

Real-world (TW) Genomic Landscape, PARP-inhibitor (PARPi) Treatment Patterns and Adverse Events (AE) in Advanced Prostate Cancer: The National CanaDNA Study

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Background

- The genomic landscape of advanced prostate cancer continues to be defined.
- The landscape of biomarker-driven treatment approaches in advanced prostate cancer is rapidly evolving.
- Thus, the national genomic profiling, targeted treatment patterns and their associated adverse events were examined and analyzed in this real-world study.

Methods

- Western Canadian (Alberta, British Columbia) patients with advanced prostate cancer (metastatic castration-sensitive [mCSPC] or castration-resistant prostate cancer [mCRPC]) who underwent comprehensive genomic profiling were included.
- For Alberta patients, cell-free concordance analysis was conducted (130-gene custom in-house NGS panel vs. FoundationOne Liquid CDx).
- Clinical characteristics, treatment patterns, and outcomes were derived from the POET Clinical Database.
- Clinical and genomic data were used to examine:
 - * Impact of presence (+) or absence (0) of HRR alterations on outcomes (e.g. time to castration resistance).
 - * Treatment patterns, use of PARPi therapy, and response to therapy including adverse events, progression-free survival (PFS), and overall survival (OS), in patients with HRR+ disease of patients in Alberta, Canada.



This real-world analysis offers valuable insights into the genomic landscape PARPi-treatment outcomes for advanced prostate cancer in Western Canada.

- 1 in 4 RW patients had an HRR+ detected on two testing panels (100% concordance for clinical actionability).
 - In RW patients, PARPi use achieved survival outcomes comparable to those of HRR- patients, despite often poorer prognosis.
 - In real-world HRR+ patients, PARPi were used infrequently and later in the treatment course, yielding poorer clinical outcomes (response to therapy; duration of disease control) compared to clinical trial findings, despite similar duration of PARPi use.
- Equitable standard-of-care genomic testing in advanced prostate cancer is necessary to enable earlier PARPi access and improve outcomes.**

