

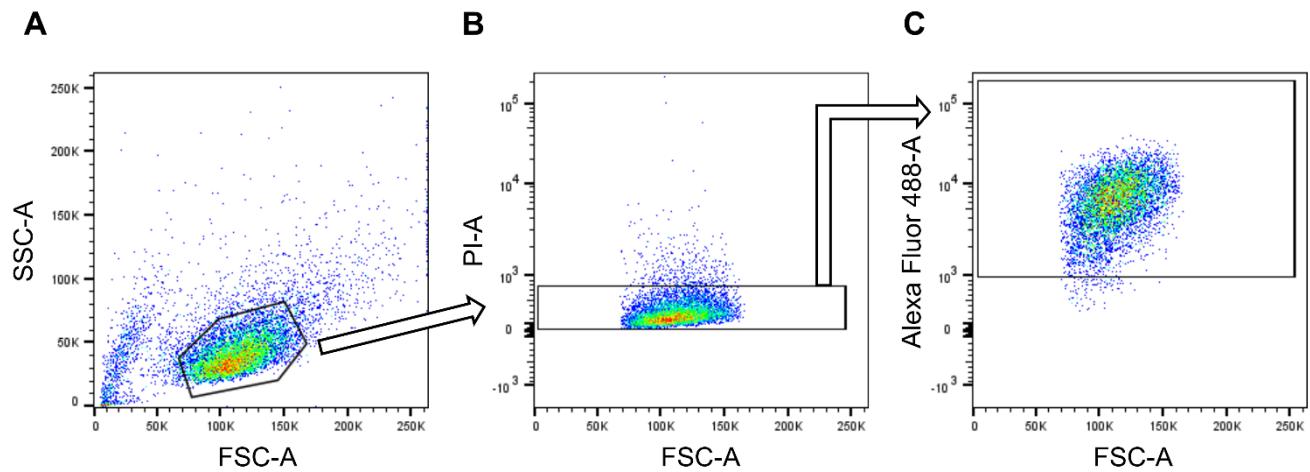
## **Supplementary Material**

### **Exploring the role of combined external beam radiotherapy and targeted radioligand therapy with [<sup>177</sup>Lu]Lu-PSMA-617 for prostate cancer – from bench to bedside**

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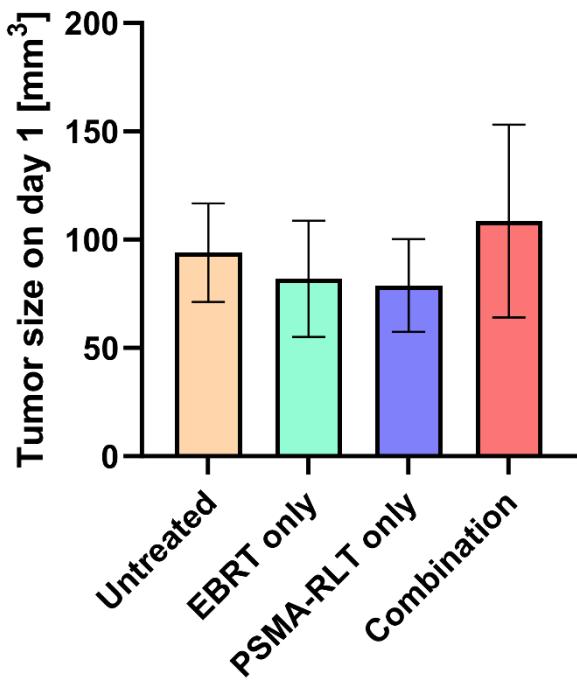
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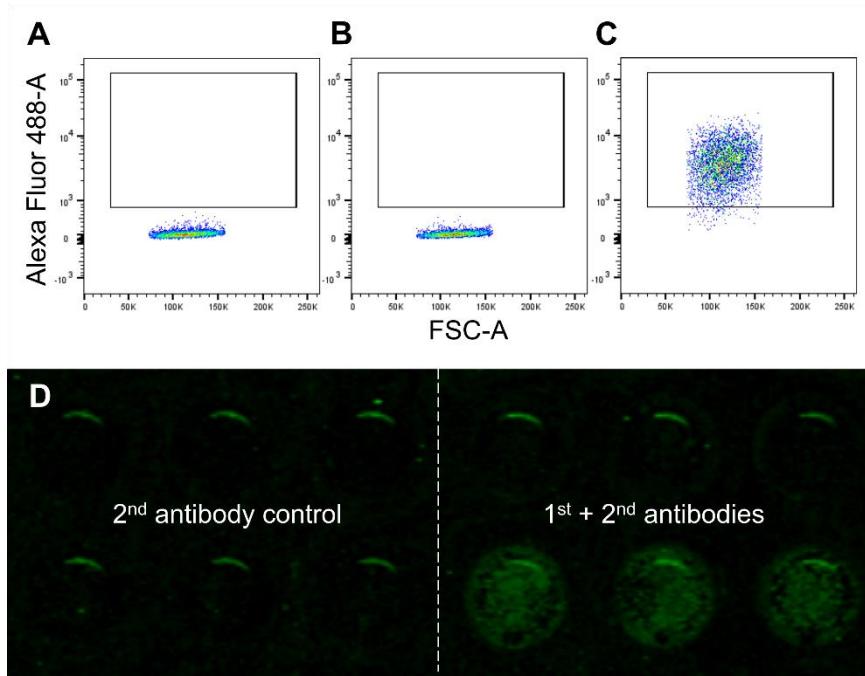
**Figure S1:** Gating strategy in the flow cytometry experiments. The main cell population was selected (A), dead cells were excluded by PI-staining (B) and the median fluorescence intensity in the Alexa Fluor™ 488 channel was analyzed (C).

