

# Preliminary Results from TRITON2: A Phase 2 Study of Rucaparib in Patients with Metastatic Castration-Resistant Prostate Cancer (mCRPC) Associated with Homologous Recombination Repair (HRR) Gene Alterations

Wassim Abida,<sup>1</sup> Alan H. Bryce,<sup>2</sup> Nicholas J. Vogelzang,<sup>3</sup> Robert J. Amato,<sup>4</sup> Ivor Percent,<sup>5</sup> Jeremy Shapiro,<sup>6</sup> Ray McDermott,<sup>7</sup> Arif Hussain,<sup>8</sup> Akash Patnaik,<sup>9</sup> Daniel Petrylak,<sup>10</sup> Charles J. Ryan,<sup>11</sup> Thomas Stanton,<sup>12</sup> Jingsong Zhang,<sup>13</sup> Andrew D. Simmons,<sup>14</sup> Darrin Despain,<sup>14</sup> Melanie Collins,<sup>14</sup> Tony Golsorkhi,<sup>14</sup> Howard I. Scher,<sup>1</sup> Simon Chowdhury<sup>15</sup>

<sup>1</sup>Memorial Sloan Kettering Cancer Center, New York, NY, USA; <sup>2</sup>Mayo Clinic, Phoenix, AZ, USA; <sup>3</sup>Comprehensive Cancer Centers of Nevada, Las Vegas, NV, USA; <sup>4</sup>University of Texas Health Science Center, Houston, TX, USA; <sup>5</sup>Florida Cancer Specialists, Port Charlotte, FL, USA; <sup>6</sup>Cabrini Hospital, Malvern, VIC, Australia; <sup>7</sup>Adelaide and Meath Hospital (Incorporating the National Children's Hospital), Dublin, Ireland; <sup>8</sup>University of Maryland Greenebaum Cancer Center, Baltimore, MD, USA; <sup>9</sup>University of Chicago Comprehensive Cancer Center, Chicago, IL, USA; <sup>10</sup>Yale University, Yale Cancer Center, New Haven, CT, USA; <sup>11</sup>University of Minnesota, Minneapolis, MN, USA; <sup>12</sup>St. Joseph Health Cancer Center, Santa Rosa, CA, USA; <sup>13</sup>H. Lee Moffitt Cancer Center, Tampa, FL, USA; <sup>14</sup>Clovis Oncology, Inc., Boulder, CO, USA; <sup>15</sup>Guy's Hospital and Sarah Cannon Research Institute, London, UK

