

Electronic Supplementary Information

Elucidating the effect of sequence and degree of polymerization on antimicrobial properties for block copolymers

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Table S1. Polymer characterization data

Polymer	M_n (g/mol)^a	<i>D</i>
1a	2600	1.09
1b	2450	1.13
2a	2650	1.15
2b	2630	1.09
3a	2370	1.10
3b	2630	1.11
4a	2460	1.10
4b	2750	1.07
5b	3340	1.09
6b	3290	1.09
7b	3990	1.08
8b	1910	1.13
9b	1700	1.11

^aMeasured relative to polystyrene standards.

Table S2. Antimicrobial results for polymers **1a–4a** and **1b–9b** in mmol/mL.

Polymer	<i>S. aureus</i> ^a	<i>E. coli</i> ^a	<i>K. pneumoniae</i> ^a	<i>A. baumannii</i> ^a	<i>P. aeruginosa</i> ^a	HEK293 ^b
Gram +ve/–ve	+	–	–	–	–	
1a: C ₂ -MA ₁₂ - <i>b</i> -Amine ₆	>197	>197	>197	>197	>197	>98
2a: C ₂ -Amine ₆ - <i>b</i> -MA ₁₂	>193	97	193	97	193	>97
3a: C ₂ -MA ₆ - <i>b</i> -Amine ₆ - <i>b</i> -MA ₆	>216	108	216	108	54	>108
4a: C ₂ -Amine ₃ - <i>b</i> -MA ₁₂ - <i>b</i> -Amine ₃	>208	52	104	26	104	>104
1b: C ₁₂ -MA ₁₂ - <i>b</i> -Amine ₆	>209	52	>209	52	104	>104
2b: C ₁₂ -Amine ₆ - <i>b</i> -MA ₁₂	97	12	24	12	49	29-45
3b: C ₁₂ -MA ₆ - <i>b</i> -Amine ₆ - <i>b</i> -MA ₆	195	24	195	49	97	63-97
4b: C ₁₂ -Amine ₃ - <i>b</i> -MA ₁₂ - <i>b</i> -Amine ₃	93	3	23	3	23	10-18
5b: C ₁₂ -Amine ₁ - <i>b</i> -MA ₁₂ - <i>b</i> -Amine ₁	>153	38-77	77-153	38-77	>153	33-75
6b: C ₁₂ -Amine ₂ - <i>b</i> -MA ₁₂ - <i>b</i> -Amine ₂	>156	19	39	19-39	78-156	17-38
7b: C ₁₂ -Amine ₆ - <i>b</i> -MA ₁₂ - <i>b</i> -Amine ₆	64	16	16	16	16	24-30
8b: C ₁₂ -Amine ₃ - <i>b</i> -MA ₆ - <i>b</i> -Amine ₃	8	8-4	17-8	8-4	8	46-103
9b: C ₁₂ -Amine ₃ - <i>b</i> -MA ₃ - <i>b</i> -Amine ₃	9	9	38-19	19-9	19-9	83-135

^aMinimum inhibitory concentration listed in mmol/mL. ^bCC₅₀ listed in mmol/mL

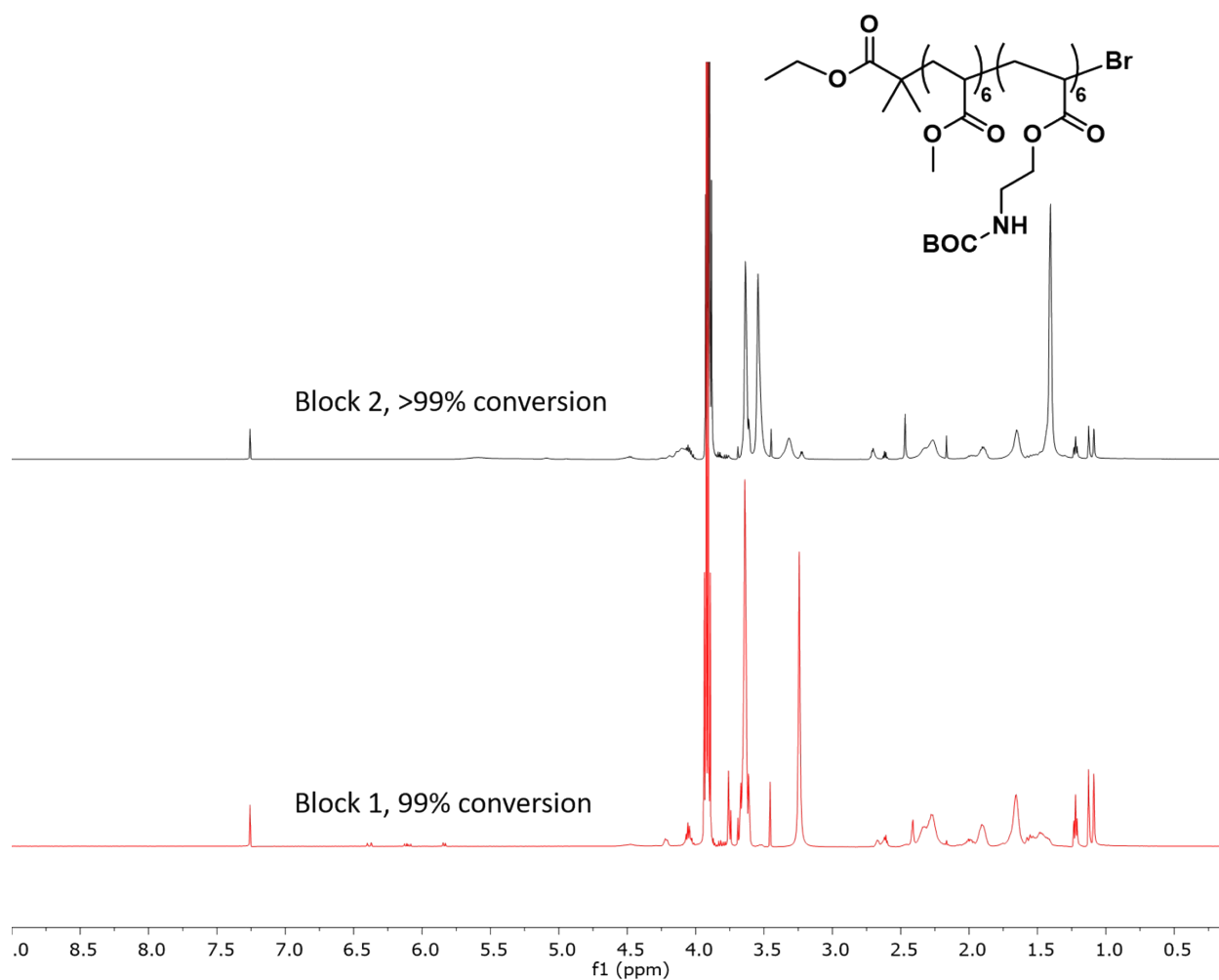


Figure S1. ^1H NMR spectrum of **1a** [EBiB-MA₁₂Am₆] block 1 in CDCl_3 showing high conversion throughout all monomer additions.

