

## General Information

Analytical and preparative high-performance liquid chromatography (HPLC) was performed using Shimadzu gradient systems (Neufahrn, Germany) equipped with an SPD-20A UV/Vis detector. The columns for analytical (MultoKrom 100C18, 150×4.6 mm, 5 μm), radio-analytical (Multospher 100RP18, 125×4.6 mm, 5 μm) and preparative (MultoKrom 100C18, 250×20 mm, 5 μm) HPLC were purchased from CS Chromatographie Service (Langerwehe, Germany). Eluents for all HPLC operations were water (solvent A) and acetonitrile with 2 vol.% water (solvent B), both containing 0.1 vol.% trifluoroacetic acid (TFA). Radioactivity was detected via a HERM LB 500 NaI detector (Berthold Technologies, Bad Wildbad, Germany). Radio-thin layer chromatography (TLC) was carried out with a Scan-RAM detector (LabLogic Systems, Sheffield, United Kingdom). Electrospray ionization-mass spectra were acquired on an expression<sup>+</sup> CMS (Advion, Harlow, UK).

## Synthesis of PSMA Ligands

The uncomplexed radiohybrid ligands rhPSMA-7.1, -7.2, -7.3 and -7.4 were prepared by applying a mixed solid phase/solution phase synthetic strategy, according to the literature protocols (1). rhPSMA-10.1 and -10.2 were obtained in analogy to the rhPSMA-7 isomers, by substitution of the DOTA-GA chelator with DOTA. PSMA I&T was prepared according to the published procedure (2) and PSMA-617 was purchased from MedChemExpress LLC (Monmouth Junction, USA).

For complexation with non-radioactive lutetium for *in vitro* studies, a 2 mM solution of the PSMA inhibitor (1.0 eq.) in DMSO was combined with a 20 mM aqueous solution of LuCl<sub>3</sub> (2.5 eq.) and heated to 95°C for 30 min. Analytical data of the Lu-chelated PSMA ligands are summarized in the supporting information.

## Analytical data of Lu-complexed PSMA inhibitors:

**Lu-rhPSMA-7.1:** RP-HPLC (10 to 70% B in 15 min):  $t_R = 9.7$  min,  $K' = 3.85$ . Calculated monoisotopic mass (C<sub>63</sub>H<sub>96</sub>FLuN<sub>12</sub>O<sub>25</sub>Si): 1642.6; found:  $m/z = 1643.5$  [M+H]<sup>+</sup>, 822.5 [M+2H]<sup>2+</sup>.

**Lu-rhPSMA-7.2:** RP-HPLC (10 to 70% B in 15 min):  $t_R = 9.4$  min,  $K' = 3.70$ . Calculated monoisotopic mass (C<sub>63</sub>H<sub>96</sub>FLuN<sub>12</sub>O<sub>25</sub>Si): 1642.6; found:  $m/z = 1642.9$  [M+H]<sup>+</sup>, 822.0 [M+2H]<sup>2+</sup>.

**Lu-rhPSMA-7.3:** RP-HPLC (10 to 70% B in 15 min):  $t_R = 9.6$  min,  $K' = 3.80$ . Calculated monoisotopic mass (C<sub>63</sub>H<sub>96</sub>FLuN<sub>12</sub>O<sub>25</sub>Si): 1642.6; found:  $m/z = 1643.4$  [M+H]<sup>+</sup>, 822.3 [M+2H]<sup>2+</sup>.

**Lu-rhPSMA-7.4:** RP-HPLC (10 to 70% B in 15 min):  $t_R = 9.6$  min,  $K' = 3.80$ . Calculated monoisotopic mass (C<sub>63</sub>H<sub>96</sub>FLuN<sub>12</sub>O<sub>25</sub>Si): 1642.6; found:  $m/z = 1643.0$  [M+H]<sup>+</sup>, 822.3 [M+2H]<sup>2+</sup>.

