

Copyright Wiley-VCH Verlag GmbH & Co. KGaA, 69451 Weinheim, 2004  
*Chem. Eur. J.* **2004**

Supporting Information

for

**Synthesis and Esterolytic Activity of Catalytic Peptide Dendrimers**

By

David Lagnoux, Estelle Delort, Céline Douat-Casassus, Annamaria Esposito and Jean-Louis

Reymond\*

*Department of Chemistry and Biochemistry, University of Bern, Freiestrasse 3, 3012 Bern,*

*Switzerland, FAX +41 31 631 80 57*

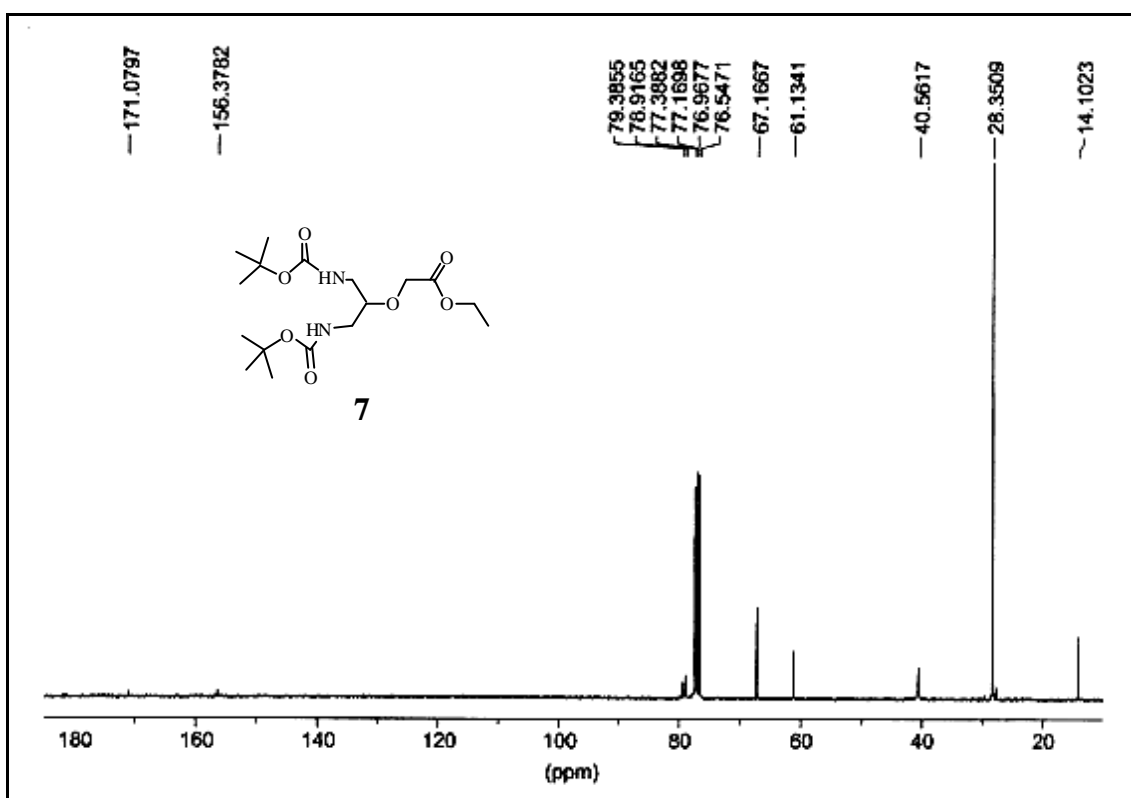
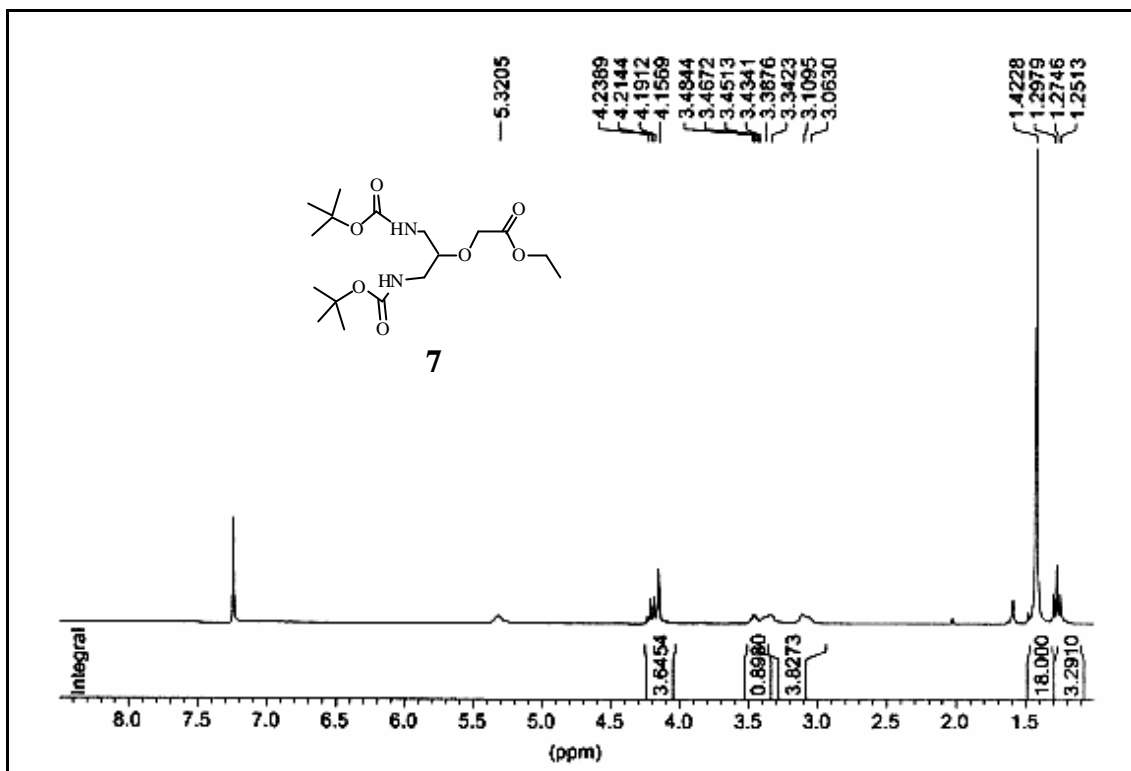
e-mail: [reymond@ioc.unibe.ch](mailto:reymond@ioc.unibe.ch)

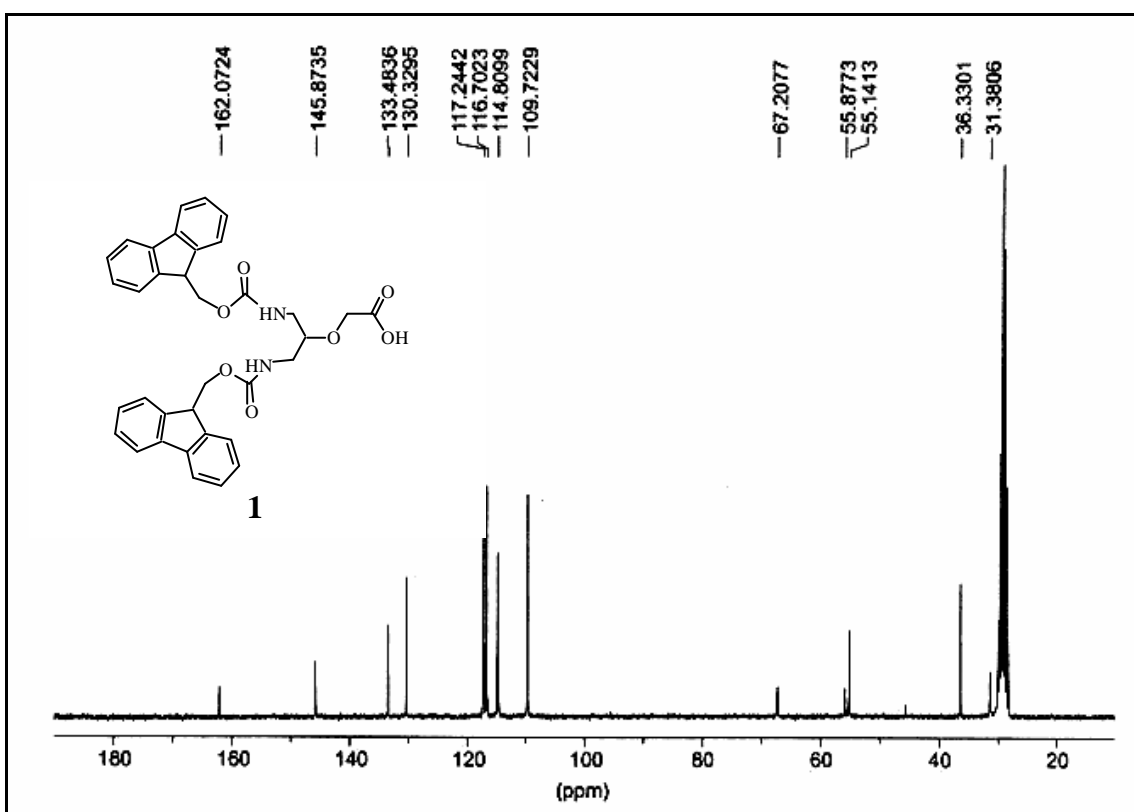
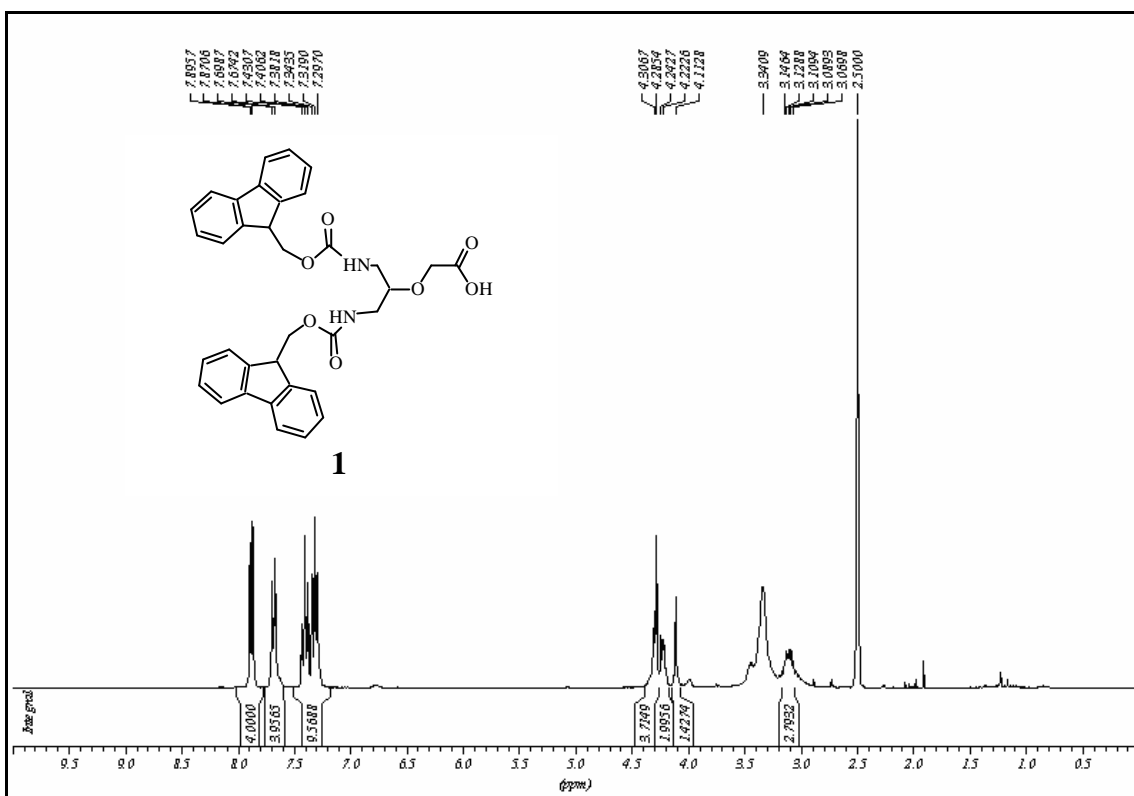
**<sup>1</sup>H and <sup>13</sup>C NMR for intermediates and linkers 1, 2; capping unit 4; quinolinium substrates 15-17 and 19-21.**

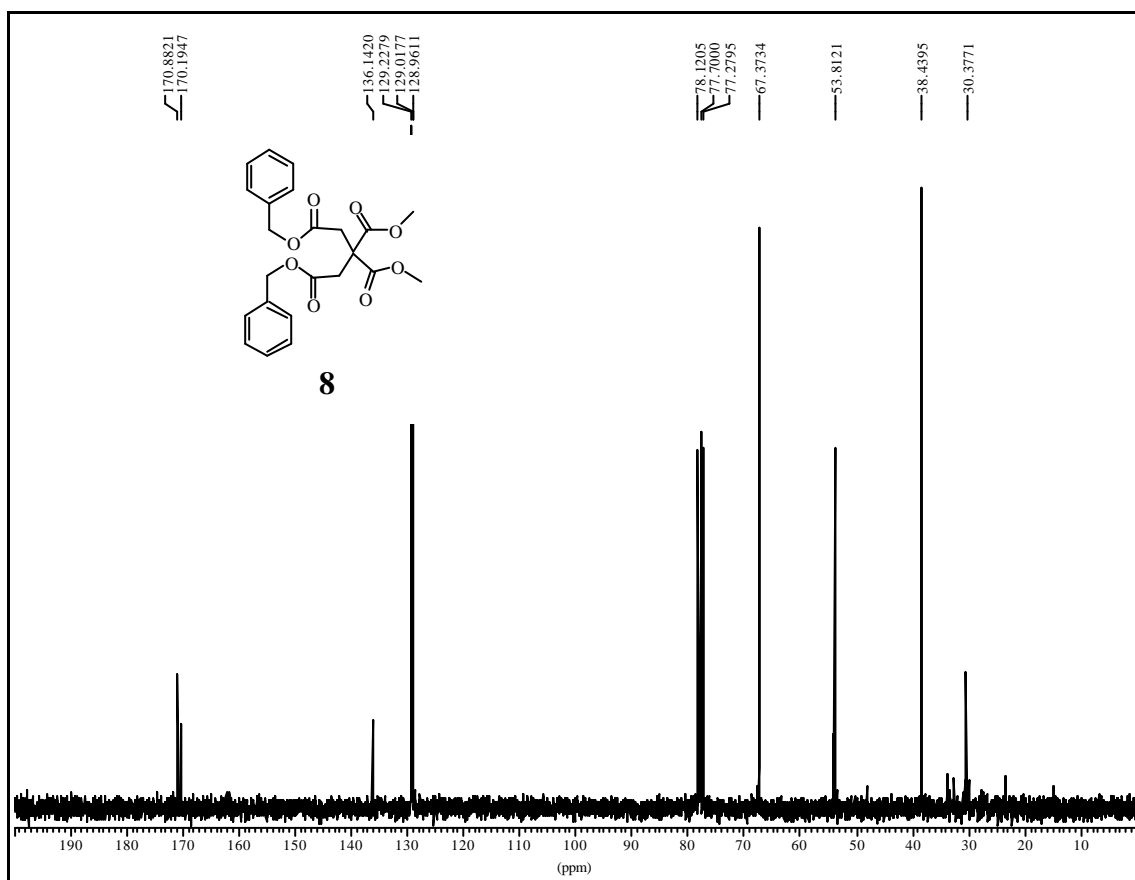
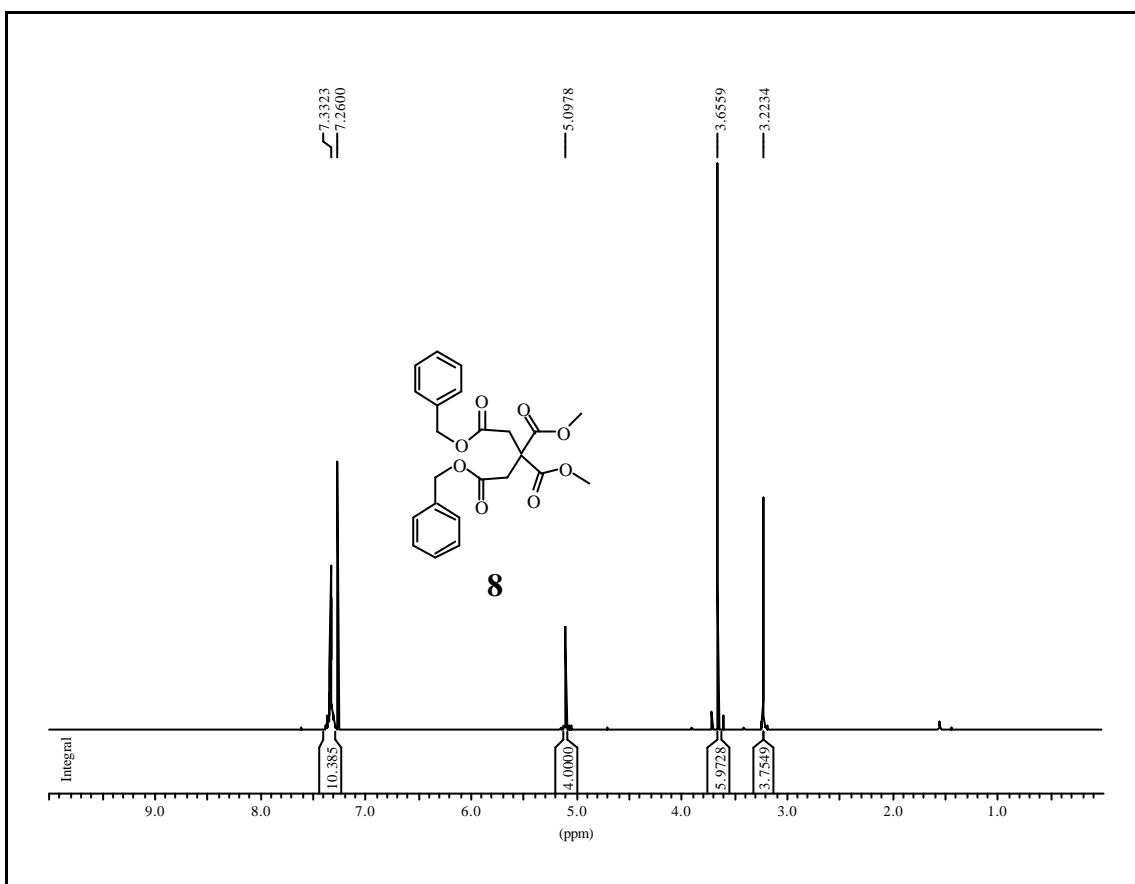
**Analytical HPLC profiles and Mass spectra (ESI MS) data for the monomeric and dimeric peptide dendrimers**