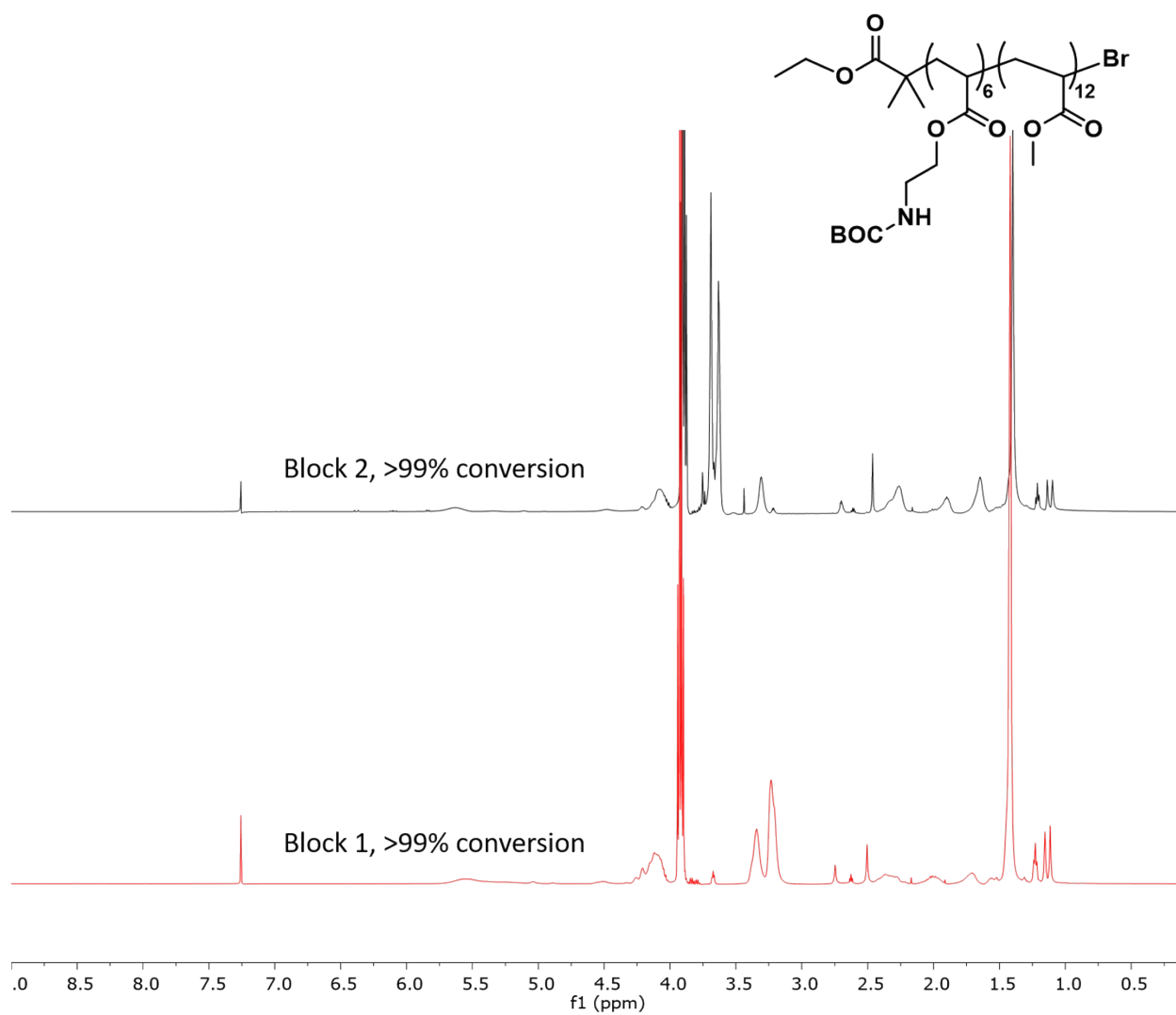


Figure S2. ¹H NMR spectrum of **2a** [EBiB-Am₆MA₁₂] block 1 in CDCl₃ showing high conversion throughout all monomer additions.

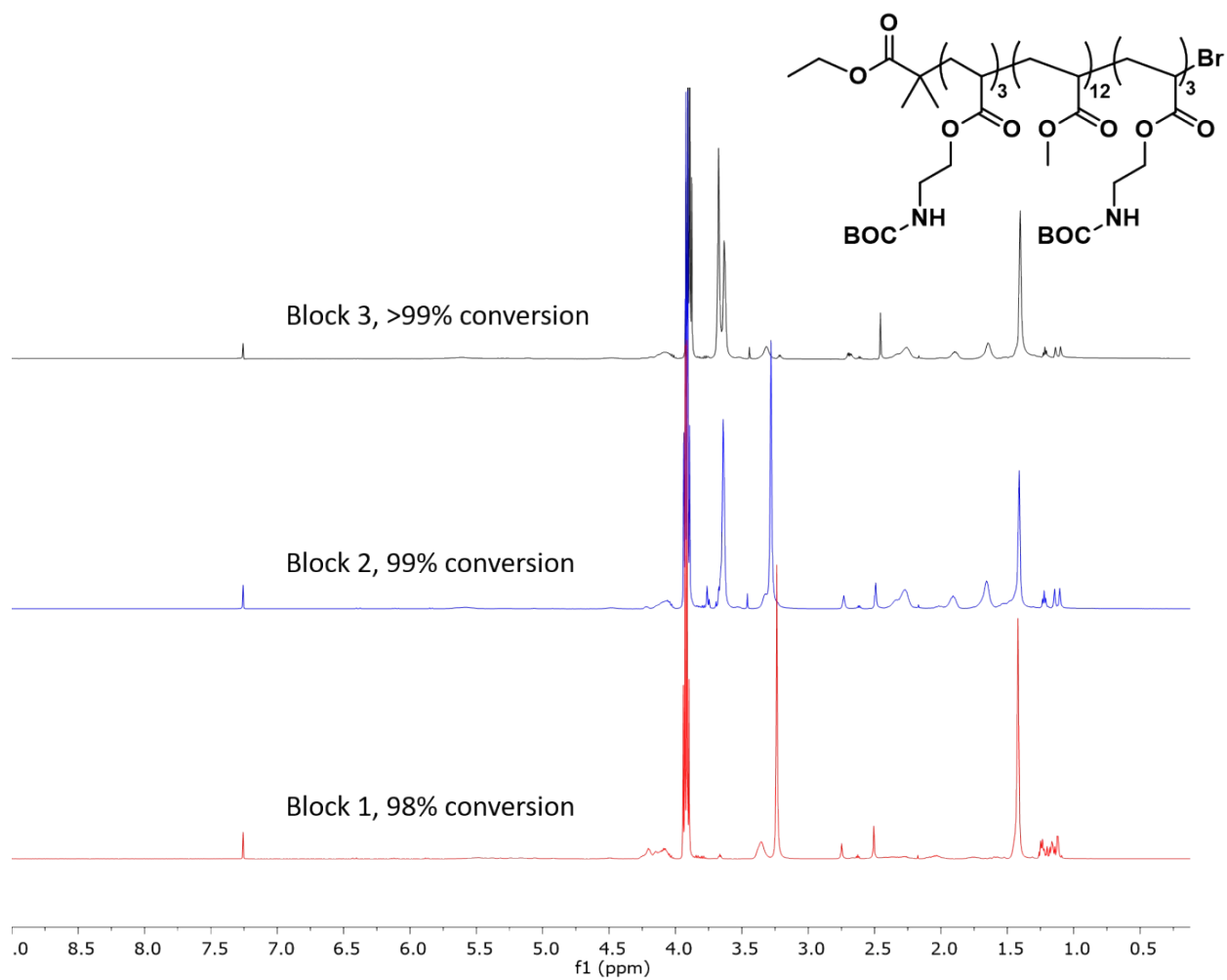


Figure S4. ^1H NMR spectrum of **4a** [EBiB-Am₃MA₁₂Am₃] block 1 in CDCl_3 showing high conversion throughout all monomer additions.

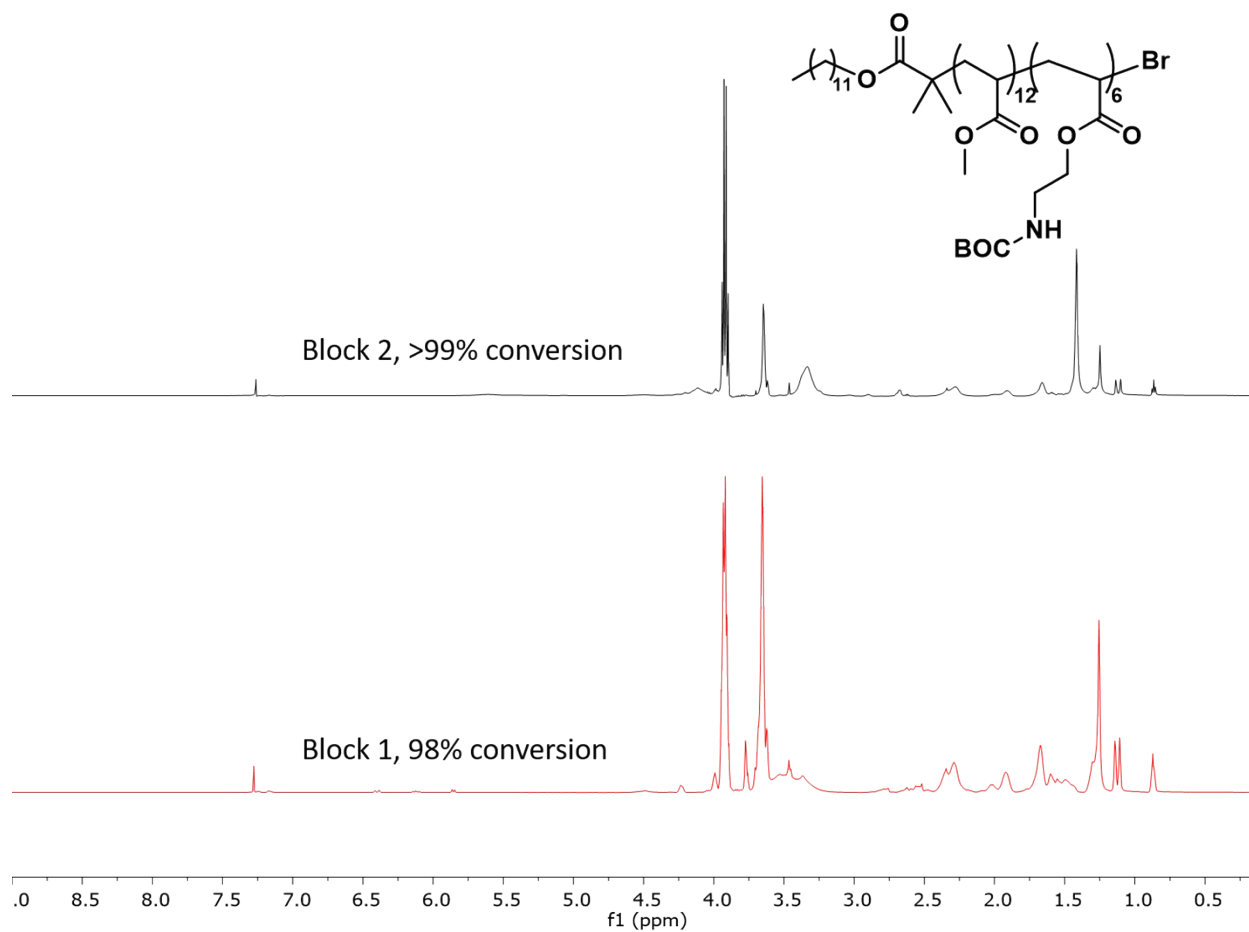


Figure S5. ¹H NMR spectrum of **1b** [DBiB-MA₁₂Am₆] block 1 in CDCl₃ showing high conversion throughout all monomer additions.

