

Supplementary Materials: Amphiphilic Anionic Oligomer-Stabilized Calcium Phosphate Nanoparticles with Prospects in siRNA Delivery via Convection-Enhanced Delivery

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Structural analysis of pristine oligomer

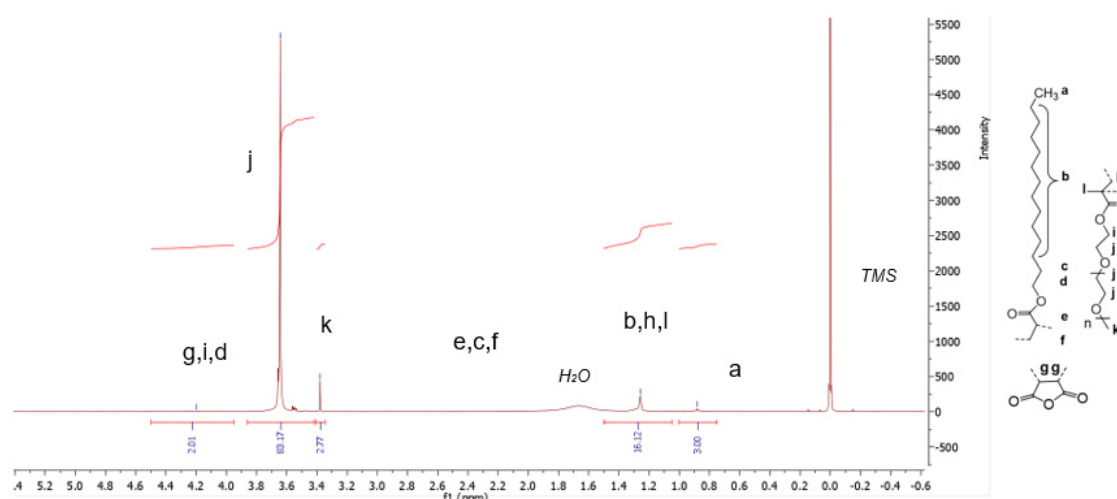


Figure S1. ^1H -NMR of o14PEGMA (04/04/10) in CDCl_3 with 0.03% TMS, $c = 0.5 \text{ mg/ml}$
 δ [ppm] 0.75-1.00 ; 1.05-1.50; 3.35-3.40; 3.42-3.86; 3.95-4.50