Abbreviations: Alexa Fluor 488-A: Alexa Fluor 488 – pulse area; FSC: forward scatter – pulse area; PI: propidium iodide; SSC: side scatter – pulse area.

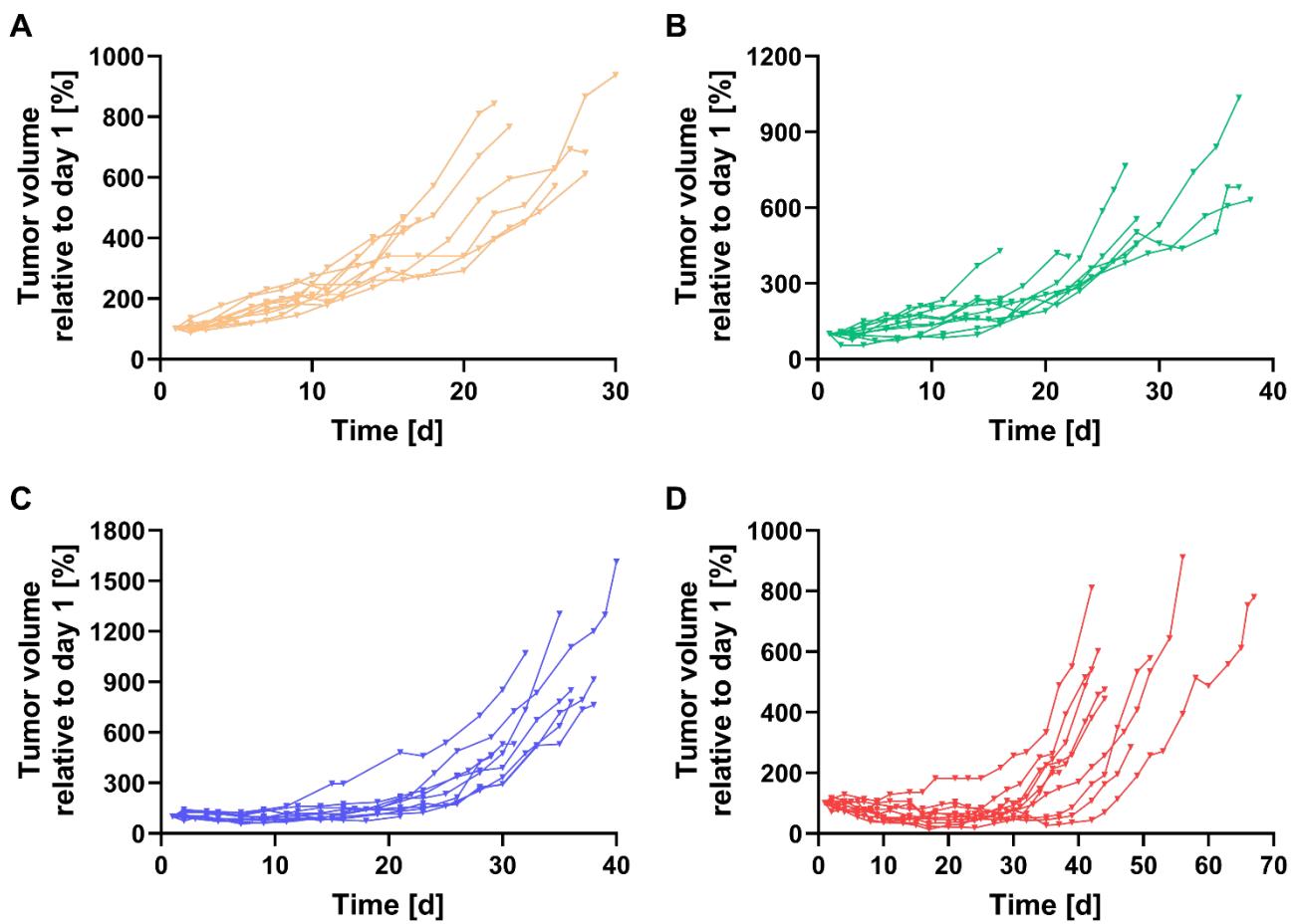


**Figure S2:** Mean tumor size with SD on the day of treatment (= day 1) for each group. There were no significant differences between groups (one-way ANOVA, p = 0.1410).

Abbreviations: ANOVA: analysis of variance; EBRT: external beam radiotherapy; PSMA-RLT: PSMA-targeted radioligand therapy; SD: standard deviation.



**Figure S3:** Negative controls for the analysis of PSMA protein levels. **A-C:** Controls for flow cytometry. Signal intensities in the Alexa Fluor™ 488 channel (Alexa Fluor 488-A) were measured for unstained cells (**A**), IgG1 isotype control-stained cells (**B**), and cells stained with the PSMA-reactive antibody (**C**). **D:** Negative controls for OCW included LNCaP cells stained only with the secondary antibody (bottom row, left) and PC-3 cells (top row) with both the primary and secondary antibodies. No signals were detected as compared to LNCaP cells stained with both antibodies (bottom row, right). Abbreviations: Alexa Fluor 488-A: Alexa Fluor 488 – pulse area; FSC: forward scatter – pulse area; IgG1: immunoglobulin G1; OCW: On-Cell Western Assay; PSMA: prostate-specific membrane antigen.



