

Retrospective Analysis of Germline and Somatic Aberrations and Corresponding Outcomes Associated with the Use of Poly(ADP-ribose) Polymerase (PARP) Inhibitors

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Disclosure

• The speaker has no actual or potential conflict of interest in relation to this presentation.

• I will be discussing off-label indications



Objective

 Describe patient populations which may benefit from poly(ADP-ribose) polymerase (PARP) inhibitor therapy based on germline and somatic aberrations.



PARP Inhibitors

- Indicated for treatment and/or maintenance therapy for ovarian or breast cancer
 - Olaparib
 - Niraparib
 - Rucaparib
 - Talazoparib
 - Veliparib (investigational)

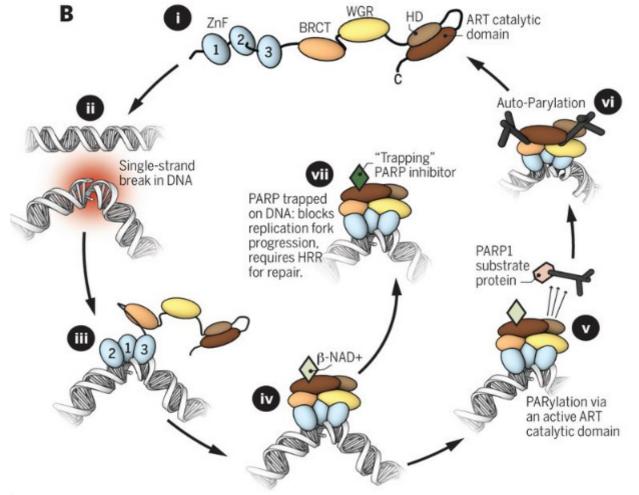
Zejula [package insert] 2018 Talzenna [package insert] 2018 Lynparza [package insert] 2018 Rubraca [package insert] 2018

- Generally well-tolerated
 - Decreased blood counts
 - Nausea/vomiting/ diarrhea
 - Fatigue
 - Abdominal pain



PARP: poly(ADP-ribose) polymerase

PARP Inhibitor Mechanism of Action



Adapted from: Lord Science 2017

PARP: poly(ADP-ribose) polymerase





PARP Inhibitors



Primary Cancer	Treatment Arms	Patient Cohorts	Outcome
Metastatic breast (n=302)	Olaparib vs standard therapy	gBRCA	PFS: 7.0 vs 4.2 months
Recurrent	Niraparib vs placebo for maintenance	gBRCA	PFS: 21.0 vs 5.5 months
ovarian (n=533)		HR deficiency	PFS: 12.9 vs 3.8 months
Recurrent ovarian (n=196)	Rucaparib vs placebo for maintenance	gBRCA	PFS: 16.6 vs 5.4 months
		HR deficiency	PFS: 13.6 vs 5.4 months
Advanced breast (n=431)	Talazoparib vs standard therapy	gBRCA	PFS: 8.6 vs 5.6 months
Ovarian, breast, pancreas, & prostate (n=298)	Olaparib (single arm)	gBRCA	ORR: 26.2% Stable disease: 42%

Robson NEJM 2017; Mirza NEJM 2016; Coleman Lancet 2017; Litton NEJM 2018; Kaufman J Clin Oncol 2014

PARP: poly(ADP-ribose) polymerase; PFS: progression-free survival; gBRCA: germline breast cancer susceptibility gene; HR: homologous repair; ORR: overall response rate

Rationale for Study

- PARP inhibitors have demonstrated efficacy in patients with germline BRCA mutations in several tumor types
- Preclinical data has suggested efficacy of PARP inhibitors in patients with genetic mutations in the DNA damage repair system
- Further clinical data are needed to assess efficacy of PARP inhibitors in patients with other DNA mutations which may confer PARP inhibitor sensitivity via homologous repair deficiency



