

Table of Contents Entry

Results

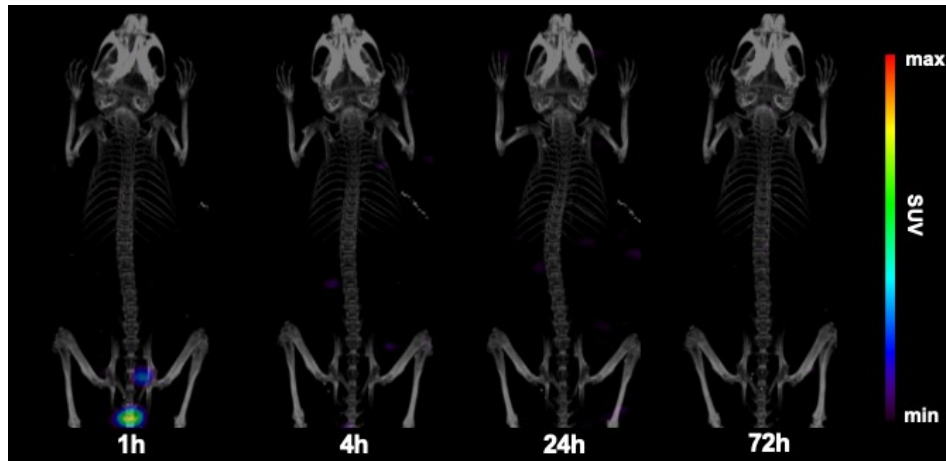
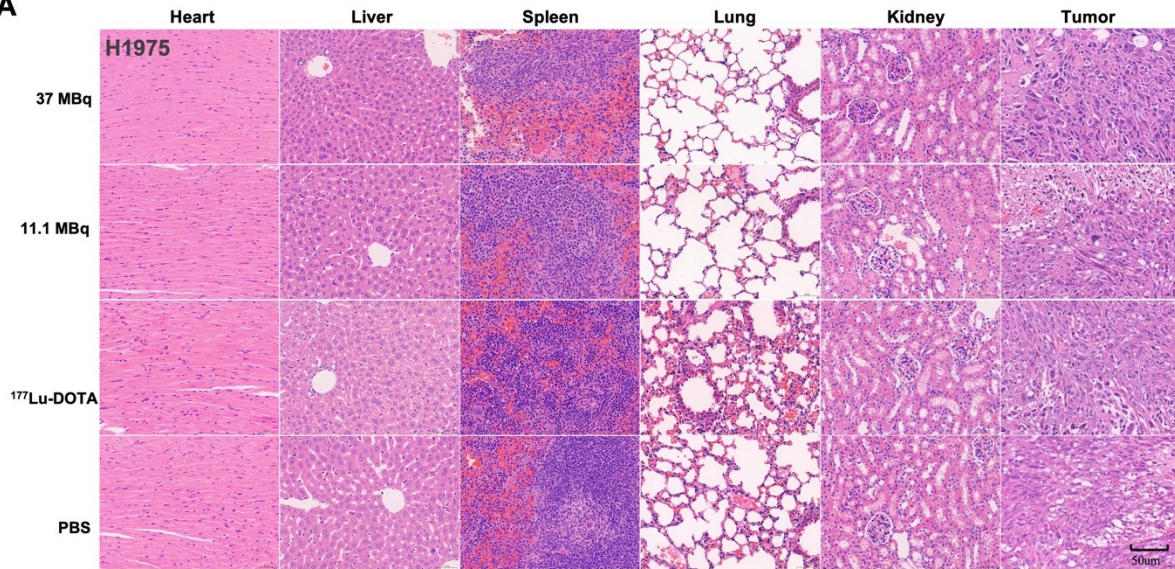


Figure S1. SPECT imaging of $[^{177}\text{Lu}]\text{Lu-DOTA}$ on H1975 xenograft models.