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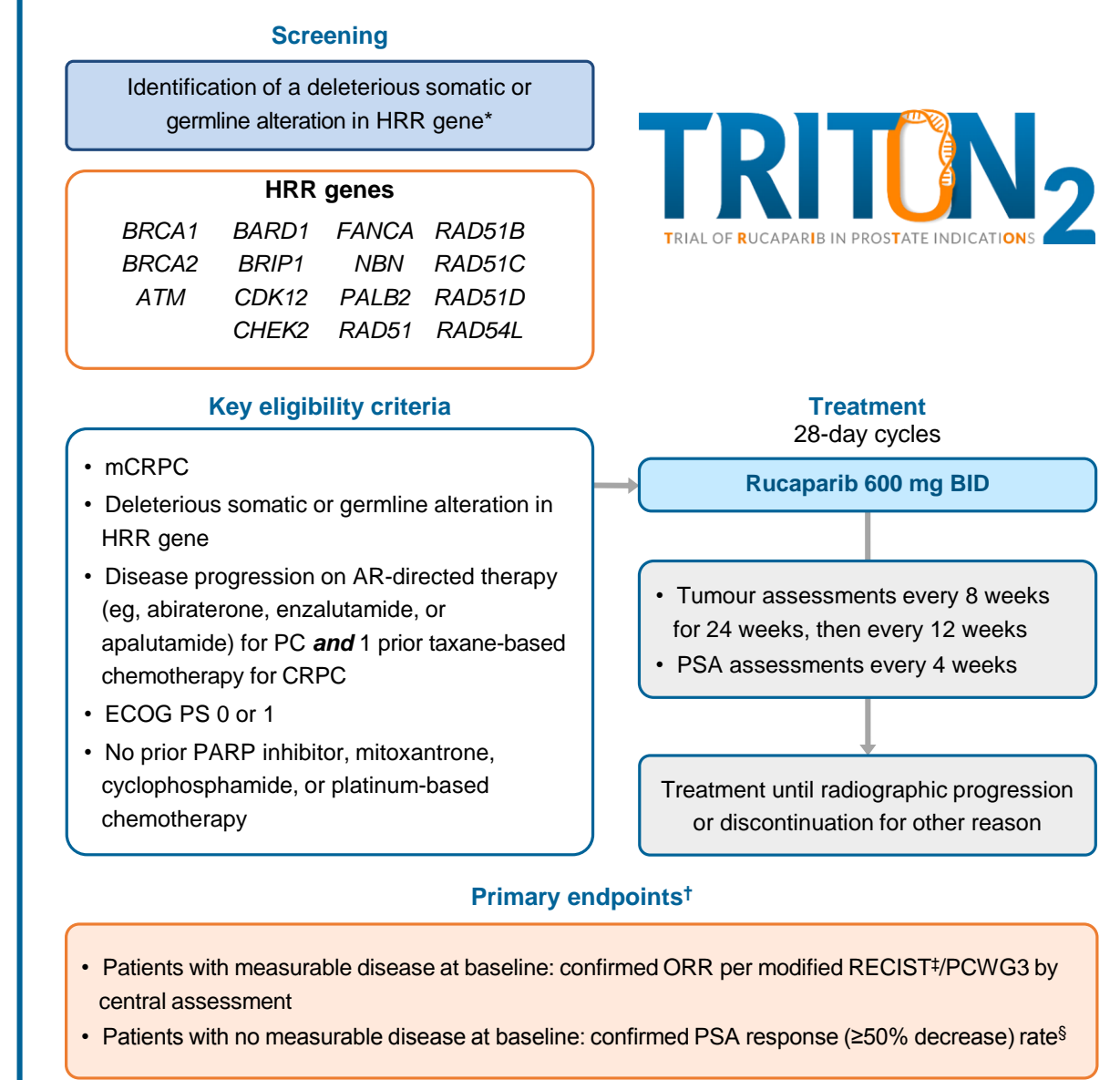
## INTRODUCTION

- There are limited treatment options available for patients with mCRPC following androgen receptor (AR)-directed therapy and taxane chemotherapy.<sup>1,2</sup>
- In metastatic prostate cancer, up to 25% of patients harbour a deleterious germline and/or somatic alteration in *BRCA1*, *BRCA2*, *ATM*, or other HRR genes.<sup>3-5</sup>
- In cells with HRR deficiency, inhibition of poly(ADP-ribose) polymerase (PARP) results in cell death via synthetic lethality.<sup>6-8</sup>
- Preclinical and limited clinical data have shown that PARP inhibitors have antitumour activity in HRR-deficient prostate cancer
  - In preclinical studies, rucaparib demonstrated activity in prostate cancer cell lines with reduced levels of *BRCA1*, *BRCA2*, or *ATM*<sup>9</sup>
  - In the phase 2 study TOPARP-A (NCT01682772), 16 of 49 (33%) evaluable patients with mCRPC progressing on ≥1 prior chemotherapy responded to olaparib treatment, using a composite endpoint of radiographic, prostate-specific antigen (PSA), or circulating tumour cell response; of the 16 responders, 14 had a tumour alteration in *BRCA1* (n=1), *BRCA2* (n=7), *ATM* (n=4), or other HRR gene (n=2)<sup>10</sup>
- Here we present interim data from TRITON2 (CO-338-052; EudraCT 2016-003162-13; NCT02952534), a phase 2, international, multicentre, open-label study evaluating rucaparib in patients with mCRPC associated with an HRR gene alteration

## METHODS

- Eligible patients were screened for the presence of a deleterious germline or somatic alteration in *BRCA1*, *BRCA2*, *ATM*, or other prespecified HRR gene<sup>11</sup> (Figure 1)
  - Deleterious alterations were defined as protein-truncating mutations, large protein-truncating rearrangements, splice site mutations, deleterious missense mutations, and homozygous deletions
- Enrolled patients received oral rucaparib 600 mg twice daily (BID)

