# Weill Cornell Medicine 🗧 NewYork-Presbyterian

# TITLE: Phase I/II dose-escalation trial of combination fractionateddose <sup>177</sup>Lu-J591 and <sup>177</sup>Lu-PSMA-617 in patients with metastatic castration-resistant prostate cancer

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#### **Confidentiality Statement**

This document is confidential and is to be distributed for review only to investigators, potential investigators, consultants, study staff, and applicable independent ethics committees or institutional review boards. The contents of this document shall not be disclosed to others without written authorization from WCM.

# List of Abbreviations

ADT	Androgen Deprivation Therapy
AE	Adverse Event
AR	Androgen Receptor
CFR	Code of Federal Regulations
CRF	Case Report Form
CRPC	Castration Resistant Prostate Cancer
CTCAE	Common Terminology Criteria for Adverse Events
СТЕР	Cancer Therapy Evaluation Program
CTSC	Clinical Translational Science Center
DLT	Dose Limiting Toxicity
DSMB	Data and Safety Monitoring Board
DSMP	Data Safety Monitoring Plan
FDA	Food and Drug Administration
<sup>68</sup> Ga	Gallium–68
GCP	Good Clinical Practice
HIPAA	Health Insurance Portability and Accountability Act of 1996
HRBAF	Human Research Billing Analysis Form
ICF	Informed Consent Form
IND	Investigational New Drug
IRB	Institutional Review Board
<sup>177</sup> Lu	Lutetium-177
mCRPC	Metastatic Castration Resistant Prostate Cancer
MTD	Maximum tolerated dose
NA	Not Applicable
NCI	National Cancer Institute
PC	Prostate Cancer
PHI	Protected Health Information
PI	Principal Investigator
PSMA	Prostate Specific Membrane Antigen
<sup>223</sup> Ra	Radium-223
REDCap	Research Electronic Data Capture
RP2D	Recommended Phase 2 dose
SAE	Serious Adverse Event
SUSAR	Suspected Unexpected Serious Adverse Reaction
UAP	Unanticipated Problem
WCM	Weill Cornell Medicine

# **Protocol Summary**

Full Title:	Phase I/II dose-escalation trial of combination fractionated dose $^{177}$ Lu–J591 and $^{177}$ Lu–PSMA–617 in patients with metastatic
Short Title:	castration-resistant prostate cancer <sup>177</sup> Lu-J591 and <sup>177</sup> Lu-PSMA-617 combination for mCRPC
<b>Clinical Phase:</b>	Phase I
Principal Investigator:	Scott T. Tagawa, MD, MS
Sample Size:	N = 6 - 57 (Phase I: $3 + 3$ dose–escalation study design up to 6
	dose–escalation / de-escalation cohorts for minimum of 6 and maximum of 30; Phase II minimum of 16 and maximum of 27)
Accrual Ceiling:	This study will enroll up to 57 subjects who receive treatment.
Study Population:	Adult male patients of $\geq 18$ years with documented progressive metastatic CRPC
Accrual Period:	3 years (approximately $1-2$ patients per month with gaps between
	dose-escalation cohorts and between stage 1 and 2 of the phase II
	portion).
Study Design:	Phase I dose escalation study with combination of <sup>177</sup> Lu–J591 and
wind 2 wight	$^{177}Lu-PSMA-617$ using a dose-fractionated regimen will be
	nerformed in nations with documented progressive metastatic CRPC
	The cumulative $^{177}$ Lu–1591 dose for each subject will be 2.7 GBa/m <sup>2</sup>
	$(73 \text{ mCi/m}^2)$ of $^{177}\text{Lu}$ with 20 mg [591 and the cumulative
	$\frac{177}{10}$ u=PSMA=617 dose for each subject will vary (depending on the
	Cohort) from 3.7 GBq (100 mCi) to 18.5 GBq (500 mCi). The
	<sup>177</sup> Lu-PSMA-617 dose will be escalated in up to 6 different dose
	levels (3+3 dose–escalation study/de-escalation design). For the phase
	II portion, a minimum number of 14 patients will be enrolled at MTD
	(including those enrolled at MTD in Phase I) and a maximum of 24.
Study Duration:	Approximately 3 months after enrollment then transition to long-term
,	follow up
Study Agent:	1) $^{177}$ Lu-PSMA-617 [1 85 GBa (50 mCi) – 9 25 GBa (250 mCi)] x2
	doses 2 weeks apart (Treatment Visit #1 and #2) IV administration
	2) $^{177}$ Lu–J <b>591</b> [1 35 GBa/m <sup>2</sup> or 36 5 mCi/m <sup>2</sup> ] x <sup>2</sup> doses 2 weeks apart
	(Treatment Visit #1 and #2) IV administration
	3) $^{68}C_9 - PSMA - HBFD - CC [185 + 74 MBg or 5 + 2 mCi] intravenous$
	during screening and at 12 weeks $(+1 \text{ week})$ with standard imaging
Primary Objectives.	• (Phase I) Determine the dose limiting tovicity (DLT) of
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	$^{177}$ Lu-PSMA-617

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# Schema

**Screening:** Written informed consent, history, physical examination, conventional imaging modalities (CT/MRI, Bone scan), PSMA–ligand based PET/CT imaging



# Protocol #1802018988 v2 Version Date: 02/19/2019

# **Table of Contents**

1. ST	TUDY OBJECTIVES	
1.1	Primary ObjectivesErro	r! Bookmark not defined.
1.2	Secondary Objectives	
1.3	Exploratory Objectives	
2 B/	ACKGROUND	7
21	Disease	7
2.2	PSMA	
2.3	Anti-PSMA Monoclonal humanized anitbody (J591)	
2.4	Risk/Benefit Assessment	
2.5	Radionuclinde-chelating PSMA ligand	
2.6	Investigational Agent	
2.7	Rationale	
2.8	Rationale for the appropriate radioisotope	
2.9	Risk/Benefit Assessment	
2.4	Risk/Benefit Assessment	
2.10	Correlative Studies Background	
3. SU	JBJECT SELECTION	
3.1	Study Population	
3.2	Inclusion Criteria	
3.3	Exclusion Criteria	
4. O	VERVIEW OF STUDY DESIGN AND METHODOLOGY	23
4.1	Study Design	
4.2	Rationale for Dose-Escalation Strategy	
5 RI	EGISTRATION PROCEDURES	17
5. Ki	Identification of subjects	
5.1	Consent process	
5.3	Central Patient Registration	
6 97		47
0. 51	Schedule of Evaluations	
0.1 6.2	Treatment Administration	
6.2	General Concomitant Medication and Supportive Care Guide	
6.3	Duration of Therapy and Criteria for Removal from Study	35
6.5	Duration of Follow Up	
7 D		MADZ NOT DEFINED
7. D	USING DELA I 5/DUSE MUDIFICATIONS EKKOR! BOOK	MAKK NUT DEFINED.
8. PH	IARMACEUTICAL INFORMATION	
8.1	Investigational Agent	

8.2	Availability	Error! Bookmark not defined.
8.3	Agent Ordering	
8.4	Agent Accountability	
9. CC	DRRELATIVE/SPECIAL STUDIES	
9.1	Laboratory Correlative Studies	
9.2	Special Studies	
10. MI	EASUREMENT OF EFFECT	
10.1	CTC count response	
10.2	Biochemical (PSA) response	
10.3	Duration of PSA response	
10.4	Duration of CTC response	
10.5	PSA Progression	
10.6	PSA Stabilization	
10.7	Time to PSA Progression	
10.8	Change in lesion size	
10.9	PSMA Expression on CTCs	
9.2	Special Studies	
11. DA	ATA REPORTING / REGULATORY CONSIDERAT	TIONS 41
11.1	Data Collection	
11.2	Regulatory Considerations	
12. ST	ATISTICAL CONSIDERATIONS	
12.1	Study Design/Endpoints	
12.2	Sample Size/Accrual Rate	
12.3	Stratification Factors	
12.4	Analysis of Endpoints	
12.5	Interim Analysis	
12.6	Reporting and Exclusions	
REFER	ENCES	
APPEN	DIX	

### **1. STUDY OBJECTIVES**

The objectives of this clinical trial are as follows:

#### 1.1 **Primary Objectives**

- (Phase I) Determine the dose limiting toxicity (DLT) of the combination of <sup>177</sup>Lu–J591 and <sup>177</sup>Lu–PSMA–617 in a 2–week dose–fractionation regimen
- (Phase I) Determine the cumulative maximum tolerated dose (MTD) and/or recommended phase II dose (RP2D) of the combination of <sup>177</sup>Lu–J591 and <sup>177</sup>Lu–PSMA–617 in a 2–week dose–fractionation regimen
- (Phase II) To assess the proportion with PSA decline following treatment with the combination of <sup>177</sup>Lu–J591 and <sup>177</sup>Lu–PSMA–617 in a 2-week dose-fractionation regimen

### 1.2 Secondary Objectives

- To assess radiographic response rate by RECIST 1.1 with PCWG3 modifications
- To assess biochemical and radiographic progression-free survival by PCWG3 criteria
- To assess overall survival following treatment with the combination of <sup>177</sup>Lu–J591 and <sup>177</sup>Lu–PSMA–617 in a 2–week dose–fractionation regimen
- To assess safety of treatment with the combination of <sup>177</sup>Lu–J591 and <sup>177</sup>Lu–PSMA–617 in a 2–week dose–fractionation regimen as assessed by CTCAE 4.0
- To assess changes in CTC count as measured by CellSearch and the rate of favorable CTC count and LDH at 12 weeks following treatment with the combination of <sup>177</sup>Lu–J591 and <sup>177</sup>Lu–PSMA–617 in a 2–week dose–fractionation regimen
- To assess patient reported outcomes using FACT-P and the Brief Pain Inventory short form

### **1.3 Exploratory Objectives**

- Disease assessment with PSMA-ligand based imaging prior to and following investigational treatment
- To assess immune effects of PSMA-targeted radionuclide therapy
- To assess genomic alterations in relationship to outcome following treatment with the combination of <sup>177</sup>Lu–J591 and <sup>177</sup>Lu–PSMA–617 in a 2-week dose–fractionation regimen
- To estimate whole body distribution and radiation dosimetry of <sup>177</sup>Lu–J591 and <sup>177</sup>Lu–PSMA–617

# 2. BACKGROUND

#### 2.1 Disease

Prostate cancer (PC) is a significant health burden, with 180,890 new diagnoses and 26,120 deaths in the United States in 2016(1). Despite advances in diagnostic technology and treatment strategies, up to 40% of patients treated with primary therapy with curative intent will experience disease progression. PC deaths are typically the result of metastatic castration-resistant prostate cancer (mCRPC), and historically the median survival for men with mCRPC has been less than two years (2). The recent availability of novel treatments for mCRPC has given a resurgence of hope for these men as studies now demonstrate improved survival with a variety of new agents. However, the unfortunate reality is that mCRPC remains an incurable disease, and it is against this backdrop that we look to the future with cautious optimism and new hope for scientific discovery.

First line therapy for advanced PC is androgen deprivation with a mean duration of efficacy of 12–18 months, although there is a wide variation in response in this heterogeneous disease. Upon progression, the disease becomes castration–resistant and subsequently many such patients develop frank metastasis. Exact mechanism of transformation from castration–sensitive prostate cancer to castration–resistant disease is still not fully understood, but with recent scientific innovations in basic research, there is now a better understanding. Now we know that despite castrate levels of androgens, the androgen receptor (AR) remains active and continues to drive prostate cancer progression (3, 4). This understanding has led to the development of novel agents aimed at further decreasing androgen production or blocking AR function. However, there are also many other biologic pathways that function independent of androgen signaling resulting in CRPC. With a greater understanding of the tumor biology, there is hope for continued development of innovative treatment options that improve survival for men with mCRPC.

Treatment of mCRPC has drastically changed over the past decade. Metastatic castration-resistant prostate cancer (mCRPC) poses a particular clinical challenge in need of additional therapeutic approaches beyond classic androgen deprivation therapies. Currently, the chemotherapy compounds docetaxel and cabazitaxel, the androgen receptor signaling inhibitor enzalutamide, CYP-17-inhibitor abiraterone, autologous cellular immuno therapy with sipuleucel-T, and the bone-seeking  $\alpha$ -emitter <sup>223</sup>Ra have shown improved overall survival (OS) and most have demostrated quality of life advantages as well (5-13). These agents have been tested in multiple "disease states" of CRPC to determine if or when patients might benefit from each treatment. In all cases, however, these now established therapies become ineffective in controlling tumor progression over time(14). Novel therapies are urgently needed in order to further ameliorate the course of the disease. Other treatments for men with mCRPC have been shown to improve outcomes, but remain to be approved by the FDA (15).

### 2.2 PSMA

In PC, the most well established, prostate-restricted, cell surface antigen yet identified is prostate specific membrane antigen (PSMA)(16-20). PSMA is a trans-membrane protein with a

707-amino-acid extracellular portion. The PSMA gene (FOLH1) is located on the short arm of chromosome 11.

Although first thought to be entirely prostate-specific, subsequent studies have demonstrated that PSMA is also expressed by cells of the small intestine, proximal renal tubules and salivary glands (20). PSMA is expressed in the apical region of normal prostatic cells, the epithelium surrounding prostatic ducts(21). Dysplastic changes in the prostate result in the expression of PSMA on the luminal surface of prostatic ducts (22, 23). Increasing prostate cancer stage and grade result in higher cell membrane PSMA expression (24, 25). The eventual progression to advanced prostate cancer and castrate resistance corresponds to further increases in PSMA expression (26). PSMA expression in prostate cancer cell membranes is 100- to 1000-fold that in normal cells (24, 25). Thus, PSMA represents a promising target for imaging and therapy of prostate cancer.

### 2.3 Anti–PSMA Monoclonal humanized antibody (J591)

J591 is a de-immunized monoclonal antibody directed at the extracellular domain of human PSMA. Initially J591 was developed as a mouse antibody by Dr. Neil Bander's lab at Weill Cornell Medical Center. Humanized J591 was derived from murine J591 using Biovation's DeImmunization technology (Biovation, Aberdeen, Scotland, UK). In this technology, individual amino acids in predicted B and T cell epitopes were replaced with alternative amino acids such that the murine epitopes are no longer immunogenic to the human immune system.(27) The modification results in a non-immunogenic antibody that may be administered to patients on multiple occasions and over long periods without inducing an immune response. J591 is produced from NS0 cells by Lonza (Lonza Biologics, Slough, UK).

The humanized version of J591 has provided promising results in imaging both localized PC and metastatic disease. Initial phase I/II studies using J591 trace–labeled with <sup>111</sup>Indium (<sup>111</sup>In) and <sup>177</sup>Lutitium (<sup>177</sup>Lu) using a DOTA chelate showed that repetitive dosing was well tolerated without the development of a human anti–humanized antibody (HAHA) response (28-33). No dose limiting toxicity occurred at imaging doses of radionuclide conjugates and the maximum tolerated dose was not reached. Excellent tumor targeting could be detected at all dose levels of the mAb. No mAb targeting to sites other than those involved by PC was observed although, as seen in other trials using radiometals, the liver is the primary site of radiometal metabolism. Percent injected dose in the liver diminished with increasing dose of cold J591, and higher doses were associated with longer plasma clearance times (34-36). The future efforts are to explore further therapeutic options by radiolabeling J591 with more cytotoxic radionuclides like  $\alpha$ – particle emitter Actenium–225 that holds the potential to induce a much stronger anti–tumor response.

# 2.3.1 Prior clinical experience with J591 and radiolabeled-J591 in men with prostate cancer:

### 2.3.1.1 Pilot study with huJ591

Initial phase I studies using huJ591 trace-labeled with <sup>111</sup>In using a DOTA chelate showed that

repetitive dosing was well tolerated with total doses of up to 500 mg/m<sup>2</sup> without the development of a human anti-humanized (de-immunized) antibody (HAHA) response(37). No dose limiting toxicity occurred and the maximum tolerated dose was not reached. Excellent tumor targeting could be detected at all dose levels of mAb. No mAb targeting to non-prostate cancer sites was observed although, as seen in other trials using radiometals, the liver is the primary site of excretion.

#### 2.3.1.2 Phase I and II studies with single dose radiolabeled J591

Two independent phase I clinical trials were initally performed at WCMC using a single–dose of <sup>177</sup>Lu or <sup>90</sup>Y linked via a DOTA chelate to huJ591 in subjects with metastatic, hormone–refractory prostate cancer (28, 31). The primary objectives of these trials were to define the maximum tolerated doses (MTD) of the isotopes as well as to further define dosimetry, pharmakokinetics, and HAHA of the radiolabeled mAb conjugates. Anti–tumor responses were assessed as a secondary endpoint. The design and entry criteria of the 2 trials were identical. Eligible subjects had a prior histologic diagnosis of prostate cancer and evidence of progressing, recurrent or metastatic disease defined by at least 3 serially rising PSAs and/or radiographic studies. As prior studies had demonstrated that all prostate cancers were PSMA–positive (38), no determination of PSMA expression was done.

# i. Phase I Trial of <sup>177</sup>Lutetium–Labeled J591 in subjects with metastatic castration–resistant prostate cancer (CRPC)(28):

<sup>177</sup>Lu–J591 was evaluated in patients with metastatic CRPC demonstrating acceptable toxicity (MTD=70 mCi/m<sup>2</sup>), excellent targeting of metastatic sites and biologic activity. Thirty–five subjects received <sup>177</sup>Lu–J591, of whom 16 received up to three doses. Myelosuppression was dose limiting at 75mCi/m<sup>2</sup>, and the 70mCi/m<sup>2</sup> dose level was determined to be the single–dose MTD. Repeat dosing at 45 to 60mCi/m<sup>2</sup> was associated with dose–limiting myelosuppression; however, up to three doses of 30mCi/m<sup>2</sup> could be safely administered. Nonhematologic toxicity was not dose limiting. Targeting of all known sites of bone and soft tissue metastases was seen in all 30 subjects with positive bone scan, computed tomography, or magnetic resonance images. No subject developed a human anti–J591 antibody response to deimmunized J591 regardless of number of doses. Biologic activity was seen with four subjects experiencing  $\geq$ 50% declines in PSA levels lasting from 3+ to 8 months. An additional 16 subjects (46%) experienced PSA stabilization for a median of 60 days (range, 1 to 21+ months).

# ii. Phase 2 trial of <sup>177</sup>Lutetium (<sup>177</sup>Lu) radiolabeled J591 (<sup>177</sup>Lu–J591) in subjects with metastatic castration–resistant prostate cancer (CRPC) (39):

In a phase II trial patients with progressive metastatic CRPC received a single dose of <sup>177</sup>Lu–J591 in two cohorts (65 and 70mCi/m<sup>2</sup>). Cohort 1: 15 patients; Cohort 2: 17 patients. The primary endpoint was PSA and measurable disease response assessed at week 12 and the secondary endpoint was to evaluate toxicity. One <sup>177</sup>Lu imaging study was done at 1 week post–treatment to

confirm tumor targeting. Three patients achieved PSA declines of >50%; 31% had at least 30% decrease in PSA (the cutoff associated with survival benefit in chemotherapy trials). Though the phase I MTD was 70 mCi/m<sup>2</sup>, based upon FDA restrictions, the initial cohort was treated at 65 mCi/m<sup>2</sup>. In an exploratory analysis, there was a dose–response relationship. The  $\geq$ 30% PSA response rates in the 65 mCi/m<sup>2</sup> and 70 mCi/m<sup>2</sup> cohorts were 13% and 47% respectively (p=0.06); and any PSA decrease in 46% vs. 71% respectively. Hematological toxicity was similar to that in phase I trials; no significant drug–related non–heme toxicity occurred. Grade (Gr) 4 thrombocytopenia occurred in 42% and 9 patients received platelet transfusions. Targeting of known sites of PC metastasis was observed in 30 of 32 (94%) patients.

# iii. Phase I trial of fractionated dose <sup>177</sup>Lu–J591 in men with metastatic castration–resistant prostate cancer (40, 41):

Men with progressive metastatic CRPC received 2 fractionated doses two weeks apart. Initially, 6 cohorts of 3-6 pts got 2 doses of <sup>177</sup>Lu-J591 2 wks apart (20 mCi/m2, escalating to 45 mCi/m2 x2). Subsequently, pts enrolled in 2 expansion cohorts at the recommended phase 2 doses (RP2D). Planar<sup>177</sup>Lu–J591 imaging was semi-quantitatively scored. The endpoints were PSA changes and survival (OS); as well as CTC count (CellSearch) changes in the expansion cohorts. 49 patients, with median age 74.1 years (range 55–95), median PSA of 44.9 ng/mL (1.9–766.5); 83.7% with bone, 61.2% with lymph node, 40.8% with visceral metastasis. 8.2% were CALGB (Halabi) low, 34.7% were intermediate, 57.1% were in high-risk group. RP2D's of fractionated <sup>177</sup>Lu-J591 were 40 mCi/m2 x2 or 45 mCi/m2 x2 with option for GCSF. PSA changes for the low dose group were reported as 6.3% showing >50% PSA decline, 12.5% reporting >30% PSA decline, and 37.5% with any PSA decline. within RP2D group 21.2% showing >50% PSA decline, 42.4% reporting >30% PSA decline, and 66.7% with any PSA decline. The median overall survival for low dose group was 14.6 months and for RP2D group was 27.7 months. Accurate targeting of <sup>177</sup>Lu–J591 was seen in 79.6%. Patients with lower PSMA expression by imaging were less likely to respond (p=0.07). Of 25 with CTC counts, 14 declined, 8 stably favorable, and 3 increased. RP2D was associated with more PSA declines (p=0.036) and longer OS (p=0.004), even after controlling for CALGB prognostic grouping (adjusted HR 0.42 [95% CI 0.21, 0.84] p=0.01). Predictable, reversible myelosuppression was seen. 36 (73.5%) patients had grade 3/4 heme toxicities; 19 (57.6%) had Grade 4 heme toxicities in RP2D cohorts with 45.4% receiving prophylactic platelet transfusions (median 1, range 1-4) and 6 GCSF. 14 (28.6%) had infusion reactions (without pre-meds), with 1 patient having Grade 2 infusion reaction leading to withdrawing from the study prior to his 2<sup>nd</sup> dose. 5 (10.2%) had transient Gr 1/2 AST/ALT. This study concluded that fractionated <sup>177</sup>Lu–J591 is well tolerated with predictable, reversible myelosuppression and PSA and CTC declines. Additionally, with dose-fractionation, the cumulative dose MTD is 14–28% higher than single dose MTD with similar toxicity.