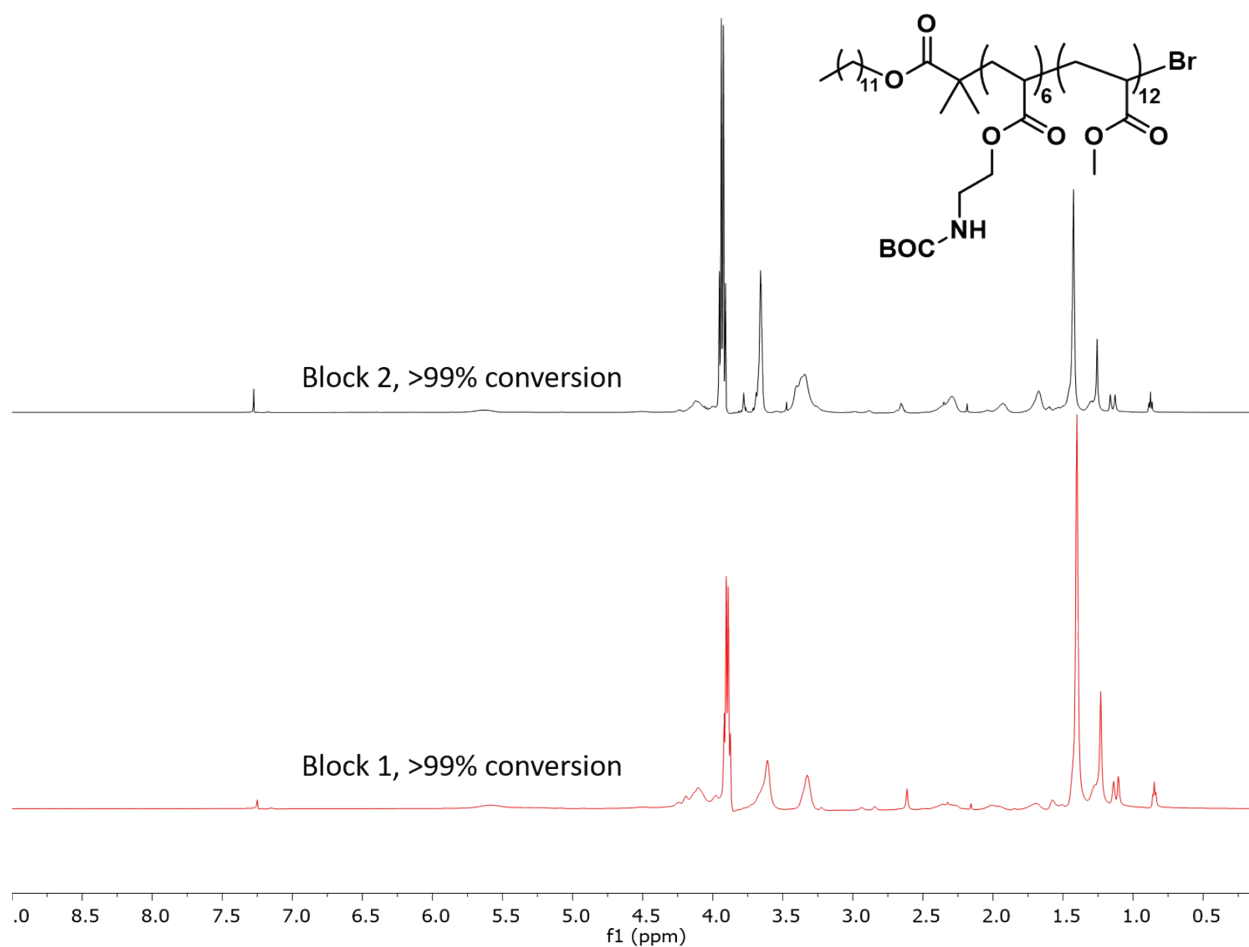
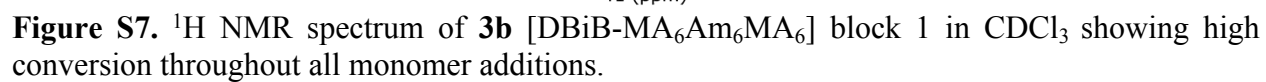


Figure S6. ¹H NMR spectrum of **2b** [DBiB-Am₆MA₁₂] block 1 in CDCl₃ showing high conversion throughout all monomer additions.



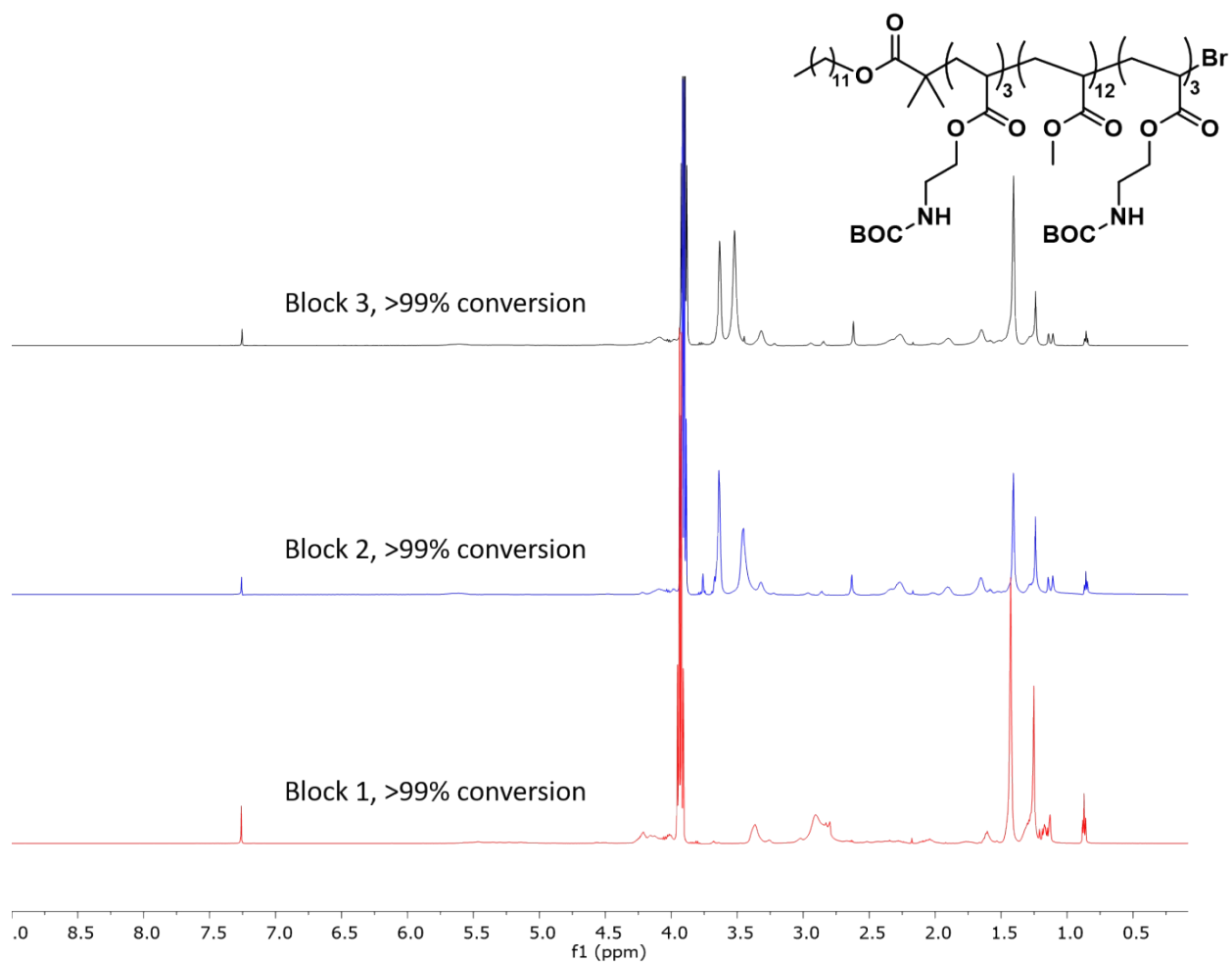


Figure S8. ^1H NMR spectrum of **4b** [DBiB- $\text{Am}_3\text{MA}_{12}\text{Am}_3$] block 1 in CDCl_3 showing high conversion throughout all monomer additions.