**Lu-rhPSMA-10.1:** RP-HPLC (10 to 70% B in 15 min):  $t_R = 9.9$  min,  $K' = 3.95$ . Calculated monoisotopic mass ( $C_{60}H_{92}FLuN_{12}O_{23}Si$ ): 1570.6; found:  $m/z = 1571.8 [M+H]^+$ ,  $786.2 [M+2H]^{2+}$ .

**Lu-rhPSMA-10.2:** RP-HPLC (10 to 70% B in 15 min):  $t_R = 9.6$  min,  $K' = 3.80$ . Calculated monoisotopic mass ( $C_{60}H_{92}FLuN_{12}O_{23}Si$ ): 1570.6; found:  $m/z = 1571.9 [M+H]^+$ ,  $786.6 [M+2H]^{2+}$ .

**Lu-PSMA-I&T:** RP-HPLC (10 to 70% B in 15 min):  $t_R = 7.2$  min,  $K' = 3.32$ . Calculated monoisotopic mass ( $C_{63}H_{89}ILuN_{11}O_{23}$ ): 1669.5; found:  $m/z = 1670.5 [M+H]^+$ ,  $1113.8 [2M+3H]^{3+}$ .

**Lu-PSMA-617:** RP-HPLC (10 to 70% B in 15 min):  $t_R = 6.5$  min,  $K' = 2.82$ . Calculated monoisotopic mass ( $C_{49}H_{68}LuN_9O_{16}$ ): 1213.4; found:  $m/z = 1213.6 [M+H]^+$ ,  $607.5 [M+2H]^{2+}$ .

**Supplemental Table 1:** Radiochemical purity (RCP) of  $^{177}Lu$ -labeled rhPSMA isomers, determined by radio-TLC before lipophilicity studies, using either the citrate buffer system (0.1 M sodium citrate buffer on iTLC-SG chromatography paper) or acetate buffer system (1.0 M  $NH_4OAc$ /DMF buffer (1/1; v/v) on TLC Silica gel 60 F254 plates).

$[^{177}Lu]Lu-$	<b>citrate-TLC RCP</b>	<b>acetate-TLC RCP</b>
rhPSMA-7.1	99.5%	99.7%
rhPSMA-7.2	99.8%	99.8%
rhPSMA-7.3	99.8%	99.9%
rhPSMA-7.4	99.8%	99.8%
rhPSMA-10.1	99.8%	99.8%
rhPSMA-10.2	99.7%	99.7%

**Supplemental Table 2:** Binding affinities ( $IC_{50}$  [nM], 1 h, 4°C) of Lu-rhPSMA-7.1 to -7.4 (n=3), Lu-rhPSMA-10.1, -10.2 (n=3) and the references Lu-PSMA-617 and Lu-PSMA-I&T (n=3); PSMA-mediated internalization of [ $^{177}\text{Lu}$ ]Lu-rhPSMA-7.1 to -7.4 (n=3), [ $^{177}\text{Lu}$ ]Lu-rhPSMA-10.1, -10.2 (n=3) and the references [ $^{177}\text{Lu}$ ]Lu-PSMA-617 and [ $^{177}\text{Lu}$ ]Lu-PSMA I&T (n=3) by LNCaP cells (1 h, 37°C) as a percentage of the reference ligand ([ $^{125}\text{I}$ ]IBA)KuE); Lipophilicity of [ $^{177}\text{Lu}$ ]Lu-rhPSMA-7.1 to -7.4 (n=6), [ $^{177}\text{Lu}$ ]Lu-rhPSMA-10.1, -10.2 (n=6) and the references [ $^{177}\text{Lu}$ ]Lu-PSMA-617 and [ $^{177}\text{Lu}$ ]Lu-PSMA I&T (n=6), expressed as partition coefficient ( $\log D_{7.4}$ ) using the *n*-octanol/PBS (pH 7.4) distribution system ; Apparent molecular weight ( $MW_{\text{app}}$ ) of [ $^{177}\text{Lu}$ ]Lu-rhPSMA-7.1 to -7.4, [ $^{177}\text{Lu}$ ]Lu-rhPSMA-10.1, -10.2 and the references [ $^{177}\text{Lu}$ ]Lu-PSMA-617 and [ $^{177}\text{Lu}$ ]Lu-PSMA I&T, as determined by human serum albumin mediated size exclusion chromatography (AMSEC).

