (C) Other locations. Bev, bevacizumab; DBD, DNA-binding domain; NC, not calculated; Ola, olaparib; Pbo, placebo; PFS, progression-free survival; RAD51-BD, RAD51-binding domain.

BRCA2

Association of location of *BRCA1* and *BRCA2* mutations with benefit from olaparib and bevacizumab maintenance in high-grade ovarian cancer: phase III PAOLA-1/ENGOT-ov25 trial subgroup exploratory analysis

S. L. Labidi-Galy^{1,2,3}, M. Rodrigues^{4,5,6}, J. L. Sandoval^{7,8}, J. E. Kurtz^{9,10}, F. Hellz^{11,12}, A. M. Moconi^{13,14}, J. Romero^{15,16}, U. Denison^{17,18}, S. Nagao^{19,20}, L. Vergote^{21,22}, G. Parma^{23,24}, T. J. Natrup^{25,26}, E. Rouleau²⁷, G. Garnier^{28,29}, A. B. Balat^{30,31,32}, C. Zamagni³³, C. Martin-Lorente^{34,35}, E. Pujade-Lauraine³⁶, A. Révet³⁷ & L. L. Ray-Coquard^{38,39,40}

Table 2. PFS according to the location of mutations in *BRCA1* and *BRCA2*

	Region (AA)	Median PFS, placebo (95% CI)	Median PFS, olaparib (95% CI)	24-month PFS, placebo (95% CI)	24-month PFS, olaparib (95% CI)	Placebo events (cases)	Olaparib events (cases)	HR (95% CI)	P
Gene									
<i>BRCA1</i> (n = 159)		17.6	36	0.2 (0.11-0.39)	0.7 (0.61-0.79)	34 (47)	37 (112)	0.26 (0.16-0.41)	<0.001
<i>BRCA2</i> (n = 74)		22.2	NR	0.5 (0.34-0.73)	0.84 (0.73-0.96)	17 (29)	7 (45)	0.22 (0.09-0.54)	0.001
Functional domain of <i>BRCA1</i>									
RING (n = 18)	8-96	11	27.4	0.5 (0.19-1)	0.66 (0.43-1)	2 (5)	6 (13)	0.38 (0.07-2.13)	0.273
DBD (n = 40)	452-1092	16	NR	0.15 (0.04-0.51)	0.89 (0.76-1)	14 (16)	4 (24)	0.08 (0.02-0.28)	<0.001
BRCT (n = 33)	1646-1736	19.9	29.6	0.3 (0.1-0.88)	0.59 (0.42-0.84)	6 (9)	10 (24)	0.55 (0.2-1.56)	0.265
Other (n = 68)	1760-1855	16.6	37.2	0.18 (0.05-0.59)	0.67 (0.55-0.82)	12 (17)	17 (51)	0.24 (0.11-0.51)	<0.001
Functional domain of <i>BRCA2</i>									
RADS1-8D (n = 36)	900-2000	21.7	NR	0.33 (0.15-0.75)	0.74 (0.57-0.97)	10 (15)	5 (21)	0.31 (0.11-0.92)	0.034
DBD (n = 13)	2459-3190	NR	NR	1 (1-1)	0.9 (0.73-1)	0 (3)	1 (10)	NC	NC
Other (n = 25)		24	NR	0.55 (0.32-0.94)	0.93 (0.8-1)	7 (11)	1 (14)	0.09 (0.01-0.75)	0.025
Exon 11 mutation									
Yes (n = 123)		17.6	37.2	0.24 (0.14-0.43)	0.78 (0.68-0.89)	34 (47)	18 (76)	0.2 (0.11-0.36)	<0.001
No (n = 110)		19.9	NR	0.45 (0.3-0.68)	0.7 (0.6-0.81)	17 (29)	26 (81)	0.41 (0.22-0.75)	0.004

AA, amino acid; BRCT, C-terminal domain of *BRCA1*; CI, confidence interval; DBD, DNA-binding domain; HR, hazard ratio; NC, not calculated; NR, not reached; PFS, progression-free survival; RADS1-8D, RADS1-binding domain; RING, Really Interesting New Gene.

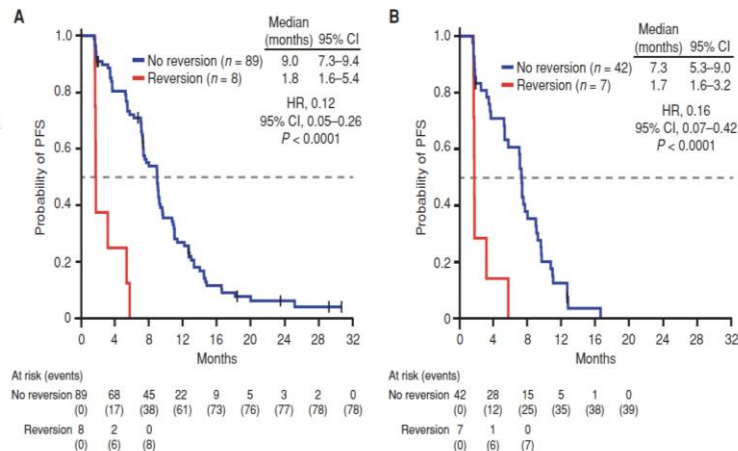
RESEARCH BRIEF

BRCA Reversion Mutations in Circulating Tumor DNA Predict Primary and Acquired Resistance to the PARP Inhibitor Rucaparib in High-Grade Ovarian Carcinoma

Kevin K. Lin¹, Maria I. Harrell², Amit M. Oza³, Ana Daknin⁴, Isabelle Ray-Coquard⁵, Anna V. Tinker⁶, Elena Helman⁷, Marc R. Radke⁸, Carmen Say⁹, Lan-Thanh Vo⁹, Elaine Mann⁹, Jeffrey D. Isaacson⁹, Lara Maloney¹⁰, David M. O'Malley¹¹, Setsuko K. Chambers¹², Scott H. Kaufmann¹³, Clare L. Scott¹⁴, Gottfried E. Konecny¹⁵, Robert L. Coleman¹⁶, James X. Sun¹⁷, Heidi Giordano¹⁸, James D. Brenton¹⁹, Thomas C. Harding¹, Iain A. McNeish¹⁹, and Elizabeth M. Swisher²

BRCA Reversion Mutations and Resistance to PARP Inhibitor

RESEARCH BRIEF



Pacientes BRCA mutadas del estudio ARIEL2 (fase II) tratadas con rucaparib, se les recogio muestra de tumor y ctDNA pre-tto y post-tto

En la muestra de ctDNA pretto: 18% de las platino-refractarias 13% de las platino-resistentes 2% de las platino-sensibles