Size exclusion chromatography of pristine oligomer stabilizer

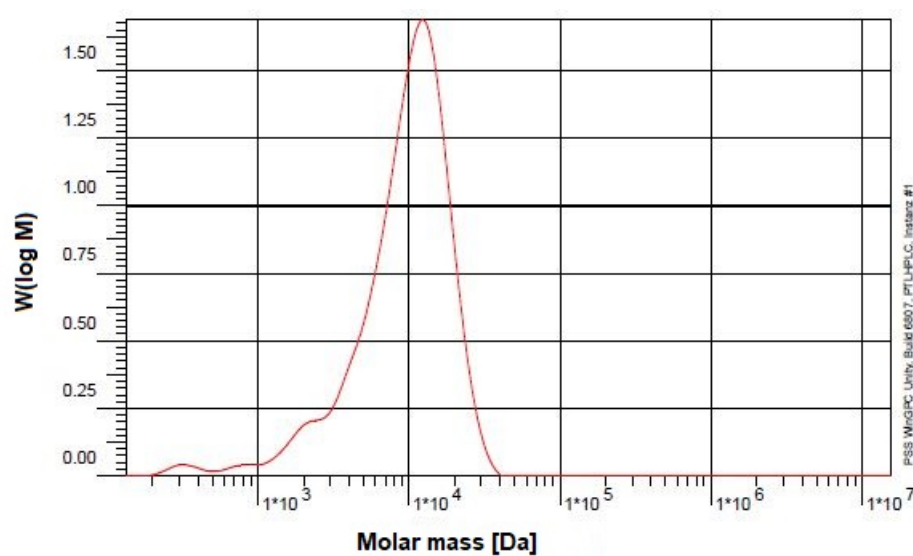
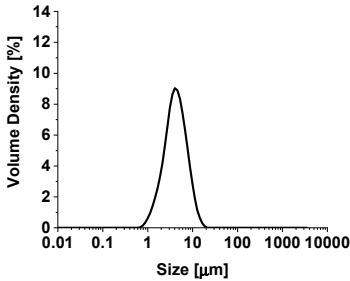
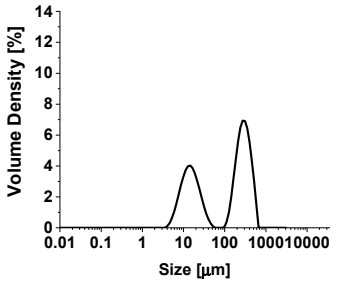
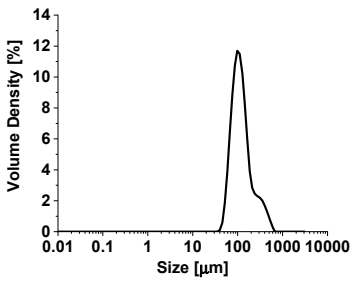
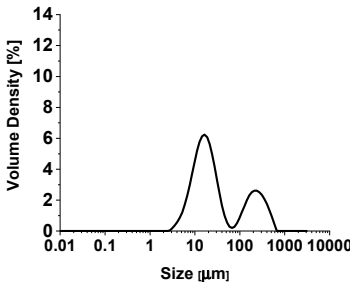
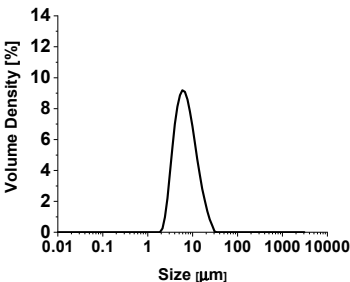
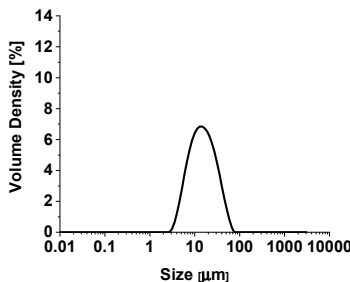
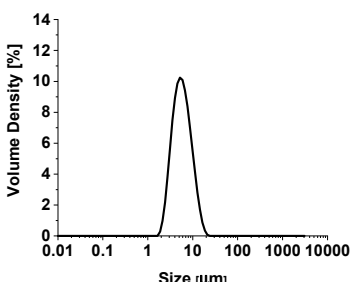
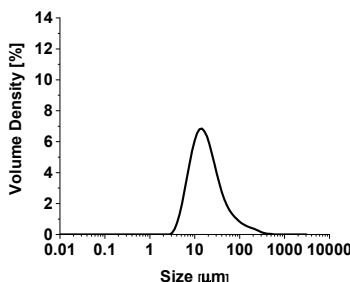
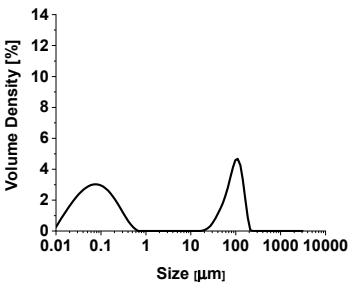
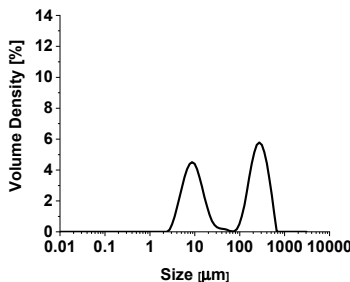
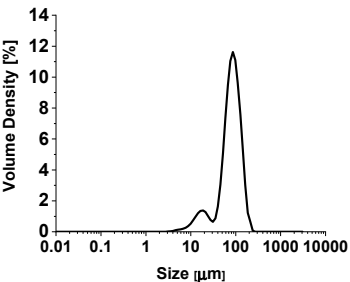
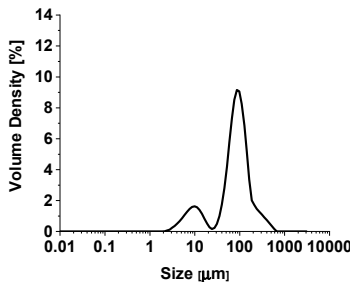
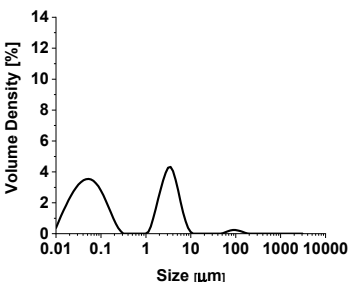
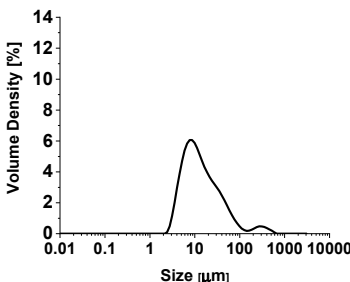
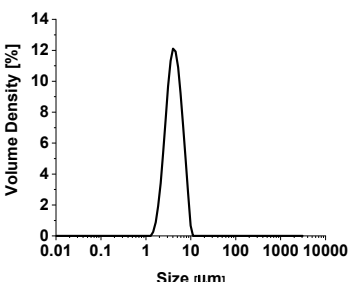
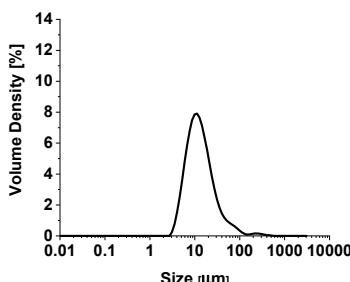


Figure S2. Gel permeation chromatography chromatograms of o14PEGMA(1:1:2.5) to determine size distribution
c = 10mg/ml in THF ; V = 40 μ l; flow rate=1ml/min; n=3

Impact of oligomer concentration on CaP-NP stabilization.

