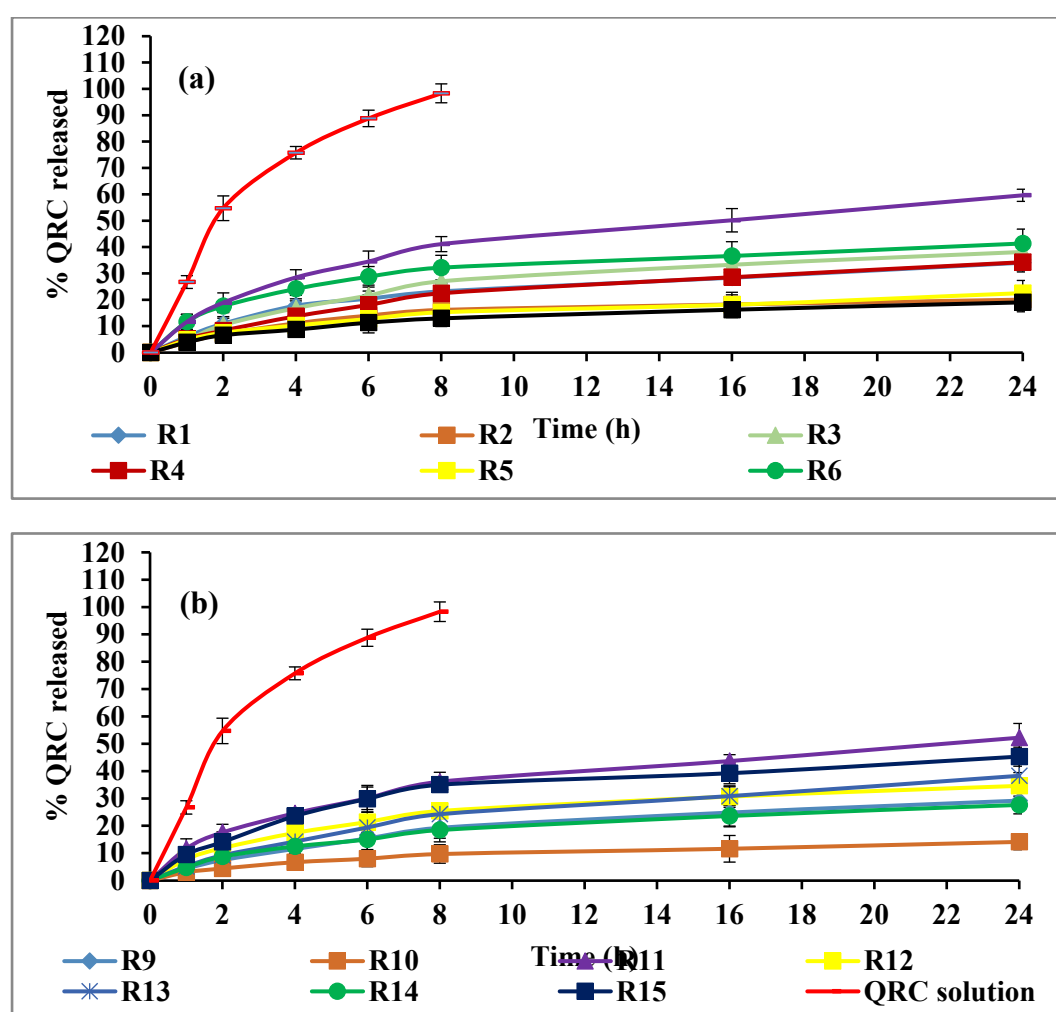


# Supplementary Materials: Tailoring of Selenium-Plated Novasomes for Fine-Tuning Pharmacokinetic and Tumor Uptake of Quercetin: In Vitro Optimization and In Vivo Radiobiodistribution Assessment in Ehrlich Tumor-Bearing Mice

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**Figure S1.** In vitro release profiles of QRC from drug solution and different NOV runs: (a) R1–R8 and (b) R9–R15.

**Table S1.** QRC Biodistribution after i.v. administrations of <sup>99m</sup>Tc-QRC solution, <sup>99m</sup>Tc-QRC-NOVs, and <sup>99m</sup>Tc-QRC-SeNOVs to Ehrlich tumor-bearing mice at various time intervals.