Study Design

• Single-center, retrospective review

- Assessment of safety and efficacy of PARP inhibitors comparing:
 - Patients with known deleterious germline BRCA mutations
 - Patients with other known germline or somatic mutations



Study Criteria

Inclusion

- Adults age 18 or older
- Patients receiving at least one dose of a PARP inhibitor for a solid tumor malignancy at the Avera Cancer Institute at any time from January 1st, 2013 – June 30th, 2018

Exclusion

 Patients who had neither germline nor somatic tumor mutation testing results



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Objectives

• Primary

 Compare PFS in patients treated with a PARP inhibitor with germline BRCA mutations against those without germline BRCA mutations

PFS: progression-free survival; PARP: poly(ADP-ribose) polymerase; BRCA: breast cancer susceptibility gene



Objectives

- Secondary
 - Compare PFS in patients treated with a PARP inhibitor with germline BRCA against those with other known genetic mutations
 - Compare best response between cohorts
 - Assess the rate of adverse effects which required discontinuation of PARP inhibitor therapy
 - Report the tolerability of PARP inhibitors as monotherapy and in combination with other agents



PFS: progression-free survival; PARP: poly(ADP-ribose) polymerase; BRCA: breast cancer susceptibility gene

METHODS



Data Collection and Analysis

Patient charts reviewed to find PARP inhibitor utilization Further data collected to assess efficacy and tolerability of PARP inhibitor therapy Patient charts reviewed for germline and somatic tumor mutation testing results Pharmacogenomic data compared with efficacy data to determine association with specific mutations



Methods

- Patient information collected
 - Diagnosis
 - PARP inhibitor received
 - Purpose of therapy
 - Treatment
 - Maintenance
 - Prescribed dose & maximum tolerated dose
 - Duration of therapy/progression data
 - Best response to therapy
 - Germline and somatic genetic mutations



Methods

- Germline testing
 - Multigene panel
 - Individual gene testing
- Somatic tumor testing
 - Biopsy
 - Samples may be taken from either original tumor specimen or metastatic site
 - FoundationOne[®]
 - Cell-free DNA test
 - FoundationOne ACT[®], Guardant360[®]



Methods

- Mutations selected for analysis
 - ATM BRIP1 NF1
 - -BRCA1/2 -CDH1 -RAD50
 - CHEK2 CDKN2A ARID1A
 - PALB2 MSH2 ARID1B
 - PTEN
- Excluded variants of unknown significance