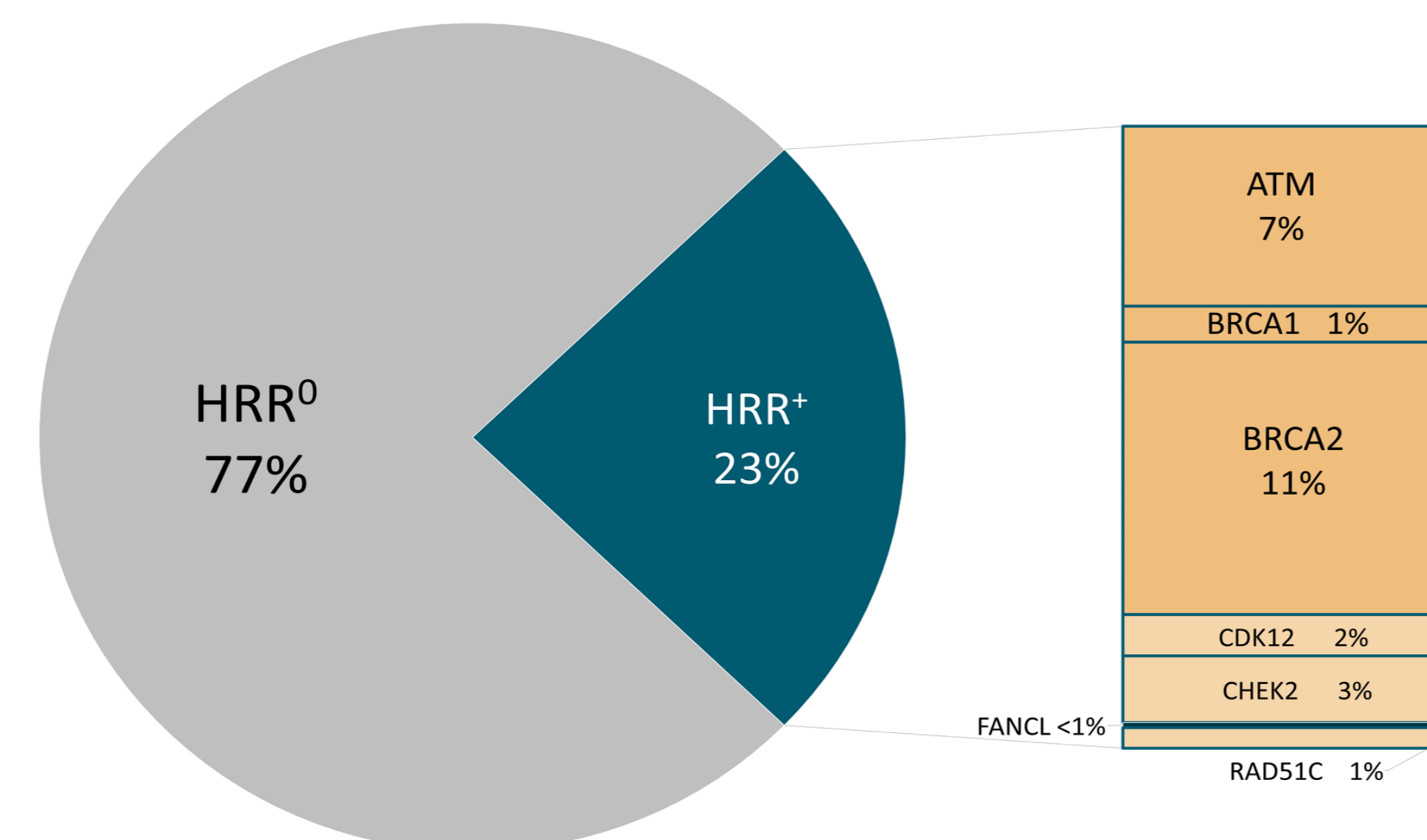
Results

A cohort of 486 (Alberta n=236; British Columbia: n=250) were identified.

Median age was 68 years, 62% had a Gleason Score ≥ 8 .
41% had high volume disease (15% visceral involvement).
58% had de novo mCSPC. 57% developed mCRPC a median 28.5 months post-androgen therapy initiation.

Genomic Landscape and Impact on Outcomes

HRR alterations were identified in 23% of the Western Canadian cohort



High concordance was observed between an in-house NGS panel and FoundationOne Liquid CDx

98% Analytical Concordance
100% Clinical Actionability

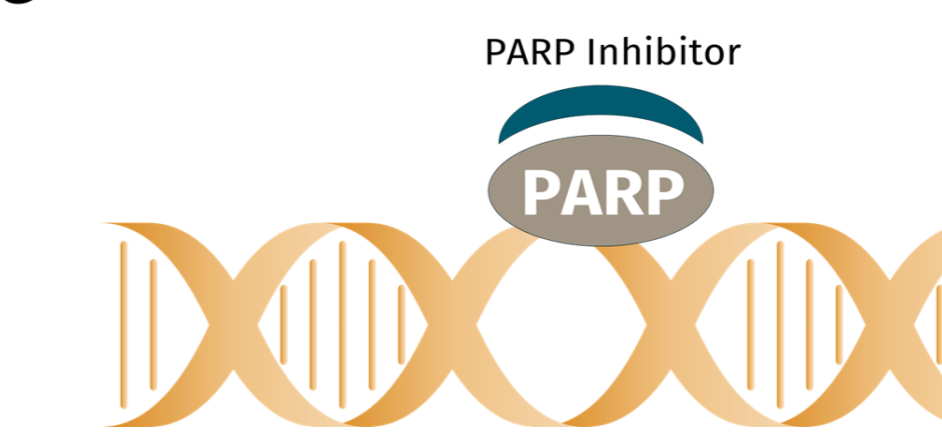
Patients with HRR alterations progressed to mCRPC significantly faster than those without (log rank p < 0.001; HR 1.52 [95% CI: 1.2 – 2.0])