			Size distribution	
			t = 5 min	t = 30 min
2 $\mu\text{mol/L}$ o14PEGMA(1:1:2.5)_NH ₃	1.5 mmol/L PO ₄ ³⁻	Method 1	 <p>D[4;3] = 5.05 μm</p>	 <p>D[4;3] = 194 μm</p>
		Method 2	 <p>D[4;3] = 151 μm</p>	 <p>D[4;3] = 88.4 μm</p>
	6.0 mmol/L PO ₄ ³⁻	Method 1	 <p>D[4;3] = 8.45 μm</p>	 <p>D[4;3] = 19.1 μm</p>
		Method 2	 <p>D[4;3] = 6.72 μm</p>	 <p>D[4;3] = 29.8 μm</p>

			Size distribution	
			t = 5 min	t = 30 min
5 $\mu\text{mol/L}$ o14PEGMA(1:1:2.5)_NH ₃	1.5 mmol/L PO ₄ ³⁻	Method 1	 D[4;3] = 39.6 μm	 D[4;3] = 166 μm
		Method 2	 D[4;3] = 89.2 μm	 D[4;3] = 104 μm
	6.0 mmol/L PO ₄ ³⁻	Method 1	 D[4;3] = 3.21 μm	 D[4;3] = 31.4 μm
		Method 2	 D[4;3] = 4.77 μm	 D[4;3] = 20.6 μm