Compound	[ $^{177}\text{Lu}$ ]Lu-rhPSMA-7.1	[ $^{177}\text{Lu}$ ]Lu-rhPSMA-7.2	[ $^{177}\text{Lu}$ ]Lu-rhPSMA-7.3	[ $^{177}\text{Lu}$ ]Lu-rhPSMA-7.4	[ $^{177}\text{Lu}$ ]Lu-rhPSMA-	[ $^{177}\text{Lu}$ ]Lu-rhPSMA-	[ $^{177}\text{Lu}$ ]Lu-PSMA-617	[ $^{177}\text{Lu}$ ]Lu-PSMA I&T
$IC_{50}$ [nM]	3.11 ± 0.64	2.88 ± 1.06	3.29 ± 1.00	3.06 ± 1.51	2.76 ± 0.51	3.61 ± 0.59	3.27 ± 0.19	4.17 ± 0.78
Internalization [%IBA-KuE]	137.4 ± 5.8	197.3 ± 15.0	184.4 ± 11.8	190.4 ± 10.5	177.4 ± 14.6	205.6 ± 8.3	203.2 ± 10.1	145.4 ± 13.8
$\log D_{7.4}$	-4.27 ± 0.24	-4.25 ± 0.29	-4.12 ± 0.11	-4.10 ± 0.14	-3.78 ± 0.06	-3.83 ± 0.10	-4.1 ± 0.1	-4.1 ± 0.1
$MW_{\text{app}}$ [kDa]	26.3	31.7	30.4	35.7	25.1	21.8	13.7	5.3

**Supplemental Table 3:** Biodistribution of [<sup>177</sup>Lu]Lu-rhPSMA-7.1 to -7.4 at 24 h p.i. in male LNCaP tumor-bearing SCID mice. Data are expressed as a percentage of the injected dose per gram (% ID/g), mean ± standard deviation (SD; n=4).

uptake in % ID/g	[ <sup>177</sup> Lu]Lu- rhPSMA-7.1 24 h p.i., n = 4		[ <sup>177</sup> Lu]Lu- rhPSMA-7.2 24 h p.i., n = 4		[ <sup>177</sup> Lu]Lu- rhPSMA-7.3 24 h p.i., n = 4		[ <sup>177</sup> Lu]Lu- rhPSMA-7.4 24 h p.i., n = 4	
	mean	SD	mean	SD	mean	SD	mean	SD
<b>blood</b>	<b>0.0023</b>	<b>0.0006</b>	<b>0.0090</b>	<b>0.0015</b>	<b>0.0032</b>	<b>0.0007</b>	<b>0.0052</b>	<b>0.0026</b>
heart	0.036	0.009	0.032	0.005	0.031	0.009	0.032	0.006
lung	0.050	0.006	0.049	0.009	0.049	0.011	0.056	0.014
<b>liver</b>	<b>0.28</b>	<b>0.06</b>	<b>0.24</b>	<b>0.06</b>	<b>0.22</b>	<b>0.04</b>	<b>0.16</b>	<b>0.03</b>
spleen	0.24	0.04	0.79	0.13	0.73	0.23	0.72	0.29
pancreas	0.019	0.003	0.076	0.071	0.025	0.005	0.027	0.006
stomach	0.071	0.018	0.246	0.236	0.059	0.028	0.068	0.029
intestine	0.15	0.05	0.34	0.31	0.13	0.08	0.15	0.07
<b>kidney</b>	<b>4.10</b>	<b>0.81</b>	<b>19.04</b>	<b>4.45</b>	<b>9.82</b>	<b>2.74</b>	<b>15.66</b>	<b>8.92</b>
adrenals	0.12	0.04	0.61	0.13	0.56	0.22	0.34	0.18
muscle	0.011	0.001	0.010	0.002	0.009	0.003	0.010	0.003
bone	0.030	0.009	0.034	0.008	0.048	0.010	0.023	0.009
<b>tumor</b>	<b>12.40</b>	<b>1.27</b>	<b>12.73</b>	<b>2.76</b>	<b>11.63</b>	<b>2.24</b>	<b>12.56</b>	<b>3.52</b>
parotid gl.	0.083	0.020	0.114	0.030	0.131	0.042	0.111	0.057
submand. gl.	0.051	0.004	0.072	0.005	0.056	0.010	0.075	0.019