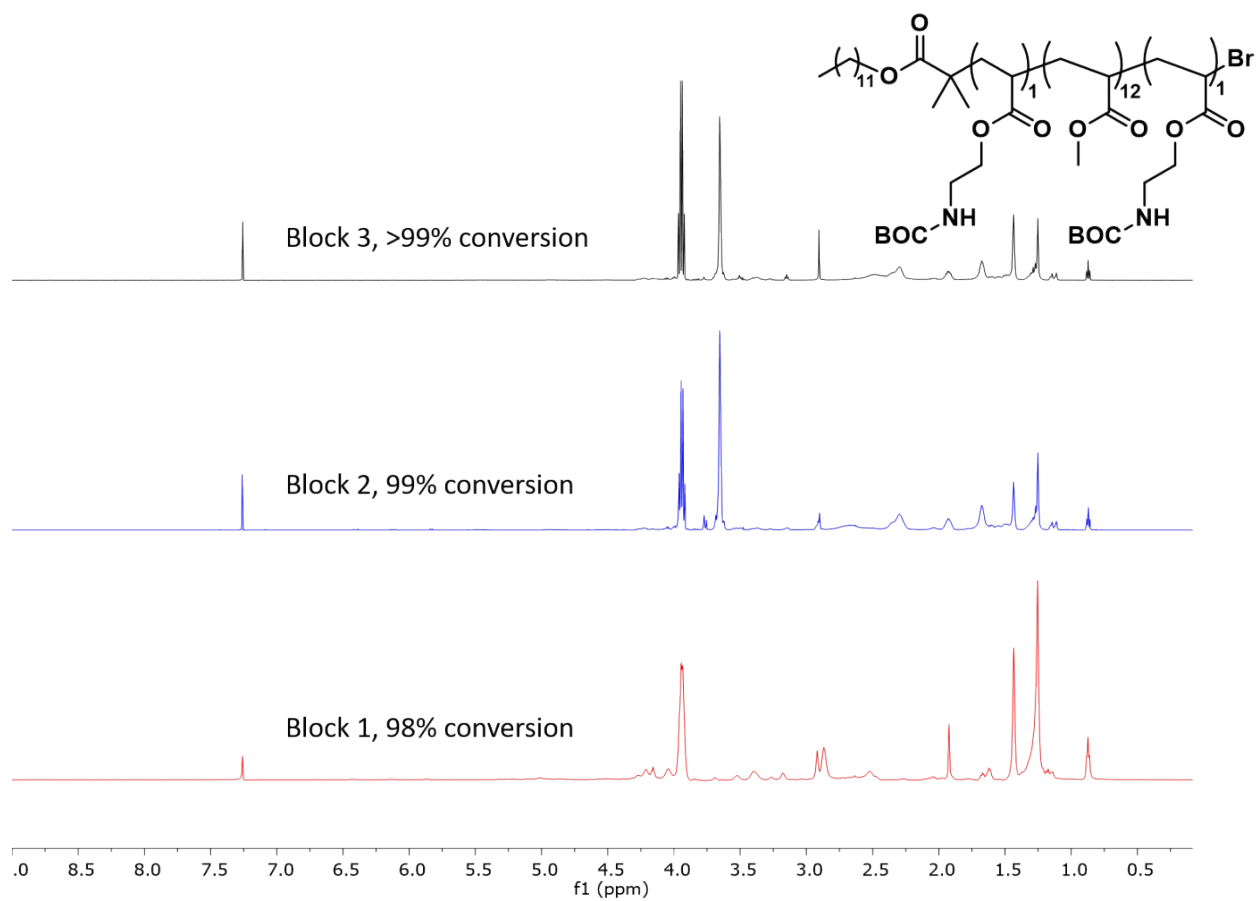


Figure S9. ^1H NMR spectrum of **5b** [DBiB-Am₁MA₁₂Am₁] block 1 in CDCl₃ showing high conversion throughout all monomer additions.

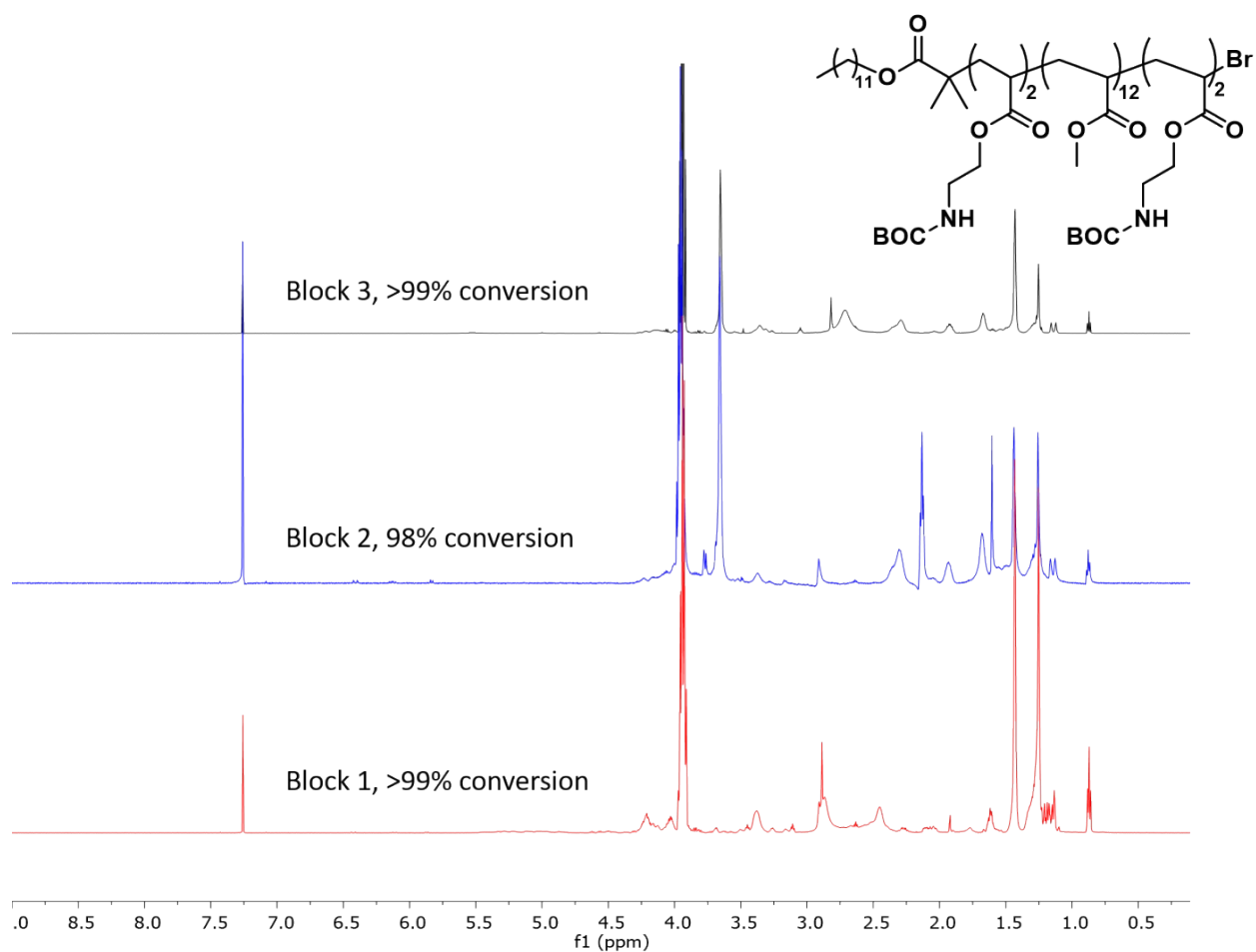


Figure S10. ^1H NMR spectrum of **6b** [DBiB- $\text{Am}_2\text{MA}_{12}\text{Am}_2$] block 1 in CDCl_3 showing high conversion throughout all monomer additions.