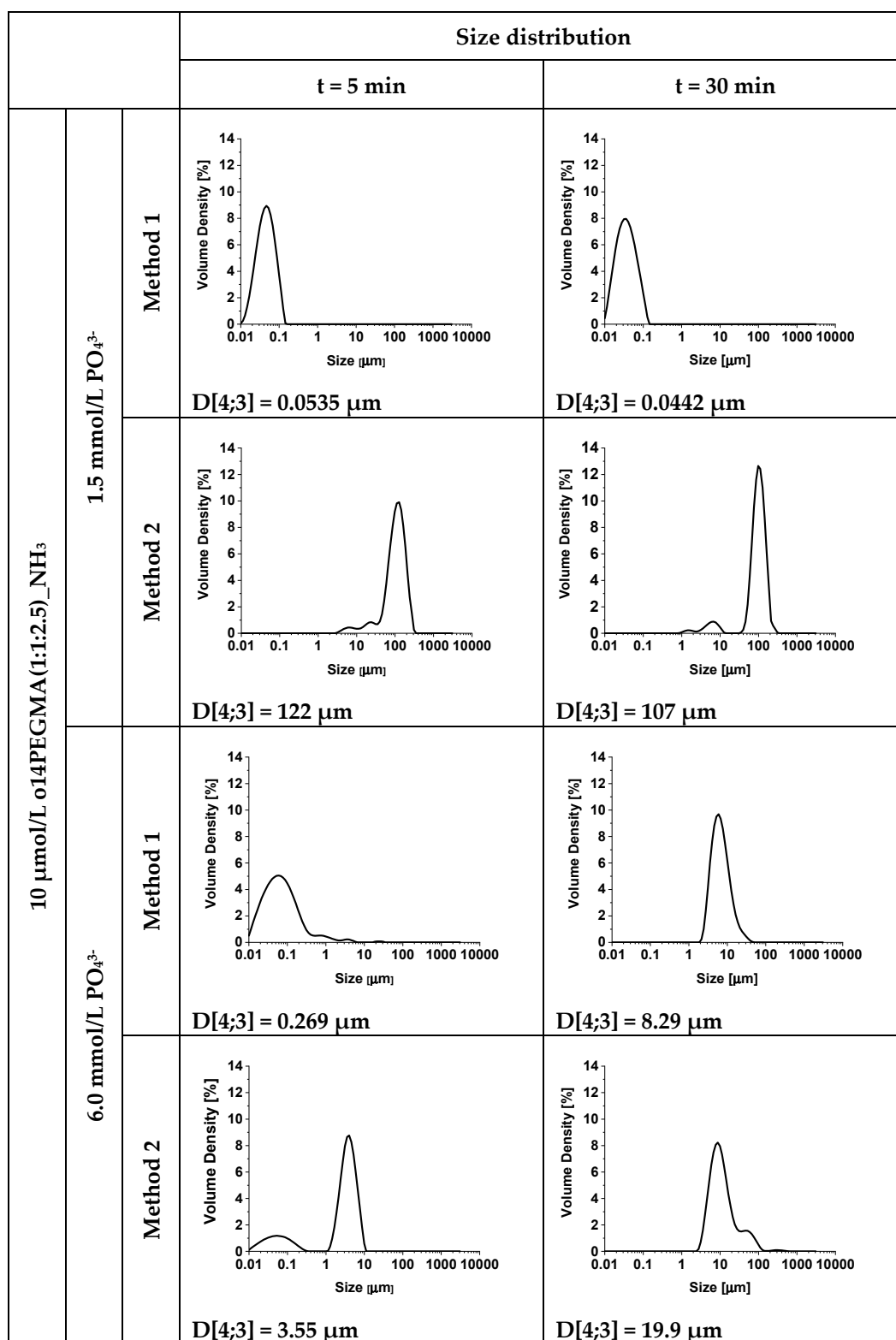
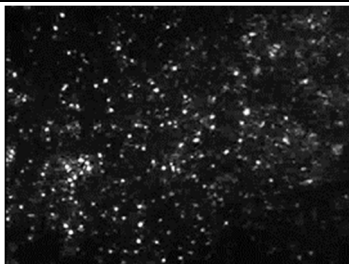
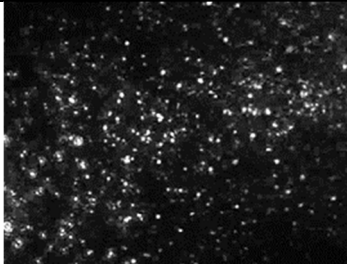
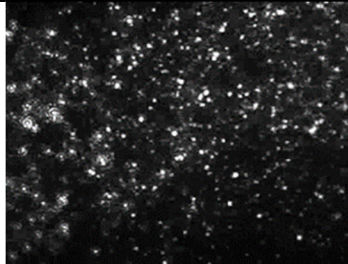
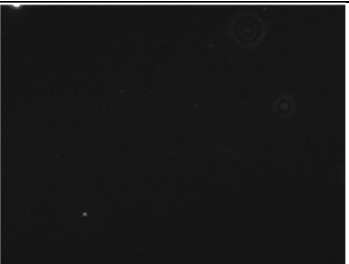
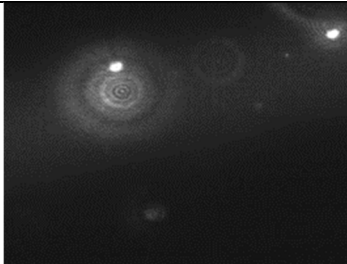
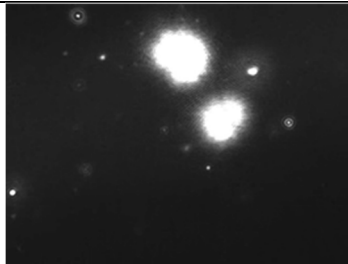
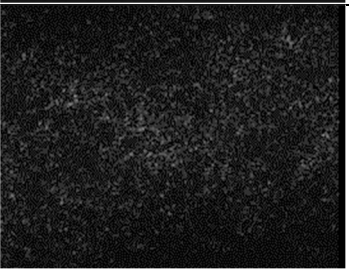
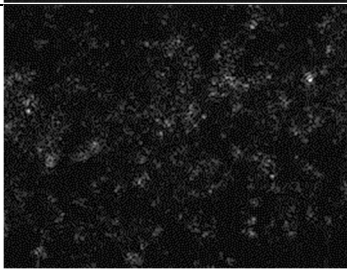
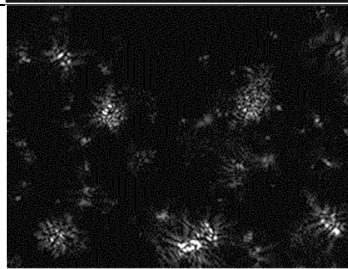
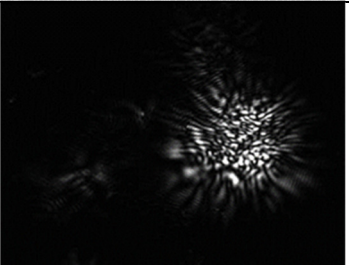
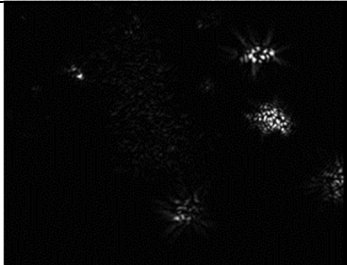
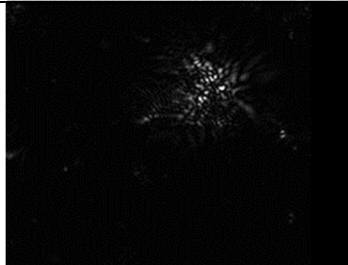
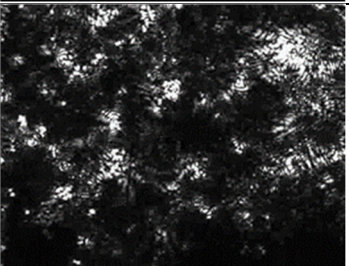
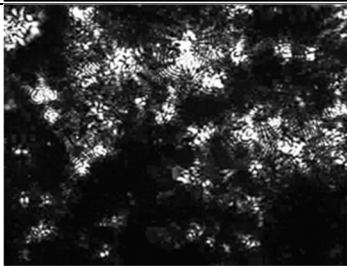
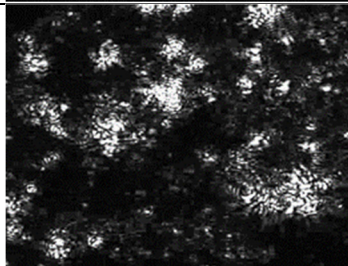
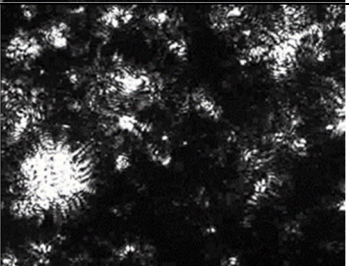
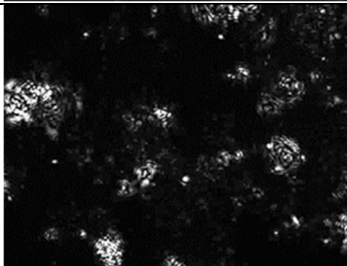
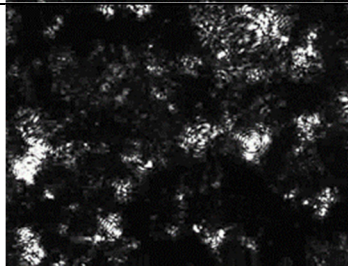


Figure S3. Influence of oligomer concentration on size distribution and aggregation of CaP particles determined by Laser diffraction analysis.

