A general strategy for degradable single-chain nanoparticles via cross-linker mediated chain collapse of radical copolymers

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1. Materials and Methods

All reagents and solvents were purchased from Sigma Aldrich and used as received. 2-Methylene-1,3-dioxepane (MDO) was prepared as previously reported.¹ Dry THF was prepared using a Glass Contour - Solvent Purification System. Deionized (DI) water was prepared using an AquaMAX-Basic 321 water purification system. ¹H NMR spectra were recorded on a Bruker 400 Ultra Shield spectrometer. Dynamic light scattering (DLS) was conducted on a Malvern Zetasizer Nanoseries at 25 °C. Size exclusion chromatography (SEC) was conducted on a Viscotek TDAmax consisting of a GPCmax integrated solvent and sample delivery module, a TDA 302 Triple Detector Array, and OmniSEC software. 2 x PLgel 5 µm Mixed-C (200-2,000,000) columns were applied in sequence for separation. THF was used as the eluent at 1.0 mL/ min with column and detector temperature at 30 °C, molecular weight values were determined against polystyrene standards.

2. Experimental

2.1 Synthesis of Poly(MDO)² and Treatment with Diamine



Figure S1. Synthesis of Poly(MDO) and subsequent control experiment to determine stability in the presence of primary amine.

Poly(MDO) was prepared and treated with 1,5-diaminopentane and Et₃N (under identical conditions to (**DSCNP6**) confirming that main-chain ester hydrolysis or aminolysis does not occur in the presence of primary amines (Figure S1). A 15 mL Schlenk tube was fitted with a magnetic stirrer bar, rinsed with Et₃N and dried under high vacuum. 2-Methylene-1,3-dioxepane (10.0 g, 87.61 mmol, 100 eq) and azobisisobutyronitrile (143.8 mg, 0.87 mmol, 1 eq) were added to the Schenk Tube. The reaction mixture was degassed via five freeze-pump-thaw cycles and backfilled with N₂ before heating to 65 °C and stirring at 500 rpm for 24 h. After this time the reaction was quenched by rapid cooling, and the polymer isolated as a white waxy solid (8.60 g, 86 % yield) by three precipitations from CHCl₃ into hexane. **Poly(MDO)** was analysed via ¹H NMR (Fig. S2a) and SEC (Fig. S3). Subsequently, a 250 mL round bottom flask was fitted with a magnetic stirrer bar, thoroughly rinsed with acetone and dried at 100 °C overnight. **Poly(MDO)** (100 mg), 1,5-diaminopentane (16.75 mg), Et₃N (40 μ L) and dry THF (100 mL) were transferred into the round bottom flask and the solution obtained was stirred at room temperature for 24 h. After evaporation under reduced pressure at 28 °C, the resulting polymer dried under high vacuum at room temperature. Sample obtained was analysed via ¹H NMR (Fig. S2b) and SEC (Fig. S3).





Figure S3. (a) SEC traces of **Poly(MDO)** (blue line) and Poly(MDO) after treatment with 1,5-diaminopentane and Et_3N in THF (red line). Peak at 21.0 – 23.0 min = system peak indicating the transition from entropy-dominated size-based separation to enthalpy-dominated interaction chromatography. (b) Expansion of polymer traces to clearly observe no degradation.

Control experiment SEC analysis

Poly(MDO)	Mn = 6,090 Da Mw = 20,135 Da Mp = 17,725 Da	Poly(MDO) treated with 1,5-diaminopentane	M _n = 5,635 Da M _w = 19,465 Da M _p = 17,315 Da
	М _р = 17,725 Da Ð _M = 3.31		М _р = 17,315 Da Ð _M = 3.45

