Supplementary Materials For

Synthesis of Large Dendrimers with the Dimensions of Small Viruses

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General Synthetic Procedures

All chemicals were purchased from Aldrich and Acros and used without further purification. All solvents were ACS grade and used without further purification. HPLC was carried out using an Agilent Technologies 1260 Infinity system and an Agilent Technologies 1260 Infinity DAD detector. NMR spectra were recorded on a Mercury 300 MHz spectrometer in CDCl₃. All mass spectral analyses were carried out by an Agilent Technologies 6224 TOF LC/MS system.

Scheme S1. Synthesis of the Dendrimers

Details of Synthetic Procedures

Compound 1 (Boc-protected G3). A solution of 14 (1.20 g, 0.61 mmol), 16 (0.12 g, 0.074 mmol), and DIPEA (0.30 mL, 1.71 mmol) in THF (3 mL), methanol (0.3 mL), and H_2O (0.3 mL) was stirred at 75 °C in a capped vessel

for 4 d. The solution was evaporated under vacuum. The residue was dissolved in dichloromethane, washed with brine, dried over MgSO₄, filtered, and evaporated under vacuum. The crude product was purified by silica gel chromatography (from EA:DCM:MeOH = 6:6:1 to DCM:MeOH = 7:1) to give **1** (0.60 g, 86%) as a white wax.

¹H NMR (300 MHz, CDCl₃) δ 4.05 (br, 2H, HC=CCH₂), 3.51-3.29 (m, 448H, CH₂OCH₂CH₂OCH₂CH₂OCH₂CH₂OCH₂, C₃N₃-NHCH₂CH₂CH₂O), 3.08 (br m, 32H, BocNHCH₂), 2.17 (br, 1H, HC=CCH₂), 1.69-1.60 (m, 120H, OCH₂CH₂CH₂), 1.31 (s, 144H, C(CH₃)₃);

¹³C NMR (75 MHz, CDCl₃) δ 165.7 (**C**₃N₃), 155.9 (**C**O), 81.1 (not found, HC=CCH₂), 78.6 (**C**(CH₃)₃), 70.5 (not found, HC=CCH₂), 70.4 (OCH₂CH₂O), 70.0 (two lines, OCH₂CH₂O), 69.3 (CH₂CH₂CH₂O), 69.1 (CH₂CH₂CH₂O), 69.0 (CH₂CH₂CH₂O), 38.3 (CH₂CH₂CH₂O), 37.8 (CH₂CH₂CH₂O), 30.1 (not found, HC=CCH₂), 29.5 (NHCH₂CH₂CH₂O), 28.3 (C(CH₃)₃); MS (ESI-TOF) calcd for C₄₂₈H₈₀₈N₁₀₆O₁₂₂ 9386.03, found 9395.31 (M + H)⁺.

Compound 2 (deprotected G3). A solution of 1 (0.27 g, 0.029 mmol) in concentrated HCl (2 mL) and methanol (4 mL) was stirred for 16 h at room temperature and then evaporated under vacuum. The residue was dissolved in chloroform, washed with 5 M NaOH (aq), dried over MgSO₄, filtered, and evaporated under vacuum to give 2 (0.22 g, quantitative) as a white wax. ¹H NMR (300 MHz, CDCl₃) δ 4.12 (br, 2H, HC=CCH₂), 3.60-3.38 (m, 448H, CH₂OCH₂CH₂OCH₂CH₂OCH₂, C₃N₃-NHCH₂CH₂CH₂O), 2.78 (br, 32H, OCH₂CH₂CH₂OH₂), 2.22 (br, 1H, HC=CCH₂), 1.78-1.69 (m, 120H, OCH₂CH₂CH₂); ¹³C NMR (75 MHz, CDCl₃) δ 166.1 (C₃N₃), 81.1 (not found, HC=CCH₂), 70.6 (OCH₂CH₂O), 70.5 (HC=CCH₂), 70.3 (OCH₂CH₂O), 70.2 (OCH₂CH₂O) 69.5 (CH₂CH₂CH₂O), 69.4 (CH₂CH₂CH₂O), 69.3 (CH₂CH₂CH₂O), 39.6 (CH₂CH₂CH₂O), 38.1 (CH₂CH₂CH₂O), 32.9 (OCH₂CH₂CH₂NH₂), 30.1 (not found, HC=CCH₂), 29.7 (NHCH₂CH₂CH₂O); MS (ESI-TOF) calcd for C₃₄₈H₆₈₀N₁₀₆O₉₀ 7785.19, found 7789.62 (M + H)⁺.

Compound 3 (Boc-protected G5). A solution of 14 (1.0 g, 0.50 mmol), 2 (0.11 g, 0.014 mmol), and DIPEA (0.30 mL, 1.71 mmol) in THF (5 mL), methanol (0.5 mL), and H₂O (0.5 mL) was stirred at 75 °C in a capped vessel for 4 d. The solution was evaporated under vacuum. The residue was dissolved in dichloromethane, washed with brine, dried over MgSO₄, filtered, and evaporated under vacuum. The crude product was dissolved in dichloromethane and precipitated by adding diethyl ether. The precipitation step was repeated until macromonomer 14 was completely removed, which was monitored by thin layer chromatography (DCM:MeOH = 14:1). The pure product 3 (0.44 g, 80%) was obtained as a white wax. 1 H NMR (300 MHz, CDCl₃) δ 4.05 (not found, 2H, HC=CCH₂), 3.53-3.32 (m, 1888H, CH₂OCH₂CH₂OCH₂CH₂OCH₂CH₂OCH₂, C₃N₃-NHCH₂CH₂CH₂O), 3.11 (br m, 128H, BocNHCH₂), 2.17 (not found, 1H, HC=CCH₂), 1.72-1.63 (m, 504H, OCH₂CH₂CH₂), 1.33 (s, 576H, C(CH₃)₃); 13 C NMR (75 MHz, CDCl₃) δ 165.1 (br, C₃N₃), 156.0 (CO), 81.1 (not found, HC=CCH₂), 78.7 (C(CH₃)₃), 70.5 (not found, HC=CCH₂), 70.4 (OCH₂CH₂O), 70.1 (three lines, OCH₂CH₂O), 69.4 (CH₂CH₂CH₂O), 69.1 (CH₂CH₂CH₂O), 38.4 (CH₂CH₂CH₂O), 38.0 (CH₂CH₂CH₂O), 30.1 (not found, HC=CCH₂), 29.4 (NHCH₂CH₂CH₂O), 28.4 (C(CH₃)₃); MS (ESI-TOF) calcd for C₁₇₇₂H₃₃₅₂N₄₄₂O₅₆ 38925.0, found 38952.4 (M + H)⁺.

Compound 4 (deprotected G5). A solution of 3 (0.40 g, 0.0103 mmol) in concentrated HCl (3 mL) and methanol (6 mL) was stirred for 16 h at room temperature and then evaporated under vacuum. The residue was dissolved in chloroform, washed with 5 M NaOH (aq), dried over MgSO₄, filtered, and evaporated under vacuum to give 4 (0.30 g, 90%) as a white wax. 1 H NMR (300 MHz, CDCl₃) δ 4.12 (not found, 2H, HC \equiv CCH₂), 3.55-3.34 (m, 1888H, CH₂OCH₂CH₂OCH₂CH₂OCH₂, C₃N₃-NHCH₂CH₂CH₂O), 2.71 (br, 128H, OCH₂CH₂CH₂CH₂NH₂), 2.22 (not found, 1H, HC \equiv CCH₂), 1.75-1.62 (m, 504H, OCH₂CH₂CH₂); 13 C NMR (75 MHz, CDCl₃) δ 166.0 (C₃N₃), 81.1 (not found, HC \equiv CCH₂), 70.5 (HC \equiv CCH₂, OCH₂CH₂O), 70.2 (OCH₂CH₂O), 69.4 (CH₂CH₂CH₂O), 69.2 (CH₂CH₂CH₂O), 39.5 (CH₂CH₂CH₂O), 38.0 (CH₂CH₂CH₂O), 33.3 (OCH₂CH₂CH₂NH₂), 30.1 (not found, HC \equiv CCH₂), 29.6 (NHCH₂CH₂CH₂O); MS (ESI-TOF) calcd for C₁₄₅₂H₂₈₄₀N₄₄₂O₃₇₈ 32521.7, not found.

Compound 5 (Boc-protected G7). A solution of 14 (1.0 g, 0.504 mmol), 4 (0.10 g, 3.07 μmol), and DIPEA (0.30 mL, 1.71 mmol) in THF (5 mL), methanol (0.5 mL), and H₂O (0.5 mL) was stirred at 75 °C in a capped vessel for 4 d. The solution was evaporated under vacuum. The residue was dissolved in dichloromethane, washed with brine, dried over MgSO₄, filtered, and evaporated under vacuum. The crude product was dissolved in dichloromethane and precipitated by adding diethyl ether. The precipitation step was repeated until macromonomer 14 was completely removed, which was monitored by thin layer chromatography (DCM:MeOH = 14:1). The pure product 5 (0.39 g, 81%) was obtained as a white wax. 1 H NMR (300 MHz, CDCl₃) δ 4.05 (not found, 2H, HC=CCH₂), 3.58-3.36 (m, 7648H, CH₂OCH₂CH₂OCH₂CH₂OCH₂CH₂OCH₂, C₃N₃-NHCH₂CH₂CH₂O), 3.16 (br m, 512H, BocNHCH₂), 2.17 (not found, 1H, HC=CCH₂), 1.76-1.67 (m, 2040H, OCH₂CH₂CH₂), 1.37 (s, 2304H, C(CH₃)₃); 13 C NMR (75 MHz, CDCl₃) δ 165.9 (C₃N₃), 156.1 (CO), 81.1 (not found, HC=CCH₂), 78.8 (C(CH₃)₃), 70.6 (OCH₂CH₂O), 70.5 (not found, HC=CCH₂), 70.3 (OCH₂CH₂O), 70.2 (two lines, OCH₂CH₂O), 69.5 (CH₂CH₂CH₂O), 69.3 (CH₂CH₂CH₂O), 69.2 (CH₂CH₂CH₂O), 38.5 (CH₂CH₂CH₂O), 38.1 (CH₂CH₂CH₂O), 30.1 (not found, HC=CCH₂), 29.6 (NHCH₂CH₂CH₂O), 28.5 (C(CH₃)₃); MS (ESI-TOF) calcd for C₇₁₄₈H₁₃₅₂₈N₁₇₈₆O₂₀₄₂ 157081.0, not found.

Compound 6 (deprotected G7). A solution of 5 (0.34 g, 2.16 μmol) in concentrated HCl (4 mL) and methanol (8 mL) was stirred for 16 h at room temperature and then evaporated under vacuum. The residue was dissolved in chloroform, washed with 5 M NaOH (aq), dried over MgSO₄, filtered, and evaporated under vacuum to give 6 (0.26 g, 92%) as a white wax. ¹H NMR (300 MHz, CDCl₃) δ 4.12 (not found, 2H, HC=CCH₂), 3.57-3.34 (m, 7648H, CH₂OCH₂CH₂OCH₂CH₂OCH₂CH₂CH₂CH₂CH₂CH₂O), 2.72 (t, J = 6.6, 512H, OCH₂CH₂CH₂NH₂), 2.22 (not found, 1H, HC=CCH₂), 1.75-1.63 (m, 2040H, OCH₂CH₂CH₂CH₂); ¹³C NMR (75 MHz, CDCl₃) δ 166.1 (C₃N₃), 81.1 (not found, HC=CCH₂), 70.6 (OCH₂CH₂O), 70.5 (HC=CCH₂, OCH₂CH₂O), 70.2 (OCH₂CH₂O), 70.1 (OCH₂CH₂O), 69.4 (CH₂CH₂CH₂O), 69.3 (CH₂CH₂CH₂O), 69.2 (CH₂CH₂CH₂O), 39.5 (CH₂CH₂CH₂O), 38.0 (CH₂CH₂CH₂O), 33.4 (OCH₂CH₂CH₂NH₂), 30.1 (not found, HC=CCH₂), 29.6 (NHCH₂CH₂CH₂O); MS (ESI-TOF) calcd for C₅₈₆₈H₁₁₄₈₀N₁₇₈₆O₁₅₃₀ 131467.5, not found.