NTA screenshots for analysis of aggregation status

10 μmol/L o14PEGMA(1:1:2.5) _NH ₃			0 μmol/L siRNA	1 μmol/L siRNA	10 μmol/L siRNA
1.5 mmol/L PO ₄ ³⁻	Method 1				
	Method 2				
3.75 mmol/L PO ₄ ³⁻	Method 1				
	Method 2				
6.0 mmol/L PO ₄ ³⁻	Method 1				
	Method 2				
			0 μmol/L siRNA	1 μmol/L siRNA	10 μmol/L iRNA

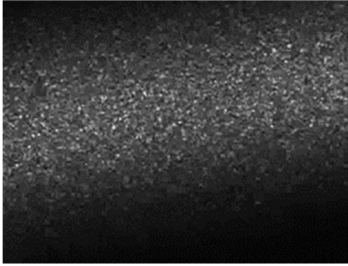
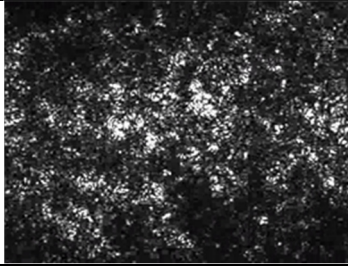

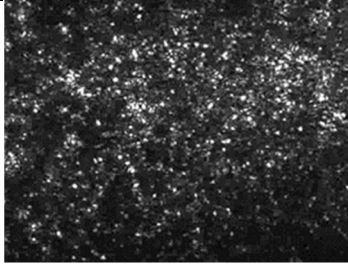
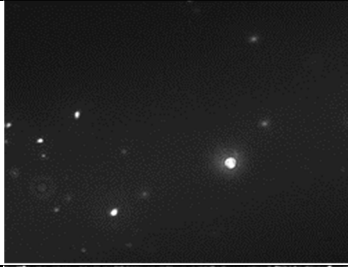
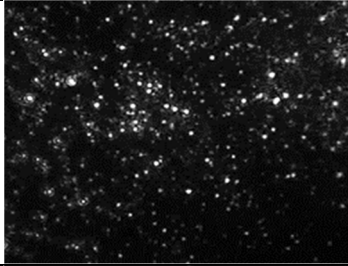
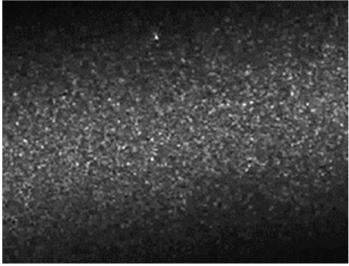
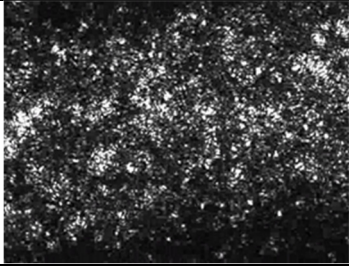
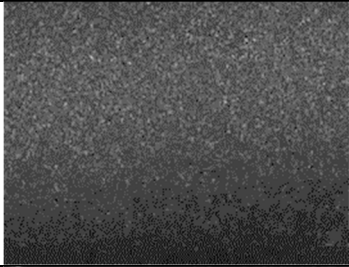
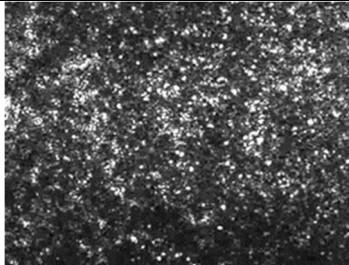

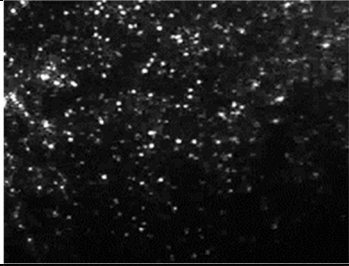
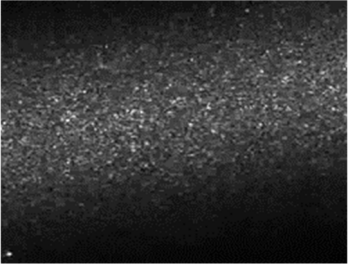
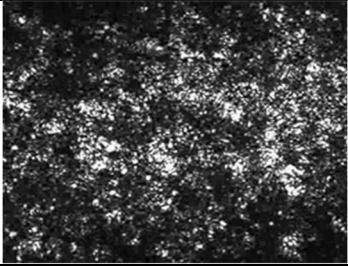
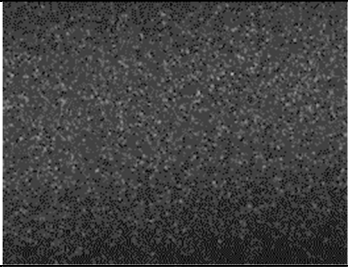
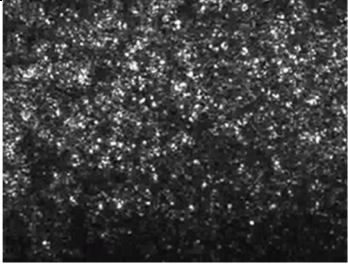

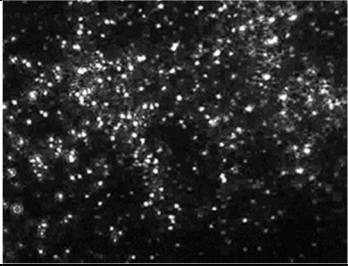
40 $\mu\text{mol/L}$ o14PEGMA(1:1:2.5)_NH ₃					
6.0 mmol/L PO ₄ ³⁻		3.75 mmol/L PO ₄ ³⁻		1.5 mmol/L PO ₄ ³⁻	
Method 2	Method 1	Method 2	Method 1	Method 2	Method 1
					
					
					

Figure S4. NTA screenshots of CaP-NP formulation 30 min after fabrication.