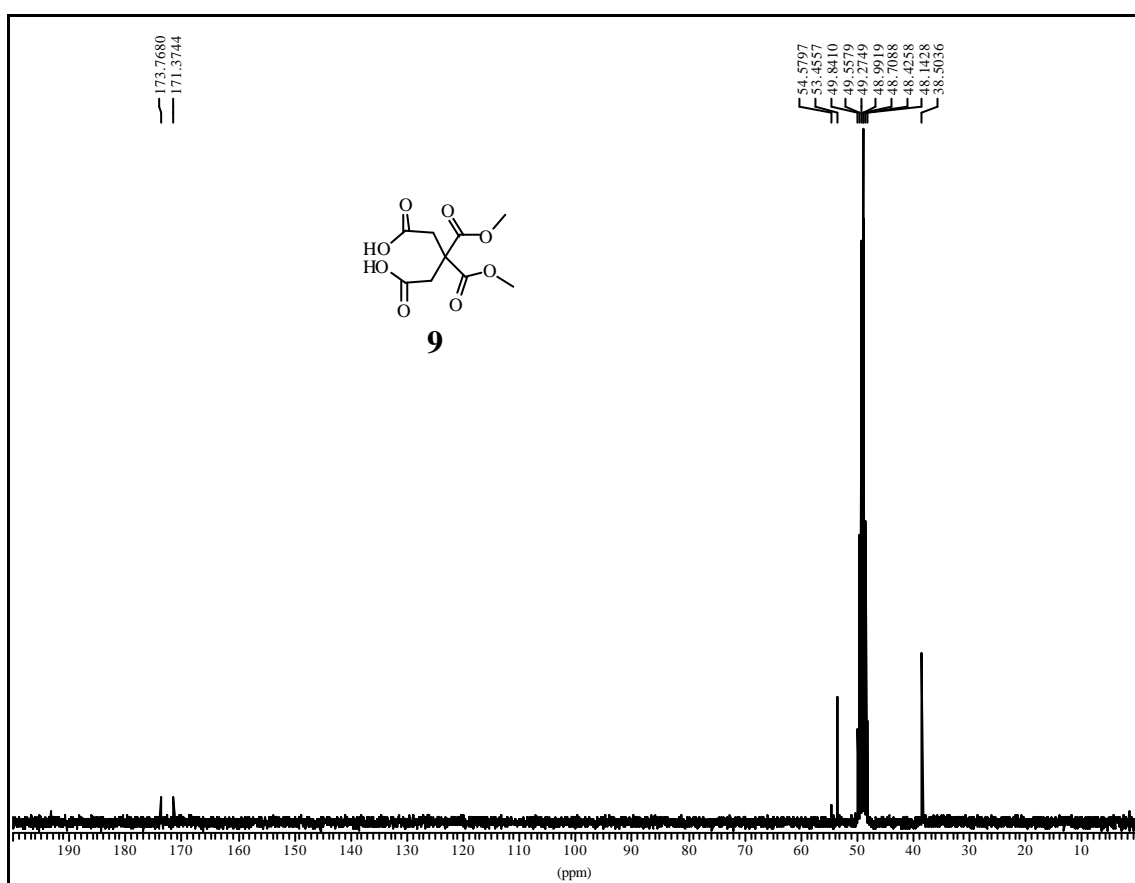
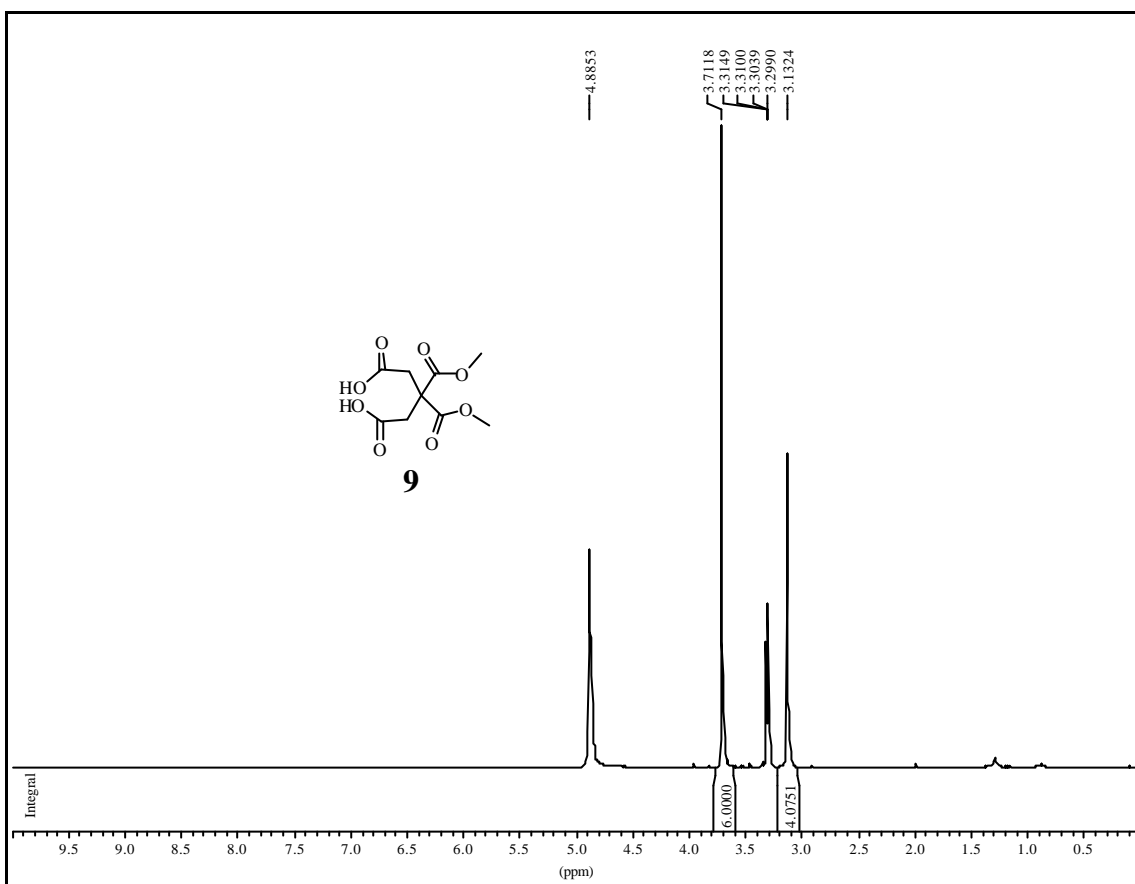
Analytical HPLC columns: Vydac 218TP-54 C<sub>18</sub>, pore size 300 Å, 0.45 × 22 cm, flow 1.5 mL.min<sup>-1</sup> or Chromolith Performance RP-18e, 0.46 × 10 cm, flow 5 mL.min<sup>-1</sup>; detection by UV at 214 nm, solvent systems: A = 0.1% TFA in H<sub>2</sub>O; B = 40% H<sub>2</sub>O / 60% CH<sub>3</sub>CN, 0.1% TFA.

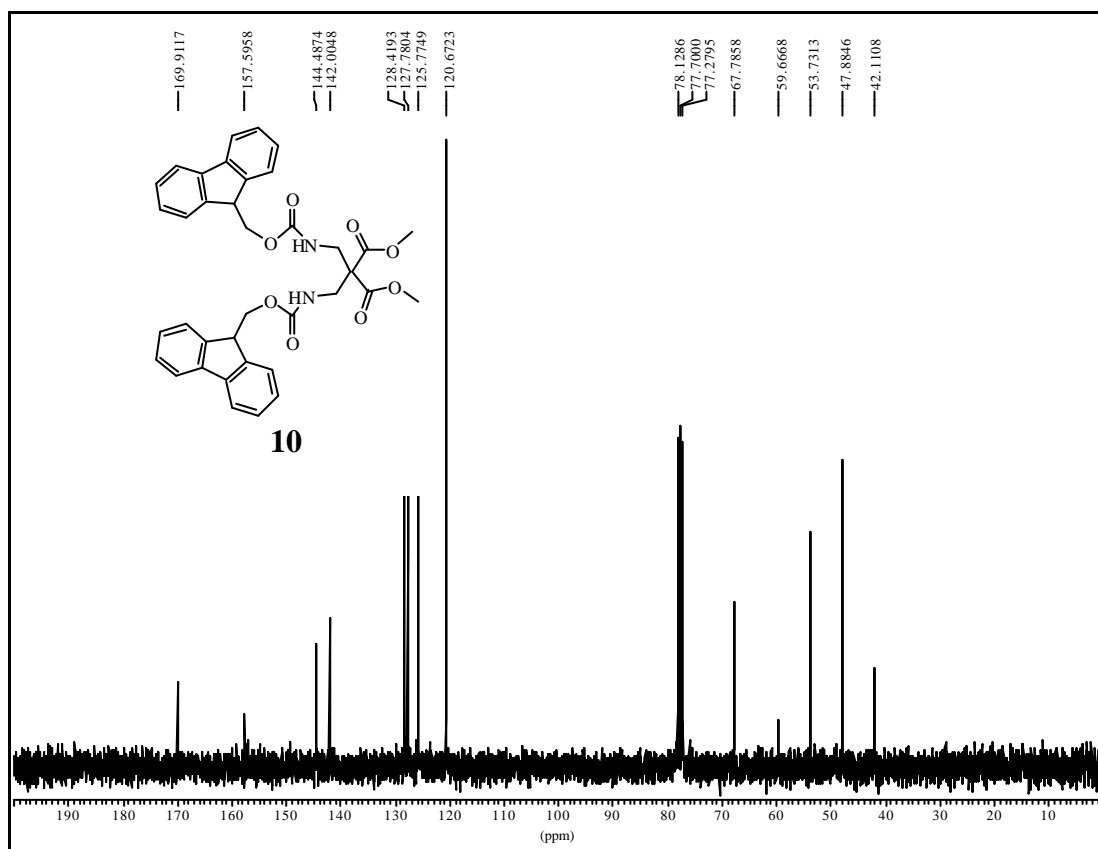
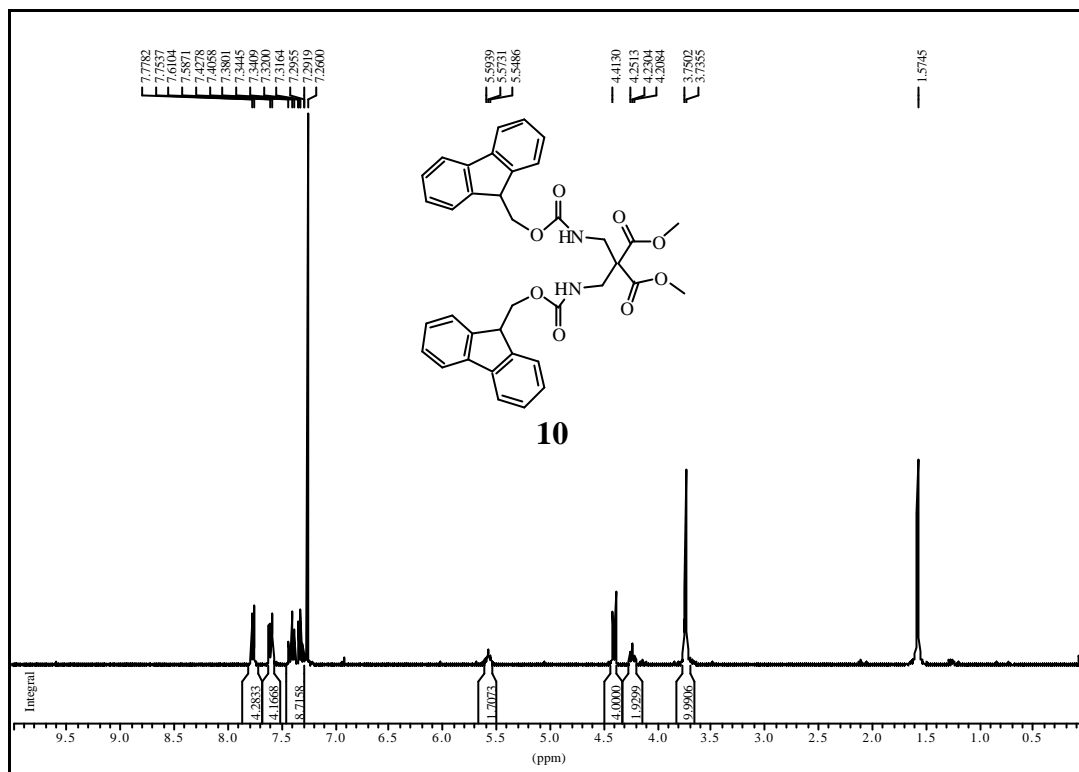
Starting conditions: for dendrimers with linkers **1, 2, 3**: 80% A / 20% B, gradient +1% B / min; for hydrophobic dendrimers: 20% B to 100% B in 5 min.