Tenian reversion de la mutación BRCA

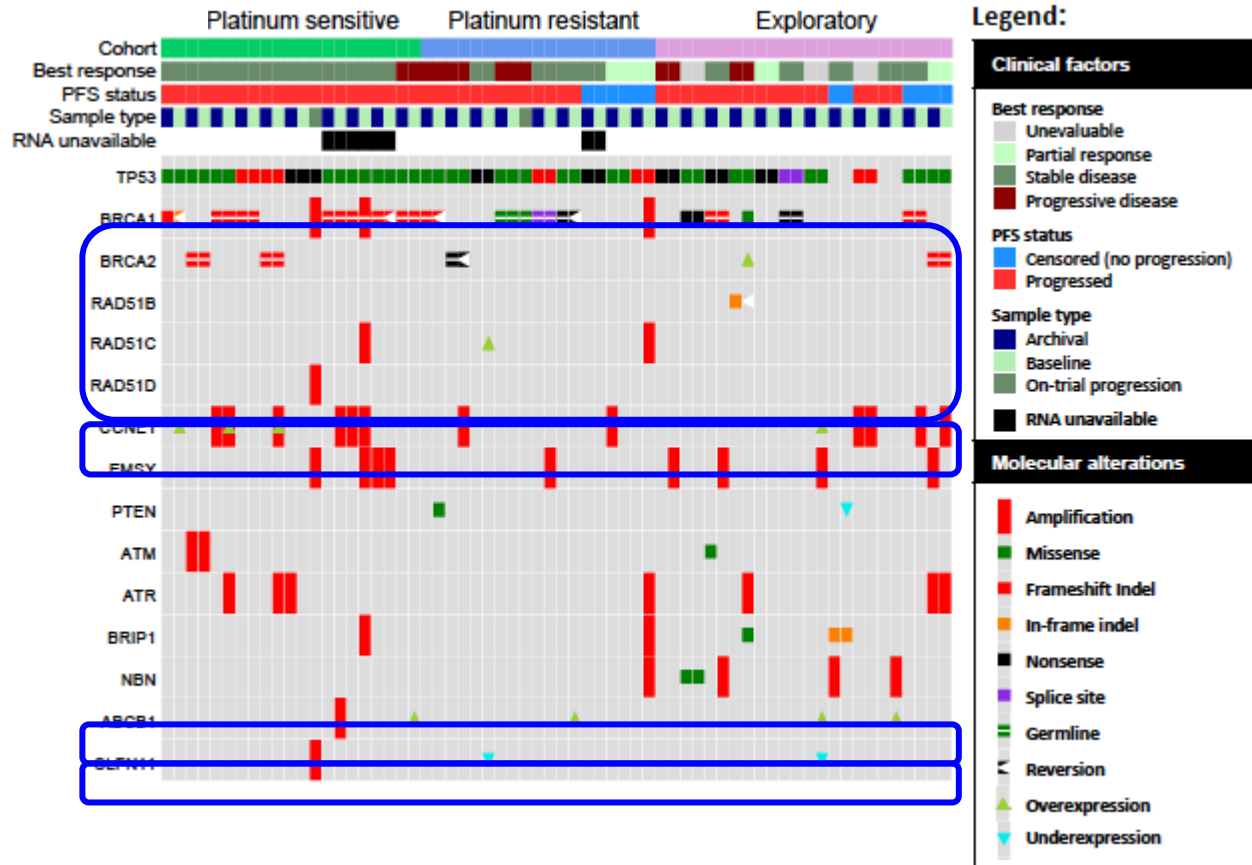
Las pacientes con reversion tenían una PFS mucho más corta a rucaparib que las que no habían revertido BRCA antes de iniciar rucaparib

Clinical Cancer Research

EVOLVE: A Multicenter Open-Label Single-Arm Clinical and Translational Phase II Trial of Cediranib Plus Olaparib for Ovarian Cancer after PARP Inhibition Progression

Stephanie Lheureux, Ana Oaknin, Swati Garg, et al.

Clin Cancer Res. Published OnlineFirst May 22, 2020.



