

Supplementary Material

for

Tuning the envelope structure of enzyme nanoreactors for *in vivo* detoxification of organophosphates

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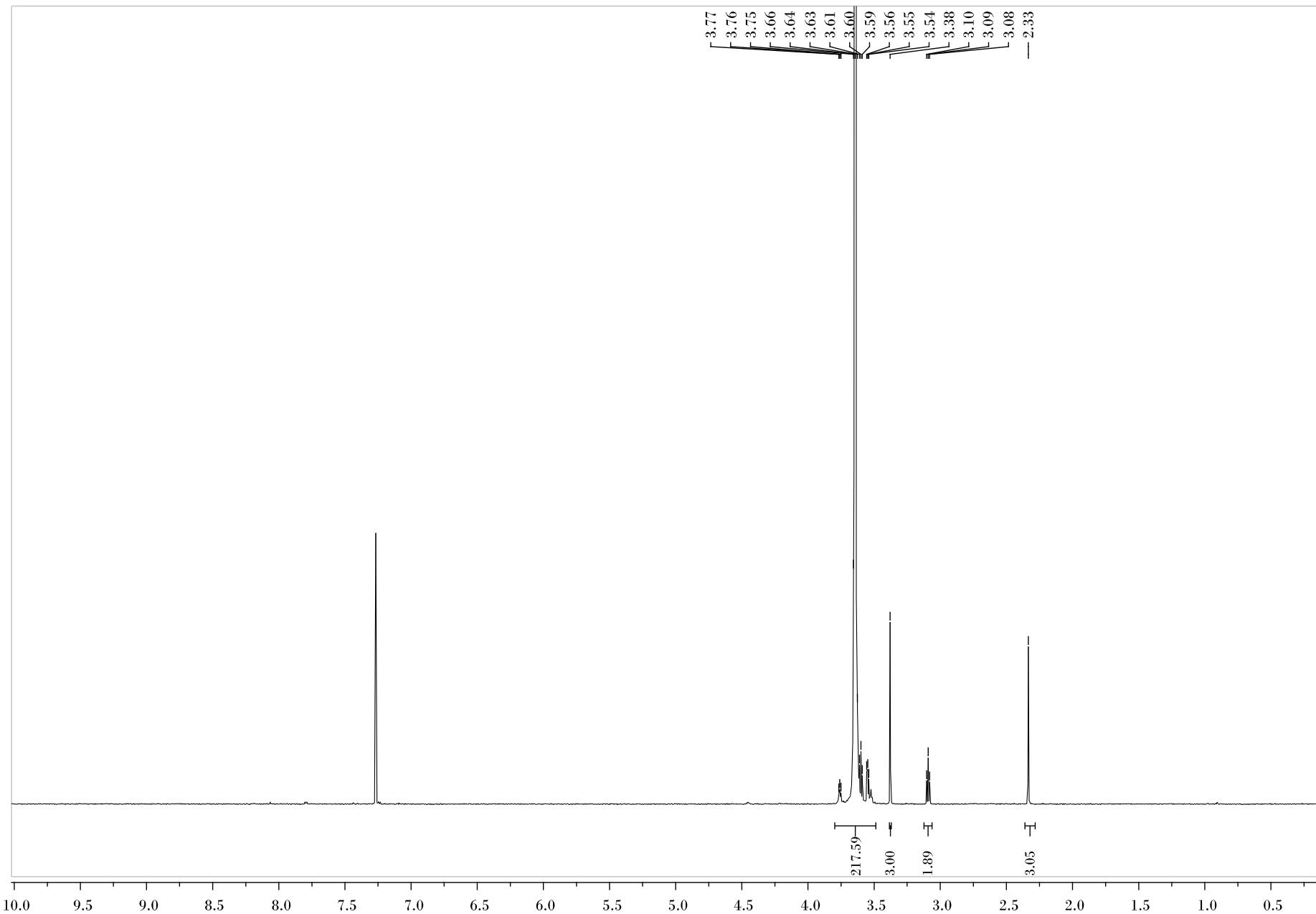