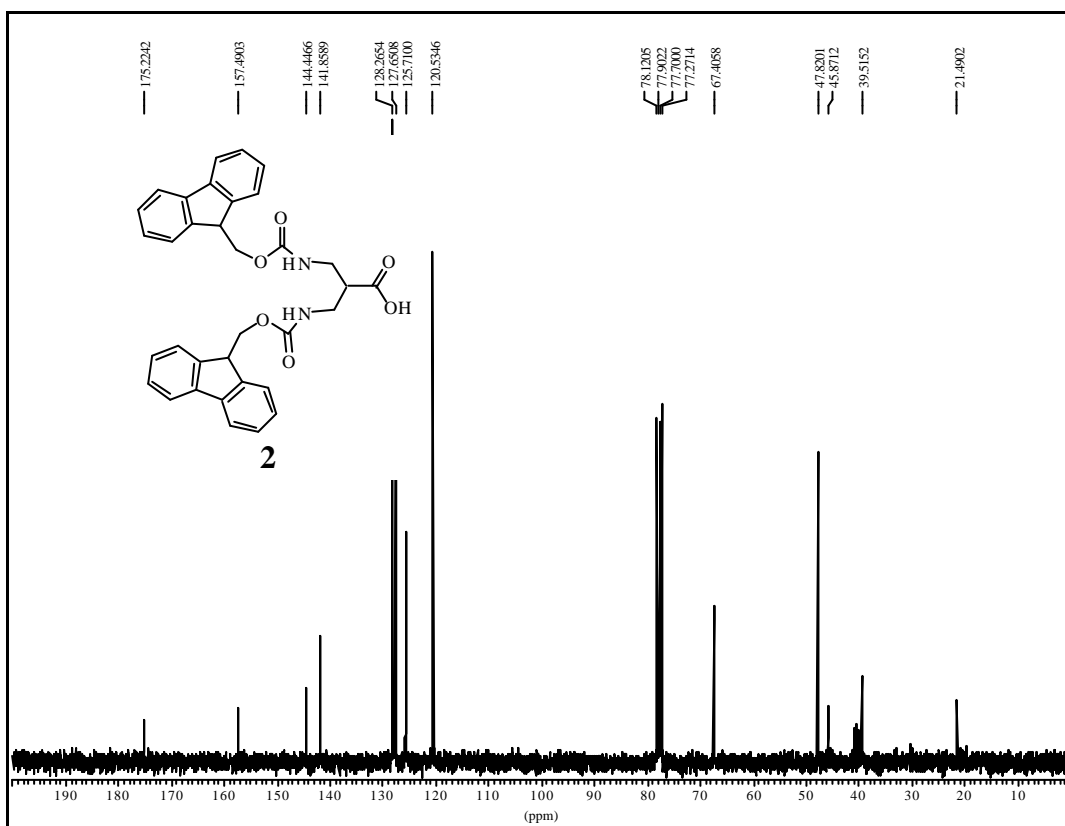
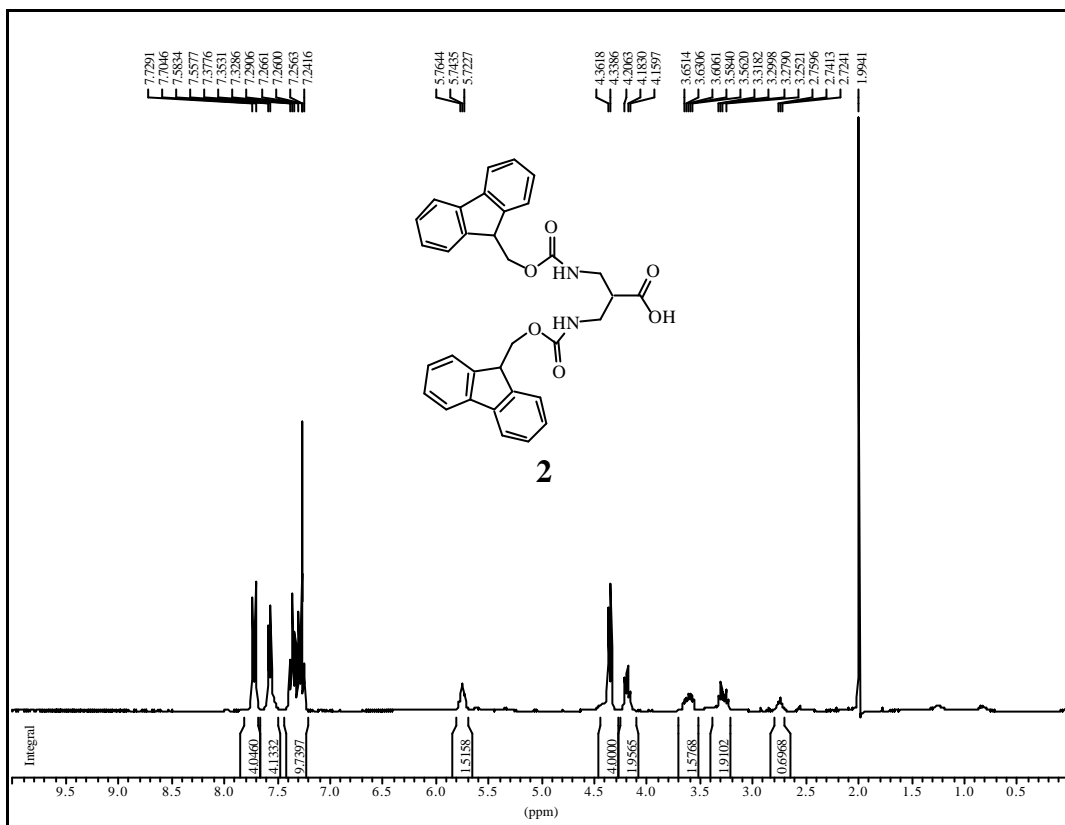




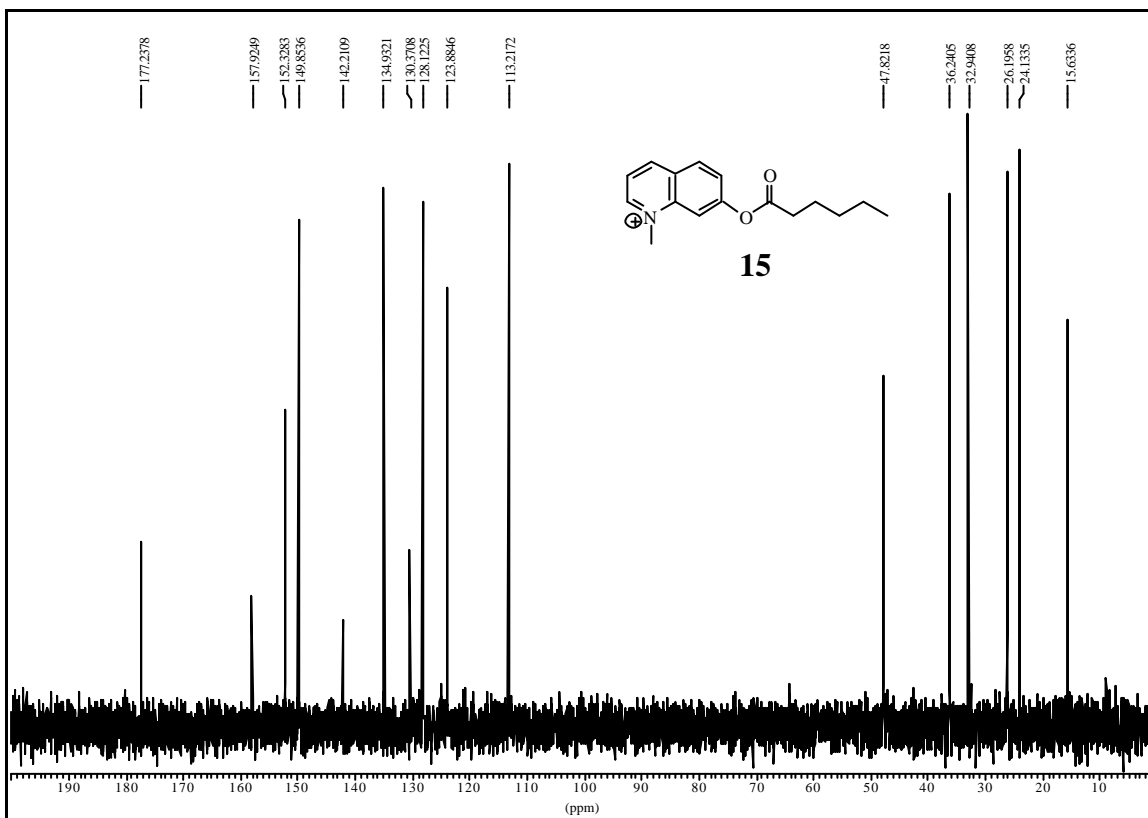
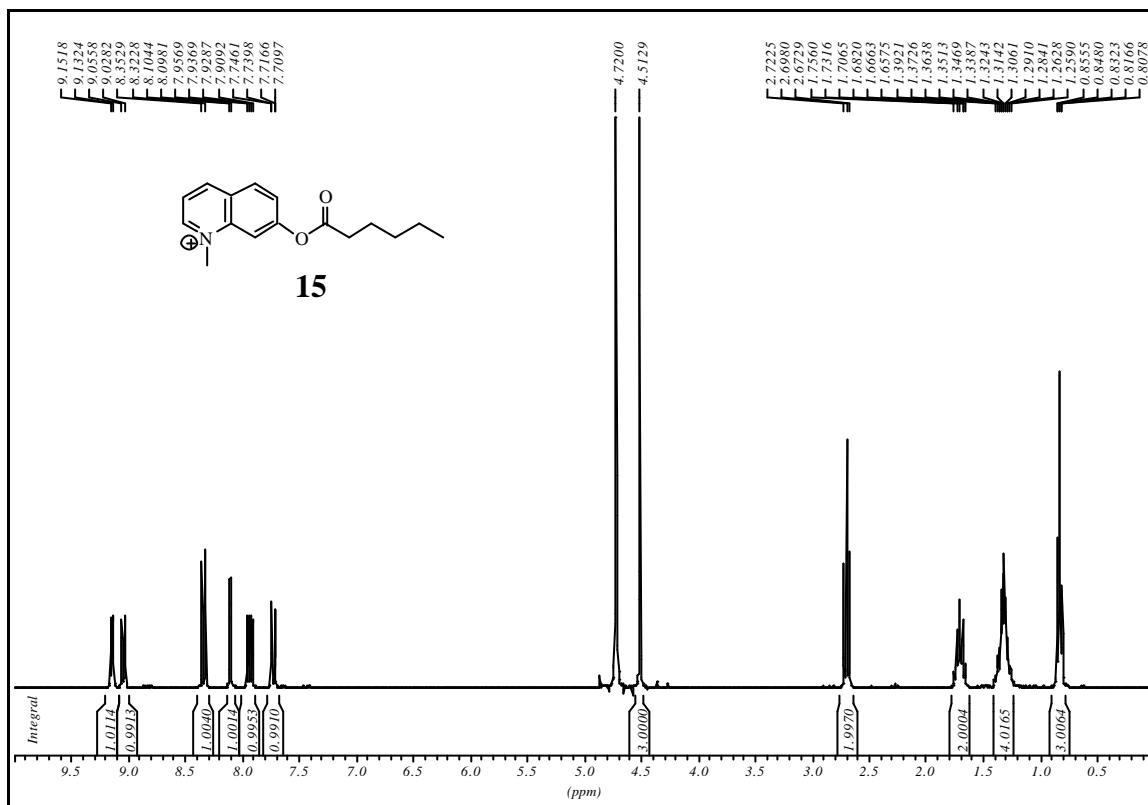


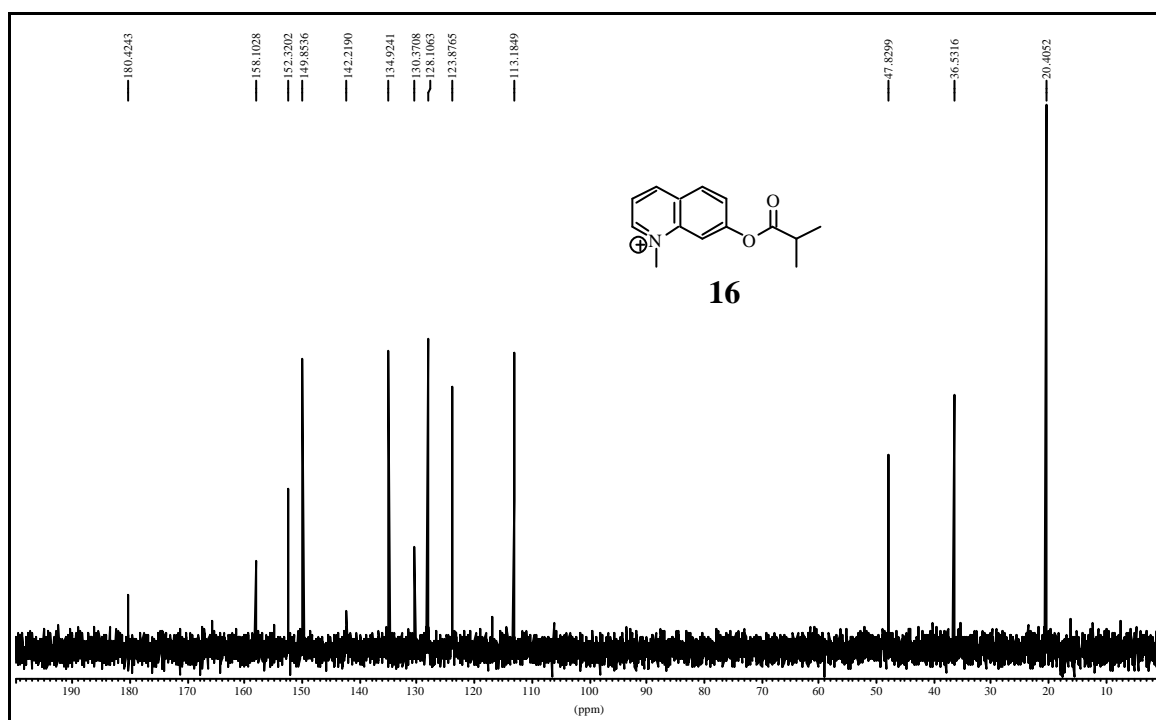
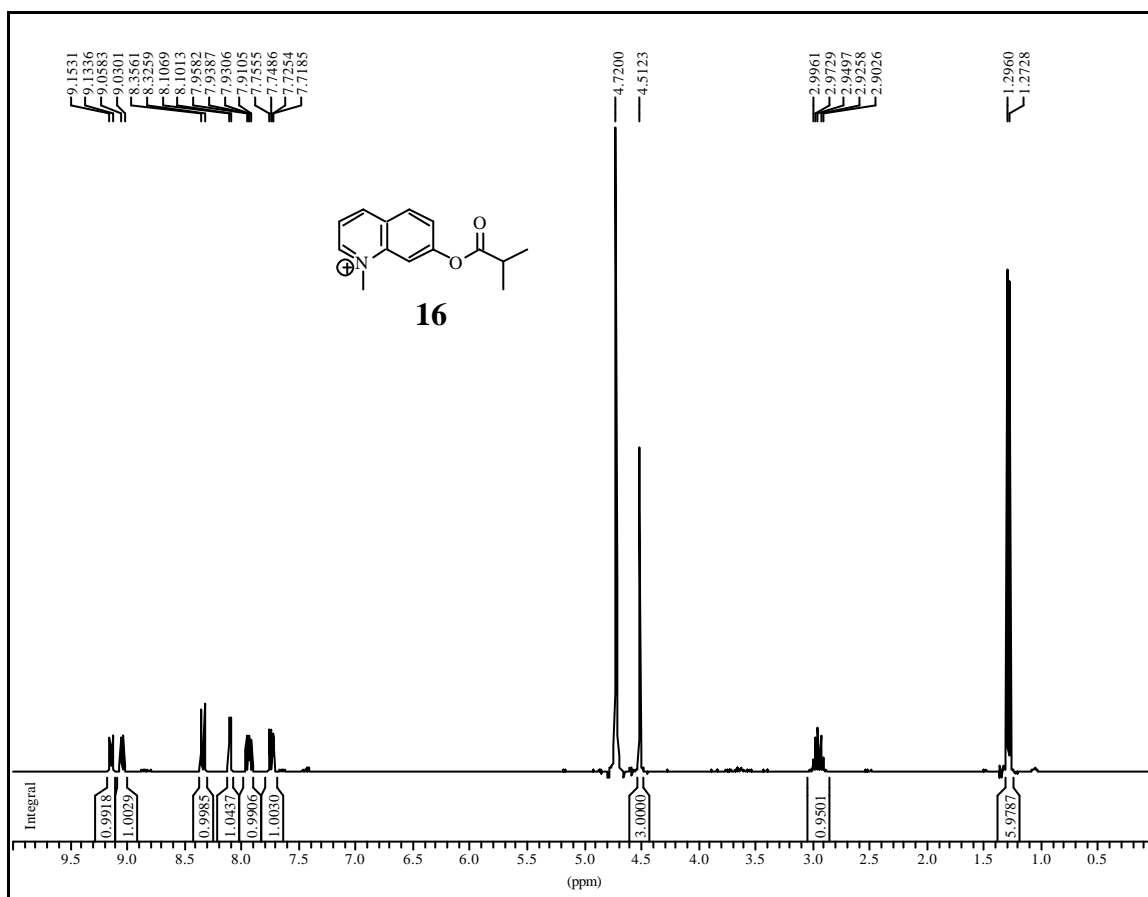


