Abstract 3115: Talazoparib in Patients with Solid Tumors with BRCA1/2 Mutations: **Results from the Targeted Agent and Profiling Utilization Registry (TAPUR) Study**

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Background:

- TAPUR is a phase II basket study that evaluates antitumor activity of commercially available targeted agents in patients (pts) with advanced cancers with specific genomic alterations.
- Results of a cohort of pts with various solid tumors with BRCA1/2 mutations (mut) treated with talazoparib (Tala) are reported.

Methods:

Study Design:

- Eligible pts: Advanced solid tumors, ECOG performance status (PS) 0-2, adequate organ function, measurable disease and no standard treatment (tx) options. Tx was assigned according to pre-specified protocol matching rules based on genomic testing performed in CLIA-certified, CAP-accredited labs selected by clinical sites.
- Pts received 1 mg of Tala orally daily until disease progression, unacceptable toxicity or pt or physician choice to discontinue.
- **Primary endpoint:** Disease control (DC) determined by investigator assessment of objective response (OR) or stable disease (SD) of at least 16 weeks (wks) duration (SD16+) per RECIST v1.1. Confirmation of response was not required.
- Secondary endpoints: Progression-free survival (PFS), overall survival (OS), duration of response, duration of SD, and toxicity per CTCAE v4.0. Grade 3-5 adverse events (AEs) or serious adverse events (SAEs) at least possibly related to tx are reported.

Statistical Methods:

 Inferences are based on a one-sided exact binomial test with a null hypothesis of DC rate ≤15%; power and alpha were 82% and 0.10, respectively. Two-sided 95% Cls are used for other efficacy endpoint estimates.

Results:

- 28 pts enrolled from December 2019 to September 2021. All pts were evaluable and included in efficacy analysis.
- **Demographics**: Median age 66 (range, 32-80); 50% female; 93% self-identified as White, 4% as Black/African American, 4% as prefer not to answer; 93% as not Hispanic or Latino, and 7% preferred not to answer.
- **Clinical characteristics:** 96% PS 0-1, 4% PS 2; 61% received ≥3 prior systemic regimens. Primary tumor type (# pt): NSCLC (5), breast (3), pancreas (2), uterus (2), anus (1), biliary tract (1), cervix (1), cholangiocarcinoma (1), esophagus (1), GE junction (1), HCC (1), leiomyosarcoma (1), melanoma (1), mesothelioma/peritoneal (1), non-melanoma skin (1), ovary (1), prostate (1), SCLC (1), stomach (1), vagina
- Outcomes: 16 pts had CR (1), PR (9), or SD16+ (6) for a DC rate of 57% (90% CI: 43 to 100) (Tables 1 and 2). The null DC rate of 15% was rejected (p<0.001).
- Safety: 13 pts (46%) had ≥1 SAE or grade 3 AE at least possibly related to Tala. All were consistent with the drug label except bilirubin increase and hyponatremia (both grade 3 AEs).

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<u>Conclusion</u>: Talazoparib demonstrated antitumor activity in patients with advanced solid tumors and BRCA1/2 mutations.

<u>Future Direction</u>: Additional study is warranted to confirm the efficacy of talazoparib in pts with non-breast/ovarian solid tumors with BRCA1/2 mutations.