A

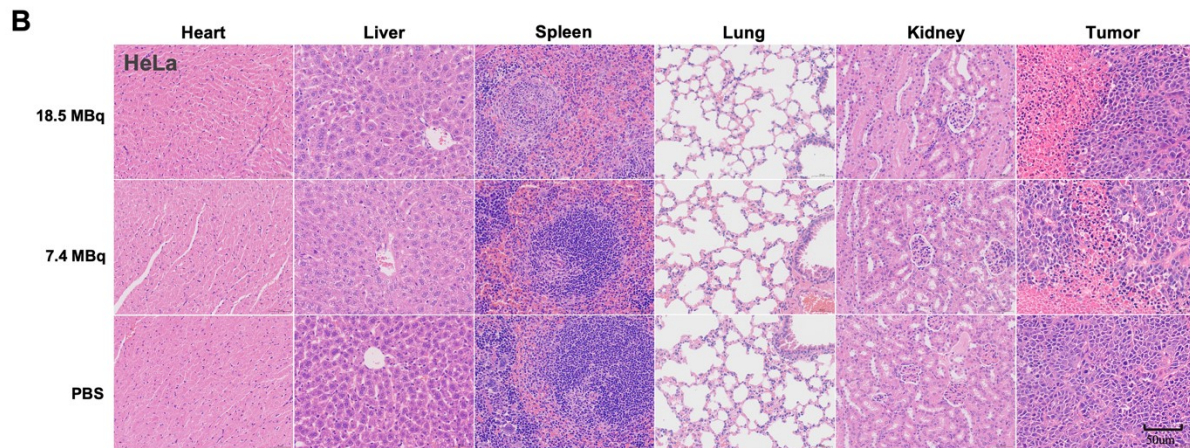


Figure S2. H&E staining was performed on major organs (heart, lung, liver, spleen, and kidney) and tumors for H1975 (A) and HeLa (B) xenograft models. Scale bar: 50 μm .

Animal SPECT image acquisition and processing

Table S1 summarizes the performance metrics of the animal SPECT scanner.

Table S2 provides detailed information on animal SPECT image acquisition and processing parameters.

Table S1. Characteristics of the Novel Medical animal SPECT scanner

Imaging module	Performance metrics	
SPECT	Cross-sectional view	70 mm
	Axial effective field of view	250 mm
	Spatial resolution	≤ 1.5 mm
	Central detection efficiency	$\geq 0.4\%$ (Tc99m)
	Energy Range	80 keV–400 keV
	Image Uniformity	$< 15\%$
CT	Maximum cross-sectional field of view	70 mm
	Maximum axial field of view	250 mm
	High contrast resolution	≥ 6.25 lp/mm
	Low contrast resolution	$\leq 3\%$

Table S2. Image acquisition and processing data

Imaging module	Scanning metrics		Reconstruction metrics	
CT	Voltage	50 kV	Displaying Matrix	512 x 512
	Current	0.5 mA	Slices	500
	Total frames	360	Filter	Soft
	Single frame acquisition time	2000 ms	Pixel size	0.136722 mm
	Step angle	1.00°	Slice thickness	0.18000 mm
SPECT	Scout image	Adopt CT scout image	Algorithm	OSEM
	Acquisition time	25 min	Displaying Matrix	60 x 60
			Slices	114
			Iteration ordinal number	40
			Pixel size	0.70000 mm
			Slice thickness	0.70000 mm

Table S3. Fluorescence intensity-based quantitative analysis of DOTA-DZ-HX

		Time (h)			
		4	24	48	96
Tumor	P/s/cm ² /sr (e+009)	1.75 ± 0.10	1.83 ± 0.23	1.04 ± 0.13	0.49 ± 0.06
Tumor/ background	ratio	98.95 ± 2.89	108.15 ± 10.75	80.12 ± 10.97	65.22 ± 8.01

Table S4. Organ biodistribution of [¹⁷⁷Lu]Lu-DOTA-DZ-HX in H1975 xenograft models*