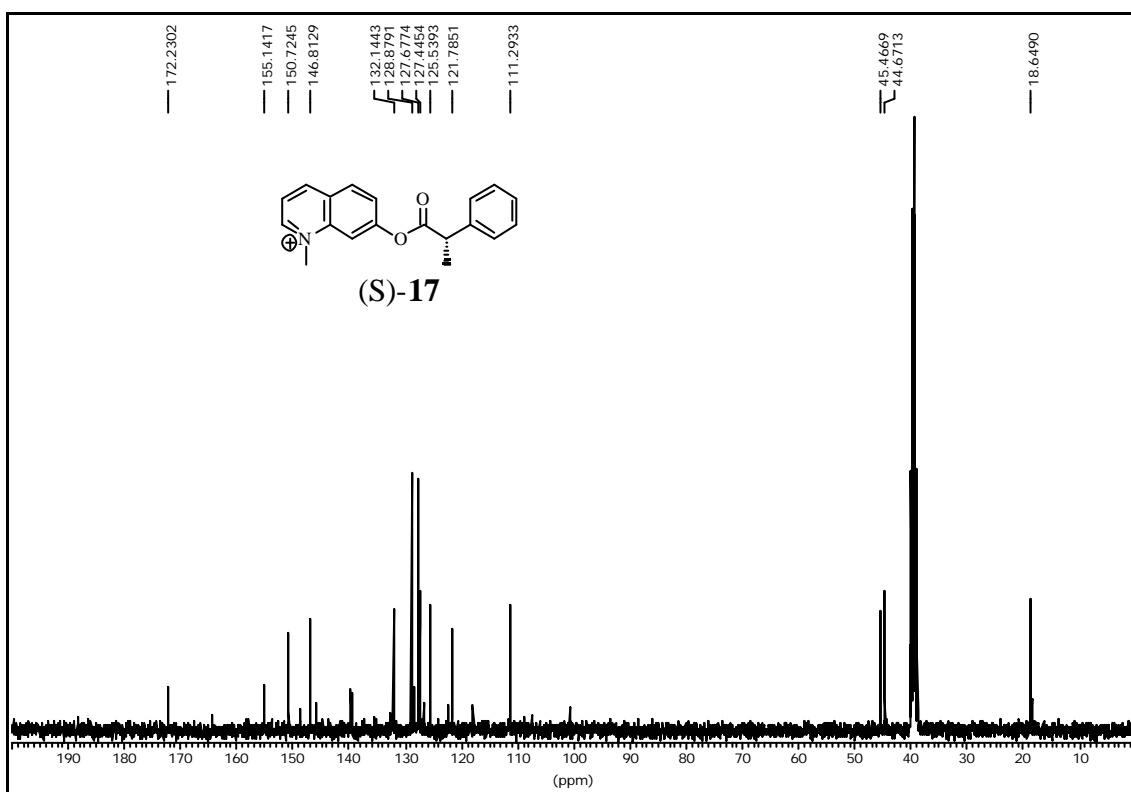
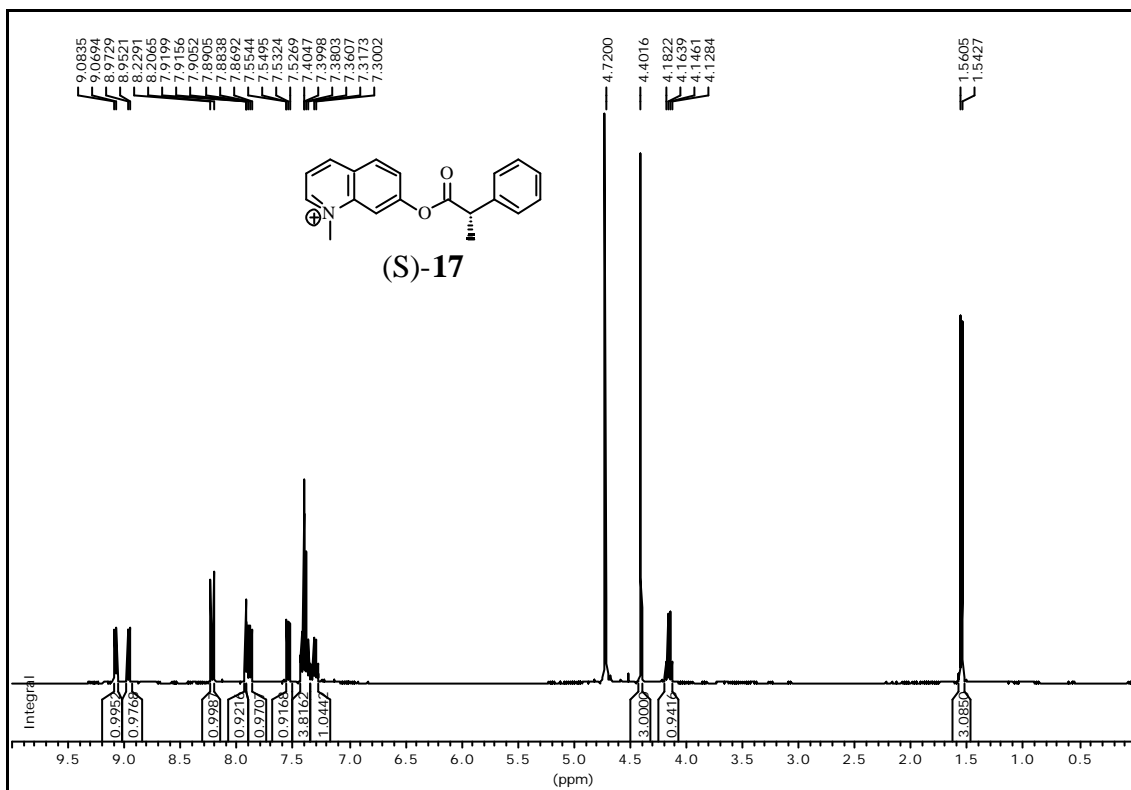


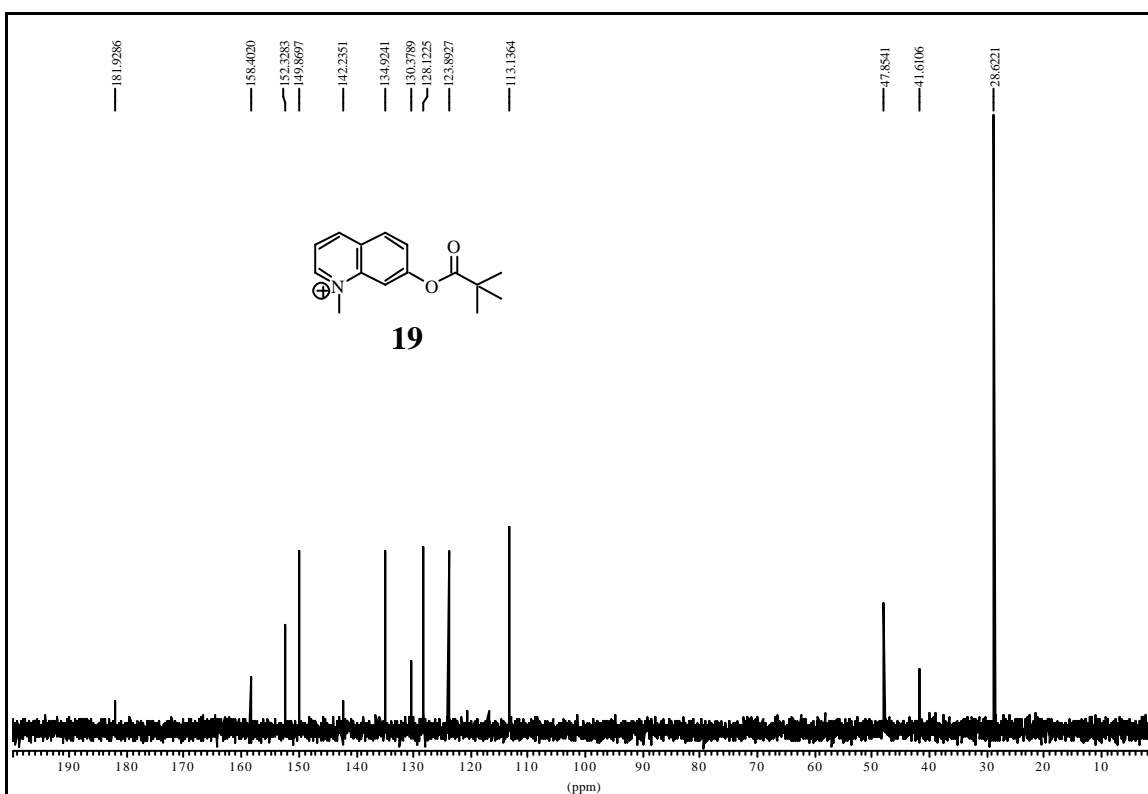
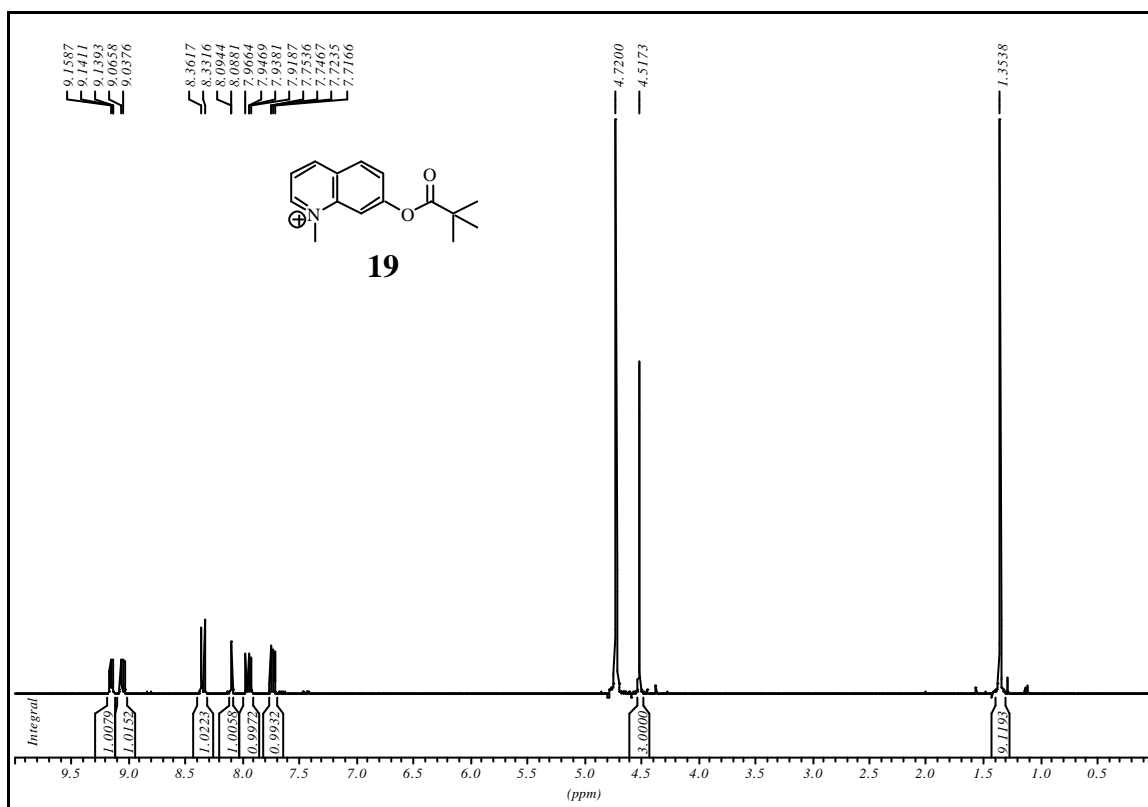


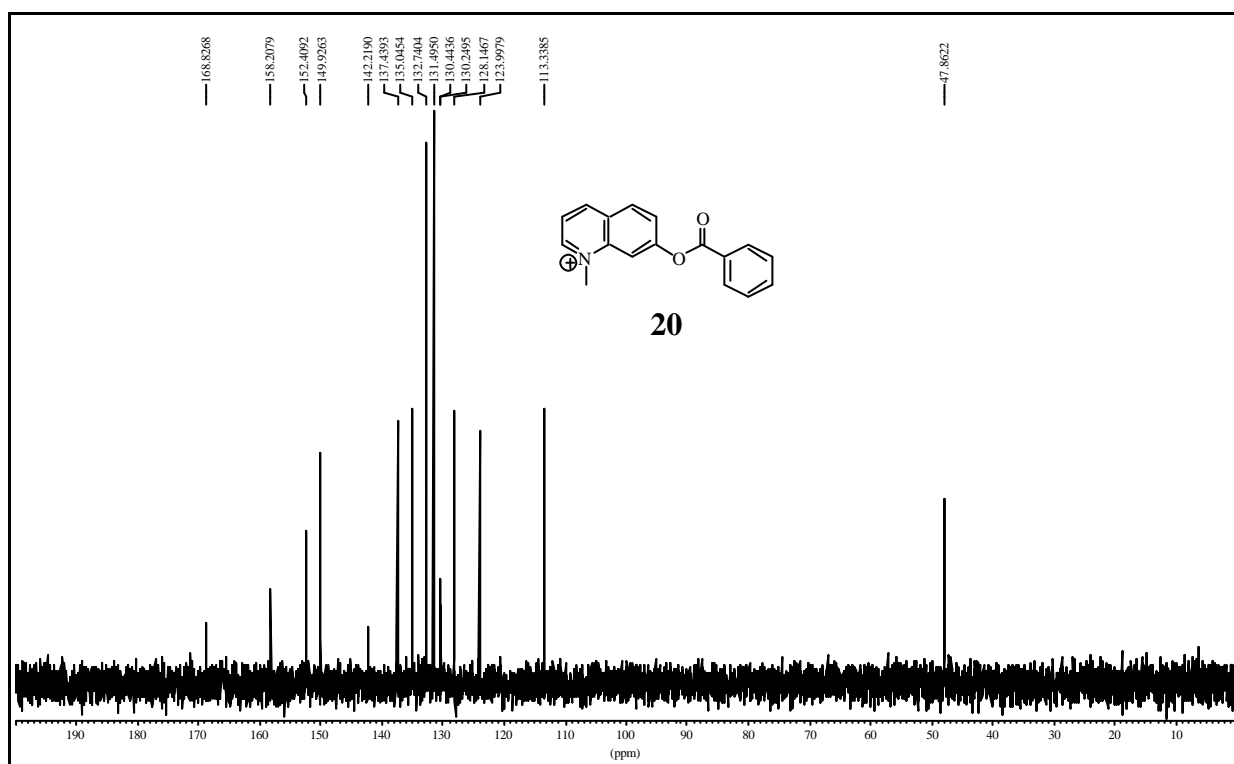
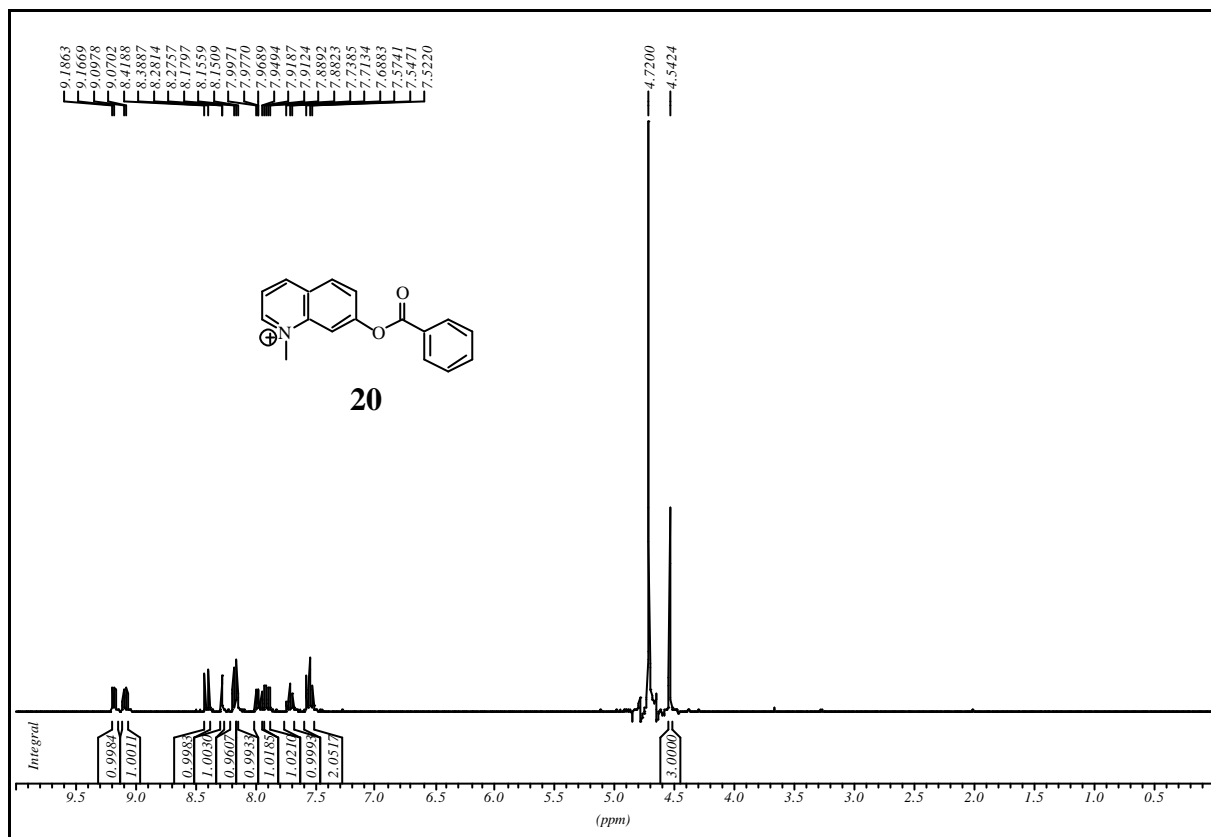