2.2 Synthesis of Poly(MDO-c-NHSMA)

A 15 mL Schlenk tube was fitted with a magnetic stirrer bar, rinsed with Et₃N and dried under high vacuum. 2-Methylene-1,3-dioxepane (3.88 g, 34 mmol, 85 eq), methacrylic acid *N*-hydroxysuccinimide ester (1.10 g, 6 mmol, 15 eq), azobisisobutyronitrile (6.6 mg, 40 µmol, 0.1 eq) and 1,4-dioxane (1.66 g, 25 wt %) were added to the Schenk Tube. A small aliquot (*time = 0 h*) was then extracted for monomer conversion determined by ¹H NMR (CDCl₃ passed over Na₂CO₃) spectroscopy. The reaction mixture was degassed via three freeze-pump-thaw cycles and backfilled with N₂ before heating to 70 °C and stirring at 500 rpm for 6 h. After this time the reaction was quenched by rapid cooling, and a second aliquot (*time = 6 h*) extracted for monomer conversion analysis. Monomer conversions were determined by comparing the 1,4-dioxane protons present at δ 3.71 ppm with the residual MDO vinyl protons at δ 3.42 ppm and the residual NHSMA vinyl protons at δ 6.37 and 5.84 ppm. Quenching the reaction before complete NHSMA consumption eliminates the formation of homoMDO polymer chains. The polymer was isolated as a white solid (1.50 g, 83 % yield) by three precipitations from THF into hexane.

2.3 Synthesis of Degradable Single-Chain Nanoparticles (DSCNP1-6)

The reagents and quantities for each degradable single-chain nanoparticle (**DSCNP1-6**) preparation are provided in Table S1. Six 250 mL round bottom flasks were fitted with magnetic stirrer bars, thoroughly rinsed with acetone and dried at 100 °C overnight. **Poly(MDO-***c***-NHSMA)** (100 mg) was transferred into each round bottom flask followed by the stated quantity of dry THF, each solution was left to stir at room temperature for 1 h to ensure complete polymer dissolution.

Separately, a stock solution (1 mg/ mL) of diamine cross-linker was prepared by adding 50 mg of 1,5-diaminopentane to 50 mL of dry THF. The stated quantity of stock solution was added to each round bottom flask, and the solutions obtained allowed to stir at room temperature for 30 mins. After this time Et₃N (40 μ L) was added to each of the six solutions. Amide bond formation was allowed to proceed for 24 h at room temperature. After this time precipitation of *N*-hydroxysuccinimide was observed in **DSCNP5** and **DSCNP6**, both of these solutions were filtered to remove the precipitated by-product. All six reactions were then separated into two portions (40 mL and 60 mL). The 40 mL portion was reduced under rotary evaporation to approximately 4 mL (10 mg/ mL polymer conc.) before accurate dilution with dry THF to 8 mL (5 mg/ mL polymer conc.) for SEC analysis. The second 60 mL portion was directly analysed by DLS analysis, then extensively evaporated to dryness before ¹H NMR analysis (20 mg in 0.4 mL CDCl₃), leaving 40 mg available for degradation study.

Sample ID	Poly(MDO- <i>c</i> - NHSMA)	NHSMA Quantity	Target Cross- Linking (%)	1,5-DAP (m _r = 102.18 Da)	Et₃N (<i>m</i> r = 101.19 Da)	Dry THF	1,5-DAP Solution (1 mg/ mL)
DSCNP1	100 mg 48 mol % NHSMA 60 wt % NHSMA	1 eq 60 mg 0.328 mmol	0	-	1 eq - 33.19 mg/ 46 μL 0.328 mmol	100 mL	-
DSCNP2			10	0.05 eq 1.68 mg 0.0164 mmol		98.3 mL	1.7 mL
DSCNP3			25	0.125 eq 4.20 mg 0.0410 mmol		95.8 mL	4.2 mL
DSCNP4			50	0.250 eq 8.38 mg 0.0820 mmol		91.6 mL	8.4 mL
DSCNP5			75	0.375 eq 12.57 mg 0.1230 mmol		87.4 mL	12.6 mL
DSCNP6			100	0.50 eq 16.76 mg 0.1640 mmol		83.2 mL	16.8 mL

Table S1. Reagents and quantities employed during synthesis of degradable single-chain nanoparticles (**DSCNP1-6**). 1,5-DAP = 1,5-diaminopenate, $Et_3N = trimethylamine$.



2.4 Synthesis of Water Soluble Degradable Single-Chain Nanoparticle (DSCNP7)

Figure S4. (a) ¹H NMR (CDCl₃) spectra of **DSCNP7**. (b) DLS (number) analysis of **Poly(MDO-***c***-NHSMA)** and **DSCNP7** in THF (3 mg/ mL at 25 °C). (c) DLS analysis of **DSCNP7** in H₂O (3 mg/ mL at 25 °C).