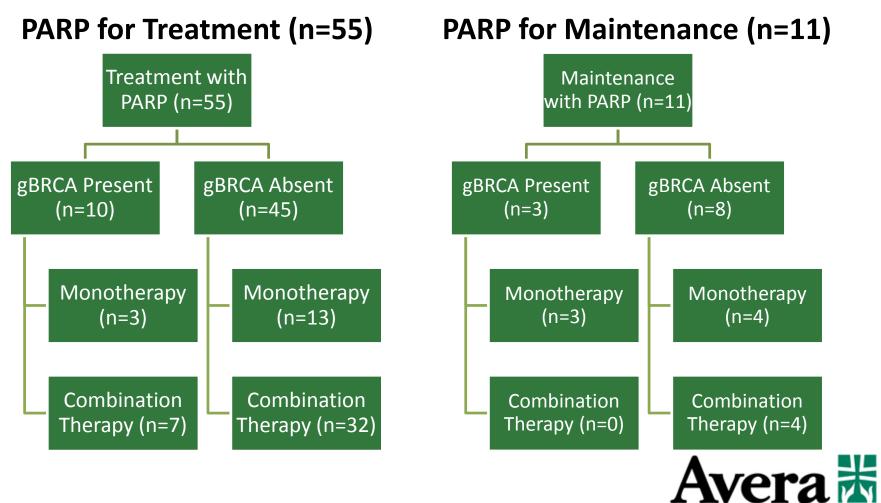


Konstantinopoulos Cancer Discov. 2015 Shimelis J Natl Cancer Inst. 2018

RESULTS



Patient Characteristics

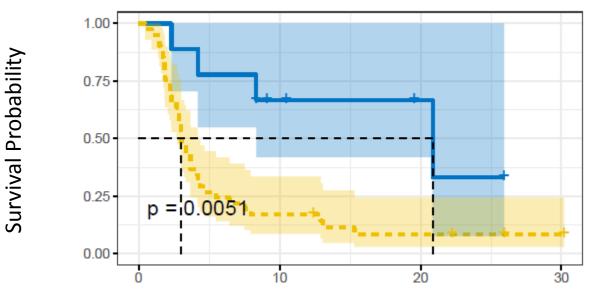


PARP: poly(ADP-ribose) polymerase



Primary Outcome

• Progression-free survival

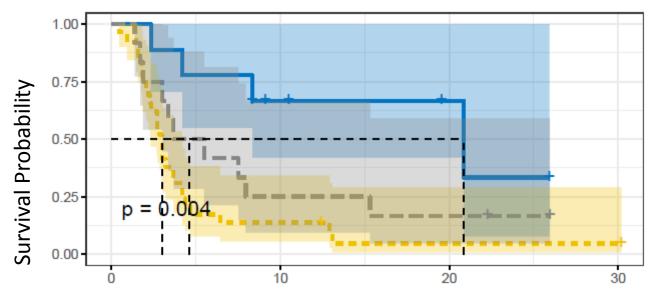


Time (months)

Cohort	PFS (months)	95% CI
Germline BRCA Mutation (n=9)	20.9	8.34-NR
Mo Germline BRCA Mutation (n=41)	3.0	2.73-4.21



• Progression-free survival by BRCA mutation status



Time (months)

Cohort	PFS (months)	95% CI
Germline BRCA Mutation (n=9)	20.9	8.34-NR
Somatic BRCA Mutation (n=12)	4.6	3.02-NR
No BRCA Mutation (n=29)	3.0	2.33-4.17

Secondary Outcome Avera

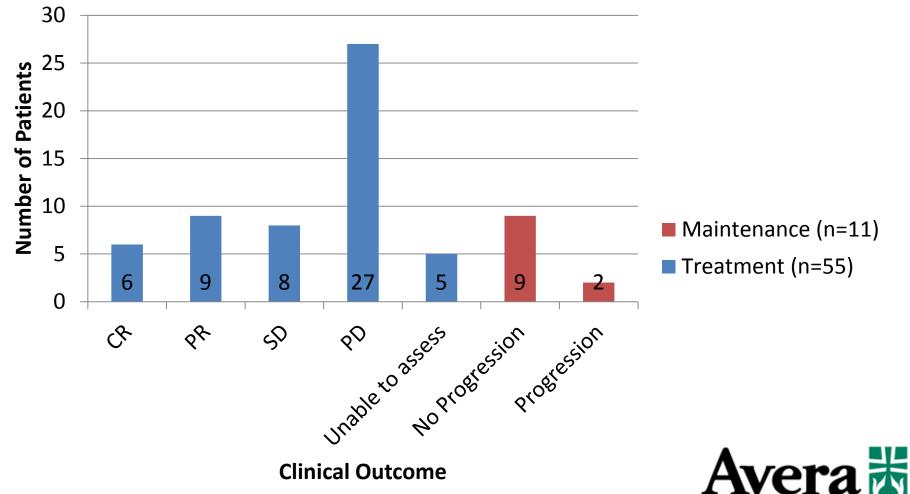
• Progression-free survival: exploratory analyses