**Supplemental Table 4:** Biodistribution of [<sup>177</sup>Lu]Lu-rhPSMA-10.1, -10.2 and the references [<sup>177</sup>Lu]Lu-PSMA-617 and [<sup>177</sup>Lu]Lu-PSMA I&T at 24 h p.i. in male LNCaP tumor-bearing SCID mice. Data are expressed as a percentage of the injected dose per gram (% ID/g), mean ± standard deviation (SD; n=4-5). Values of [<sup>177</sup>Lu]Lu-PSMA I&T were taken from a previously published study by our group (3). N.d. not determined.

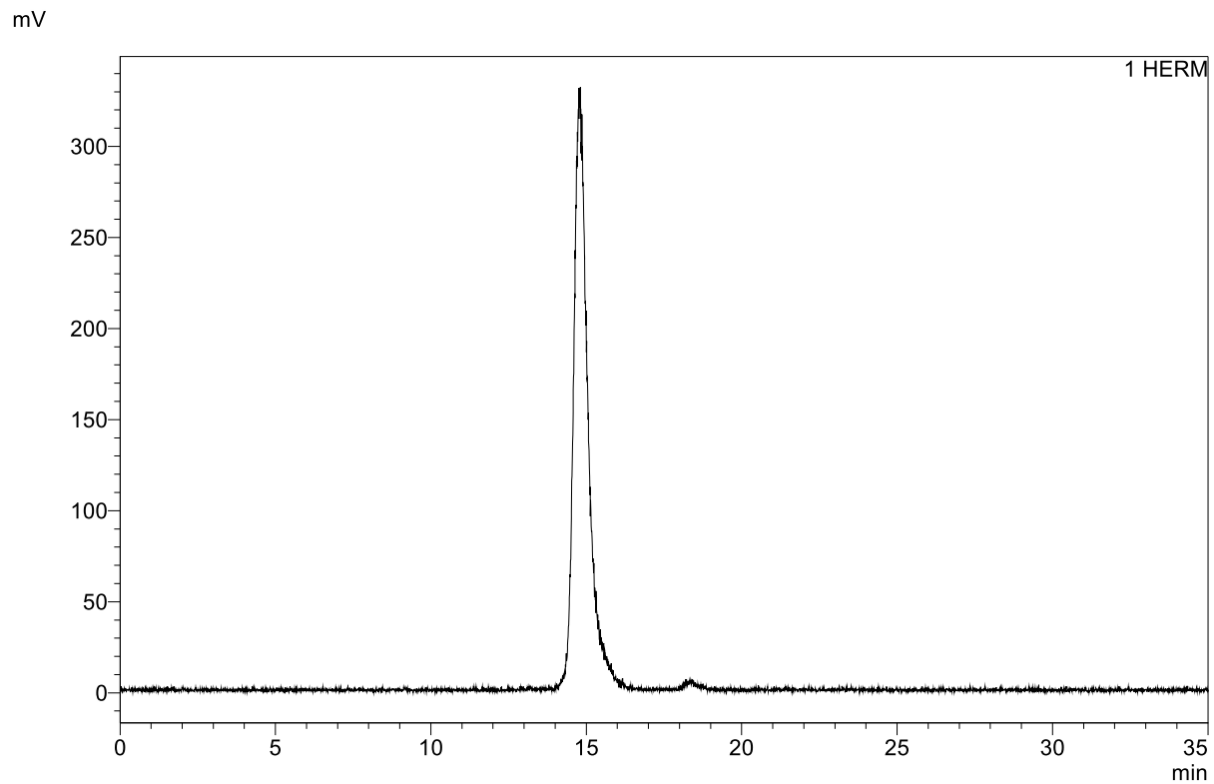
uptake in % ID/g	<b>[<sup>177</sup>Lu]Lu- rhPSMA-10.1 24 h p.i., n = 5</b>		<b>[<sup>177</sup>Lu]Lu- rhPSMA-10.2 24 h p.i., n = 4</b>		<b>[<sup>177</sup>Lu]Lu- PSMA-617 24 h p.i., n = 4</b>		<b>[<sup>177</sup>Lu]Lu- PSMA-I&amp;T 24 h p.i., n = 4</b>	
	mean	SD	mean	SD	mean	SD	mean	SD
<b>blood</b>	<b>0.0009</b>	<b>0.0001</b>	<b>0.0022</b>	<b>0.0008</b>	<b>0.0056</b>	<b>0.0026</b>	<b>0.0118</b>	<b>0.0064</b>
heart	0.017	0.001	0.020	0.004	0.012	0.004	0.049	0.034
lung	0.032	0.004	0.036	0.007	0.039	0.013	0.158	0.029