**Figure S4:** Time courses of individual tumor growth in each treatment group. All animals from the untreated (**A**), EBRT (**B**), PSMA-RLT (**C**) and combination groups (**D**) are displayed. Please note the differences in scaling of the x- and y-axes.

Abbreviations: EBRT: external beam radiotherapy; PSMA-RLT: PSMA-targeted radioligand therapy.

## Supplementary Tables

**Table S1:** Primer sequences and properties.

Target	Sequence	GC [%]	T <sub>m</sub>	Amplicon	Spanning
			[°C]	size [bp]	two exons?
<b>FOLH1</b>	Forward:	48	57.9	281	Yes
	AGTGCTCCCTTTGATTGTCG				
<b>GAPDH</b>	Reverse:	55	62.1		
	GCCTGTCTGGTAACCCTAATGG				
<b>GAPDH</b>	Forward:	45	58.4	101	No
	ACAACTTGGTATCGTGGAAGG				
	Reverse:	58	58.8		
	GCCATCACGCCACAGTTTC				

Abbreviations: bp: base pairs; FOLH1: folate hydrolase 1; GAPDH: glyceraldehyde-3-phosphate dehydrogenase; GC: guanine/cytosine content; T<sub>m</sub>: melting temperature.

**Table S2:** Changes in PSMA protein levels after irradiation (IR) measured by flow cytometry or OCW (mean values from all experiments (n = 3-5) and p-values).