Figure 1. TRITON2 Trial Schema



## RESULTS

- Enrolment is ongoing in TRITON2
- As of 16 April 2018, 85 patients were enrolled in TRITON2 (Figure 2; Table 1)
  - At the visit cutoff date (29 June 2018), median duration of follow-up was 5.7 months (range, 2.6–16.4 months)

Figure 2. Efficacy Populations

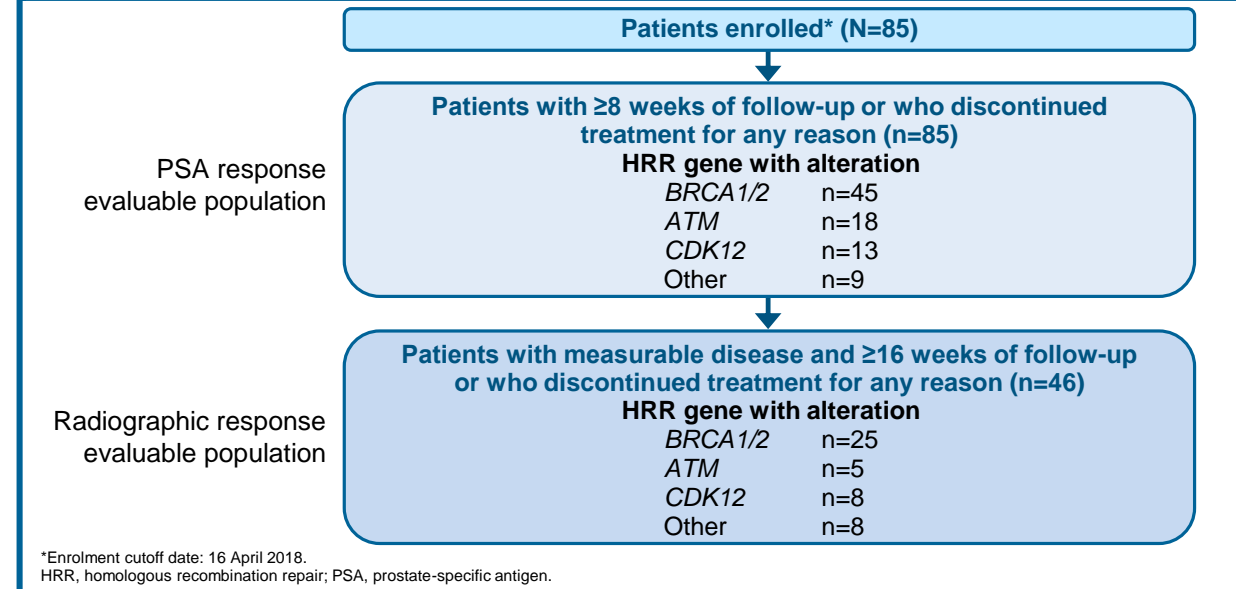


Table 1. Baseline Demographics

Characteristic	By HRR gene with alteration				Overall (N=85)
	<i>BRCA1/2</i> (n=45)	<i>ATM</i> (n=18)	<i>CDK12</i> (n=13)	Other <sup>a</sup> (n=9)	
Age, median (range), y	71 (50–88)	72.5 (62–84)	64 (49–79)	72 (60–86)	71 (49–88)
Race, n (%)					
White	35 (77.8%)	12 (66.7%)	6 (46.2%)	5 (55.6%)	58 (68.2%)
Black or African American	4 (8.9%)	3 (16.7%)	1 (7.7%)	1 (11.1%)	9 (10.6%)
Unknown	6 (13.3%)	3 (16.7%)	6 (46.2%)	3 (33.3%)	18 (21.2%)
ECOG PS, n (%)					
0	16 (35.6%)	10 (55.6%)	6 (46.2%)	1 (11.1%)	33 (38.8%)
1	28 (62.2%)	8 (44.4%)	7 (53.8%)	7 (77.8%)	50 (58.8%)
≥2	1 (2.2%)	0	0	1 (11.1%)	2 (2.4%)
Baseline PSA, median (range), ng/mL	52.0 (3.5–4782.0)	59.3 (9.2–4350.0)	57.7 (23.3–2966.5)	54.0 (8.8–798.8)	54.0 (3.5–4782.0)
Gleason score ≥8, n (%)	33 (73.3%)	6 (33.3%)	13 (100%)	6 (66.7%)	58 (68.2%)
No. prior CRPC therapies, median (range) <sup>b</sup>	2 (2–7) <sup>c</sup>	3 (2–6)	3 (2–4)	2 (2–4)	2 (2–7) <sup>c</sup>
Prior therapies, n (%) <sup>d</sup>					
Abiraterone	25 (55.6%)	16 (88.9%)	9 (69.2%)	8 (88.9%)	58 (68.2%)
Enzalutamide	33 (73.3%)	14 (77.8%)	12 (92.3%)	4 (44.4%)	63 (74.1%)
Abiraterone and enzalutamide	14 (31.1%)	12 (66.7%)	8 (61.5%)	3 (33.3%)	37 (43.5%)
Docetaxel	43 (95.6%)	17 (94.4%)	11 (84.6%)	8 (88.9%)	79 (92.9%)
Cabazitaxel	4 (8.9%)	4 (22.2%)	2 (15.4%)	1 (11.1%)	11 (12.9%)
Sipuleucel-T	6 (13.3%)	4 (22.2%)	1 (7.7%)	0	11 (12.9%)