Figure S1. ¹H NMR spectra of mPEGSAc-2000

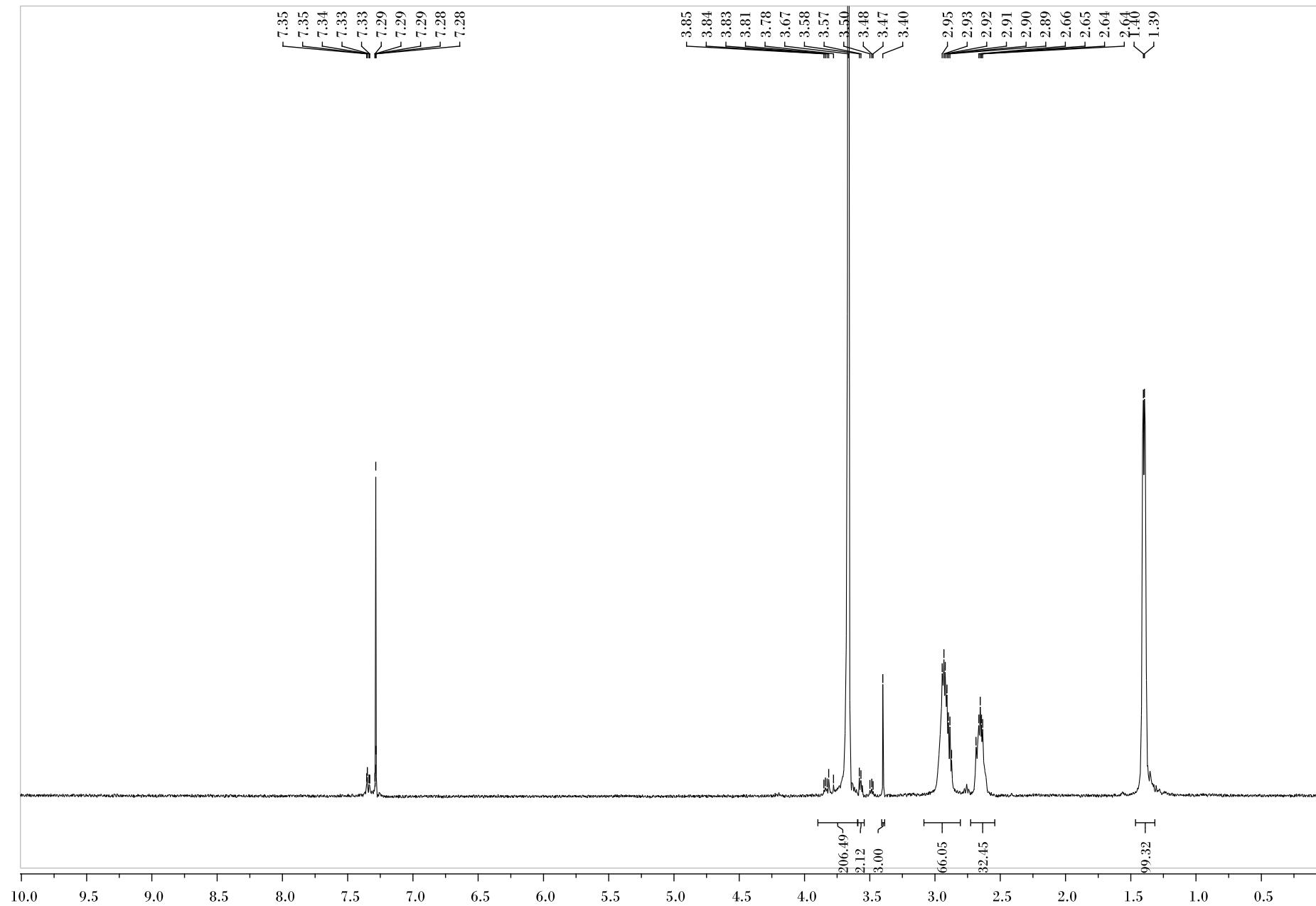


Figure S2. ¹H NMR spectra of copolymer 1a

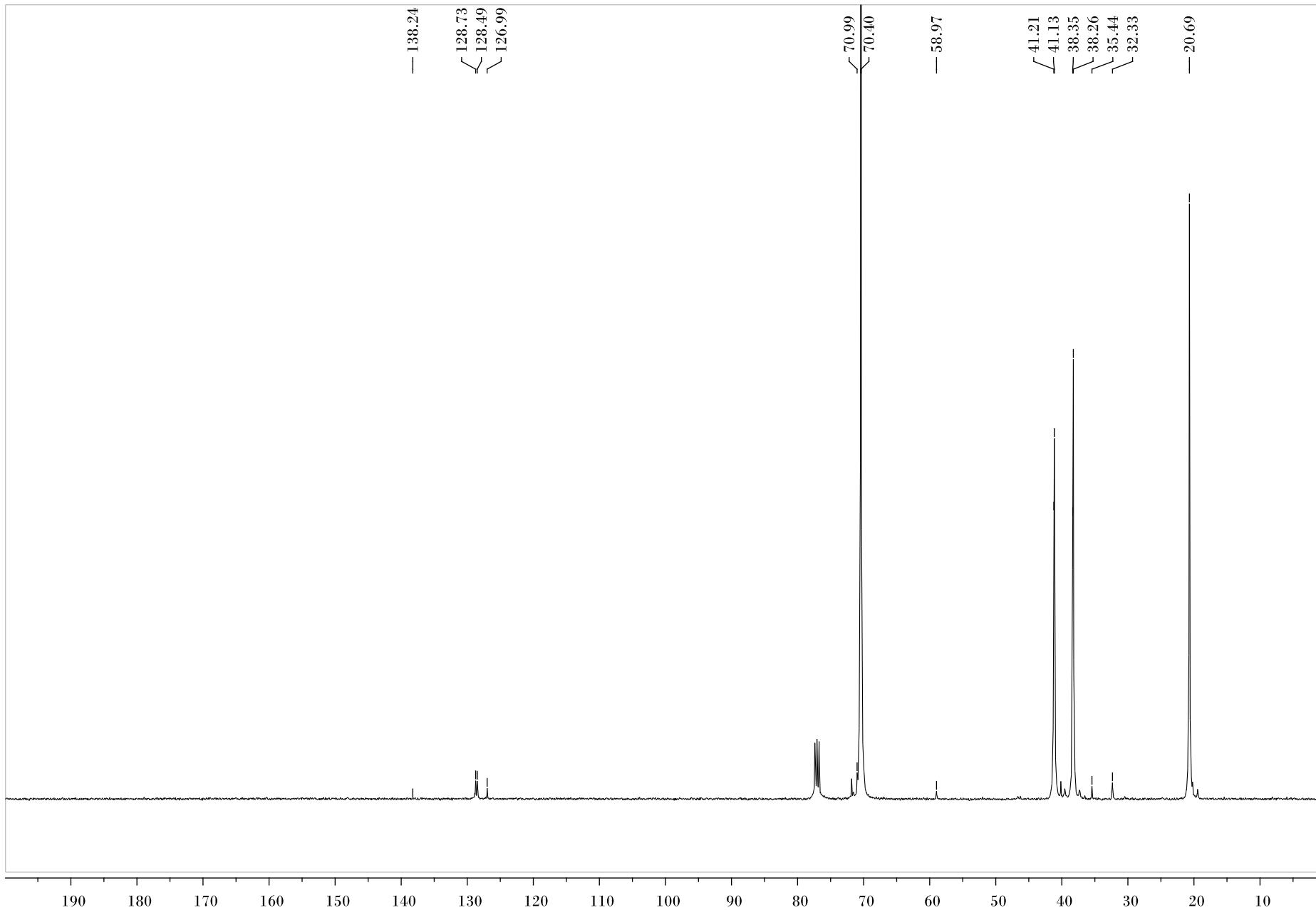


Figure S3 ^{13}C NMR spectra of copolymer **1a**

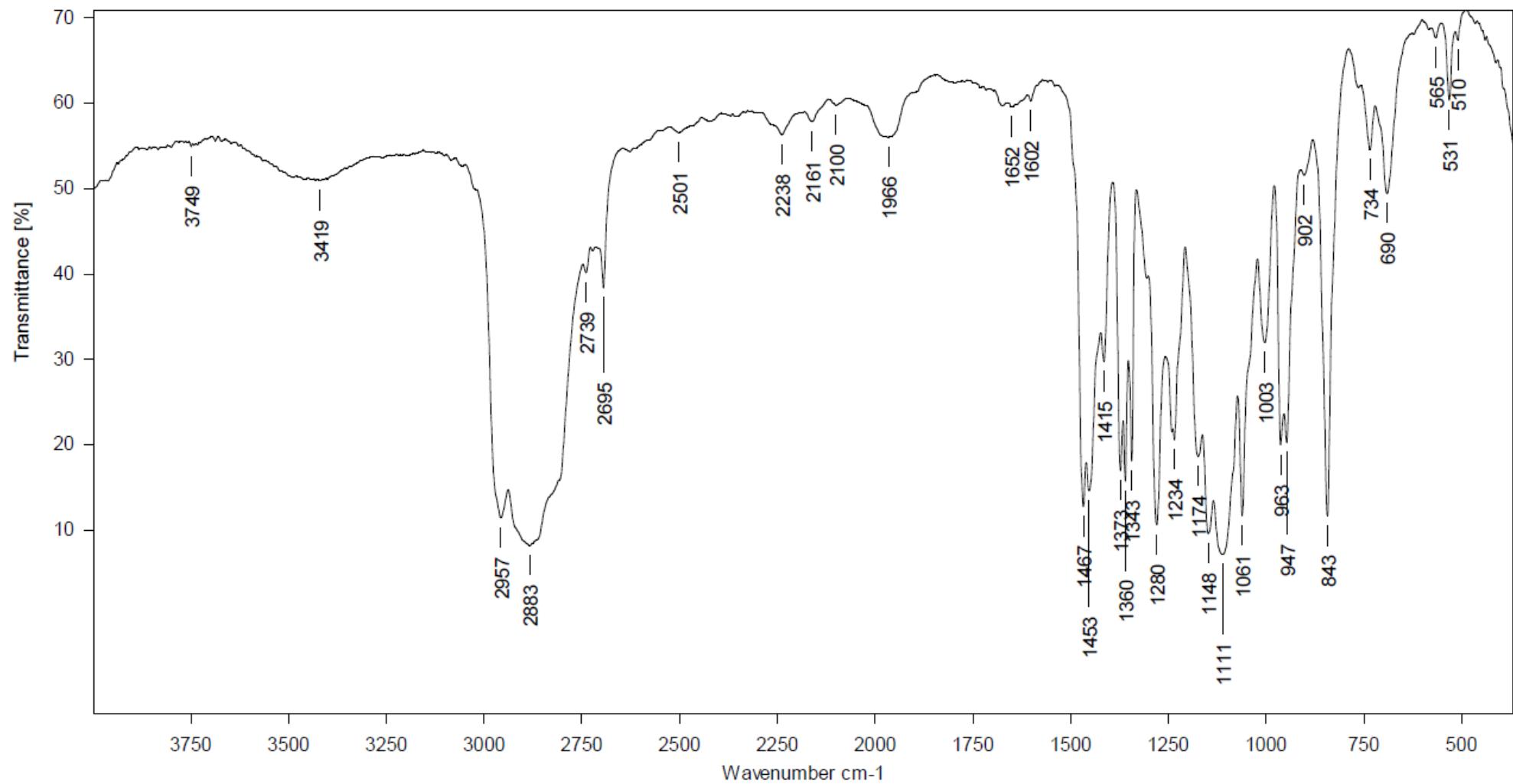


Figure S4. IR spectra of copolymer **1a**

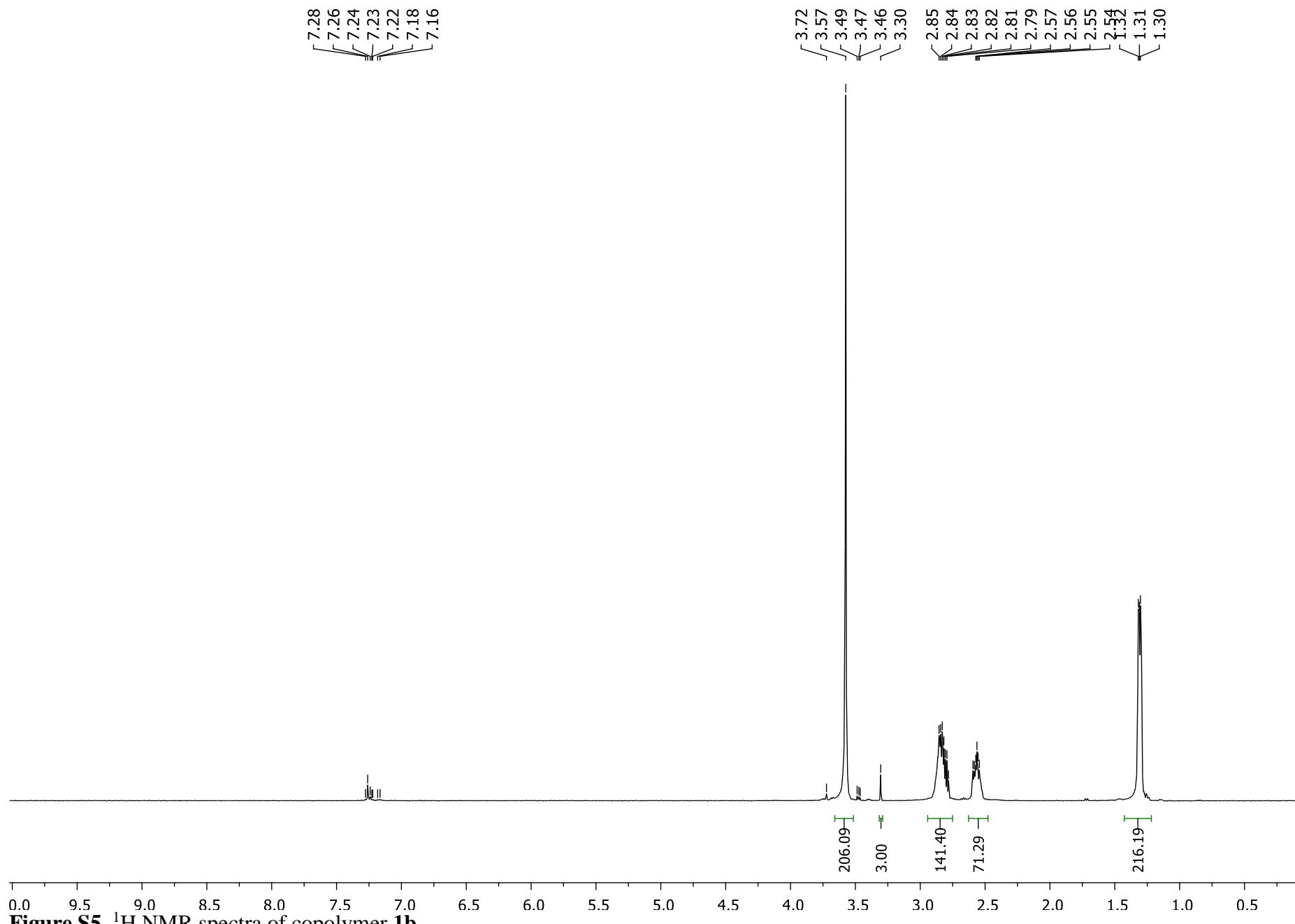


Figure S5. ^1H NMR spectra of copolymer **1b**

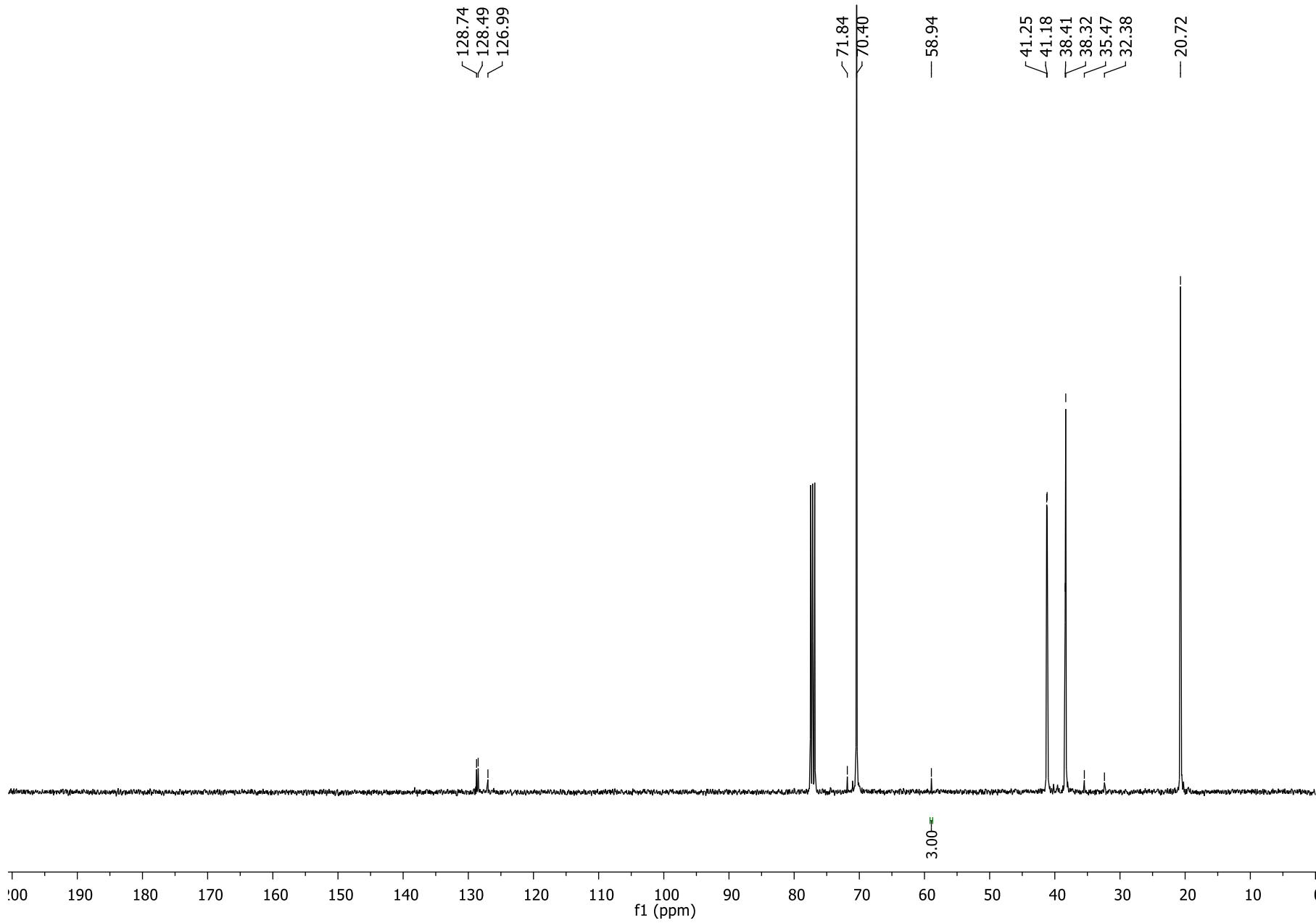


Figure S6. ¹³C NMR spectra of mPEG-PPS-mPEG copolymer **1b**