Compound 7 (Boc-protected G9). A solution of 14 (1.0 g, 0.504 mmol), 6 (0.10 g, 0.76 μmol), and DIPEA (0.30 mL, 1.71 mmol) in THF (5 mL), methanol (0.5 mL), and H₂O (0.5 mL) was stirred at 75 °C in a capped vessel for 4 d. The solution was evaporated under vacuum. The residue was dissolved in dichloromethane, washed with brine, dried over MgSO₄, filtered, and evaporated under vacuum. The crude product was dissolved in dichloromethane and precipitated by adding diethyl ether. The precipitation step was repeated until macromonomer 14 was completely removed, which was monitored by thin layer chromatography (DCM:MeOH = 14:1). The pure product 7 (0.39 g, 81%) was obtained as a white wax. ¹H NMR (300 MHz, CDCl₃) δ 4.05 (not found, 2H, HC=CCH₂), 3.58-3.36 (m, 30688H, CH₂OCH₂CH₂OCH₂CH₂OCH₂, C₃N₃-NHCH₂CH₂CH₂O), 3.16 (br m, 2048H, BocNHCH₂), 2.17 (not found, 1H, HC=CCH₂), 1.76-1.67 (m, 8184H, OCH₂CH₂CH₂CH₂), 1.38 (s, 9216H, C(CH₃)₃); ¹³C NMR (75 MHz, CDCl₃) δ 165.9 (C₃N₃), 156.1 (CO), 81.1 (not found, HC=CCH₂), 78.8 (C(CH₃)₃), 70.6 (OCH₂CH₂O), 70.5 (not found, HC=CCH₂), 70.3 (OCH₂CH₂O), 70.2 (OCH₂CH₂O), 69.5 (CH₂CH₂CH₂O), 69.3 (CH₂CH₂CH₂O), 69.2 (CH₂CH₂CH₂O), 38.5 (CH₂CH₂CH₂O), 38.1 (CH₂CH₂CH₂O), 30.1 (not found, HC=CCH₂), 29.6 (NHCH₂CH₂CH₂O), 28.5 (C(CH₃)₃); MS (ESI-TOF) calcd for C₂₈₆₅₂H₅₄₂₃₂N₇₁₆₂O₈₁₈₆ 629704.8, not found.

Compound 8 (deprotected G9). A solution of 7 (0.33 g, 0.524 μmol) in concentrated HCl (4 mL) and methanol (8 mL) was stirred for 16 h at room temperature and then evaporated under vacuum. The residue was dissolved in chloroform, washed with 5 M NaOH (aq), dried over MgSO₄, filtered, and evaporated under vacuum to give 8 (0.255 g, 92%) as a white wax. ¹H NMR (300 MHz, CDCl₃) δ 4.12 (not found, 2H, HC≡CCH₂), 3.58-3.36 (m, 30688H, CH₂OCH₂CH₂OCH₂CH₂OCH₂, C₃N₃-NHCH₂CH₂CH₂O), 2.73 (t, J = 6.6, 2048H, OCH₂CH₂CH₂NH₂), 2.22 (not found, 1H, HC≡CCH₂), 1.76-1.65 (m, 8184H, OCH₂CH₂CH₂); ¹³C NMR (75 MHz, CDCl₃) δ 166.3 (C₃N₃), 81.1 (not found, HC≡CCH₂), 70.6 (two lines, OCH₂CH₂O), 70.5 (not found, HC≡CCH₂), 70.2 (two lines, OCH₂CH₂O), 69.4 (CH₂CH₂CH₂O), 69.3 (two lines, CH₂CH₂CH₂O), 39.6 (CH₂CH₂CH₂O), 38.1 (CH₂CH₂CH₂O), 33.3 (OCH₂CH₂CH₂NH₂), 30.1 (not found, HC≡CCH₂), 29.7 (NHCH₂CH₂CH₂O); MS (ESI-TOF) calcd for C₂₃₅₃₂H₄₆₀₄₀N₇₁₆₂O₆₁₃₈ 527251.1, not found.

Compound 9 (Boc-protected G11). A solution of 14 (1.0 g, 0.504 mmol), 8 (0.10 g, 0.19 μ mol), and DIPEA (0.30 mL, 1.71 mmol) in THF (5 mL), methanol (0.5 mL), and H₂O (0.5 mL) was stirred at 75 °C in a capped vessel for 4 d. The solution was evaporated under vacuum. The residue was dissolved in dichloromethane, washed with brine, dried over MgSO₄, filtered, and evaporated under vacuum. The crude product was dissolved in dichloromethane and precipitated by adding diethyl ether. The precipitation step was repeated until macromonomer 14 was completely removed, which was monitored by thin layer chromatography (DCM:MeOH = 14:1). The pure product 9 (0.39 g, 82%) was obtained as a white wax. 1 H NMR (300 MHz, CDCl₃) δ 4.05 (not found, 2H, HC=CCH₂), 3.61-3.39 (m, 122848H, CH₂OCH₂CH₂OCH₂CH₂OCH₂, C₃N₃-NHCH₂CH₂CH₂O), 3.19 (br m, 8192H, BocNHCH₂), 2.17 (not found, 1H, HC=CCH₂), 1.80-1.71 (m, 32760H, OCH₂CH₂CH₂O), 1.41 (s, 36864H,

 $C(CH_3)_3);$ ¹³C NMR (75 MHz, CDCl₃) δ 166.0 (C_3N_3), 156.1 (CO), 81.1 (not found, HC= CCH_2), 78.9 ($C(CH_3)_3$), 70.6 (OCH_2CH_2O), 70.5 (not found, HC= CCH_2), 70.3 (two lines, OCH_2CH_2O), 70.2 (OCH_2CH_2O), 69.6 ($CH_2CH_2CH_2O$), 69.3 (two lines, $CH_2CH_2CH_2O$), 38.5 ($CH_2CH_2CH_2O$), 38.1 ($CH_2CH_2CH_2O$), 30.1 (not found, HC= CCH_2), 29.7 (NHCH₂CH₂CH₂O), 28.5 ($CCCH_3$); MS (ESI-TOF) calcd for $C_{114668}H_{217048}N_{28666}O_{32762}$ 2520199.9, not found.

Compound 10 (deprotected G11). A solution of 9 (0.33 g, 0.131 μmol) in concentrated HCl (4 mL) and methanol (8 mL) was stirred for 16 h at room temperature and then evaporated under vacuum. The residue was dissolved in chloroform, washed with 5 M NaOH (aq), dried over MgSO₄, filtered, and evaporated under vacuum to give 10 (0.25 g, 90%) as a white wax. 1 H NMR (300 MHz, CDCl₃) δ 4.12 (not found, 2H, HC≡CCH₂), 3.60-3.38 (m, 122848H, CH₂OCH₂CH₂OCH₂CH₂OCH₂, C₃N₃-NHCH₂CH₂CH₂O), 2.76 (t, J = 6.6, 8192H, OCH₂CH₂CH₂NH₂), 2.22 (not found, 1H, HC≡CCH₂), 1.78-1.67 (m, 32760H, OCH₂CH₂CH₂); 13 C NMR (75 MHz, CDCl₃) δ 166.1 (C₃N₃), 81.1 (not found, HC≡CCH₂), 70.7 (OCH₂CH₂O), 70.5 (not found, HC≡CCH₂), 70.3 (two lines, OCH₂CH₂O), 69.5 (CH₂CH₂OH₂O), 69.3 (CH₂CH₂CH₂O), 39.7 (CH₂CH₂CH₂O), 38.1 (CH₂CH₂CH₂O), 33.5 (OCH₂CH₂CH₂NH₂), 30.1 (not found, HC≡CCH₂), 29.8 (NHCH₂CH₂CH₂O); MS (ESI-TOF) calcd for C₉₄₁₈₈H₁₈₄₂₈₀N₂₈₆₆₆O₂₄₅₇₀ 2110385.2, not found.

Compound 11 (Boc-protected G13). A solution of 14 (0.70 g, 0.353 mmol), 10 (0.070 g, 0.033 μmol), and DIPEA (0.20 mL, 1.14 mmol) in THF (4 mL), methanol (0.4 mL), and H₂O (0.4 mL) was stirred at 75 °C in a capped vessel for 4 d. The solution was evaporated under vacuum. The residue was dissolved in dichloromethane, washed with brine, dried over MgSO₄, filtered, and evaporated under vacuum. The crude product was dissolved in dichloromethane and precipitated by adding diethyl ether. The precipitation step was repeated until macromonomer 14 was completely removed, which was monitored by thin layer chromatography (DCM:MeOH = 14:1). The pure product 11 (0.275 g, 82%) was obtained as a white wax. ¹H NMR (300 MHz, CDCl₃) δ 4.05 (not found, 2H, HC=CCH₂), 3.59-3.39 (m, 491488H, CH₂OCH₂CH₂OCH₂CH₂OCH₂, C₃N₃-NHCH₂CH₂CH₂CH₂O), 3.17 (br, 32768H, BocNHCH₂), 2.17 (not found, 1H, HC=CCH₂), 1.78 (br, 131064H, OCH₂CH₂CH₂), 1.39 (s, 147456H, C(CH₃)₃); ¹³C NMR (75 MHz, CDCl₃) δ 165.4 (br, C₃N₃), 156.1 (CO), 81.1 (not found, HC=CCH₂), 78.9 (C(CH₃)₃), 70.6 (OCH₂CH₂O), 70.5 (not found, HC=CCH₂), 70.3 (OCH₂CH₂O), 70.2 (OCH₂CH₂O), 69.6 (CH₂CH₂CH₂O), 69.2 (CH₂CH₂O), 38.5 (CH₂CH₂CH₂O), 38.2 (CH₂CH₂CH₂O), 30.1 (not found, HC=CCH₂), 29.6 (NHCH₂CH₂CH₂O), 28.5 (C(CH₃)₃); MS (ESI-TOF) calcd for C₄₅₈₇₃₂H₈₆₈₃₁₂N₁₁₄₆₈₂O₁₃₁₀₆₆ 10082180.6, not found.

 2.22 (not found, 1H, $HC=CCH_2$), 1.80-1.71 (br m, 131064H, $OCH_2CH_2CH_2$); ¹³C NMR (75 MHz, $CDCI_3$) δ 166.2 (C_3N_3), 81.1 (not found, $HC=CCH_2$), 70.8 (OCH_2CH_2O), 70.7 (OCH_2CH_2O), 70.5 (not found, $HC=CCH_2$), 70.4 (OCH_2CH_2O), 70.3 (OCH_2CH_2O), 69.6 ($CH_2CH_2CH_2O$), 69.4 (two lines, $CH_2CH_2CH_2O$), 39.7 ($CH_2CH_2CH_2O$), 38.2 ($CH_2CH_2CH_2O$), 33.5 ($OCH_2CH_2CH_2NH_2$), 30.1 (not found, $HC=CCH_2$), 29.8 ($NHCH_2CH_2CH_2O$); MS (ESI-TOF) calcd for $C_{376812}H_{737240}N_{114682}O_{98298}$ 8442921.6, not found.