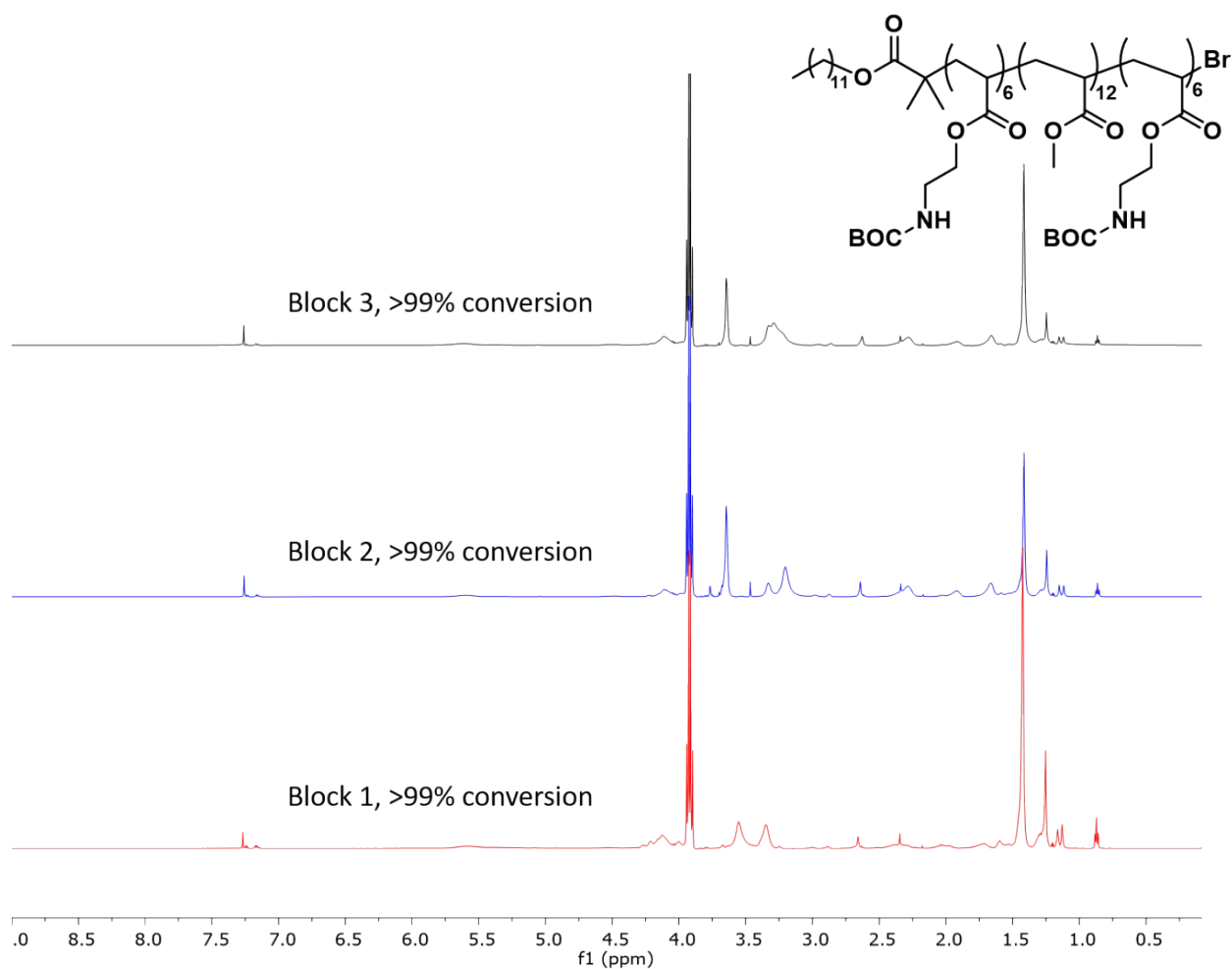


Figure S11. ¹H NMR spectrum of **7b** [DBiB-Am₆MA₁₂Am₆] block 1 in CDCl₃ showing high conversion throughout all monomer additions.

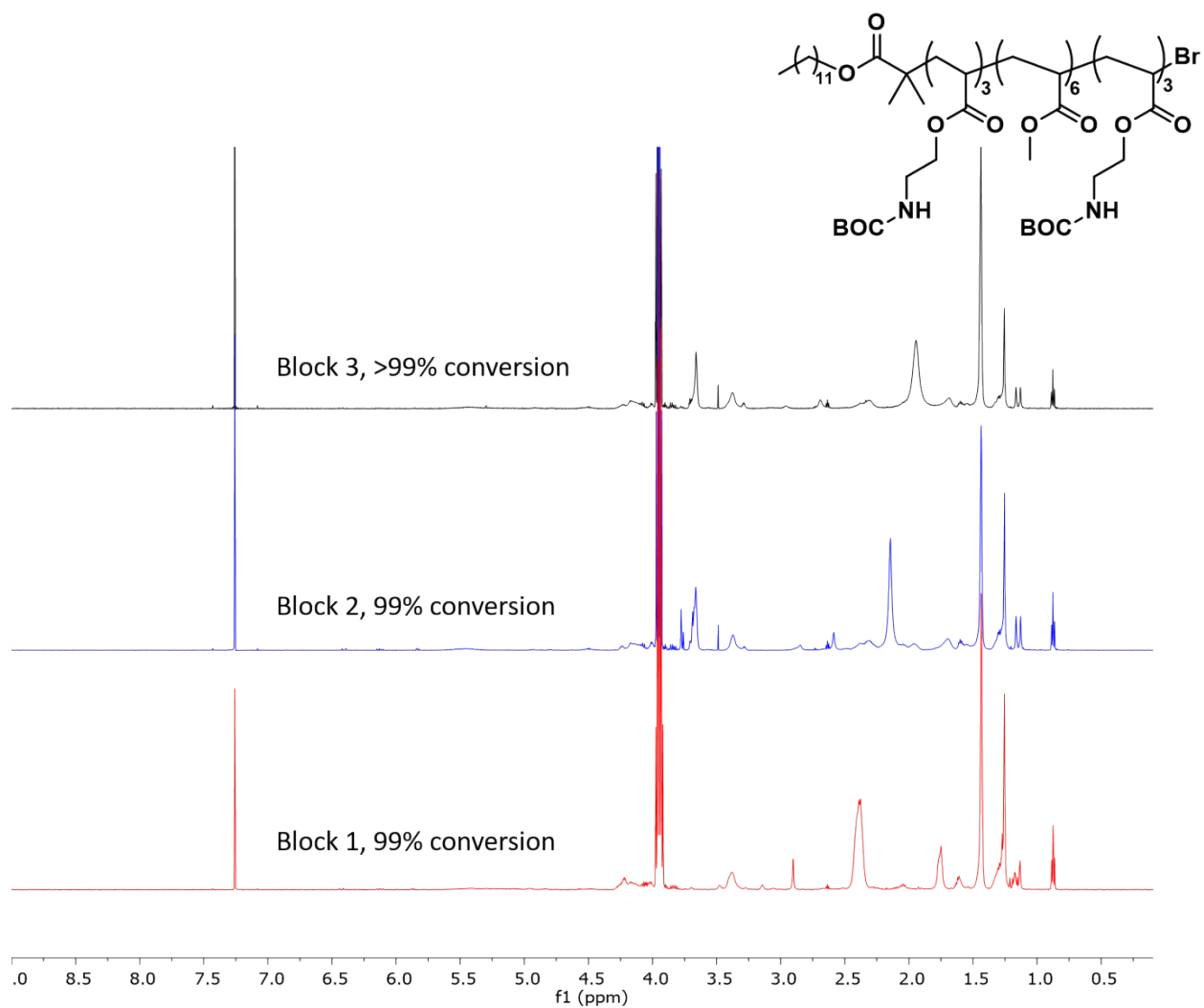


Figure S12. ^1H NMR spectrum of **8b** [DBiB-Am₃MA₃Am₃] block 1 in CDCl₃ showing high conversion throughout all monomer additions.

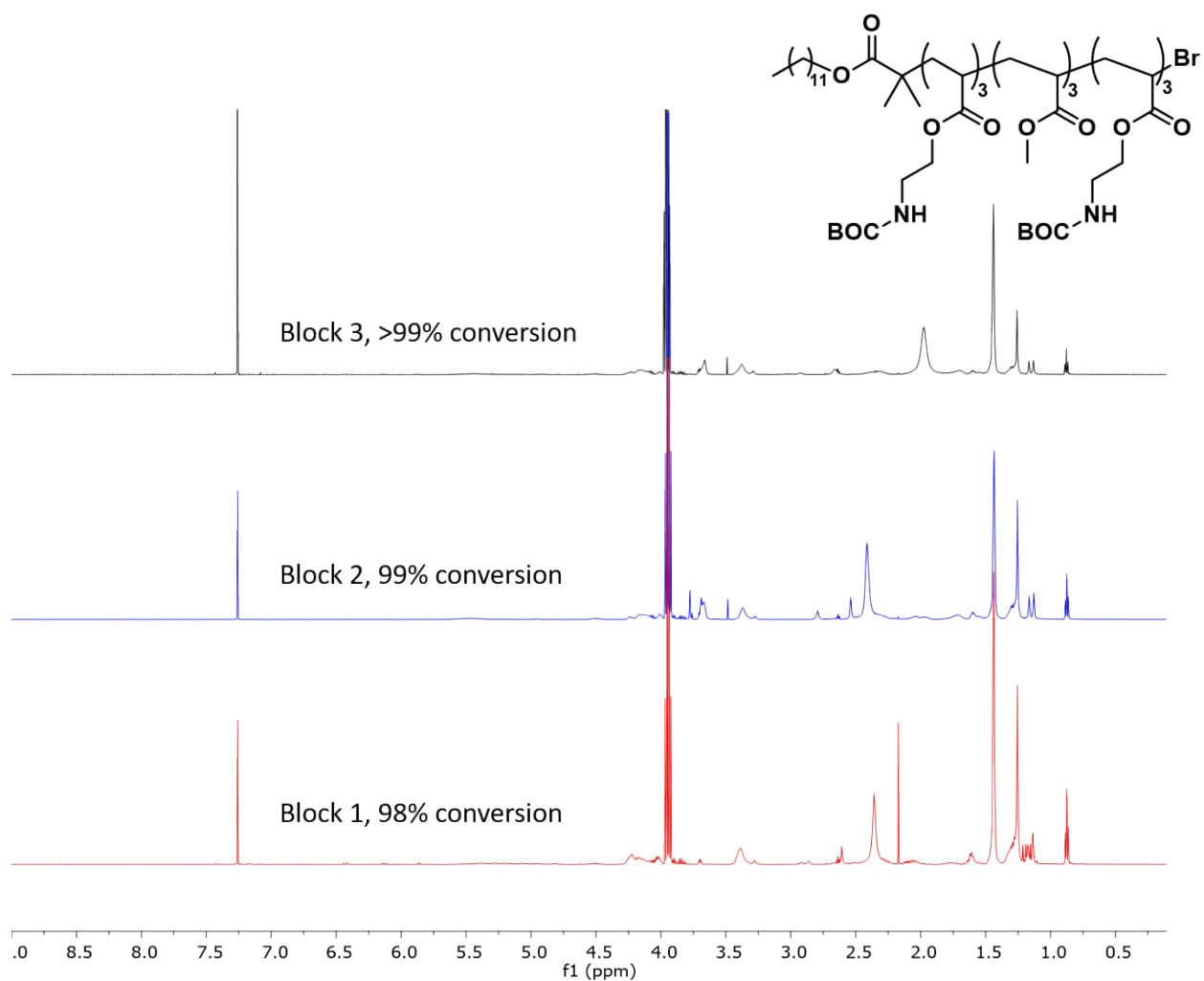


Figure S13. ¹H NMR spectrum of **9b** [DBiB-Am₃MA₆Am₃] block 1 in CDCl₃ showing high conversion throughout all monomer additions.

