A Supramolecular Approach For Liver Radioembolization

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Methods

Stability of ^{99m}Tc-Cy5_{0.5}CD₁₀PIBMA₃₉ in FCS

 99m Tc-Cy5_{0.5}CD₁₀PIBMA₃₉ was dissolved in FCS (2.5 µg/mL) and shaken in a water bath at 37 °C for 44 h. After 2, 20, and 44 h 0.1 mL samples were taken and their composition was analyzed by PD-10-SEC.

Stability of ^{99m}Tc-Maa-Ad in FCS

Lyophilized MAA (2 mg) was dissolved in 1 mL of saline (0.9% NaCl, sterile and pyrogen-free, B. Braun Medical Supplies, Inc., Oss, The Netherlands). To one portion 100 µL of a freshly eluted ^{99m}Tc-Napertechnetate solution (500 MBq/mL, Mallinckrodt Medical B.V.) was added and the mixture was gently stirred in a shaking water bath for 1 h at 37 °C. Thereafter, the solution was washed 2 times PBS by 2 centrifugation steps (3 min, 1,200 rpm). Next, 20 µL of Ad-TFP (10 mg/mL DMSO) was added. After allowing it to react in a shaking water bath for 1 h at 37 °C, the reaction mixture was washed 2 times with PBS by 2 centrifugation steps (3 min, 1,200 rpm). Next, 20 µL of Ad-TFP (10 mg/mL DMSO) was added. After allowing it to react in a shaking water bath for 1 h at 37 °C, the reaction mixture was washed 2 times with PBS by 2 centrifugation steps (3 min, 1,200 rpm) and the pellet was dissolved in 1 mL PBS. Of this solution, 0.1 mL was added to 0.9 mL of FCS and was shaken in a water bath at 37 °C up to 44 h. At 2, 20, and 44 h after incubation 0.1 mL samples were taken and their composition analysed by PD-10 SEC.

Stability of Cy5_{0.5}CD₁₀PIBMA₃₉ and MAA-Ad complexes in FCS

Mixtures of either MAA-Ad (0.2 mg/mL) with ^{99m}Tc-Cy5_{0.5}CD₁₀PIBMA₃₉ (10 µg/mL, 1 MBq) or ^{99m}Tc-MAA-Ad (0.2 mg/mL, 1 MBq) with Cy5_{0.5}CD₁₀PIBMA₃₉ (10 µg/mL) were prepared in 0.2 mL PBS and the solutions were incubated for 1 h in a shaking water bath at 37 °C. Thereafter, the formed complexes were washed twice with PBS by centrifugation (5 min, 3,000 g) and resuspended in 0.2 mL PBS. Subsequently, 0.1 mL thereof was mixed with 1 mL FCS and shaken at 37 °C in a shaking water bath up to 44 h. At 2, 20, and 44 h after incubation 0.1 mL samples were taken and diluted in 1 mL of PBS and after spinning for 5 min at 7,000 rpm, the decay corrected radioactivity of the pellet and supernatant was measured in a dose-calibrator. Hereby a reduction, in the radioactivity of the pellet represents dissociation or instability (% of binding).

Results



Figure S1. A) NMR of Ad-TFP measured in CDCL₃. B) Mass spectra of Ad-TFP, only low signals could be obtained as the compound is hard to ionize. Signals corresponding to the mass matrix are therefore clearly visible as well.

To quantify the difference of ^{99m}Tc-Cy5_{0.5}CD₁₀PIBMA₃₉ accumulation after pre-administration of nothing, MAA or MAA-Ad for all the investigated organs (Table 1 and Table S1), the relative increase of ^{99m}Tc-Cy5_{0.5}CD₁₀PIBMA₃₉ accumulation with regard to the ^{99m}Tc-Cy5_{0.5}CD₁₀PIBMA₃₉ reference distribution (Figure 3) was calculated (Figure S2). If there was a significant increase with p < 0.01 this was indicated with a *. When MAA or MAA-Ad was administered i.v. the uptake in the lungs was found highest (but this difference was not significant due to large variations). While locally administered MAA or MAA-Ad resulted in significant increases in spleen, liver and kidneys; preadministration of MAA or MAA-Ad was performed via the spleen. With the i.v. pre-administration method (Model I) the polymer accumulation increased in more organs compared to the local preadministration method (Model II), underlining once more the fact that the system works best for the clinically more relevant model i.e. local administration. The significance data can be slightly misleading since increases from e.g. 0.1 %ID/g to 0.3 %ID/g in the brain will be displayed as significant (Table 1 and Table S1).