EVOLVE trial

Olaparib + Cediranib
treatment



Array

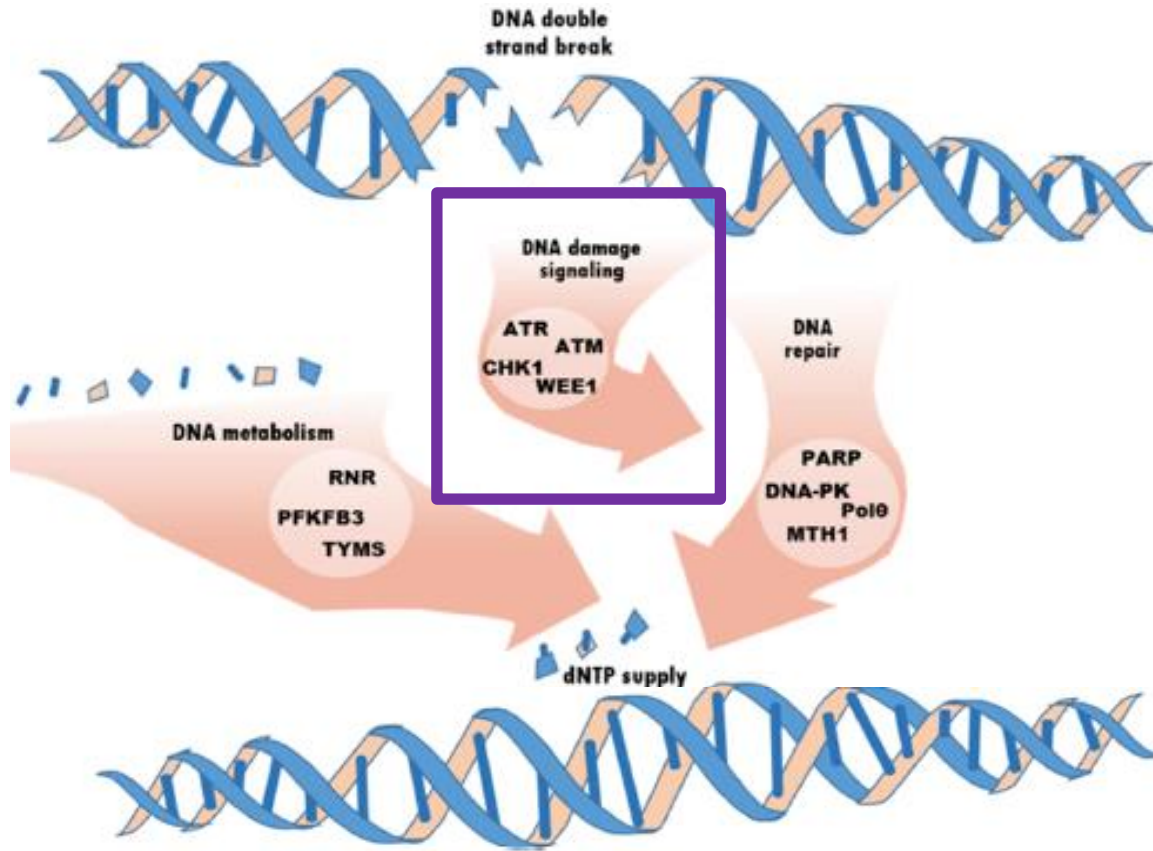
Array

**BRCA and RAD51c mutations reversion
19%**

CCNE1 amplification 16%

ABCB1 upregulation 15%

SLFN11 downregulation 11%



DNA damage signalling

Cell cycle



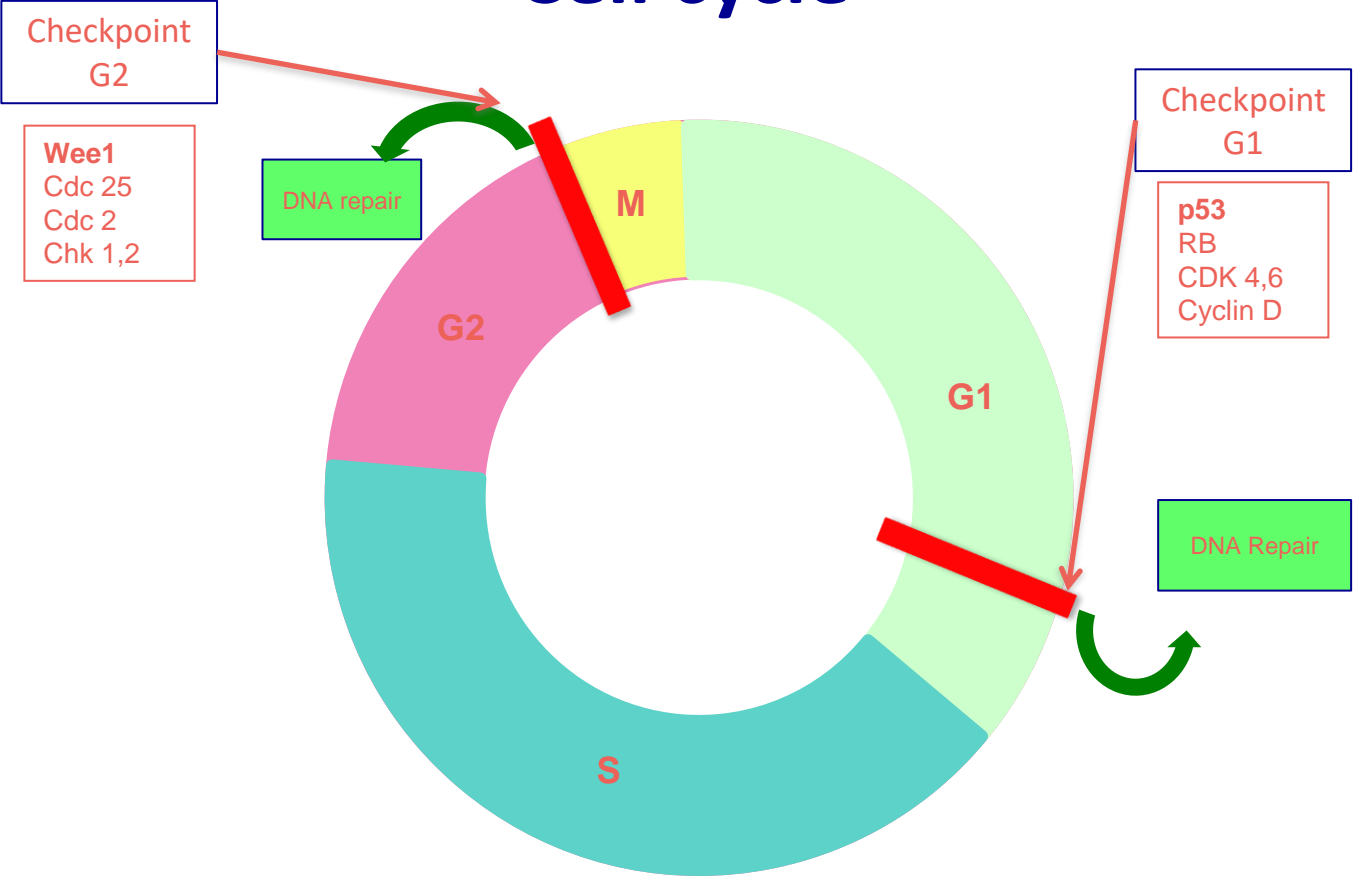
DNA damage signalling:
ATM, ATR

Cell cycle arrest
(The brake): p53,
wee1

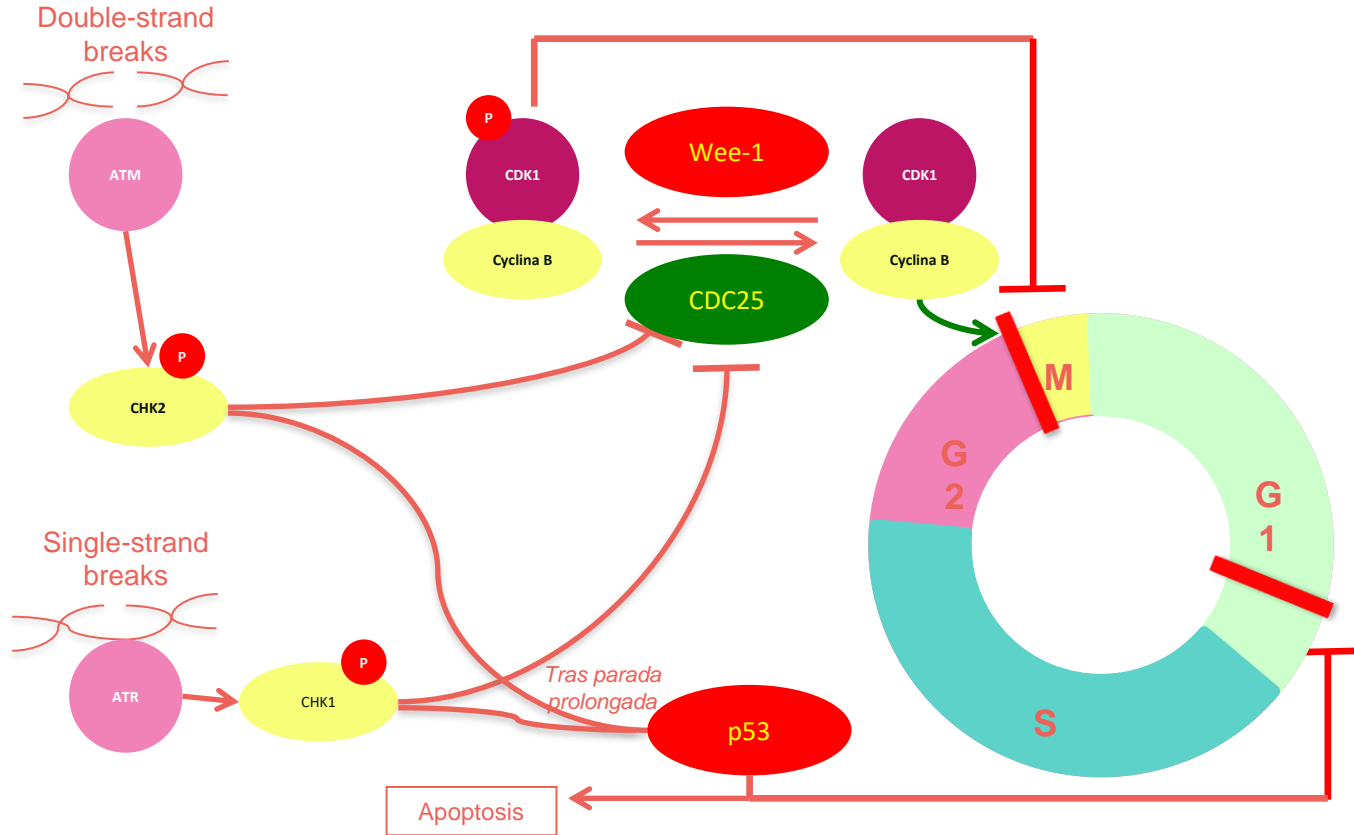