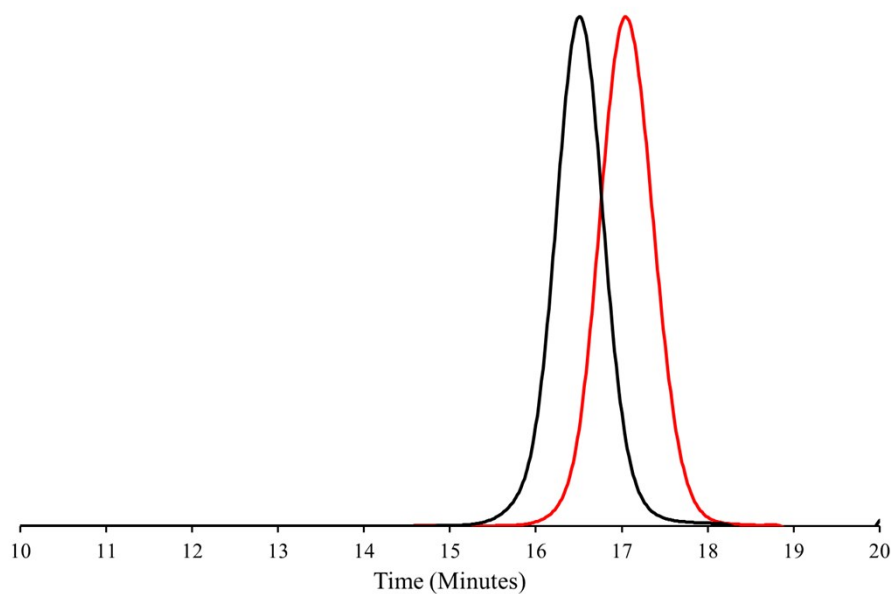


Figure S14. SEC traces of **1a** [EBiB-MA₁₂Am₆] block 1 (red) and block 2 (black).

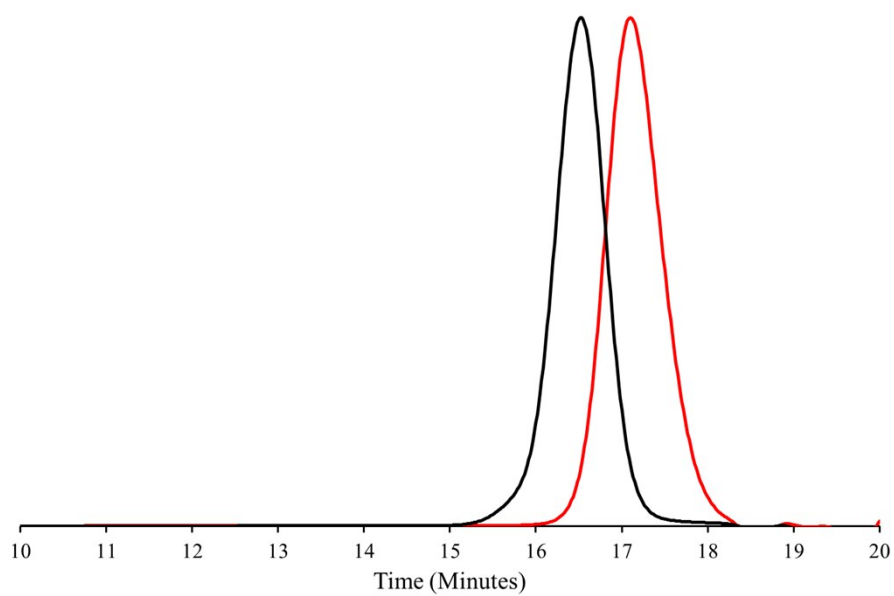


Figure S15. SEC traces of **2a** [EBiB-Am₆MA₁₂] block 1 (red) and block 2 (black).

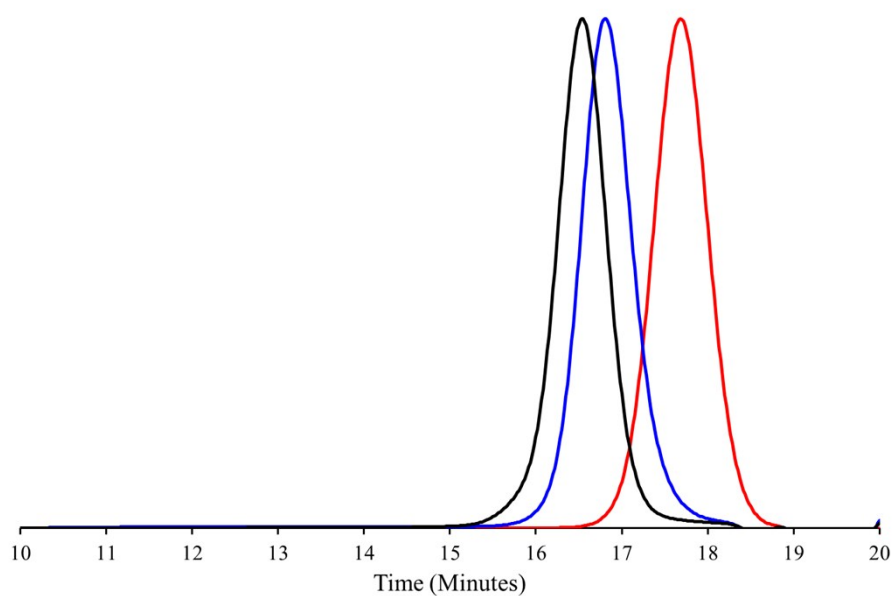


Figure S16. SEC traces of **3a** [EBiB-MA₆Am₆MA₆] block 1 (red), block 2 (blue) and block 3 (black).

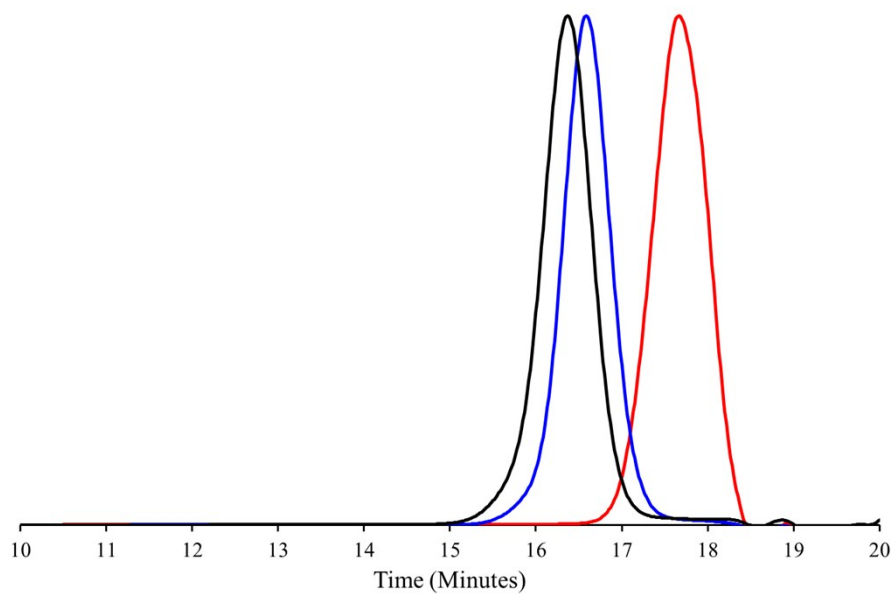


Figure S17. SEC traces of **4a** [EBiB-Am₃MA₁₂Am₃] block 1 (red), block 2 (blue) and block 3 (black).

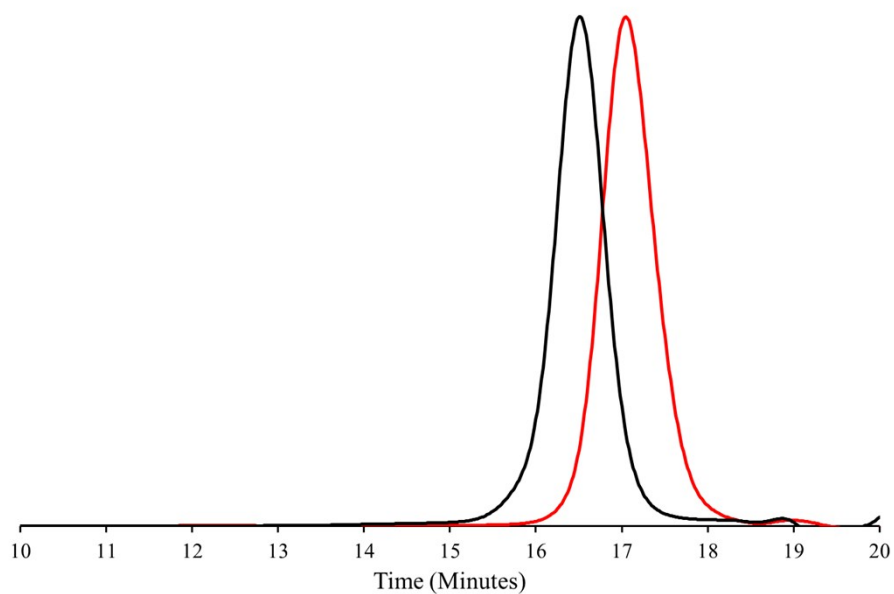


Figure S18. SEC traces of **1b** [DBiB-MA₁₂Am₆] block 1 (red) and block 2 (black).