Flow cytometry					
	<i>Difference to 0 Gy control [%]</i>				
<i>Comparison</i>	<i>1 h post-IR</i>	<i>4 h post-IR</i>	<i>8 h post-IR</i>	<i>24 h post-IR</i>	<i>48 h post-IR</i>
0 vs. 0.5 Gy	-3.22	+13.15	+5.93	-3.37	-
	-5.31	+28.06	+3.69	-0.71	-
	-2.86	-11.34	+7.98	-3.10	-
	-	+11.05	-	-	-
	-	-	-	-	-
0 vs. 2 Gy	+0.35	+9.06	-22.67	-22.99	-21.22
	+6.31	+19.91	-15.34	-22.28	+5.79
	+2.37	+23.60	-9.12	-6.11	-19.67
0 vs. 4 Gy	-7.99	+13.75	+13.09	-2.90	-
	-5.81	+12.00	+10.60	-19.49	-
	-4.00	+39.45	+1.27	-15.35	-
	-	+17.43	-	-	-
	-	+9.87	-	-	-
0 vs. 8 Gy	-6.82	+11.18	+12.42	-13.49	-
	-3.21	+11.75	+7.78	-21.71	-
	-1.93	+30.72	-12.73	-21.35	-
	-	+21.12	-	-	-
	-	+17.86	-	-	-
<i>p-values</i>					
0 vs. 0.5 Gy	0.6775, ns	0.4766, ns	0.9346, ns	> 0.9999, ns	-
0 vs. 2 Gy	> 0.9999, ns	0.1137, ns	> 0.9999, ns	0.0180, *	0.2500, ns <sup>□</sup>
0 vs. 4 Gy	0.1116, ns	0.0500, *	0.3965, ns	0.2676, ns	-
0 vs. 8 Gy	0.6775, ns	0.0244, *	> 0.9999, ns	0.0412, *	-

OCW			
	<i>Difference to 0 Gy control [%]</i>		
<i>Comparison</i>	<i>1 h post-IR</i>	<i>4 h post-IR</i>	<i>24 h post-IR</i>
0 vs. 2 Gy	-15.42	+29.22	-3.52
	+2.98	+12.16	-33.48
	+3.55	-0.61	-29.42
0 vs. 4 Gy	-1.28	+14.86	-6.30
	+9.33	-10.02	-25.86
	-10.84	+49.85	+1.39
		+28.05	
<i>p-values</i>			
0 vs. 2 Gy	> 0.9999, ns	0.9177, ns	0.0974, ns
0 vs. 4 Gy	> 0.9999, ns	0.4601, ns	0.8968, ns

Statistical analysis for each time point: Kruskal-Wallis test and Dunn's multiple comparisons test.

<sup>a</sup>Wilcoxon signed-rank test (two-tailed).

ns = not significant, \* = p < 0.05.

Abbreviations: IR: irradiation; OCW: On-Cell Western Assay; PSMA: prostate-specific membrane antigen.

**Table S3:** Changes in PSMA fold gene expression after irradiation (IR) measured by RT-qPCR (mean values from all experiments (n = 3) and p-values).

RT-qPCR			
	<i>Fold gene expression</i>		
<i>Time point</i>	<i>0 vs. 0.5 Gy</i>	<i>0 vs. 4 Gy</i>	<i>0 vs. 8 Gy</i>
4 h post-IR	1.39	1.50	2.32
	1.39	1.14	3.71
	1.63	1.24	1.88
<i>p-values</i>			
4 h post-IR	0.2028, ns	0.6268, ns	0.0061, **

Statistical analysis: Kruskal-Wallis test and Dunn's multiple comparisons test.

ns = not significant, \*\* = p < 0.01.

Abbreviations: IR: irradiation; PSMA: prostate-specific membrane antigen; RT-qPCR: reverse transcription quantitative polymerase chain reaction.

**Table S4:** Mean tumor volumes [%] by day normalized to the day of treatment = day 1. Shown are mean values  $\pm SD$ .