Organ/Tissue	Time (h)							
e	0.08	0.25	0.5	1	2	4	6	24
<sup>99m</sup> Tc-QRC solution								
blood	2.74 ± 0.71	1.75 ± 0.41	0.83 ± 0.17	0.80 ± 0.21	0.79 ± 0.22	0.91 ± 0.34	1.45 ± 0.05	0.36 ± 0.09
Muscle	0.37 ± 0.05	1.01 ± 0.18	1.36 ± 0.08	1.66 ± 0.36	2.98 ± 0.73	1.09 ± 0.04	0.66 ± 0.08	0.24 ± 0.01
Tumor	0.78 ± 0.11	1.05 ± 0.26	1.75 ± 0.09	1.77 ± 0.04	3.32 ± 0.42	1.13 ± 0.03	0.83 ± 0.01	0.45 ± 0.02
T/NT	2.11 ± 0.16	1.04 ± 0.08	1.29 ± 0.13	1.07 ± 0.11	1.11 ± 0.09	1.04 ± 0.05	1.26 ± 0.13	1.88 ± 0.21
T/blood	0.28 ± 0.02	0.60 ± 0.06	2.11 ± 0.12	2.21 ± 0.22	4.20 ± 0.28	1.24 ± 0.16	0.57 ± 0.03	1.25 ± 0.14
<sup>99m</sup> Tc-QRC-NOVs								
Blood	4.10 ± 0.71 <sup>a</sup>	3.75 ± 0.77 <sup>a</sup>	2.69 ± 0.44 <sup>a</sup>	1.30 ± 0.11 <sup>a</sup>	1.00 ± 0.29 <sup>a</sup>	2.10 ± 0.69 <sup>a</sup>	1.47 ± 0.31	0.60 ± 0.09 <sup>a</sup>
Muscle	0.88 ± 0.11 <sup>a</sup>	1.30 ± 0.12 <sup>a</sup>	1.61 ± 0.09 <sup>a</sup>	2.01 ± 0.31 <sup>a</sup>	3.01 ± 0.97	1.67 ± 0.15 <sup>a</sup>	1.09 ± 0.15 <sup>a</sup>	0.76 ± 0.01 <sup>a</sup>
Tumor	2.78 ± 0.39 <sup>a</sup>	4.19 ± 0.97 <sup>a</sup>	5.19 ± 1.37 <sup>a</sup>	7.81 ± 2.03 <sup>a</sup>	9.61 ± 3.16 <sup>a</sup>	7.00 ± 1.39 <sup>a</sup>	5.64 ± 0.94 <sup>a</sup>	2.46 ± 0.07 <sup>a</sup>
T/NT	3.16 ± 0.63 <sup>a</sup>	3.22 ± 0.71 <sup>a</sup>	3.22 ± 0.42 <sup>a</sup>	3.89 ± 0.54 <sup>a</sup>	3.19 ± 0.62 <sup>a</sup>	4.19 ± 0.31 <sup>a</sup>	5.17 ± 0.83 <sup>a</sup>	3.24 ± 0.52 <sup>a</sup>
T/blood	0.68 ± 0.15 <sup>a</sup>	1.12 ± 0.11 <sup>a</sup>	1.93 ± 0.16	6.01 ± 0.87 <sup>a</sup>	9.61 ± 1.23 <sup>a</sup>	3.33 ± 0.58 <sup>a</sup>	3.84 ± 0.63 <sup>a</sup>	4.10 ± 0.81 <sup>a</sup>
<sup>99m</sup> Tc-QRC-SeNOVs								
Blood	5.74 ± 0.61 <sup>a,b</sup>	4.75 ± 0.70 <sup>a,b</sup>	3.80 ± 0.41 <sup>a,b</sup>	2.80 ± 0.29 <sup>a,b</sup>	2.79 ± 0.20 <sup>a,b</sup>	2.91 ± 0.40 <sup>a,b</sup>	2.47 ± 0.60 <sup>a,b</sup>	1.27 ± 0.09 <sup>a,b</sup>
Muscle	0.97 ± 0.06 <sup>a</sup>	1.52 ± 0.39 <sup>a,b</sup>	2.60 ± 0.27 <sup>a,b</sup>	3.01 ± 0.97 <sup>a,b</sup>	3.67 ± 0.47 <sup>a,b</sup>	2.00 ± 0.15 <sup>a,b</sup>	1.87 ± 0.12 <sup>a,b</sup>	0.99 ± 0.02 <sup>a,b</sup>
Tumor	4.83 ± 0.47 <sup>a,b</sup>	5.61 ± 0.27 <sup>a,b</sup>	8.03 ± 2.14 <sup>a,b</sup>	15.74 ± 1.36 <sup>a,b</sup>	12.74 ± 3.79 <sup>a,b</sup>	11.00 ± 2.78 <sup>a,b</sup>	7.93 ± 0.97 <sup>a,b</sup>	4.76 ± 0.89 <sup>a,b</sup>
T/NT	4.98 ± 0.76 <sup>a,b</sup>	3.69 ± 0.72 <sup>a</sup>	3.09 ± 0.41 <sup>a</sup>	5.23 ± 0.61 <sup>a,b</sup>	3.47 ± 0.55 <sup>a</sup>	5.50 ± 0.92 <sup>a,b</sup>	4.24 ± 0.66 <sup>a,b</sup>	4.81 ± 0.83 <sup>a,b</sup>
T/blood	0.84 ± 0.26 <sup>a,b</sup>	1.18 ± 0.35 <sup>a</sup>	2.11 ± 0.31	5.62 ± 0.68 <sup>a</sup>	4.57 ± 0.84 <sup>a,b</sup>	3.78 ± 0.67 <sup>a</sup>	3.21 ± 0.56 <sup>a,b</sup>	3.75 ± 0.49 <sup>a</sup>

<sup>99m</sup>Tc: technetium-99m; QRC: quercetin; NOVs: novasomes; SeNOVs: selenium-plated novasomes; T/NT (target/non-target ratio) = tumor muscle/normal muscle. Listed data are mean values ± SD (n = 3). Using one-way ANOVA followed by Tukey post-hoc test. <sup>a</sup> *p* < 0.05 versus <sup>99m</sup>Tc-QRC solution. <sup>b</sup> *p* < 0.05 versus <sup>99m</sup>Tc-QRC-NOVs.