Impact of cell culture medium & serum proteins on size distribution of CaP-NP

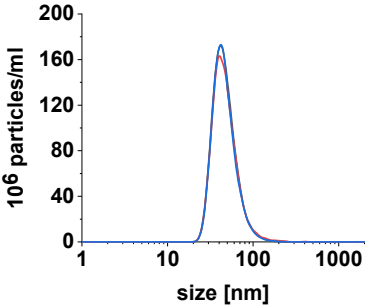
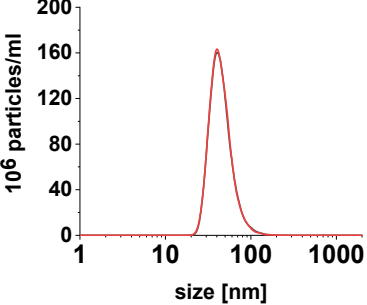


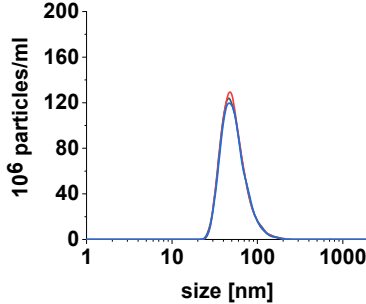
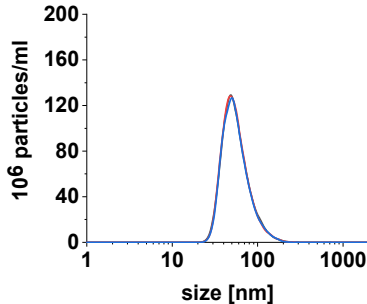


		0 $\mu\text{mol/L}$ siRNA	1 $\mu\text{mol/L}$ siRNA
Oligomer/ PO_4^{3-} -concentration/Method	40 $\mu\text{mol/L}$ /3.75 mmol/L $\text{PO}_4^{3-}/2$	 <p>Mean: 55 ± 22 nm Concentration: $57.8 \pm 1.1 \cdot 10^8/\text{ml}$</p>	 <p>Mean: 52 ± 19 nm Concentration: $49.7 \pm 0.2 \cdot 10^8/\text{ml}$</p>
			
	40 $\mu\text{mol/L}$ /6.0 mmol/L $\text{PO}_4^{3-}/2$	 <p>Mean: 63 ± 27 nm Concentration: $51.8 \pm 1.0 \cdot 10^8/\text{ml}$</p>	 <p>Mean: 69 ± 32 nm Concentration: $58.6 \pm 0.9 \cdot 10^8/\text{ml}$</p>
			

Figure S5. Effect of serum proteins and cell culture medium on stability of α 14PEGMA(1:1:2.5) $_{\text{NH}_3}$ -stabilized CaP-NP after 5 h.