Organ	¹⁷⁷ Lu]Lu-DOTA-DZ-HX					
	1 h	4 h	24 h	48 h	72 h	96 h
Blood	36.715 ± 7.213	24.936 ± 4.618	18.364 ± 6.168	7.831 ± 0.948	4.604 ± 0.732	0.685 ± 0.218
Heart	8.199 ± 1.354	6.430 ± 1.460	5.489 ± 0.668	3.705 ± 0.725	3.319 ± 0.501	0.935 ± 0.594
Liver	7.302 ± 2.060	5.788 ± 0.722	8.296 ± 0.452	8.127 ± 1.101	6.706 ± 0.724	2.335 ± 0.851
Spleen	5.111 ± 1.441	3.873 ± 0.684	5.726 ± 0.177	5.445 ± 0.916	5.313 ± 0.941	1.633 ± 0.906
Lung	9.895 ± 1.946	7.476 ± 0.863	5.391 ± 1.041	4.789 ± 0.759	3.835 ± 0.709	1.319 ± 0.557
Kidney	11.562 ± 2.412	14.228 ± 1.634	21.237 ± 2.254	22.129 ± 3.540	18.728 ± 3.240	7.099 ± 1.300
Stomach	3.550 ± 0.970	2.986 ± 0.501	2.867 ± 0.275	1.728 ± 0.184	1.446 ± 0.279	0.454 ± 0.279

Intestine	4.130 ± 1.605	3.854 ± 0.614	2.503 ± 0.436	1.984 ± 0.426	1.762 ± 0.518	0.518 ± 0.374
Muscle	0.970 ± 0.338	1.580 ± 0.588	1.768 ± 0.592	1.500 ± 0.333	1.159 ± 0.171	0.234 ± 0.028
Bone	5.754 ± 2.292	2.259 ± 0.888	1.867 ± 0.297	1.509 ± 0.495	1.089 ± 0.192	0.492 ± 0.203
Brain	0.160 ± 0.023	0.762 ± 0.222	0.569 ± 0.363	0.275 ± 0.069	0.208 ± 0.046	0.054 ± 0.044
Tumor	5.754 ± 0.292	8.697 ± 0.866	13.501 ± 1.401	12.645 ± 1.192	12.055 ± 1.387	2.764 ± 0.518
T/B	0.160 ± 0.023	0.353 ± 0.029	0.779 ± 0.189	1.625 ± 0.164	2.632 ± 0.118	4.139 ± 0.572
T/L	0.826 ± 0.182	1.517 ± 0.201	1.629 ± 0.165	1.590 ± 0.343	1.798 ± 0.076	1.235 ± 0.244
T/K	0.512 ± 0.098	0.614 ± 0.056	0.638 ± 0.067	0.584 ± 0.121	0.649 ± 0.045	0.391 ± 0.055
T/M	3.128 ± 0.495	6.294 ± 2.848	8.135 ± 2.239	8.723 ± 2.003	10.482 ± 1.165	11.794 ± 1.460

*The data was in terms of %ID/g. The data are expressed as the mean \pm SD ($n = 4$ /group; for 96h $n = 3$), T/B: Tumor/Blood, T/L: Tumor/Liver, T/K: Tumor/Kidney, T/M: Tumor/Muscle.

Table S5. Organ biodistribution of [¹⁷⁷Lu]Lu-DOTA-DZ-HX in HeLa xenograft models*

Organ	¹⁷⁷ Lu]Lu-DOTA-DZ-HX				
	4 h	24 h	48 h	72 h	96 h
Blood	37.471 \pm 3.484	21.798 \pm 2.698	10.056 \pm 2.140	6.275 \pm 0.600	4.692 \pm 1.454
Heart	10.163 \pm 1.454	7.989 \pm 1.249	4.854 \pm 0.665	4.195 \pm 0.109	3.953 \pm 0.524
Liver	8.931 \pm 1.553	11.336 \pm 1.021	9.242 \pm 1.279	9.192 \pm 0.356	8.193 \pm 1.019
Spleen	6.251 \pm 0.628	10.066 \pm 1.166	6.935 \pm 1.636	8.565 \pm 2.325	6.998 \pm 1.992
Lung	15.982 \pm 1.754	12.323 \pm 2.034	6.755 \pm 0.289	6.712 \pm 1.417	4.005 \pm 0.609
Kidney	19.447 \pm 1692	31.900 \pm 1.088	25.780 \pm 5.471	29.078 \pm 2.768	25.091 \pm 4.738

