Constellation PHARMACEUTICALS

ProSTAR: A phase 1b/2 study of CPI-1205, a small molecule inhibitor of EZH2, combined with enzalutamide (E) or abiraterone/prednisone (A/P) in patients with metastatic castration-resistant prostate cancer (mCRPC)

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BACKGROUND

- Enhancer of zeste homolog 2 (EZH2) is the catalytic subunit of Polycomb Repressive Complex 2 (PRC2) the PRC2 complex and methylates histone H3 on lysine 27, resulting in suppression of gene expression.
- EZH2 mutations and increased expression are often observed in cancer, leading to repression of genes associated with apoptosis and differentiation.^{1,2}
- Overexpression of EZH2 in metastatic castration-resistant prostate cancer (mCRPC)³ and decreased expression of target genes are correlated with disease stage and poor prognosis.⁴
- Many mCRPC tumors remain dependent on androgen receptor signaling (ARS), but eventually develop resistance to ARS inhibitors, such as E and A/P.
- Preclinical ARS-dependent prostate cancer cell models are sensitive to EZH2 inhibition.5-8
- Combining EZH2 inhibitors with ARS inhibitors results in synergistic cell growth inhibition.5-8
- CPI-1205 is a potent, selective, and cofactor-competitive EZH2 inhibitor.⁹

In this multicenter phase 1b/2 study, patients with mCRPC previously

treated with Enzalutamide (E) or Abiraterone (A/P) are enrolled in

different cohorts exploring two regimens of oral CPI-1205 on a

continuous 28-day cycle: 1) 800 mg TID or 2) 400 mg BID with cobicistat

(C), a CYP3A4 inhibitor, combined with the standard dose of E (160 mg

QD) or standard dose of A/P (1000 mg QD/5 mg BID). C is used to

increase systemic exposure of CYP3A substrates. Pre-treatment with C

successfully increased the exposure of CPI-1205 in another study, which

led to the interest to explore the addition of C to the treatment

An expansion cohort in heavily pretreated patients will be initiated with

the dose regimen that was deemed safe by the study safety committee.

After consideration of safety, PK, and PD, the RP2D will be established,

and phase 2 will be initiated. One or both of the combinations, CPI-1205

with E and with A/P, might proceed to phase 2. If both combinations are

chosen for phase 2, one combination might proceed to a randomized

phase 2, and the second combination might proceed to another

randomized phase 2 or a single-arm phase 2. NCT03480646.

- CPI-1205 monotherapy achieved objective responses and substantial target engagement at well tolerated doses in a phase 1 clinical trial in patients with non-Hodgkin's lymphoma.¹
- These observations support the clinical evaluation of CPI-1205 with standard doses of E or A/P in mCRPC. Data presented here are based on the phase 1b portion of the study with data cut of 6 Feb 2019.

STUDY DESIGN

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PHASE 1B STUDY OBJECTIVES

REFERENCES

Primary Objective:

• Determine the maximum tolerated dose (MTD) and recommended phase 2 dose (RP2D) of CPI-1205 + E and CPI-1205 + A/P in patients with mCRPC

Secondary Objectives:

combination in this study.

- Characterize the safety, tolerability, pharmacokinetic (PK) profile
- Determine preliminary signs of efficacy with each combination
- Prostate specific antigen (PSA) 50% response, circulating tumor cell (CTC) 30% response, or objective response per Prostate Cancer Working Group 3 (PCWG3) **Exploratory Objectives:**

Evaluate the pharmacodynamic effects and PK/PD relationship of CPI-1205

• Evaluate and identify potential novel predictive biomarkers utilizing genomics, transcriptomics, and protein expression technologies

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BASELINE CHARACTERISTICS Demographics Enzalutamide Abiraterone (N=20)(N=16)Age (Years) Min-Max 55-90 ECOG PS [n, (%)] 9 (56.3%) 10 (50.0%) 10 (50.0%) 7 (43.8%) PSA (ng/mL CTC Count Unfavorable [n, (%)] 2 (60.0%) ARV7+ [n, (%)] 8 (40.0%) Prior Chemotherapy [n, (%)] 5 (31.3%) Taxane-based for mCRPC [n, (%)] 2 (12.5%) 8 (40.0%) Target Lesion [n, (%)] 324.2 (140-846) 333.4 (131-758) Mean (Min-Max) Baseline abnormal [n, (%)] 7 (43.8%) 11 (55.0%) ARV7 (Androgen Receptor Splice Variant 7): protein and/or transcript levels are tested