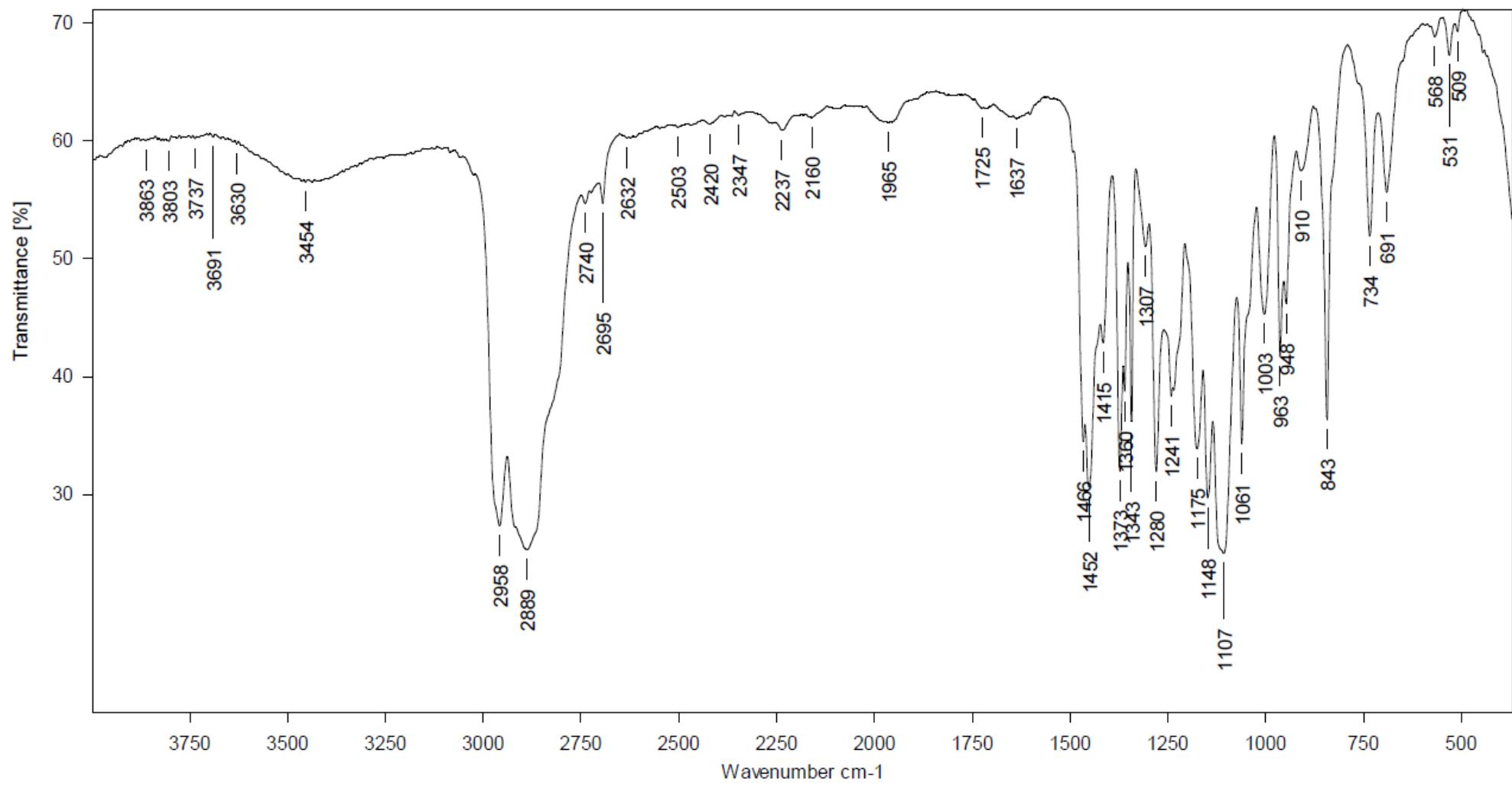


Figure S7a. IR spectra of copolymer **1b**

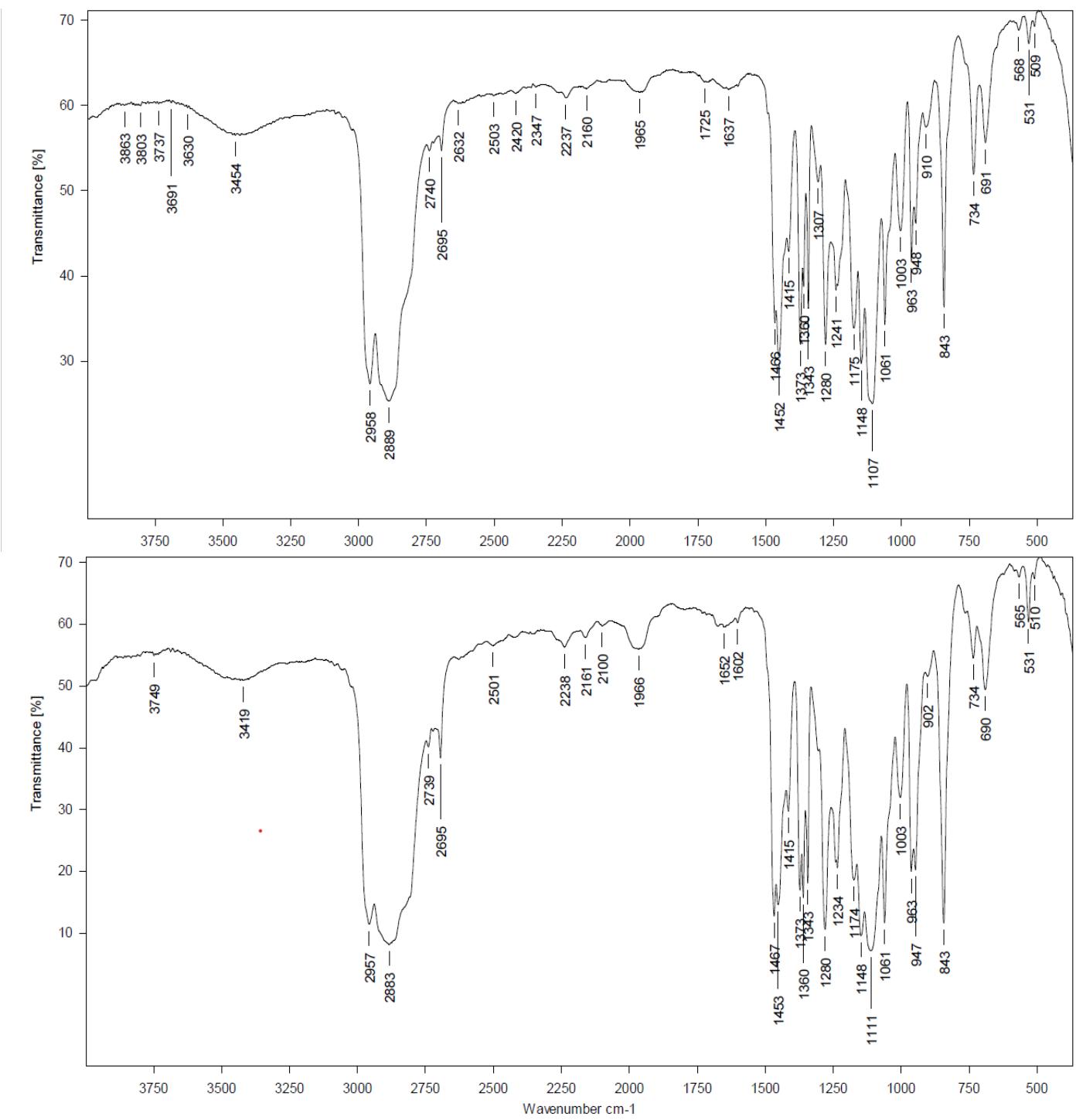


Figure S7b. Comparison of IR spectra of copolymer **1a** and **1b**

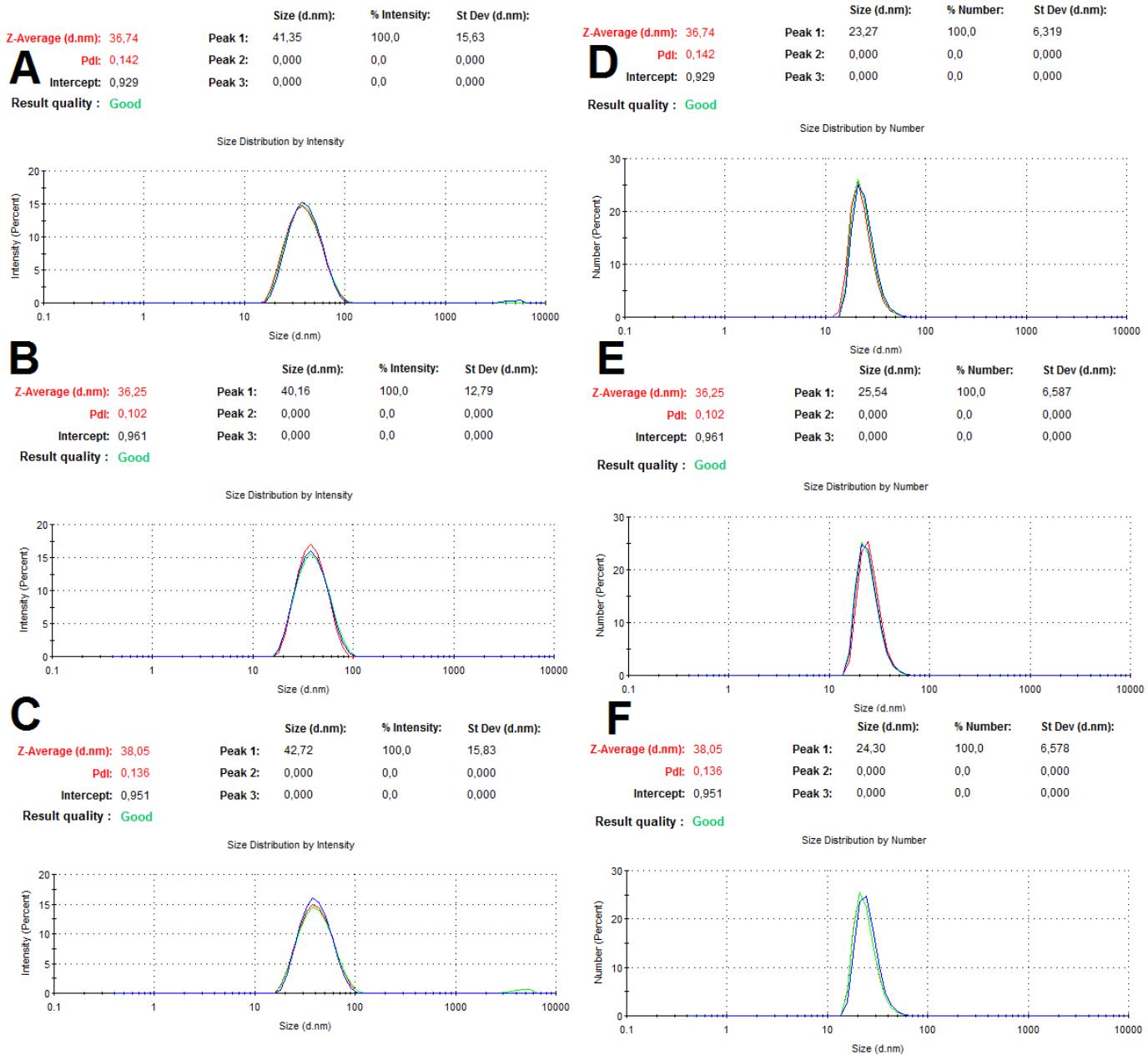
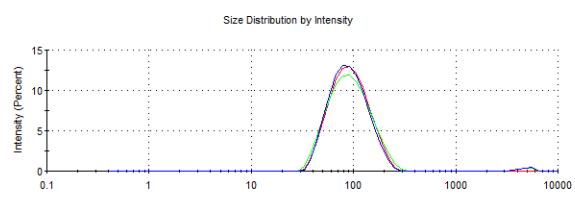


Figure S8. Screen shot of empty nanoreactor size distribution using the intensity(A-C) and number parameters (D-F) based on **1a**, copolymer concentrations 0.1, 0.2, 0.5, 0.75, 1% (wt/wt), 10 mM Tris buffer, pH=7.4, 25°C.