| Table 1. Tumor Type and Mut of Pts Meeting Response Criteria (n=16) | | | | |
|---|------------------------------|--------------|--|--|
| Response | Primary Tumor | Mut | Comutations ^a | |
| CR | Non-melanoma skin* | BRCA2 | ATM A188T ^b | |
| P R | Ovary | BRCA2 | | |
| P R | Pancreas | BRCA1, BRCA2 | MLH1 Gin391Arg ^b | |
| ۲R | Breast | BRCA1 | | |
| Ϋ́R | HCC* | BRCA1 | ATM K2383I ^b ; MRE11A R572* and I548T ^b | |
| PR | Mesothelioma, peritoneal* | BRCA2 | | |
| PR | Stomach* | BRCA2 | <i>CHEK1</i> T476I ^b ; <i>CHEK2</i> R474H ^b ; <i>PALB2</i> E426K ^b | |
| PR | Uterus* | BRCA2 | | |
| PR | Breast | BRCA2 | | |
| PR | NSCLC* | BRCA2 | ARID1A P570T ^b | |
| 5D16+ | Prostate | BRCA2 | | |
| 5D16+ | Cervix* | BRCA2 | | |
| 5D16+ | NSCLC* | BRCA1 | <i>CHEK2</i> R117G | |
| 5D16+ | Uterus* | BRCA2 | <i>ARID1A</i> Q1519fs and Q2037fs; <i>ATM</i> R2580S ^b | |
| 5D16+ | Esophagus* | BRCA1 | ATR V66M ^b | |
| 5D16+ | Leiomyosarcoma* | BRCA2 | ATRX D1525fs; NBN G136V ^b | |

^a Of the following genes examined: ARID1A, ATM, ATR, ATRX, BARD1, BRIP1, CHEK1, CHEK2, FANCA, FANCC, FANCD2, FANCE, FANCF, FANCG, FANCL, MLH1, MRE11A, NBN, PALB2, RAD51, RAD51B, RAD51D, RAD54L. ^b Variant of unknown significance. *No PARP inhibitor is currently FDA approved for this tumor type

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| ble 2: Efficacy Outcomes (N=28) | |
|---|---------------------------------|
| Crate, % (90% CI) | 57 (43, 100); p<0.001 |
| R rate, % (95% CI) | 36 (19, 56) |
| edian PFS, wks (95% CI) | 24 (8, 39) |
| edian OS, wks (95% CI) | 71 (32, inf) |
| uration of CR, wks (n=1) | 93 |
| edian duration of PR (range), wks (n=9) | 20 (11, 80) |
| edian duration of SD in pts with SD16+, wks (n=6) | 36 (19, 108) |

Figure 1: Best Percent Change from Baseline in Target Lesion Size (N=28)

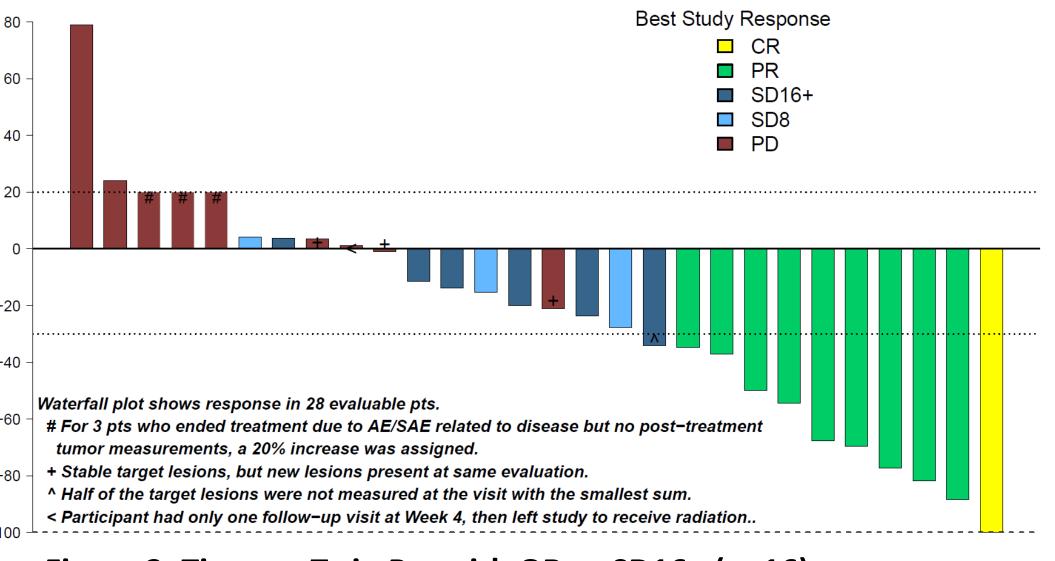


Figure 2: Time on Tx in Pts with OR or SD16+ (n=16)