<b>liver</b>	<b>0.18</b>	<b>0.06</b>	<b>0.15</b>	<b>0.05</b>	<b>0.12</b>	<b>0.06</b>	<b>0.05</b>	<b>0.01</b>
spleen	0.17	0.03	0.53	0.26	0.08	0.01	1.94	1.01
pancreas	0.013	0.002	0.021	0.002	0.011	0.004	0.048	0.029
stomach	0.056	0.014	0.104	0.069	0.020	0.003	0.048	0.021
intestine	0.11	0.05	0.24	0.13	0.12	0.08	0.12	0.06
<b>kidney</b>	<b>1.97</b>	<b>0.78</b>	<b>8.09</b>	<b>1.68</b>	<b>1.44</b>	<b>0.42</b>	<b>34.66</b>	<b>17.20</b>
adrenals	0.06	0.04	0.41	0.17	0.19	0.07	1.06	0.24
muscle	0.003	0.002	0.007	0.002	0.009	0.002	0.010	0.004
bone	0.022	0.008	0.020	0.008	0.027	0.017	0.014	0.004
<b>tumor</b>	<b>9.82</b>	<b>0.30</b>	<b>10.45</b>	<b>3.25</b>	<b>7.46</b>	<b>0.90</b>	<b>4.06</b>	<b>1.12</b>
parotid gl.	0.041	0.009	0.100	0.017	n.d.	n.d.	n.d.	n.d.
submand. gl.	0.037	0.007	0.046	0.009	n.d.	n.d.	n.d.	n.d.

**Supplemental Table 5:** Tumor-to-organ ratios of [<sup>177</sup>Lu]Lu-rhPSMA-7.1 to -7.4 at 24 h p.i. in male LNCaP tumor-bearing SCID mice. Mean ± standard deviation (SD; n=4). Tumor-to-organ ratios were calculated from individual mice and mean values were determined.