Compound 13. Cyanuric chloride (1.70 g, 9.22 mmol) was added to an ice-bath cooled solution of *N*-Boc-4,7,10-trioxa-1,13-tridecanediamine (5.80 g, 18.10 mmol) and DIPEA (4.0 mL, 22.8 mmol) in THF (40 mL). The solution was stirred for 1 h at 0 °C, warmed to room temperature, and then stirred for 16 h. After addition of a solution of 4,7,10-trioxa-1,13-tridecanediamine (14.0 g, 63.5 mmol) and DIPEA (3.0 mL, 17.1 mmol) in methanol (10 mL), the reaction solution was stirred for an additional 16 h at 70 °C and evaporated under vacuum. The residue was dissolved in dichloromethane, washed with brine, dried over MgSO₄, filtered, and evaporated under vacuum. The crude product was purified by silica gel chromatography (DCM:MeOH = 10:1 with 1% NH₄OH) to give 13 (7.6 g, 88%) as a clear oil. ¹H NMR (300 MHz, CDCl₃) δ 3.67-3.34 (m, 42H, CH₂OCH₂CH₂OCH₂CH₂OCH₂CH₂OCH₂C, 3N₃-NHCH₂CH₂CH₂O), 3.17 (m, 4H, BocNHCH₂), 2.79 (t, J = 6.6, 2H, OCH₂CH₂CH₂NH₂), 1.83-1.67 (m, 12H, OCH₂CH₂CH₂O), 1.39 (s, 18H, C(CH₃)₃); ¹³C NMR (75 MHz, CDCl₃) δ 166.0 (C₃N₃), 156.2 (CO), 78.9 (C(CH₃)₃), 70.6 (OCH₂CH₂O), 70.3 (OCH₂CH₂O), 70.2 (two lines, OCH₂CH₂O), 69.6 (CH₂CH₂CH₂O), 69.4 (CH₂CH₂CH₂O), 39.6 (CH₂CH₂O), 38.5 (CH₂CH₂CH₂O), 38.2 (CH₂CH₂CH₂O), 32.6 (OCH₂CH₂CH₂NH₂), 29.7 (NHCH₂CH₂CH₂O), 28.5 (C(CH₃)₃); MS (ESI-TOF) calcd for C₄₃H₈₅N₉O₁₃ 935.6267, found 936.6595 (M + H)⁺.

Compound 14 (macromonomer). Cyanuric chloride (0.46 g, 2.49 mmol) was added to an ice-bath cooled solution of 13 (4.70 g, 5.02 mmol) and DIPEA (2.0 mL, 11.4 mmol) in THF (40 mL). The solution was stirred for 1 h at 0 °C, warmed to room temperature, and then stirred for an additional 16 h. After concentration under vacuum, the residue was dissolved in dichloromethane, washed with brine, dried over MgSO₄, filtered, and evaporated under vacuum. The crude product was purified by silica gel chromatography (DCM:MeOH = 14:1) to give 14 (4.68 g, 95%) as a clear oil. 1 H NMR (300 MHz, CDCl₃) δ 3.64-3.41 (m, 88H, CH₂OCH₂CH₂OCH₂CH₂OCH₂CH₂OCH₂C, C₃N₃-NHCH₂CH₂CH₂O), 3.20 (br m, 8H, BocNHCH₂), 1.83-1.70 (m, 24H, OCH₂CH₂CH₂), 1.42 (s, 36H, C(CH₃)₃); 13 C NMR (75 MHz, CDCl₃) δ 165.9 (C₃N₃), 165.6 (C₃N₃), 156.1 (CO), 78.8 (C(CH₃)₃), 70.5 (OCH₂CH₂O), 70.2 (two lines, OCH₂CH₂O), 69.5 (CH₂CH₂OH₂O), 69.3 (CH₂CH₂CH₂O), 38.4 (CH₂CH₂CH₂O), 38.1 (CH₂CH₂CH₂O), 29.6 (NHCH₂CH₂CH₂O), 28.4 (C(CH₃)₃); MS (ESI-TOF) calcd for C₈₉H₁₆₈CIN₂₁O₂₆ 1982.2158, found 1983.1606 (M + H)⁺.

Compound 15 (Boc-protected G1). A solution of propargylamine (0.10 g, 1.82 mmol), 14 (0.20 g, 0.10 mmol), and DIPEA (0.20 mL, 1.14 mmol) in THF (1 mL) and methanol (0.1 mL) was stirred at 70 °C in a capped vessel for 16 h. The solution was evaporated under vacuum. The residue was dissolved in dichloromethane, washed with brine (pH 5), dried over MgSO₄, filtered, and evaporated under vacuum. The crude product was purified by silica

gel chromatography (DCM:MeOH = 10:1) to give **15** (0.18 g, 90%) as a clear oil. ¹H NMR (300 MHz, CDCl₃) δ 4.07 (br, 2H, HC \equiv CCH₂), 3.61-3.34 (m, 88H, CH₂OCH₂CH₂OCH₂CH₂OCH₂, C₃N₃-NHCH₂CH₂CH₂O), 3.12 (br m, 8H, BocNHCH₂), 2.17 (br, 1H, HC \equiv CCH₂), 1.74-1.62 (m, 24H, OCH₂CH₂CH₂), 1.34 (s, 36H, C(CH₃)₃); ¹³C NMR (75 MHz, CDCl₃) δ 165.4 (br, **C**₃N₃), 156.0 (**C**O), 81.1 (HC \equiv CCH₂), 78.7 (**C**(CH₃)₃), 70.7 (H**C** \equiv CCH₂), 70.5 (O**C**H₂CH₂O), 70.2 (O**C**H₂CH₂O), 70.1 (O**C**H₂CH₂O) 69.4 (CH₂CH₂CH₂O), 69.1 (CH₂CH₂CH₂O), 38.4 (**C**H₂CH₂CH₂O), 38.0 (**C**H₂CH₂CH₂O), 30.1 (HC \equiv CCH₂), 29.5 (NHCH₂CH₂CH₂O), 28.4 (C(**C**H₃)₃); MS (ESI-TOF) calcd for C₉₂H₁₇₂N₂₂O₂₆ 2001.2813, found 2002.2499 (M + H)⁺.

Compound 16 (deprotected G1). A solution of 15 (0.15 g, 0.075 mmol) in concentrated HCl (0.5 mL) and methanol (1.0 mL) was stirred for 16 h at room temperature and then evaporated under vacuum. The residue was dissolved in chloroform, washed with 5 M NaOH (aq), dried over MgSO₄, filtered, and evaporated under vacuum to give 16 (0.12 g, quantitative) as a clear oil. H NMR (300 MHz, CDCl₃) δ 4.09 (br, 2H, HC≡CCH₂), 3.57-3.35 (m, 88H, CH₂OCH₂CH₂OCH₂CH₂OCH₂, C₃N₃-NHCH₂CH₂CH₂O), 2.76 (br, 8H, OCH₂CH₂CH₂NH₂), 2.18 (br, 1H, HC≡CCH₂), 1.77-1.64 (m, 24H, OCH₂CH₂CH₂); 13 C NMR (75 MHz, CDCl₃) δ 166.0 (C₃N₃), 81.1 (HC≡CCH₂), 70.5 (HC≡CCH₂, OCH₂CH₂O), 70.2 (OCH₂CH₂O), 70.1 (two lines, OCH₂CH₂O) 69.4 (CH₂CH₂CH₂O), 69.2 (two lines, CH₂CH₂O), 39.4 (CH₂CH₂CH₂O), 38.0 (CH₂CH₂CH₂O), 32.4 (OCH₂CH₂CH₂NH₂), 30.1 (HC≡CCH₂), 29.6 (NHCH₂CH₂CH₂O); MS (ESI-TOF) calcd for C₇₂H₁₄₀N₂₂O₁₈ 1601.0716, found 1602.0498 (M + H)⁺.

Spectra of Compounds

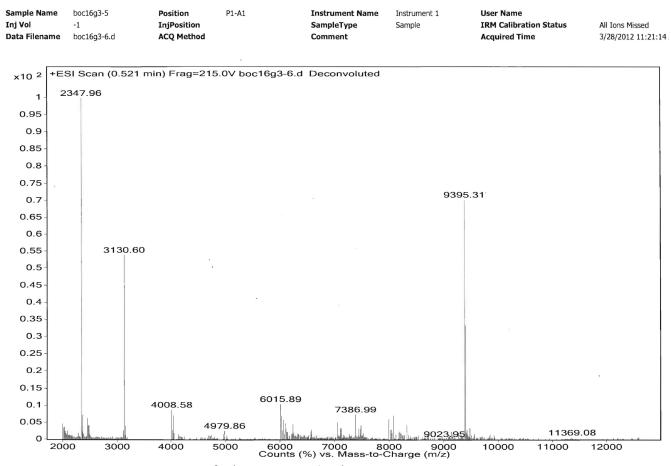


Figure S1. ESI-TOF mass spectrum of 1 (Boc-protected G3).

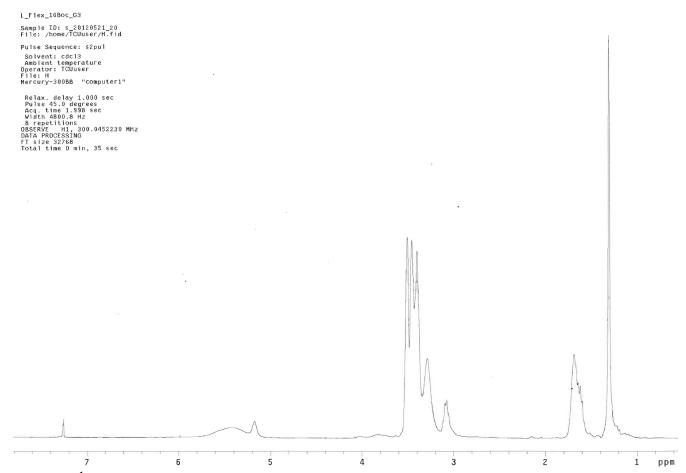


Figure S2. ¹H NMR spectrum of 1 (Boc-protected G3, 300 MHz, CDCl₃).

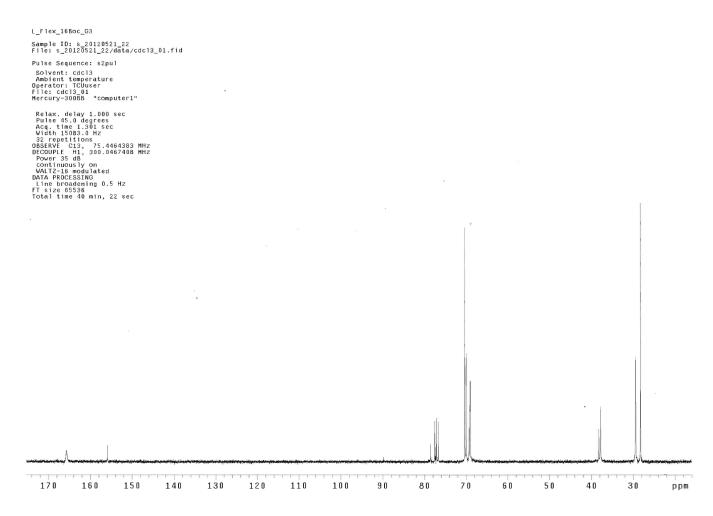


Figure S3. ¹³C NMR spectrum of 1 (Boc-protected G3, 75 MHz, CDCl₃).

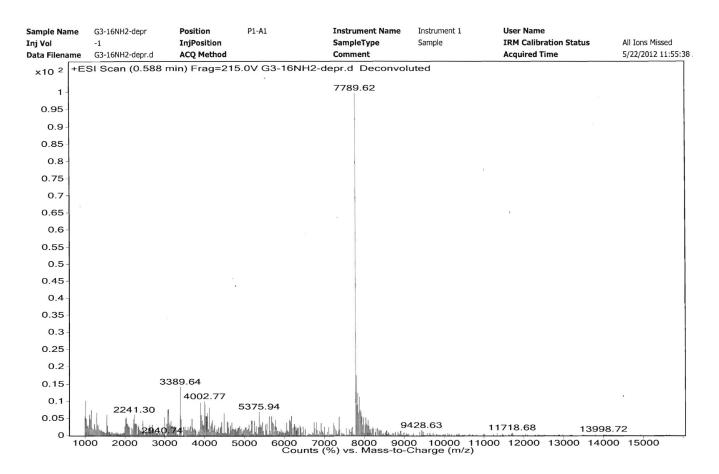


Figure S4. ESI-TOF mass spectrum of 2 (deprotected G3).