# Table 1: Toxicities data from Phase 2 single dose <sup>177</sup>Lu–J591 Study and Phase 1 fractionated <sup>177</sup>Lu–J591 study:

		Single Dose	Fractionated Dose	
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#### Protocol #1802018988 v2 Version Date: 02/19/2019

Cumulative Dose (mCi/m <sup>2</sup> )	65	70	<b>70</b> (35 + 35)	80 (40 + 40)	90 (45 + 45)
Grade 4 Thrombocytopenia	27%	56.3%	40%	50%	58.8%
Platelet Transfusion	7%	41%	0%	31.3%	52.9%
Grade 4 Neutropenia	0%	37.5%	0%	31.3%	29.4%
Febrile Neutropenia	0%	2.1%	0%	0%	5.8%

In addition, fractionated dosing allowed concurrent dosing with myelosuppressive chemotherapy:

# iv. Phase I trial of fractionated dose <sup>177</sup>Lu–J591 plus docetaxel/prednisone in men with metastatic castration-resistant prostate cancer (42):

Following progression on primary hormonal therapy, chemotherapy can offer symptomatic improvement as well as incremental survival benefit. However, responses are transient and all men eventually suffer from progression of disease as described above with single–agent anti–PSMA based radioimmunotherapy. The combination of taxane chemotherapy with radiotherapy has been used in several diseases because of the radiosensitizing effects of taxane–based chemotherapy. In addition to favorable results from fractionated RIT and the radiosensitizing effects of taxane–based chemotherapy, it is hypothesized that the additional debulking by chemotherapy will overcome some of the limits imposed by the physical characteristics of <sup>177</sup>Lu. Based upon this theory, a phase I trial of docetaxel and prednisone with escalating doses of fractionated <sup>177</sup>Lu–J591 was initiated. 15 men with median age 69.1 (49.3–80.8) were enrolled. The MTD/RP2D of <sup>177</sup>Lu–J591 was 40 mCi/m<sup>2</sup>x2 doses (delivered with cycle 3 of docetaxel), with 73.3% showing >50% PSA decline, 80.0% reporting >30% PSA decline, and 86.7% with any PSA decline. Predictable, reversible myelosuppression was seen. Even at the highest dose level, no dose limiting toxicity was observed, with short–term / reversible grade 4 neutropenia in 33% and grade 4 thrombocytopenia in 13%.

# 2.4 Radionuclide-chelating PSMA ligand

Targeted radionuclide therapy (both monoclonal antibody and/or urea based peptides) is a state of the art and rapidly developing therapy option for different cancer types. The potential advantage of targeted radionuclide therapy is saving the normal tissue while giving a high radiation dose to the tumor.

After rather unsuccessful therapy with <sup>90</sup>Y–CYT–356 monoclonal antibody (mAb) that binds to the intracellular domain of PSMA(43), phases 1 and 2 clinical trials utilizing the PSMA mAb J591, radiolabeled with <sup>90</sup>Y (Yetrium–90) or <sup>177</sup>Lu (Lutetium–177), have shown promising results(28, 34, 35, 39).

Also, PSMA has recently been discovered as a promising target for radioisotope based approaches, both for therapy, for example, using ligands labeled with either  $\beta$ - or  $\alpha$ -emitting nuclides such as <sup>131</sup>I (Iodine-131), <sup>177</sup>Lu (Lutetium-177), and <sup>225</sup>Ac (Actinium-225), and for PET imaging, for example, using <sup>68</sup>Ga-labeled PSMA ligands. Despite encouraging early results, data about intended and unintended effects of PSMA-based radioligand therapies (RLTs) are still scarce.

Results from selective clinical trials using radiolabled PSMA ligand for therapy and diagnosis of prostate cancer are reported here under:

# 2.4.1 <sup>177</sup>Lu–PSMA–617

### a) <sup>177</sup>Lu–DKFZ–PSMA–617 therapy in metastatic castration resistant prostate cancer (44):

Thirty-one mCRPC patients with progressive disease despite second-line hormonal therapy and/or docetaxel chemotherapy were recruited for the study. All underwent <sup>68</sup>Ga-PSMA-HBED-CC PET/CT, prior to therapy with <sup>177</sup>Lu-DKFZ-PSMA-617. The mean activity administered was  $5.07 \pm 1.85$  GBq ranging over one to four cycles. Study reported decline in PSA from baseline. Based on biochemical response criteria, 25/31 had either complete response, partial response or stable disease while 6 continued to have disease progression. The mean ECOG performance status improved from 2.54 to 1.78 after therapy. Two patients experienced grade I and grade II hemoglobin toxicity each, while none experienced nephrotoxicity or hepatotoxicity.

### b) <sup>177</sup>Lu–PSMA–617 as A Novel Therapeutic Option in Patients With Metastatic Castration Resistant Prostate Cancer (45):

28 consecutive patients with mCRPC who have exhausted conventional therapeutic options were treated with 1 or 2 cycles of <sup>177</sup>Lu–PSMA–617. Mean administered activity at first therapy was

 $5.92 \pm 0.44$  GBq of <sup>177</sup>Lu–PSMA–617 and  $5.86 \pm 0.73$  GBq at the second therapy. All patients underwent <sup>68</sup>Ga–PSMA–PET/CT prior to therapy. Any PSA decline occurred in 59% and 75% of patients after 1 and 2 Cycles. Similarly, a PSA decline of 50% or greater occurred in 32% and 50%. Hematologic and renal parameters changed insignificantly; permanent xerostomia or other safety–related toxicity did not occur. The estimated median survival was 29.4 weeks, which was significantly longer than survival in the historical best supportive care group.

#### c) PSMA-Targeted Radionuclide Therapy of Metastatic Castration-Resistant Prostate Cancer with <sup>177</sup>Lu-Labeled PSMA-617 (46):

The authors reported their experience with <sup>177</sup>Lu–PSMA–617–targeted radionuclide therapy in a case series of mCRPC patients resistant to other treatments. Patients were screened with either <sup>99m</sup>Tc–MIP1427 SPECT/CT (500–700 MBq) or <sup>68</sup>Ga–PSMA–11 PET/CT (150 MBq  $\pm$  20%). 30 patients received 1 to 3 cycles of <sup>177</sup>Lu–PSMA–617 (3.7 to 6 GBq per cycle) at 2 months interval. While 21 had a PSA response, 13 had a >50% PSA decrease. After 3 cycles, 8 of the 11 patients achieved a sustained PSA response (>50%) for over 24 weeks, which also correlated with radiologic response (decreased lesion number and size). Acute hematological toxicity was mild. Diffuse bone marrow involvement was a risk factor for higher grade myelosuppression but could be identified by PSMA imaging in advance. Xerostomia, nausea, and fatigue occurred sporadically (<10%). Clearance of non–tumor–bound tracer was predominantly renal and widely completed by 48 hours. Safety dosimetry revealed kidney doses of approximately 0.75 Gy/GBq, red marrow doses of 0.03 Gy/GBq, and salivary gland doses of 1.4 Gy/GBq, irrespective of tumor burden and consistent on subsequent cycles. Mean tumor–absorbed dose ranged from 6 to 22 Gy/GBq during cycle 1.

# d) Pre-therapeutic dosimetry of normal organs and tissues of <sup>177</sup>Lu-PSMA-617 in CRPC (47):

Seven patients were screened after receiving 75–150 MBq of <sup>68</sup>Ga–PSMA–11 intravenously. PET/CT images were acquired 45–60 min post–injection. All had intense tracer uptake at the lesions. They then received 177Lu–PSMA–617 activity ranged from 185 to 210 MBq with a mean of 192.6 ± 11.0 MBq. The highest radiation estimated doses were calculated for parotid glands and kidneys. Calculated radiation–absorbed doses per megabecquerel were  $1.17 \pm 0.31$  mGy for parotid glands and  $0.88 \pm 0.40$  mGy for kidneys. The radiation dose given to the bone marrow was significantly lower than those of kidney and parotid glands (p < 0.05). The calculated radiation dose to bone marrow was  $0.03 \pm 0.01$  mGy/MBq.

# Table 2: Calculated radiation-absorbed doses (mGy/MBq <sup>177</sup>Lu-PSMA-617) of organs of each patient:

Patient	Age	GS	PSA (ng/ml)	Parotid gland	Kidne y	Bone marrow	Live r	Total body	Residence time (h)
1	65	8	48	1.66	0.76	0.025	0.27	0.049	24.6

#### Protocol #1802018988 v2 Version Date: 02/19/2019

2	62	9	110	0.96	1.66	0.048	0.46	0.094	46.3
3	57	8	19	0.94	0.98	0.037	0.23	0.058	49.4
4	65	7	89	1.48	0.52	0.021	0.25	0.035	27.6
5	63	9	235	1.07	0.69	0.058	0.34	0.098	62.0
6	65	9	78	1.25	1.03	0.030	0.22	0.057	26.5
7	70	8	19	0.80	0.51	0.022	0.18	0.036	28.7
Mean	63.9	8.3	85.4	1.17	0.88	0.034	0.28	0.061	37.9
SD	3.9	0.8	74.5	0.31	0.40	0.014	0.09	0.026	14.6

\*GS Gleason score, PSA prostate-specific antigen

#### Table 3: Maximum amount of radioactivity (GBq) to reach radiation-absorbed dose limits:

Patient	Parotid gland (30 Gy)	Kidney (23 Gy)	Bone marrow (2 Gy)	Liver (32 Gy)	Total Body (2 Gy)
1	18.1	30.4	79.6	116.4	41.2
2	31.1	13.8	41.9	69.7	21.4
3	31.8	23.5	54.2	137.6	34.6
4	20.3	44.0	95.7	126.3	57.0
5	28.0	33.2	34.3	95.0	20.3
6	24.0	22.4	66.0	142.4	34.9
7	37.5	44.9	89.7	181.3	56.2
Mean	27.2	30.3	65.9	124.1	37.9
SD	6.9	11.5	23.7	35.7	14.8

\*Organ radiation-absorbed dose constraints in parentheses

#### e) <sup>177</sup>Lu–PSMA Radioligand Therapy of mCRPC: Safety and Efficacy (48):

This study analyzed the safety and efficacy of the <sup>177</sup>Lu-labeled DOTAGA-based PSMA ligand <sup>177</sup>Lu-DOTAGA-(I-y)fk(Sub-KuE) (<sup>177</sup>Lu-PSMA) in patients with mCRPC. 56 patients were <sup>177</sup>Lu-PSMA. enrolled and treated with <sup>68</sup>Ga-PSMA-(N,N'-bis-[2-hydroxy-5-(carboxyethyl)benzyl]ethylenediamine-N,N'-diacetic acid) (68Ga-PSMA) PET/CT was used for patient selection and follow-up after therapy. <sup>177</sup>Lu-PSMA demonstrated high absorbed tumor doses (median 3.3 mGy/MBq) compared with the levels in normal organs. Parotid glands received higher doses (1.3 mGy/MBq) than kidneys (0.8 mGy/MBq). All patients tolerated the therapy without any acute adverse effects. Except for mild reversible xerostomia in 2 patients, no long-term side effects were observed. A decrease in PSA levels was noted in 45 patients (80.4%). Of the 25 patients monitored for at least 6 months after 2 or more therapy cycles, <sup>68</sup>Ga-PSMA PET/CT revealed partial remission in 14, stable disease in 2, and progressive disease in 9 patients. Contrast-enhanced CT revealed partial remission in 5, stable disease in 13, and progressive disease in 7 patients. The median progression-free survival was 13.7 months, and the median overall survival was not reached during follow-up for 28 months.

#### f) German collaborative publication

The largest published experience is a retrospective case series of 145 patients with mCRPC treated at 12 centers in Germany between February, 2014 and July, 2015(49). While not a prospective research study, in general the patients included men who had tumor progression despite abiraterone (64%) and/or enzalutamide (52%) and had either received chemotherapy (54%) or were unfit/refused chemotherapy. Patients also received radium–223 if able/eligible (17%). Treatment was administered only if PSMA uptake was demonstrated on PSMA imaging. On average, 5.9 GBq was administered per dose (range 2–8 GBq). Retreatment was allowed at physician discretion. Of assessable patients (n=99), 60% had PSA decline after initial treatment, with at least 50% PSA decline occurring in 40%. Safety was assessed by retrospective chart review with charts available in all 145 and labs available in 121 (83.4%). Grade >2 hematologic AEs occurred in 12% (grade 3/4 anemia in 10%, grade 3/4 leukopenia in 3%, and grade 3/4 thrombocytopenia in 4%). Other AEs of any grade that occurred  $\geq$ 4% include AST elevation 19%, fatigue 13%, renal failure 12%, ALT elevation 8%, xerostomia 8%, nausea 6%, dysgeusia 4%. While retrospective in nature, this represents the largest published series and demonstrates some biochemical anti-tumor activity without a high occurrence of high–grade adverse events.

#### g) Preliminary data from Australian prospective phase II study

Preliminary results of the first prospective study utilizing <sup>177</sup>Lu-PSMA-617 were reported at the 2017 ESMO annual meeting. <sup>46</sup> Australian investigators enrolled 43 men with mCRPC and treated 30 (7 excluded due to imaging results, 6 others for eligibility issues). Following PSMA and FDG PET/CT, subjects received 4-8 GBq of 177Lu-PSMA-617 and were planned to receive 4 doses every 6 weeks. Seventeen (57%) had at >50% PSA decline with median PFS 6.3 and OS 12.7 months. Toxicity was deemed acceptable. In this prospective study, but adverse events were more common than previously reported in retrospective studies. Xerostomia (grade 1-2) was common at 63%, nausea 50%, vomiting 20%, and fatigue 17%. All cause grade 3/4 anemia occurred in 23%, neutropenia 10%, thrombocytopenia 27%, though attributable grade 3/4 anemia occurred in 7%, neutropenia 7%, and thrombocytopenia 13%.

Additional therapeutic studies using different radiolabled PSMA ligands are summarized in the table below:

	Radioligand Therapy	Ν	Dose	Total doses	Reference
			(GBq)		
1.	<sup>177</sup> Lu–DKFZ–PSMA–617	1	7.4	1 dose	Kratochwil et al. 2015
2.	<sup>177</sup> Lu–DKFZ–PSMA–617	10	4.1 - 6.1	1 dose	Ahmadzedehfar et al. 2015
3.	<sup>177</sup> Lu–PSMA–617	7	0.19 - 0.21	1 dose	Kabasakal et al. 2015
4.	<sup>177</sup> Lu–PSMA I&T	2	5.7-8	1 dose	Weineisen et al. 2015
5.	<sup>177</sup> Lu–DKFZ–PSMA–617	5	3.4 - 3.9	2 doses,	Delker et al. 2016
				10 weeks apart	
6.	<sup>177</sup> Lu–PSMA–617	30	3.7 - 6	1-3 dose(s),	Kratochwil et al. 2016
				2 months apart	
7.	<sup>177</sup> Lu–PSMA–617	9	5.28 - 5.77	1 dose	Hohberg et al. 2016
8.	<sup>177</sup> Lu–PSMA–617	7	5.5 - 7.4	1 dose	Das et al. 2016
9.	<sup>177</sup> Lu–PSMA–617	23	7.4	1 dose	Demir et al. 2016
10.	<sup>177</sup> Lu–PSMA–617	28	5.13 - 6.59	1 - 2  dose(s),	Rahbar et al. 2016
				2 months apart	
11.	<sup>177</sup> Lu–PSMA–617	82	5.4 - 6.4)	1 dose	Rahbar et al. 2016
12.	<sup>177</sup> Lu–PSMA–617	31	1.1 - 5.5	$1 - 4 \operatorname{dose}(s)$	Yadav et al. 2016
13.	<sup>177</sup> Lu–PSMA–617	1	NA	3 doses,	Schlenkhoff et al. 2016
				8 weeks apart	
14.	<sup>177</sup> Lu–PSMA–617	43	6	$1 - 4 \operatorname{dose}(s)$	Zimbelmann et al. 2016
15.	<sup>177</sup> Lu–PSMA–617	5	5.8 - 7.4	1 dose	Chakraborty et al. 2016
16.	<sup>177</sup> Lu–PSMA–617 +	1	1.85 + 1.3	1 dose each	Bal et al. 2016
	<sup>177</sup> Lu–EDTMP				
17.	<sup>177</sup> Lu–PSMA I&T	22	3.4 - 7.4	1 - 4  dose(s),	Heck et al. 2016
				2 weeks apart	
18.	<sup>177</sup> Lu–PSMA I&T	56	3.6 - 8.7	$1-5 \operatorname{dose}(s)$	Baum et al. 2016
19.	<sup>177</sup> Lu–PSMA I&T	18	7.4	$1-4 \operatorname{dose}(s)$	Okamoto et al. 2017
20.	177Lu-PSMA-617	15	3.7 - 6	2 doses	Fendler et al. 2017
21.	<sup>131</sup> I-MIP-1095	36	1.5 - 7.2	$1 - 3 \operatorname{dose}(s)$	Afshar–Oromeih et al. 2017
22.	<sup>177</sup> Lu–PSMA–617	1	5.9-6.4	4 doses	Roll et al. 2017

# Table 4: <sup>117</sup>Lu–PSMA–617: Pilot studies of targeted radionuclide therapy:

# h. Fractionated dose <sup>177</sup>Lu-PSMA-617

Based upon our prior experience with radiolabeled J591 (demonstrating dose and toxicity response with improved safety with dose fractionation) and the worldwide experience of radiolabeled small molecules, we initiated a dose-escalation study of fractionated dose <sup>177</sup>Lu-PSMA-617 in January, 2018. As of the time of current protocol writing, 4 cohorts have been treated with safety follow up (and the 5<sup>th</sup> / final planned cohort is enrolled), with administered/planned dose levels described in the Table below. As predicted based upon dosimetry estimations, **no dose-liming toxicity has been observed to date, including all potential dose-levels to be utilized in this protocol**.

Cohort	Treatment Dose mCi (GBq)		Total dose	# Subjects
	Day 1	Day 15	mCi (GBq)	(n)
-1	50 (1.85)	50 (1.85)	100 (3.7)	0
1	100 (3.7)	100 (3.7)	200 (7.4)	6
2	150 (5.55)	150 (5.55)	300 (11.1)	6
3	200 (7.4)	200 (7.4)	400 (14.8)	5
4	250 (9.25)	250 (9.25)	500 (18.5)	6
5	300 (11.1)	300 (11.1)	600 (22.2)	pending

Table: Dose-escalation strategy of fractionated dose <sup>177</sup>Lu-PSMA-617 protocol

# 2.4.2 <sup>68</sup>Ga-PSMA-HBED-CC (<sup>68</sup>Ga-PSMA-11)

PSMA has recently been discovered as a promising target for radioisotope based approaches for PET imaging, for example, using <sup>68</sup>Ga–labeled PSMA ligands(50-55). All of these trials report retrospective data on case reports/case series. No well–designed prospective clinical trial has been published/reported yet.

During the last two decades, many efforts have been undertaken to develop PSMA–ligands(56, 57). One of these ligands, the small molecule Glu–NH–CO–NH–Lys–(Ahx)–[<sup>68</sup>Ga(HBED–CC)], also known as PSMA–11<sup>®</sup>, PSMAHBED, Glu–CO–Lys(Ahx)–HBED–CC, DKFZ–PSMA–11, PSMA–HBED– CC, PSMA–HBED, PSMA or Prostamedix<sup>TM</sup>, developed at the German Cancer Research Center Heidelberg (DKFZ), has become the most clinically used radiotracer. This compound shows a strong binding affinity to PSMA as well as a highly efficient internalization into PCa cells(58, 59). PET/CT–imaging with <sup>68</sup>Ga–PSMA–11 has demonstrated this novel method as an important imaging modality for diagnosing recurrent PCa (50, 60-62). Perera et al., did systematic review of <sup>68</sup>Ga–PET articles and Sixteen articles involving 1309 patients were analyzed. On per–patient analysis, the summary sensitivity and specificity were 80% and 97%, respectively(63).