tumor-to-organ ratio	<b>[<sup>177</sup>Lu]Lu-rhPSMA-7.1</b> 24 h p.i., n = 4		<b>[<sup>177</sup>Lu]Lu-rhPSMA-7.2</b> 24 h p.i., n = 4		<b>[<sup>177</sup>Lu]Lu-rhPSMA-7.3</b> 24 h p.i., n = 4		<b>[<sup>177</sup>Lu]Lu-rhPSMA-7.4</b> 24 h p.i., n = 4	
	mean	SD	mean	SD	mean	SD	mean	SD
<b>blood</b>	<b>5971</b>	<b>2358</b>	<b>1460</b>	<b>495</b>	<b>3843</b>	<b>1145</b>	<b>2570</b>	<b>820</b>
heart	368	122	409	125	394	82	387	81
lung	252	53	275	113	239	24	234	87
<b>liver</b>	<b>47</b>	<b>13</b>	<b>54</b>	<b>12</b>	<b>54</b>	<b>2</b>	<b>78</b>	<b>13</b>
spleen	54	13	16	3	17	3	19	4
pancreas	671	211	285	170	487	133	489	180
stomach	191	70	187	210	262	177	228	135
intestine	95	40	150	211	129	86	104	55
<b>kidney</b>	<b>3.2</b>	<b>0.9</b>	<b>0.7</b>	<b>0.2</b>	<b>1.2</b>	<b>0.3</b>	<b>1.0</b>	<b>0.5</b>
adrenals	119	54	21	1	23	7	43	14
muscle	1167	214	1453	623	1454	601	1269	92
bone	453	190	384	66	264	111	616	309
parotid gl.	163	69	112	14	93	21	99	13
submand. gl.	244	43	177	40	207	27	169	33

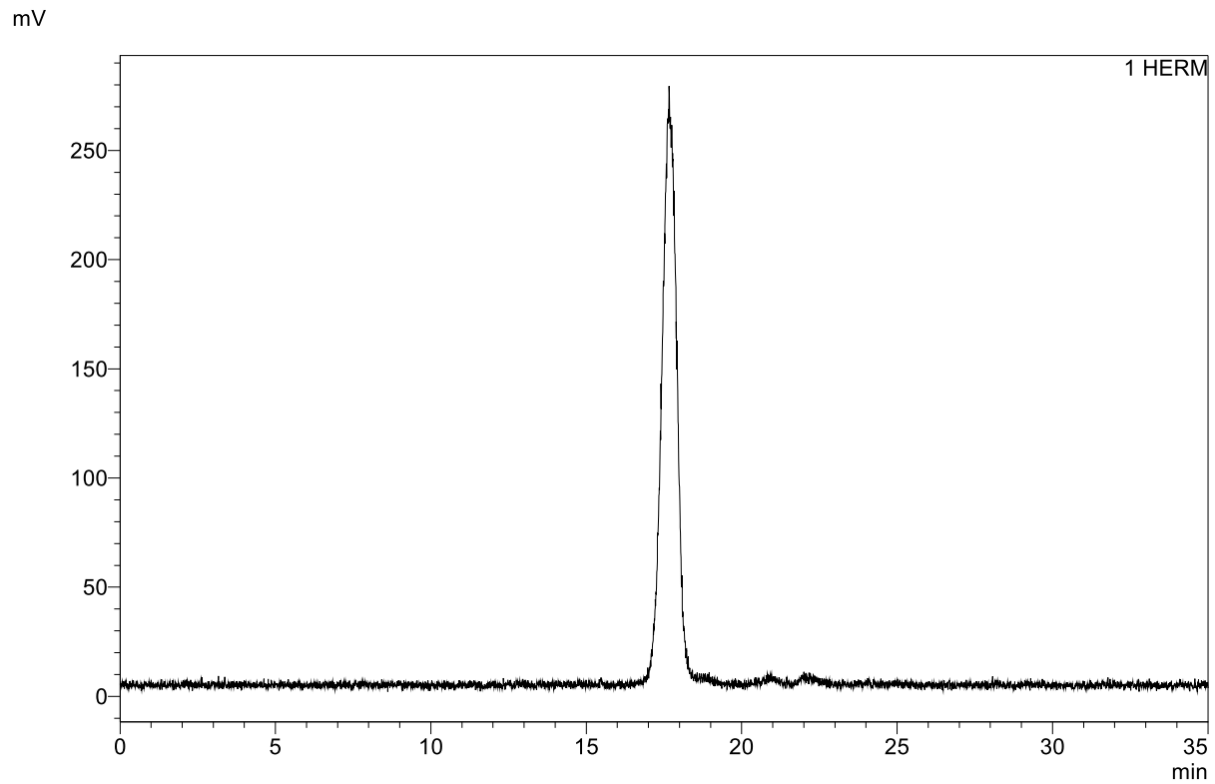
**Supplemental Table 6:** Tumor-to-organ ratios [<sup>177</sup>Lu]Lu-rhPSMA-10.1, -10.2 and the references [<sup>177</sup>Lu]Lu-PSMA-617 and [<sup>177</sup>Lu]Lu-PSMA I&T at 24 h p.i. in male LNCaP tumor-bearing SCID mice. Mean ± standard deviation (SD; n=4-5). Values of [<sup>177</sup>Lu]PSMA I&T were taken from a previously published study by our group (3). Tumor-to-organ ratios were calculated from individual mice and mean values were determined.

