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# **Supporting Information**

# Poly (2-Oxazoline)-Based Core Cross-Linked Star Polymers: Synthesis and Drug Delivery Applications

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Figure S1. <sup>1</sup>H NMR of the MeOx monomer (500 MHz, CDCl<sub>3</sub>, 25 °C).



Figure S2. <sup>1</sup>H NMR of the PhBisOx monomer (500 MHz, CDCl<sub>3</sub>, 25 °C).



Figure S3. <sup>1</sup>H NMR of the ButOx monomer (500 MHz, CDCl<sub>3</sub>, 25 °C).



CDCl<sub>3</sub>, 25 °C).



**Figure S5.** SEC traces for (a) PMeOx (black), unpurified (PMeOx)<sub>106</sub>-*b*-P(PhBisOx*cl/co*-ButOx)(red), and purified (PMeOx)<sub>106</sub>-*b*-P(PhBisOx-*cl/co*-ButOx)(blue), (DMF, 45 °C, PS standard).

#### Equation S1. The precursor of the arm percentage.

The relative precursor percentage of the PMeOx arm was calculated from the SEC chromatogram integration method by using the following equation:

 $precursor arm \% = \frac{peak area of unreacted arm}{peak area of unreacted arm + peak area of star polymer} \times 100$ 

#### Equation S2. Calculation of Number of Arms in CCS POxs

The average number of arms in the CCS POxs was calculated based on SEC, MDS-SEC, and <sup>1</sup>H NMR analysis of precursor polymers and star polymers. The number of PMeOx arms in CCS (PMeOx)<sub>n</sub>-*b*-P(PhBisOx-*cl/co*-ButOx) was estimated by molecular weights of CCS polymer and PMeOx. The purified star polymer was analyzed

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using (DA-SEC) in DMF to determine the absolute molecular weight ( $M_{w, star}$ ). The average number of arms was calculated using the following equation:

$$N_{arm} = \frac{(M_{w, star} \times arm_{wf})}{M_{n, arm, NMR}}$$

Where, N<sub>arm</sub>,  $M_{w, \text{ star}}$ , arm <sub>wf %</sub>, and  $M_{n, \text{ arm}, \text{ NMR}}$  represent the number of arms in CCS polymer, weight averaged molecular weight of CCS polymer measured by MDS-SEC, the weight fraction of PMeOx arms in CCS polymer, and weight averaged molecular weight of PMeOx arms  $M_{n, \text{ arm}, \text{ NMR}}$  calculated from NMR, respectively. The weight fraction of PMeOx was calculated as follows:

$$arm_{wf} = \frac{W(PMeOx) \times yield \ of \ CCS \ polymer}{[W(PMeOx) \times yield \ of \ CCS \ polymer + W(PhBisOx) + W \ (ButOx)]}$$

Where, W(PMeOx), W(PhBisOx), and W(ButOx) are the weights of PMeOx, PhBisOx, and ButOx in the reaction solution.



**Figure S6.** (a) DLS traces of (PMeOx)<sub>44</sub>-*b*-P(PhBisOx-*cl/co*-ButOx) (black) and (PMeOx)<sub>106</sub>-*b*-P(PhBisOx-*cl/co*-ButOx)(red) in water (concentration: 2 mg/mL) and TEM images (b) PMeOx<sub>44</sub>-*b*-P(PhBisOx-*cl/co*-ButOx) and (c) (PMeOx)<sub>106</sub>-*b*-P(PhBisOx-*cl/co*-ButOx).

#### Equation S3. Preparation of calibration Curve.

A series of free DOX solutions in water with known concentrations were used to prepare the calibration curve (Figure S7). The DOX concentration was calculated by the following equation:

Where X is the DOX concentration, and Y is the absorbance from UV-Vis.



**Figure S7.** The standard calibration curve for free DOX with different concentrations in water.

## In vitro DOX release from CCS polymer.







**Figure S8.** *In vitro* DOX release profile from CCS POxs (a)  $(PMeOx)_{44}$ -*b*-P(PhBisOx*cl/co*-ButOx), (b)  $(PMeOx)_{59}$ -*b*-P(PhBisOx-*cl/co*-ButOx), and (c)  $(PMeOx)_{62}$ -*b*-P(PhBisOx-*cl/co*-ButOx) in PBS at different pH values at 37°C. The results are presented as average data with SD (n= 2).







**Figure S9.** Cell viability of HeLa cells with different concentrations of a series of CCS POxs for 24 h. Error bars are based on SD (n=2).







**Figure S10.** Cell viability of HeLa cells with different concentrations of a series of CCS POxs for 48 h. Error bars are based on SD (n=2).



0-

0.1

0.5

1

DOX Concentration µg/mL

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**Figure S11**. Cytotoxicity of DOX-loaded series of CCS POxs and free DOX in Hela cells for 24 h. (a) (PMeOx)<sub>44</sub>-*b*-P(PhBisOx-*cl/co*-ButOx), (b) (PMeOx)<sub>59</sub>-*b*-P(PhBisOx-*cl/co*-ButOx), and (c) (PMeOx)<sub>62</sub>-*b*-P(PhBisOx-*cl/co*-ButOx). Error bars are based on SD (n=2).





**Figure S12.** Cytotoxicity of DOX-loaded a series of CCS POxs and free DOX in Hela cells for 48 h. (a) (PMeOx)<sub>44</sub>-*b*-P(PhBisOx-*cl/co*-ButOx), (b) (PMeOx)<sub>59</sub>-*b*-P(PhBisOx-*cl/co*-ButOx), and (c) (PMeOx)<sub>62</sub>-*b*-P(PhBisOx-*cl/co*-ButOx). Error bars are based on SD (n=2).