Cellular Uptake of CaP-NP in F98 cells

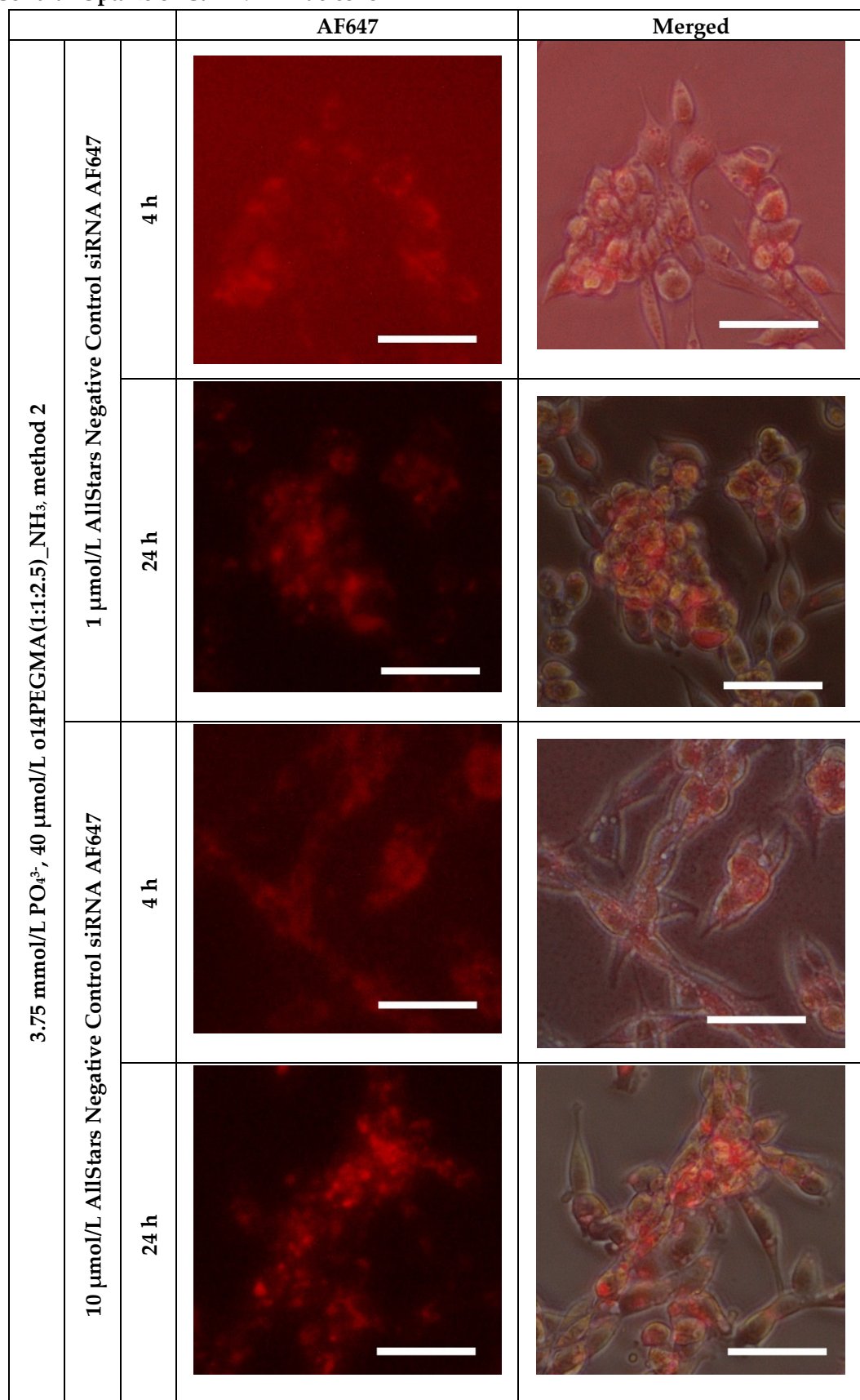
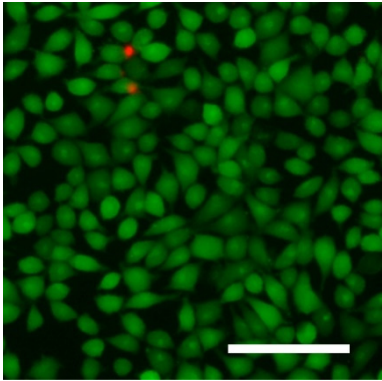
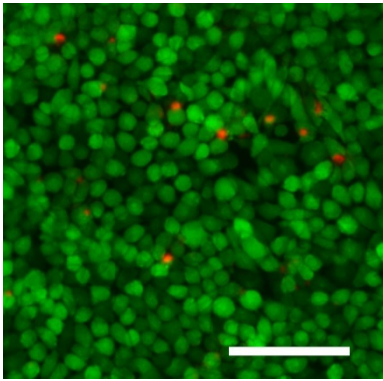
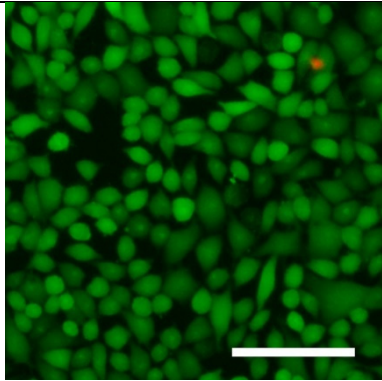
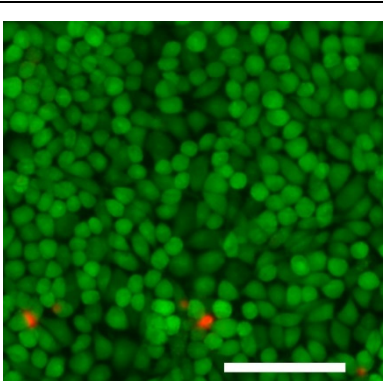
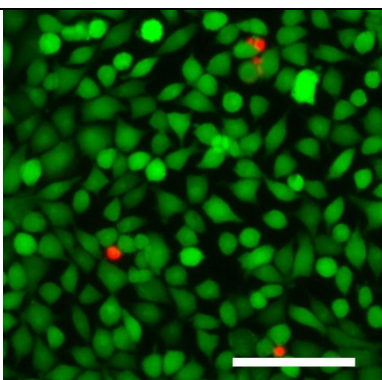
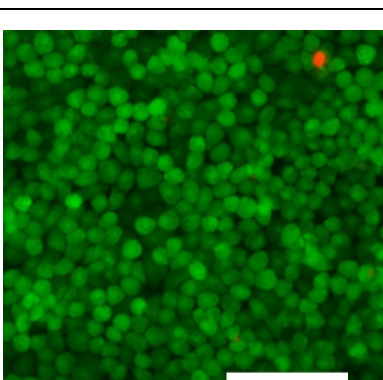






Figure S6. Uptake of siRNA (AlexaFluor™ 647 labeled)-loaded o14PEGMA(1:1:2.5) $_{\text{NH}_3}$ -stabilized CaP-NP in F98 cells. Scale bars represent 100 μm .