Results from selective clinical trials using <sup>68</sup>Ga–PSMA–HBED–CC to evaluate its diagnositic value are reported here under:

# i. Diagnostic value of <sup>68</sup>Ga-PSMA-HBED-CC PET/CT imaging in the diagnosis of recurrent prostate cancer(61):

The study reports data from a retrospective analysis of 319 patients who underwent  ${}^{68}$ Ga-PSMA-ligand PET/CT from 2011 to 2014. The  ${}^{68}$ Ga-PSMA-HBED-CC solution was applied to the respective patient via an intravenous bolus injection (mean of 172.4 MBq ±70.9, range 40 – 400 MBq, median 161 MBq). A non-contrast-enhanced CT scan was performed 1-hour post tracer injection. Histological verification was performed in 42 patients after the  ${}^{68}$ Ga-PSMA-ligand PET/CT. Tracer uptake was measured in 901 representative tumor lesions. 82.8% of the patients had at least one lesion indicative of PCa was detected. Tumor-detection was positively associated with PSA level and Androgen deprivation therapy (ADT). Gelason Score and PSA doubling time (PSA-DT) were not associated with tumor-detection. The average maximum standardized uptake value (SUVmax) of tumor lesions was 13.3 ± 14.6 (0.7-122.5). Amongst lesions investigated by histology, 30 were false-negative in 4 different patients, and all other lesions (n = 416) were true-positive or true-negative. A lesion-based analysis of sensitivity, specificity, negative predictive value (NPV) and positive predictive value (PPV) revealed values of 76.6%, 100%, 91.4% and 100%. A patient-based analysis revealed a sensitivity of 88.1%.

# ii. **PET imaging with a <sup>68</sup>Ga–PSMA ligand for the diagnosis of prostate cancer:** biodistribution in humans and first evaluation of tumor lesions (64):

Initial clinical studies with the <sup>68</sup>Ga–labeled PSMA–HBED–CC were conducted at Heidelberg University Hospital and the German Cancer Research Center to assess the biodistribution of <sup>68</sup>Ga–PSMA–HBED–CC in normal tissues and tumor lesions. A total of 37 patients with prostate cancer and rising PSA levels were subjected to <sup>68</sup>Ga–PSMA–HBED–CC PET/CT imaging. Quantitative assessment of tracer uptake was performed 1 and 3 h post–injection by analysis of mean and maximum standardized uptake values (SUVmean/max) of several organs and 65 tumor lesions. Subsequently, tumor to background ratios were calculated.

The <sup>68</sup>Ga–PSMA–HBED–CC PET/CT images showed intense tracer uptake in both kidneys and salivary glands. Moderate uptake was seen in lacrimal glands, liver, spleen and in small and large bowel. Quantitative assessment revealed excellent contrast between tumor lesions and most normal tissues. Of 37 patients, 31 (83.8 %) showed at least one lesion suspicious for cancer at a detection rate of 60 % at PSA <2.2 ng/ml and 100 % at PSA >2.2 ng/ml. Median tumor to background ratios were 18.8 (2.4–158.3) in early images and 28.3 (2.9–224.0) in late images. Within healthy organs, kidneys and salivary glands demonstrated the highest radiotracer uptake. Lesions suspicious for prostate cancer presented with excellent contrast as early as 1–hour after injection with high detection rates even at low PSA levels.

#### iii. <sup>68</sup>Ga-labeled PSMA ligand as superior PET tracer for the diagnosis of prostate

# cancer: Comparison with <sup>18</sup>F-FECH (52):

This study was also published by the Heidelberg group, compared <sup>68</sup>Ga–PSMA– HBED–CC PET/CT imaging to standard choline–based PET/CT. Thirty–seven patients with biochemical relapse of prostate cancer [mean prostate–specific antigen (PSA) 11.1  $\pm$  24.1 ng/ml, range 0.01–116] were retrospectively analyzed after <sup>18</sup>F– fluoromethylcholine and <sup>68</sup>Ga–PSMA PET/CT within a time window of 30 days. Radiotracer uptake that was visually considered as prostate cancer was semi– quantitatively analyzed by measuring the maximum standardized uptake values (SUVmax) of the scans acquired 1–hour after injection of <sup>68</sup>Ga–PSMA complex solution (median 132 MBq, range 59–263 MBq) and <sup>18</sup>F–fluoromethylcholine (median 237 MBq, range 114–374 MBq), respectively. In addition, tumor to background ratios were calculated.

The results showed a total of 78 lesions characteristic for prostate cancer that were detected in 32 patients using <sup>68</sup>Ga–PSMA–HBED–CC PET/CT imaging and 56 lesions were detected in 26 patients using choline PET/CT. The higher detection rate in <sup>68</sup>Ga–PSMA–HBED–CC PET/CT imaging was statistically significant (p=0.04). In five patients, no lesion was found with both methods. All lesions detected by <sup>18</sup>F–fluoromethylcholine PET/CT were also seen by <sup>68</sup>Ga–PSMA–HBED–CC PET/CT imaging. In <sup>68</sup>Ga–PSMA–HBED–CC PET/CT imaging SUVmax was clearly (>10 %) higher in 62 of 78 lesions (79.1 %) and the tumor to background ratio was clearly (>10 %) higher in 74 of 78 lesions (94.9 %) when compared to <sup>18</sup>F–fluoromethylcholine PET/CT.

The authors concluded that <sup>68</sup>Ga–PSMA–HBED–CC PET/CT can detect lesions characteristic for prostate cancer with improved contrast when compared to standard <sup>18</sup>F–fluoromethylcholine PET/CT, especially at low PSA levels.

# iv. Comparison of PET/CT and PET/MRI hybrid systems using a <sup>68</sup>Ga–labelled PSMA ligand for the diagnosis of recurrent prostate cancer: initial experience (60):

In a more recent publication, the Heidelberg group evaluated the feasibility of PET/MRI imaging with <sup>68</sup>Ga–PSMA–HBED–CC. Twenty patients underwent PET/CT 1–hour after injection of the <sup>68</sup>Ga–PSMA–HBED–CC followed by PET/MRI 3–hours after injection. Data from the two investigations were first analyzed separately and then compared with respect to tumor detection rate and radiotracer uptake in various tissues. To evaluate the quantification accuracy of the PET/MRI system, differences in SUVs between PET/CT and corresponding PET/MRI were

compared with differences in SUVs between PET/CT 1-hour and 3-hours after injection in another patient cohort. This cohort was investigated using the same PET/CT system. With PET/MRI, different diagnostic sequences, higher contrast of lesions and higher resolution of MRI enabled a subjectively easier evaluation of the images. In addition, four unclear findings on PET/CT could be clarified as characteristic of prostate cancer metastases by PET/MRI. However, in PET images of the PET/MRI, a reduced signal was observed at the level of the kidneys (in 11 patients) and around the urinary bladder (in 15 patients). This led to reduced SUVs in six lesions. SUVmean values provided by the PET/MRI system were different in muscles, blood pool, liver and spleen.

The authors concluded that prostate cancer was detected more easily and more accurately with <sup>68</sup>Ga–PSMA PET/MRI than with PET/CT and with lower radiation exposure. Consequently, this new technique could clarify unclear findings on PET/CT. However, scatter correction was challenging when the specific <sup>68</sup>Ga–PSMA–HBED–CC was used. Moreover, direct comparison of SUVs from PET/CT and PET/MR needs to be conducted carefully.

These encouraging results suggest <sup>68</sup>Ga–PSMA–HBED–CC to be an effective probe for the PC marker "PSMA" in metastatic prostate cancer patients and that it can provide a robust imaging scan to better detect PC tumor burden. We plan to utilize this probe to better understand its diagnostic role in prostate cancer patients with localized disease.

### 2.6 Investigational Agent

# • <sup>177</sup>Lu–J591 (IND #11613)

### i. J591

The J591 Mab is a deimmunized monoclonal antibody directed at the extracellular domain of human PSMA. J591 was derived from murine J591 using Biovation's (Aberdeen, Scotland) Deimmunisation technology in which individual amino acids in predicted B and T cell epitopes were replaced with other amino acids such that the epitope would no longer be recognized by the human immune system, thereby decreasing the likelihood of the development of an anti–mAb antibody response in humans (65). This results in a potentially non–immunogenic antibody, which can be administered to patients on multiple occasions over long periods without inducing an immune response. Furthermore, the deimmunized Mab additionally has been engineered to possess the effect of inducing antibody dependent cellular cytotoxicity (ADCC) with human immune effector cells. J591 is produced from NS0 cells by Lonza Biologics (Slough, UK). The molecular weight of J591 is approximately 147,000 daltons as determined by Matrix Assisted Laser Desorption Mass Spectrometry (MALDI–TOF). The naked antibody is formulated in a 50 mM sodium phosphate, pH 5.5, containing 100 mM sodium chloride and 2 mM EDTA at a nominal concentration of 5 mg/mL.

## ii. DOTA-J591

In order to radiolabel J591 with  $\beta^-$  emitting radionuclides such as <sup>177</sup>Lu, the J591 antibody molecule is first conjugated to a macrocyclic chelating agent, 1,4,7,10–tetraazacyclododecane–N,N',N'',N'''–tetraacetic acid (DOTA) by direct coupling of one of the four carboxylic acid groups of DOTA to the primary amines present in the protein structure. The DOTA conjugated antibody is formulated in 0.3 M ammonium acetate, pH 7.0 at a nominal concentration of 8 mg/mL.

# iii. Radiolabeled J591 (<sup>177</sup>Lu–J591)

The DOTA–J591 antibody molecule is labeled with radiometal <sup>177</sup>Lu in 1.0 M ammonium acetate buffer at pH 7.0. The radiolabeled J591 complex is then purified using size–exclusion column chromatography. To avoid the effects of autoradiolysis on the antibody, reaction time is minimized and the radiolabeled antibody preparations are formulated in phosphate buffered saline containing not more than 1% Human Serum Albumin (HSA). All reagents used in the conjugation and purification of J591 are made with pyrogen–free water. Radiolabeled J591 is periodically tested for sterility and pyrogenicity.

Each patient dose will be supplied to the study site on the day of or day before treatment by the central radiopharmacy. The J591 mAb is a deimmunized monoclonal antibody directed at the extracellular domain of human PSMA. For additional details, see Section 1 and/or Investigator Brochure. Radiolabeled antibody preparations are formulated in phosphate buffered saline containing not more than 1% Human Serum Albumin (HSA).In order to radiolabel J591 with  $\beta^-$  emitting radionuclides, the antibody molecule is first conjugated to a macrocyclic chelating agent, 1,4,7,10–tetraazacyclododecane– N,N',N'',N'''-tetraacetic acid (DOTA) by direct coupling of one of the four carboxylic acid groups of DOTA to the primary amines present in the protein structure. The radiolabeling of J591 with <sup>177</sup>Lu is achieved by adding the radionuclide to the ammonium acetate buffered J591. To avoid the effects of autoradiolysis on the antibody, reaction time is minimized and the reaction mixture is purified with a size exclusion column prior to administration. All reagents used in the conjugation and purification of J591 are made from pyrogen–free water. Radiolabeled J591 is tested periodically for sterility and pyrogenicity.

### iv. Pharmacokinetics

The pharmacokinetics of J591 antibody was studied in patients with prostate cancer using <sup>111</sup>In– J591 as a radiotracer. In a Phase 1 dose escalation study (see above) 4 groups of patients received different amounts of J591 antibody (25, 50, 100, 200 mg/m<sup>2</sup>) mixed with a trace amount of <sup>111</sup>In– J591 (5 mCi/1–2 mg). In a different Phase 1 dose–escalation study using <sup>90</sup>Y–J591 (see above), a group of patients received 20 mg of total J591 antibody mixed with a trace amount of <sup>111</sup>In–J591 (5 mCi/1–2 mg) for imaging studies prior to <sup>90</sup>Y treatment dose. These studies demonstrated that the plasma clearance kinetics of <sup>111</sup>In–J591 antibody is dose–dependent at low doses (<50 mg/m<sup>2</sup>). It should be noted that, in these studies, plasma clearance kinetics was determined based on the measurement of radioactivity in the plasma over time. The actual J591 antibody levels have not been measured; there is an assumption that the clearance of radiolabeled antibody does indeed reflect the J591 antibody clearance. The plasma clearance kinetics, based on bi–exponential fitting of plasma–time activity curves indicate that the terminal half–life ( $T_{1/2}\beta$ ) is about 44.2 ± 13.9 hrs at 11 mg/m<sup>2</sup> antibody dose and 59.3±16.6 hrs at 25 mg/m<sup>2</sup> antibody dose. At higher doses of antibody mass, the plasma clearance is relatively slower, but stable. In summary, the plasma clearance of antibody increases at lower doses of antibody (<25 mg/m<sup>2</sup>). In addition, repeat administration of J591 antibody over a 4–week period does not affect plasma clearance of radiolabeled J591 in patients with prostate cancer (data not shown). In a Phase 1 dose–escalation study using <sup>177</sup>Lu–J591 (see above), a group of patients received J591 antibody mixed with a <sup>177</sup>Lu–J591 (10–70 mCi/m<sup>2</sup> and total J591 10mg/m<sup>2</sup>). The plasma clearance kinetics (shown in **Table–5**) demonstrate that there is no difference between <sup>177</sup>Lu and <sup>111</sup>In labeled J591 antibody.

Table–5:	<sup>177</sup> Lu–J591 Vs.	$^{111}$ In–J591:	Plasma	Clearance Kinetics:
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Pts.	mg/m <sup>2</sup>	T½ α (hr)	T½ β (hr)	Vd at To	Clearance
n=18 <sup>#</sup>	10	3.61±5.70	46.8±12	3949±1067	72±34
n=16*	11	2.37±1.94	44.2±14	4042±863	94±34
n=3 <sup>\$</sup>	25	3.36±2.17	59.3±17	3698±224	56±16
n=3\$	50	7.83±9.5	99.7±32	4156±256	47±6.7
n=3 <sup>\$</sup>	100	5.7±7.5	105±38	4373±566	40±13

<sup>#</sup> = In a Phase 1 Dose escalation protocol, 18 patients were treated with  $^{177}$ Lu–J591

\* = In a Phase 1 Dose escalation protocol, 16 Patients were treated with  ${}^{90}$ Y–J591 and one week prior to treatment,  ${}^{111}$ In–J591 kinetics were studied.

 $^{$}$  = In a Phase 1 Dose escalation protocol with huJ591 cold antibody treatment, 3 patients in each group received different amounts of cold huJ591 mixed with <sup>111</sup>In–J591.

#### v. Biodistribution

The imaging studies with <sup>111</sup>In–J591 in patients with prostate cancer showed that liver is the principal organ with significant uptake of radiolabeled antibody. The liver uptake of J591 is a function of J591 antibody mass administered (**Table–6**): the higher the antibody mass, the lower the relative proportional uptake of radiolabeled antibody by the liver. In addition, at 20 mg of J591 administered in Phase 1 dose escalation trials with <sup>111</sup>In–DOTA–J591, the liver uptake increases with time and is at a maximum by day 6 ( $27.0 \pm 1.7$  % of injected dose). No significant localization was seen in other major organs, such as spleen and kidneys.

Biodistribution: <sup>111</sup>In–J591 vs. <sup>177</sup>Lu–J591. The imaging studies were done 5 times over a period of one week with <sup>111</sup>In and 2 weeks with <sup>177</sup>Lu. The uptake of radioactivity in various source

organs (% injected dose/organ) was determined. The imaging studies clearly document that the in vivo distribution of radiolabeled J591 during the first week is quite similar with both <sup>111</sup>In and <sup>177</sup>Lu radionuclides (**Table-7**).

Time	<u>11 mg/m<sup>2</sup></u>	<u>25 mg/m<sup>2</sup></u>	<u>50 mg/m<sup>2</sup></u>
1	$13.1 \pm 2.5$	9.6 ± 1.0	$7.9 \pm 0.8$
24	$19.0 \pm 3.2$	$12.9\pm2.1$	$11.2 \pm 1.2$
48	$20.8 \pm 2.8$	ND	$11.0 \pm 0.4$
72	$22.5\pm2.4$	$14.5 \pm 1.2$	$12.1 \pm 1.0$
120	$\textbf{24.6} \pm \textbf{2.8}$	ND	ND
144	$\textbf{27.0} \pm \textbf{1.7}$	$17.1 \pm 3.2$	$13.9\pm1.0$

Table–6:	<sup>111</sup> In–J591:	Liver Uptake	(% injected	dose) as a Funct	tion of J591 Antibod	ly Mass:
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Table-7: Biodistribution of Radiolabeled J591 Antibody:

	<sup>111</sup> In–J591			<sup>177</sup> Lu–J591				
Organ	DAY 0		DAY 6		DAY 0		DAY 7	
Whole Body	100.0	± 0.0	66.1	± 18.4	100.0	± 0.0	69.8	± 9.5
Remainder	52.3	± 5.7	24.9	± 16.8	57.5	± 6.4	34.7	± 6.9
Heart Contents	8.9	± 2.3	3.4	± 1.2	8.8	± 2.2	3.1	± 0.9
Kidneys	2.4	± 0.8	2.5	± 0.7	2.2	± 0.5	2.5	± 0.5
Liver	14.7	± 3.2	28.0	± 7.8	15.0	± 3.8	23.7	± 7.0
Lungs	9.0	± 2.0	4.4	± 1.7	6.9	± 1.9	3.1	± 0.7
Red Marrow	10.9	± 5.0	1.2	± 1.1	7.4	± 2.4	0.6	± 0.4
Spleen	1.8	± 0.6	1.7	± 0.8	2.2	± 1.0	2.1	± 0.8

#### vi. Radiation Dosimetry: <sup>177</sup>Lu–J591

Based on imaging studies and plasma clearance with <sup>177</sup>Lu–J591, the residence times ( $\tau$ ) of radioactivity in various source organs (as shown in **Table**–7) were calculated. Using "S" factors from MIRD tables, radiation dosimetry (cGy/mCi) for <sup>177</sup>Lu–J591 was estimated (**Table–8**). Following administration of <sup>177</sup>Lu–J591, the liver is the critical organ, receiving 7.77± 2.23 rads/mCi, and the kidneys and spleen receive <7 rads/mCi. The bone marrow dose was estimated to be 1.17± 0.37 rads/mCi based on the assumption that 36% of bone marrow volume represents plasma volume. Hematopoietic toxicity is usually the dose–limiting factor in RIT(66). Radiation dose of 9.0 Gy to the bone marrow is generally regarded as an "organ injury dose" and the tolerable

absorbed doses were usually estimated to be 65% of LD50 (50%) organ injury dose. Therefore, the tolerable absorbed dose for bone marrow is about 3.0 Gy (330 rads). The maximum tolerable radiation doses to kidneys, liver and spleen are approximately 20–25 Gy (2000–2500 rads). Dosimetric studies using radiolabeled antibodies have been effectively used by many investigators to determine patient–specific radiation dosimetry of radiolabeled antibody treatment(67, 68). Based on the dosimetry data shown in **Table–8**, the maximum bone marrow dose following administration of 70mCi/m<sup>2</sup> of <sup>177</sup>Lu–J591 would deliver 164 rads to an average person with a BSA of 2.0 m<sup>2</sup>. The maximum dose the liver would be about 1100 rads, well below the maximum tolerable liver dose.

Organ	Mean		S.D.	
Adrenals	0.52	±	0.08	
Brain	0.40	±	0.07	
Gall Bladder	0.55	±	0.08	
LLI Wall	0.42	±	0.07	
Small Intestine	0.43	±	0.07	
Stomach	0.45	±	0.07	
ULI Wall	0.44	±	0.07	
Heart Wall	3.50	±	0.70	
Kidneys	5.20	±	1.29	
Liver	7.77	±	2.23	
Lungs	2.79	÷	0.80	
Muscle	0.42	±	0.07	
Pancreas	0.52	±	0.08	
Red Marrow	1.17	±	0.37	
Bone Surfaces	0.70	±	0.14	
Skin	0.42	±	0.07	
Spleen	7.28	±	3.41	
Testes	0.36	±	0.13	
Thymus	0.44	±	0.07	
Thyroid	0.41	±	0.07	
Urin. Bladder Wall	0.97	±	0.22	
Total Body	0.71	±	0.10	
Effective Dose Equiv.	2.14	±	0.35	
Effective Dose	1.33	±	0.19	

#### Table-8: Radiation Dosimetry (rads/mCi) of <sup>177</sup>Lu-J591 in Patients with Prostate Cancer:

### • 68Ga-PSMA-HBED-CC (IND # 124495)

PSMA has recently been discovered as a promising target for radioisotope based approaches for PET imaging, for example, using <sup>68</sup>Ga–labeled PSMA ligands(50-55). All of these trials report retrospective data on case reports/case series. No well–designed prospective clinical trial has been published/reported yet.

During the last two decades, many efforts have been undertaken to develop PSMA–ligands(56, 57). One of these ligands, the small molecule Glu–NH–CO–NH–Lys–(Ahx)–[<sup>68</sup>Ga(HBED–CC)], also known as PSMA–11<sup>®</sup>, PSMAHBED, Glu–CO–Lys(Ahx)–HBED–CC, DKFZ–PSMA–11, PSMA–HBED– CC, PSMA–HBED, PSMA or Prostamedix<sup>TM</sup>, developed at the German Cancer Research Center Heidelberg (DKFZ), has become the most clinically used radiotracer. This compound shows a strong binding affinity to PSMA as well as a highly efficient internalization into PCa cells(58, 59). PET/CT–imaging with <sup>68</sup>Ga–PSMA–11 has demonstrated this novel method as an important imaging modality for diagnosing recurrent PCa(50, 60-62). Perera et al., did systematic review of <sup>68</sup>Ga–PET articles and Sixteen articles involving 1309 patients were analyzed. On per–patient analysis, the summary sensitivity and specificity were 80% and 97%, respectively(63).

ABX Advanced Biochemical Compounds, Radeberg, Germany produces PSMA–HBED–CC for clinical use. Upon purchase and shipment to Weill Cornell Medicine, the peptide will be labeled with <sup>68</sup>Ga with the final product being <sup>68</sup>Ga–PSMA–HBED–CC. Radionuclide conjugation will be done at the Division of Nuclear Medicine, Department of Radiology, Weill Cornell Medicine.