A

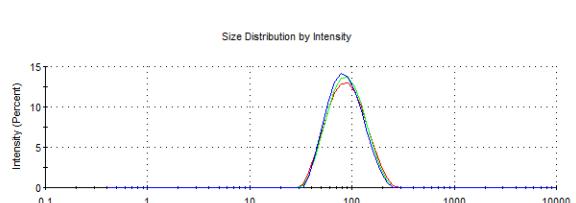
	Size (d.nm):	% Intensity:	St Dev (d.nm):
Z-Average (d.nm):	83,91	Peak 1: 98,14	100,0 41,56
Pdt:	0,157	Peak 2: 0,000	0,0 0,000
Intercept:	0,956	Peak 3: 0,000	0,0 0,000

Result quality : Good

**B**

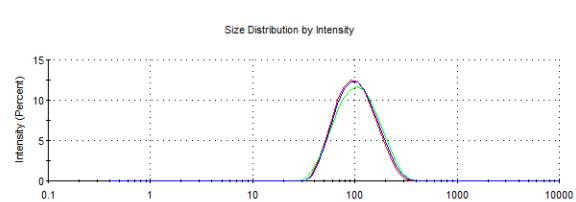
	Size (d.nm):	% Intensity:	St Dev (d.nm):
Z-Average (d.nm):	80,89	Peak 1: 94,92	100,0 39,90
Pdt:	0,135	Peak 2: 0,000	0,0 0,000
Intercept:	0,968	Peak 3: 0,000	0,0 0,000

Result quality : Good

**C**

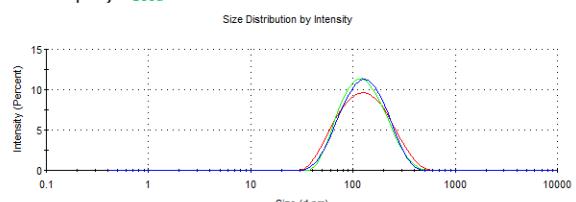
	Size (d.nm):	% Intensity:	St Dev (d.nm):
Z-Average (d.nm):	92,06	Peak 1: 109,1	100,0 48,57
Pdt:	0,164	Peak 2: 0,000	0,0 0,000
Intercept:	0,934	Peak 3: 0,000	0,0 0,000

Result quality : Good

**D**

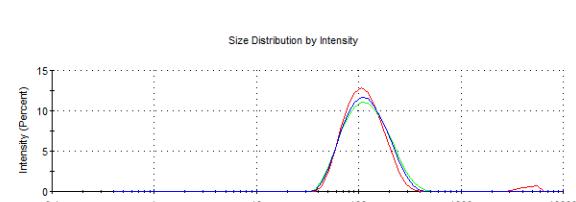
	Size (d.nm):	% Intensity:	St Dev (d.nm):
Z-Average (d.nm):	110,9	Peak 1: 146,0	100,0 81,55
Pdt:	0,211	Peak 2: 0,000	0,0 0,000
Intercept:	0,964	Peak 3: 0,000	0,0 0,000

Result quality : Good

**E**

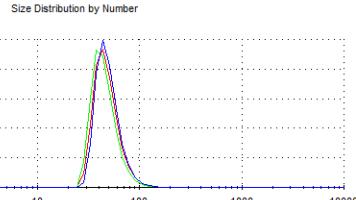
	Size (d.nm):	% Intensity:	St Dev (d.nm):
Z-Average (d.nm):	106,1	Peak 1: 120,3	97,8 51,33
Pdt:	0,191	Peak 2: 4553	2,2 849,1
Intercept:	0,951	Peak 3: 0,000	0,0 0,000

Result quality : Good

**F**

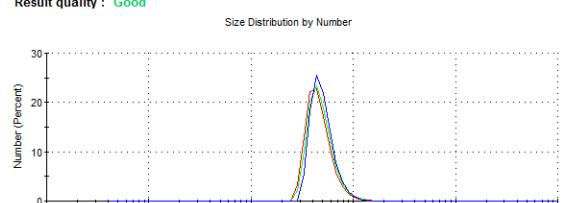
	Size (d.nm):	% Number:	St Dev (d.nm):
Z-Average (d.nm):	83,91	Peak 1: 48,52	100,0 15,03
Pdt:	0,157	Peak 2: 0,000	0,0 0,000
Intercept:	0,956	Peak 3: 0,000	0,0 0,000

Result quality : Good

**G**

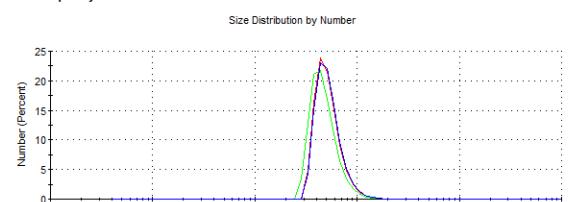
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Z-Average (d.nm):	80,89	Peak 1: 47,06	100,0 14,48
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Intercept:	0,968	Peak 3: 0,000	0,0 0,000

Result quality : Good

**H**

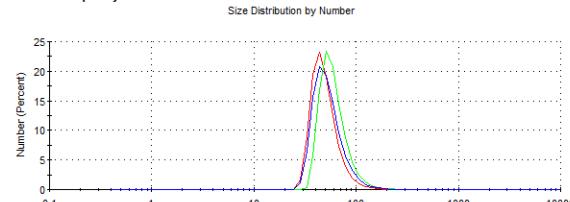
	Size (d.nm):	% Number:	St Dev (d.nm):
Z-Average (d.nm):	92,06	Peak 1: 52,61	100,0 16,49
Pdt:	0,164	Peak 2: 0,000	0,0 0,000
Intercept:	0,934	Peak 3: 0,000	0,0 0,000

Result quality : Good

**I**

	Size (d.nm):	% Number:	St Dev (d.nm):
Z-Average (d.nm):	110,9	Peak 1: 50,47	100,0 17,99
Pdt:	0,211	Peak 2: 0,000	0,0 0,000
Intercept:	0,964	Peak 3: 0,000	0,0 0,000

Result quality : Good

**J**

	Size (d.nm):	% Number:	St Dev (d.nm):
Z-Average (d.nm):	106,1	Peak 1: 61,60	100,0 19,53
Pdt:	0,191	Peak 2: 0,000	0,0 0,000
Intercept:	0,951	Peak 3: 0,000	0,0 0,000

Result quality : Good

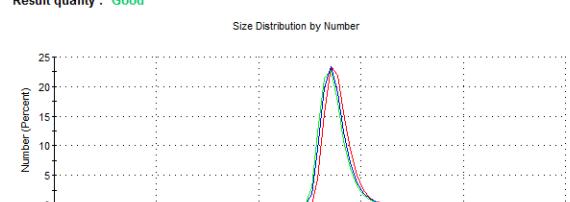
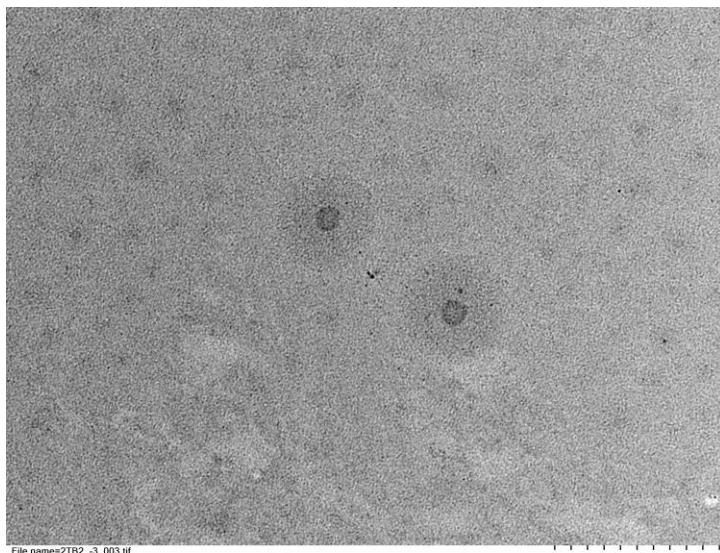
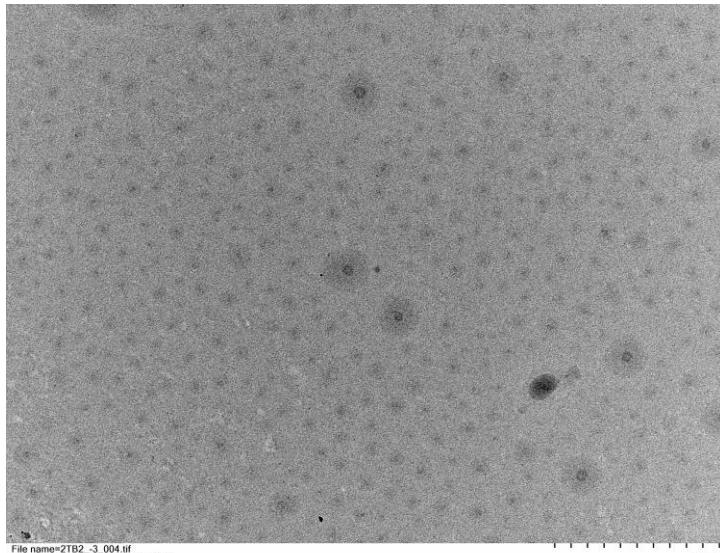


Figure S9. Screen shot of empty nanoreactor size distribution using the intensity(A-E) and number parameters (F-K) based on **1b**, copolymer concentrations 0.1, 0.2, 0.5, 0.75, 1% (wt/wt), 10 mM Tris buffer, pH=7.4, 25°C.



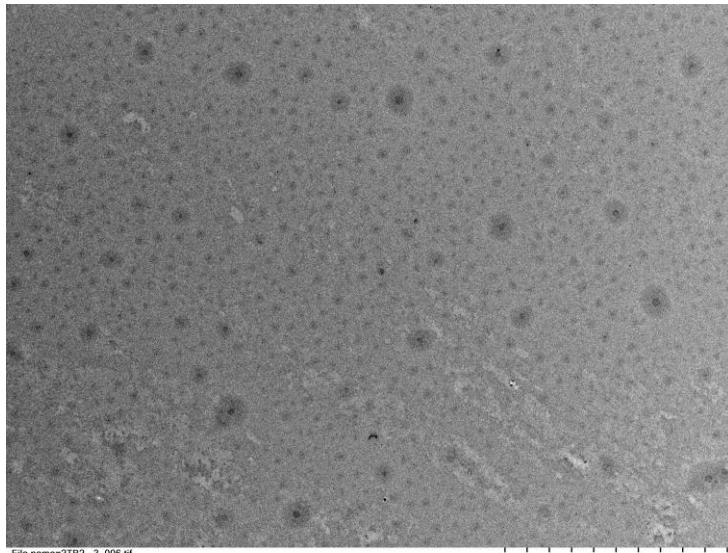
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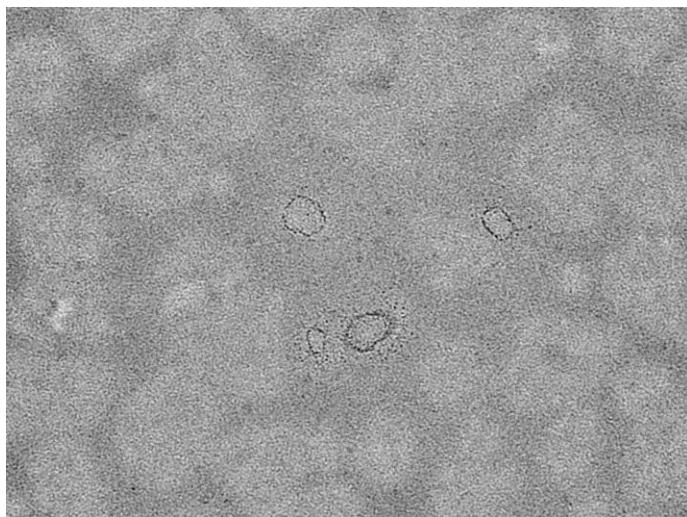
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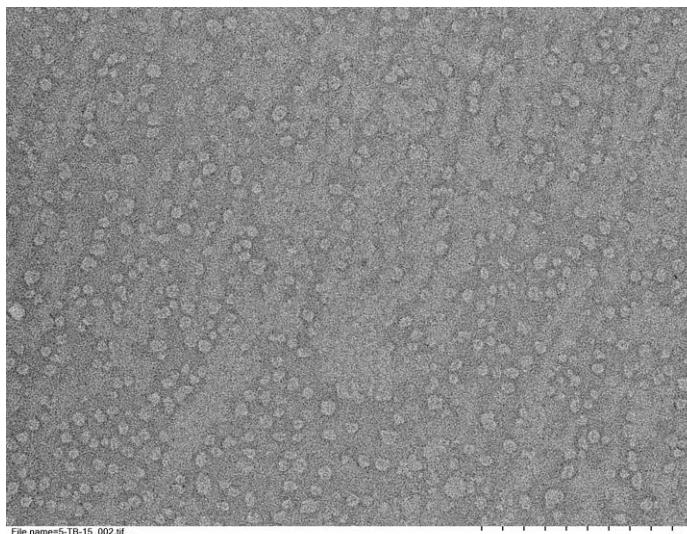
Figure S10. TEM Images of empty nanoreactors based on **1a**, C_{1a} = 10 μg/mL, Tris-Buffer, pH = 7.4, 25 °C.