tumor-to-organ ratio	<b>[<sup>177</sup>Lu]Lu-rhPSMA-10.1</b> 24 h p.i., n = 5		<b>[<sup>177</sup>Lu]Lu-rhPSMA-10.2</b> 24 h p.i., n = 4		<b>[<sup>177</sup>Lu]Lu-PSMA-617</b> 24 h p.i., n = 4		<b>[<sup>177</sup>Lu]Lu-PSMA-I&amp;T</b> 24 h p.i., n = 4	
	mean	SD	mean	SD	mean	SD	mean	SD
<b>blood</b>	<b>11498</b>	<b>1953</b>	<b>6125</b>	<b>4185</b>	<b>1424</b>	<b>455</b>	<b>408</b>	<b>209</b>
heart	568	52	585	321	704	270	128	77
lung	313	33	295	100	210	74	26	9
<b>liver</b>	<b>57</b>	<b>13</b>	<b>69</b>	<b>17</b>	<b>91</b>	<b>73</b>	<b>86</b>	<b>31</b>
spleen	58	11	27	19	93	22	3.0	2.0
pancreas	774	189	511	198	799	293	144	115
stomach	187	47	221	231	376	65	105	60
intestine	115	75	111	162	90	50	49	33
<b>kidney</b>	<b>5.7</b>	<b>2.0</b>	<b>1.4</b>	<b>0.6</b>	<b>5.9</b>	<b>3.0</b>	<b>0.15</b>	<b>0.08</b>
adrenals	171	111	33	22	44	15	4.1	1.5
muscle	2441	373	1798	792	855	279	496	242
bone	595	462	589	280	472	378	315	144
parotid gl.	250	64	107	37	n.d.	n.d.	n.d.	n.d.
submand. gl.	275	49	223	45	n.d.	n.d.	n.d.	n.d.