The material produced are subjected to quality assurance testing as outlined in the Food and Drug Administration "Points to Consider". All production data and supporting documents, together with the results of all quality assurance testing, have been provided to and reviewed by the Food and Drug Administration (<sup>68</sup>Ga–PSMA–HBED–CC, IND# 124495).

### i. Radiation Dosimetry: <sup>68</sup>Ga–PSMA–HBED–CC

Radiation dose estimates were provided by Professor Marcus Schwaiger of Technische Universitat Miinchen (TUM), Munich, Germany. Balb1–c mice were used to determine the biodistribution at 4 time points after administration. Tissues were dissected after sacrifice and the percent of the administered dose was determined per organ and per gram of organ weight. The data of the biodistribution were extrapolated from mouse to man using OLINDA/EXM 1.0 Dosimetry Software (64; Stabin MG, et al. J Nucl Med. 2005 Jun;46(6):1023–7.).

Administered ddose (Mbq)	150 +/20	185 +/20		
Administered dose (mCi)	(4.0)	(5.0)		
organ	[mSv] per organ	[mSv] per organ		
Brain	1.64E-01	2.05E-01		
Lung	8.17E-01	1.02E+00		
Liver	1.30E+00	1.63E+00		
Kidneys	1.11E+02	1.39E+02		
Stomach Wall	3.32E+00	4.15E+00		
Spleen	7.96E+00	9.95E+00		
Pancreas	2.80E+00	3.50E+00		
Red Bone Marrow	1.35E+00	1.69E+00		
Bladder Wall	1.41E+00	1.76E+00		
Effective Dose	1.94E+00	2.43E+00		

Table 9: Radiation dose estimates for Humans administered 148 or 185 MBq (4 or 5 mCi) of 68Ga–PSMA–HBED–CC:

For comparison purposes, the average per capita annual exposure from natural background in the continental USA is 3.0 mSv. (*Ionizing radiation exposure of the population of the United States. Bethesda, Md.: National Council on Radiation Protection and Measurements. 2009. ISBN* 978–0–929600–98–7. NCRP No. 160.)

### • <sup>177</sup>Lu–PSMA–617 (IND # 131966)

Modifications of PSMA-11 have resulted in the development of another novel small molecule PSMA-ligand, PSMA-617. In preclinical studies, this ligand showed a significantly improved binding affinity to PSMA as well as a highly efficient internalization into PCa cells(69).

PSMA-617 can be labeled with radionuclides like <sup>68</sup>Ga(Gallium-68), <sup>177</sup>Lu(Lutetium-177), <sup>111</sup>In(Indium-111), and <sup>90</sup>Y(Yetrium-90) and, therefore, be used for PET-imaging as well as for radioligand-based therapy. Preclinical assays of PSMA-617 showed Ki values of  $2.3 \pm 2.9$  nM, demonstrating a significant improvement compared to PSMA-11 ( $12.0 \pm 2.8$  nM). Based on these results, PSMA-617 has one of the highest binding affinities to the PSMA receptor, which have been published, so far. In preclinical studies, tumor-to-background ratios of up to 1,058 were observed at 24h post infusion. In addition, the internalization of the PSMA-617 into the PCa cells is highly effective: Internalized fraction:  $17.67 \pm 4.34$  % IA/106 LNCaP cells (PSMA-11: 9.47 ± 2.56 % IA/106 LNCaP cells)(59, 69). Since 2013, <sup>177</sup>Lu-PSMA-617 has been increasingly used for radioligand therapy of metastatic PCa patients in several centers (Bad Homburg, Bonn, Cologne, Freiburg, Heidelberg, Istanbul, Melbourne, LMU Munich, Münster)(70).

Radiation dosimetry and organ specific dosage details are described in section 2.4.1 and Table 2 & 3 above. Toxicity data from various studies on <sup>177</sup>Lu–PSMA–617 is described in section 2.7 Table 12.

Diagnostic evaluation with <sup>68</sup>Ga–PSMA–HBED–CC followed by therapy with <sup>177</sup>Lu–PSMA–617 is thus opening a new theranostic potential of nuclear medicine. Initial studies in patients have demonstrated the utility of <sup>177</sup>Lu–PSMA–617 for the treatment of disseminated prostate cancer(70, 71). Dosimetric evaluation has shown that the critical organs are the parotid glands, and that up to 27 GBq (~700 mCi) of <sup>177</sup>Lu–PSMA–617 can be administered without crossing the 30 Gy critical dose threshold to the parotid glands(47).

# 2.7 Rationale

Prostate cancer (PC) is the most common solid male cancer. For those patients with clinically organ-confined disease, radical prostatectomy (RP) remains the most commonly used treatment modality (72). However, even after surgical treatment, high risk PC is prone to recur and often becomes resistant to conventional hormonal therapy (73). For this reason, efforts have been made to optimize patient outcome in the adjuvant and neoadjuvant settings (74-76). Thus far, neoadjuvant therapies have yet to demonstrate a survival benefit (77). Therefore, newer strategies to prevent or rescue PC recurrences are highly desirable.

It is now recognized that evasion of an effective immune response is a hallmark of cancer (78). Immunotherapy may induce tumor cell death with a relatively high degree of specificity unlike hormonal or chemotherapy, which has a non-specific action. PC is an ideal target for immunotherapy because of its indolent nature which allows adequate time to activate the immune system (79), and in fact, the first therapeutic cancer vaccine to be approved was for PC (10). Identification of PC tumor associated antigens (TAA) has provided ideal targets for immunotherapy, with prostate-specific membrane antigen (PSMA) as one of the most relevant targets in oncology as a highly restricted cell-surface membrane protein that is not secreted (18, 19, 80, 81).

Based on the knowledge that certain tumor types express significant amounts of one or several

specific cell–surface proteins, the development of drugs aimed at these targets is becoming an increasingly important component in the treatment of cancer. One example is the successful use of the mAb rituximab in patients with non–Hodgkin's lymphoma (NHL). Rituximab targets CD20 which is expressed on the surface of 95% of B cells in NHL (82). Another clinically validated target is human epidermal growth factor receptor 2 (HER2), which in breast cancer is targeted by trastuzumab (83). In PC, several studies, which will be addressed throughout this protocol, have emphasized PSMA as both a diagnostic and therapeutic target. Similar to prostate–specific antigen (PSA), PSMA is a highly expressed, prostate–specific biomarker. While PSMA is not a novel discovery, it remains a highly relevant target with the knowledge that attenuated AR signaling even prior to hormonal therapy is the hallmark of lethal PC and recent studies have confirmed the reciprocal relationship between PSMA and AR signaling in the era of highly potent AR–directed therapy(26, 84-88).

#### A. Rationale for the appropriate radioisotope:

As described earlier, targeted radionuclide therapy as a treatment choice for mCRPC patients is being actively studied at various centers in the world (especially Europe and Australia). When compared to monoclonal antibodies, the low molecular weight compounds have a higher permeability into solid tumors, offer a significant advantage in achieving higher tumor uptake as well as a high percentage of specific binding. Additionally, small molecules display more rapid tissue distribution and faster blood clearance compared with intact immunoglobulins. These properties often lead to an enhanced target to non-target tissue ratio, which is important not just for imaging but also for successful application of therapeutic absorbed doses.

Following tumor localization, the radiometal is trapped within the cell, leading to higher accretion of radionuclide by the tumor. <sup>131</sup>I has lower energy  $\beta^{-}$  particles and longer physical half–life (max  $\beta^{-} 0.61 \text{ MeV}$ ;  $T_{\frac{1}{2}} = 8.04 \text{ d}$ ) compared to that with <sup>90</sup>Y(max  $\beta^{-} 2.28 \text{ MeV}$ ;  $T_{\frac{1}{2}} = 2.67 \text{ d}$ ). Because of its higher energy, <sup>90</sup>Y may be appropriate for larger tumors while <sup>131</sup>I may be more cytotoxic for smaller, micro–metastatic lesions. Since <sup>90</sup>Y does not emit any  $\gamma$  photons, the corresponding <sup>111</sup>In labeled antibodies are generally used as chemical and biological surrogates to study biodistribution and estimate radiation dosimetry of <sup>90</sup>Y labeled antibodies. In addition to <sup>90</sup>Y, the lanthanide radiometal, <sup>177</sup>Lu (max  $\beta^{-} 0.497 \text{ MeV}$ ;  $T_{\frac{1}{2}} = 6.74 \text{ d}$ ) is potentially useful for RIT. Also, <sup>177</sup>Lu has useful gamma photons ( $\gamma = 0.21 \text{ MeV}$ ) for biodistribution and dosimetry studies. Therefore, <sup>177</sup>Lu may have the advantages of both <sup>131</sup>I and <sup>90</sup>Y and may be the most appropriate radioinouclide for therapy. Type of radiation and energy associated with some of the prevalent radioisotopes is summarized in the table hereunder:

Radioisotope	T ½		E. Max	Range Max
-	Hours	Emission	KeV	mm
I-131	193	β-	610	2.9
		γ	364	
Lu-177	162	β-	498	2.0
		γ	208	
Y-90	64	β-	2250	11.0
Bi-213	0.76	α	8400	0.1
		γ	440	
At-211	7.2	α	5870 & 7450	0.055-0.08
		X-	77-92	
Ac-225	240	α		
		γ		
Ra-223	273.6	α	5871	
Th-227	449	α		
		γ		

#### Table 10: Radionuclides for Therapy:

#### **B.** Rationale for Dose–Fractionated regimen:

Pre-clinical models do not always predict toxicity in humans, but based upon the available data, the pre-clinical models combined with animal and human dosimetry data have predicted the toxicity seen with anti-PSMA radioimmunotherapy (i.e. radiolabeled monoclonal antibody) and anti-PSMA radioligand therapy (i.e. radiolabeled peptides).

With radioimmunotherapy, one would predict that the dose-limiting organ is bone marrow, based upon the kinetics / circulation time and exposure of the bone marrow to circulating radiolabeled antibody before it gets into tumors days later. With radioligand therapy, one would predict that there is comparatively very little bone marrow exposure, but there would be more exposure to sites of PSMA expression that are not exposed to (bulky) full-length antibodies. Based upon the available data, this has been born out in human subjects (for radioimmunotherapy) and human patients (for radioligand therapy).

We have conducted a series of overlapping, but sequential phase I and II clinical trials evaluating radiolabeled anti–PSMA monoclonal antibody J591. These studies are summarized immediately below with additional details about dose–fractionation (as requested) following the summary.

- Initial (1st in human) studies of trace-labeled <sup>111</sup>In-DOTA-J591: demonstrated safety and targeting of J591 in men with mCRPC(32, 89, 90)
- Phase I (single-dose) <sup>90</sup>Y-J591: demonstrated safety, targeting, and preliminary efficacy in mCRPC(31)
- Phase I (single-dose) <sup>177</sup>Lu-J591: demonstrated safety, targeting, and preliminary efficacy in

mCRPC(28)

- Phase II (single dose) <sup>177</sup>Lu–J591: demonstrated efficacy in initial 32 subjects, subsequent analysis with significant dose–response and preliminary biomarkers(91)
- Phase I fractionated-dose <sup>177</sup>Lu-J591 dose-escalation study: demonstrated safety, confirming hypothesis that higher cumulative doses could be adminsterd as a split dose(92)
- Phase II (single dose) <sup>177</sup>Lu–J591 expansion phase: confirmed efficacy and dose–response with exploratory biomarkers(39)
- Phase I fractionated-dose <sup>177</sup>Lu-J591 plus docetaxel: demonstrated the ability to safely combine fractionated <sup>177</sup>Lu-J591 with docetaxel 75 mg/m<sup>2</sup> in men with mCRPC(93)
- Phase I fractionated-dose <sup>177</sup>Lu-J591 expansion phase: provided efficacy data, confirmed dose-response, continued analysis of exploratory biomarkers(41)

Taken together, with >250 patients treated on these sequential clinical trials, we have drawn the following conclusions: (i) there is a clear dose-response with more responses and longer survival at higher doses; (ii) there is clear dose-toxicity response in terms of myelosuppression (which is both predictable and reversible); (iii) dose-fractionation allows higher cumulative dose with comparatively less toxicity (including allowing concurrent chemotherapy); and (iv) without pre-selection for PSMA expression, approximately 90% have accurate tumor targeting, though the small minority with low/no PSMA expression by imaging have a lower likelihood of response. While  $^{177}Lu-J591$  mAb demonstrated significant efficacy, the major limitation in the optimization of therapeutic dose is the hematological toxicity (described in greater detail below).

With a single-dose, 70 mCi/m<sup>2</sup> of  $^{177}$ Lu-J591 was the MTD, with dose-limiting myelosuppression in those receiving 75 mCi/m<sup>2</sup> (with 2 of 3 with dose-limiting hematologic toxicity)(28). In the phase II study, this was confirmed with 70 mCi/m<sup>2</sup> of  $^{177}$ Lu-J591 being tolerable with reversible myelosuppression, but with a significant amount of manageable grade 4 myelosuppression (including approximately 40% platelet transfusion)(39).

In radiotherapy, the anti-tumor response has been thought to be primarily due to induction of apoptosis by radiation(94-97). However, the degree of anti-tumor response following the administration of targeted radionuclides depends on several variables, especially total (cumulative) radiation dose to the tumor, dose-rate and tumor radiosensitivity. Single-agent RIT, although potentially useful for slowing solid tumor growth, has not been effective in controlling aggressive tumors, which often have p53 and other mutations and are less susceptible to apoptosis, the apparent mechanism of cell death from low dose-rate radiation(98). Bone marrow is the dose-limiting organ in RIT in the absence of marrow reconstitution(99, 100). **Dose-fractionation** is a practical strategy to decrease the dose to bone marrow while increasing the cumulative radiation dose to the tumor at an optimal dose-rate(97, 101, 102). The main reason for using dose fractionation is to take advantage of the difference between early-responding and late-responding tissues. The radiation effect on early-responding tissue can be reduced by prolonging the treatment time and dose fractionation. The radiation effect on late-responding tissues will not be changed significantly if the total dose is not changed(97, 102). Preclinical data have shown that dose fractionation or multiple low dose treatments can decrease toxicity while increasing the

efficacy(96, 103, 104). Similarly, there is some clinical evidence that bone marrow toxicity can be reduced with some modest increase in the cumulative maximum tolerated dose(105-107).

Based upon these data, we performed two sequential studies of fractionated dose <sup>177</sup>Lu–J591.

As described in in the paragraph above, dose–fractionation of external–beam radiation as well as radioimmunotherapy has been demonstrated to have less toxicity and might be more efficacious by delivering active radiation over a more prolonged period of time to tumor cells. Based upon the half–life of <sup>177</sup>Lu–J591 combined with the kinetics of radiolabeled intact monoclonal antibody, we chose a 2–week interval between <sup>177</sup>Lu–J591 doses:

# *i)* Phase I trial of fractionated dose <sup>177</sup>Lu–J591 in men with metastatic castration–resistant prostate cancer(41, 92)

Men with progressive metastatic CRPC received 2 fractionated doses two weeks apart. Initially, 6 cohorts of 3-6 pts got 2 doses of <sup>177</sup>Lu-J591 2 weeks apart (20 mCi/m2, escalating to 45 mCi/m2 x2). Subsequently, pts enrolled in 2 expansion cohorts at the recommended phase 2 doses (RP2D). Planar<sup>177</sup>Lu–J591 imaging was semi-quantitatively scored. The endpoints were PSA changes and survival (OS); as well as CTC count (CellSearch) changes in the expansion cohorts. 49 patients, with median age 74.1 years (range 55–95), median PSA of 44.9 ng/mL (1.9–766.5); 83.7% with bone, 61.2% with lymph node, 40.8% with visceral metastasis, 8.2% were CALGB (Halabi) low, 34.7% were intermediate, 57.1% were in high-risk group. RP2D's of fractionated <sup>177</sup>Lu-J591 were 40 mCi/m2 x2 or 45 mCi/m2 x2 with option for GCSF. PSA changes for the low dose group were reported as 6.3% showing >50% PSA decline, 12.5% reporting >30% PSA decline, and 37.5% with any PSA decline, within RP2D group 21.2% showing >50% PSA decline, 42.4% reporting >30% PSA decline, and 66.7% with any PSA decline. The median overall survival for low dose group was 14.6 months and for RP2D group was 27.7 months. Accurate targeting of <sup>177</sup>Lu–J591 was seen in 79.6%. Patients with lower PSMA expression by imaging were less likely to respond (p=0.07). Of 25 with CTC counts, 14 declined, 8 stably favorable, and 3 increased. RP2D was associated with more PSA declines (p=0.036) and longer OS (p=0.004), even after controlling for CALGB prognostic grouping (adjusted HR 0.42 [95% CI 0.21, 0.84] p=0.01). Predictable, reversible myelosuppression was seen. 36 (73.5%) patients had grade 3/4 heme toxicities; 19 (57.6%) had Grade 4 heme toxicities in RP2D cohorts with 45.4% receiving prophylactic platelet transfusions (median 1, range 1-4) and 6 GCSF. 14 (28.6%) had infusion reactions (without pre-meds), with 1 patient having Grade 2 infusion reaction leading to withdrawing from the study prior to his 2<sup>nd</sup> dose. 5 (10.2%) had transient Gr 1/2 AST/ALT. This study concluded that fractionated <sup>177</sup>Lu–J591 is well tolerated with predictable, reversible myelosuppression and PSA and CTC declines. Additionally, with dose-fractionation, the cumulative dose MTD is 14–28% higher than single dose MTD with similar toxicity.

	Single Dose		Fractionated Dose		
Cumulative Dose $(mCi/m^2)$	65	70	70 (35 + 35)	80 (40 + 40)	90 $(45 + 45)$
Platelets Grade	27%	56.3%	40%	50%	58.8%
Platelet Transfusion	7%	41%	0%	31.3%	52.9%
Neutropenia Grade 4	0%	37.5%	0%	31.3%	29.4%
Febrile Neutropenia	0%	2.1%	0%	0%	5.8%

Table 11: Toxicities data from Phase 2 single dose <sup>177</sup>Lu–J591 Study and Phase 1 fractionated 177Lu–J591 study:

In addition, fractionated dosing allowed concurrent dosing with myelosuppressive chemotherapy:

# ii) Phase I trial of fractionated dose <sup>177</sup>Lu–J591 plus docetaxel/prednisone in men with metastatic castration–resistant prostate cancer (42):

Following progression on primary hormonal therapy, chemotherapy can offer symptomatic improvement as well as incremental survival benefit. However, responses are transient and all men eventually suffer from progression of disease as described above with single–agent anti–PSMA based radioimmunotherapy. The combination of taxane chemotherapy with radiotherapy has been used in several diseases because of the radiosensitizing effects of taxane–based chemotherapy. In addition to favorable results from fractionated RIT and the radiosensitizing effects of taxane–based chemotherapy will overcome some of the limits imposed by the physical characteristics of <sup>177</sup>Lu. Based upon this theory, a phase I trial of docetaxel and prednisone with escalating doses of fractionated <sup>177</sup>Lu–J591 was initiated. 15 men with median age 69.1 (49.3–80.8) were enrolled. The MTD/RP2D of <sup>177</sup>Lu–J591 was 40 mCi/m<sup>2</sup>x2 doses (delivered with cycle 3 of docetaxel), with 73.3% showing >50% PSA decline, 80.0% reporting >30% PSA decline, and 86.7% with any PSA decline. Predictable, reversible myelosuppression was seen. Even at the highest dose level, no dose limiting toxicity was observed, with short–term / reversible grade 4 neutropenia in 33% and grade 4 thrombocytopenia in 13%.

Aside from the overall rates of dose limiting toxicity, the combined studies have also consistently demonstrated the kinetics of myelosuppression, with nadir neutrophil and platelet counts 28–32 days following the last administered dose of <sup>177</sup>Lu–J591.

With small-molecule radioligand therapy, both the predicted and reported toxicities are much different. One would predict very little myelosuppression, but if it were to occur, it would be
expected to have kinetics similar to (or more quickly than) typical cytotoxic chemotherapy, as the clearance of the myelotoxic drug is quick (minutes to hours rather than days to weeks for antibodies). In fact (with the known caveats of retrospective studies), this is the reported human experience. Essentially no grade 4 neutropenia or thrombocytopenia has been reported within months of exposure to <sup>177</sup>Lu–PSMA radioligands (see details below).

While we have published on the long-term follow up with <sup>177</sup>Lu–J591 with no cases of attributable permanent myelosuppression and/or myelodysplasia/leukemia, results are unknown with <sup>177</sup>Lu–PSMA RLT. The available data with up to 6 months of reported post-treatment follow up are:

1) no attributable grade 4 hematologic (or other) toxicity has been reported at any time (the one case of grade 4 thrombocytopenia is reported with a patient with grade 4 platelet count at pre-treatment baseline)

2) repeat dosing has been performed in many patients at 8-12 weeks after initial dosing and in all cases, white blood cell count and platelet counts were acceptable prior to subsequent doses (up to 5 doses, with dose #5 approximately 40 weeks after dose #1)

The Table below is a summary of the prior summary anti–PSMA RLT Table which includes the severe leukocyte/neutrophil and platelet count rates following <sup>177</sup>Lu–radiolabeled anti–PSMA peptides. In addition to the Table below, a multicenter review of <sup>177</sup>Lu–PSMA–617 data of 82 patients receiving a single dose (mean 5.9 GBq) revealed no grade 4 hematologic or non–hematologic toxicity(108).