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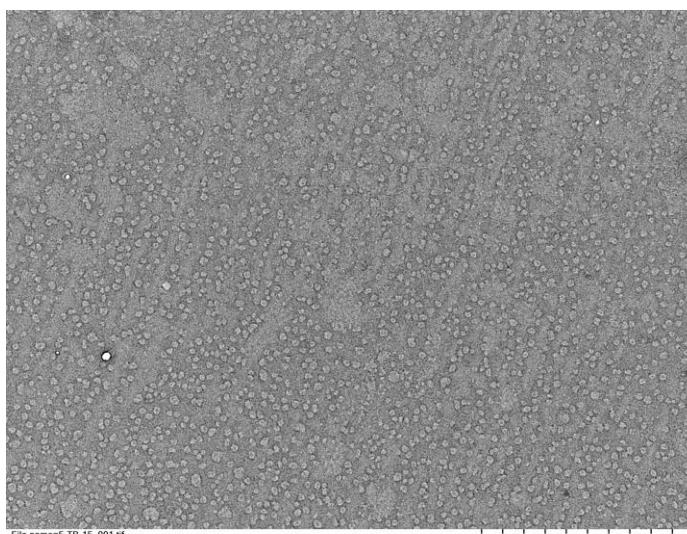
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1.0μm



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2.0μm

Figure S11. TEM Images of empty nanoreactors based on **1b**, $C_{1b} = 10 \mu\text{g/mL}$, Tris-Buffer, pH = 7.4, 25 °C.

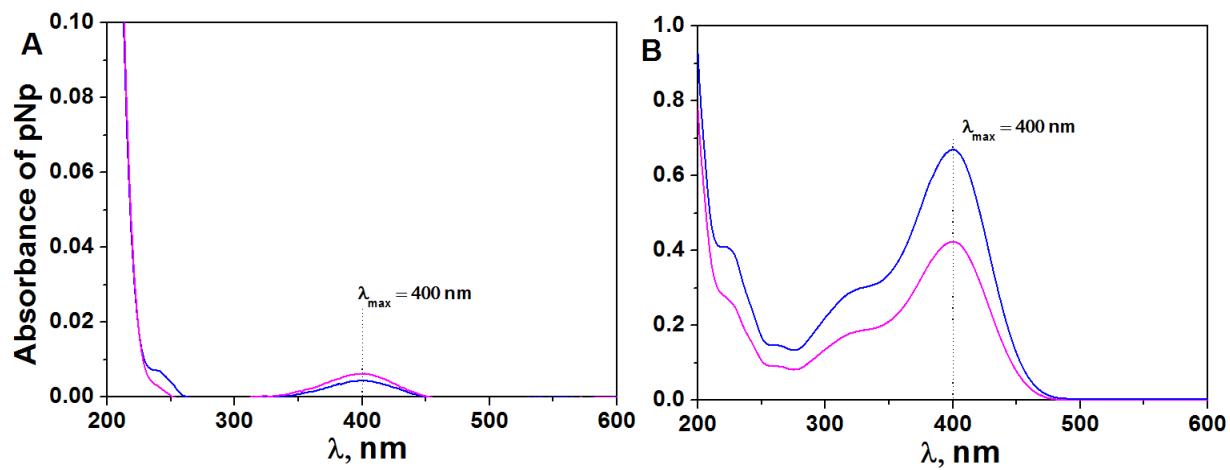


Figure S12. UV Absorbance spectra of pNp for encapsulation efficiency of nanoreactors based on **1b** (A) and **1a** (B), 10mM Tris buffer, pH=7.4, 25°C, C_{polymers} (wt/wt) = 0.5%.

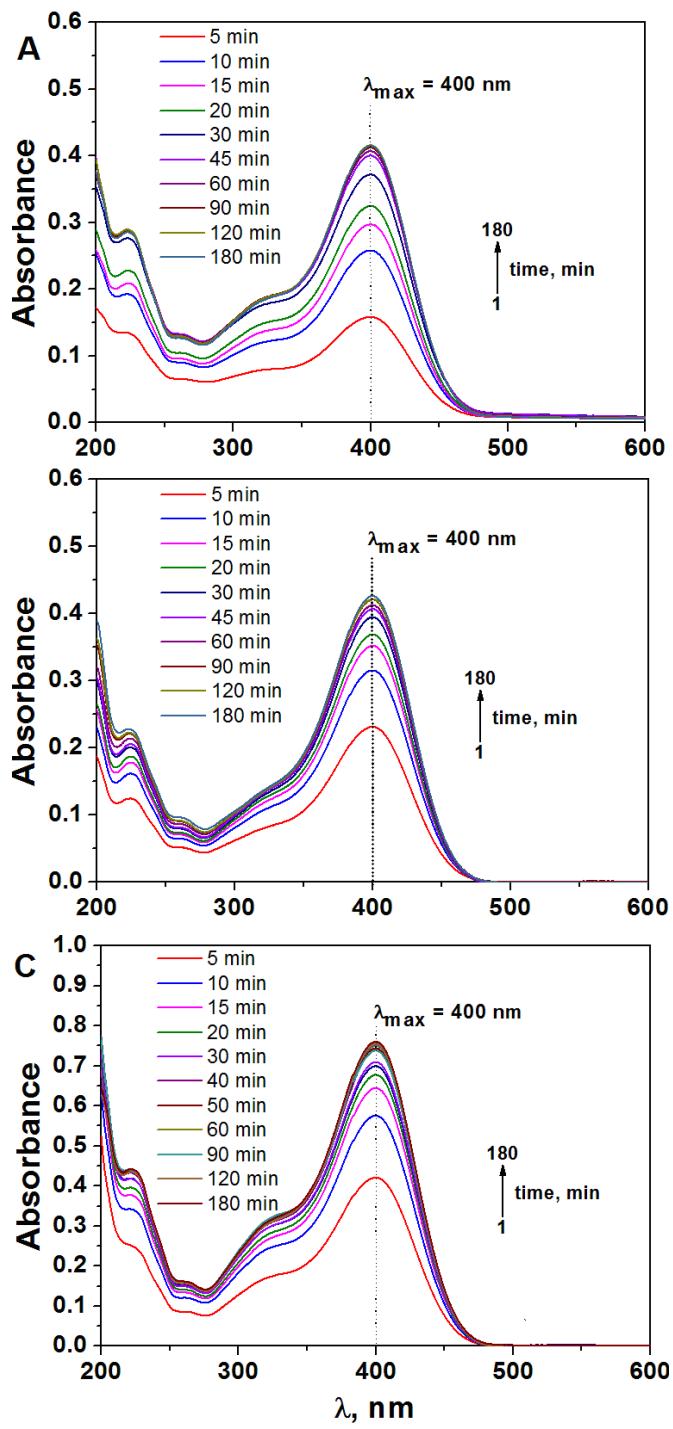


Figure S13. UV Absorbance spectra of pNp from nanoreactors **1a** (**B**) and **1b** (**C**), where control is in the absence of nanoreactors (**A**), $C_{1a} = C_{1b.} = 0.5\% \text{ (wt/wt)}$, $C_{pNp} = 0.1\% \text{ (wt/wt)}$ 37°C , 10 mM Tris-Buffer, pH = 7.4.

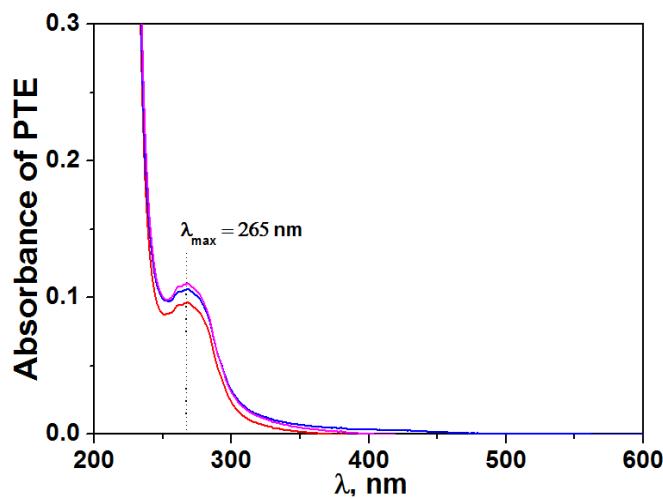
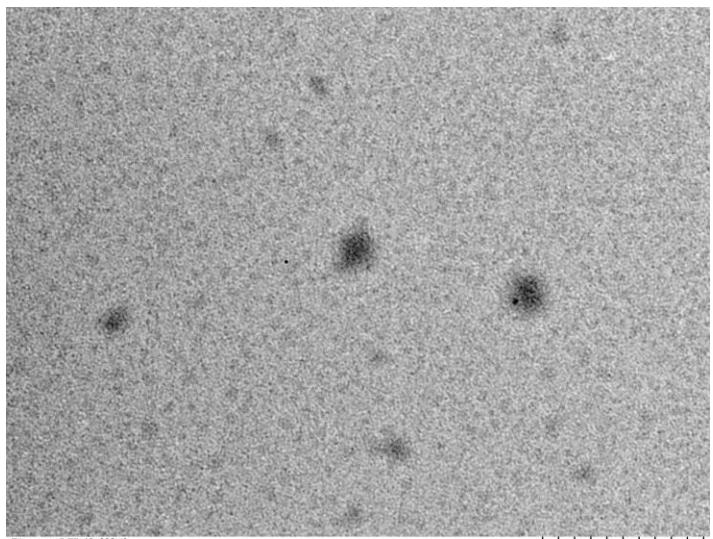
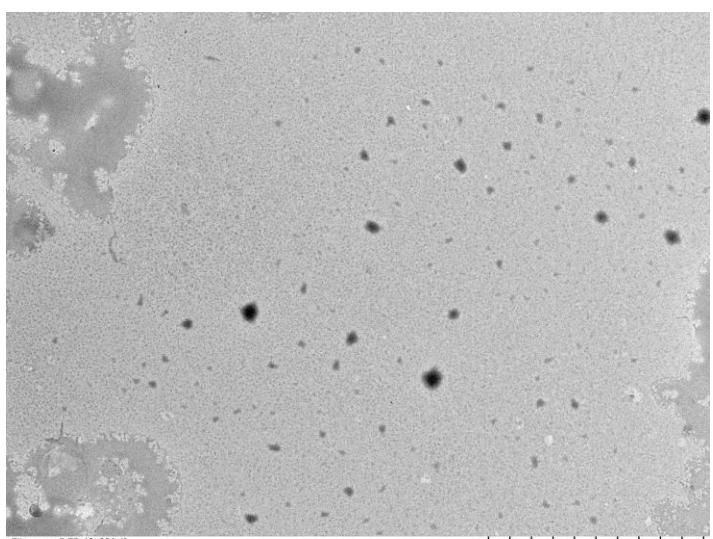


Figure S14. UV Absorbance spectra of enzyme for encapsulation efficiency of nanoreactors, 10mM Tris buffer, pH=7.4, 25°C, C_{1b} (wt/wt) = 0.5%.