Repair mechanisms:
HR / BER / NHEJ



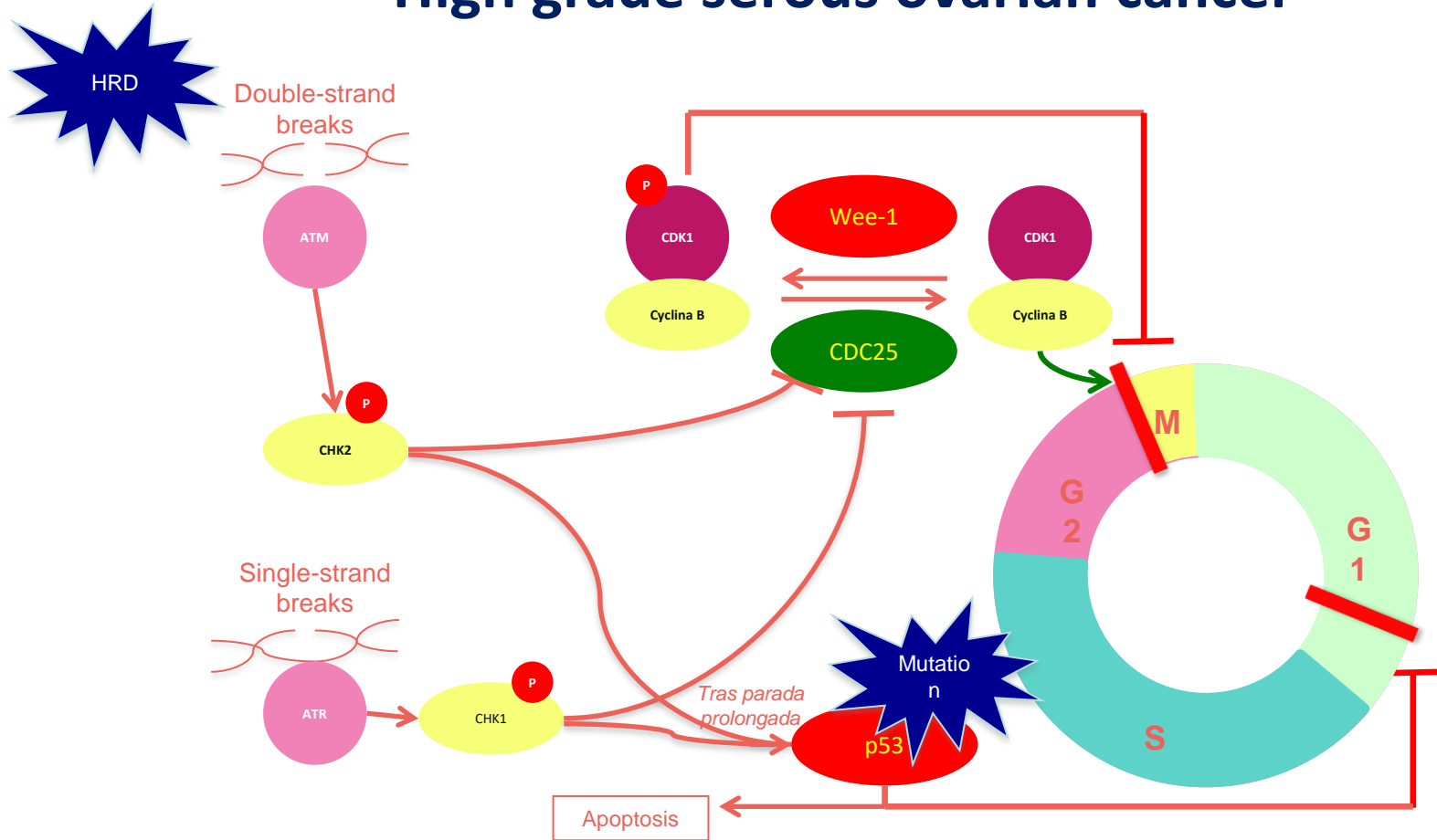
Cell cycle



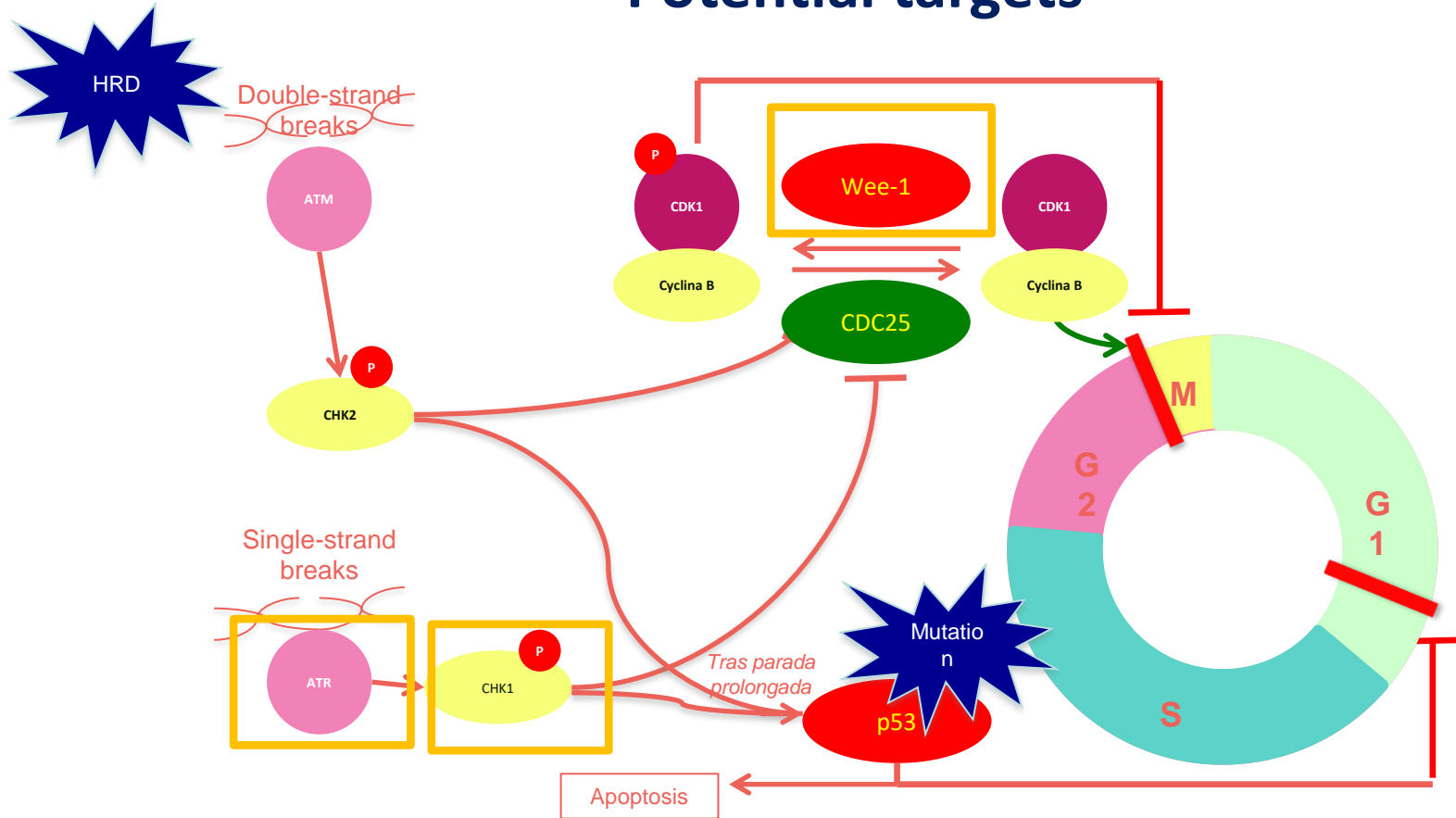
Cell cycle and DNA repair system

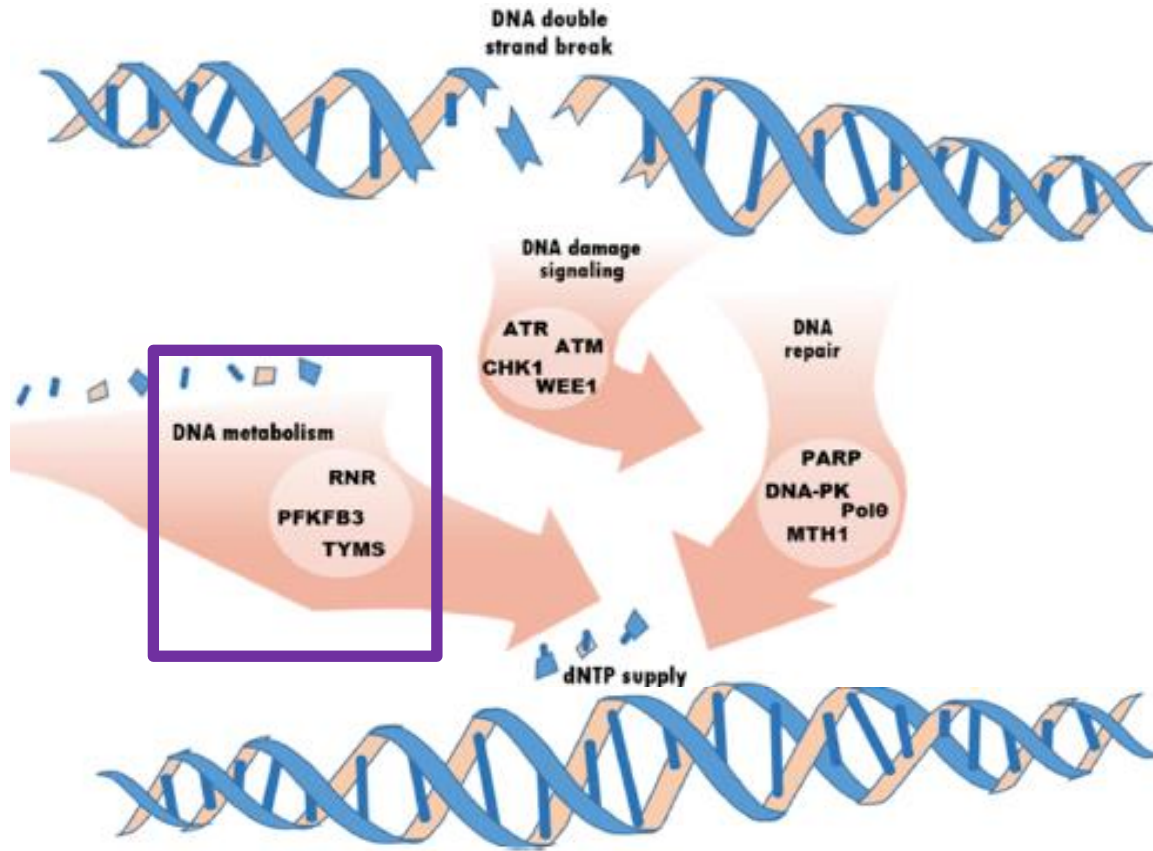


High grade serous ovarian cancer



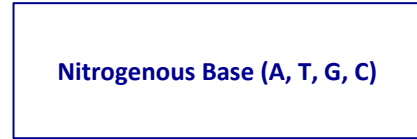
Potential targets



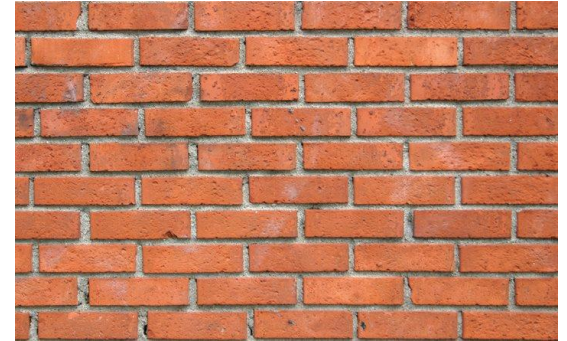


dNTP deoxyriboNucleosideTriPhosphate