Stomach	4.975 ± 0.554	4.377 ± 0.226	2.416 ± 0.634	2.234 ± 0.083	2.476 ± 1.044
Intestine	5.912 ± 1.470	6.352 ± 3.061	2.719 ± 0.970	2.795 ± 0.714	2.846 ± 0.409
Muscle	2.800 ± 0.896	2.259 ± 0.406	1.533 ± 0.237	1.706 ± 0.202	1.525 ± 0.309
Bone	3.949 ± 0.751	2.875 ± 0.498	1.750 ± 0.456	2.157 ± 0.383	1.438 ± 0.301
Brain	0.804 ± 0.174	0.547 ± 0.049	0.313 ± 0.073	0.306 ± 0.182	0.208 ± 0.058
Tumor	11.892 ± 1.012	17.647 ± 4.098	16.522 ± 3.029	14.409 ± 2.743	11.938 ± 1.003
T/B	0.319 ± 0.032	0.804 ± 0.123	1.659 ± 0.220	2.297 ± 0.372	2.672 ± 0.549
T/L	1.350 ± 0.163	1.545 ± 0.271	1.798 ± 0.338	1.563 ± 0.247	1.463 ± 0.073
T/K	0.617 ± 0.096	0.551 ± 0.116	0.647 ± 0.097	0.494 ± 0.065	0.485 ± 0.063

T/M	4.600 ± 1.542	7.772 ± 1.061	11.079 ± 3.119	8.439 ± 1.081	8.012 ± 1.267
------------	---------------	---------------	----------------	---------------	---------------

* The data was in terms of %ID/g. The data are expressed as the mean ± SD ($n = 4/\text{group}$), T/B: Tumor/Blood, T/L: Tumor/Liver, T/K: Tumor/Kidney, T/M: Tumor/Muscle.

Table S6. The H1975 tumor volume of each group*

H1975 Group	Tumor volume (mm ³)		Tumor Growth Inhibition (%) at the end of the therapy
	at the beginning of the therapy	at the end of the therapy	
37 MBq [¹⁷⁷ Lu]Lu-DOTA-DZ-HX	33.59 ± 8.76	10.01 ± 2.75	96.18
11.1 MBq [¹⁷⁷ Lu]Lu-DOTA-DZ-HX	30.77 ± 3.55	15.48 ± 7.51	
37 MBq [¹⁷⁷ Lu]Lu-DOTA	32.91 ± 7.23	196.16 ± 77.37	
PBS	35.81 ± 11.42	279.37 ± 115.53	

*The data are expressed as the mean ± SD (*n* = 5/group).

Table S7. The HeLa tumor volume and weight of each group*

HeLa Group	Tumor volume (mm ³)		Tumor weight	Tumor Growth Inhibition (%) at the end of the therapy
	at the beginning of the therapy	at the end of the therapy		
18.5 MBq [¹⁷⁷ Lu]Lu-DOTA-DZ-HX	80.86 ± 11.19	71.52 ± 28.86	0.04 ± 0.01	89.74
7.4 MBq [¹⁷⁷ Lu]Lu-DOTA-DZ-HX	73.66 ± 7.52	86.41 ± 23.76	0.06 ± 0.02	
PBS	86.97 ± 6.55	749.67 ± 154.30	0.64 ± 0.15	

*The data are expressed as the mean ± SD (*n* = 5/group).

Table S8. Quantitative Evaluation of Tumor Tissue Apoptosis and Immunohistochemical Profiling*

Tumor	Apoptosis rate (%)			% DAB Positive Tissue	
	High	Low	PBS	HIF-1α	OATP2B1
H1975	60.92 ± 7.60	18.57 ± 7.37	2.02 ± 0.41	38.86	32.15
HeLa	27.30 ± 10.49	13.28 ± 0.03	0.00 ± 0.00	58.87	79.38

*The data are expressed as the mean ± SD (*n* = 3/group).