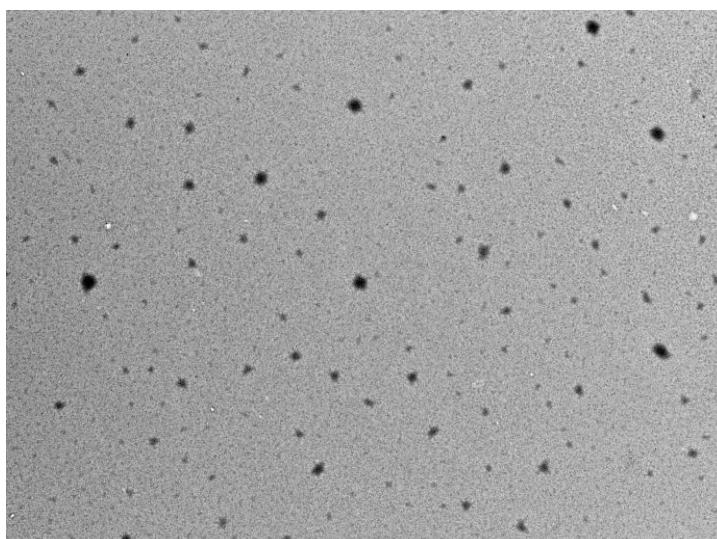
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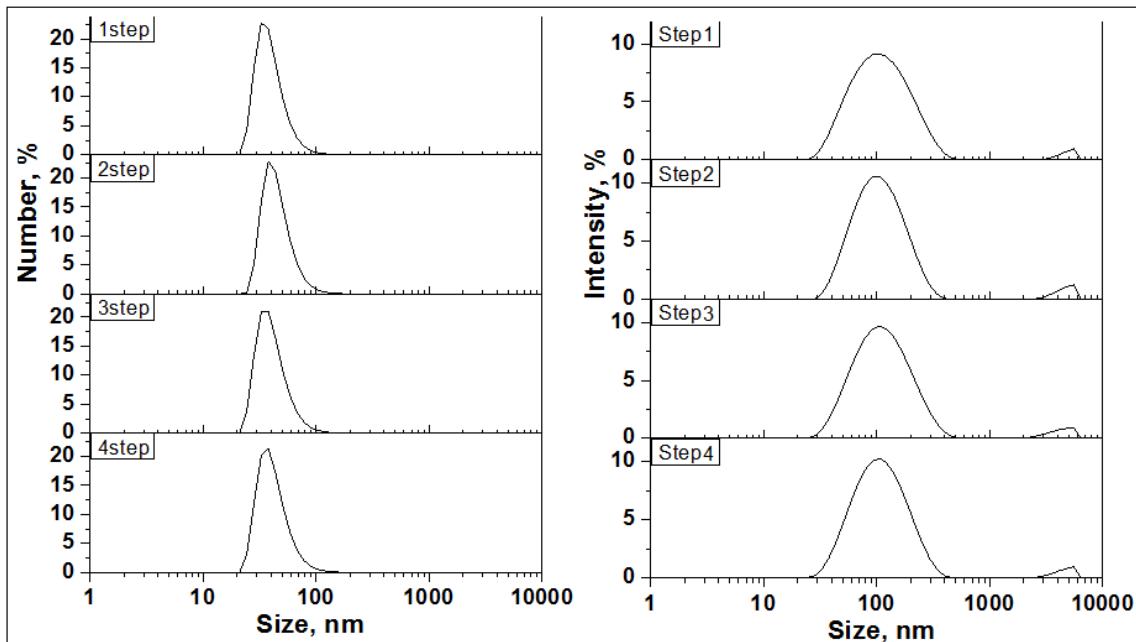
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Figure S15. TEM Images of enzyme-loaded nanoreactors based on **1b**, Tris-Buffer, pH = 7.4, 25 °C.



Figures S16. The size distribution using number (A-E) and intensity (F-J) parameters for nanoreactors based on **1b**, copolymer concentrations 0.5%, (wt/wt), 10 mM Tris buffer, pH=7.4, 25°C, where 1 step is keeping the blood at +37°C for 1 hour, 2 step is keeping the blood at +4°C for 1 hour, 3 step – centrifuge the blood (15min, 2500rpm, +4°C), 4 step – freezing the serum at -20°C – for 7 days.

Table S1. *In vitro* stability of nanoreactors **1b** in blood.

Sample	Medium	Z average, nm	PDI
nR-PTE	Tris-buffer	95.7±0.1	0.16±0.01
1 step	blood	94±2	0.29±0.04
2 step	blood	92±2	0.27±0.03
3 step	serum	99±1	0.27±0.01
4 step	serum	92±2	0.26±0.01

where 1 step is keeping the blood at +37°C for 1 hour, 2 step is keeping the blood at +4°C for 1 hour, 3 step – centrifuge the blood (15min, 2500rpm, +4°C), 4 step – freezing the serum at -20°C – for 7 days.

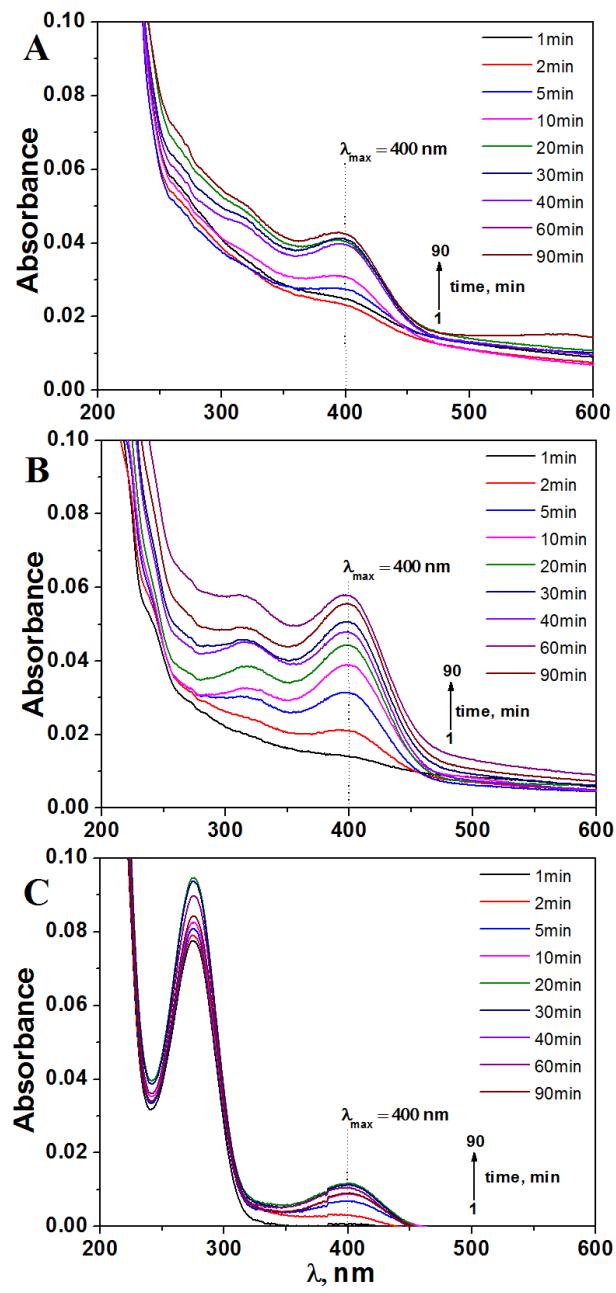


Figure S17. UV Absorbance spectra of pNp after the neutralization of POX by enzyme (A) and enzyme-loaded nanoreactors (B), control - pNp solution (pNp, 5 μM) (C), $C_{\text{POX}} = 5 \mu\text{M}$, $C_{\text{enzyme}} = 1 \mu\text{M}$, 10 mM Tris buffer, pH=7.4, 37°C.

Table S2. Prophylaxis and post-exposure treatment of paraoxon s.c. acute toxicity by *i.v.* administration of enzyme-loaded nanoreactors in mice, where Control 1 is the EtOH 10% in sodium chloride 0.9% solution (*s.c.*) and Control 2 is empty nanoreactor solution (*i.v.*).

Dose, mg/kg	Animals dead / total
Paraoxon acute toxicity study	
Control 1	0/3
Control 2	0/3
0.5	0/3
0.625	0/3
0.65	1/3
0.7	3/3
0.75	3/3
1.25	3/3
enzyme-loaded nanoreactors prophylactic <i>i.v.</i> administration	
5	0/3
10	1/3
15	4/6
25	3/3
enzyme-loaded nanoreactors <i>i.v.</i> treatment	
2.5	0/3
5	1/3
10	3/3

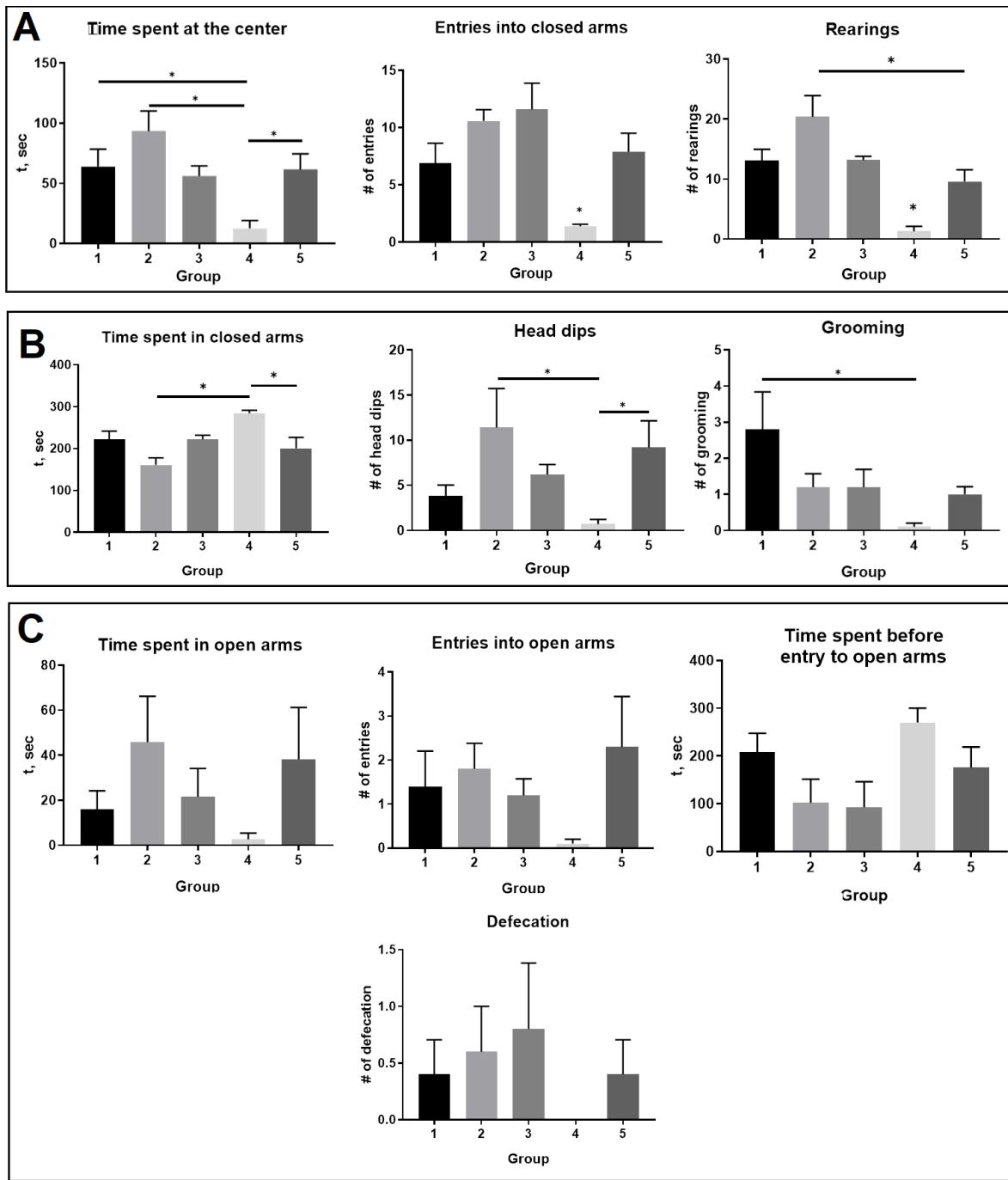


Figure S18. Elevated cross maze parameters on the 1st day of experiment, * p < 0.05