dNTP



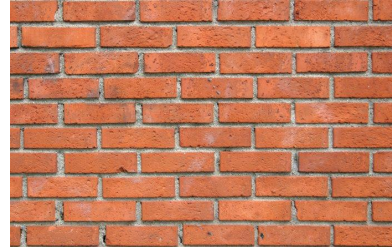
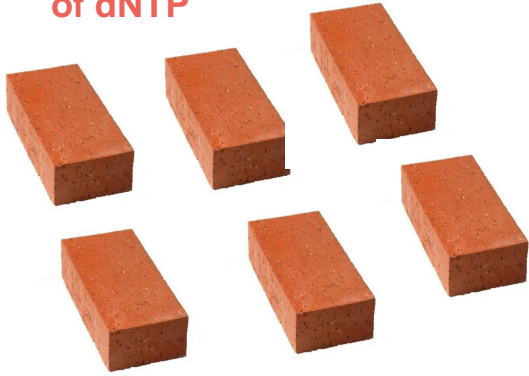
dNTP



DNA

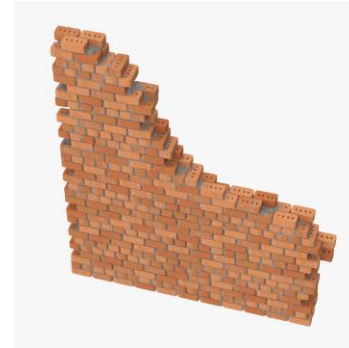
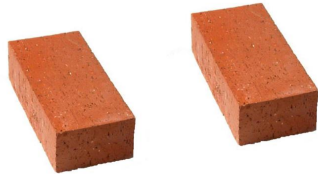
dNTP pool

Increased pool
of dNTP



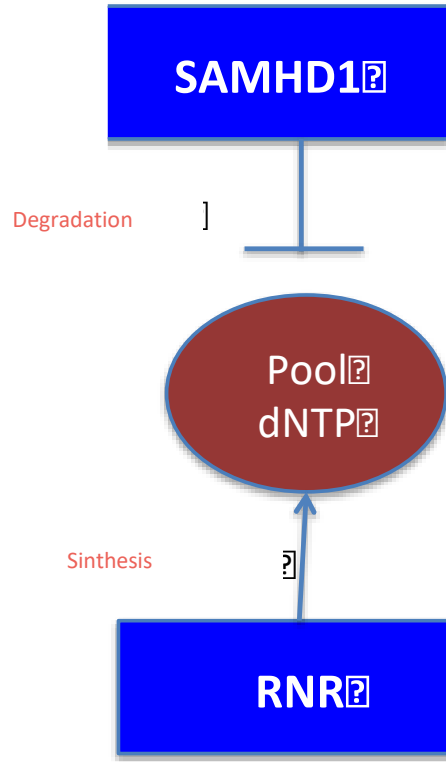
DNA

Decreased pool
of dNTP



DNA

dNTP pool balance



*RNR: Ribonucleotide Reductase
SAMHD1: Sterile Alpha Motive and HD domain protein 1*