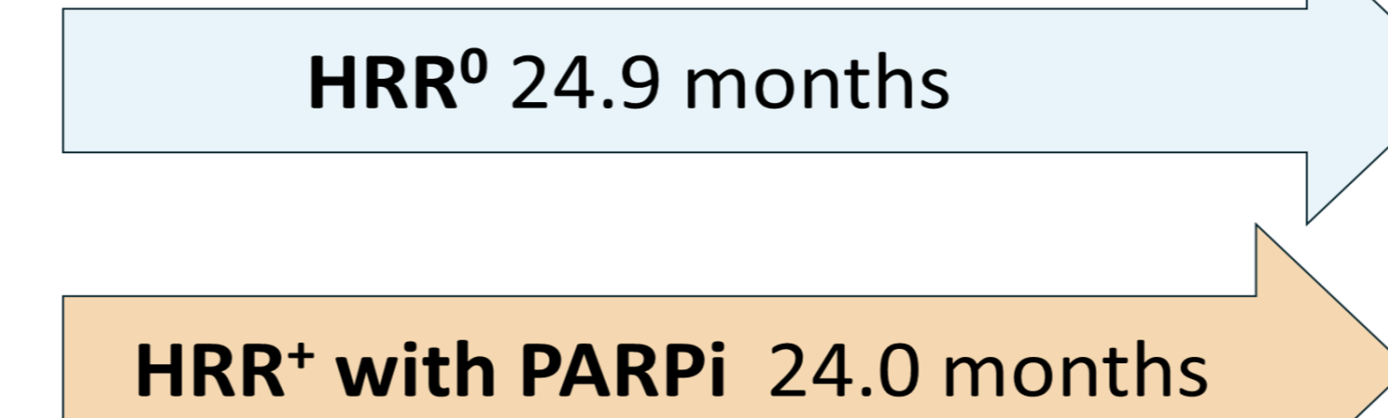
PARPi Therapy Outcomes

PARPi was used in 46% (51/111) of HRR+ patients in the 2nd (28% 20/51), 3rd 23% (12/51), 4th 23% (12/51) or 5th (13% 7/51) palliative intent systemic therapy line.

47% (24/51) received a taxane-based cytotoxic chemotherapy prior to PARPi.



Median survival post-mCRPC was similar between HRR⁰ patients and HRR⁺ patients receiving PARPi



Log-rank p = 0.4; HR: 1.2; [95% CI: 0.8 – 1.7]

Among HRR+ patients without PARPi, 46% (11/24) received ≥ 1 line of cytotoxic chemotherapy.



PARPi Received for HRR+ Disease (n=51)	n (%) or months [95% CI]
ECOG at PARPi-initiation	
≤ 1	38 (75)
≥ 2	13 (25)
PSA decline $\geq 50\%$	13 (25)
Objective Response Rate	3 (6)
Disease Control Rate	24 (47)
Progression-Free Survival	
Biochemical	4.0 [2.7 – 6.3]
Radiographic	5.4 [3.3 – 7.7]
Duration of Tx	4.8
PARPi ≥ 30 days post radiological progression	29 (58)
PARPi-related AE	24 (50)
Tx modification required	17/24 (71)
Tx termination required	7/24 (29)
Management(s) post-PARPi	23 (45)
Taxane-based CTx	20/23 (87)
Platinum-based CTx	3/23 (13)

Median survival post-PARPi initiation: 7.0 months [3.7 – 9.1]

Future Directions

The CanaDNA Study is ongoing and aims to enroll 1000 patients to build a pan-Canadian database characterizing the genomic landscape of advanced prostate cancer.

In addition to genomic profiles, the CanaDNA study's collection of clinical management and outcomes will help establish a knowledge base on advanced prostate cancer care prior to the widespread adoption of standard-of-care genomic testing in Canada.

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