Table S1. The biodistribution of 99m Tc-Cy5_{0.5}CD₁₀PIBMA₃₉ following injection of: none (reference distribution), MAA, or MAA-Ad and the biodistribution of 99m Tc-MAA-Ad administered via Models I and II. Data (expressed as the mean ± SD of the percentage of the injected dose per gram tissue (%ID/g) of 5 observations) are calculated from radioactivity counts in various tissues at 2 h post-injection of the tracer.

	Reference distribution host	Distribution of host (^{99m} Tc-Cy5 _{0.5} CD ₁₀ PIBMA ₃₉) following Reference distribution of indicated guest					bution guest
Tissue	^{99m} TC- Cy5 _{0.5} CD ₁₀ PIBMA	Model I: MAA	Model I: MAA-Ad	Model II: MAA	Model II: MAA-Ad	Model I: ^{99m} Tc-MAA-Ad mean	Model II: ^{99m} Tc-MAA-Ad mean
	mean	Mean	Mean	Mean	mean		
Salivary gland	3.5 ± 0.4	9.1 ± 3.3	17.3 ± 3.5	3.0 ± 0.6	3.9 ± 0.1	N.A.	8.9 ± 1.9
Stomach	4.4 ± 0.8	19.8 ± 7.6	10.8 ± 1.4	3.0 ± 0.6	2.3 ± 0.8	11.5 ± 9.1	12.7 ± 3.1
Intestines	0.8 ± 0.3	1.5 ± 0.6	1.5 ± 0.3	0.9 ± 0.0	0.7 ± 0.3	0.6 ± 0.3	1.8 ± 0.5



Figure S2. Relative increase of ^{99m}Tc-Cy5_{0.5}CD₁₀PIBMA₃₉ after i.v. (A) or Local (B) administration of MAA or MAA-Ad with respect to ^{99m}Tc-Cy5_{0.5}CD₁₀PIBMA₃₉ accumulation in the indicated organ when no particles are preadministered. With significance of difference (p < 0.01) indicated by *.

Both the individual components and the complexes formed demonstrated a high serum stability (Figure S3). No clear metabolites of the individual components could be defined. This said some dissociation of ^{99m}Tc was observed both from the individual components as from the complexes formed. Nevertheless, the complex yielded around a 80% stability at 44 h.

As the Cy5_{0.5}CD₁₀PIBMA₃₉ polymer was not optimized for ^{99m}Tc chelation, but merely provided coordination sides by its free –COOH moieties, some dissociation of ^{99m}Tc was observed *in vivo* (Table S1, Figure S3). When this occurred, characteristic uptake in the salivary glands and stomach could be observed. As these findings did not complicate the assessment of the pre-targeting ability, no attempts were made to optimize the chelation stability.











C Stability of complexed ${}^{99m}\text{Tc-Cy5}_{0.5}\text{CD}_{10}\text{PIBMA}_{39}$ with MAA-Ad

D Stability of complexed Cy50.5CD10PIBMA39 with 99mTc-MAA-Ad



Figure S3 A) Serum stability of ^{99m}Tc-Cy5_{0.5}CD₁₀PIBMA₃₉ (peak around fraction 18 indicated smaller fragments), B) Serum stability of ^{99m}Tc-MAA (peak around fraction 22 indicated some free ^{99m}Tc). C) Serum stability of [^{99m}Tc-Cy5_{0.5}CD₁₀PIBMA₃₉ * MAA-Ad] complexes, D) Serum stability of [Cy5_{0.5}CD₁₀PIBMA₃₉ * ^{99m}Tc-MAA-Ad] complexes.