EVOLVE trial

Olaparib + Cediranib
treatment



Array

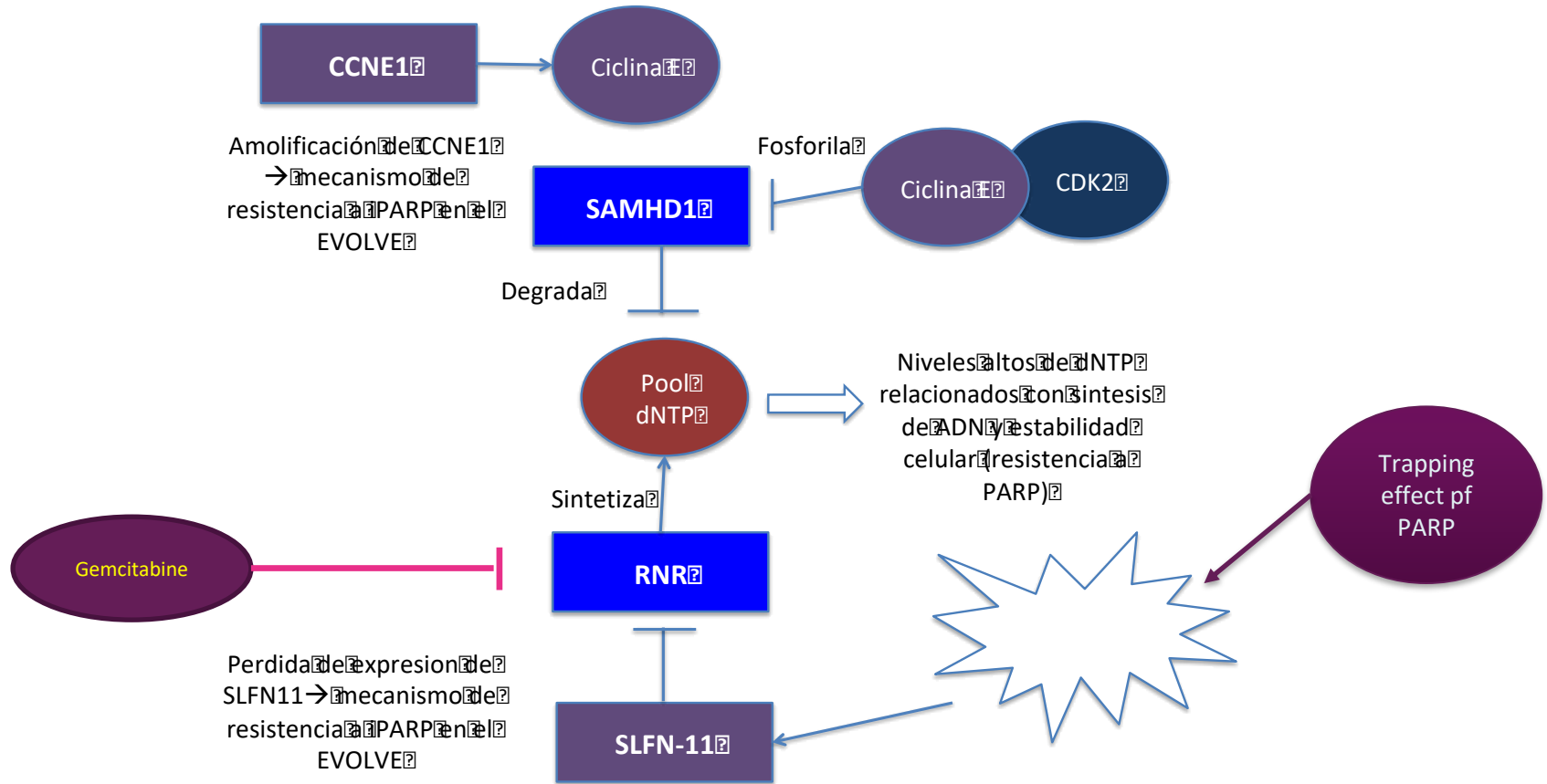
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BRCA and RAD51c mutations reversion
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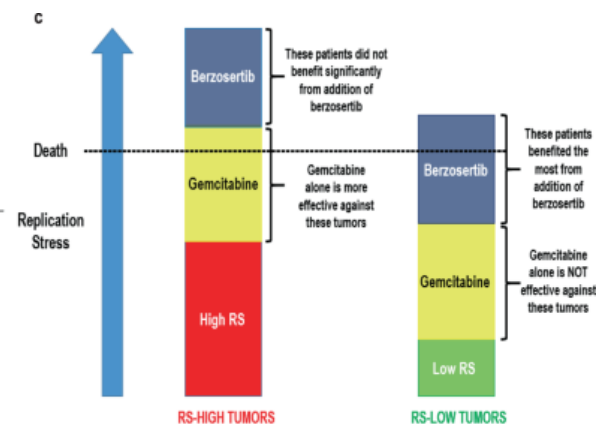
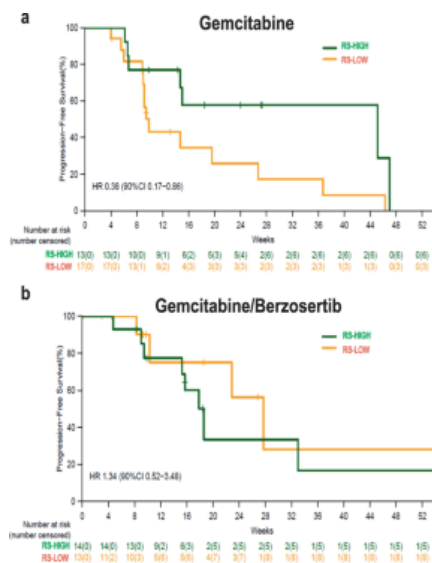
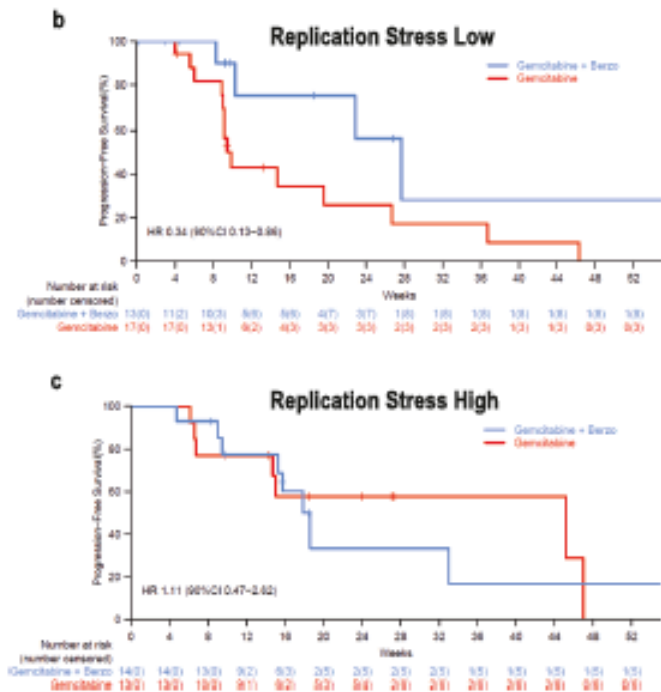
ABCB1 upregulation 15%

SLFN11 downregulation 11%



A Replication stress biomarker is associated with response to gemcitabine versus combined gemcitabine and ATR inhibitor therapy in ovarian cancer

Panagiotis A. Konstantinopoulos¹✉, Alexandre André B. da Costa^{2,10}, Doga Gulhan^{3,10}