Day	Untreated	EBRT only	PSMA-RLT only	Combination
1	100.00 (0)	100.00 (0)	100.00 (0)	100.00 (0)
2	106.77 (16.99)	94.75 (19.58)	114.52 (21.76)	104.76 (17.77)
3	109.65 (10.42)	95.65 (16.31)	85.71	83.83 (6.83)
4	133.91 (26.20)	109.36 (36.61)	117.15 (13.77)	95.65 (22.75)
5	116.67	73.33	95.27 (24.51)	104.00
6	163.18 (38.31)	141.48 (26.39)		77.04 (22.41)
7	174.27 (35.26)	118.55 (41.58)	93.93 (21.75)	84.18 (21.07)
8	186.21 (35.50)	158.36 (34.71)	85.71	60.19 (20.99)
9	195.51 (37.29)	138.19 (52.59)	99.20 (31.00)	83.23 (21.42)
10	234.30 (32.35)	169.46 (40.59)	79.12	44.44 (9.28)
11	222.60 (48.21)	147.45 (59.02)	110.93 (38.39)	85.80 (33.45)
12	206.88 (9.73)	190.35 (40.74)	109.93 (31.41)	86.67
13	296.87 (46.29)	181.22 (27.55)		41.77 (8.00)
14	316.58 (66.01)	203.56 (99.87)	111.24 (32.17)	87.71 (32.71)
15	316.67 (33.00)	189.33 (31.19)	159.07 (118.40)	39.26 (10.53)
16	387.25 (90.00)	218.63 (102.58)	139.70 (66.13)	76.64 (30.89)
17	356.27 (94.69)	205.89 (33.47)	144.23	29.93 (14.96)
18	444.54 (144.16)	217.88 (53.57)	119.95 (39.02)	86.77 (54.41)
19	392.86	240.00	141.68 (26.29)	96.00
20	315.96 (34.00)	223.19 (45.65)		34.77 (14.10)
21	592.23 (191.27)	289.73 (79.52)	192.57 (115.37)	91.18 (49.29)
22	573.19 (238.06)	317.26 (76.28)	230.77	36.76 (14.59)
23	598.61 (166.68)	313.73 (58.18)	219.64 (107.05)	86.23 (49.78)
24	478.33 (40.07)	342.01 (24.70)	356.04	42.71 (18.82)

25	486.11	448.83 (122.24)	311.91 (198.86)	86.99 (54.74)
26	599.86 (40.60)	528.92 (199.73)	288.65 (123.17)	68.57
27	693.33	516.73 (214.80)	371.43	66.70 (34.21)
28	719.58 (132.08)	492.20 (47.11)	384.36 (144.63)	90.26 (64.25)
29		418.18	496.90 (64.52)	90.69 (44.87)
30	937.50	493.96 (52.06)	451.61 (198.95)	101.09 (78.01)
31		439.09	627.74 (137.94)	101.98 (47.22)
32		437.14	759.46 (298.79)	109.10 (93.77)
33		740.38	636.53 (150.30)	95.33
34		566.29		184.89 (54.49)
35		670.71 (239.41)	793.43 (301.00)	135.71 (121.72)
36		643.81 (52.53)	911.28 (172.07)	230.30 (28.83)
37		857.98 (250.35)	764.05 (39.66)	173.76 (159.34)
38		630.00	959.18 (221.80)	306.62 (82.79)
39			1298.08	198.05 (216.16)
40			1613.32	170.67
41				456.11 (77.87)
42				323.38 (273.46)
43				529.58 (102.95)
44				266.88 (161.25)
46				219.34 (118.94)
47				335.24
48				284.45
49				377.51 (173.69)
51				457.12 (173.89)
53				271.49
54				643.81

56	652.82 (365.73)
58	514.12
60	487.06
63	558.82
65	611.76
66	752.58
67	780.45

Abbreviations: EBRT: external beam radiotherapy; PSMA-RLT: prostate-specific membrane antigen-targeted radioligand therapy; SD: standard deviation.