TREATMENT DURATION



9/36 patients are still ongoing • 2 patients on treatment with CPI-1205 + E • 7 patients on treatment with CPI-1205 + A



Study-drug-related TEAEs: Reported as related to CPI-1205.



CPI-1205 + A

[*: transcript or protein]

RECIST 1.1 TARGET LESION PERCENT CHANGE (n=13)



6/13 (46%) experienced reduction in tumor burden.

- Among 13 patients with measurable disease. 9 (69%) patients had PR/SD as best response per RECIST 1.1. • 7/13 (54%) patients had DCR \geq 3 months.
- SDDCRDCR(any duration)(any duration)(≥ 3 mont CPI-1205 + E 1/5 (20%) 2/5 (40%) 3/5 (60%) 3/5 (60%) CPI-1205 + A/P 0/8 (0%) 6/8 (75%) 6/8 (75%) 4/8 (50%)

DCR: Disease Control Rate includes patients with best response of SD or PR

PR: Partial Response; SD: Stable Disease

*By central review.

PSA ≥80 AND RECIST 1.1 RESPONSE BY ARV7 STATUS

	ARV7 Positive (n=13)				ARV7 Negative (n=23)			
	PSA ≥80 Response	RECIST 1.1 Response**			PSA ≥ 80	RECIST 1.1 Response**		
		CR	PR	SD	Response	CR	PR	SD
CPI -1205 + E	0/5	0	0/2	1/2 (50%)	3/11 (27%)	0	1/3 (34%)	1/3 (34%)
CPI -1205 + A/P*	0/8	0	0/0	0/0	2/10 (20%)	0	0/8	6/8 (75%)

² patients with ARV7 negative status are not evaluable for PSA. **11 ARV7 negative and 2 ARV7 positive patients had measurable disease at baseline

SUMMARY OF ctDNA MUTATION ANALYSIS

- Genomic aberrations were detected from cell free DNA (cfDNA). Germline variants and suspected clonal hematopoiesis associated variants were not filtered from presented dataset. Genes with previous published frequency¹¹ of >5% c detected in >2 samples in current study are presented
- Current study shows variation in frequency of aberrations identified compared to published reports, including under-representation tion of AR amplification and ETS fusions and over-representation of AR mutations TP53 mutations, and RB1 copy number loss (poor prognostic indicators).
- 19% (5/27) patients had inactivating alterations in 2, or all 3, of the PTEN, RB1, or TP53 genes (commonly observed of aggressive and/or neuroendocrine disease).
- Additional analyses of tumor biopsi ongoing



quencies calculated from mCRPC cohort with prior E or A treatment. Grayed out boxes indicate gene was not included in genomics panel

CONCLUSIONS

- A significant number of patients had multiple indicators of poor prognosis at baseline (56%) unfavorable CTC enumeration, 50% high LDH, and 36% ARV7 expression).
- The combination of CPI-1205 800 mg TID or 400 mg BID with C administered with E or A/P were generally well tolerated.
- Co-administration with C did not completely reverse the reduction of CPI-1205 exposure in patients who received CPI-1205 in combination with E.
- The CPI-1205 TID dosing regimen was sufficient to elicit a substantial PD response with similar results observed when combined with either E or A/P.
- CPI-1205 800 mg TID in combination with standard doses of E or A/P was selected as the RP2D.
- Several patients in both arms demonstrated PSA responses, RECIST responses, and/or CTC reductions in this advanced mCRPC second-line setting.
- Based on these encouraging results, we are conducting a phase 2 single-arm trial with CPI-1205 given in combination with A/P and a randomized trial of CPI-1205 given in combination with E in second-line mCRPC. An expansion cohort in heavily pre-treated patients has also been initiated.

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