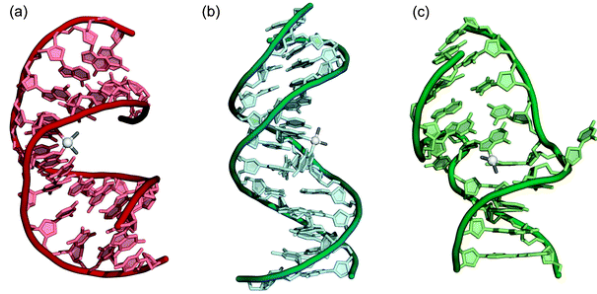




**¿Existe una resistencia
cruzada PARPi-platino?**

Carboplatin + PARPi

Platinum DNA damage



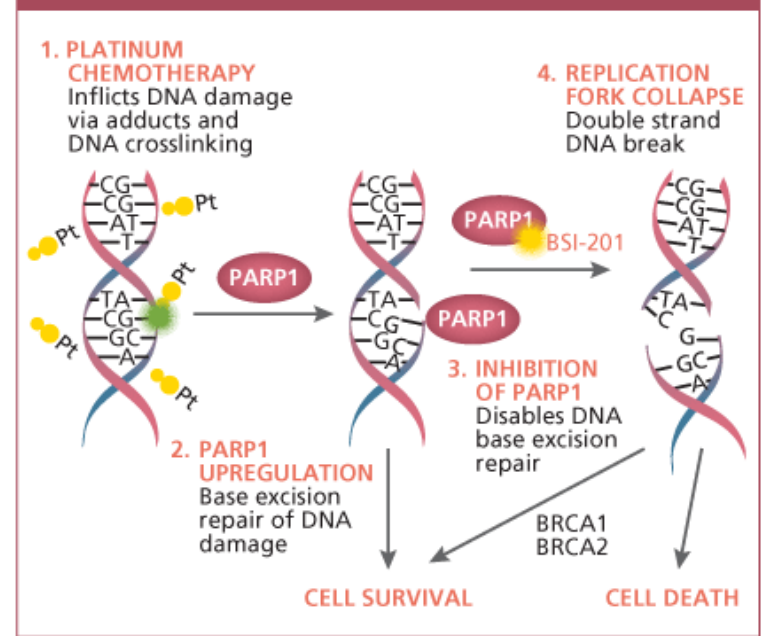
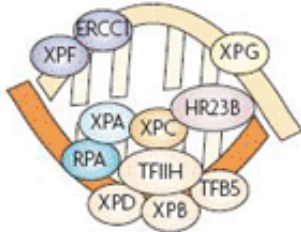
BER and Homologous recombination repair systems



Nucleoside excision repair system

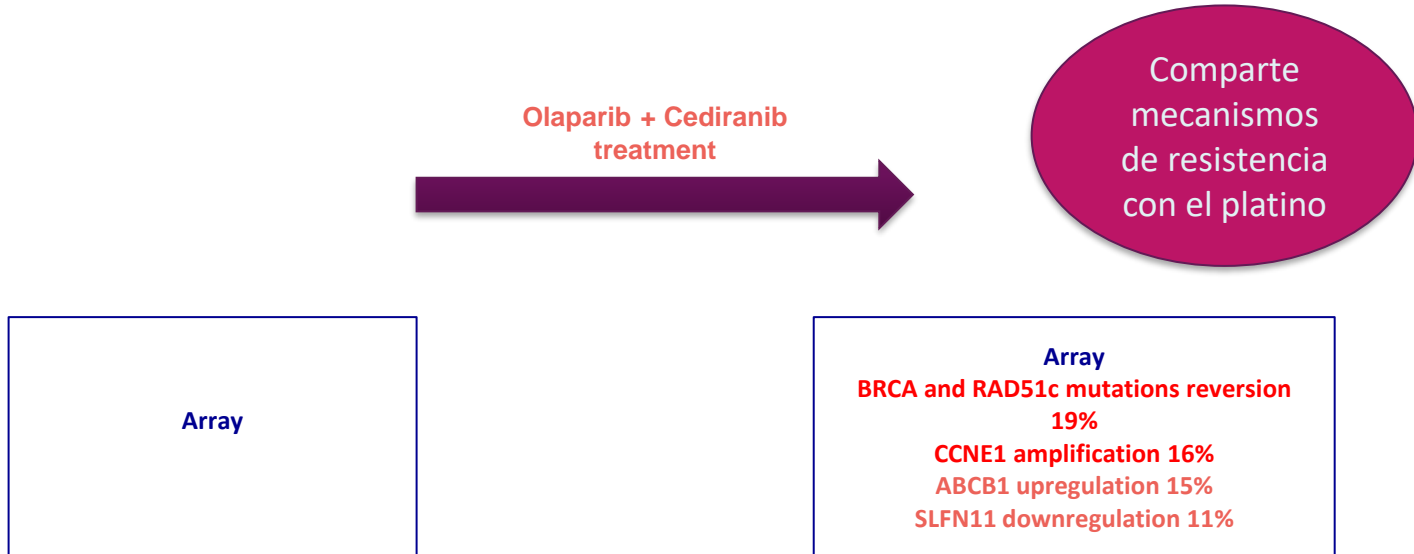


Increased nucleotide-excision repair (for example, increased ERCC1)



Kelland LI Nat Rev Cancer 2007
Reedijk J Platinum Metals Rev 2008
O'Shaughnessy J et al. JCO 2009 (abstract 3)

EVOLVE trial



¿Existe una resistencia cruzada PARPi-platino?

PARPi mechanism of resistance (EVOLVE)	Is a cross-resistant mechanism with Platinum?
HRD phenotype - BRCA mutation reversion - RAD51c reversion mutations	Yes yes
Multidrug resistance: - ABCB upregulation	No?
Cell-cycle regulation: - CCNE1 upregulation - SLFN11 downregulation	Yes ?

Efficacy of subsequent chemotherapy for patients with *BRCA1/2*-mutated recurrent epithelial ovarian cancer progressing on olaparib versus placebo maintenance: *post-hoc* analyses of the SOLO2/ENGOT Ov-21 trial

J. S. Frenel^{1,2}, J. W. Kim³, N. Aryal⁴, R. Asher⁵, D. Berton¹, L. Vidal¹, P. Pautier⁶, J. A. Ledermann⁷, R. T. Penson⁷, A. M. Oza⁸, J. Korach⁹, T. Huzarski¹⁰, S. Pignata¹¹, N. Colombo¹², T. W. Park-Simon¹³, K. Tamura¹⁴, G. S. Sonke¹⁵, A. E. Freimund¹⁶, C. K. Lee¹⁷ & E. Pujade-Lauraine¹⁷

Table 1. Characteristics of patients who received chemotherapy as subsequent therapy regimen, with comparison between olaparib- and placebo-treated patients