Radium	5 (11.1%)	4 (22.2%)	0	1 (11.1%)	10 (11.8%)
Metastases, n (%) <sup>§</sup>					
Bone	40 (88.9%)	17 (94.4%)	10 (76.9%)	7 (77.8%)	74 (87.1%)
Nodal	28 (62.2%)	5 (27.8%)	11 (84.6%)	8 (88.9%)	52 (61.2%)
Visceral	19 (42.2%)	4 (22.2%)	4 (30.8%)	1 (11.1%)	28 (32.9%)
Hepatic	8 (17.8%)	2 (11.1%)	1 (7.7%)	1 (11.1%)	12 (14.1%)
Measurable disease at baseline per investigator, n (%)					
Yes	27 (60.0%)	5 (27.8%)	8 (61.5%)	8 (88.9%)	48 (56.5%)
No	18 (40.0%)	13 (72.2%)	5 (38.5%)	1 (11.1%)	37 (43.5%)
Gene alteration status, n (%)					
Germline	15 (33.3%)	5 (27.8%)	0	0	20 (23.5%)
Somatic	30 (66.7%)	10 (55.6%)	0	0	40 (47.1%)
Not available	0	3 (16.7%)	13 (100%)	9 (100%)	25 (29.4%)

<sup>a</sup>Includes patients with an alteration in *FANCA*, and 1 patient each with an alteration in *BRIP1*, *CHEK2*, *NBN*, *PALB2*, *RAD51*, *BRIP1*, *CHEK2*, or *CHEK2/CDK12*. <sup>b</sup>Does not include luteinizing hormone-releasing hormone analogues, first-generation antiandrogens, hormones, corticosteroids, bone-targeted agents, haematopoietic growth factors, or docetaxel given for hormone-sensitive disease. <sup>c</sup>Data unavailable for 1 patient. <sup>d</sup>Categories are not mutually exclusive. <sup>§</sup>CRPC, castration-resistant prostate cancer; ECOG PS, Eastern Cooperative Oncology Group Performance Status; HRR, homologous recombination repair; PSA, prostate-specific antigen.

## Radiographic Response in Patients with Measurable Disease

- Of patients with a *BRCA1/2* alteration and measurable disease at baseline, 44.0% (11/25) had a confirmed objective response by investigator assessment (Table 2; Figure 4)
- A confirmed objective response by investigator assessment was also observed in 1 patient with a *BRIP1* alteration and 1 patient with a *FANCA* alteration (Table 2)