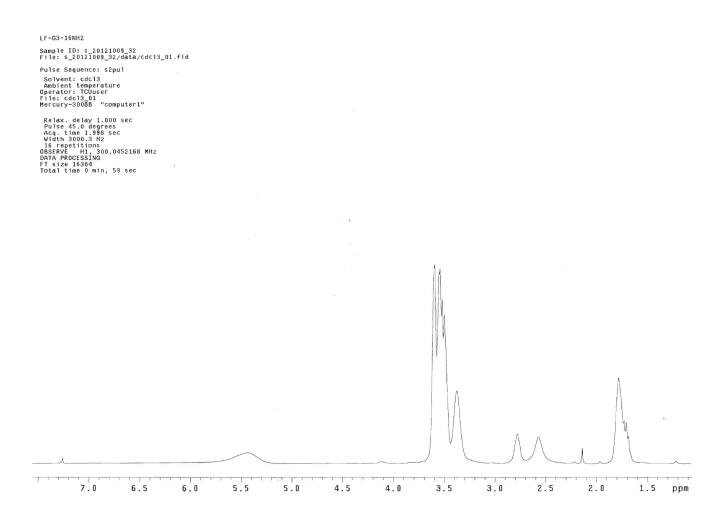


Figure S5. ¹H NMR spectrum of **2** (deprotected G3, 300 MHz, CDCl₃).

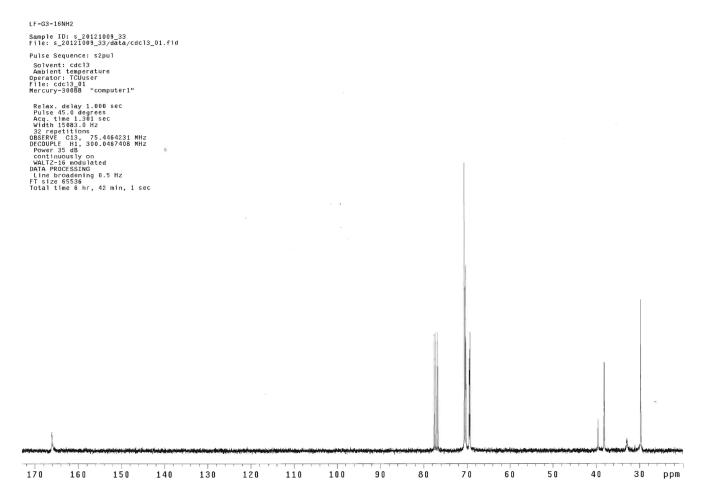


Figure S6. 13 C NMR spectrum of 2 (deprotected G3, 75 MHz, CDCl $_3$).

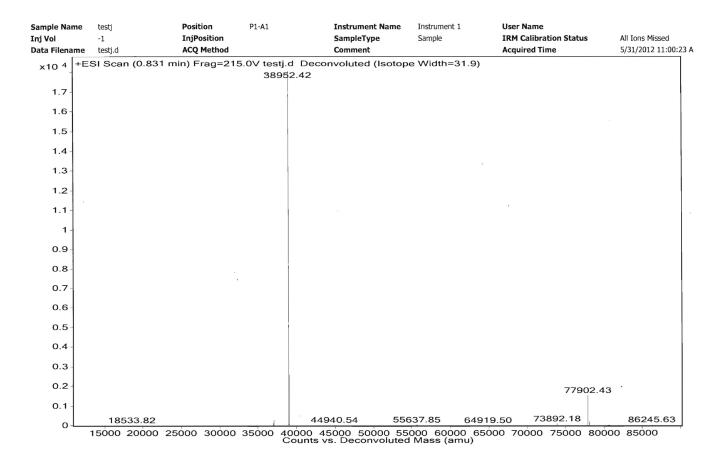


Figure S7. ESI-TOF mass spectrum of **3** (Boc-protected G5). The peak at 77902 is attributed to noncovalent dimer seen in the deconvolution, and not a synthetic impurity.

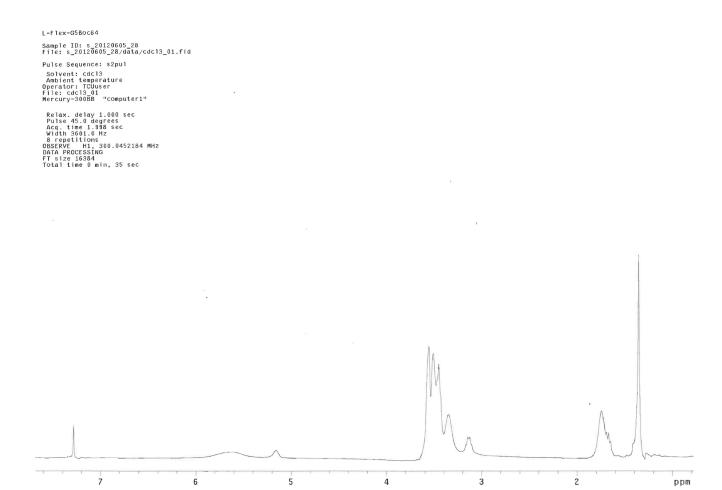


Figure S8. ¹H NMR spectrum of 3 (Boc-protected G5, 300 MHz, CDCl₃).

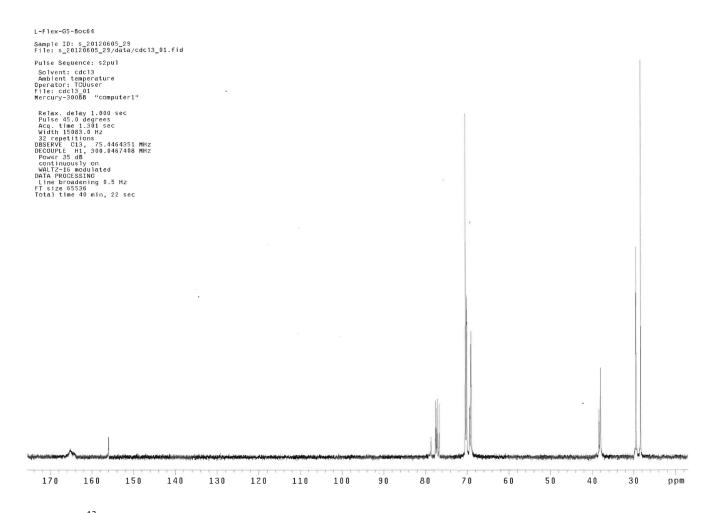


Figure S9. ¹³C NMR spectrum of **3** (Boc-protected G5, 75 MHz, CDCl₃).

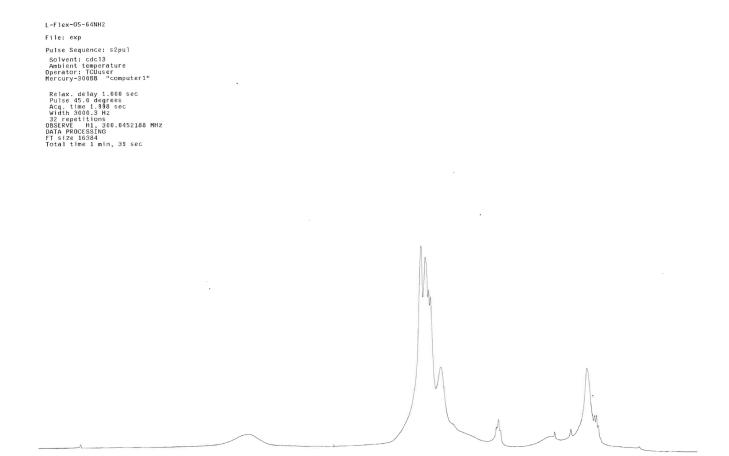


Figure S10. ¹H NMR spectrum of 4 (deprotected G5, 300 MHz, CDCl₃).

 ppm

L-Flex-G5-64NH2

Sample ID: s_20120612_02
File: s_20120612_02/data/cdc13_01.fid

Pulse Sequence: s2pul
Solvent: cdc13
Ambient temperature
Operator: TCUUser
File: cdc13_01
Mercury-300BB "computer1"

Relax. delay 1.000 sec
Pulse 45.0 degrees
Acq. time 1.301 sec
Width 15083.0 Hz
64 repetitions
OBSERVE C13, 75.4464304 MHz
DECOUPLE H1, 300.0467408 MHz
Power 35 dB
continuously on
WALTZ-16 modulated
DATA PROCESSING
Line broadening 0.5 Hz
FT size 65536
Total time 40 min, 13 sec

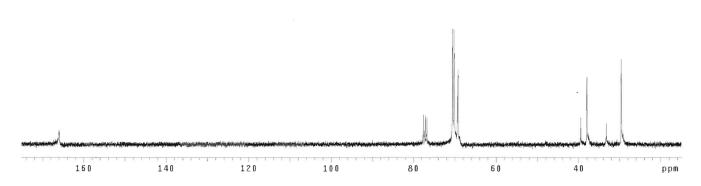


Figure S11. ¹³C NMR spectrum of 4 (deprotected G5, 75 MHz, CDCl₃).

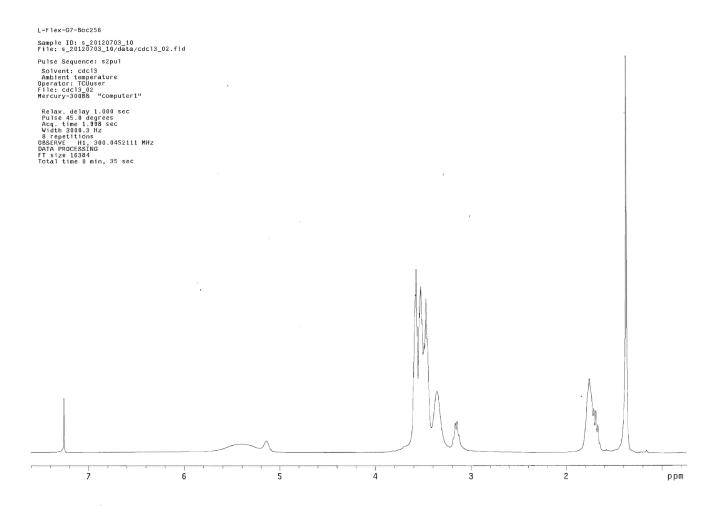


Figure S12. ¹H NMR spectrum of 5 (Boc-protected G7, 300 MHz, CDCl₃).

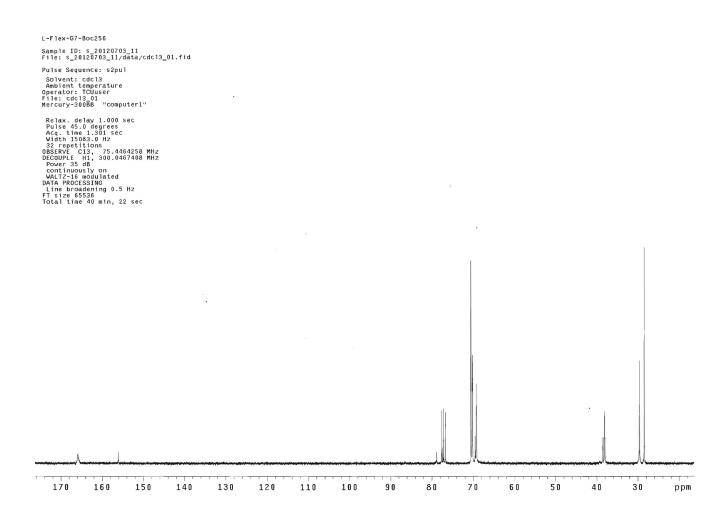


Figure S13. ¹³C NMR spectrum of 5 (Boc-protected G7, 75 MHz, CDCl₃).

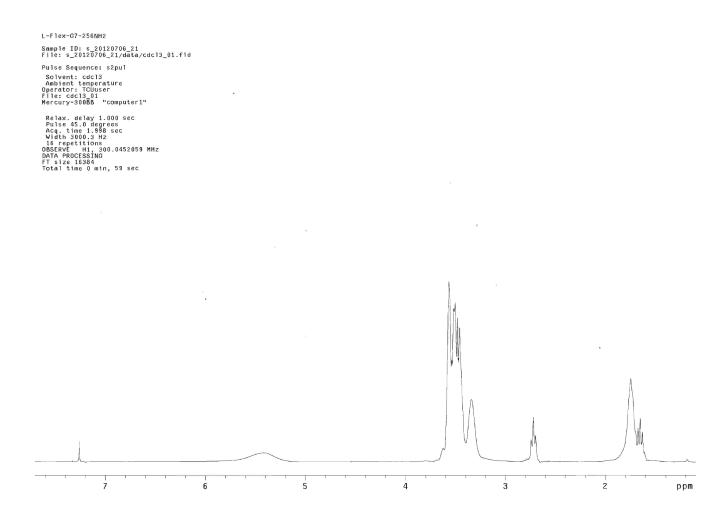


Figure S14. ¹H NMR spectrum of 6 (deprotected G7, 300 MHz, CDCl₃).