**Table S5:** Survival time [d] for each animal within a treatment group.

<b>Animal</b>	<b>Untreated</b>	<b>EBRT only</b>	<b>PSMA-RLT only</b>	<b>Combination</b>
1	12	38	21	44
2	28	28	36	37
3	26	12	35	42
4	17	37	40	43
5	30	27	38	51
6	16	28	38	48
7	28	22	29	67
8	22	37	31	42
9	23	16	36	44
10	13	17	32	56
<b>Mean</b>	21.5	26.2	33.6	47.4
<b>SD</b>	6.6	9.4	5.6	8.7

Abbreviations: EBRT: external beam radiotherapy; PSMA-RLT: prostate-specific membrane antigen-targeted radioligand therapy; SD: standard deviation.

**Table S6:** p-values for comparisons of differences regarding tumor control and survival outcomes between treatment groups in the *in vivo* therapy study.

Parameter	Tumor control		
	EBRT only vs.	EBRT only vs.	PSMA-RLT only vs.
	PSMA-RLT only	Combination	Combination
Mean tumor growth	0.9672, ns	< 0.0001, ****	< 0.0001, ****
Time to 2-fold tumor increase	0.0058, **	< 0.0001, ****	< 0.0001, ****
Time to 5-fold tumor increase	0.9047, ns	< 0.0001, ****	< 0.0001, ****

### Survival

Survival curve	0.1239, ns	< 0.0001, ****	< 0.0001, ****
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Statistical analysis: Mixed-Effects Model and Tukey's multiple comparisons testing (tumor growth); one-way ANOVA and Tukey's multiple comparisons testing (time to 2-fold or 5-fold increase); Kaplan-Meier survival analysis and log-rank test (survival).

ns = not significant, \*\* =  $p < 0.01$ , \*\*\*\* =  $p < 0.0001$ .

Abbreviations: ANOVA: analysis of variance; EBRT: external beam radiotherapy; PSMA-RLT: prostate-specific membrane antigen-targeted radioligand therapy.

**Table S7:** Tumor-absorbed doses [Gy/MBq] for each animal that received [<sup>177</sup>Lu]Lu-PSMA-617. Doses were calculated for different time periods: Day of injection until day 21 ( $t = 0\text{-}21$ ); day of injection until infinity ( $t = 0\text{-infinity}$ ); day of injection until the last measurement ( $t = 0\text{-last data point}$ ).

<b>PSMA-RLT only</b>			
<b>Animal</b>	<b><math>t = 0\text{-}21</math></b>	<b><math>t = 0\text{-infinity}</math></b>	<b><math>t = 0\text{-last data point}</math></b>
<b>1</b>	0.59	0.68	0.58
<b>2</b>	0.42	0.46	0.40
<b>3</b>	0.46	0.49	0.45
<b>4</b>	0.22	0.22	0.20
<b>5</b>	0.38	0.40	0.33
<b>6</b>	0.43	0.43	0.39
<b>7</b>	0.25	0.26	0.23
<b>8</b>	0.38	0.39	0.32
<b>9</b>	0.27	0.28	0.24
<b>10</b>	0.25	0.25	0.24
<b>Mean</b>	<b>0.36</b>	<b>0.39</b>	<b>0.34</b>
<b>SD</b>	<b>0.12</b>	<b>0.14</b>	<b>0.12</b>
<b>Combination</b>			
<b>Animal</b>	<b><math>t = 0\text{-}21</math></b>	<b><math>t = 0\text{-infinity}</math></b>	<b><math>t = 0\text{-last data point}</math></b>
<b>1</b>	0.23	0.25	0.21
<b>2</b>	0.17	0.18	0.16
<b>3</b>	0.16	0.19	0.14
<b>4</b>	0.18	0.24	0.14
<b>5</b>	0.59	0.62	0.54
<b>6</b>	0.49	0.51	0.46
<b>7</b>	0.90	0.94	0.82
<b>8</b>	0.45	0.46	0.43

<b>9</b>	0.40	0.40	0.28
<b>10</b>	0.57	0.59	0.52
<b>Mean</b>	<b>0.41</b>	<b>0.44</b>	<b>0.37</b>
<b>SD</b>	<b>0.24</b>	<b>0.24</b>	<b>0.22</b>

Abbreviations: [<sup>177</sup>Lu]Lu-PSMA-617: lutetium-177 (<sup>177</sup>Lu)-labeled PSMA-617; PSMA-RLT: prostate-specific membrane antigen-targeted radioligand therapy; SD: standard deviation.