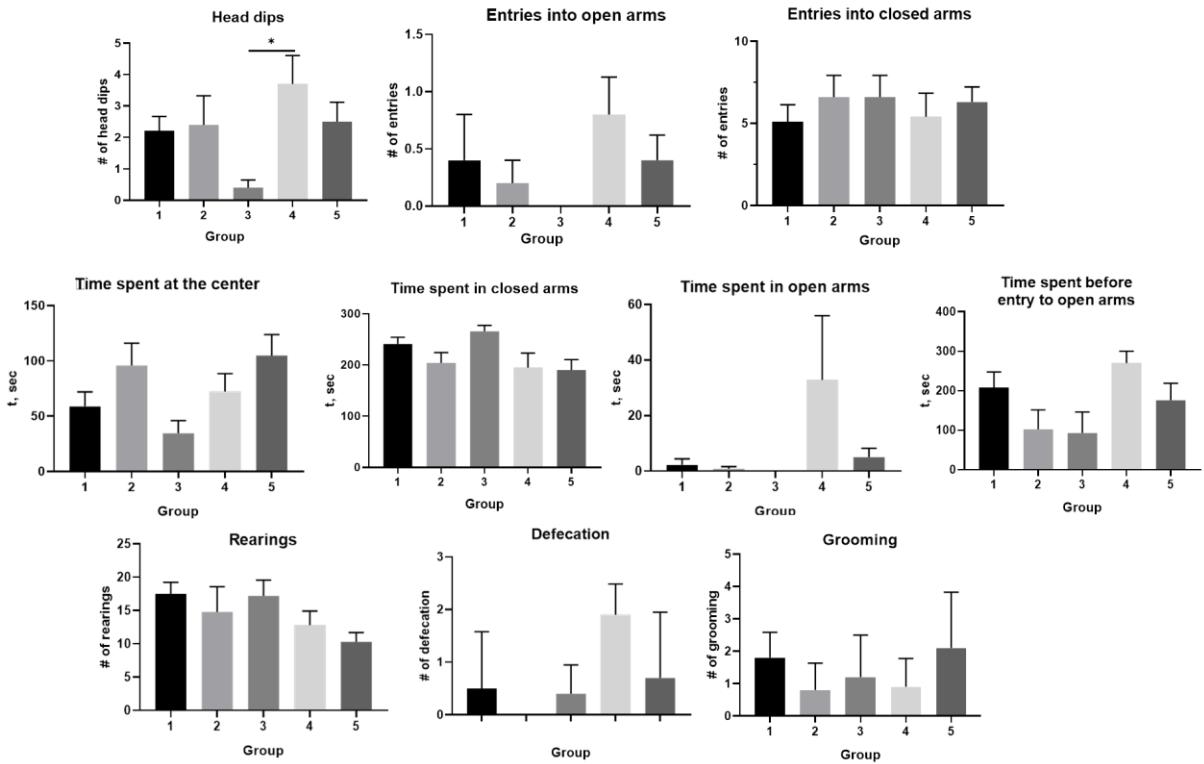


Figure S19. Elevated cross maze parameters on the 30th day of experiment, * p < 0.05.

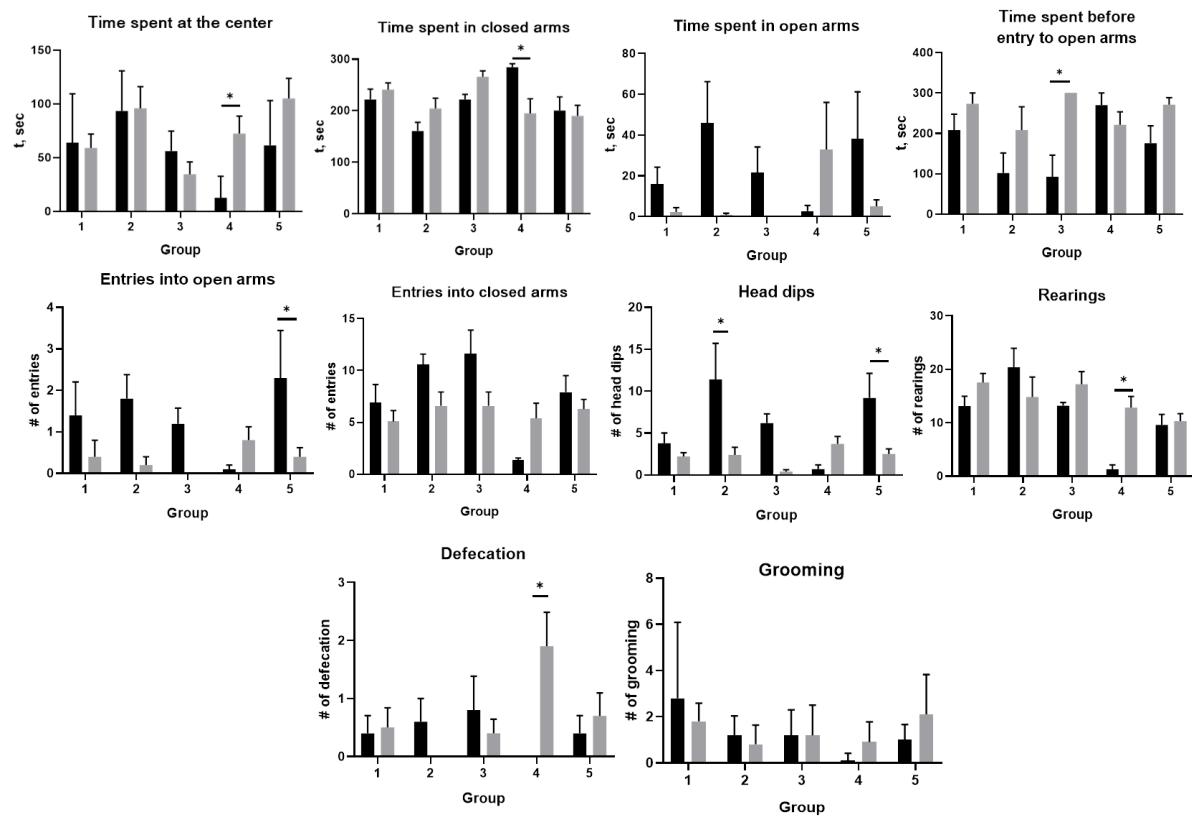


Figure S20. Elevated cross maze parameters on the 1st (black column) vs 30th (grey column) days of experiment, * p < 0.05.

Two sample t Test (16/06/23 20:31:10)

Notes	X-Function	Two sample t Test		
	User Name	Òàòùýia Ìàøèðiâà		
	Time	16/06/23 20:31:10		
Input Data	Data	Range		
	1st Data Range	[Book1]Sheet1!A [1:3]		
	2nd Data Range	[Book1]Sheet1!B [1:3]		
Descriptive Statistics	N	Mean	SD	SEM
A	3	21.96226	3.15392	1.82092
B	3	26.15995	2.83388	1.63614
Difference		-4.19769		
t-Test Statistics		t Statistic	DF	Prob> t
Equal Variance Assumed		-1.71475	4	0.16154
Equal Variance NOT Assumed		-1.71475	3.95506	0.16235
Null Hypothesis: mean1=mean2 = 0				
Alternative Hypothesis: mean1=mean2 <> 0				
At the 0.05 level, the difference of the population means is NOT significantly different with the test difference(0)				

Figure S21. Student's t-test for free enzyme (1) and nanoreactor based on **1b** (2) 1st injection.

Two sample t Test (16/06/23 21:21:10)

Notes

X-Function	Two sample t Test
User Name	Òàòùyìà Ìàøèðiâà
Time	16/06/23 21:21:10

Input Data

	Data	Range
1st Data Range	[Book2]Sheet1!A	[1*:3*]
2nd Data Range	[Book2]Sheet1!B	[1*:3*]

Descriptive Statistics

	N	Mean	SD	SEM
A	3	26.15995	2.83388	1.63614
B	3	25.12116	2.2397	1.29309
Difference		1.03879		

t-Test Statistics

	t Statistic	DF	Prob> t
Equal Variance Assumed	0.49812	4	0.64454
Equal Variance NOT Assumed	0.49812	3.79727	0.64586

Null Hypothesis: mean1-mean2 = 0
Alternative Hypothesis: mean1-mean2 <> 0
At the 0.05 level, the difference of the population means is NOT significantly different with the test difference(0)

Figure S22. Student's t-test for nanoreactor 1st injection and nanoreactor 2nd injection (after 30 days) based on **1b**.

ANOVAOneWay (16/06/23 21:21:37)

+ Notes ▾

+ Input Data ▾

- Descriptive Statistics ▾

	Sample Size	Mean	Standard Deviation	SE of Mean
A	3	21.96226	3.15392	1.82092
B	3	26.15995	2.83388	1.63614
C	3	25.12116	2.2397	1.29309

- One Way ANOVA ▾

+ Overall ANOVA ▾

	DF	Sum of Squares	Mean Square	F Value	Prob>F
Model	2	28.67837	14.33919	1.87079	0.23365
Error	6	45.98857	7.66476		
Total	8	74.66694			

Null Hypothesis: The means of all levels are equal.
Alternative Hypothesis: The means of one or more levels are different.
At the 0.05 level, the population means are not significantly different.

+ Fit Statistics ▾

R-Square	Coeff Var	Root MSE	Data Mean
0.38408	0.1134	2.76853	24.41446

Figure S23. One Way ANOVA for free enzyme 1st injection, nanoreactor 1st injection and nanoreactor 2nd injection (after 30 days) based on **1b**.

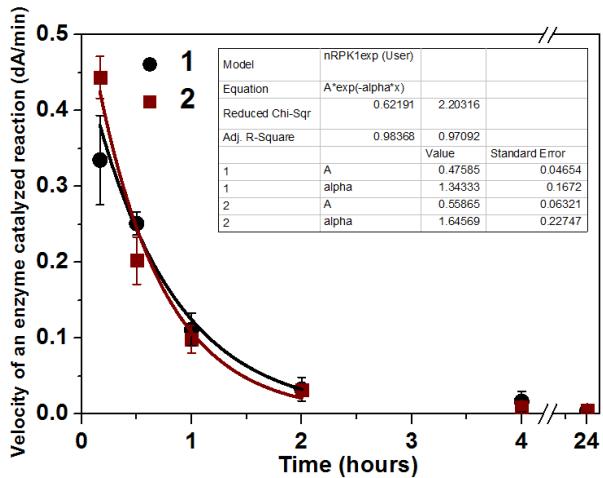


Figure S24. Pharmacokinetic profiles in mouse plasma after intravenous injection of free enzyme (1), enzyme-loaded nanoreactors made by three-block copolymers (2) after second injection (30-day). The dose of enzyme is 5.76 mg/kg. Each point represents the mean \pm SD in 3 mice.

Table S3. Pharmacokinetic parameters observed in mouse after intravenous injection of free enzyme and enzyme-loaded nanoreactors second time after 1 month of 1st injections. The dose of enzyme 5.76 mg/kg. Results represent the mean ± SD for three mice, three-block copolymer nanoreactor.