The hydrophilic cross-linker 1,8-diamino-3,6-dioxaoctane (DOA) was utilized to prepare the water-soluble **DSCNP7** targeted a cross-linking degree of 100 % (conditions detailed below). A reaction duration of 48 h was required to reach a 69 % calculated degree of cross-linking, determined by ¹H NMR (Fig. S4a). This increase in cross-linking duration is likely due to the additional bulkiness of DOA. SEC (Table 1 in manuscript) confirmed an apparent decrease in molecular weight for **DCSNP7** (M_p = 22,400 Da). Dynamic light scattering (DLS) analysis in THF (Fig. S4b) further confirmed a decrease in volume when comparing the linear precursor **Poly(MDO-***c***-NHSMA)** (D_h = 16.6 nm) with **DSCNP7** (D_h = 9.8 nm). This difference in D_h (6.8 nm) corresponds to a chain compaction of 41 %. Making a direct comparison in H₂O isn't possible as **Poly(MDO-***c***-NHSMA)** is not water soluble. However, after dispersion in H₂O via solvent evaporation DLS analysis of

DSCNP7 (Fig. S4c) displayed a $D_h = 15.8$ nm. This nanoparticle solution was very stable with no changes in DLS after one month. The size difference observed for **DCNP7** in THF ($D_h = 9.8$ nm) and H₂O ($D_h = 15.8$ nm) could be due to variations in solvation within the nanoparticle interior, or a result of **DSCNP7** aggregation in H₂O. **DSCNP7** successfully underwent chemical degradation into branched oligomers ($M_p = 710$ Da) via main-chain ester hydrolysis (Table 1 in manuscript).

Detailed conditions for the synthesis of **DSCNP7** are as follows:

Poly(MDO-c-NHSMA) (100 mg) was transferred into a 250 mL round bottom flask fitted with a magnetic stirrer bar (previously rinsed with acetone and dried at 100 °C overnight). Dry THF (76.6 mL) was added and the solution left to stir at room temperature for 1 h to ensure complete polymer dissolution. 1,8-Diamino-3,6-dioxaoctane (0.5 eq relative to NHSMA content, 0.1640 mmol, 23.4 mg) in 23.4 mL of dry THF was added, and the solutions obtained allowed to stir at room temperature for 30 mins. After this time Et₃N (40 μ L) was added and amide bond formation was allowed to proceed for 48 h at room temperature. After this time, samples were prepared for ¹H NMR analysis (20 mg in 0.4 mL CDCl₃), SEC analysis (5 mg/ mL in THF), DLS analysis (3 mg/ mL in THF and H₂O) and degradation study.

2.5 Preparation of DSCNP7 Aqueous Solution

DSCNP7 (20 mg, obtained after NMR sample evaporation to dryness) was transferred into a 25 mL round bottom flask fitted with a magnetic stirrer bar. Acetone (5 mL) was added and this solution was stirred for 15 mins, which resulted in complete nanoparticle dissolution. H_2O (10 mL) was then added dropwise over 30 mins while stirring. The obtained solution was left open to the atmosphere and was stirred at room temperature for 24 h to allow acetone evaporation. After this time, the solution obtained was analysed directly via DLS (3 mg/ mL at 25 °C).

2.6 Degradation of Poly(MDO-c-NHSMA) and DSCNP1-7

The degradation of **Poly(MDO-***c***-NHSMA)** and **DSCNP1-7** was performed under accelerated alkali conditions. Each sample (40 mg) was transferred into a small vial fitted with a magnetic stirrer bar. THF (2 mL) was added, followed by KOH (80 mg) in H₂O (2 mL). The obtained solutions were stirred at room temperature for 48 h, to ensure complete main-chain ester hydrolysis. After this time the solutions were acidified with conc. $HCl_{(aq)}$ aliquots and subsequently evaporated to dryness. CHCl₃ (5 mL) was then added and the solutions stirred for 2 h, after this time KCl precipitated was removed by filtration and the degraded oligomers obtained after evaporation to dryness. Samples were prepared for SEC analysis (5 mg/ mL in THF).