Table 2. Confirmed Investigator-Assessed ORR in Evaluable Patients

Characteristic	By HRR gene with alteration			
	<i>BRCA1/2</i> (n=25)	<i>ATM</i> (n=5)	<i>CDK12</i> (n=8)	Other (n=8)
ORR, n (%) [95% CI] <sup>a</sup>	11 (44.0%) [24.4–65.1]	0 [0.0–52.2]	0 [0.0–36.9]	2 (25.0%) [3.2–65.1]
Complete response, n (%)	0	0	0	0
Partial response, n (%)	11 (44.0%)	0	0	2 (25.0%) <sup>b</sup>
Stable disease, n (%)	9 (36.0%)	4 (80.0%)	5 (62.5%)	5 (62.5%)
Progressive disease, n (%)	4 (16.0%)	1 (20.0%)	2 (25.0%)	1 (12.5%)
Not evaluable, n (%)	1 (4.0%)	0	1 (12.5%)	0

Visit cutoff date: 29 June 2018. Includes patients who had measurable disease at baseline per the investigator and ≥16 weeks of follow-up or who discontinued treatment. <sup>a</sup>Per modified RECIST/PCWG3 criteria. <sup>b</sup>One patient had a *BRIP1* alteration and 1 patient had a *FANCA* alteration. CI, confidence interval; HRR, homologous recombination repair; ORR, objective response rate; PCWG3, Prostate Cancer Clinical Trials Working Group 3; RECIST, Response Evaluation Criteria In Solid Tumors version 1.1.

Figure 3. Best Change from Baseline in Sum of Target Lesions (n=46)

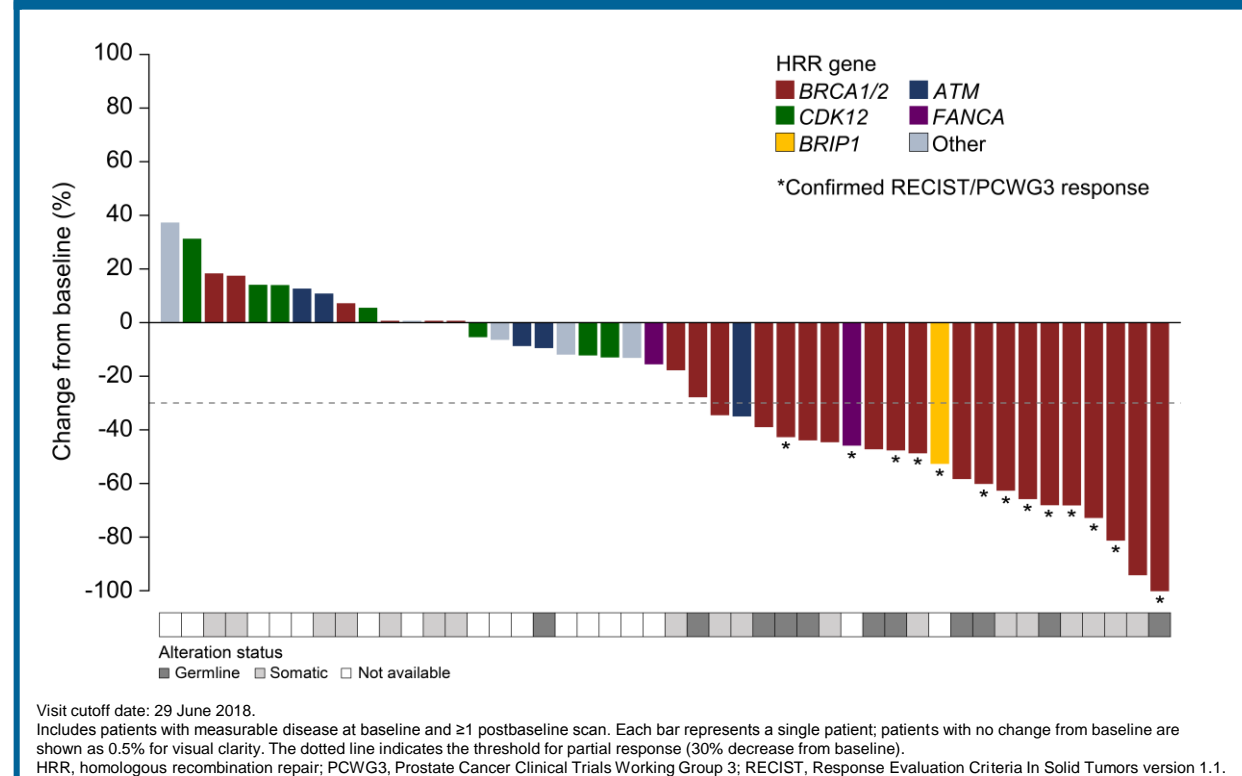
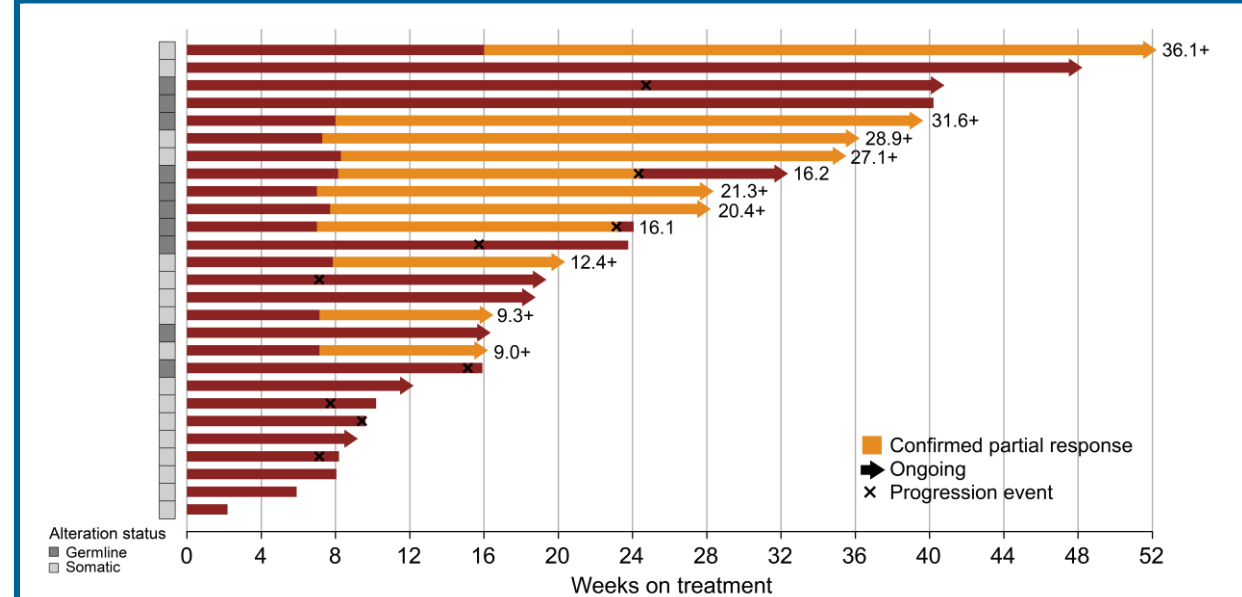