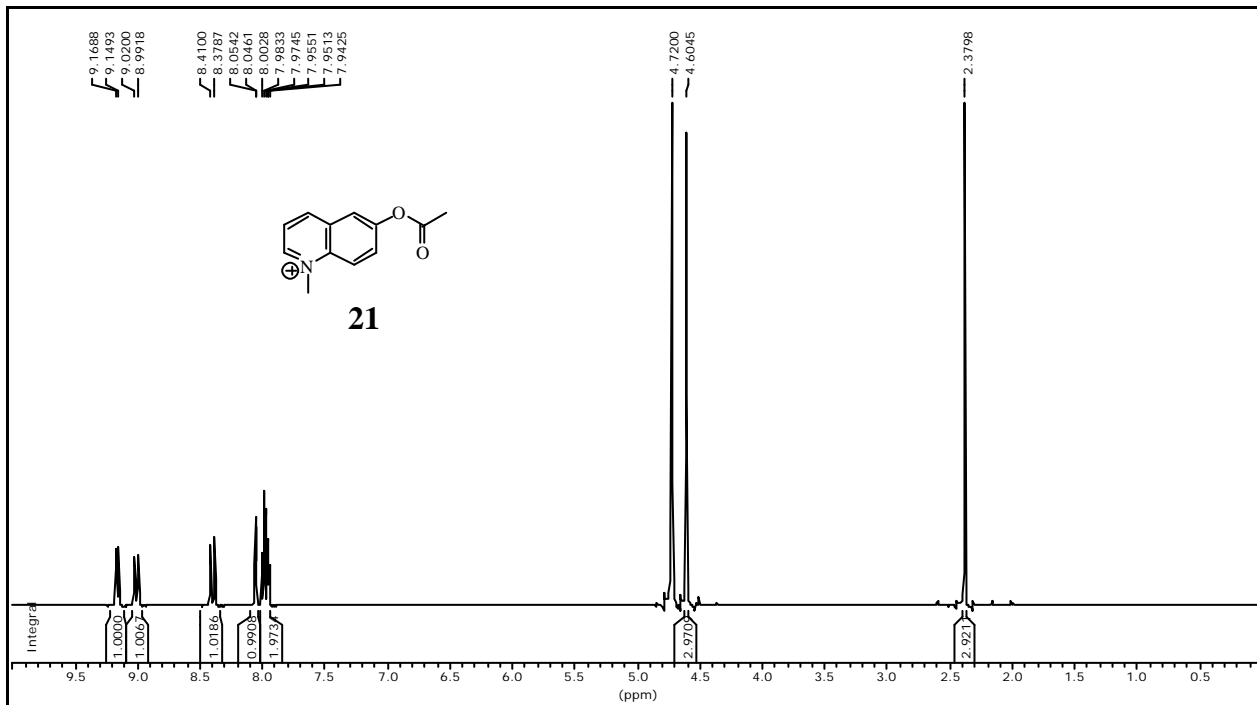
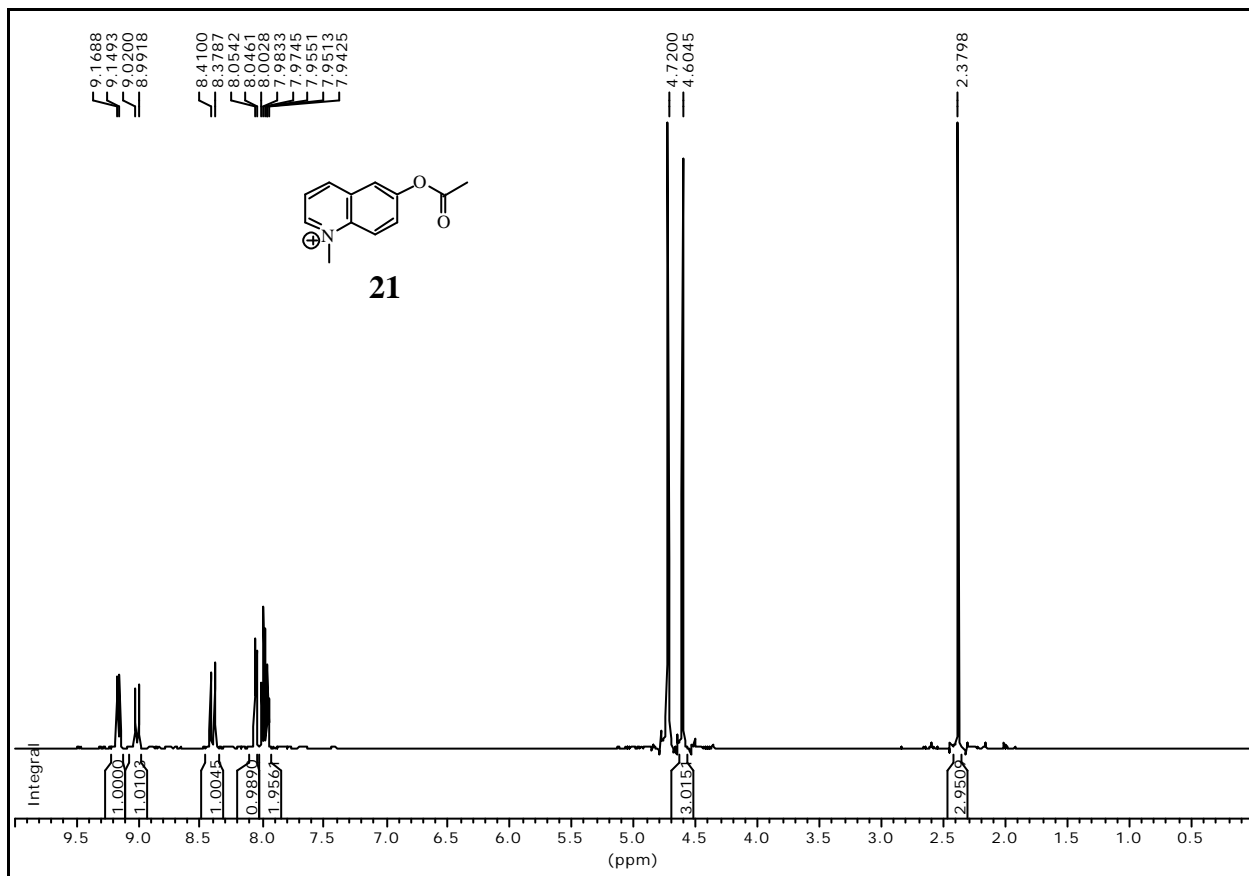


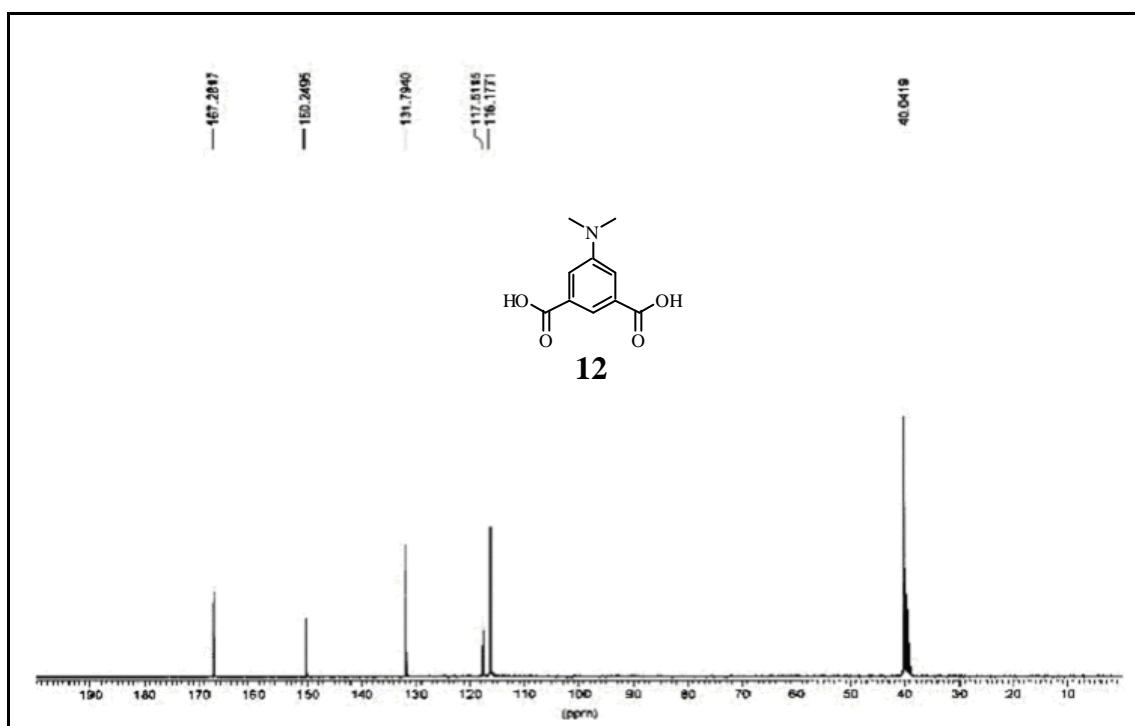
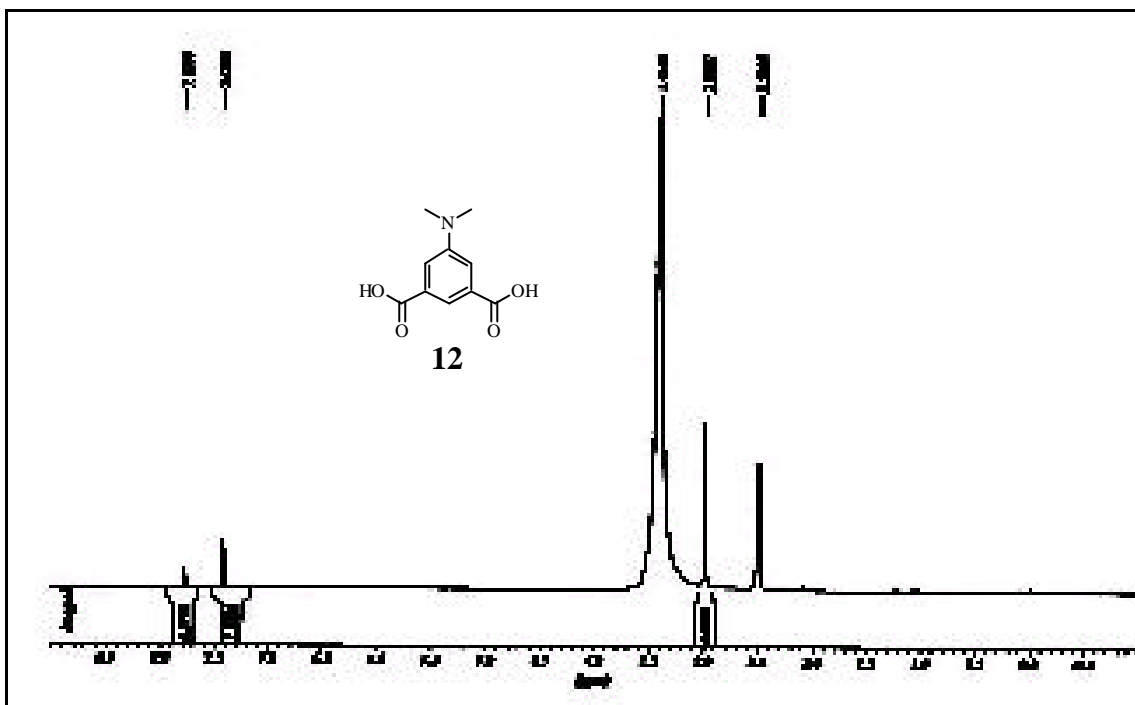


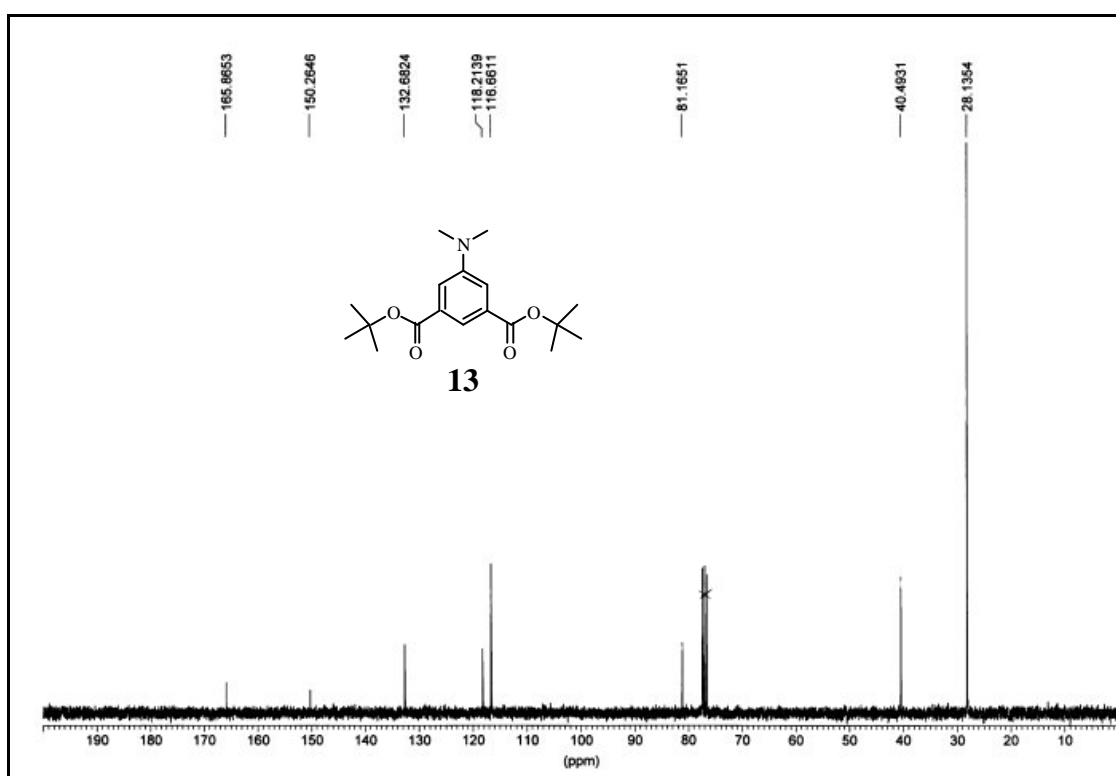
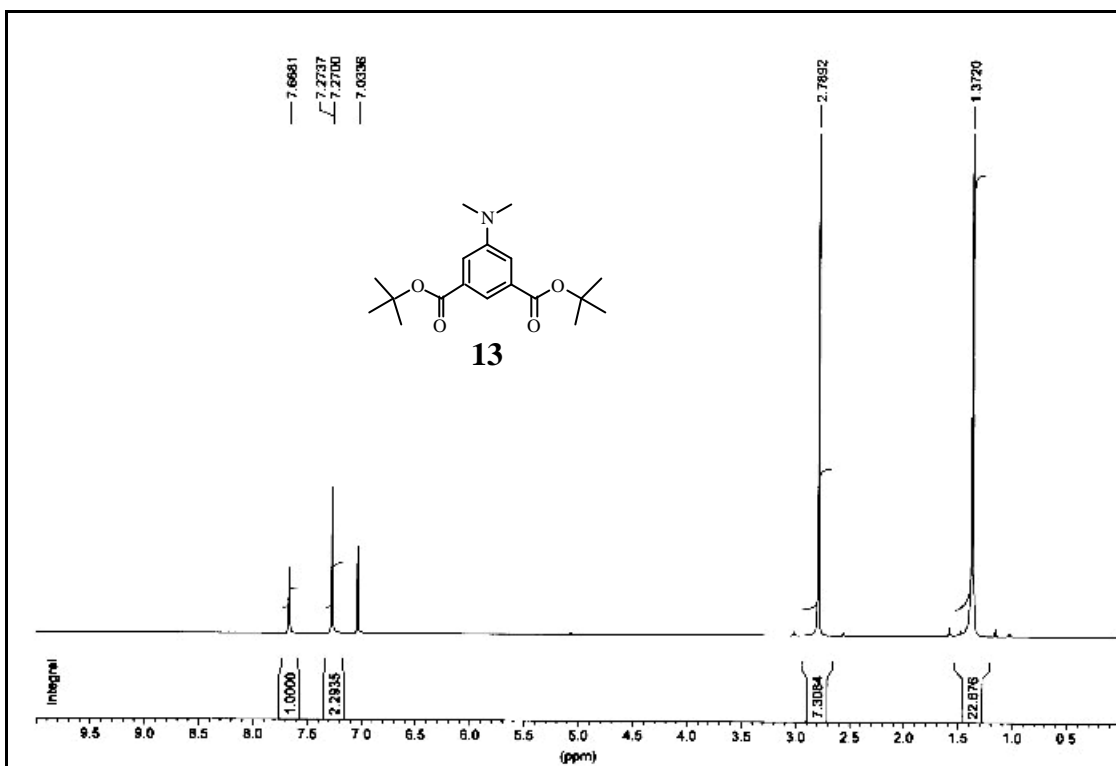