Sample	number of injections	δ (min ⁻¹)	t _{1/2δ} [*] (min)
Free enzyme	2, 30-day	0.0224±0.0028	30.96±3.85
Enzyme-loaded nanoreactors	2, 30-day	0.0274±0.0038	25.27±3.49

*t_{1/2α} = ln2/α, in which α is the distribution rate (min⁻¹) from blood. Data were analyzed via two-tailed Student's t test (*p ≤ 0.05)

Two sample t Test (19/06/23 16:22:33)

Notes	X-Function	Two sample t Test		
	User Name	Òàòüÿìà Ìàøèðiâà		
	Time	19/06/23 16:22:33		
Input Data	Data	Range		
	1st Data Range	[Book1]Sheet1!1 [1*:3*]		
	2nd Data Range	[Book1]Sheet1!2 [1*:3*]		
Descriptive Statistics	N	Mean	SD	SEM
"1"	3	30.9595	3.85343	2.22478
"2"	3	25.27136	3.49305	2.01671
Difference		5.68814		
t-Test Statistics		t Statistic	DF	Prob> t
Equal Variance Assumed		1.89428	4	0.1311
Equal Variance NOT Assumed		1.89428	3.96205	0.13178
Null Hypothesis: mean1-mean2 = 0 Alternative Hypothesis: mean1-mean2 <> 0 At the 0.05 level, the difference of the population means is NOT significantly different with the test difference(0)				

Figure S25. Student's t-test for free enzyme and three-block copolymer nanoreactor 2nd injection.

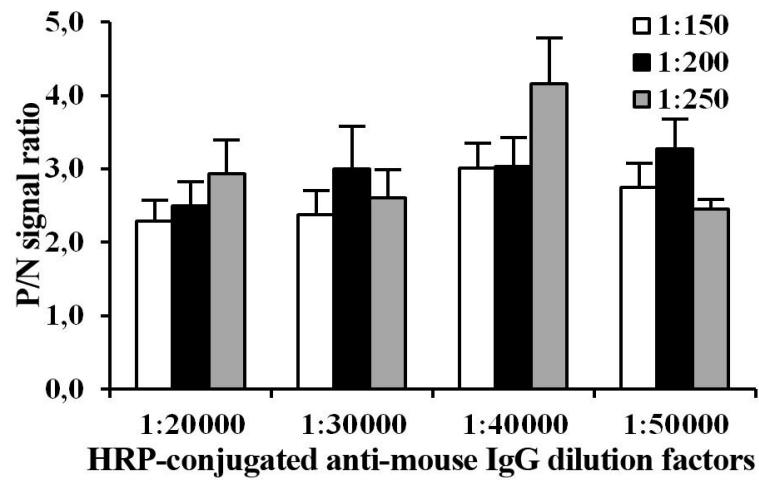


Figure S26. The ratio of the signals from positive and negative control samples (P/N signal ratio) with tested dilution factors.

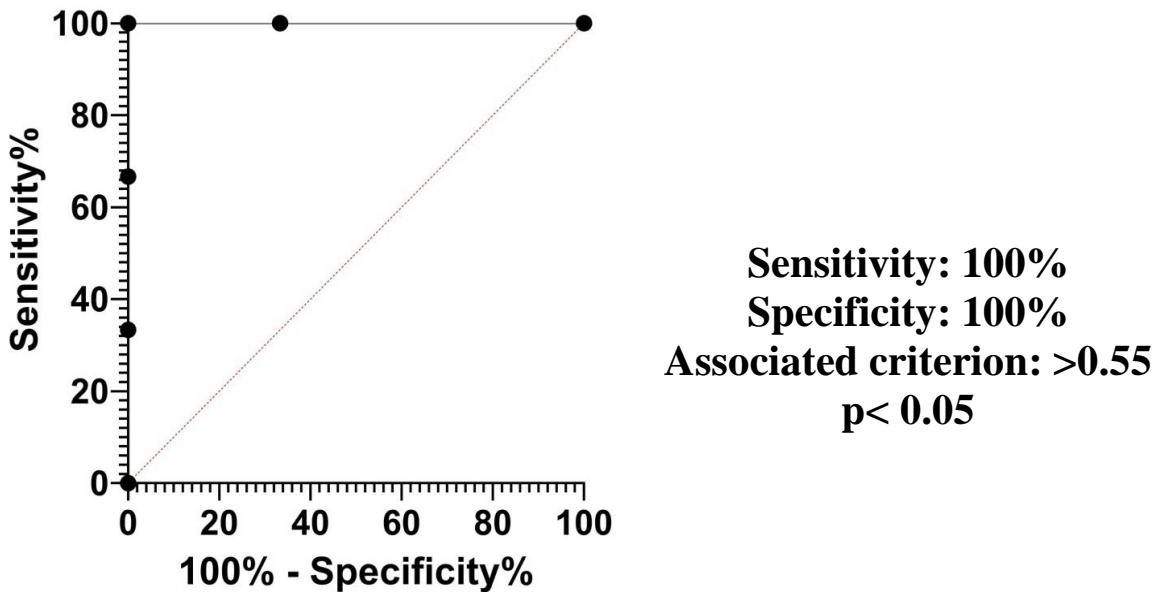


Figure S27. The receiver operating characteristic (ROC) analysis of ELISA showing sensitivity versus specificity for discrimination of positive and negative serum samples.

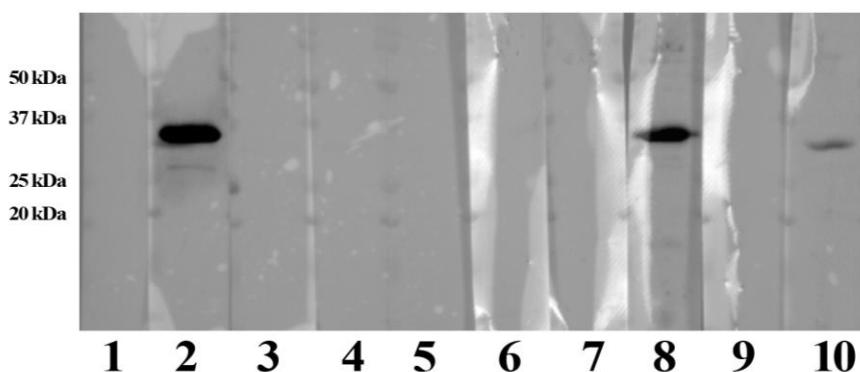
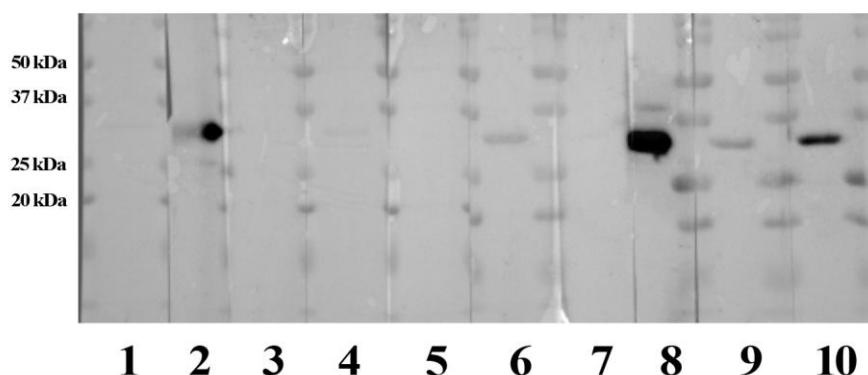
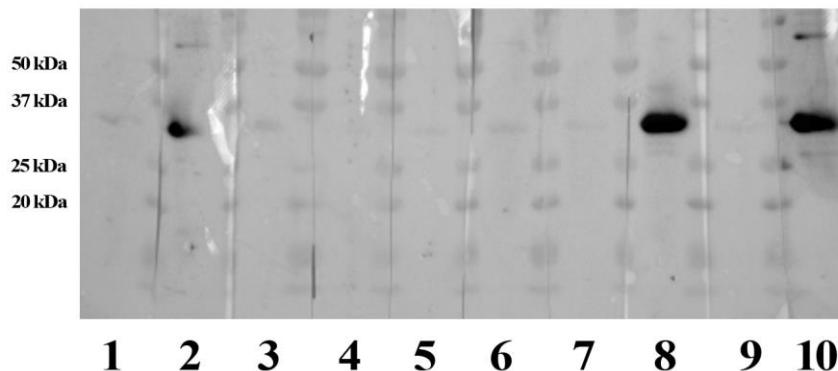


Figure S28. The western blot membranes for anti-PTE-IgG from 3 independent experiments: 1 - negative control, 2 – positive control, 3 – POX, 4 – empty nanoreactor, 5 - solvent, 6 – PTE, 7 – PTE-mPEG–PPS–mPEG nanoreactor (1 injection), 8 - PTE-mPEG–PPS–mPEG nanoreactor (2 injections), 9 - PTE-mPEG–PPS nanoreactor (1 injection), 10 - PTE-mPEG–PPS nanoreactor (2 injections).

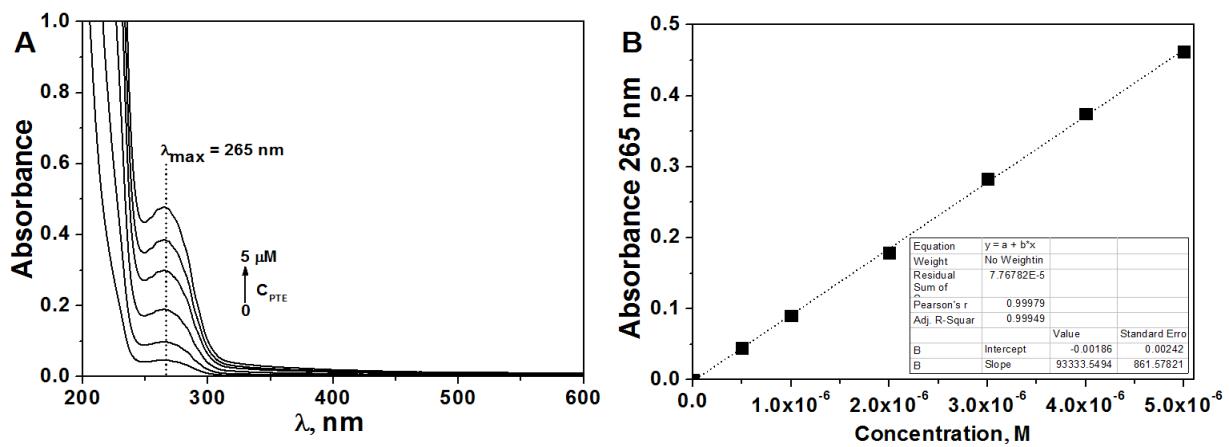


Figure S29. UV Absorbance spectra (A) and calibration curve (B) of Phosphotriesterase, 10mM Tris buffer, pH=7.4, 25°C.

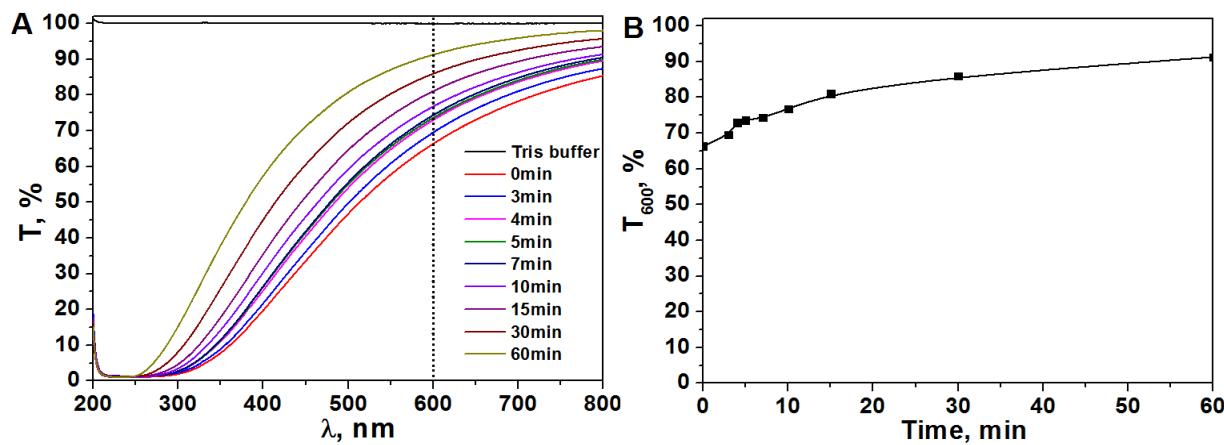


Figure S30. Transmittance of empty nanoreactors under centrifugation conditions at 5000 rpm, using centrifuge over time, C_{1b} (wt/wt) = 0.5%, 10 mM Tris buffer, pH=7.4.