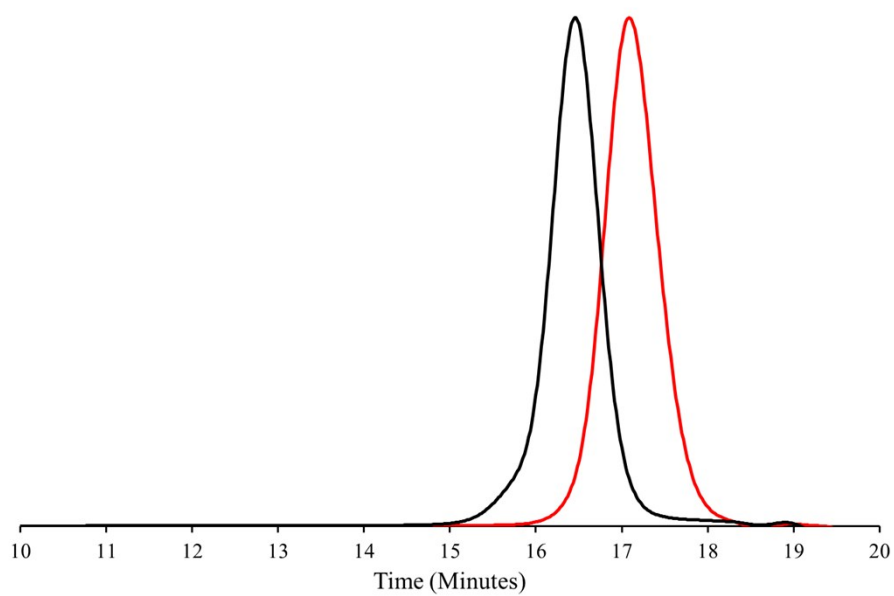


Figure S19. SEC traces of **2b** [DBiB-Am₆MA₁₂] block 1 (red) and block 2 (black).

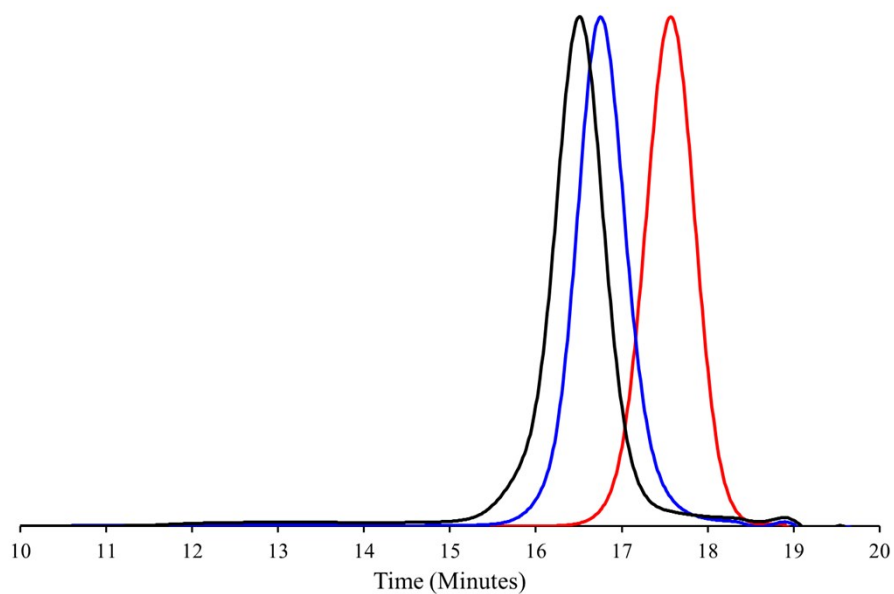


Figure S20. SEC traces of **3b** [DBiB-MA₆Am₆MA₆] block 1 (red), block 2 (blue) and block 3 (black).

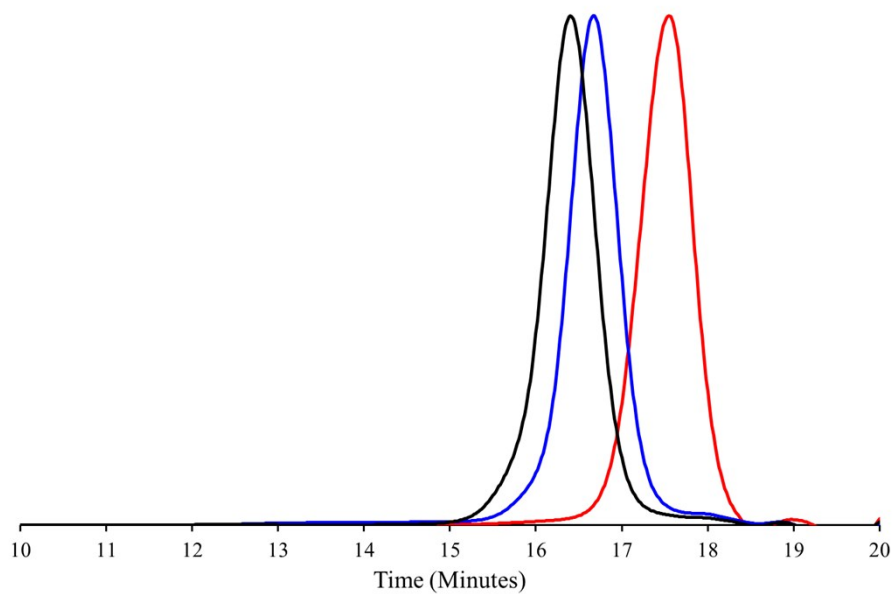


Figure S21. SEC traces of **4b** [DBiB-Am₃MA₁₂Am₃] block 1 (red), block 2 (blue) and block 3 (black).

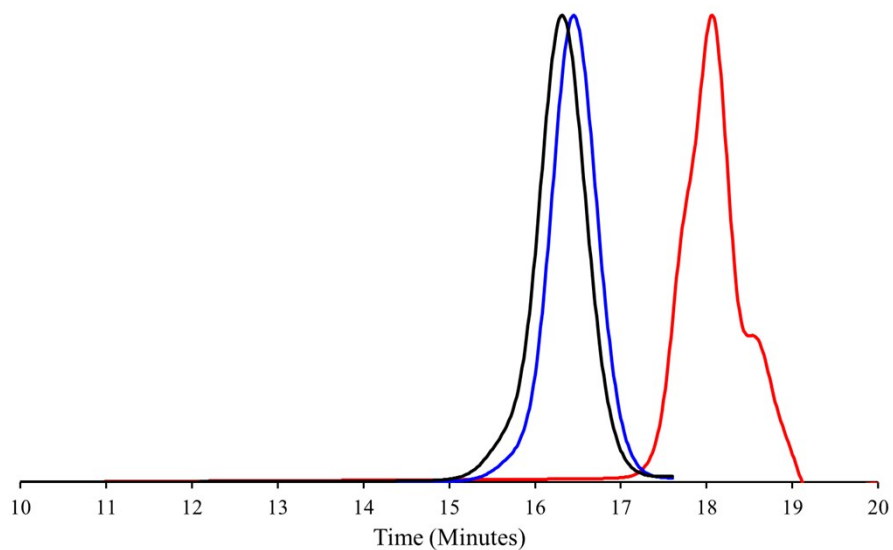


Figure S22. SEC traces of **5b** [DBiB-Am₁MA₁₂Am₁] block 1 (red), block 2 (blue) and block 3 (black).

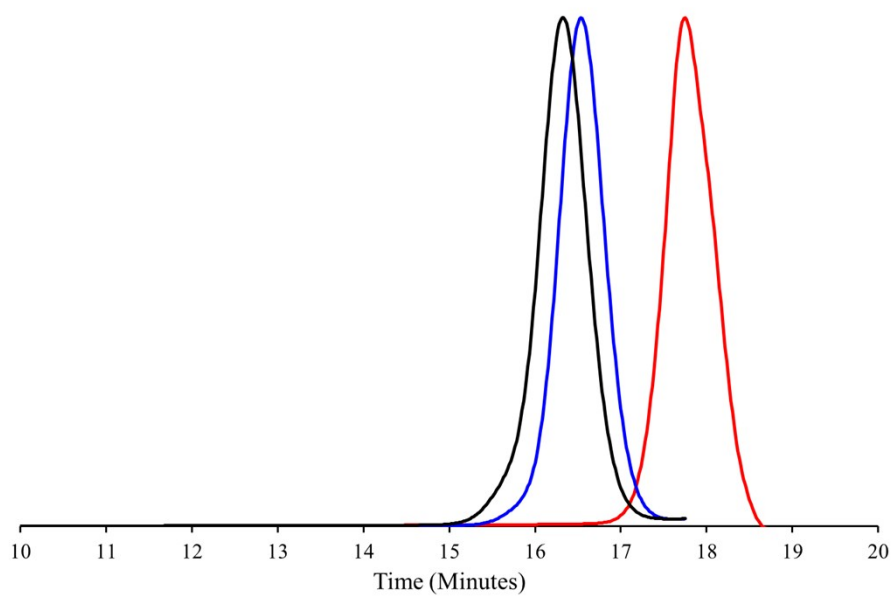


Figure S23. SEC traces of **6b** [DBiB-Am₂MA₁₂Am₂] block 1 (red), block 2 (blue) and block 3 (black).