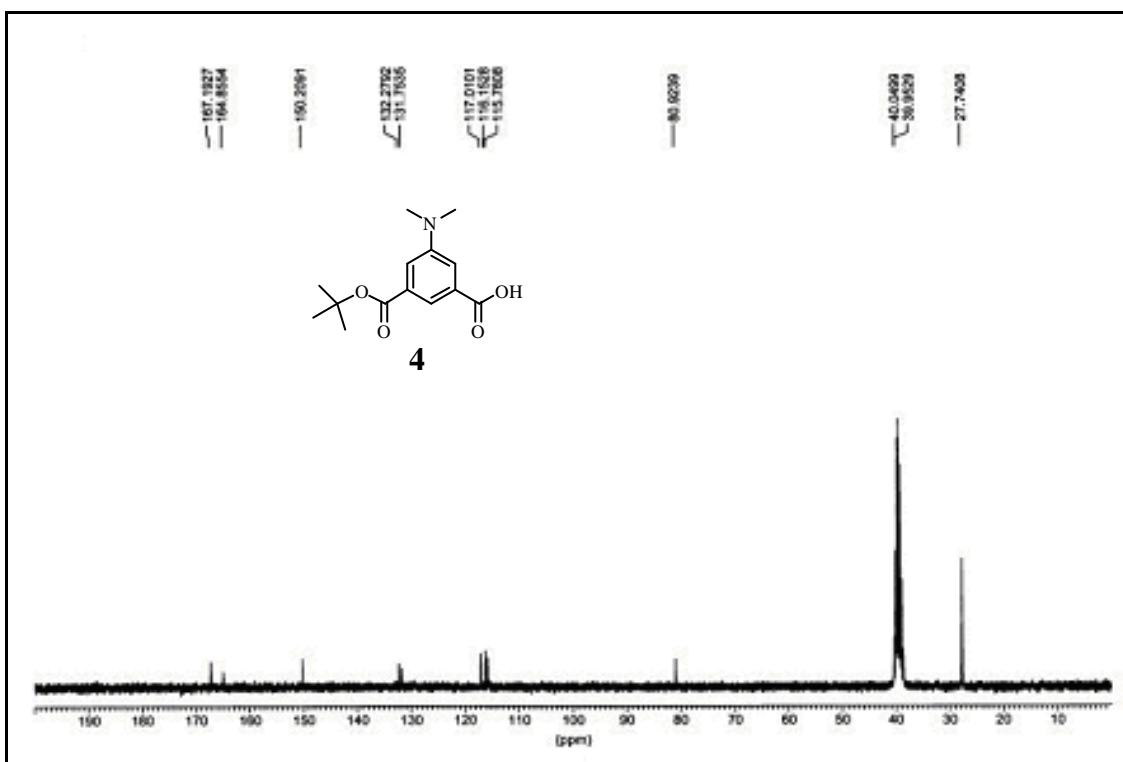
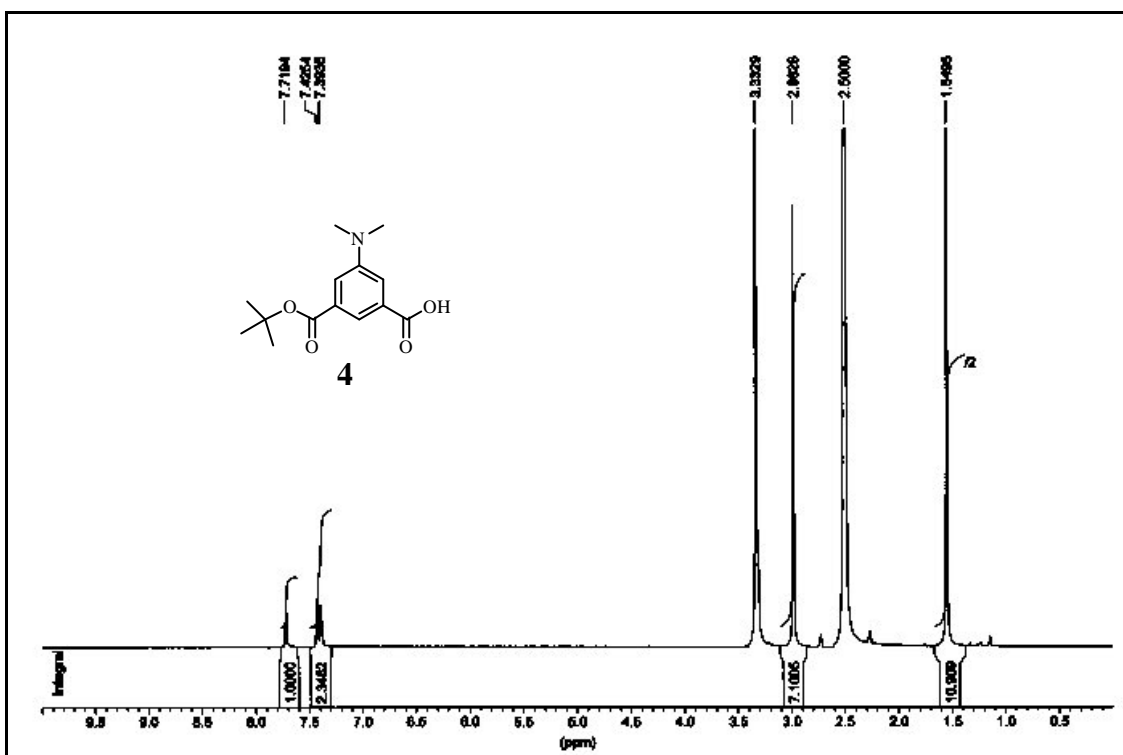


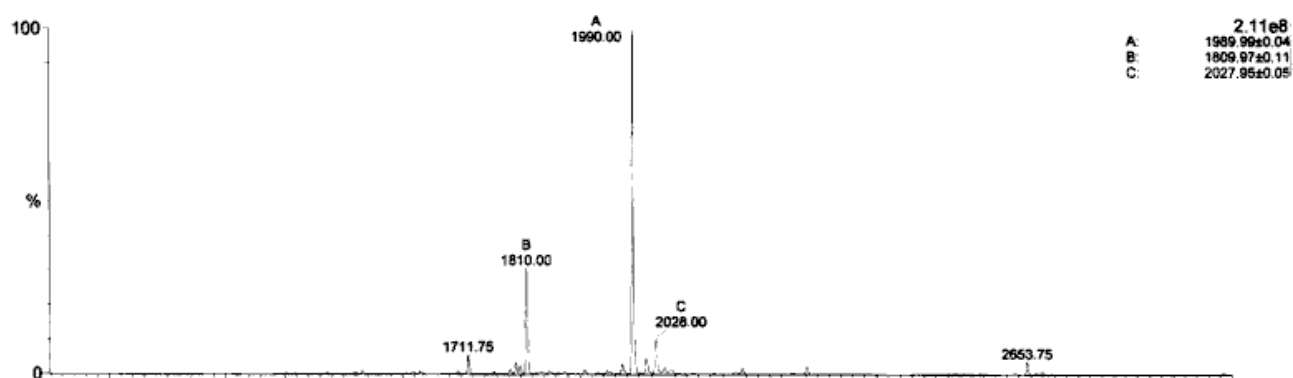
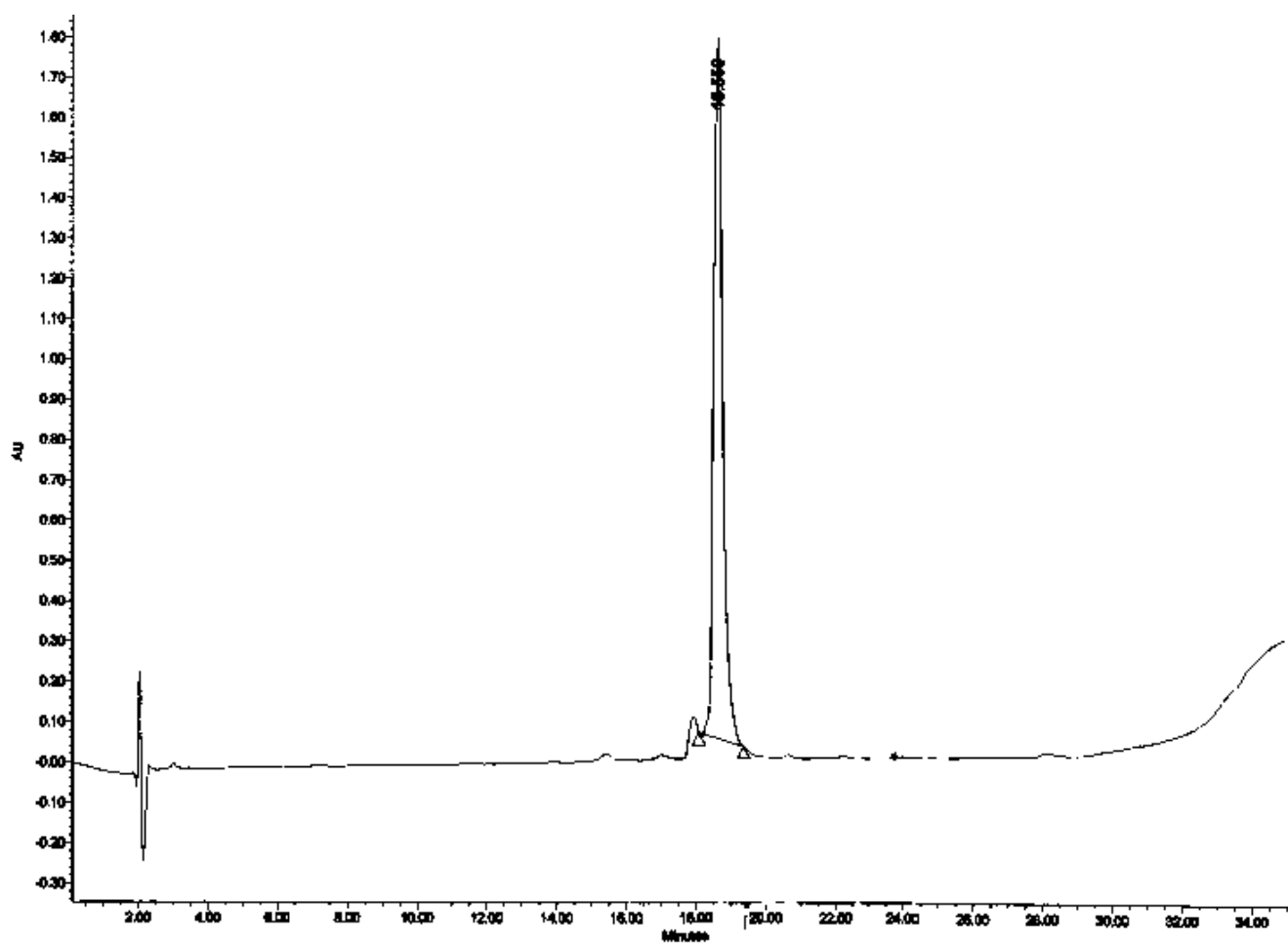




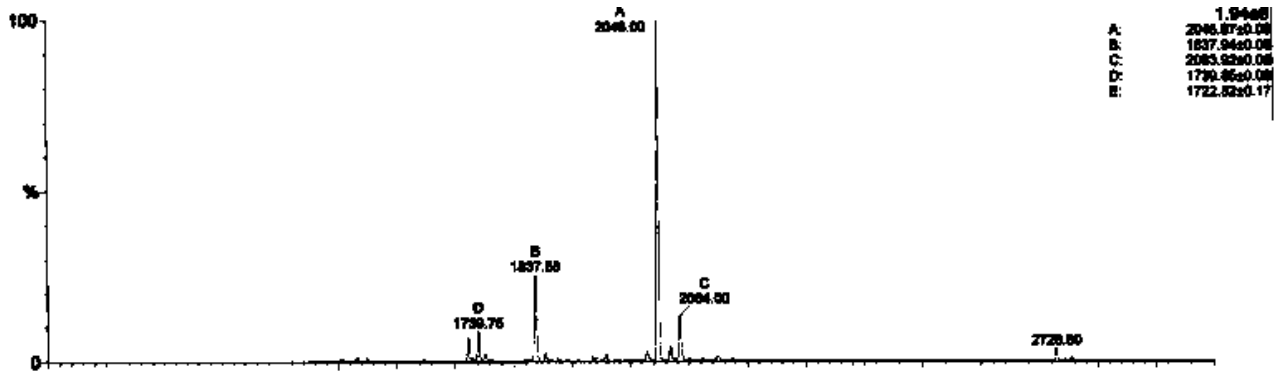
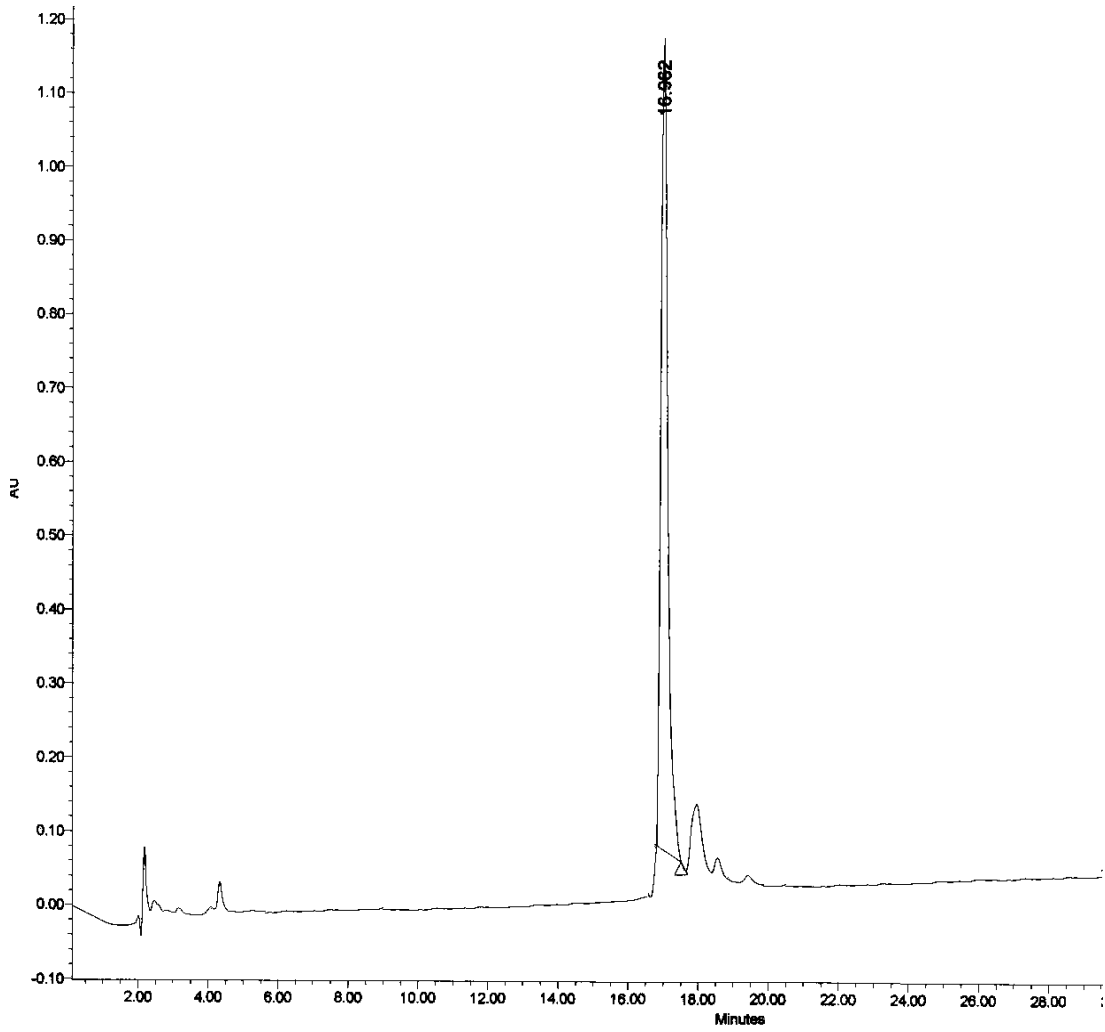