Radioligand Therapy	Ν	Dose (GBq), Range	Total doses	Grade 4 neutropenia	Grade 4 platelets	Reference
<sup>177</sup> Lu–PSMA–617	30	3.7 - 6	1-3	0	0*	Kratochwil et al 2015
<sup>177</sup> Lu–PSMA I&T	56	3.6 - 8.7	1-5	0	0	Baum et al 2016
<sup>177</sup> Lu–PSMA–617	9	5.28 – 5.77	1	NR	NR	Hohberg et al 2016
<sup>177</sup> Lu–DKFZ–PSMA–617	5	3.4 - 3.9	2	NR	NR	Delker et al 2015
<sup>177</sup> Lu–DKFZ–PSMA–617	10	4.1-6.1	1	0	0	Ahmadzedehfar et al 2016
$^{177}$ Lu-PSMA-617	7	5.5 - 7.4	1	NR	NR	Das et al 2016
$^{177}$ Lu-PSMA-617	23	7.4	1	NR	NR	Demir et al 2016
<sup>177</sup> Lu–PSMA–617	28	6	1-2	0	0	Rahbar et al 2016
$^{177}$ Lu-PSMA-617	31	1.1-5.5	1-4	0	0	Yadav et al 2016
<sup>177</sup> Lu–PSMA I&T	2	5.7 - 8	1	0	0	Weineisen et al 2015
<sup>177</sup> Lu–PSMA I&T	22	3.4 - 7.4	1-4	0	0	Heck et al 2016
<sup>177</sup> Lu–PSMA–617	43	6	1-4	0	0	Zimbelmann et al 2016
<sup>177</sup> Lu–PSMA–617	5	5.8-7.4	1	0	0	Chakraborty et al 2016

Table 12: Toxicities data following anti–PSMA radioligand therapy:

\*One patient had pre-treatment baseline platelet count of grade 4 NR = not reported

A dose fractionated regimen provides the theoretic benefit of higher cumulative radiation dose to the target tissue/organ while limiting toxicities secondary to non-specific binding of the radiolabeled ligand. In addition to the toxicity data provide above, our group has consistently seen evidence of dose-response (see Table below), justifying a dose-escalation study.

Group	Ν	Any PSA decline	$\geq$ 30% PSA decline	> 50% PSA decline	Median OS mo [95% CI]*
Total	49	55.1%	32.7%	16.3%	22.9 [16.2, 29.7]
Low doses	16	37.5%	12.5%	6.3%	14.6 [9.9, 19.4]
RP2D	33	66.7%	42.4%	21.2%	27.7 [15.8, 39.6]
40 x2 cohort	16	50.0%	25.0%	12.5%	19.6 [9.1, 30.2]
45 x2 cohort	17	87.5%	58.8%	29.4%	48.3 [16.0, 80.6]

Table 13: Dose response from the Phase-I fractionated <sup>177</sup>Lu-J591 study:

#### C. Rationale for combination regimen:

PSMA represents an example of a cellular target that can be co-targeted by 2 different agents that bind to different sites on the PSMA molecule and where each agent has significantly different properties. The crystal structure of PSMA has previously been determined. The binding site of the J591 antibody has been mapped to the apical domain of PSMA, furthest away from the cell membrane (see figure below).



In this exemplary case, the antibody and small molecule (enzymatic inhibitor) have very different properties as shown in the table below.

Table	14:	Differential	characteristics	of	anti-PSMA	antibody	(J591)	and	anti-PSMA
peptid	es (P	PSMA-617):							

Characteristic	Antibody	Peptide
Size (Molecular weight)	150 kd	1 kd
Kinetics (blood half-life)	3–7 <b>days</b>	3-7 hours
Bio-distribution	Tumor, Blood	<b>Tumor,</b> Salivary glands, Kidneys
Side effects	Transient ↓ blood counts	Dry mouth/eyes; ↓ kidney function
Route of elimination from body	Via liver	Via urinary tract

As a result of the differing properties, the predicted side effects caused by the 2 targeting agents and their cytotoxic payloads are virtually mutually exclusive and the actual dose-limiting toxicity of the individual agents in humans is also consistent. This allows co-administration of the 2 targeting agents with their payloads to provide additive payload dose to the tumor without increasing side effects.

Our group has recently conducted both in-vitro and in-vivo experiments to confirm:

1.







_		
2.		



#### Summary:

Taken together, these data show that co–administering 2 agents that bind different sites of a target molecule (e.g., a cell surface receptor molecule) present on a population of cells results in additive binding of those agents. Where the target molecule/receptor is internalized, additive amounts of the 2 agents will, therefore, be internalized. Furthermore, if the 2 targeting agents have different properties (e.g., molecular mass, charge, hydrophilicity/hydrophobicity, pharmacokinetics, bio–distribution, etc.) such that their respective side effects differ, and are non–overlapping, the 2 agents can be co–administered to result in additive binding/uptake by the targeted cells without causing any added toxicity. It is clear across our prior PSMA-targeted radionuclide therapy studies that higher administered doses lead to more responses and longer survival, but also increased toxicity. As a corollary, the dose of each agent can be modestly reduced (by 5–50%) below its

respective maximum tolerated dose whereby the co–administration of the 2 targeting agents still provides an additive dose to the target cells but can substantially or completely reduce the toxicity experienced by the subject.

# 2.8 Risk/Benefit Assessment

This is a phase I clinical trial with the primary endpoints to find DLT and MTD. Based on the prior clinical studies using <sup>177</sup>Lu–J591 by our group the dose–limiting organs/tissue is expected to be red marrow. The expected toxicities can be leukopenia, thrombocytopenia, anemia, fatigue, renal failure, and transaminitis. All these toxicities have been transient and reversible. mCRPC patients treated with RIT using <sup>177</sup>Lu–J591 have shown improved survival compared to historical controls.

Based on findings by retrospective publications, the most common side effects and/or toxicities reported for <sup>177</sup>Lu–PSMA–617 or <sup>68</sup>Ga–PSMA–HBED–CC are fatigue, renal failure, xerostomia, leukopenia, anemia, thrombocytopenia (so far none of the published studies have reported any significant hematological toxicities) and transaminitis. All of these toxicities are reportedly mild and transient. mCRPC patients treated with RLT have higher response rates than any available standard drug in the refractory setting by historical standards.

#### 2.9 Correlative Studies Background

#### i. Archival tissue and cell-free plasma DNA to assess genomic alterations:

We will incorporate tissue-based and circulating tumor DNA (ctDNA) approaches to assess prostate cancer pathway changes including AR signaling pathway activation (mRNA) and genomic alterations including DNA repair pathways and will correlate with PSMA heterogeneity on imaging. We will evaluate the association of molecular biomarkers with imaging and treatment response. Mechanisms of resistance will be assessed in a subset of patients enrolled at WCM through integrative genome-wide analysis of pre-treatment and progression tumor biopsies. At the end of this Aim, we will have defined the molecular landscape of CRPC patients enrolled on prospective trials of PSMA-targeted radionuclide therapy and will have identified biomarkers associated with outcome to be evaluated in a larger study to determine whether they are predictive or prognostic.

PC is a clinically and molecularly heterogeneous disease with marked variability in patient outcomes. Defining the genomic alterations in PC has improved classification of tumors: for instance, 50% harbor TMPRSS2-ERG gene fusion, 10% SPOP mutation, and germline or somatic alterations involving DNA repair pathways (eg., BRCA2, ATM, MMR defects) occur in 20% of CRPC. Based on recent data from our group and others, we predict that these genomic alterations may help identify patients most likely to respond to DNA damaging therapy with PSMA-targeted radionuclide therapy. In PC models, ERG expression results in relative resistance to ionizing radiation. Conversely, SPOP mutant prostate cancers and/or those with CHD1 loss show increased

genomic instability, play a role in double strand DNA break repair, and result in increased sensitivity to DNA damage. Germline or somatic alterations involving DNA damage repair genes (such as BRCA2 and ATM) have also been shown to be preferentially sensitive to DNA damaging therapy including PARP inhibitors and platinum chemotherapy. We therefore hypothesize that distinct molecular phenotypes of prostate cancer may have predictable and exploitable differences in sensitivity to DNA damaging agents such as <sup>177</sup>Lu. We and others have also described prostate cancer subtypes based upon AR signaling and neuroendocrine differentiation. We hypothesize that the NEPC / AR low tumors would express low PSMA and respond less well to PSMA-targeted therapy. We will test these hypotheses using tissue and ctDNA collected from our prospective cohorts.

We will evaluate archival tumor tissue (preferentially from metastatic biopsies) with the plan for utilization of our prostate cancer specific targeted platform covering DNA (for mutation and copy number) of 74 genes/regions commonly aberrant in prostate cancer and 42 control genes. Prostate cancer genes including AR, SPOP, CHD1, DNA repair, ERG fusion, and other CRPC-relevant alterations. We will also collect a single-sample of blood for germline analysis of DNA repair defects. In addition, we will collect plasma samples for ctDNA analysis prior to treatment, at 3-months, and at progression using the prostate cancer specific PCF SELECT platform developed in collaboration with others as part of a PCF Challenge Award. The panel is also being used for other prospective therapeutic trials which will allow for cross-comparison of data.

Gene expression of androgen signaling genes and other relevant pathways will be assessed using a custom panel Nanostring platform developed and validated by Dr. Beltran's lab. This panel includes 350 genes including the AR signaling signature genes (n=30), the AR V7 splice variant, neuroendocrine prostate cancer genes, epithelial mesenchymal transition, cell cycle- genes, TMPRSS2-ERG fusion transcript, and control/housekeeper genes. As discussed above, we will describe findings from this exploratory endpoint in each study with the plan to analyze identified biomarkers to determine their association with efficacy and toxicity across studies, in particular at doses found to have efficacy.

# ii. CTC Count:

Circulating tumor cell (CTC) counts via the FDA-cleared, standard of care CellSearch platform were demonstrated to be prognostic in men with advanced prostate cancer prior to systemic therapy and a "conversion" from an unfavorable count ( $\geq$ 5 CTCs/7.5 mL) was associated with a median similar to those starting with favorable counts, leading to clearance of this particular test by the FDA (109). More recently, the combination of CellSearch CTC enumeration and serum LDH have been demonstrated to have prognostic value, meeting Prentice criteria for survival surrogacy in the setting of abiraterone/prednisone treatment in men with mCRPC previously treated with docetaxel (110)

#### iii. Immune Assays:

We and others have observed some durable responses following radiolabeled-PSMA therapy. Radiation may lead to an induction of or increase in immune responses to locally expressed antigens, but unlike external beam radiation, PSMA-targeting generally delivers radiation to all sites of disease (by imaging) and therefore observing an abscopal response is not possible. In prostate cancer, humoral immune response has already been documented with immunotherapy. *We hypothesize that prostate cancer is immunogenic following PSMA-targeted radiation, and that the serological repertoire in patients may be useful as a pharmacodynamic, prognostic, or even predictive tool.* 

We propose to test the hypothesis that radiolabeled anti-PSMA treatment of advanced prostate tumors leads to induction or increase in immune responses to locally expressed antigens by assessing serological changes pre- and post-treatment. Indeed, one surrogate to test whether these tumors become more immunogenic is to measure naturally occurring antibodies in the serum or plasma of patients. Autologous serotyping, i.e., serum antibody recognizing tumor-derived antigens, have resulted in a growing list of human cancer antigens that are immunogenic in the host of origin. With regard to cancer antigens recognized by the humoral immune system, over 2700 sequences have been identified by SEREX analysis of sera from patients with a wide range of cancers; these antigens include mutational, overexpressed, oncogenic viral, differentiation, and cancer-testis (CT) antigens. These can be rapidly tested in high-throughput ELISA using recombinant antigens (Grand Serology). In some cases, such as the cancer/testis antigen NY-ESO-1, serum antibodies are associated with spontaneous T cell responses measurable from peripheral blood. It should be noted that most of the antigens defined as targets of autoantibodies are intracellular, rather than surface antigens, and reflect the capacity of the immune system to detect abnormalities associated with neoplasia. The prevalent hypothesis is that antibodies are generated as antigens are released from dying tumor cells, either because of spontaneous necrosis or because of external treatments. The presence of antibodies to intracellular targets can be used as a marker of tumor existence or progression, but can also have a functional role to help generating T cell responses through the formation of immune complexes with their cognate antigen. Today, protein microarrays allow for the detection of specific serum antibodies against a very large number of targets simultaneously. Arrays can be used to determine patterns of antigens recognized by autoantibodies during the course of diseases. These tools are also useful to measure changes in antibody responses with treatment such as immunotherapies. The opportunity to define "seromic" changes in these prostate cancers, i.e., the scope of antigens recognized with increased specificity and frequency by post-treatment patient sera vs. baseline, is now within reach, provided adequate analyses of the vast amount of data generated by these microarray can be properly interpreted.

**Grand serology:** A series of known tumor antigens will be assessed in a hypothesis-driven manner for their capacity to elicit autoantibodies in advanced prostate cancer patients, at baseline and after treatment with PSMA-TRT. Using ELISA as previously described, we propose to test a series of 20-25 tumor antigens, including mutational, stem-cell, endogenous retroviral, and cancer-testis antigens such as TP53, NY-ESO-1, GAG-HERV-K, SOX2, PRAME, WT1, MAGE-A3, SSX2, CT45, etc., most of which already have demonstrated spontaneous immunogenicity various solid tumors, including advanced prostate cancer. Briefly, sera will be

tested in 4x serial dilutions, starting from 1/100, for IgG reactivity against full-length E. coli- or yeast-produced proteins of tumor and control antigens to assess specificity. Positive and negative control sera with known reactivity will be used on each plate to validate the assays. Reciprocal titers will be extrapolated and considered significant if >100. In samples with multiple serum collection time points available, significant changes in humoral response will be defined as antigen-specific antibody going from undetectable (titer <100) to detectable (titer >100), or with titers at least 4-fold different between time points. For antigens showing humoral reactivity, mapping of linear epitopes may be assessed using overlapping peptide series covering the sequence of the antigen, to assess polyclonality and potential spreading over time. In a subset of patients, other immunoglobulin isotype or subclass usage may be assessed as well. We propose that systemic tumor antigen-specific antibody responses in plasma or serum may be useful as a marker of immunological response to radiolabeled anti-PSMA treatment in advanced prostate cancer patients, and that increases or induction in immune responses will occur with treatment in comparison to baseline. These experiments are expected to shed light on both mechanisms of action of the drug, including occurrence of antigen spreading, and on potential predictive biomarkers.

**Seromics:** While Grand Serology offers a high-throughput way to quickly assess changes of antibody specificity in serum or plasma sample to known tumor antigens, it is limited by the small number of tumor targets identified to date, especially in prostate cancer. To circumvent this limitation, an alternative is to use a broader array of self-antigens instead as potential targets of immune responses generated by immunotherapy.

Seromics is an exploratory, non-biased, hypothesis-generating platform that allows the testing of thousands of human proteins simultaneously as potential targets of autoantibodies from patient serum or plasma. The method only requires a few microliters of material, and may be customized for applications other than IgG antibody detection. The seromic platform can be used to define biomarkers or sets of antigens present at baseline in specific patient populations, and that could be used as prognostic or predictive markers. It has been previously described that prostate cancer spontaneously elicits humoral responses to a small subset of autoantigens that are not immunogenic in healthy donors. Accordingly, seromics is suited to comprehensively look for serum antibody changes in prostate cancer patients at the antigen-specific level between consecutive time points, before and after treatment with radiolabeled anti-PSMA therapy. We plan to use either ProtoArrays (Invitrogen) or HuProt (CDI), protein microarrays spotted with 9000 to 20000 cancer- and non-cancer associated proteins and controls, to define serum autoantibody changes in a subset of proposed treated patients (pre-post) with favorable outcomes, and compare them to changes in age-matched patients with no response.

**Immunogenic cell death**: we hypothesize that the PSMA TRT will lead to generation of an immune response that will induce immunogenic cell death.

High mobility group box 1 (HMGB1) is an important mediator of immunogenic cell death, and it has been shown that it is released by dying tumor cells following radiation. HMGB1 may promote dendritic cell activation and migration and antigen presentation; following certain anti-neoplastic therapy (such as radiation) higher levels of HMGB1 have been associated with improved

outcomes. We will assess HMBG1 via ELISA in serum obtained pre-and post-treatment with PSMA-TRT and analyze for associations with response, progression-free survival, and overall survival outcomes. We have stored serum (previously collected for HAHA assessment) from prior studies with long-term outcome data and continue to prospectively collect serum from patients being enrolled in current PSMA-TRT studies.

Among the several markers of immunogenic cell death, calreticulin is one of the most investigated. Calreticulin exposure on tumor cell surface is widely accepted as an "eat-me" signal, which is required for phagocytosis of dying tumor cells. Mainly an endoplasmic reticulum (ER)-associated chaperone protein, CTR translocation to the plasma membrane is triggered by reactive oxygen species (ROS)-induced ER stress, dictating the immunogenic cell death. We hypothesize that calreticulin tumor cell expression and cellular localization is associated with tumor response to the immunogenic response-inducing PSMA TRT. Unfortunately, the availability of pre- and post-treatment tumor tissue is often a logistical problem in patients with prostate cancer. We expect to have a subset with pre- and post-treatment tumor tissue available and we will assess membrane expression of CRT in that subset.



#### **3.** SUBJECT SELECTION

**3.1 Study Population** 

Subjects who have documented progressive metastatic CRPC disease, who meet the inclusion and exclusion criteria will be eligible for participation in this study.

#### 3.2 **Inclusion** Criteria

- 1. Histologically or cytologically confirmed adenocarcinoma of prostate
- 2. Documented progressive metastatic CRPC based on Prostate Cancer Working Group 3 (PCWG3) criteria, which includes at least one of the following criteria:
  - i. PSA progression
  - ii. Objective radiographic progression in soft tissue
  - iii. New bone lesions
- 3. ECOG performance status of 0-2
- 4. Have serum testosterone  $\leq$  50 ng/dL. Subjects must continue primary and rogen deprivation with an LHRH/GnRH analogue (agonist/antagonist) if they have not undergone bilateral orchiectomy.
- 5. Have previously been treated with at least one of the following:
  - Androgen receptor signaling inhibitor (such as enzalutamide) ٠
  - CYP 17 inhibitor (such as abiraterone acetate)
- 6. Have previously received taxane chemotherapy, been determined to be ineligible for taxane chemotherapy by their physician, or refused taxane chemotherapy.
- 7. Age > 18 years
- 8. Patients must have normal organ and marrow function as defined below:

•	Absolute neutrophil count	$\geq$ 2,000 cells/mm <sup>3</sup>
•	Hemoglobin	$\geq 9 \text{ g/dL}$
•	Platelet count	$\geq$ 150,000 x 10 <sup>9</sup> / $\mu$ L
•	Serum creatinine	$\leq$ 1.5 x upper limit of normal (ULN) or calculated creatinine clearance $\geq$ 60 mL/min/1.73 m <sup>2</sup> by Cockcroft–Gault

- Serum total bilirubin  $\leq 1.5 \times \text{ULN}$  (unless due to Gilbert's syndrome in which case direct bilirubin must be normal)
- Serum AST and ALT  $\leq 1.5 \times ULN$  in the absence of liver metastases;  $<3 \times ULN$  if due to liver metastases (in both circumstances bilirubin must meet entry criteria)
- 9. Ability to understand and the willingness to sign a written informed consent document.

#### **3.3** Exclusion Criteria

- 1. Implantation of investigational medical device ≤4 weeks of Treatment visit #1 (Day 1) or current enrollment in oncologic investigational drug or device study
- 2. Use of investigational drugs ≤4 weeks or <5 half-lives of Treatment visit # 1(Day 1) or current enrollment in investigational oncology drug or device study
- 3. Prior systemic beta-emitting bone-seeking radioisotopes
- 4. Known active brain metastases or leptomeningeal disease
- 5. History of deep vein thrombosis and/or pulmonary embolus within 1 month of Treatment visit #1
- 6. Other serious illness(es) involving the cardiac, respiratory, CNS, renal, hepatic or hematological organ systems which might preclude completion of this study or interfere with determination of causality of any adverse effects experienced in this study
- 7. Radiation therapy for treatment of PC  $\leq 4$  weeks of Treatment visit #1
- 8. Patients on stable dose of bisphosphonates or denosumab, which have been started no less than 4 weeks prior to treatment start, may continue on this medication, however patients are not allowed to initiate bisphosphonate/Denosumab therapy during the DLT-assessment period of the study.
- 9. Having partners of childbearing potential and not willing to use a method of birth control deemed acceptable by the principle investigator and chairperson during the study and for 1 month after last study drug administration

- 10. Currently active other malignancy other than non-melanoma skin cancer. Patients are considered not to have "currently active" malignancy if they have completed any necessary therapy and are considered by their physician to be at less than 30% risk of relapse.
- 11. Known history of known myelodysplastic syndrome

# 4. OVERVIEW OF STUDY DESIGN AND METHODOLOGY

#### 4.1 Study Design

This is an open–label, single–center Phase I dose–escalation study designed to determine the dose–limiting toxicity (DLT) and the maximum tolerated dose (MTD) of combination of <sup>177</sup>Lu–J591 and <sup>177</sup>Lu–PSMA–617 in a two–week dose–fractionation regimen. <sup>177</sup>Lu–J591 will be given at a moderate dose previously demonstrated to be safe x2 infusions two weeks apart. For <sup>177</sup>Lu–PSMA–617 the dose escalation will start at 3.7 GBq (100 mCi) and escalate in increments of 1.85 GBq (50 mCi) for each dose to a planned maximum of 9.25 GBq (250 mCi) x2 doses, 2 weeks apart. Should there be unacceptable toxicity at the initial dose level, we will de-escalate to dose level -1 (1.85 GBq / 50 mCi per dose). After the phase I study has established a MTD, the Phase II, single-arm trial will start.