L-Flex_07-250NH2

Sample 10: s_20120706_24
File: s_20120706_24/data/cdc13_01.fid
Pulse Sequence: s2pul
Solvent: cdc13
Abblent temperature
Operator: TCUuser
File: cdc13_01
Mercury-30086 "computer1"

Relax. delay 1.000 sec
Pulse 45.0 degrees
Acq. time 1.301 sec
Vidth 15083.0 Hz
32 repetitions
OBSERVE C13, 75.4464254 MHz
DECOUPLE H1, 300.0467408 MHz
Power 30 Mb on
VALTZ-1s nodulated
DATA PROCESSING
Line broadening 0.5 Hz
FT size 85536
Total time 40 min, 22 sec

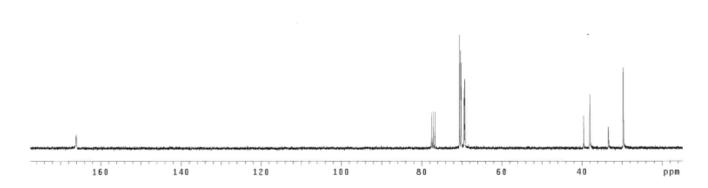


Figure S15. ¹³C NMR spectrum of 6 (deprotected G7, 75 MHz, CDCl₃).

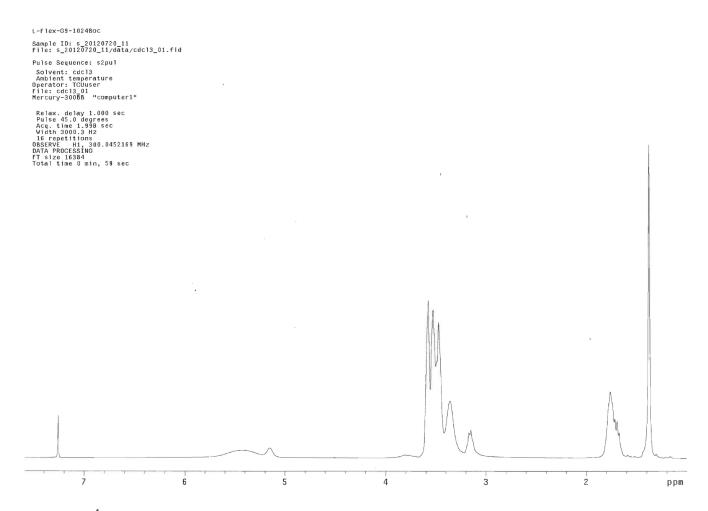


Figure S16. ¹H NMR spectrum of **7** (Boc-protected G9, 300 MHz, CDCl₃).

L-Flex-G9-1024Boc

Sample ID: s_20120720_12
file: s_20120720_12/data/cdc13_01.fid
Pulse Sequence: s2pul

Solvent: cdc13
Amblent temperature
floreated 13_01
Mercury-300BB "computer1"

Relax. delay 1.000 sec
Pulse 45.0 degrees
Acquetime 1.301 sec
Width 1598.0 htz
32 repetitions
0BSERWE C13, 75.4464263 MHz
DECQUPLE H1, 300.0467408 MHz
POWER 35 dB
continuously on
WALTZ-16 modulated
DATA PROCESSING
Line broadening 0.5 Hz
FT size 65536

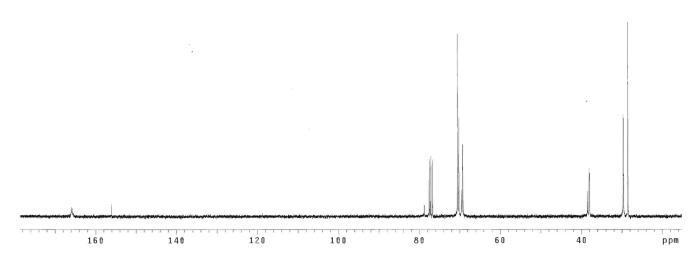


Figure S17. ¹³C NMR spectrum of **7** (Boc-protected G9, 75 MHz, CDCl₃).

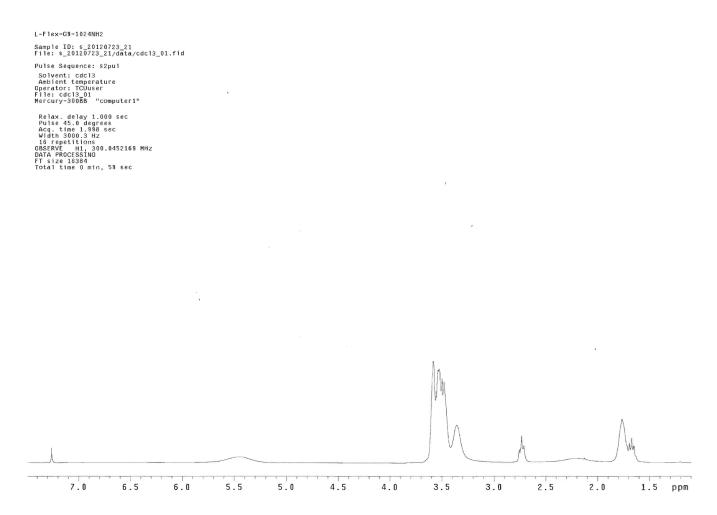
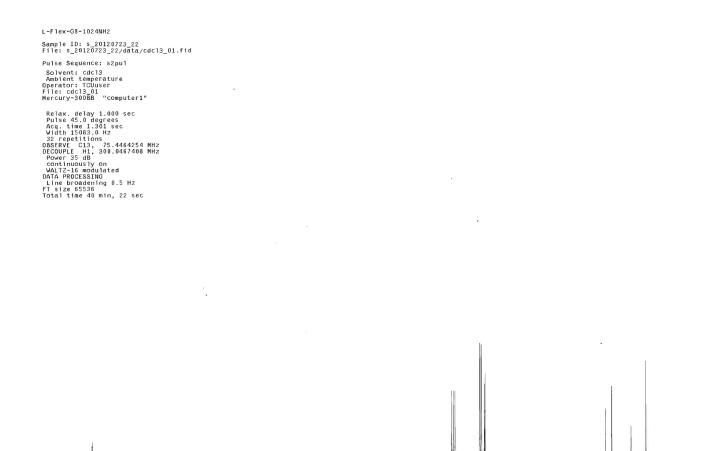


Figure S18. ¹H NMR spectrum of 8 (deprotected G9, 300 MHz, CDCl₃).



ppm

Figure S19. ¹³C NMR spectrum of 8 (deprotected G9, 75 MHz, CDCl₃).

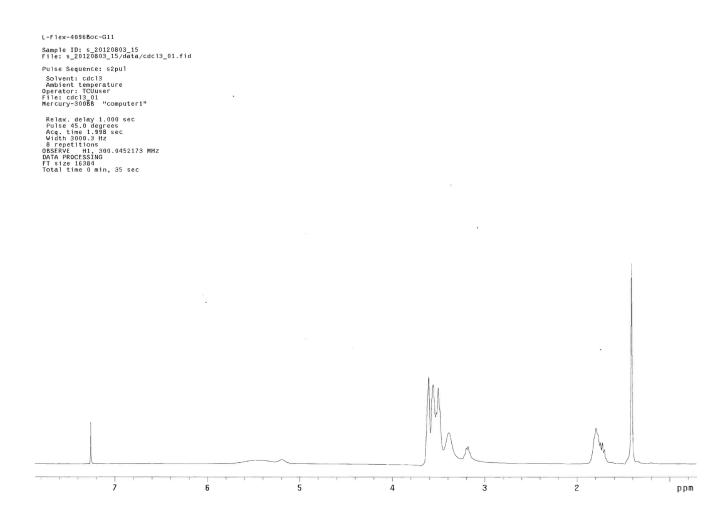


Figure S20. ¹H NMR spectrum of 9 (Boc-protected G11, 300 MHz, CDCl₃).

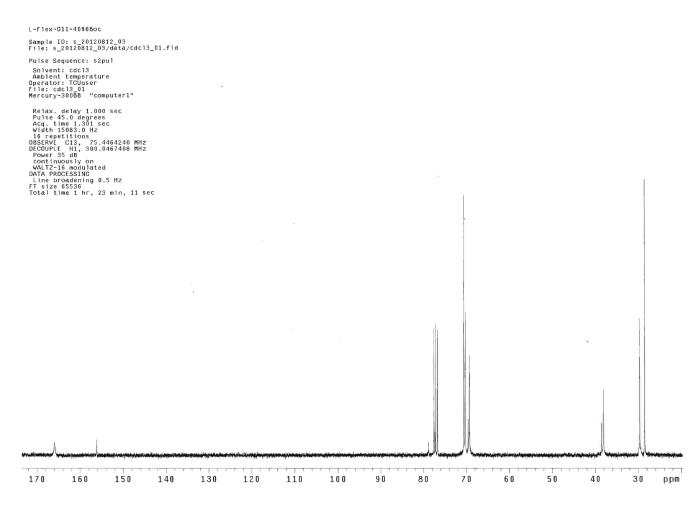


Figure S21. ¹³C NMR spectrum of 9 (Boc-protected G11, 75 MHz, CDCl₃).

L-Fix-011-4098MN2
pad-01 run vit findz0 before acquisition

Sample ID: =_01120818_31
File: =_02120818_31/ada_ccis_0.0.fid

Pulse Saquence: \$2pul

Solvent: cdol: 30

Solvent: cdol: 30

Hercury-3080 "Computer1"

Relax. delay 1.000 sec
Acq. time 1.989 sec

Viths Solo. 31:2

OSSERVE H1, 300.0452173 MH2

ORIGINAL H2, 300.0452173 MH2

Total line 0 min, 59 sec

Figure S22. ¹H NMR spectrum of **10** (deprotected G11, 300 MHz, CDCl₃).

ppm

L-Flex-G11-4096NH2

Sample ID: s 20120814_27
file: /home/TCUuser/L_Flex_G11_4096NH2_C.fid

Pulse Sequence: s2pul

Solvent: cdc13
 Ambient temperature
Operator: TCUuser
File: L_Flex_G11_4096NH2_C
Mercury-300BB "Computer!"

Relax. delay 1.000 sec
Pulse 45.0 degrees
Acq. time 1.301 sec
Width 15083.0 H2
16 repetitions
OBSERVE C13, 75.4467425 MHz
DECOUPLE H1, 300.0467408 MHz
Power 35 dB
continuously on
WALTZ-16 modulated
DATA PROCESSING
Line broadening 0.5 Hz
Fi size 65536
Total time 41 min, 42 sec

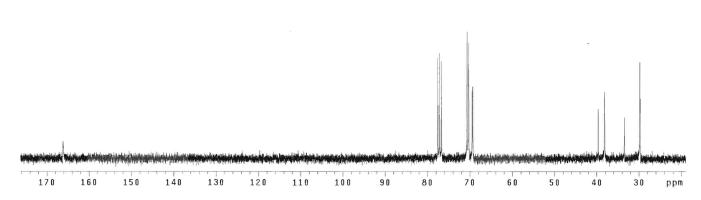


Figure S23. ¹³C NMR spectrum of **10** (deprotected G11, 75 MHz, CDCl₃).