4. Size Exclusion Chromatography (DSCNP1-6 and DSCNP1-6 After Hydrolysis)



Figure S8. SEC analysis (in THF) of **DSCNP1-6**. Peak at 21.0 - 23.0 min = system peak indicating the transition from entropy-dominated size-based separation to enthalpy-dominated interaction chromatography. In order to prevent inter-chain cross-linking in the dry state **DSCNP1-6** samples are not completely evaporated to dryness, therefore additional peaks are present in these traces. Peak at 19.5 - 20 min = residual 1,5-diaminopentane and Et₃N. Potential nanoparticle column interactions are observed between 18.0 - 19.5 min, which increase at higher degrees of cross-linking.



Figure S9. SEC analysis (in THF) of **DSCNP1-6** after main-chain ester hydrolysis. Peak at 21.0 – 23.0 min = system peak indicating the transition from entropy-dominated size-based separation to enthalpy-dominated interaction chromatography.

5. Dynamic Light Scattering Analysis (DSCNP1-6)

Comula ID	Run	Hydrodynamic Diameter (D _h)				
Sample ID		Intensity (nm)	Volume (nm)	Number (nm)	Z-Average (nm)	וטץ
	1	38.53	22.64	16.85	31.74	0.179
DCCNID4	2	38.66	21.86	15.69	32.20	0.195
DSCNPI	3	39.65	22.15	16.14	31.87	0.176
0% Cross Linking	4	39.85	21.90	16.06	31.86	0.180
0 % Cross-Linking	5	38.59	22.65	16.53	31.69	0.195
	Average	39.06	22.24	16.25	31.87	0.185
	1	36.41	19.46	13.65	29.10	0.179
DCCNID2	2	36.52	19.11	13.30	29.07	0.188
DSCNPZ	3	36.42	19.67	14.19	29.13	0.186
0.% Cross Linking	4	36.98	19.03	13.07	29.27	0.185
9 % Cross-Linking	5	36.17	19.98	14.49	29.15	0.180
	Average	36.50	19.45	13.74	29.14	0.184
	1	33.15	18.23	12.85	26.82	0.172
	2	33.22	18.55	13.57	27.05	0.184
DSCNP3	3	33.75	18.82	13.79	27.10	0.177
22.0/ Cross Linking	4	33.20	18.91	13.98	26.94	0.170
23 % Cross-Linking	5	33.15	18.43	13.24	26.82	0.170
	Average	33.29	18.59	13.49	26.95	0.175
	1	29.67	16.88	12.26	24.29	0.163
DCONDA	2	29.70	17.13	12.60	24.27	0.164
DSCNP4	3	29.68	16.80	12.12	24.31	0.162
45 % Cross Linking	4	29.87	16.78	12.22	24.30	0.166
45 % Cross-Linking	5	29.70	17.29	12.67	24.40	0.161
	Average	29.72	16.98	12.37	24.31	0.163
	1	26.98	16.06	11.85	22.49	0.150
DCONDE	2	26.49	16.62	12.42	22.37	0.143
DSCNP5	3	26.59	16.29	12.15	22.37	0.144
C2 0/ Crease Linking	4	27.22	16.23	12.15	22.64	0.161
63 % Cross-Linking	5	27.18	15.93	11.50	22.56	0.154
	Average	26.89	16.23	12.01	22.49	0.150
	1	25.88	15.38	11.54	21.45	0.161
DECNIDE	2	26.40	14.86	10.88	21.44	0.169
DSCNPb	3	26.13	15.06	11.05	21.51	0.169
90% Crocs Linking	4	25.94	15.39	11.61	21.67	0.174
00 % Cross-Linking	5	26.32	14.82	10.82	21.70	0.175
	Average	26.13	15.10	11.18	21.55	0.170

Table S2. Dynamic light scattering analysis of DSCNP1-6 in THF (1 mg/ mL) at 25 °C.



Figure S10. DLS overview of DSCNP1-6 by intensity, volume, number and Z-Average.







Figure S12. DLS analysis of DSCNP2 by (a) intensity, (b) volume and (c) number.



Figure S13. DLS analysis of DSCNP3 by (a) intensity, (b) volume and (c) number.





Figure S16. DLS analysis of DSCNP6 by (a) intensity, (b) volume and (c) number.























6. References

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