Figure 4. Treatment Duration and Duration of Radiographic Response in Patients with a *BRCA1/2* Alteration and Measurable Disease at Baseline (n=27)



## PSA Response

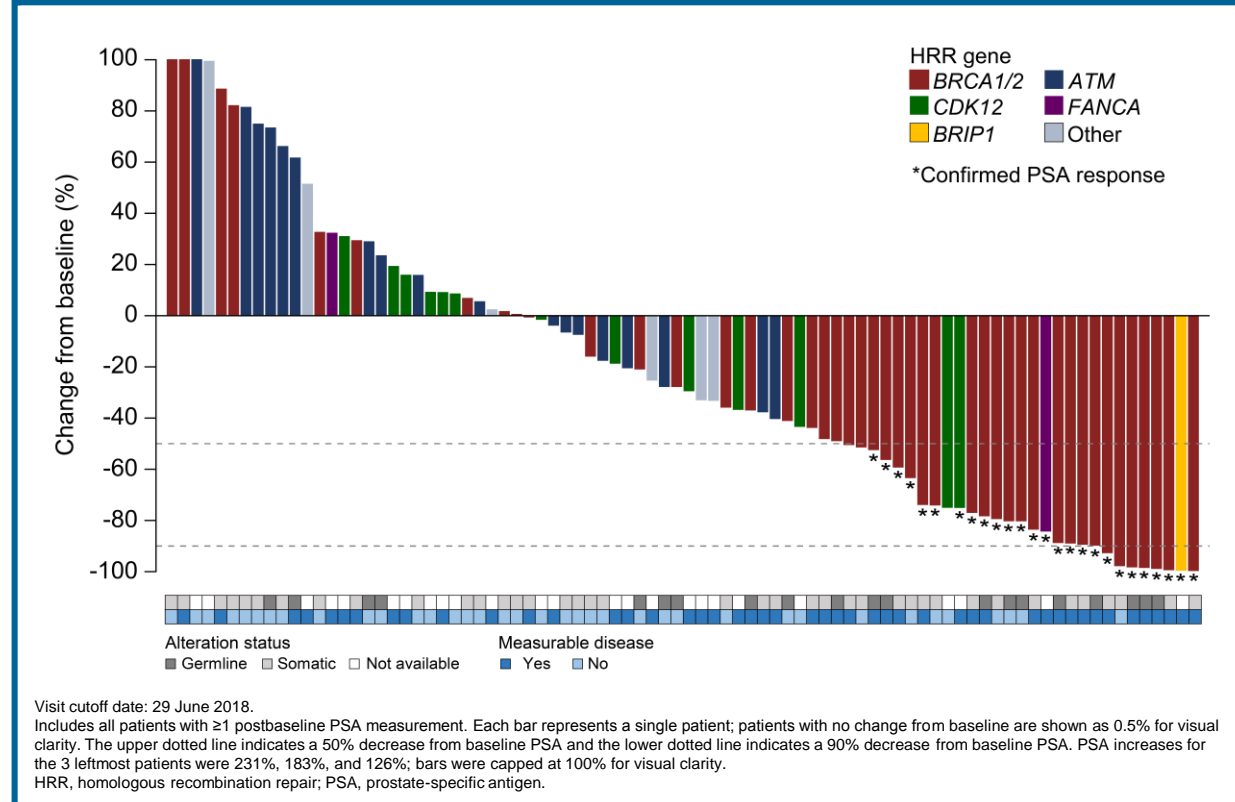
- Among patients with a *BRCA1/2* alteration, 51.1% (23/45) had a confirmed PSA response (Table 3)
- A confirmed PSA response was also observed in 1 patient with a *CDK12* alteration, 1 patient with a *BRIP1* alteration, and 1 patient with a *FANCA* alteration (Table 3)

Table 3. Confirmed PSA Response Rates

PSA response rate	By HRR gene with alteration, n/N (%) [95% CI]			
	<i>BRCA1/2</i>	<i>ATM</i>	<i>CDK12</i>	Other
All evaluable patients	23/45 (51.1%) [35.8–66.3]	0/18 (0%) [0.0–18.5]	1/13 (7.7%) <sup>a</sup> [0.2–36.0]	2/9 (22.2%) <sup>b</sup> [2.8–60.0]
With measurable disease	17/27 (63.0%) [42.4–80.6]	0/5 (0%) [0.0–52.2]	1/8 (12.5%) <sup>a</sup> [0.3–52.7]	2/8 (25.0%) <sup>b</sup> [3.2–65.1]
With no measurable disease	6/18 (33.3%) [13.3–59.0]	0/13 (0%) [0.0–24.7]	0/5 (0%) [0.0–52.2]	0/1 (0%) [0.0–97.5]

Visit cutoff date: 29 June 2018. Includes patients who had ≥8 weeks of follow-up or who discontinued treatment. <sup>a</sup>This patient did not demonstrate a confirmed objective radiographic response. <sup>b</sup>One patient with a *BRIP1* alteration and 1 patient with a *FANCA* alteration; both patients demonstrated a confirmed objective radiographic response. CI, confidence interval; HRR, homologous recombination repair; PSA, prostate-specific antigen.