L-Tier-013-16364BC

Sample ID: g_2012082_12

Firs: s_201252_12/arta/cdc13_01.fld

Pulse Sequence: 52µ1

Solvent: cdc13 ere

Operator: Coluser

First case Sequence: 62 pul

Servent: cdc3 ere

Operator: Coluser

First case Sequence: 62 pul

March 1:00 sec

Palse. delay 1:00 sec

Aca. time 1:980 sec

Id dich 2003.02

OSSEWE 11: 300.0152136 PHZ

OSSEWE 11: 300.0152136

Figure S24. ¹H NMR spectrum of 11 (Boc-protected G13, 300 MHz, CDCl₃).

ppm

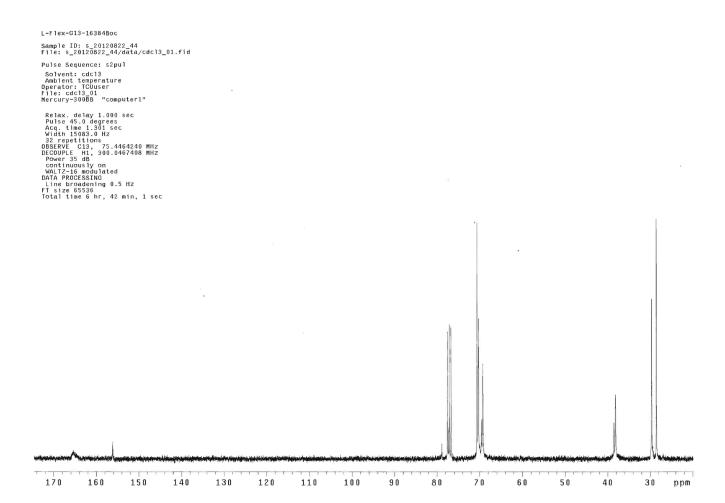


Figure S25. ¹³C NMR spectrum of 11 (Boc-protected G13, 75 MHz, CDCl₃).

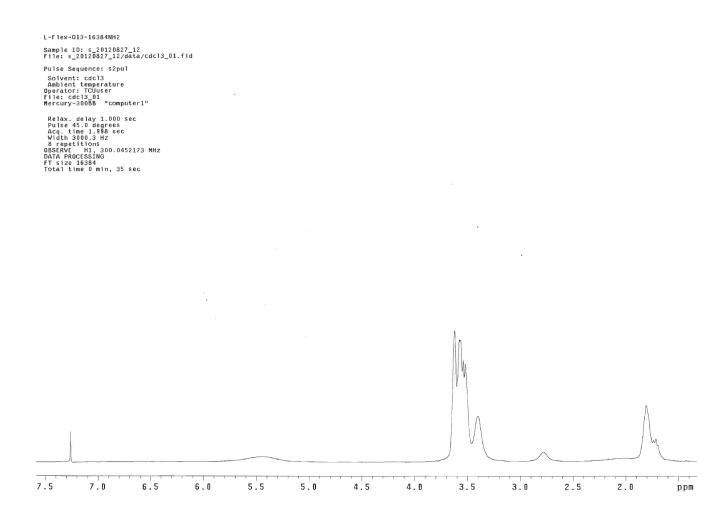


Figure S26. ¹H NMR spectrum of 12 (deprotected G13, 300 MHz, CDCl₃).

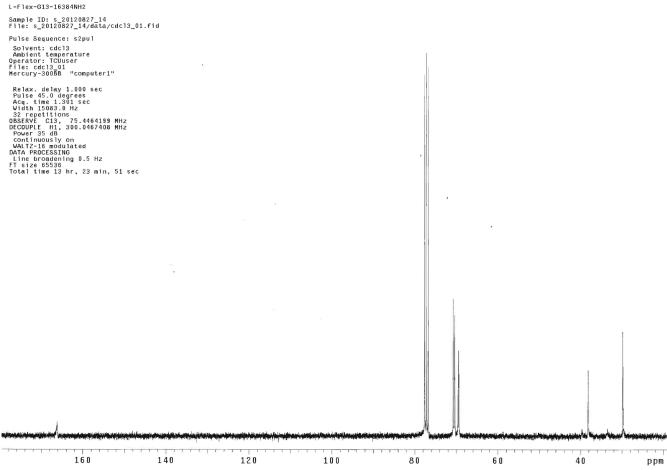


Figure S27. ¹³C NMR spectrum of 12 (deprotected G13, 75 MHz, CDCl₃).

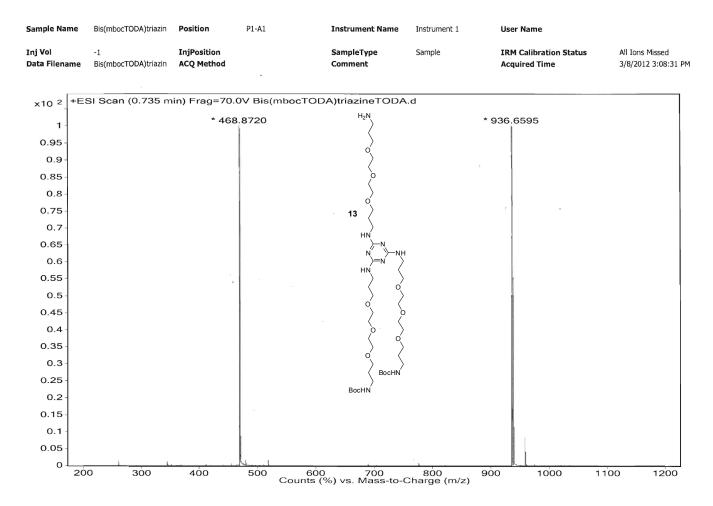


Figure S28. ESI-TOF mass spectrum of 13.

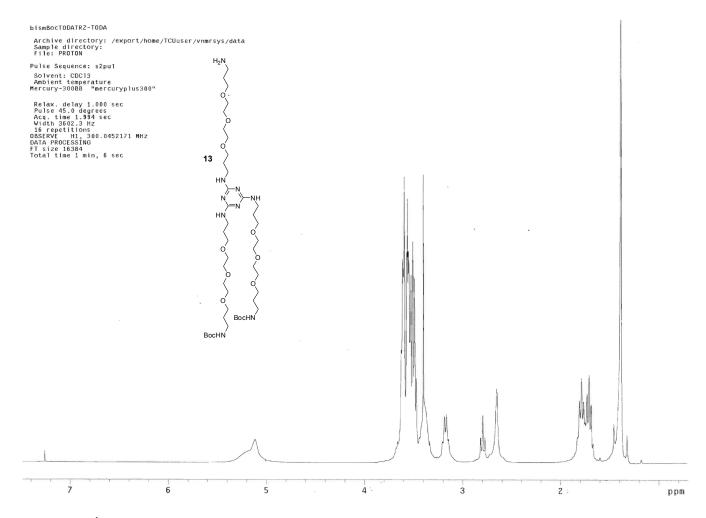


Figure S29. ¹H NMR spectrum of 13 (300 MHz, CDCl₃).

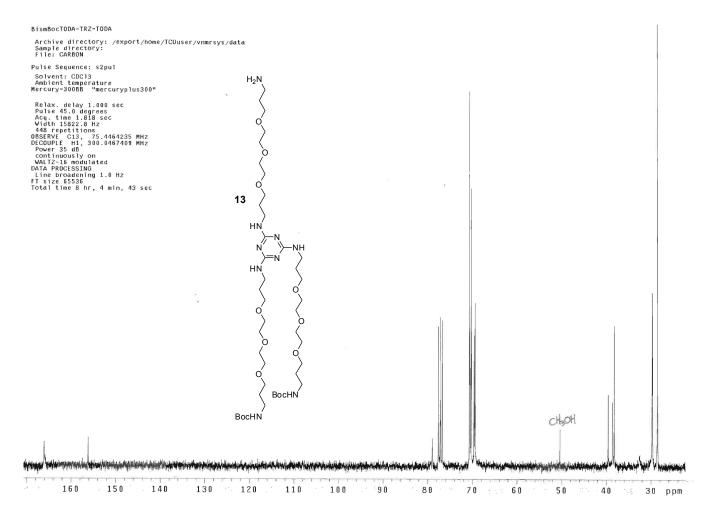


Figure S30. ¹³C NMR spectrum of 13 (75 MHz, CDCl₃).

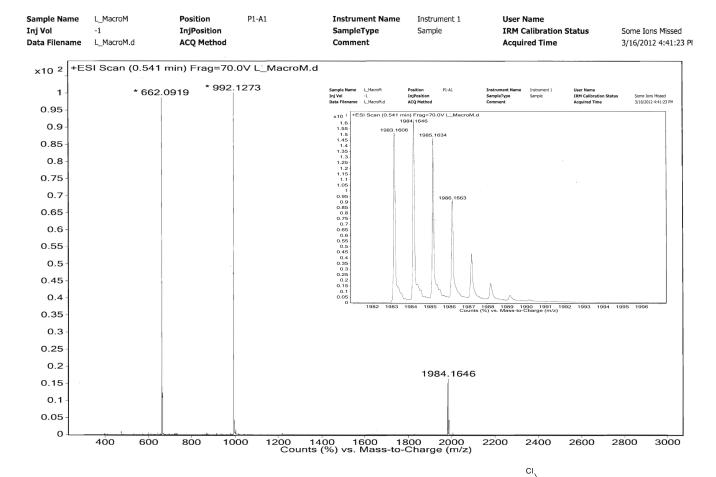


Figure S31. ESI-TOF mass spectrum of 14 (macromonomer).

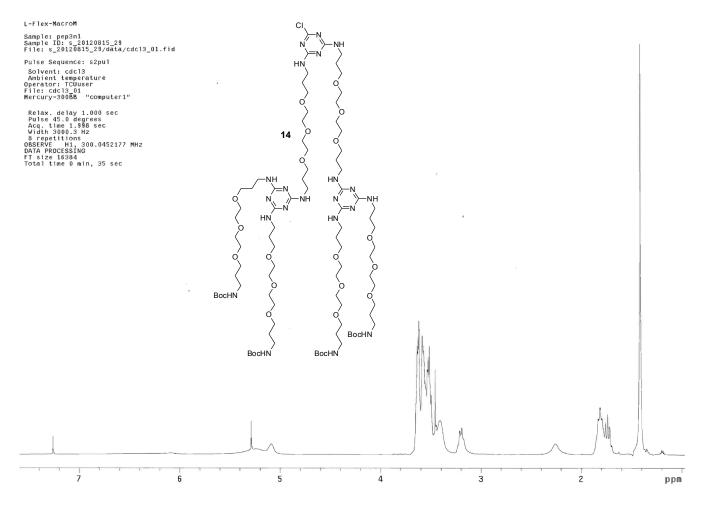


Figure S32. ¹H NMR spectrum of 14 (macromonomer, 300 MHz, CDCl₃).

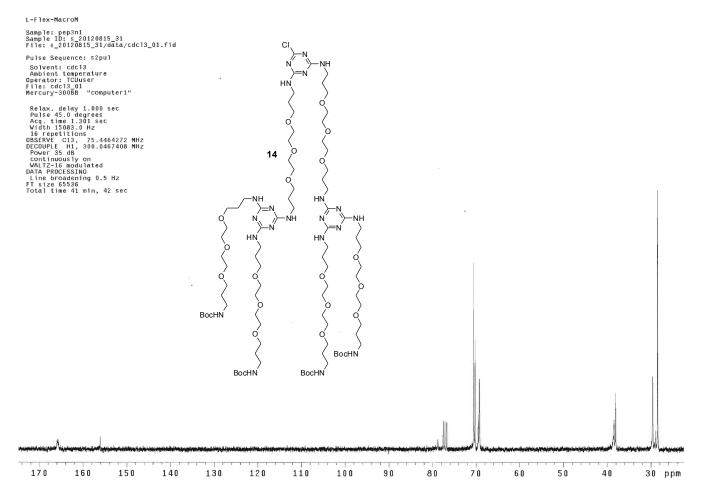


Figure S33. ¹³C NMR spectrum of 14 (macromonomer, 75 MHz, CDCl₃).

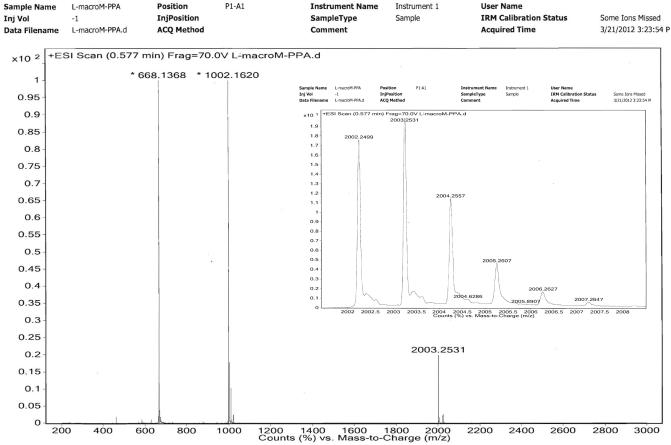


Figure S34. ESI-TOF mass spectrum of 15 (Boc-protected G1).