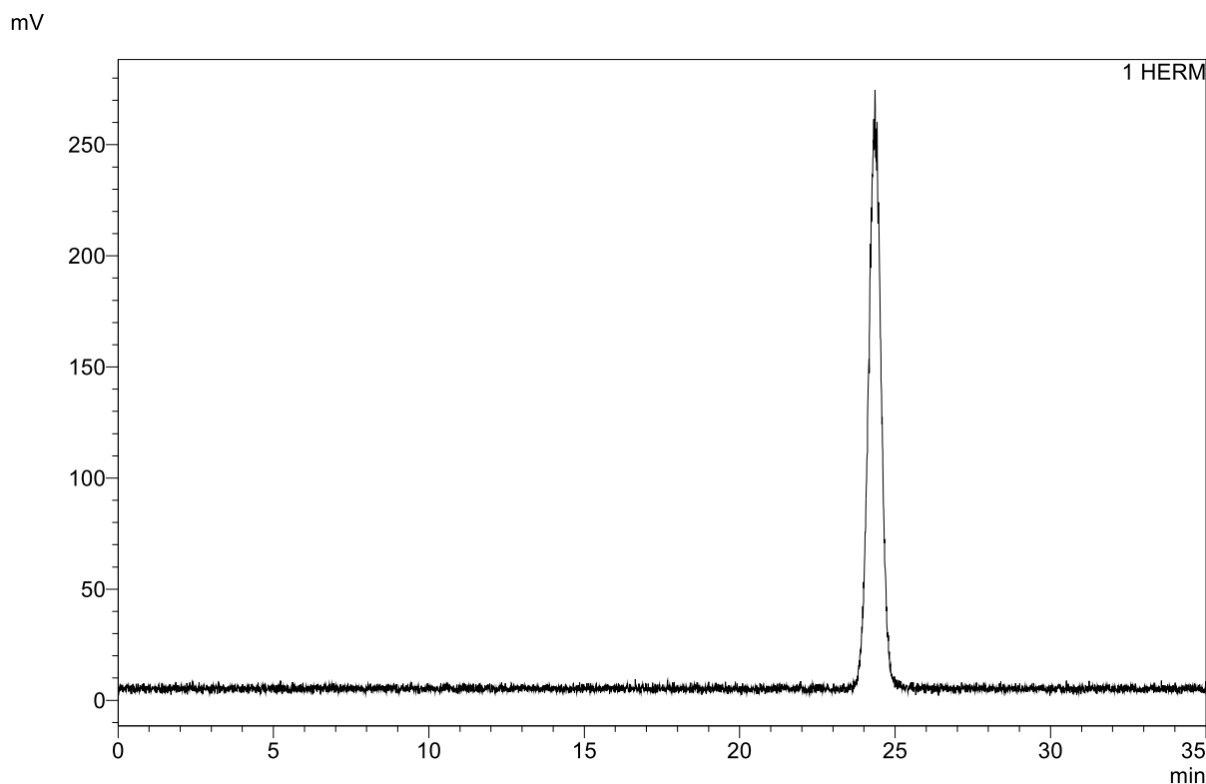


**Supplemental Figure 1:** Exemplary radio-chromatogram of [<sup>177</sup>Lu]Lu-rhPSMA-7.3 in the AMSEC experiment. The retention time of 14.8 min measured in this particular experiment correlates with an AMW of 30.3 kDa (AMW of 30.4 ± 0.5 kDa (mean ± SD) for n = 10 independent measurements).





**Supplemental Figure 2:** Radio-chromatogram of [ $^{177}\text{Lu}$ ]Lu-PSMA-617 in the AMSEC experiment. The retention time of 17.7 min measured in this particular experiment correlates with an AMW of 13.7 kDa.



**Supplemental Figure 3:** Radio-chromatogram of [ $^{18}\text{F}$ ]fluoride in the AMSEC experiment. Due to the absence of interaction of fluoride with HSA, [ $^{18}\text{F}$ ]fluoride is eluted after 24.4 min, as expected for a small inorganic ion, and the corresponding AMW of 2.1 kDa lies below the fractionation range of the gel filtration column (70 – 3 kDa).

## References

1. Wurzer A, Parzinger M, Konrad M, et al. Preclinical comparison of four [(18)F, (nat)Ga]rhPSMA-7 isomers: influence of the stereoconfiguration on pharmacokinetics. *EJNMMI Res.* 2020;10:149.
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3. Wirtz M, Schmidt A, Schottelius M, et al. Synthesis and in vitro and in vivo evaluation of urea-based PSMA inhibitors with increased lipophilicity. *EJNMMI Res.* 2018;8:84.