Table S1: Enhancement of the oncolytic activity of magnetic viral complexes relative to naked virus and the influence of magnetic field and FCS. Data are graphically presented in figure 9C.

Sample	magnet	ic field	no magnetic field						
odinpic	7.5 % FCS	50 % FCS	7.5 % FCS	50 % FCS					
	IC ₅₀ virus /IC ₅₀ MNP-VP in McA cells								
vsv	1	1	1	1					
PEI-Mag2-VSV	2.5	7.9	1.6	6.0					
SO-Mag6-11.5-VSV	1.6	7.0	1.8	5.7					
	IC ₅₀ virus /IC ₅₀ MNP-VP in RDB cells								
Ad	1	1	1	1					
PEI-Mag2-Ad	11.8	27.8	1.2	3.8					
SO-Mag6-11.5-Ad	27.5	43.6	2.8	4.5					

Table S2: $\mathbf{r_2}$ and $\mathbf{r_2}^*$ relaxivities of the MNP assemblies. The table shows the $\mathbf{r_2}$ and $\mathbf{r_2}^*$ relaxivity data of the free MNPs and all magnetic viral complex assemblies in liver-mimicking agarose phantoms, and the ratios of the nanoassembly relaxivity normalized to the relaxivity of free MNPs ($\mathbf{r_2}/\mathbf{r_2}$ (MNP) and $\mathbf{r_2}^*/\mathbf{r_2}^*$ (MNP), respectively, green) and the $\mathbf{r_2}^*/\mathbf{r_2}$ ratios (red). All measurements were performed using a clinical 3 T MRI system from Philips.

	PEI-Mag2				SO-Mag6-11.5					
Sample	r ₂	ratio	r ₂ *	ratio	r ₂ */r ₂	r ₂	ratio	r ₂ *	ratio	r ₂ */r ₂
	[mM ⁻¹ Fe s ⁻¹]	[%]	[mM ⁻¹ Fe s ⁻¹]	[%]		[mM ⁻¹ Fe s ⁻¹]	[%]	[mM ⁻¹ Fe s ⁻¹]	[%]	
MNP	238 ± 11	100	476 ± 58	100	2.0	274 ± 10	100	857 ± 12	100	3.1
MNP in McA	55 ± 5 (4.5 pg Fe/cell)	23	584 ± 17 (4.5 pg Fe/cell)	123	10.6	37 ± 3 (5.4 pg Fe/cell)	14	788 ± 12 (5.4 pg Fe/cell)	92	21.3
MNP-VSV MNP-VSV in McA	23 ± 3 (3.2 pg	10	518 ± 8 (3.2 pg	109	22.5	57 ± 2 (2.5 pg	21	1494 ± 42 (2.5 pg	174	26.2
MNP in RDB	Fe/cell) 19 ± 1 (17.3 pg Fe/cell)	8	Fe/cell) 560 ± 23 (17.3 pg Fe/cell)	118	29.5	Fe/cell) 38 ± 2 (15.9 pg Fe/cell)	14	Fe/cell) 1203 ± 118 (15.9 pg Fe/cell)	140	31.7
MNP-Ad	124 ± 13	52	542 ± 19	114	4.4	18 ± 2	7	1004 ± 32	117	55.8
MNP-Ad in RDB	21 ± 3 (6.7 pg Fe/cell)	9	757 ± 22 (6.7 pg Fe/cell)	159	36.0	8 ± 6 (7.4 pg Fe/cell)	3	412 ± 26 (7.4 pg Fe/cell)	48	51.5

Figure S1: $\mathbf{r_2}$ and $\mathbf{r_2}^*$ relaxivity plots. The panels **E** and **F** are overlays of the panels in figure 8 to demonstrate the comparability of the data acquired on clinical 1.5 T and 3 T systems. Panel **E** is the overlay of the panels **A** and **C**, the transverse relaxation and panel **F** is the overlay of the spin-spin relaxation in panels **B** and **D** of figure 8. Free MNPs are plotted in black, cell internalized MNPs in red, and free complexes in blue and cell internalized complexes in purple.