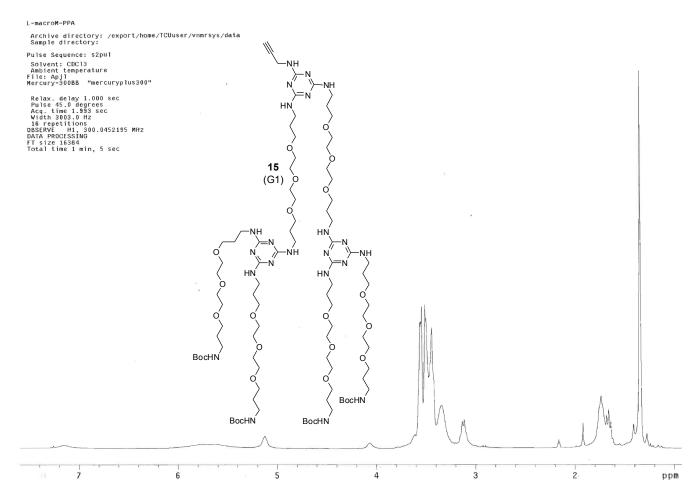


Figure S35. ¹H NMR spectrum of 15 (Boc-protected G1, 300 MHz, CDCl₃).

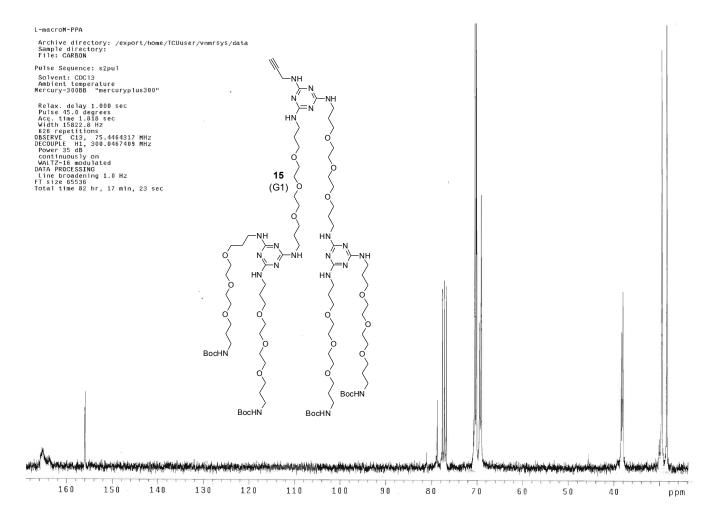


Figure S36. 13 C NMR spectrum of 15 (Boc-protected G1, 75 MHz, CDCl $_3$).

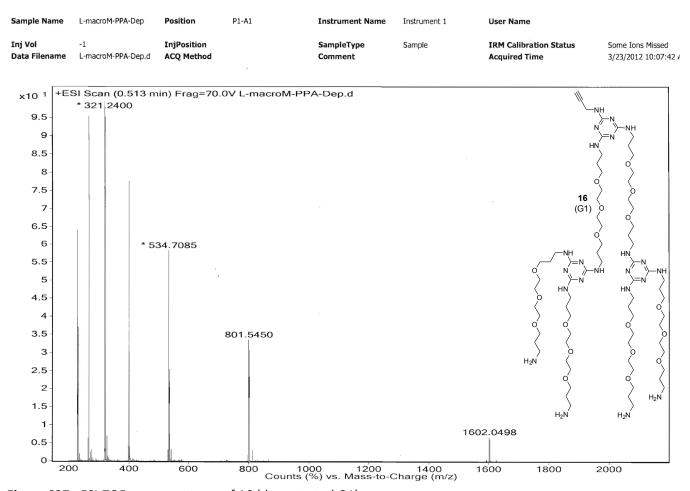


Figure S37. ESI-TOF mass spectrum of 16 (deprotected G1).

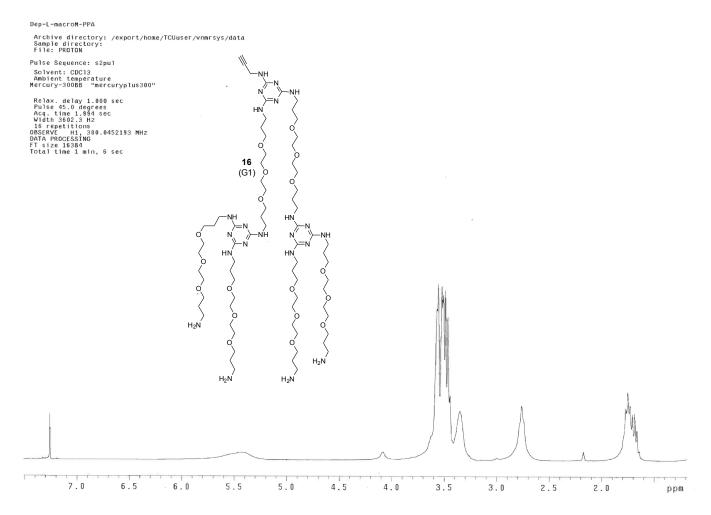


Figure S38. ¹H NMR spectrum of 16 (deprotected G1, 300 MHz, CDCl₃).

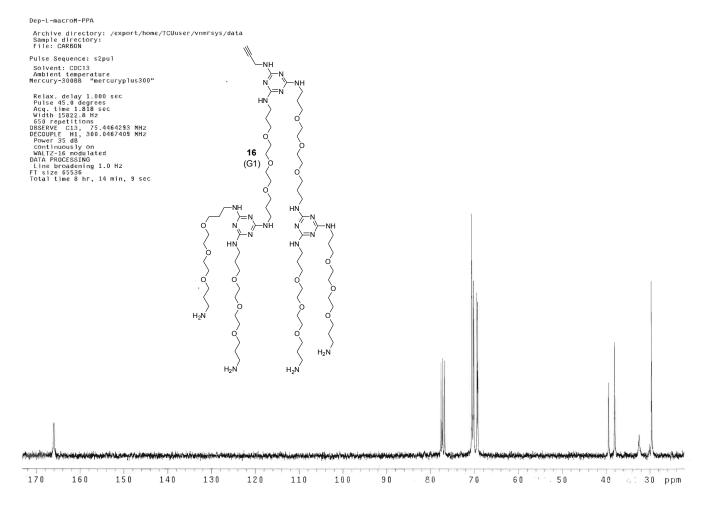


Figure S39. ¹³C NMR spectrum of 16 (deprotected G1, 75 MHz, CDCl₃).

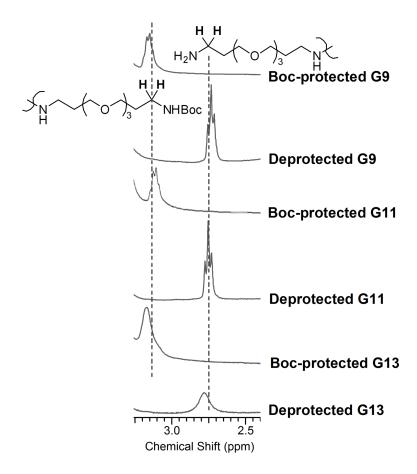


Figure S40. ¹H NMR Spectra of the large generation dendrimers (G9-G13) display the finger print region for monitoring the reiterative addition and deprotection: The vicinal proton signals of NHBoc groups appear around at 3.2 ppm, while the vicinal proton signals of NH₂ groups appear around at 2.75 ppm.

HPLC Analysis

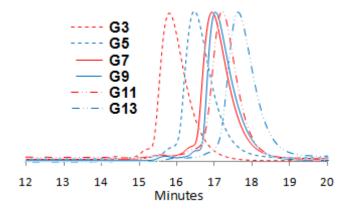


Figure S41. HPLC traces of the dendrimers (G3-G13). For analytic HPLC of the dendrimers, a ZORBAX 300SB-C8 column (1.0 x 150 mm, 3.5 μ m) was used with a gradient elution: 70% A to 30% A over 20 min and then keep 30% A (A = water with 0.1% TFA, B = acetonitrile with 0.1% TFA) with a flow rate of 80 μ L/min. UV detection was performed at 230 nm.

Transmission Electron Microscopy

Experimental. Transmission electron microscopy micrographs (TEM) were recorded on a Tecnai 12 Bio Twin instrument operating at a 120 kV accelerating voltage . Samples (dendrimer: 1 mg mL $^{-1}$ in Milli-Q water, CCMV: 50 mg L $^{-1}$ in 10 mM NaAc, 1 mM EDTA, 1 mM NaN $_3$, pH 5) were prepared on formvar carbon-coated copper grids by placing a 3 μL drop of the sample solution on the grid. The sample drop was left standing for 1 min after which time the excess solution was blotted away with filter paper. Samples were negatively stained by applying 3 μL of stain (0.5% uranyl acetate in Milli-Q water) onto the grid and removing the excess stain with filter paper after 15 s. The samples were dried under air flow for at least 5 min before imaging.

Cryo-TEM samples were prepared from the same aqueous (dendrimer) or buffer (virus) solutions. Prior to sample deposition the TEM grids (Quantifoil R 3.5/1, holey carbon film, Cu 200 mesh) were treated with Gatan Solarus 950 plasma system. 3 μ L of sample was placed on the grid, which was consecutively blotted for 1-2 s (100 % relative humidity, -2 mm blot offset), with Fei Vitrobot Mk3 followed by immediate vitrification with a mixture of liquid ethane and propane (~1:1) at -180 °C. Vitrified samples were cryo-transferred to the microscope. Images were obtained with a JEOL JEM-3200 FSC field emission cryo electron microscope operating at a 300 kV accelerating voltage and specimen temperature of 86 K.

Dynamic Light Scattering

Measurements of dynamic light scattering (DLS) were performed at 25.0 \pm 0.1 °C. All dendrimer samples were filtered through a 0.2 mm filter (Anotop 10, Whatman) and placed in a test tube. The experiments were performed on a light scattering apparatus built using the following main components: He-Ne laser (35 mW, 632.8 nm, Coherent Radiation), manual goniometer and thermostat (Photocor Instruments), multi-tau correlator, APD detector and software (PD4042, Precision Detectors). All measurements were performed at a scattering angle of 90°. The dynamic-light-scattering correlation functions were analyzed using a regularization algorithm (Precision Deconvolve 32, Precision Detectors). Light-scattering distributions were bimodal. The calculated z-average diffusion coefficient, D, corresponds to the peak at low apparent radii (fast diffusion mode). For dilute solutions, diffusion coefficient values can be converted into the corresponding hydrodynamic radius, $R_{\rm h}$, using the Stokes-Einstein equation: $R_{\rm h} = k_{\rm B}T/(6\pi\eta D)$ for a sphere, where $k_{\rm B}$ is the Boltzmann constant, $T=298.2\,{\rm K}$ the absolute temperature, and η the corresponding viscosity of water. The viscosity value of water, $\eta=0.890\times10^{-3}\,{\rm kg}\,{\rm m}^{-1}{\rm s}^{-1}$, was used to calculate $R_{\rm h}$.

For **G9** and **G11**, DLS measurements were performed in phosphate buffered saline (PBS) solutions. For **G13**, filtering of phosphate buffered saline solutions virtually removed all dendrimer material. Thus, DLS measurements were performed in NaCl 0.01 M and water. Our results are summarized in **Table S1**. Representative light-scattering distributions for **G9**,**G11** and **G13** are shown in **Figure S42**.