Characteristics	Overall population (N = 347)			Platinum-based cohort (N = 96)			Non-platinum-based cohort (N = 51)			SOLO2 population (N = 295)	
	Olaparib n = 78	Placebo n = 69	P value	Olaparib n = 54	Placebo n = 42	P value	Olaparib n = 24	Placebo n = 27	P value	Olaparib n = 196	Placebo n = 99
Mean (SD) age, years	57 (40-83)	56 (39-75)	0.41	57 (40-83)	57 (40-75)	0.58	56 (45-66)	55 (39-70)	0.65	56 (51-63)	56 (49-63)
ECOG, n (%)											
Normal activity	62 (81)	54 (78)	0.61	46 (87)	31 (74)	0.11	16 (67)	23 (85)	0.12	162 (84)	77 (78)
Restricted activity	15 (19)	15 (22)		7 (13)	11 (26)		8 (33)	4 (15)		32 (16)	22 (22)
Missing	1			1						2	0
Primary tumor location, n (%)											
Ovary	65 (83)	59 (86)	0.86	45 (83)	36 (86)	0.67	20 (83)	23 (85)	0.55	164 (84)	86 (87)
Fallopian	5 (6)	3 (4)		5 (9)	2 (5)		0 (0)	1 (4)		13 (7)	4 (4)
Other	8 (10)	7 (10)		4 (7)	4 (10)		4 (17)	3 (11)		19 (9)	9 (9)
Histology, n (%)											
Serous	75 (96)	63 (91)	0.37	53 (98)	38 (90)	0.22	22 (92)	25 (93)	0.90	183 (93)	86 (87)
Endometrioid	3 (4)	5 (7)		1 (2)	3 (7)		2 (8)	2 (7)		9 (5)	8 (8)
Others	0 (0)	1 (1)		0 (0)	1 (2)		0 (0)	0 (0)		4 (2)	5 (5)
Myriad BRCA status, n (%)											
BRCA1	53 (71)	43 (63)	0.43	36 (69)	24 (59)	0.28	17 (74)	19 (70)	0.78	132 (69)	61 (64)
BRCA2	22 (29)	25 (37)		16 (31)	17 (41)		6 (26)	8 (30)		58 (31)	35 (36)
Missing	3	1		2	1		1	12		6	3
Previous platinum-free interval, n (%)											
6-12 months	40 (51)	33 (48)	0.68	28 (52)	20 (48)	0.68	12 (50)	14 (52)	0.89	79 (40)	40 (40)
>12 months	38 (49)	36 (52)		26 (48)	22 (52)		12 (50)	13 (48)		117 (60)	59 (60)
Previous platinum-based regimen, n (%)											
2	41 (53)	37 (54)	0.15	30 (56)	23 (55)	0.49	11 (46)	14 (52)	0.25	110 (56)	62 (63)
3	28 (36)	17 (25)		18 (33)	11 (26)		10 (42)	6 (22)		60 (31)	20 (20)
>3	9 (12)	15 (22)		6 (11)	8 (19)		3 (13)	7 (26)		25 (13)	17 (17)
Disease status at inclusion in the SOLO2 trial, n (%)											
Partial response	60 (77)	41 (59)	0.02	40 (74)	23 (55)	0.05	20 (83)	18 (67)	0.17	91 (46)	47 (47)
Complete response	18 (23)	28 (41)		14 (26)	19 (45)		4 (17)	9 (33)		105 (54)	52 (53)
Prior use of bevacizumab, n (%)											
Yes	14 (18)	18 (26)	0.23	12 (22)	14 (33)	0.22	2 (8)	4 (15)	0.47	33 (17)	20 (20)
No	64 (82)	51 (74)		42 (88)	28 (77)		22 (92)	23 (85)		163 (83)	79 (80)

Bold values are statistically significant.
 ECOG, Eastern Cooperative Oncology Group; SD, standard deviation.

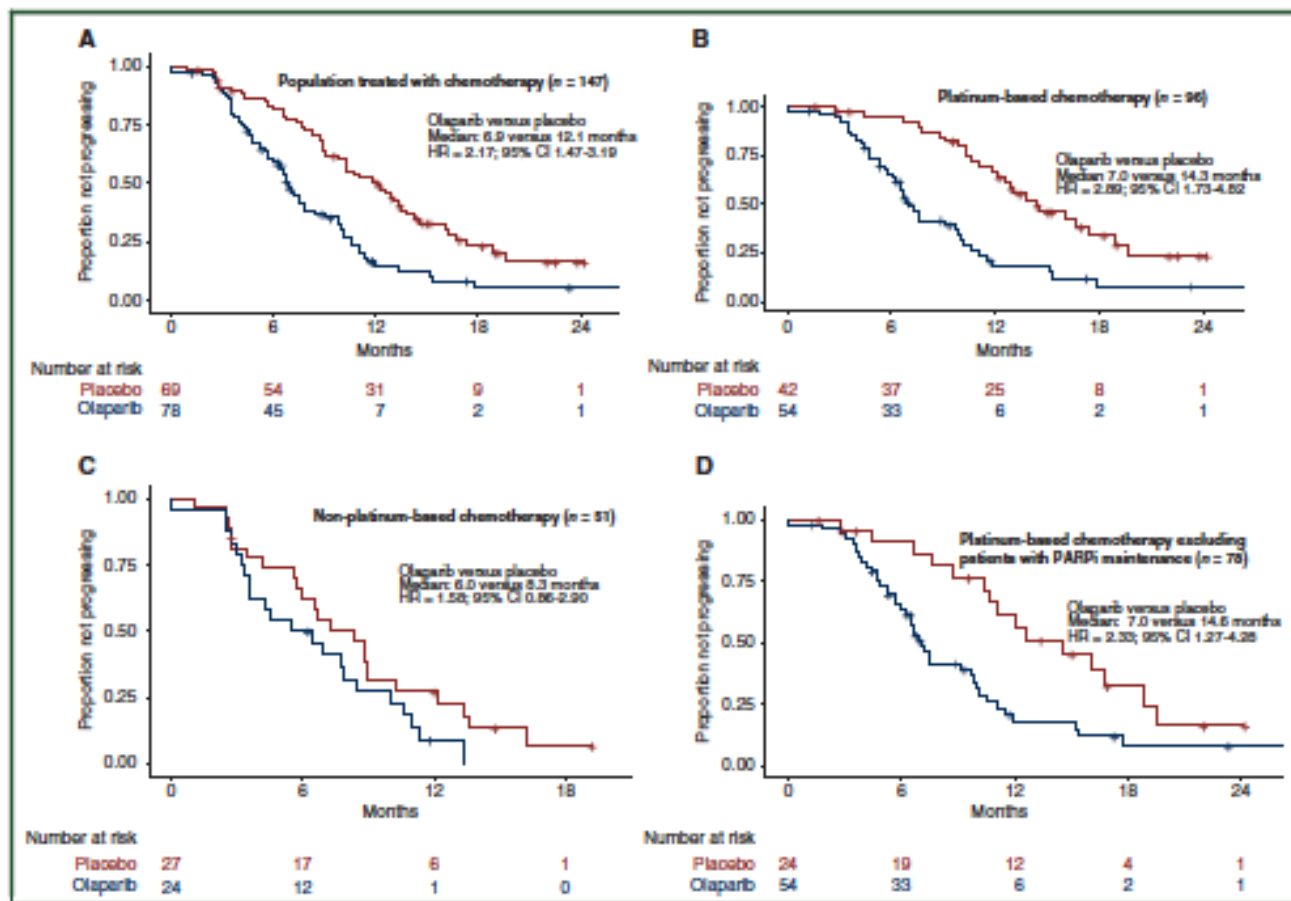


Figure 2. Time to second progression according to subsequent therapy type. CI, confidence interval; HR, hazard ratio; PARPi, poly (ADP-ribose) polymerase inhibitor.

¿Qué podemos hacer tras progresión platinosensible a PARPi?

¿Evitar reintroducción de platinos?	¿Reintroducir platino?
Trabectedina-DLP	Reintroducir platino y luego reintroducir iPARP (OREO)
Nuevas drogas?: <ul style="list-style-type: none">- Mirvetuximab- Otros	Reintroducir platino y asociar Bevacizumab (Carbo-Caelyx-Bev o Carbo-Gem-Beva)
Si oligometastática: citorreducción secundaria y mantener mismo iPARP???	Citorreducción secundaria → platino + una de las dos anteriores