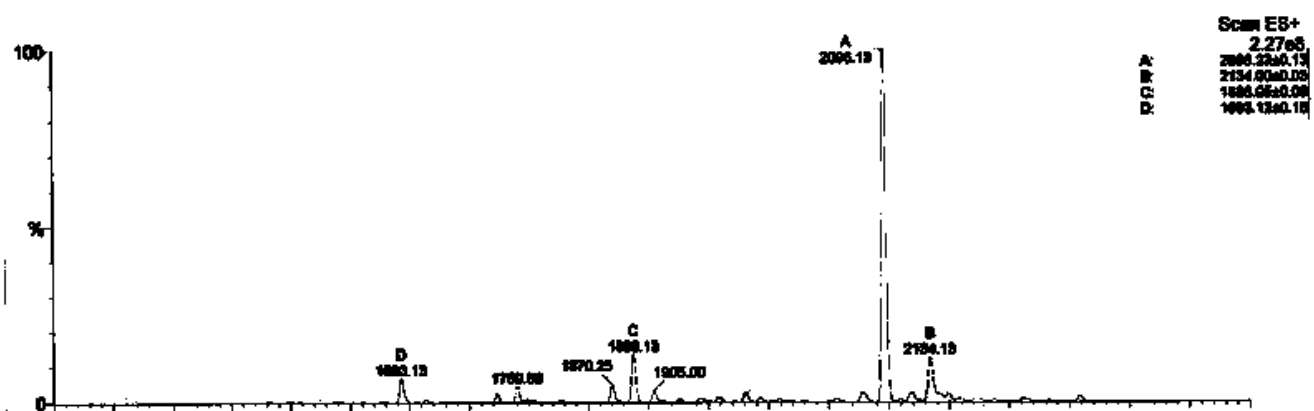
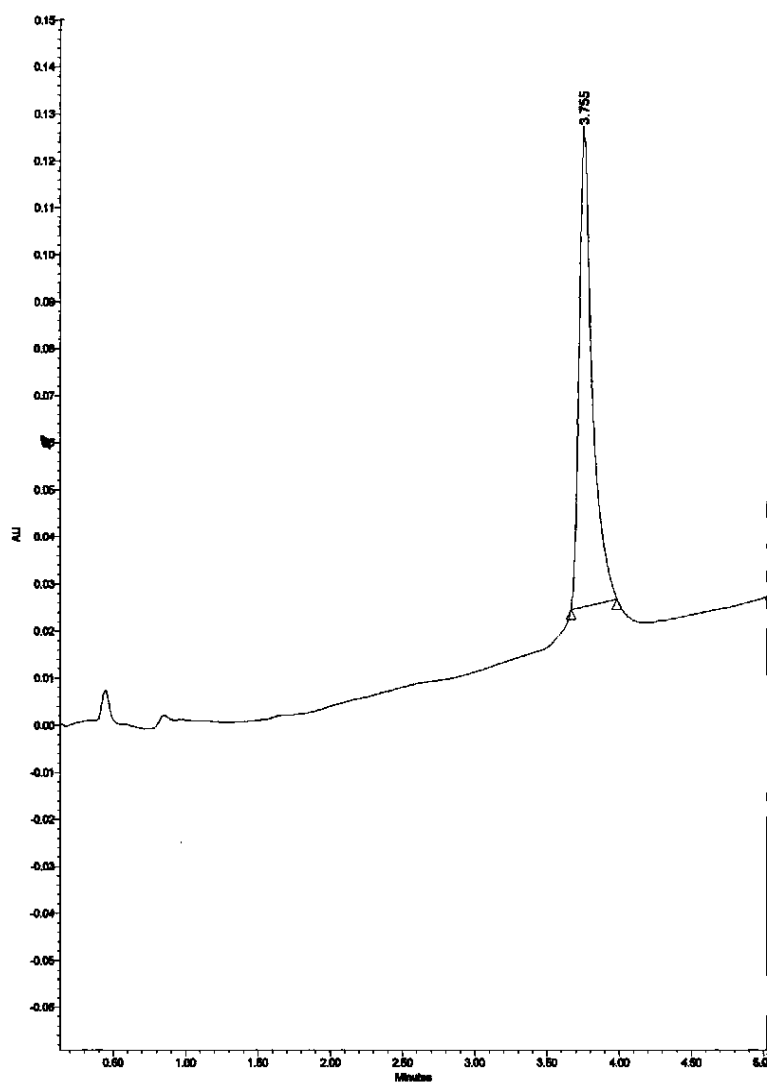


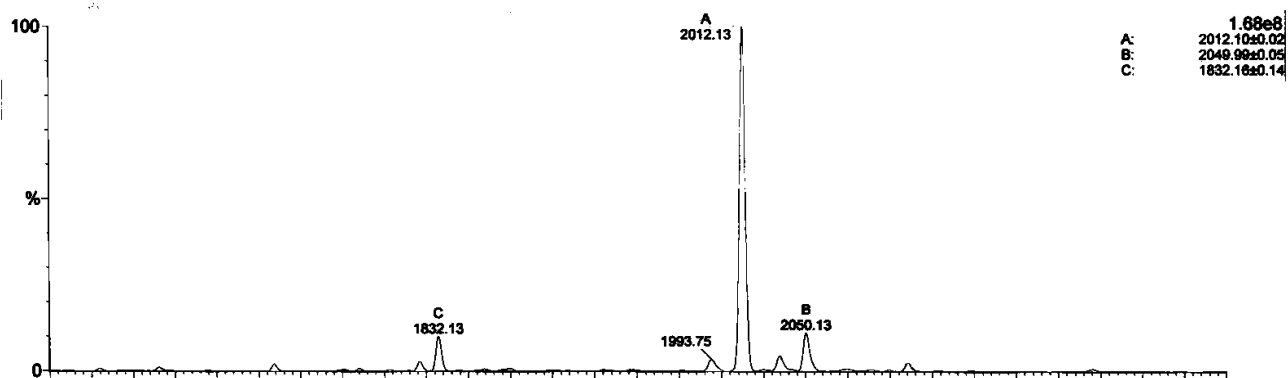
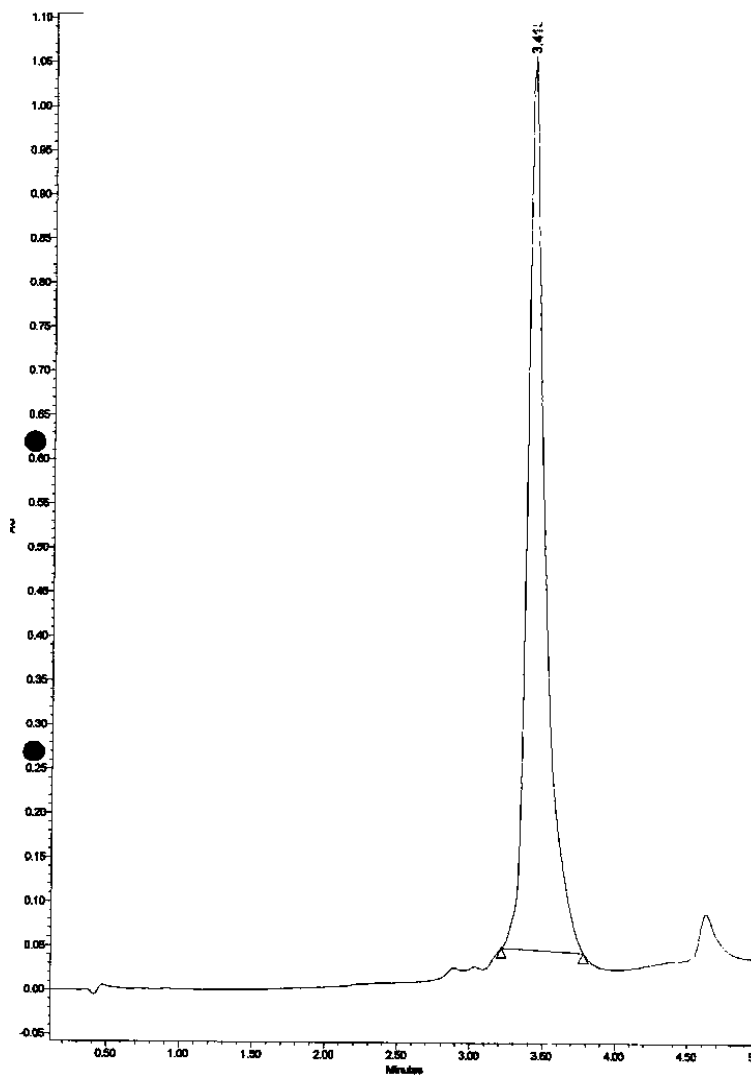


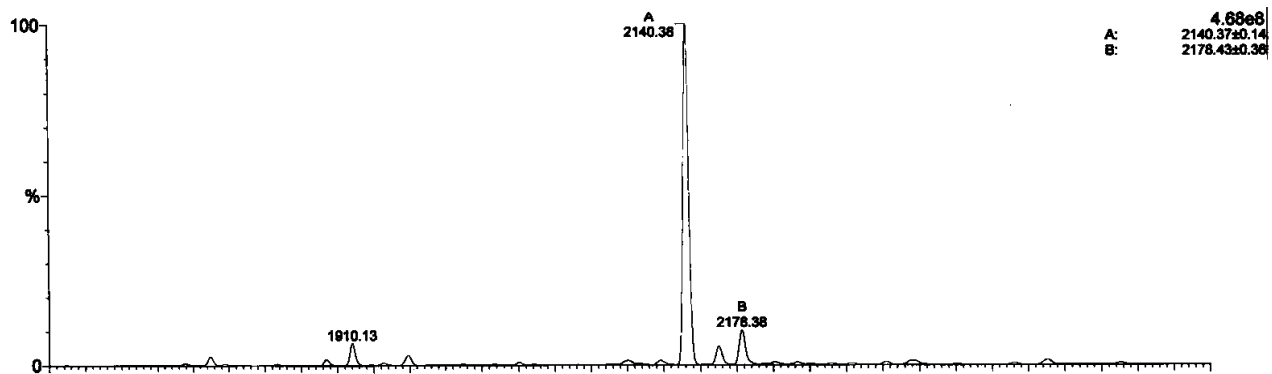
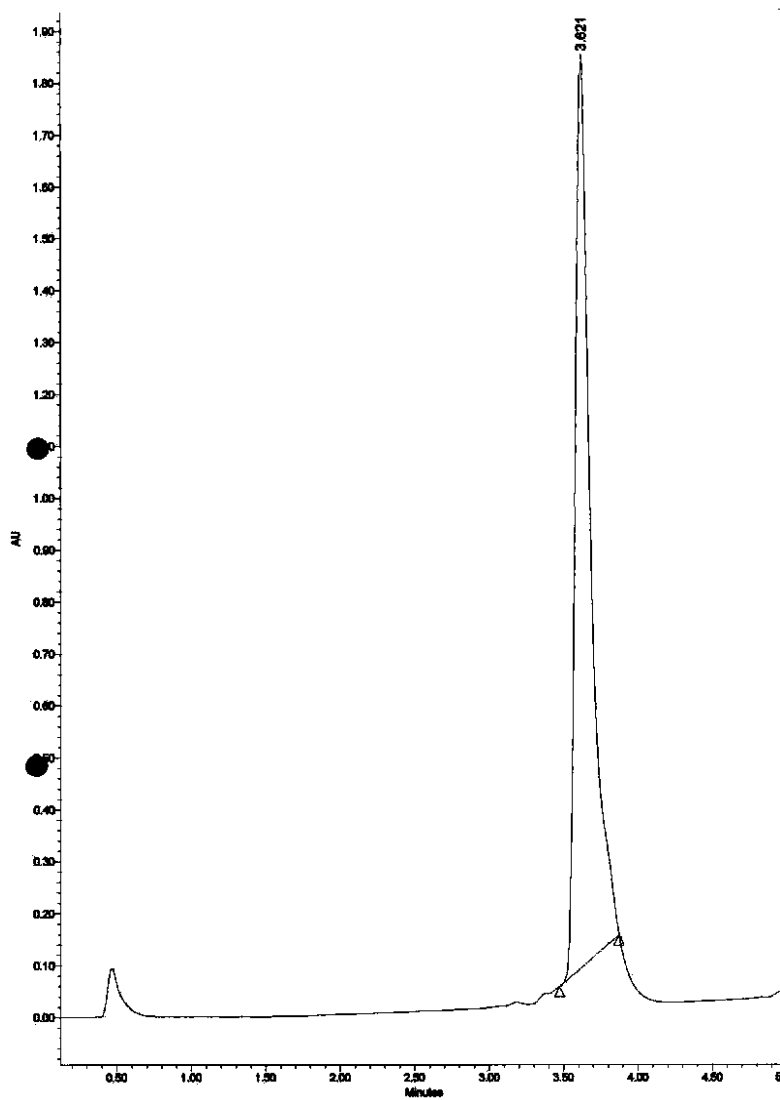
Dendrimer A<sub>1</sub>