Live/Dead staining

	Group		24 h	72 h
Non-concentrated	Control	Untreated		
				
	3.75 mmol/L PO ₄ ³⁻ , 40 μM o14PEGMA(1:1:2.5)_NH ₃ , method 2	0 μmol/L siRNA		
		1 μmol/L siRNA		
				

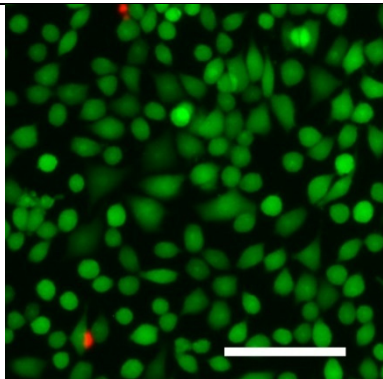
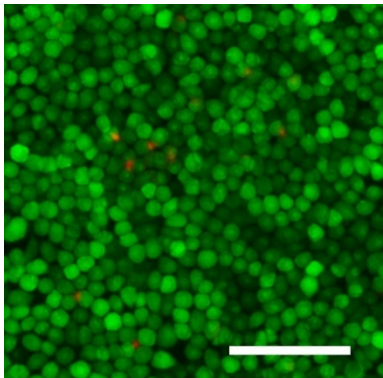
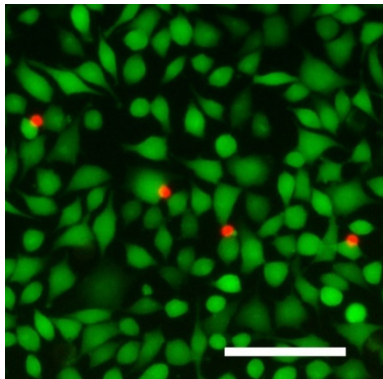
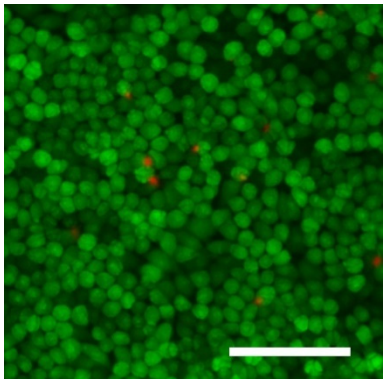
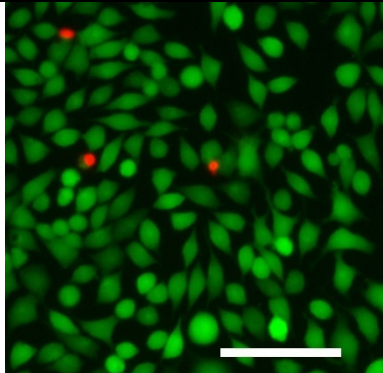
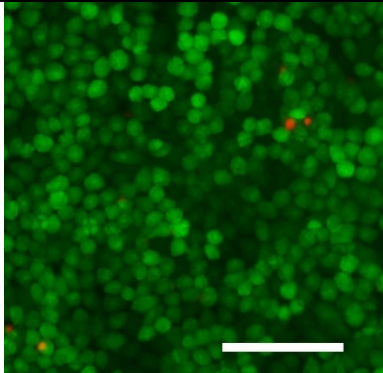
	Group		24 h	72 h
	Concentrated	Control		
		Untreated		
	3.75 mmol/L PO ₄ ³⁻ , 40 μM o14PEGMA(1:1:2.5)_NH ₃ , method 2	0 μmol/L siRNA		
		10 μmol/L siRNA		

Figure S7. *In vitro* biocompatibility testing of CaP-NP in L929 cells: Live/Dead staining. Scale bars represent 100 μm.

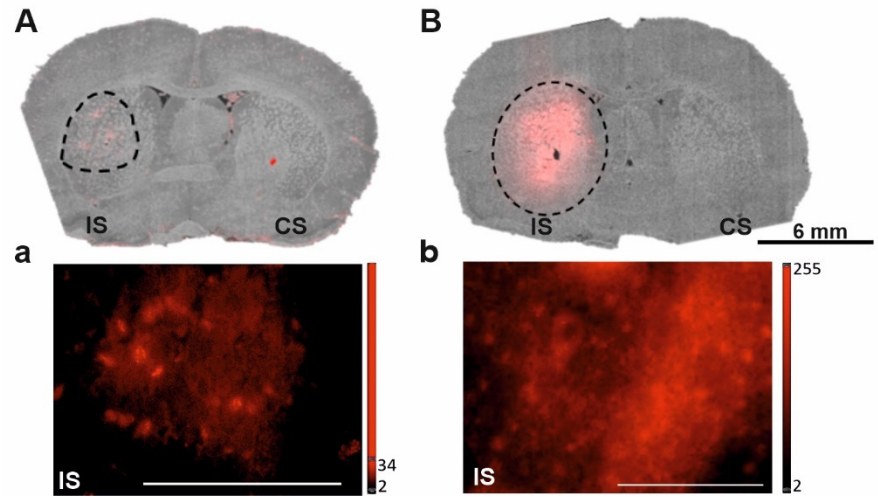


Figure S8. *In vivo* biodistribution of CaP-NP loaded with 10 $\mu\text{mol/L}$ AllStars Negative Control siRNA AF647 (A, a) or AllStars Negative Control siRNA AF647 alone (B, b) 1 hour post-CED in healthy brain ($n=2$). Representative brain sections of fluorescence distribution in the whole slide (A, B) and fluorescence of the AllStars Negative Control siRNA AF647 at the injection site within the CaP-NP (a) and alone (b). IS: ipsilateral side (CED injection), CS: contralateral side (control) (scale bars represent 150 μm , $\times 40$ magnification).