Figure 5. Best Change from Baseline in PSA (n=84)



- Enrolment of patients with a *CDK12* alteration has been halted per protocol based on the lack of responses observed in these patients to date

## Safety

- In the safety population (defined as all patients who received ≥1 dose of rucaparib; N=85), median treatment duration in the overall population was 3.7 months (range, 0.5–12.9 months)
  - Median treatment duration in patients with a *BRCA1/2* alteration was 4.4 months (range, 0.5–12.0 months)

Table 4. Summary of TEAEs

	Overall safety population (N=85), n (%)
At least 1 TEAE	81 (95.3%)
At least 1 TEAE grade ≥3	45 (52.9%)
Treatment interruption and/or dose reduction due to TEAE	45 (52.9%)
Treatment interruption due to TEAE	41 (48.2%)
Dose reduction due to TEAE	25 (29.4%) <sup>a</sup>
TEAE leading to discontinuation	5 (5.9%) <sup>b</sup>
Death due to TEAE	1 (1.2%) <sup>c</sup>

Visit cutoff date: 29 June 2018. <sup>a</sup>The most common were asthenia/fatigue (8.2%), anaemia/decreased haemoglobin (7.1%), thrombocytopenia/decreased platelet count (5.9%), and nausea (4.7%). <sup>b</sup>One patient each (1.2%) discontinued due to anaemia/decreased haemoglobin, asthenia/fatigue, decreased appetite, and general physical health deterioration; 1 patient (1.2%) discontinued due to TEAEs of asthenia/fatigue, decreased appetite, and weight loss. <sup>c</sup>Due to disease progression. TEAE, treatment-emergent adverse event.

Table 5. Most Common (≥10%) TEAEs of Any Grade in All Patients Regardless of Causality

	Overall safety population (N=85)	
	Any grade, n (%)	Grade ≥3, n (%)
Asthenia/fatigue	38 (44.7%)	4 (4.7%)
Nausea	36 (42.4%)	3 (3.5%)
Anaemia/decreased haemoglobin	24 (28.2%)	13 (15.3%)
Decreased appetite	24 (28.2%)	3 (3.5%)
Constipation	19 (22.4%)	1 (1.2%)
ALT/AST increased	18 (21.2%)	4 (4.7%)
Vomiting	17 (20.0%)	0
Diarrhoea	16 (18.8%)	1 (1.2%)
Arthralgia	11 (12.9%)	1 (1.2%)
Dizziness	11 (12.9%)	0
Back pain	10 (11.8%)	2 (2.4%)
Oedema peripheral	10 (11.8%)	0
Weight decreased	10 (11.8%)	0
Dysgeusia	9 (10.6%)	0
Dyspnoea	9 (10.6%)	0
Haematuria	9 (10.6%)	3 (3.5%)

Visit cutoff date: 29 June 2018. ALT, alanine aminotransferase; AST, aspartate aminotransferase; TEAE, treatment-emergent adverse event.

## Pharmacokinetic (PK) Data

- Data for trough rucaparib plasma concentration were collected from 79 patients enrolled as of the 16 April 2018 enrolment cutoff date
- The plasma concentration of rucaparib in men enrolled in TRITON2 was comparable to that observed in women with ovarian cancer also treated with rucaparib 600 mg BID<sup>12</sup>

## CONCLUSIONS

- Rucaparib has encouraging antitumour activity in patients with a deleterious alteration in *BRCA1* or *BRCA2*
  - Among evaluable patients with a *BRCA1/2* alteration, 44.0% (11/25) had a confirmed radiographic response and 51.1% (23/45) had a confirmed PSA response
- Two patients with *BRIP1* or *FANCA* alterations also had confirmed radiographic and PSA responses
- Confirmed radiographic responses have not yet been observed in patients with *ATM* or *CDK12* gene alterations
  - Some reductions in target lesion diameters and PSA measurements have been observed in these patients
- Preliminary safety and PK data for rucaparib in men with mCRPC are consistent with those observed in patients with ovarian cancer and other solid tumours<sup>13-17</sup>
- Based on initial data from TRITON2, on 2 October 2018, the U.S. Food and Drug Administration granted Clovis Oncology Breakthrough Therapy designation for rucaparib as a monotherapy treatment of adult patients with *BRCA1/2*-mutated mCRPC who have received at least 1 prior AR-directed therapy and taxane-based chemotherapy<sup>18</sup>
- Enrolment in TRITON2 is ongoing, and the safety and efficacy of rucaparib as treatment for patients with mCRPC associated with an HRR gene alteration continues to be evaluated

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