Table S1. Diffusion coefficients and calculated hydrodynamic radii. ^a concentration values do not take into account dendrimer material removed by filtering. ^b errors are standard deviations. ^c pH≈7.4

Sample description ^a	$D/10^{-9} m^2 s^{-1}$	R _h /nm
G3 3.9 g/L in PBS	0.124±0.01 b	-
G3 7.0 g/L in PBS	0.117±0.01	-
G3 9.0 g/L in PBS	0.112±0.01	-
G3 0.0 g/L in PBS °	0.132±0.01	1.85±0.02
G5 4.9 g/L in PBS	0.0614±0.006	3.99±0.04
G7 1.0 g/L in PBS	0.0358±0.002	6.85±0.03
G9 1.0 g/L in PBS	0.0229±0.003	10.7±0.2
G11 0.48 g/L in PBS	0.0168±0.002	14.6±0.2
G13 0.25 g/L in NaCl 0.01 M d	0.0162±0.005	15.1±0.4
G13 0.11 g/L in water d	0.0187±0.003	-
G13 0.40 g/L in water d	0.0207±0.006	-
G13 2.0 g/L in water d	0.0317±0.033	-

Comments. Diffusion coefficients of **G13** in water were found to be higher than that in NaCl 0.01 M. This can be explained by considering dendrimer charge. In the absence of electrostatic screening, dendrimer diffusion coefficients are expected to be higher than their intrinsic Brownian mobilities. Thus, the Stokes-Einstein equation was not applied in this case.

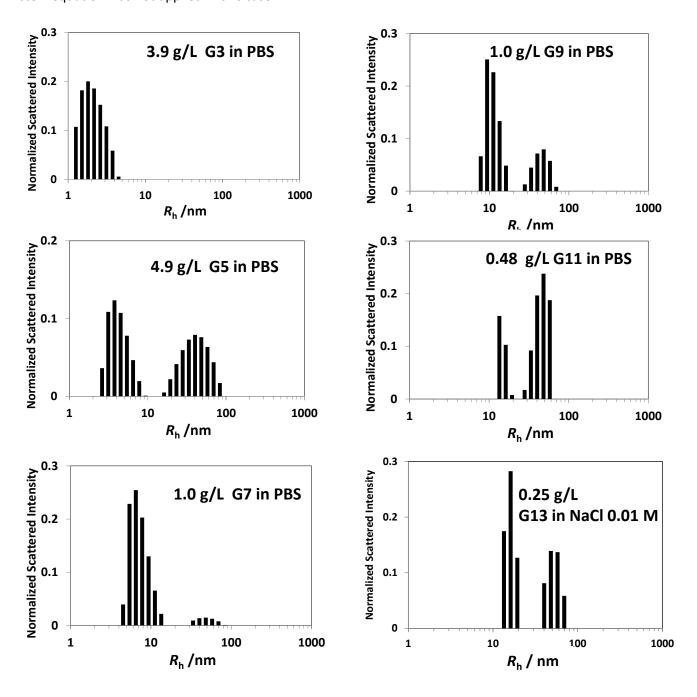


Figure S42. Normalized scattered-intensity distributions.

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Sample Preparation. For AFM analysis on air, G13 dendrimers (2 mg/ml water solution filtered through the 0.2 μm filter (Whatman GD/X))) were dropped (5 μl) on freshly cleaved circular mica disc (diameter 1 cm) and solution was let to evaporate at ambient conditions (30 min.). Samples for liquid AFM were prepared by drop casting of G13 solution (15 μl) of the same composition on mica disc fixed in liquid measurement cell. After 10 minutes of incubation (without drying) the surface of mica was gently layered with 1ml of ultrapure (Milli-Q) water and AFM analysis performed.

AFM analysis. Atomic force microscopy analysis was performed with an AFM Integra Probe Nanolaboratory (NT-MDT, Russia). Analysis of dry samples was performed in semi-contact mode with a 100×100 μm closed-loop scanner (scanning by sample). Samples were analyzed by high accuracy noncontact composite (HA_NC) ETALON silicon tip cantilevers (NT-MDT, Russia) with a typical resonant frequency of 280 kHz, a tip radius of 10 nm and a force constant of 11.5 N/m in air, and at ambient temperature and humidity. Analysis of hydrated G13 was performed in liquid cell (total volume 1ml) fixed on 100×100 μm closed-loop scanner (scanning by sample) in soft contact mode AFM. Triangular silicon nitride cantilevers (NanoAndMore Inc.) with tip radius bellow 10 nm and force constant 0.08 N/m were used. A scan rate of 0.5–1 Hz was used for the best resolution. Data were collected from at least three different samples, with two different tips and in a minimum of 10 different positions on each sample.

Image processing and height (Z_{max}) analysis. The images were analyzed by Scanning Probe Image Processor software (Image Metrology A/S, Denmark). Raw images were corrected for the tilting of the sample stage and zero leveled based on the dominant height value in the distribution histogram. No other image filtering was used. Objects (at least 100 entities) were analyzed for the Z_{max} (the maximum Z value of all points inside the shape contour) by threshold method using embedded grain analysis module and values obtained presented as height distribution histogram.

Comments. Liquid AFM analysis is very precise method suitable to measure the diameter (Z_{max} value) of symmetric (spherical) hydrated nanoobjects as are dendrimers. The important prerequisite for such analysis in liquid is the immobilization of sample on atomically flat surface. Fortunately, the **G13** triazine dendrimers are positively charged at neutral pH and therefore able to be immobilized electrostatically on negatively charged mica surface. AFM analysis of the hydrated dendrimers revealed the predominant presence (\approx 66%) of subpopulation of **G13** with dimensions within the

interval 29-36 nm (see histogram **Fig S43**) with average $Z_{max} = 31.5 + /- 1.9$ nm (n = 93)). Although there were present also other subpopulations of dendrimers in lower range than expected (lower or incomplete generations), their overall percentage is much lower compared to largest subpopulation in histogram. Based on the HPLC, the purity of sample has been predicted more than 95%. We therefore expect that the presence of **G13** (largest subpopulation) will be significantly higher than $\approx 66\%$ as obtained from AFM analysis. **G13** tend to aggregate compared to lower generations (as observed by DLS). This aggregation is further increased during the AFM sample preparation (preconcentration step on mica surface). The relative proportion of **G13** could be underestimated compared to its presence in solution due to aggregation.

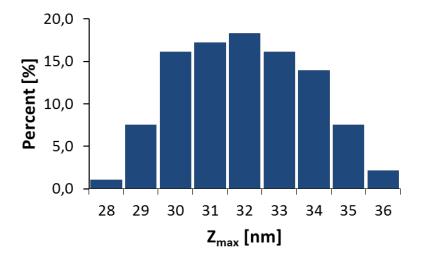


Fig. S43. Histogram of Z_{max} distributions (n = 93) of G13 dendrimers as obtained by AFM in liquid

Fig. S45 shows AFM analysis of G13 on air (dry sample). Based on the cross section profile and histogram is apparent, that the desiccation process leads to significant changes in volume and Z_{max} values of dendrimers, absorbed electrostatically on the surface of mica. Since the diameter of the dendrimers measured on dry samples by TEM is approx. 25-30 nms (measured at x-y scale), we may assume that the dendrimer shrinks predominantly in the Z-scale (average final value 9.8 +/- 1.9 nm) towards the mica surface resulting in disc shape. Volume and Z_{max} changes can be explained by gradual loss of water molecules during the drying process and possibly also by attractive forces between the negatively charged mica and positively charged amino groups of dendrimer which may deform the expected spherical shape of G13. The partial aggregation of G13 observed on Fig. S44 is the most probably the result of the preconcentration of dendrimers on mica surface during the drying process. The extent of the volume changes observed suggests natural high hydration state of internal space of dendrimer.

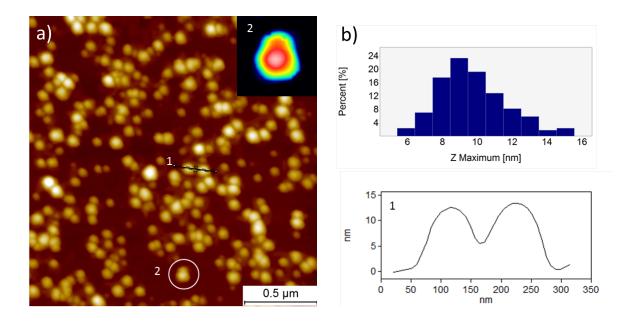


Figure S44. Atomic force microscopy (AFM) analysis of G13 on air (dry sample). a) G13 imaged on mica surface. Cross section profile (average of 3 lines) of two dendrimers (marked 1) is shown on the right panel. Inset: close view of G13 dendrimer marked by cycle (numbered 2) in the lower part of the image (image size 200×200 nm). b) histogram of Z_{max} values obtained from AFM analysis.

Computation

Experimental. The molecular models for the whole dendrimer series were constructed and parametrized according to a validated procedure. S1-S3 The dendrimers were built iteratively in consecutive generations from G1 to G13 by using the Material Studio software (Accelrys, San Diego, CA, USA). Each built generation underwent to preliminary gas-phase minimization and short molecular dynamics (MD) simulation in order to relax the structure and to eliminate bad contacts within the dendritic scaffold before growing with the next generation. For the larger generations – i.e., G9, G11 and G13 – this procedure was extremely delicate and time consuming, since complexity increases exponentially with the dendritic generation. For this reason, it has been very hard to obtain an atomistic model for G13, which required many steps of minimizations and of MD simulations with increasing temperature and time step. In particular, T was increased from 0 to 1000 K and then the system was cooled down again to 300 K, with increasing time step from 0.5 to 2 femtoseconds (fs) due to high atom velocity in the earlier steps of this phase. The final G13 model, subject of this study, is composed by different residues - a central (COR) core unit, the repetitive units that compose the flexible branches (BRA) and the terminal groups which constitute the dendrimer's surface (END). G13 has 16384 surface groups bearing primary amino-groups that at neutral pH (\approx 7.4) are assumed to be protonated. The entire parametrization, simulation work and data analysis were carried out using the AMBER 12 suite of programs. S4 The partial charges for all of the non-standard residues that constitute G13 were obtained using the AM1-BCCS5 calculation method within the antechamber^{S6} module of AmberTools 12 (AMBER 12). Parameters and force field types were assigned consistently with the "general AMBER force field (GAFF)" (qaff.dat)^{S7} - such parameters already demonstrated to be well-consistent for the parameterization and the simulation of dendrimers s1-52,58 and dendrons. S3,S9

Comments. Due to the enormous size of the atomistic model of **G13** (1343416 atoms), for this generation it was not possible to run a molecular dynamics (MD) simulation in explicit solvent, as the number of water molecules introduced in the system largely exceed the maximum size limit that it is currently possible to

simulate. Even the use of a simplified coarse-grained (CG) model to decrease the system size was not practicable in this case. In fact, first, CG models that account the presence of explicit solvent did not allow for the size reduction necessary to run the simulation – the system was still too large. Secondly, regarding the CG simulation of G13 in absence of explicit solvent, a very coarse model would have been necessary to reduce substantially the size of G13 to make the use of a CG model convenient. This was incompatible with the highly flexible branches that compose the dendritic scaffold. In fact, the approximation the linear monomers of G13 into a single, or two-bound beads would not reproduce correctly the flexibility of the molecule nor the surface roughness and irregularity. For this reason, a fully atomistic MD simulation of G13 was conducted *in vacuum* to obtain a single-molecule view of this large dendrimer. G13 was initially minimized and then preliminary 100 ps of MD simulation a 1 fs time step were run at 300 K of temperature. G13 was then equilibrated for 4 ns at 300K, using a time step of 2 fs, the Langevin thermostat and a 12 Å cutoff. The SHAKE algorithm was used on the bonds involving Hydrogen atoms. The reduced simulation time was sufficient to obtain a molecular picture of the dendrimer, as G13 reached rapidly the equilibrium due to the high structural crowding and rigidity. Size (radius of gyration) and radial distribution functions (RDF) for G13 were extracted from the equilibrated phase of the MD trajectories using the *ptraj* module within AMBER 12.

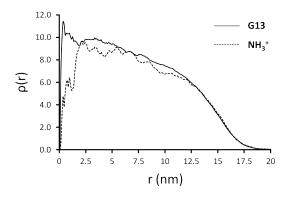


Figure S45. Radial distribution function of **G13** showing peripheral amine groups reveal extensive backfolding. observed.

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