Patients must have documented progressive metastatic CRPC disease based on Prostate Cancer Working Group 3 (PCWG3) criteria in order to be eligible for enrollment. Upon meeting the inclusion and exclusion criteria and signing the informed consent and HIPPA form, subjects will undergo the screening. As part of the screening, subjects will get a single dose of <sup>68</sup>Ga–PSMA–HBED–CC and will have a PET/CT. Nuclear Medicine physician(s) will review the PET/CT scans to document PSMA expression at tumor site(s).

Subjects will have Lutetium–177 Planar/SPECT Imaging on Day 8 ( $\pm 1$  day) after the first dose of <sup>177</sup>Lu–J591 + <sup>177</sup>Lu–PSMA–617. Optimal images will be performed on selected consenting subjects between the initial treatment visit #1 on Day 1 and Day 4 and prior to treatment visit #2 on D15  $\pm 1$ . Subjects will be closely monitored for AEs (weekly x2 weeks, then every 2 weeks for one month, at 8 and 12 weeks, and then every 4 weeks for next 3 months).

Upon completion of investigational treatment with dose–fractionation regimen of the combination of <sup>177</sup>Lu–J591 + <sup>177</sup>Lu–PSMA–617, subjects will undergo <sup>68</sup>Ga–PSMA–HBED–CC injection and same day PET/CT at the end of study visit to document treatment response. Subsequently survival data and additional treatment(s) information will be captured from their routine Standard of care (SOC) visits.

 Table 14: Phase I dose-escalation plan for combination of <sup>177</sup>Lu–J591 + <sup>177</sup>Lu–PSMA–617:

<sup>177</sup> Lu–J591 + <sup>177</sup> Lu–PSMA–617: Dose escalation (per dose)									
	<sup>177</sup> Lu–J591*	<sup>177</sup> Lu–PSMA–617							
	Dose: GBq (mCi) / m <sup>2</sup>	Dose: GBq (mCi)							
Cohort									
-1	2.22 (60)	1.85 (50)							
1	2.22 (60)	3.7 (100)							
1a	1.35 (36.5)	3.7 (100)							
2	1.35 (36.5)	5.55 (150)							
3	1.35 (36.5)	7.4 (200)							
4	1.35 (36.5)	9.25 (250)							

\*Note: BSA will be capped at 2 for 177Lu-J591 dosing.

#### **Rationale for dosing amendment:**

Prior studies of 177Lu- and 90Y-J591 have utilized individualize dosing per body surface area (BSA). As a single agent, 177Lu-J591 when delivered in a 2-dose fractionated regimen is suggested to be dosed at 45 mCi/m2 x2 (or 40 mCi/m2 x2with less toxicity and less efficacy). The dose in combination with docetaxel is 40 mCi/m2 x2). In the initial version of the protocol, an estimate for fixed dosing based upon prior safe dose levels and average BSA was utilized. One subject in the initial cohort had prolonged myelosuppression and on analysis, had a low BSA. Therefore based upon the wealth of prior safety data with BSA dosing and the wish to primarily assess one rather than 2 dosing variables, the dosing amendment will revert to BSA dosing. In addition, we will cap BSA at 2 m<sup>2</sup> with the intent of minimizing myelosuppression from the RIT.

#### 4.2 Phase I:

The Phase I portion will use a 3+3 dose-escalation study design, with the planned initial and subsequent cohorts described in Table 14. This study design is well-described and accepted by the clinical and scientific community, so we will not provide complete details here. Briefly, this design is constructed to reduce the chance of escalating the dose when the probability of DLT is high, and increase the chance of escalating the dose when the probability of DLT is low. The MTD is defined as the highest dose level with an observed incidence of DLT in no more than one out of six patients treated at a particular dose level. The dose escalation scheme provides the following probabilities of escalation based on the true chances of DLT at a specific dose level. One can see that the probability of escalation is high if the toxicity risks are low:

True Probability of Toxicity	y: .05	.10	.20	.30	.40	.50	.60
Probability of Escalation:	.97	.91	.71	.49	.31	.17	.08

As there is a reasonably large pool of short- and long-term experience with PSMA-directed <sup>177</sup>Lu, a phase II trial using the MTD to will be launched to obtain preliminary efficacy data.

# 4.3 Phase II:

We will enroll up to 27 subjects in the Phase II trial utilizing a Simon two-stage design (including the 3-6 patients used for establishing the MTD). For the phase II portion, we will define evaluable patients as those who met eligibility requirements, have initiated therapy, and were not removed from the study for non-compliance or withdrawal. Sample size recommendations for the two-stage design are determined according to Simons two-stage minimax design. The first stage will accrue a total of 16 patients (including those who were treated at the MTD in the phase I portion), if 6 or fewer of the first 16 evaluable patients do not experience a 30% decline in PSA, the study will be terminated and declared to have a negative result. If 7 or more patients out of the first 16 evaluable patients experience a 30% decline in PSA, 9 more patients will be accrued in Stage 2 (total sample size of 27). The new regimen will be declared to have activity and be worthy of further testing if 13 or more patients experience a 30% PSA decline among the 27 patients who are part of the phase II trial.

# 5. **REGISTRATION PROCEDURES**

# 5.1 Identification of subjects:

Patients diagnosed with documented progressive metastatic CRPC disease who are visiting Oncology Clinic at NYPH–Cornell Campus for their standard of care visit, will be approached for recruitment for this study. Investigators or delegates under their direct supervision may perform pre–screening of these potential subjects.

# 5.2 Consent process

Potential subjects will have a discussion with the investigator/delegate including the rationale for the study, investigational nature of the protocol and study drug and the voluntary nature of participation, potential risks and benefits, alternatives to participation, and study procedures. Individuals will have the opportunity to read the written informed consent document at their leisure (preferably outside of the clinical area for > 1 day) and the opportunity to have questions answered in a private location with the understanding that should they decide not to participate, they will still be able to receive any available standard of care therapy. Potential subjects will also have the opportunity to obtain the advice of their treating physician. Investigators or delegates under their direct supervision will verify the subject's understanding of the investigational and voluntary nature of the study, the potential risks and benefits, study procedures, and alternatives prior to signing of the written informed consent.

# 5.3 Central Patient Registration

Subjects will be assigned a sequence number for the protocol and will be centrally registered with the Weill Cornell Medicine (WCM), Division of Hematology and Medical Oncology Clinical Research Office with the following documents:

- a. WCM Patient registration form
- b. First and last page of the fully executed informed consent form (including HIPPA), plus additional pages if checkboxes for correlative studies are required.
- c. Eligibility checklist signed and dated by investigator and research nurse
- d. Documentation of any eligibility waivers granted
- e. Entry of screening information into WCM web-based system

Central registration information is reviewed and entered into the HemOnc centralized research database. Documentation of patient registration will be confirmed prior to release of study agent by nuclear medicine.

#### 6. STUDY PROCEDURES

Screening assessments and study procedures outlined in this section can only be performed after obtaining informed consent. All on-study visits and dosing should be scheduled from Day 1 (date of the first infusion) on the study. It is very important that protocol procedures are performed at the time-points stipulated below. When it is not possible to perform, the study visit at the exact time-point, the visit maybe performed within the acceptable visit window as defined in the visit-specific section below.

After obtaining informed consent from enrolled subject(s), screening and study related treatment procedures will be performed as outlined in **Table 15** and described in detail in **Section 6.1**.

# 6.1 Schedule of Evaluations

# Table 15: Schedule of trial events:

	Screening	Treatment Visit 1	Imaging visit	Treatment Visit 2	F/U Visit 1	F/U Visit 2	F/U Visit 3	F/U Visit 4	F/U Visit 5	Scan / Efficacy Visit	Short- Term F/U <sup>k</sup>	Long- Term F/U
Informed Consent	Х											
Demographics	Х											
Medical History	Х	X	Х	х	X	Х		Х		X	Х	
Physical Exam	Х	X	Х	х	X	Х		Х		X	Х	
Performance Status	Х	X	Х	Х	X	Х		Х		X	X	
Vital Signs <sup>a</sup>	Х	X	Х	Х	Х	Х		Х		X	Х	
PRO <sup>1</sup>	Х	X	Х	Х	X	Х		Х		X	x <sup>l</sup>	
CBC with diff, plts <sup>b</sup>	Х	X	Х	x <sup>c</sup>	Х	Х	Х	Х	X	x	Х	
Serum Chemistry <sup>d</sup>	Х	X	Х	X	Х	Х	Х	Х	Х	X	Х	
PSA	Х	X		х		Х		Х		X	Х	
LDH		X		х		Х		Х		X	Х	
Testosterone	Х									X		
CTC Count <sup>e</sup>	Х	x <sup>e</sup>								X		
CTC for Research <sup>f</sup>	Х	xf								X		
Cell-Free DNA Research sample <sup>g</sup>	X	X <sup>g</sup>								x		
Immune research sample <sup>m</sup>	X	x <sup>m</sup>								X		
<sup>68</sup> Ga-PSMA infusion & PET CT	X									X		
Radiographic evaluation <sup>h</sup>	X									X	x <sup>h</sup>	x <sup>h</sup>
<sup>177</sup> Lu-J591 infusion		Х		X								
<sup>177</sup> Lu-PSMA-617 infusion		x		x								

<sup>177</sup> Lu-												
Planar/SPECT			x <sup>i</sup>	$\mathbf{x}^{i}$								
imaging <sup>i</sup>												
Archival Tissu	e x											
Adverse Event		v	v	v	v	v		v		v	v	
Monitoring		А	Λ	Λ	А	Λ		А		А	Λ	
Concurrent	v	v	v	v	v	v		v		v	v	
Medications	Λ	А	Λ	Λ	А	Λ		А		А	Λ	
Survival												v
Assessment <sup>j</sup>												А
a: Vital signs will include height and weight during screening and at least weight thereafter												
b: Any subject	b: Any subject with grade >2 neutropenia or thrombocytopenia at any time point will be followed at least weekly until resolution to grade 1											
c: Any subject with grade $\geq$ 2 neutropenia or thrombocytopenia at Treatment 2 will initially not be eligible to receive the second <sup>177</sup> Lu-J591 / <sup>177</sup> Lu-PSMA-617 infusions. The												
dose will be he	eld until the toxicity	has resolved to	o grade 1 for	a maximum o	f 2 weeks. I	f the toxicity	has not resol	ved after 2	weeks, the d	ose will be di	scontinued.	
d: CMP (with direct bilirubin in subjects with known Gilbert's syndrome)												
e: CTC enume	eration via CellSearc	h methodolog	y may be obt	ained during s	screening or	prior to treat	ment C1D1					
f: CTC collect	tion for research pur	poses may be	performed du	iring screening	g or prior to	treatment C	D1: two 2.39	% sodium ci	trate "light l	olue top" tube	S	
g: Cell-Free D	NA BCT <sup>®</sup> may be	obtained durin	ng screening of	or prior to trea	tment C1D1	and will be	whole blood	collection in	n a tube inte	nded for colle	ction, stabil	ization and
transportation	of cell-free plasma I	DNA. 10 mL to	ube will be u	sed for collect	ing whole b	lood for rese	arch purposes	5				
h: Radiographi	c evaluation will inc	lude bone sca	n, CT/MRI o	f abdomen/pe	lvis, and Ch	est x-ray (CZ	KR waived if	CT/MRI in	cludes chest	). Following n	nandatory re	epeat
imaging with S	Scan/Efficacy visit, c	continued repe	at imaging is	recommended	d approxima	tely 12 week	s until radiog	graphic prog	ression as p	art of standard	l care.	
i: Planar/SPEC	T imaging one week	c after the <sup>177</sup> L	u-PSMA-61	7 1 <sup>st</sup> dose infu	sion in all su	ibjects. Opti	onal imaging	will be per	formed on c	onsenting sub	jects betwee	en D1-D4
and/or prior to	treatment #2 on D15	5.										
j: Survival asse	essment and relevant	t medical histo	ory to be colle	ected until dea	th							
k: Short-term f	follow up to be comp	oleted q4 week	ks following t	he scan visit u	intil 6 montl	hs from the 1	<sup>st</sup> treatment v	isit. Subjec	ts who are p	progression-fre	ee at 6 mont	hs will
continue with	q4-week labs and q1	2-week standa	ard imaging p	er guidelines	(as in h abov	ve).						
1: PRO = patie	nt reported outcomes	s = BPI-SF and	d FACT-P; to	be performed	d at time poi	nts specified	during treatr	nent phase,	then q12 we	eks (at imagir	ng time poin	ts) during
short-term foll	ow up until radiogra	phic progressi	on									
m: serum for in	m: serum for immune assays; 1 sample to be collected at any time during screening prior to treatment #1, then again during scan/efficacy visit											

1

### 6.1.1. Screening Visit

The following procedures must be completed no more than 1 month prior to enrollment and no more than 4 weeks following enrollment.

- Informed Consent
- Demographics
- Medical History
- Previous therapy
- Surgical report will include date and type of surgery +/- lymphadenectomy
- Radiotherapy report will include modality of therapy with prescribed dose and field and dates of therapy
- Previous systemic (hormonal, chemo, other) therapy drugs, doses, dates of therapy
- Complete Physical Exam including height and weight
- Brief Pain Inventory
- Vital Signs
- ECOG Performance Status
- CBC with differential and platelet count
- CMP (with direct bilirubin with known Gilbert's syndrome)
- PSA
- Testosterone
- CTC count via CellSearch methodology (may be collected prior to treatment C1D1, do not need to repeat if has known result within 1 month of C1D1)
- CTC for Research (may be collected prior to treatment C1D1)
- Cell-Free DNA BCT<sup>®</sup> Research blood sample (may be collected prior to treatment C1D1)
- Immune research blood sample (may be collected prior to treatment C1D1)
- CT or MRI (abdomen-pelvis), up to 1 month prior to enrollment
- Bone scan, up to 1 month prior to enrollment
- Any confirmatory tests to assess equivocal results of bone scan should also be completed within a month of enrollment
- CXR, up to 1 month prior to enrollment (CXR waived if CT chest performed)
- Single intravenous dose of <sup>68</sup>Ga-PSMA-HBED-CC (5±2mCi or 185±74MBq) at least 1 week prior to <sup>177</sup>Lu-PSMA-617 infusion
- <sup>68</sup>Ga-PSMA-HBED-CC PET/CT scan to confirm the PSMA expression at the tumor site(s). PET/CT scans will be obtained 1 to 3 hours after the infusion of <sup>68</sup>Ga-PSMA-HBED-CC
- Archival tissue for PSMA expression and genomic studies to identify DNA damage

Screening (except <sup>68</sup>Ga-PSMA-HBED-CC PET/CT scan) and visit #1 may occur on the same day provided results are available, all entry criteria are met, and the subject is registered on the study prior to dosing. In this instance, duplicate procedures do not need to be performed.

# 6.1.1.1. Re-screening

Subjects who are unable to complete the initial screening or are not initially eligible will be permitted undergo repeat screening (with repeat written informed consent).

#### 6.1.2. Treatment Phase

The treatment and early monitoring phase comprises of 8 visits spanning over approximately 12 weeks.

Absolute neutrophil counts, platelet counts, bilirubin, transaminases, and serum creatinine must be performed with results available and within range of eligibility criteria within 1 week prior to treatment initiation on C1D1. Details for each visit are listed below:

# 6.1.2.1 <sup>177</sup>Lu-J591 and <sup>177</sup>Lu-PSMA-617 infusion (Treatment 1 Day 1) and (Treatment 2 Day 15 ±1 day)

The following procedures must be completed on the day of treatment:

- Targeted Physical Examination with vital signs and weight
- Medical History
- Brief Pain Inventory, FACT-P
- ECOG Performance Status
- CBC with differential and platelet count
- CMP (with direct bilirubin with known Gilbert's syndrome)
- PSA
- LDH
- Intravenous dose of <sup>177</sup>Lu-PSMA-617 (Dose based on the Cohort in which the subject is enrolled) and <sup>177</sup>Lu-J591
- Adverse event evaluation
- Concomitant medications
- Concomitant procedures
- Planar/SPECT imaging (on D15 prior to treatment in consenting subjects)

As stated above, Absolute neutrophil counts, platelet counts, bilirubin, transaminases, and renal function assessment must be available within 1 week prior to treatment on D1 and meet eligibility criteria.

For treatment #2 (D15), grade  $\geq$  2 thrombocytopenia or neutropenia within 24 hours of the second dose of <sup>177</sup>Lu-PSMA products (Visit 3) requires a subject's treatment to be held until the toxicity has resolved to least grade 1. If the toxicity has not resolved after 2 weeks of monitoring, the subject's <sup>177</sup>Lu-PSMA product treatment will be permanently discontinued and will be termed dose-limiting toxicity

# 6.1.2.2 Imaging visit (Day 8 ±1) [optional D1, D4, D15 in consenting subjects]

- Planar/SPECT imaging
- Targeted Physical Examination with vital signs and weight
- Medical History
- Brief Pain Inventory, FACT-P
- ECOG Performance Status
- CBC with differential and platelet count
- CMP (with direct bilirubin with known Gilbert's syndrome)
- Adverse event evaluation
- Concomitant medications

# 6.1.2.2 Follow-up visits: Follow up Visit 1 (Day 22 ±3), Follow up Visit 2 (Day 29 ±3), Follow up Visit 4 (Day 57 ±3)

- Targeted Physical Examination with vital signs and weight
- Medical History
- Brief Pain Inventory, FACT-P
- ECOG Performance Status
- CBC with differential and platelet count
- CMP (with direct bilirubin with known Gilbert's syndrome)
- PSA (Follow up visits 2 & 4)
- LDH (Follow up visits 2 & 4)
- Adverse event evaluation
- Concomitant medications

# 6.1.2.3 Follow-up lab visits: Follow up Visit 3 (Day 43 ±3); Follow up Visit 5 (Day 71 ±3)

- CBC with differential and platelet count
- CMP (with direct bilirubin with known Gilbert's syndrome)
- PSA
- LDH

Lab visits may be performed at any CLIA certified lab

6.1.3. Efficacy Evaluation (scan) Visit (Day  $85 \pm 7$ ; note that these study procedures are not expected to occur on a single day)

The following procedures must be completed during the efficacy visit(s):

- Targeted Physical Examination with vital signs and weight
- Medical History
- Brief Pain Inventory, FACT-P
- ECOG Performance Status
- CBC with differential and platelet count
- CMP (with direct bilirubin with known Gilbert's syndrome)
- PSA
- LDH
- Testosterone
- CTC count via CellSearch methodology
- CTC for research
- Cell-Free DNA BCT<sup>®</sup> Research blood sample
- Immune research blood sample
- Same standard radiographic imaging modality as baseline (CT/MRI and bone scan +/- CXR)
- Single intravenous dose of <sup>68</sup>Ga-PSMA-HBED-CC (5±2mCi or 185±74MBq)
- PET/CT scan will be obtained between 1 and 3 hours post-injection of <sup>68</sup>Ga-PSMA-HBED-CC
- Adverse event evaluation
- Concomitant medications
- Concomitant procedures

# 6.1.4 Short term follow up (q4 week visits x3 following Efficacy/Scan visit)

Unless subjects have radiographic progression, withdraw from study, or initiate new treatment, the following procedures must be completed every 4 weeks (+/- 7 days) after the initial efficacy/scan visit:

- Targeted Physical Examination with vital signs and weight
- Interim Medical History
- ECOG Performance Status
- CBC with differential and platelet count
- CMP (with direct bilirubin with known Gilbert's syndrome)
- PSA
- LDH

These q4-week lab/clinic visits may be performed with any licensed provider at any CLIA certified lab, provided that the results are available/provided to the Investigator within allowable windows. Visits associated with repeat CT and bone scan (q12 weeks until radiographic progression) should be performed at the WCM investigative site.

# 6.1.5 Long Term Follow Up

Following cancer progression, survival assessment and relevant medical history to be collected until death. Information may be collected from external providers. Information will be updated at least every 6 months following disease progression.

#### 6.2 Treatment Administration

Treatment will be administered on an outpatient basis. Reported adverse events and potential risks are described in **Section 13**.

#### 6.2.1 Agent Administration

Treatment will be administered only to eligible subjects under the supervision of the investigator or identified co-investigator(s). Treatment will be administered on an outpatient basis. Reported adverse events and potential risks are described in **Section 13**. Appropriate dose modifications/delays of the study drug are described in **Section 7**. No investigational or commercial agents or therapies other than those described below may be administered with the intent to treat the subject's malignancy.

#### 6.2.2. Study drug preparation

#### A. <sup>177</sup>Lu–J591

The humanized monoclonal antibody-huJ591 is provided by Dr. Neil Bander. Radionuclide conjugation will be done at the Division of Nuclear Medicine, Department of Radiology, Weill Cornell Medicine.

The DOTA–J591 antibody molecule is labeled with radiometal <sup>177</sup>Lu in 1.0 M ammonium acetate buffer at pH 7.0. The radiolabeled J591 complex is then purified using size–exclusion column chromatography. To avoid the effects of autoradiolysis on the antibody, reaction time is minimized and the radiolabeled antibody preparations are formulated in phosphate buffered saline containing not more than 1% Human Serum Albumin (HSA). All reagents used in the conjugation and purification of J591 are made with pyrogen–free water. Radiolabeled J591 is periodically tested for sterility and pyrogenicity.