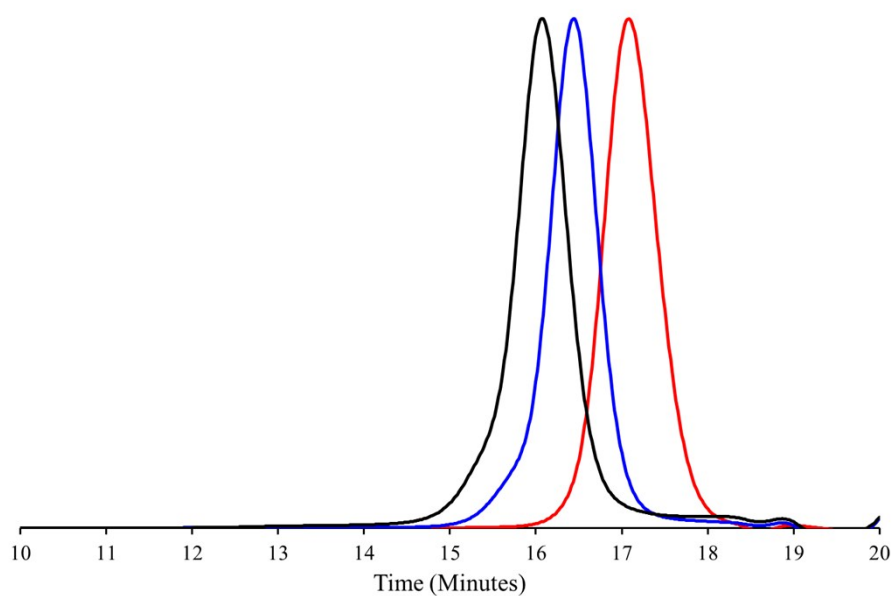


Figure S24. SEC traces of **7b** [DBiB-Am₆MA₁₂Am₆] block 1 (red), block 2 (blue) and block 3 (black).

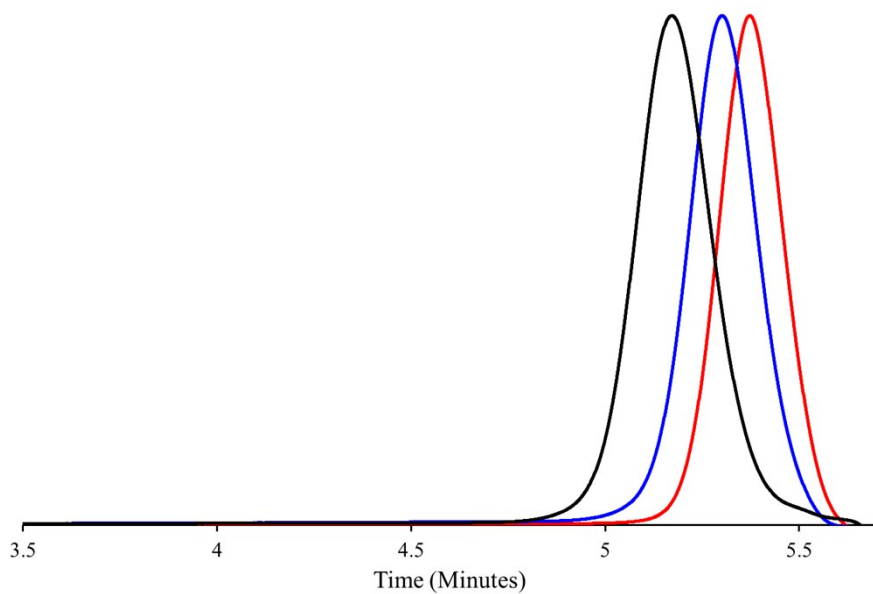


Figure S25. SEC traces of **8b** [DBiB-Am₃MA₃Am₃] block 1 (red), block 2 (blue) and block 3 (black).

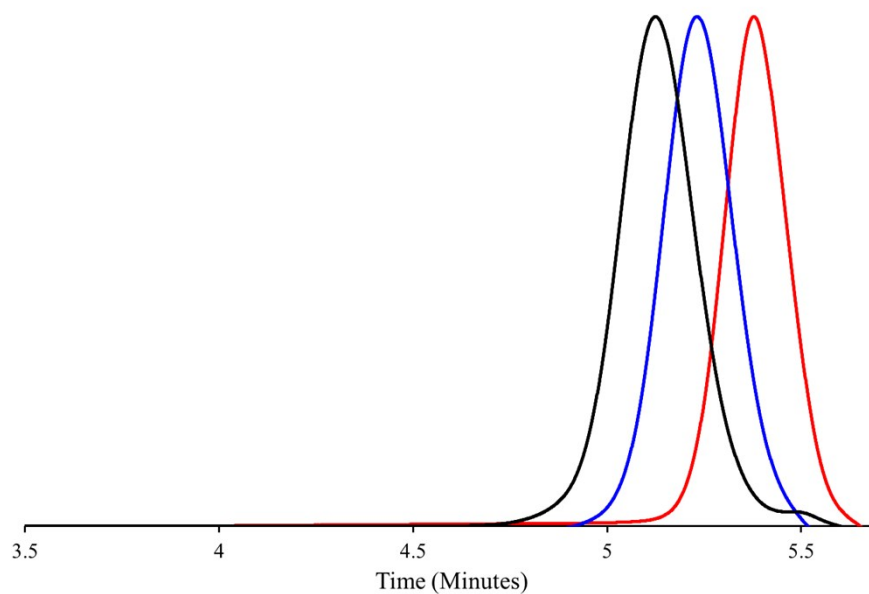


Figure S26. SEC traces of **9b** [DBiB-Am₃MA₆Am₃] block 1 (red), block 2 (blue) and block 3 (black).

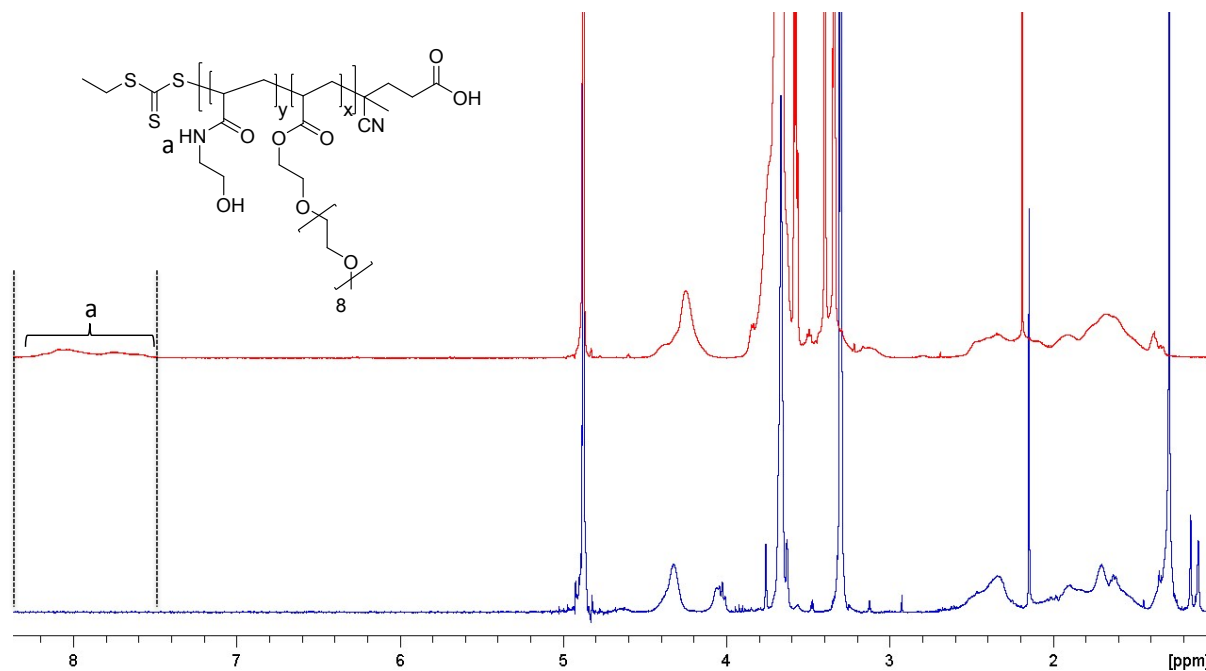


Figure S27. ¹H NMR of P(PEGA-co-HEAA)-SC(=S)SC₂H₅ (published in Polym. Chem., 2015, 6, 3865–3874) (top red) and polymer **1b** (DBiB-MA₁₂Am₆) reported in this work (bottom blue), indicating no transamination has occurred.