Cohort	PFS (months)	95% CI	P-value
BRCA1/BRCA2 Analysis			
Germline BRCA2 Mutation (n=6)	20.86	NA-NA	0.0094
Germline BRCA1 Mutation (n=3)	4.21	2.33-NA	
Somatic BRCA Mutation (n=12)	4.58	3.02-NA	
No Germline BRCA Mutation (n=41)	2.99	2.33-4.17	
Monotherapy versus Combination Therapy			
mBRCA, Combination therapy (n=13)	15.31	3.68-NA	0.0024
mBRCA, Monotherapy (n=8)	6.72	3.02-NA	
wtBRCA, Combination therapy (n=23)	2.99	2.69-4.37	
wtBRCA, Monotherapy (n=6)	1.81	1.58-NA	

mBRCA: mutant BRCA, either germline or somatic; wtBRCA: wild-type BRCA



Secondary Outcome: Best Response



CR: complete response; PR: partial response; SD: stable disease; PD: progressive disease

Secondary Outcomes Avera

Early Discontinuation of Therapy				
Therapy	Concomitant Agents	Diagnosis	Duration of Therapy	Discontinuation Reason
Niraparib 200 mg daily	None	IIIC recurrent ovarian	2 weeks	Thrombocytopenia
Olaparib 200 mg capsule BID given days 1-10 of a 21-day cycle	Paclitaxel 80 mg/m ² weekly & carboplatin AUC 5 every 3 weeks	IA triple- negative invasive ductal carcinoma	3 days	Rash which resolved after drug discontinuation
Olaparib 200 mg tablet BID given days 2-8 of a 21- day cycle	Paclitaxel 80 mg/m ² & carboplatin AUC 1.5 weekly	IA triple- negative invasive ductal carcinoma	8 weeks	Thrombocytopenia, neutropenia
Niraparib 300 mg daily	None	IIIC recurrent ovarian	3 days	Malaise
Niraparib 100 mg daily	Nab-paclitaxel 62.5 mg/m ² & gemcitabine 500 mg/m ² weekly	IV pancreatic	39.5 weeks	Fatigue

Secondary Outcomes Avera

- Average dose intensity
 - Monotherapy: Percentage of FDA-recommended dose which was tolerated by patients
 - Combination therapy: Percentage of prescribed dose which was tolerated by patients

	Monotherapy	Combination Therapy	Total
Olaparib (n=40)	87%	94%	92%
Niraparib (n=22)	81%	86%	84%
Rucaparib (n=4)	83%	N/A	83%

Discussion

Strengths

- Provides an assessment of clinical outcomes for patients with genomic mutations other than germline BRCA
- Assessed tolerability of PARP inhibitors in a clinical setting

Limitations

- Insufficient patient population to assess outcomes for individual PARP inhibitors
- Retrospective data
- Single-center analysis



Discussion

- Progression-free survival (PFS)
 - Patients with germline BRCA mutations had a significantly longer PFS
 - Patients with somatic BRCA mutations had longer PFS than patients without BRCA mutations
 - Analyses of other mutations of interest were performed, but were not found to be statistically significant

- Secondary outcomes
 - Patients with germline BRCA2 mutations may receive more benefit from PARP inhibitors than other groups
 - Limitations: small sample size and patient selection
 - Patients treated with PARP inhibitors had an overall response rate of 27%
 - Prescribed doses of PARP inhibitors were well tolerated as monotherapy and in combination



PARP: poly(ADP-ribose) polymerase; BRCA: breast cancer susceptibility gene; PFS: progression-free survival

Conclusions

- In this cohort of 66 patients treated with PARP inhibitors, patients with germline BRCA mutations had a significantly longer PFS
- Patients with germline BRCA2 mutations had improved PFS over those with other BRCA mutations, though external validity for this data is limited
- PARP inhibitor therapy was generally well tolerated



PARP: poly(ADP-ribose) polymerase; PFS: progression-free survival

Future Directions

- An expanded review is planned which will include patients without known genetic mutations
- The expanded review will also include patients who received talazoparib or veliparib
- Further studies with larger cohorts are needed to assess other genetic mutations which may be associated with improved outcomes



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