Each patient dose will be supplied to the study site on the day of or day before treatment by the central radiopharmacy. The J591 mAb is a deimmunized monoclonal antibody directed at the

extracellular domain of human PSMA. For additional details, see Section 1. Radiolabeled antibody preparations are formulated in phosphate buffered saline containing not more than 1% Human Serum Albumin (HSA).In order to radiolabel J591 with  $\beta^-$  emitting radionuclides, the antibody molecule is first conjugated to a macrocyclic chelating agent, 1,4,7,10– tetraazacyclododecane– N,N',N'','-tetraacetic acid (DOTA) by direct coupling of one of the four carboxylic acid groups of DOTA to the primary amines present in the protein structure. The radiolabeling of J591 with <sup>177</sup>Lu is achieved by adding the radionuclide to the ammonium acetate buffered J591. To avoid the effects of autoradiolysis on the antibody, reaction time is minimized and the reaction mixture is purified with a size exclusion column prior to administration. All reagents used in the conjugation and purification of J591 are made from pyrogen–free water. Radiolabeled J591 is tested periodically for sterility and pyrogenicity. Labeling efficiency and radiochemical purity will be determined using ITLC. Immunoreactivity of radiolabeled J591 mAb preparations will be determined using PSMA+ LNCaP tumor cells.

The material produced are subjected to quality assurance testing as outlined in the Food and Drug Administration "Points to Consider". All production data and supporting documents, together with the results of all quality assurance testing, have been provided to and reviewed by the Food and Drug Administration (<sup>177</sup>Lu–J591, IND# **11613**).

# B. <sup>177</sup>Lu–PSMA–617 and <sup>68Ga</sup>–PSMA–HBED–CC

ABX Advanced Biochemical Compounds, Radeberg, Germany produces both peptides/ligands, PSMA–617 and PSMA–HBED–CC, with PSMA-617 licensed to Endocyte (West Lafayette, LA). Upon purchase and shipment to Weill Cornell Medicine, they will be labeled with either <sup>177</sup>Lu or <sup>68</sup>Ga with final product being <sup>177</sup>Lu–PSMA–617 and <sup>68</sup>Ga–PSMA–HBED–CC, respectively. Radionuclide conjugation will be done at the Division of Nuclear Medicine, Department of Radiology, Weill Cornell Medicine.

The material produced are subjected to quality assurance testing as outlined in the Food and Drug Administration "Points to Consider". All production data and supporting documents, together with the results of all quality assurance testing, have been provided to and reviewed by the Food and Drug Administration (<sup>177</sup>Lu–PSMA–617, IND # 131966 and <sup>68</sup>Ga–PSMA–HBED–CC, IND# 124495).

# 6.2.3 Route of Administration

<sup>177</sup>Lu–J591, <sup>177</sup>Lu–PSMA–617 and <sup>68</sup>Ga–PSMA–HBED–CC will be administered intravenously.

# 6.2.4 Dose levels for <sup>177</sup>Lu–J591 + <sup>177</sup>Lu–PSMA–617

Subject(s) enrollment will be done as 3+3 study design at each dose level. Initially at least 3 subject(s) will be enrolled in Cohort 1 and will receive <sup>177</sup>Lu–PSMA–617 (2 GBq or 54 mCi x2 doses 2 weeks apart) with dose escalation as described above in **Section 4.1 (Table 14)**.

Simultaneously, all subjects will receive in combination a fixed dose of <sup>177</sup>Lu–J591 (1.35 GBq/m2 or 36.5 mCi/m2 capped at BSA of 2) x2 doses, 2 weeks apart. The enrollment ceiling of the dose escalation portion of the study is up to 30 treated study participants (up to five groups, up to 6 at each dose level). Each patient would receive two doses of both investigational agents (<sup>177</sup>Lu–J591 and <sup>177</sup>Lu–PSMA–617) in combination 2 weeks apart. The total dose received by a subject will vary from 4 GBq/m<sup>2</sup> (108 mCi/m<sup>2</sup>) to 20 GBq/m<sup>2</sup> (540 mCi/m<sup>2</sup>) for <sup>177</sup>Lu–PSMA–617. The total dose received by a subject for <sup>177</sup>Lu–J591 will be 2.7 GBq/m2 (73 mCi/m2), with BSA capped at 2.

# 6.2.5 Dose levels for <sup>68</sup>Ga–PSMA–HBED–CC

All subjected enrolled will receive  $5 \pm 2mCi$  or  $185 \pm 74MBq$  of  ${}^{68}Ga-PSMA-HBED-CC$  during the screening visit and scan/efficacy visit.

#### 6.2.6 Study drug premedication

No pre-medication is required, but is recommended prior to J591 dosing. Individual subjects may receive pre-medication at the discretion of their treating physician provided that they are not prohibited medications per study. The recommended pre-medication for <sup>177</sup>Lu–J591 is a H1 blocker (recommended 25–50 mg diphenhydramine) and acetaminophen (recommended 325–650 mg). No pre-medications will be administered for <sup>68</sup>Ga–PSMA–HBED–CC.

#### 6.2.7 Study drug administration

Intravenous access must be well established prior to initiating infusion. At the time of dosing, the IV line will be connected to an infusion container containing the prepared volume of <sup>177</sup>Lu–J591, <sup>177</sup>Lu–PSMA–617 or <sup>68</sup>Ga–PSMA–HBED–CC. <sup>177</sup>Lu–J591 will be infused at a rate of no faster than 5 mg/minute. <sup>177</sup>Lu–PSMA–617 and <sup>68</sup>Ga–PSMA–HBED–CC may be administered as slow IV push.

# 6.2.8. Monitoring Vital signs pre/post <sup>177</sup>Lu–J591 + <sup>177</sup>Lu–PSMA–617 and <sup>68</sup>Ga–PSMA–HBED–CC

The infusion of <sup>177</sup>Lu–J591, <sup>177</sup>Lu–PSMA–617 and <sup>68</sup>Ga–PSMA–HBED–CC and subsequent monitoring will occur in a facility that is equipped for cardio–pulmonary resuscitation. The dispensed dose will be infused under the supervision of nuclear medicine physician or designee under the supervision of a nuclear medicine physician. Infusion–related reactions (fever, rigors) will be treated with acetaminophen, meperidine and diphenhydramine hydrochloride as clinically appropriate. Other allergic events will be managed as follows: rash, pruritus, urticaria and wheezing will be treated with diphenhydramine hydrochloride, meperidine and/or steroids as clinically appropriate. Anaphylaxis or anaphylactoid signs or symptoms will be treated with steroids and/or epinephrine as clinically indicated. Vital signs will be monitored during the infusion. Systolic and diastolic blood pressure (mm Hg), temperature, pulse rate (beats/minute), and respiratory rate (breaths/minute), will be recorded with the patient in sitting position. Any

clinically significant change in the vital signs will be recorded as AEs. Serial vital signs including temperature, BP, and heart rate will be monitored within 30 minutes before the infusion, within 30 minutes and at 60 minutes ( $\pm 10$  minutes) [additionally, at 90  $\pm 10$  minutes and 120  $\pm 10$  minutes for <sup>177</sup>Lu–J591 + <sup>177</sup>Lu–PSMA–617] after the infusion. If any subject has any adverse reaction at 60 minutes for <sup>68</sup>Ga–PSMA–HBED–CC and at 120 minutes for <sup>177</sup>Lu–J591 + <sup>177</sup>Lu–PSMA–617, they will stay longer until it is resolved.

#### 6.2.9. Imaging Plan

#### A. <sup>68</sup>Ga-PSMA-HBED-CC PET/CT Scan

Patient preparation should be according to the policies and procedures of the local imaging site (CBIC). The patient does not need to be fasting for either the infusion or the scans. The use of intravenous or oral contrast will not be permitted. Specifications for acquiring the <sup>68</sup>Ga–PSMA–HBED–CC PET/CT scans will be provided in study specific documentation by the study chair or the co–investigators from the Division of Nuclear Medicine. PET/CT should be obtained during the screening visit as well as at the end of study visit(efficacy/scan visit). The images are acquired between 1 and 3 hours after the <sup>68</sup>Ga–PSMA–HBED–CC infusion. Image acquisition will be from vertex of skull to mid thighs.

#### B. Lutetium-177 Planar/SPECT imaging

Patient preparation should be according to the policies and procedures of the local imaging site. The patient does not need to be fasting for either the infusion or the scans. The use of intravenous or oral contrast will not be permitted. Specifications for acquiring the <sup>177</sup>Lu–PSMA–617 planar/SPECT images will be provided in study specific documentation by the study chair or the co–investigators from the Division of Nuclear Medicine. Planar/SPECT images should be obtained during the Imaging Visit (Day 8 ±1) on all subjects after the <sup>177</sup>Lu–J591 + <sup>177</sup>Lu–PSMA–617 infusion. Image acquisition will be from vertex of skull to mid thighs. Optional images will be performed on selected consenting subjects between initial Treatment Visit (between D1 – D4) and prior to Treatment Visit 2 (Day 15 ±1).

#### 6.2.10 Managing toxicity

NCI CTCAE version 4.0 is used to grade all adverse events.

Grade ≥ 2 thrombocytopenia or neutropenia within 24 hours of the second dose of <sup>177</sup>Lu–J591 + <sup>177</sup>Lu–PSMA–617 (Treatment Visit #2) requires a subject's treatment to be held until the toxicity has resolved to least grade 1. If the toxicity has not resolved after 2 weeks of monitoring, the subject's <sup>177</sup>Lu–J591 + <sup>177</sup>Lu–PSMA–617 treatment will be permanently discontinued and will be termed dose–limiting toxicity.

### Dose-limiting toxicity (DLT) is defined as:

- Grade 4 hematologic toxicity that is deemed at least possibly related to <sup>177</sup>Lu–J591 and/or <sup>177</sup>Lu–PSMA–617 will be termed dose–limiting toxicity
- Any grade > 2 non-hematologic toxicity deemed to be at least possibly related to <sup>177</sup>Lu–J591 and/or <sup>177</sup>Lu–PSMA–617 will require a subject's <sup>177</sup>Lu–J591 + <sup>177</sup>Lu–PSMA–617 treatment to be permanently discontinued and will be termed dose–limiting toxicity.
  - A 1-grade increase in a pre-existing from grade 2 to grade 3 will only be considered a DLT if probably or definitely related to study drug

# Maximum Tolerated Dose (MTD) is defined as:

• The dose that produces an "acceptable" level of toxicity or that, if exceeded, would put subjects at "unacceptable" risk for toxicity. Definition of the MTD usually relies on the sample, as MTD is defined as the dose level at which no more than two patients out of six experienced dose–limiting toxicity (DLT).

<u>Note:</u> Toxicities as described above will be considered DLT if they are at least possibly related to  $^{177}$ Lu–J591,  $^{177}$ Lu–PSMA–617 or  $^{68}$ Ga–PSMA–HBED–CC as judged by the investigator (see note above re: 1-grade increases in which case only probably/definitely related toxicities will be deemed DLT). Attribution will be reviewed by the study chair and discussed with the medical monitor if there are questions about severity or attribution.

#### 6.3 General Concomitant Medication and Supportive Care Guidelines

All medications that are administered during the study must be recorded in the patient's CRF and in the source documents. Concomitant medications for other medical conditions are permitted as clinically indicated subject to approval by the study chair.

Subjects will be advised to use contraceptive precautions to avoid pregnancy during the treatment phase since the effects of investigational agents on sperms and embryos are unknown.

#### 6.4 Duration of Therapy and Criteria for Removal from Study

Duration of treatment portion of the study (excluding screening time) will be approximately 85 Days ( $\pm$ 7 days). By signing the informed consent form and agreeing to participate in this study, the subjects are required to participate in entirety up to the completion of all scheduled visits and study procedures, or until one of the following criteria applies:

- A protocol violation occurs
- Disease progression occurs

- A serious or intolerable adverse event occurs (that in the opinion of the Investigator, requires the subject's discontinuation)
- The Investigator withdraws the subject (at the Investigator's discretion for reasons other than the adverse event)
- The Principle Investigator terminates the protocol
- The subject requests to be discontinued from the protocol
- The subject is lost to follow–up
- Intercurrent illness that prevents administration of <sup>177</sup>Lu–J591 or <sup>177</sup>Lu–PSMA–617
- Previous anaphylactic reaction to any J591 product or PSMA peptides/ligands

The investigators or physicians may stop the protocol or terminate a subject's participation in the protocol at any time should they judge:

- That it is not in the subject's best interest to continue
- If the subject experiences a protocol-related injury
- If the subject needs life-saving medications/procedures/treatment
- If the subject does not comply with the study plan
- General or specific changes in the patient's condition render the patient unacceptable for further treatment in the judgment of the investigator

They may also remove the subject from the study for various other administrative and medical reasons. They can do this without the patient's consent.

#### 6.5 **Duration of Follow Up**

Patients will be followed until death for survival assessment. Patients removed from study for unacceptable adverse events will be followed until resolution or stabilization of the adverse event.

# 7. DOSING DELAYS/DOSE MODIFICATIONS

There will be no dosing modifications. Unless there are specific reasons not to do so, all patients who are eligible for the trial will receive  $5 \pm 2 \text{ mCi} (185 \pm 74 \text{ MBq})$  of  ${}^{68}\text{Ga}\text{-PSMA}\text{-HBED}\text{-CC}$  during the screening visit and the Efficacy/Scan visit. During the treatment phase, the subject will receive  ${}^{177}\text{Lu}\text{-J591} + {}^{177}\text{Lu}\text{-PSMA}\text{-}617 \text{ x } 2$  doses, 2-weeks apart based on the dose-level/Cohort to which he is assigned. The study chair must clear any dosing delays due to logistical issues (e.g. subject scheduling and/or radionuclide shipping that fall outside of the treatment window).

#### 8. PHARMACEUTICAL INFORMATION

A list of the adverse events and potential risks associated with *Investigational Agent* can be found in **Section 13**.

#### 8.1. Investigational Agent

#### A. <sup>177</sup>Lu–J591:

The humanized monoclonal antibody-huJ591 is provided by Dr. Neil Bander. Radionuclide conjugation will be done at the Division of Nuclear Medicine, Department of Radiology, Weill Cornell Medicine.

The DOTA–J591 antibody molecule is labeled with radiometal <sup>177</sup>Lu in 1.0 M ammonium acetate buffer at pH 7.0. The radiolabeled J591 complex is then purified using size–exclusion column chromatography. To avoid the effects of autoradiolysis on the antibody, reaction time is minimized and the radiolabeled antibody preparations are formulated in phosphate buffered saline containing not more than 1% Human Serum Albumin (HSA). All reagents used in the conjugation and purification of J591 are made with pyrogen–free water. Radiolabeled J591 is periodically tested for sterility and pyrogenicity.

Each patient dose will be supplied to the study site on the day of or day before treatment by the central radiopharmacy. The J591 mAb is a deimmunized monoclonal antibody directed at the extracellular domain of human PSMA. For additional details, see Section 1. Radiolabeled antibody preparations are formulated in phosphate buffered saline containing not more than 1% Human Serum Albumin (HSA). In order to radiolabel J591 with  $\beta^-$  emitting radionuclides, the antibody molecule is first conjugated to a macrocyclic chelating agent, 1,4,7,10–tetraazacyclododecane– N,N',N'',N'''-tetraacetic acid (DOTA) by direct coupling of one of the four carboxylic acid groups of DOTA to the primary amines present in the protein structure. The radiolabeling of J591 with <sup>177</sup>Lu is achieved by adding the radionuclide to the ammonium acetate buffered J591. To avoid the effects of autoradiolysis on the antibody, reaction time is minimized and the reaction mixture is purified with a size exclusion column prior to administration. All reagents used in the conjugation and purification of J591 are made from pyrogen–free water. Radiolabeled J591 is tested periodically for sterility and pyrogenicity. Labeling efficiency and radiochemical purity will be determined using ITLC. Immunoreactivity of radiolabeled J591 mAb preparations will be determined using PSMA+ LNCaP tumor cells.

The material produced are subjected to quality assurance testing as outlined in the Food and Drug Administration "Points to Consider". All production data and supporting documents, together with the results of all quality assurance testing, have been provided to and reviewed by the Food and Drug Administration (<sup>177</sup>Lu–J591, IND# **11613**).

# B. <sup>177</sup>Lu-PSMA-617 and <sup>68</sup>Ga-PSMA-HBED-CC:

PSMA-617 will be obtained from Endocyte (West Lafayette, LA) and PSMA-HBED-CC will be obtained from ABX Advanced Biochemical Compounds, Radeberg, Germany and will be shipped to and stored at WCM as per manufacturer's guidelines. Upon subject's enrollment and confirmation of date of infusions, these peptides will be labeled with Lutetium-177 or Gallium-68, respectively.

The material produced are subjected to quality assurance testing as outlined in the Food and Drug Administration "Points to Consider". All production data and supporting documents, together with the results of all quality assurance testing, have been provided to and reviewed by the Food and Drug Administration (<sup>177</sup>Lu–PSMA–617, IND # 131966 and <sup>68</sup>Ga–PSMA–HBED–CC, IND# 124495).

#### 8.2. Availability

<sup>177</sup>Lu–J591 (IND# 11613), <sup>177</sup>Lu–PSMA–617 (IND# 131966) and <sup>68</sup>Ga–PSMA–HBED–CC (IND# 124495) are investigational agents supplied to investigators by Weill Cornell Medicine.

#### 8.3. Agent Ordering

Upon enrollment of the subject, the WCM Study Coordinator will be notified about the <sup>177</sup>Lu–J591, <sup>177</sup>Lu–PSMA–617 and <sup>68</sup>Ga–PSMA–HBED–CC infusion dates. The WCM Study Coordinator will arrange with the staff of Nuclear Medicine Division at WCM for the timely labeling and delivery of the radiolabeled PSMA peptides to the site of infusion. The dates for study visits will be confirmed with the subject, site of infusion, and site of imaging.

#### 8.4. Agent Accountability

The investigator, or a responsible party designated by the investigator, will maintain a careful record of the inventory and disposition of all agents received from Sponsor on a Drug Accountability Record Form (DARF).

# 9. CORRELATIVE/SPECIAL STUDIES

# 9.1 Laboratory Correlative Studies

9.1.1 CTC immune analysis:

Whole blood samples (2.7 mL in each vial) in two 4.5 mL 2.3% sodium citrate tubes (light blue top tube) will be collected for research analysis at specified time points (Screening/baseline and efficacy visit). These samples are collected for research purposes only and will not be billed to the subject. CTC will be stained for calreticulin analyzed for research purposes only.

9.1.2 Cell–Free Plasma DNA:

Peripheral blood will be collected in two 10 mL Cell–Free DNA BCT<sup>®</sup> tubes (Streck) at the specified time points (screening or pre-treatment and efficacy visit). These samples are collected for research purposes only and will not be billed to the subject. The cell–free plasma DNA will be analyzed for genomic studies and to identify DNA repair and AR-pathway alterations.

#### 9.1.3 Immune Assay:

Serum samples will be obtained during screening prior to treatment and at the main efficacy visit for immune biomarkers and processed as per the lab manual. Results of these assays will not be billed to the patient and are for research purposes only.

#### 9.1.4 Archival tissue:

Archival tissue will be requested during the screening visit. Fifteen unstained slides containing tumor material from archival paraffin–embedded tissue should be obtained. If available, metastatic tissue is preferred to prostate biopsy/prostatectomy specimens. The tissue will be analyzed for PSMA expression and DNA damage repair pathways for research purposes only.

#### 9.2. Special Studies

Conventional Imaging studies as well as <sup>68</sup>Ga–PSMA–HBED–CC PET/CT imaging and Lutetium–177 Planar/SPECT imaging will be performed as described in schedule as well as clinically indicated to assess disease response. In subjects enrolled in this study, the measurable disease response will be calculated using Response Evaluation Criteria in Solid Tumors (RECIST Version 1.1) with PCWG3 modifications.

#### **10. MEASUREMENT OF EFFECT**

#### **10.1 CTC count response**

All subjects in this study will get blood samples drawn (at screening and EOS visit) for CTC enumeration by CellSearch methodology. Our primary analysis will assess those whose CTC counts drop to less than 5 or stay below 5 (responders) vs. those who remain at least 5 or above (non-responders) at EOS visit.

In addition, it appears that decreases in CTC counts with therapy are a favorable marker even if the count remains at least 5. We will also analyze % changes in CTC counts with at least 50% decline in CTC count from baseline considered a response at 12 weeks. Best response will also be analyzed (% increase or best % decrease at any point will also be reported in a waterfall plot).

As reported in phase III studies of men with mCRPC, a favorable CTC count and LDH level at 12 weeks has been associated with overall survival. We will report the proportion of subjects who have CTC count <5 and normal LDH at the efficacy (scan) visit time point.

# **10.2** Biochemical (PSA) response

PSA response will be determined by comparing the PSA levels after therapy to the baseline, pre-treatment PSA. Declines of  $\geq$  30% confirmed by a second PSA value  $\geq$  2 weeks later, will be reported. Subjects must not demonstrate clinical or radiographic (CT and/or MR) evidence of disease progression during this time period.

#### **10.3 Duration of PSA response**

Duration of PSA response is defined as the time from the first 25% PSA decline until the PSA value is confirmed to increase by 25% above the nadir, provided that the increase is at least 2 ng/mL above the nadir.

#### **10.4 Duration of CTC response**

Duration of CTC response is defined from the first CTC count drop to <5 (or  $\le50\%$  of baseline) to the time CTC count increases to  $\ge5$  (or  $\ge25\%$  increase from nadir).

#### **10.5 PSA Progression**

PSA progression will be defined as a rise of > 25% above either the pretreatment level or the nadir PSA level (whichever is lowest). PSA must increase by > 2 ng/ml to be considered progression. Confirmation requires a second consecutive rising PSA at least 2 weeks apart.

#### **10.6 PSA Stabilization**

PSA stabilization is referred to as any set of PSA values that do not meet the criteria for PSA response or PSA progression.

#### **10.7** Time to PSA Progression

Time to PSA progression is defined as the interval between initiating treatment until the PSA rises 25% above nadir provided that the increase is at least 2 ng/mL.

#### **10.8** Change in lesion size

In subjects with measurable disease, complete response (CR) is defined as complete disappearance of all measurable and evaluable lesions by physical examination or imaging studies and normalization of PSA with no appearance of new lesions for > 1 month. Partial response (PR) is defined as a 30% or greater reduction in the sum longest uni–dimensional diameter of all measurable lesions. There may be no new lesions. Stable Disease (SD) is characterized by subjects who do not meet the criteria of PR and who are without signs of progressive disease for at least 1 month. Disease Progression (DP) is defined as a greater than 20% increase in the sum longest uni–dimensional diameters of the indicator lesions or the appearance of new lesions. Bone scan progression (evaluable disease only) is requires at least 2 new lesions seen on a scan subsequent to baseline followed by a repeat scan at least 6 weeks later with at least one new additional lesion.

Conventional Imaging studies (MRI, CT, Bone Scan) along with optional but recommended investigational images (<sup>68</sup>Ga–PSMA–HBED–CC PET/CT imaging and Lutetium–177 Planar/SPECT imaging) will be performed during the study visits or as clinically indicated to assess disease response. In subjects enrolled in this study, the measurable disease response will be calculated using Response Evaluation Criteria in Solid Tumors (RECIST Version 1.1) with PCWG3 modifications.

Radiographic scans will be used to assess best overall response and radiographic progression based on modified RECIST criteria for soft-tissue lesions and protocol-specific criteria for bone lesions Baseline images should be taken during Screening as close as possible to, and never more than 28 days before treatment visit #1. Every effort must be made to ensure the same radiographic method is used before and after treatment at scheduled visits. Radiographic progression free survival will be evaluated based on these results.

#### Analysis of Efficacy:

Median bPFS, rPFS, and OS, including survival curves, will be estimated using Kaplan–Meier methodology. Greenwood's formula will be used to calculate 95% confidence intervals for the Kaplan–Meier estimates. Percent change in PSA from baseline will be described by mean/median and standard deviation/inter–quartile range, as appropriate, depending on the distribution of percent change from baseline. Modified RECIST response (i.e., CR, PR, and CR/PR proportions), CTC count response proportion, defined favorable LDH/CTC proportion, and associated 95% confidence intervals, will be estimated via binomial proportions.

All subjects will be offered optional but recommended baseline and subsequent <sup>68</sup>Ga–PSMA–HBED–CC PET/CT, and those treated with <sup>177</sup>Lu–labeled products will undergo Planar/SPECT imaging. We will analyze the data for associations with response to treatment as well as survival by chi–square tests/Fisher's exact tests and log–rank tests, respectively. For pre/post imaging comparisons, McNemar's chi–square test and paired t–tests/Wilcoxon signed–rank tests will be used, as appropriate. We plan to report data for each individual study as well as across the studies.

#### **10.9** Calreticulin Expression on CTCs (immune measure)

We will analyze for associations between calreticulin expression as a categorical variable (membrane vs cytoplasmic vs mixed vs no expression) and response to treatment and survival using chi–square tests/Fisher's exact tests and log–rank tests, respectively. We plan to report data for each individual study as well as across PSMA studies.

#### **10.10** Patient reported outcomes

Initial bone-targeted  $\beta$ -emitters were approved for the control of painful bone metastases. Radium-223 dichloride appears to be associated with improved/preserved patient reported outcomes as well. We will assess pain prior to and following treatment with the brief pain inventory (and will also collect pain medication data). We will assess global and prostate cancer specific patient reported outcomes with BPI-SF and FACT-P questionnaire. As with the other correlative studies, we will plan to report results for this individual study as well as across current and prior studies, particularly examining subjects that receive what we believe are efficacious doses.

# 11. DATA REPORTING / REGULATORY CONSIDERATIONS

#### 11.1. Data Collection

The data collection plan for this study is to utilize REDCap to capture all treatment, toxicity, efficacy, and adverse event data for all enrolled patients.

#### 11.1.1. REDCap

REDCap (Research Electronic Data Capture) is a free data management software system that is fully supported by the Weill–Cornell Medical Center CTSC. It is a tool for the creation of customized, secure data management systems that include Web–based data–entry forms, reporting tools, and a full array of security features including user and group based privileges, authentication using institution LDAP system, with a full audit trail of data manipulation and export procedures. REDCap is maintained on CTSC–owned servers that are backed up nightly and support encrypted (SSL–based) connections. Nationally, the software is developed, enhanced and supported through a multi–institutional consortium led by the Vanderbilt University CTSA.

#### 11.2. Regulatory Considerations

All protocol amendments and consent form modifications will be made by the Principal Investigator. Should an external sponsor be identified, they will have the opportunity to review
and approve the changes prior to submission of these changes to the local IRB and distribution to participating sites.

## **12. STATISTICAL CONSIDERATIONS**

#### 12.1 Phase I Study Design/Endpoints

The phase I portion is a study of subjects with documented progressive metastatic CRPC with the primary endpoint of determination of MTD (or recommended phase II dose). In general, if there are sufficient numbers of subjects, descriptive statistics (e.g., number of observations, means, standard deviations, medians, and ranges) will be used to summarize data and selected endpoints may be summarized by dosing regimen. Otherwise, subject listings will be provided. No formal hypothesis testing is planned.

The dose-escalation schedule (using 3+3 modified Fibonacci escalation), definitions of DLT, and determination of MTD are defined above (Section 6.2.10). The design is constructed to reduce the chance of escalating the dose when the probability of DLT is high, and increase the chance of escalating the dose when the probability of DLT is low. The maximum tolerated dose is defined as the highest dose level with an observed incidence of DLT in no more than one out of six patients treated at a particular dose level. The dose escalation scheme provides the following probabilities of escalation based on the true chances of DLT at a specific dose level. One can see that the probability of escalation is high if the toxicity risks are low.

True Probability of Toxicity	0.05	0.10	0.20	0.30	0.40	0.50	0.60
Probability of Escalation	0.97	0.91	0.71	0.49	0.31	0.17	0.08

#### 12.2. Phase II Study Design/Endpoints

We will enroll up to 27 subjects into the Phase II portion (including the 3-6 used for establishing the MTD). We will define evaluable patients as patients who met eligibility requirements, have initiated therapy, and were not removed from the study for non-compliance or patient withdrawal.

Sample size recommendations for the two-stage design are determined according to Simon's two-stage minimax design. We project a 30% PSA decline proportion of 35%, below which the regimen will be unacceptable and a 30% PSA decline proportion of 60%, above which the regimen will be considered worthy of further exploration. The null hypothesis that the 30% PSA decline proportion is less than or equal to 35% will be tested against the alternative hypothesis that the 30% PSA decline proportion is greater than or equal to 60%.

The sample size computations were performed assuming a 0.10 one-sided level of significance and 90% power assuming a historical control value for PSA decline of 35% and the alternative

hypothesis of 60%. If 6 or fewer of the first 16 evaluable patients do not experience a 30% decline in PSA (stage 1), the study will be terminated and declared to have a negative result. If 7 or more patients out of the first 16 evaluable patients experience a 30% decline in PSA, ongoing accrual will proceed to the target sample size of 27 patients (stage 2). The new regimen will be declared active in this patient population and worthy of further testing if 13 or more patients experience a 30% PSA decline among the 27 patients entered. This two-stage design yields a 0.90 probability of a positive result if the true 30% PSA decline proportion is 35%. An exact 95% binomial confidence interval will be constructed for the proportion of patients with PSA decline.

## 12.3 Sample Size/Accrual Rate

The planned sample size for this dose escalation Phase I study is between 6 and 24 total patients and the accrual for phase II will be between 16 and 27 patients and will include patients treated at the MTD from the phase I portion. The accrual rate is anticipated to be 1-2 subjects every month on average (with expected pauses in enrollment between dose-escalation cohorts and between the stage 1 and stage 2 of the phase II trial).

## **12.4** Stratification Factors

This is not a randomized trial and so there are no stratification factors.

#### 12.5 Analysis of Endpoints

#### 12.5.1 Analysis of Primary Endpoints

The primary endpoint will be the proportion of subjects with DLT from Treatment Visit #1 through End of Study (Efficacy/scan) Visit. The MTD is the highest dose amongst the different dose–level cohorts in this study at which no more than 1 out of 6 subjects in a cohort experience DLT.

#### 12.5.2 Analysis of Secondary Endpoints

For PSA response, CTC response and imaging response, descriptive analysis will be done. For imaging response RECIST (Version 1.1) criteria with PCWG3 modifications will be applied. Biomarker and patient reported outcome data will be analyzed within this study as well as across additional radiolabeled–PSMA studies.

#### 12.6 Interim Analysis

No interim analysis is planned for the phase I portion (though safety analyses will occur in real-time and prior to each dose-escalation). For the phase II portion, accrual will be suspended

after 16 patients have been enrolled in stage 1. If at least 7 patients out of the 16 patients experienced a PSA decline > 30%, 9 more patients will be accrued as part of stage 2.

## **12.6** Reporting and Exclusions

## **12.6.1** Evaluation of toxicity

All subjects will be evaluable for toxicity from the time of their first infusion with  ${}^{68}$ Ga-PSMA-HBED-CC as well as when they receive their first treatment with  ${}^{177}$ Lu-J591 +  ${}^{177}$ Lu-PSMA-617 and up to 30 days after there last treatment. The distributions of the maximum observed grade toxicity will be tabulated for each type of toxicity and presented by dose level and overall. Results will be summarized with descriptive statistics.

#### **12.6.2** Evaluation of response

Subjects who complete <sup>177</sup>Lu–J591 + <sup>177</sup>Lu–PSMA–617 treatment combination (2 doses of each investigational agents) and at least 12 weeks of subsequent follow–up evaluations (as described in the Schedule Calendar–**Section 6.1**) will be considered evaluable (consistent with PCWG3 guidelines). As the timing of response is not known, should the anticipated time to response be determined to be earlier than with other therapies using alternative markers of response, we may not replace subjects who are not evaluable through week 12 on study.

# **13.** ADVERSE EVENT REPORTING REQUIREMENTS

Adverse event (AE) monitoring and reporting is a routine part of every clinical trial. The investigator will be required to provide appropriate information concerning any findings that suggest significant hazards, contraindications, side effects, or precautions pertinent to the safe use of the drug or device under investigation. Safety will be monitored by evaluation of adverse events reported by patients or observed by investigators or research staff, as well as by other investigations such as clinical laboratory tests, x-rays, electrocardiographs, etc.

#### **13.1** Adverse Event Definition

An adverse event (also referred to as an adverse experience) can be any unfavorable and unintended sign (e.g., an abnormal laboratory finding), symptom, or disease temporally associated with the use of a drug, and does not imply any judgment about causality. An adverse event can arise with any use of the drug (e.g., off–label use, use in combination with another drug) and with any route of administration, formulation, or dose, including an overdose.

# 13.1.1. Investigational Agent Risks

There are no known contraindications for <sup>177</sup>Lu–J591, <sup>177</sup>Lu–PSMA–617 or <sup>68</sup>Ga–PSMA–HBED–CC. Because of the potential for infusion or allergic reaction, the subject should be monitored for safety for one to two hours following the infusion.

Based on prior studies, the known side effects, risks, and hazards associated with the administration of <sup>177</sup>Lu–J591, <sup>177</sup>Lu–PSMA–617 or <sup>68</sup>Ga–PSMA–HBED–CC include: infusion reaction (fever, chills, rash, hypotension, and/or hypertension after injection), xerostomia, renal insufficiency, hematological/bone marrow toxicity (thrombocytopenia, neutropenia) and transient hepatic enzyme elevations. In addition, allergic reactions including anaphylaxis, hypotension, hypertension, difficulty breathing, rash, human anti–humanized antibody (HAHA) response and CNS toxicity are a rare possibility.

#### **Precautions/monitoring:**

The infusion of <sup>177</sup>Lu–J591, <sup>177</sup>Lu–PSMA–617 or <sup>68</sup>Ga–PSMA–HBED–CC and subsequent monitoring will occur in a facility that is equipped for cardio–pulmonary resuscitation. The dispensed dose will be infused under the supervision of nuclear medicine physician or designee under the supervision of a nuclear medicine physician. Infusion–related reactions (fever, rigors) will be treated with acetaminophen, meperidine and diphenhydramine hydrochloride as clinically appropriate. Vital signs will be monitored before/after the <sup>177</sup>Lu–J591 + <sup>177</sup>Lu–PSMA–617 or <sup>68</sup>Ga–PSMA–HBED–CC infusion. Systolic and diastolic blood pressure (mm Hg), temperature, pulse rate (beats/minute), and respiratory rate (breaths/minute), will be recorded with the patient in sitting position. Any clinically significant change in the vital signs will be recorded as AEs. Serial vital signs including temperature, BP, and heart rate will be monitored within 30 minutes of the infusion, and within 30 minutes for <sup>177</sup>Lu–J591 + 177Lu–PSMA–617). If any subject has any adverse reaction at 60 minutes for <sup>68</sup>Ga–PSMA–HBED–CC and 120 minutes for <sup>177</sup>Lu–J591 + 177Lu–PSMA–617, they will stay longer until it is resolved.

#### 13.1.2. Adverse Event Characteristics and Related Attributions

**CTCAE term (AE description) and grade:** The descriptions and grading scales found in the revised NCI Common Terminology Criteria for Adverse Events (CTCAE) version 4.0 will be utilized for AE reporting. A copy of the CTCAE version 4.0 can be downloaded from the CTEP web site (<u>http://ctep.cancer.gov</u>).

#### Attribution of the AE:

**Definite** – The AE *is clearly related* to the study treatment. A reaction that follows a plausible temporal sequence from administration of the study drug and follows a known response pattern to the suspected study drug. The reaction can be confirmed with a positive re–challenge test or supporting laboratory data.

**Probable** – The AE *is likely related* to the study treatment. A reaction that follows a plausible temporal sequence from administration of the study drug and follows a known response pattern to the suspected study drug. The reaction cannot be reasonably explained by the known characteristics of the patient's clinical state or other modes of therapy administered to the patient.

**Possible** – The AE *may be related* to the study treatment. A reaction that follows a plausible temporal sequence from administration of the study drug and follows a known response pattern to the suspected study drug. The reaction might have been produced by the patient's clinical state or other modes of therapy administered to the patient.

**Unlikely** – The AE *is doubtfully related* to the study treatment. The current state of knowledge indicates that a relationship is unlikely.

**Unrelated** – The AE *is clearly NOT related* to the study treatment. No relationship between the experience and the administration of study drug; related to other etiologies such as concomitant medications or patient's clinical state.

#### **13.1.3. Recording of Adverse Events**

All adverse events will be recorded on a patient specific AE log. The AE log will be maintained by the research staff and kept in the patient's research chart.

# 13.1.4. Reporting of AE to WCM IRB

All AEs occurring on this study will be reported to the IRB according to the IRB policy, which can be accessed via the following link:

http://researchintegrity.weill.cornell.edu/forms and policies/forms/Immediate Reporting Policy .pdf

#### **13.2.** Definition of SAE

SAE's include death, life threatening adverse experiences, hospitalization or prolongation of hospitalization, disability or incapacitation, overdose, congenital anomalies and any other serious events that may jeopardize the subject or require medical or surgical intervention to prevent one of the outcomes listed in this definition.

# 13.2.1. Reporting of SAE to IRB

All SAEs occurring on this study will be reported to the IRB according to the IRB policy, which can be accessed via the following link:

http://researchintegrity.weill.cornell.edu/forms\_and\_policies/forms/Immediate\_Reporting\_Policy .pdf

## 13.2.2. Reporting of SAE to FDA

If an SAE occurs on this study, the event will be filed on a MedWatch form with the FDA. The investigator must notify the FDA of any SAE's as soon as possible but no later than 7 calendar days after the initial receipt of the information

Food and Drug Administration Center for Drug Evaluation and Research Central Document Room 5901–B Ammendale Road Beltsville, MD 20705–1266

#### 13.3. AE/SAE Follow Up

All SAEs and AEs reported during this study will be followed until resolution or until the investigator confirms that the AE/SAE has stabilized and no more follow–up is required. This requirement indicates that follow–up may be required for some events after the patient discontinues participation from the study.

#### 14. DATA AND SAFETY MONITORING PLAN (DSMP)

This study will utilize the Weill Cornell Medicine (WCM) Institutional Data Safety Monitoring Board (DSMB) and follow its policies and procedures for monitoring this study for safety concerns, with ongoing updates from the Study Chair on a continuous basis.

The Weill Cornell's DSMB is comprised of medical specialists and advisors on human rights issues in human subjects research. The DSMB currently has 9 members, meets at quarterly intervals during the year, and carries out ongoing review of protocols submitted throughout the year. Once a protocol has been submitted and approved by the Institutional Review Board (IRB) and is recommended for oversight by the DSMB, the Board determines if the protocol will be reviewed quarterly, semi–annually, or annually.

The DSMB evaluates the accumulated data from the study in order to monitor the safety of subjects throughout the trial and reviews the risks and benefits, as well as the efficacy, of the study. The DSMB will also evaluate the overall trial conduct and progress. Ultimately, the DSMB validates the continuation of the trial or determines if a study needs modification or termination.

Reports to the DSMB will include the following items for review:

- Completed DSMB Periodic Review Form.
- Synopsis of the study to date.
- IRB approved consent form.
- IRB current protocol.
- Summary table of study results.
- Adverse event table.
- Data safety monitoring plan.

Safety monitoring is carried out to ensure and maintain the scientific integrity of human subject research projects and to protect the safety of human subjects. Safety monitoring can be viewed as any process during a clinical trial that involves the review of accumulated outcome data for groups of patient–subjects to determine if any of the treatment procedures practiced should be altered or stopped. NIH Guidelines (1998, 2000) specify that all clinical trials should have a system in place for appropriate oversight and monitoring to ensure the safety of participants and the validity of the data.

Monitoring activities will be commensurate with the nature, size, and complexity of the trial in accordance with institutional policies and will be determined after IRB and DSMB review of the protocol immediately prior to study activation. For a small, single–center study, usually a statistician in conjunction with a Safety Officer performs the monitoring. For that single–site, high–risk trials, a DSMB may be appropriate. For larger, single or multi–site studies, a committee, often called a Data Safety Monitoring Board (DSMB), usually performs the monitoring. Ongoing review of the data by an independent individual or committee assures the investigators, the IRB, the study's sponsor, and the funding agency that the trial can continue without jeopardizing subjects' safety.

Weill Cornell Medicine requires that all research approved by the WCMC IRB include an appropriate plan for the monitoring of data to ensure the safety of human subjects. Research supported by Federal agencies will be monitored according to all regulations and guidelines of the relevant Federal agency.

For this study, the DSMB will be notified after each cohort has been completed prior to dose escalation to the next cohort and prior to transition from phase I to phase II. In addition, a report will be made to the DSMB every 6 months.

#### 14.1 Medical Monitor

The medical monitor is required to review all unanticipated problems involving risk to subjects or others, serious adverse events and all subject deaths associated with the protocol and provide an unbiased written report of the event. At a minimum, the medical monitor must comment on the outcomes of the event or problem and in case of a serious adverse event or death, comment on the relationship to participation in the study. The medical monitor must also indicate whether he/she concurs with the details of the report provided by the principal investigator. Reports for events

determined by either the investigator or medical monitor to be possibly or definitely related to participation and reports of events resulting in death must be promptly forwarded to the appropriate committees/agencies. This individual will be a qualified physician, other than the principal Investigator, not associated with this particular study, able to provide medical care to research subjects for conditions that may arise during the conduct of this study, and will monitor the subjects during the conduct of the study.

Sarah Rutherford, MD will serve as the Medical Monitor for this study.

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Performance Status Criteria						
ECOG Performance Status Scale		Karnofsky Performance Scale				
Grade	Descriptions	Percent	Description			
0 Normal activity. Fully active, able to carry on all pre–disease performance without restriction.	Normal activity. Fully active, able to	100	Normal, no complaints, no evidence of disease.			
	without restriction.	90	Able to carry on normal activity; minor signs or symptoms of disease.			
Symptoms, but ambulatory. Restricted physically strenuous activity, but ambulatory and able to carry out work a light or sedentary nature (e.g., light housework, office work).	Symptoms, but ambulatory. Restricted in physically strenuous activity, but	80	Normal activity with effort; some signs or symptoms of disease.			
	a light or sedentary nature (e.g., light housework, office work).	70	Cares for self, unable to carry on normal activity or to do active work.			
2 In bed <50% of the time. Ambulatory and capable of all self–care, but unable to carry out any work activities. Up and about more than 50% of waking hours.	60	Requires occasional assistance, but is able to care for most of his/her needs.				
	about more than 50% of waking hours.	50	Requires considerable assistance and frequent medical care.			
In bed >50% of the limited self-care, co more than 50% of v	In bed >50% of the time. Capable of only	40	Disabled, requires special care and assistance.			
	more than 50% of waking hours.	30	Severely disabled, hospitalization indicated. Death not imminent.			
4	100% bedridden. Completely disabled. Cannot carry on any self-care. Totally	20	Very sick, hospitalization indicated. Death not imminent.			

# APPENDIX A

# Protocol #1802018988 v2 Version Date: 02/19/2019

	confined to bed or chair.	10	Moribund, fatal processes progressing rapidly.
5	Dead.	0	Dead.

# **APPENDIX B**

WCMC IRB SAE Reporting Forms

http://researchintegrity.weill.cornell.edu/institutional review board/irb adv.html