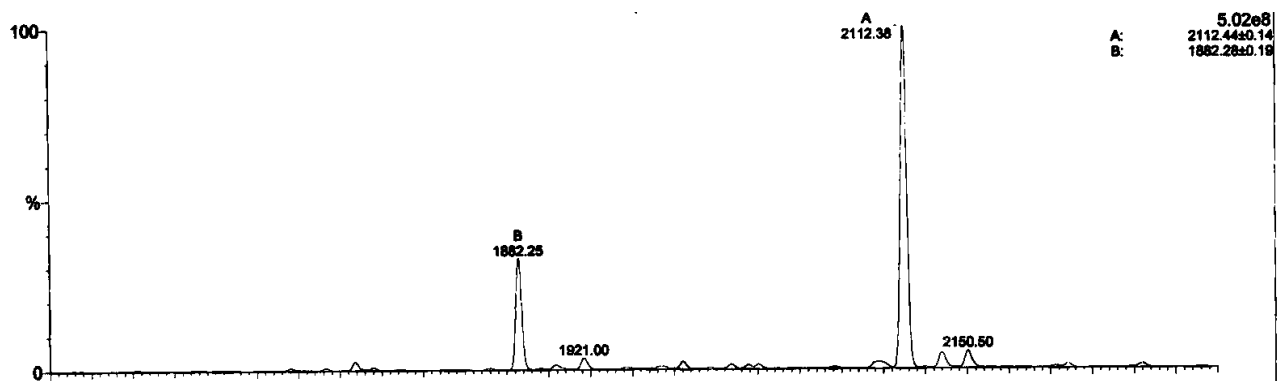
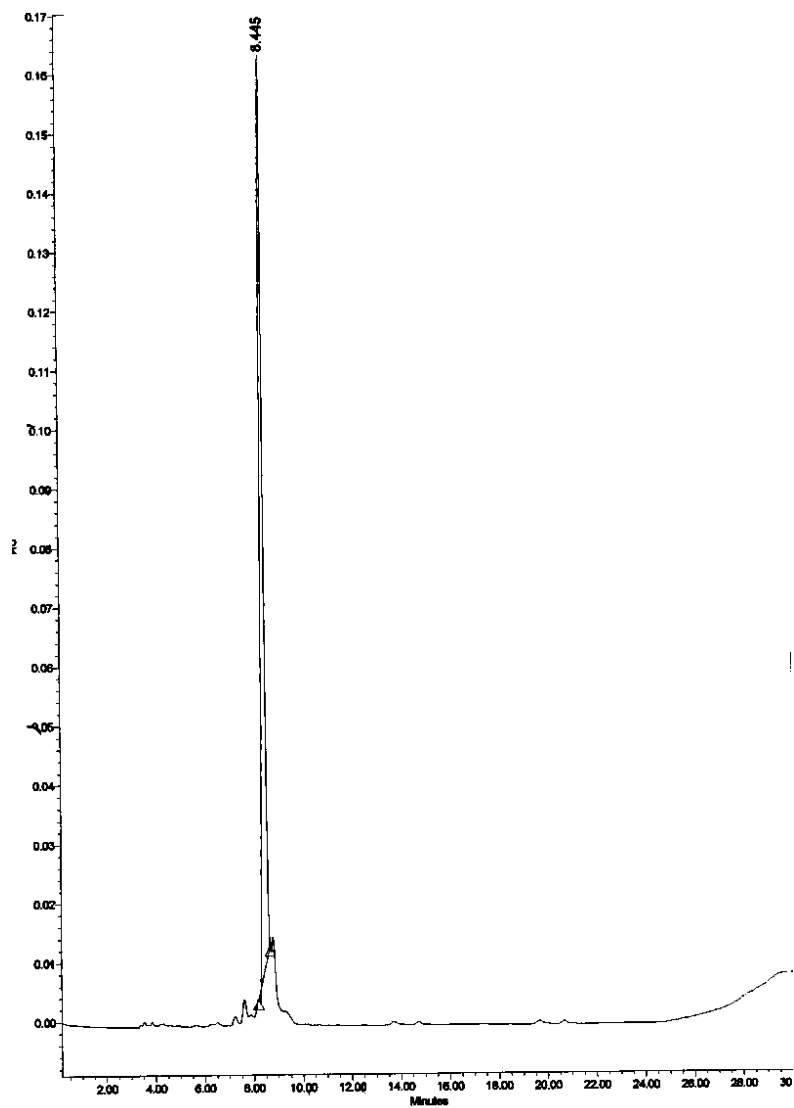
Dendrimer B<sub>1</sub>

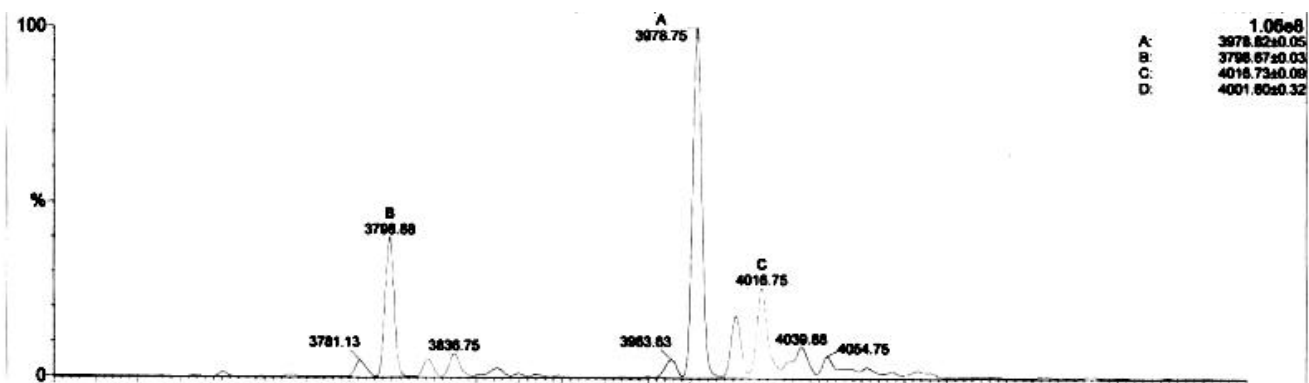
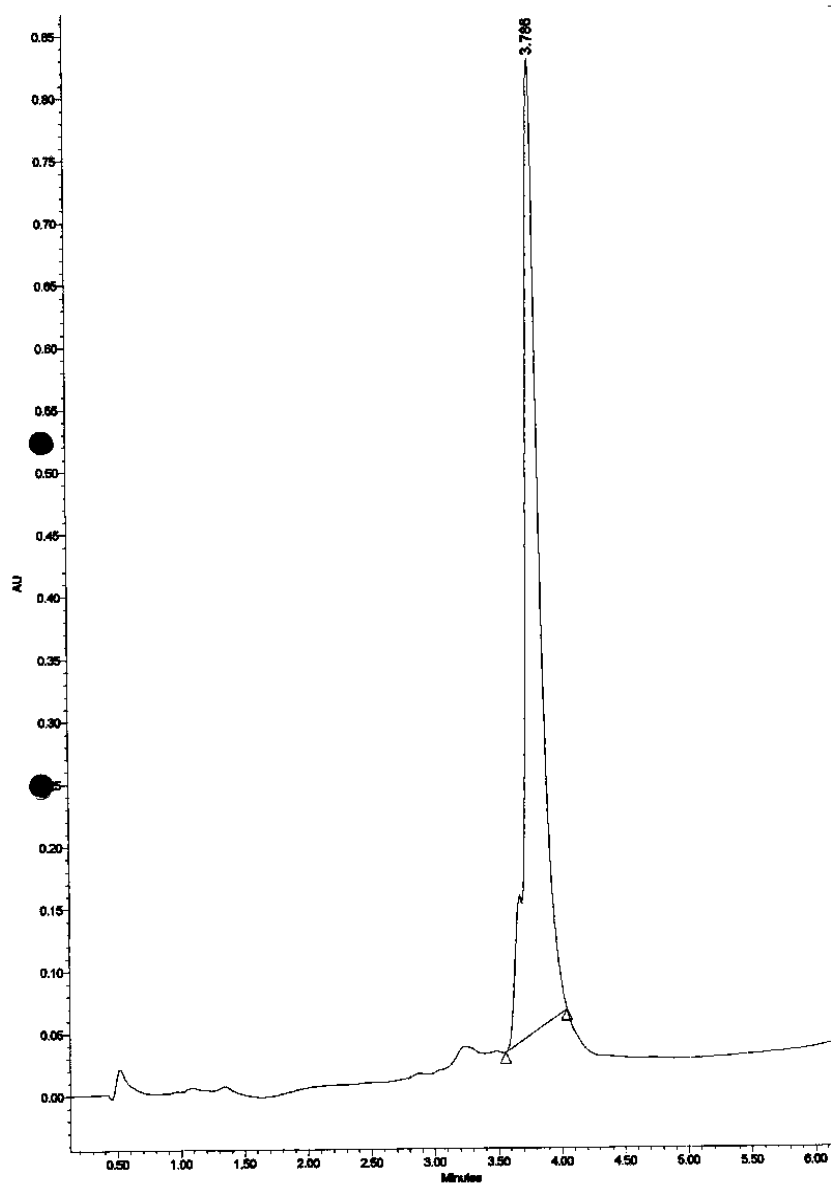


Dendrimer C<sub>1</sub>

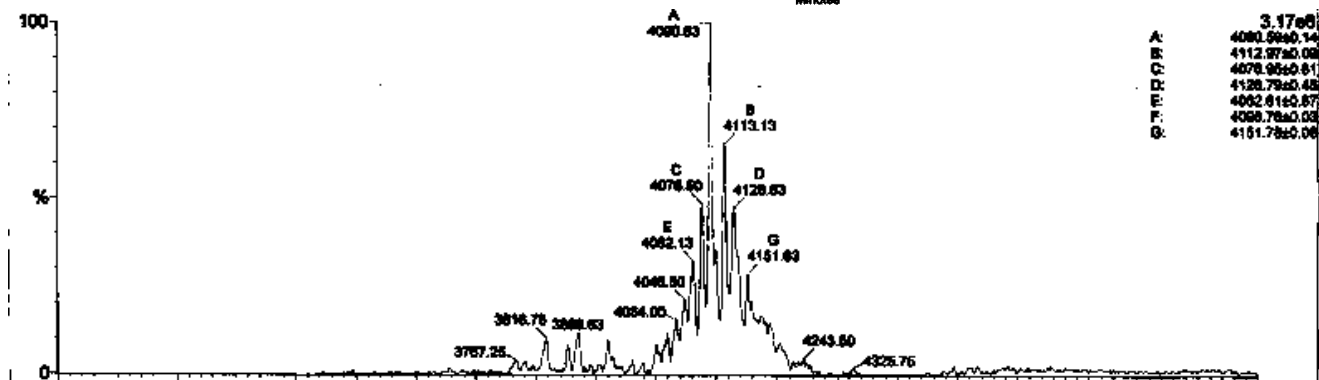
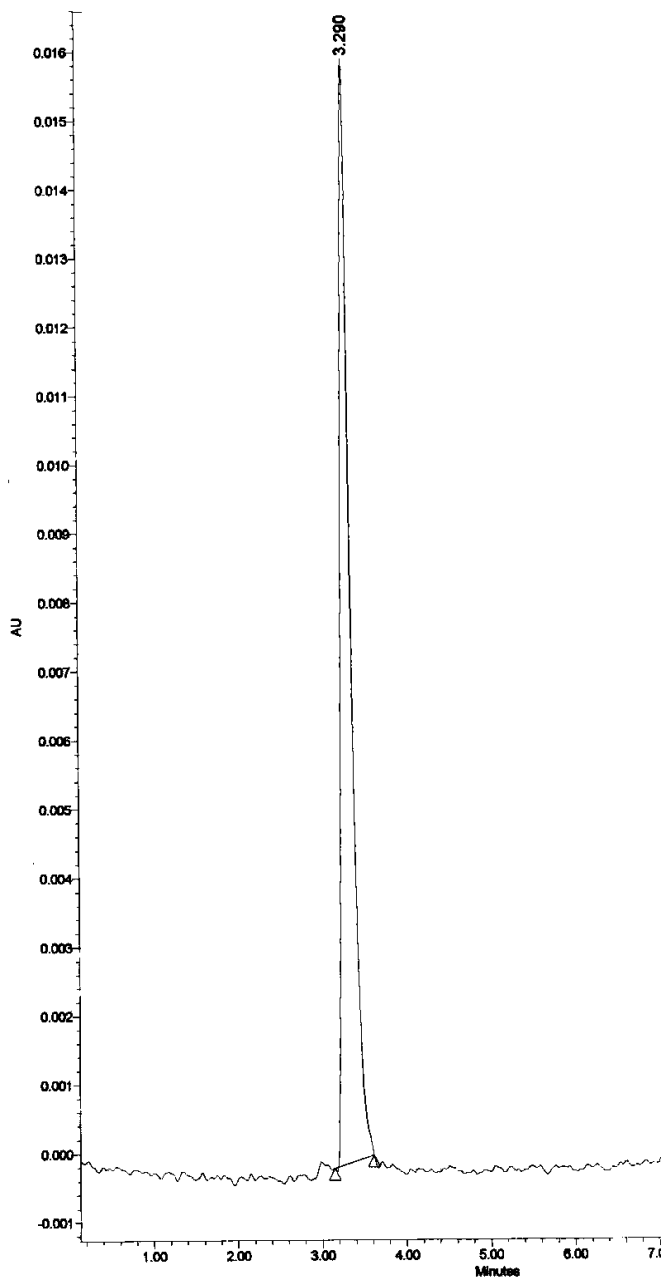
Dendrimer D<sub>1</sub>

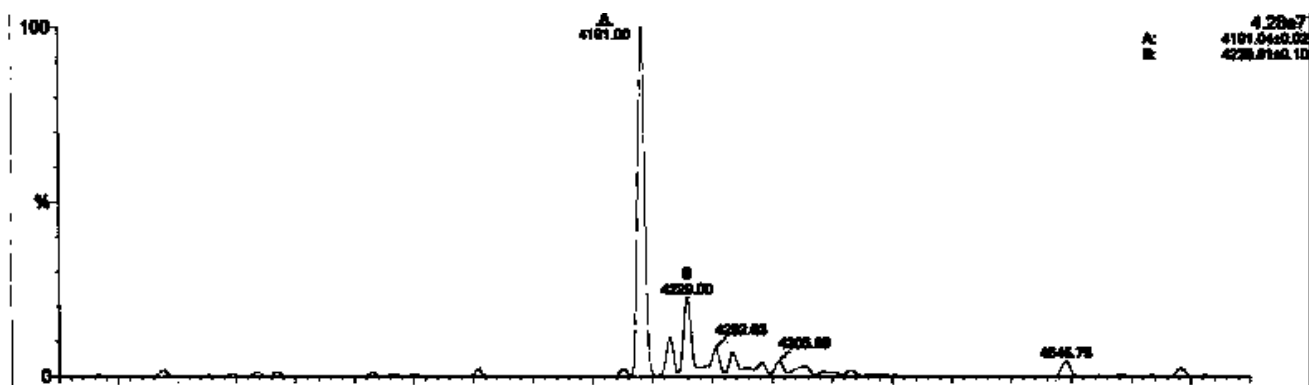
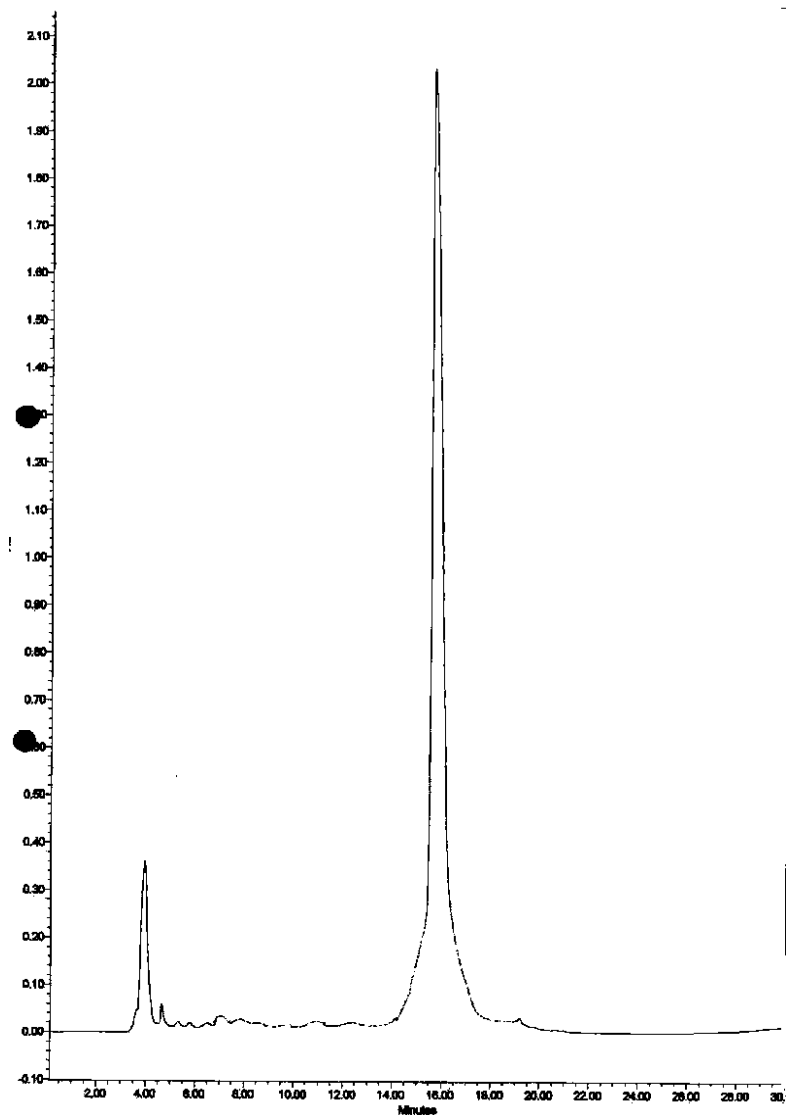
Dendrimer E<sub>1</sub>

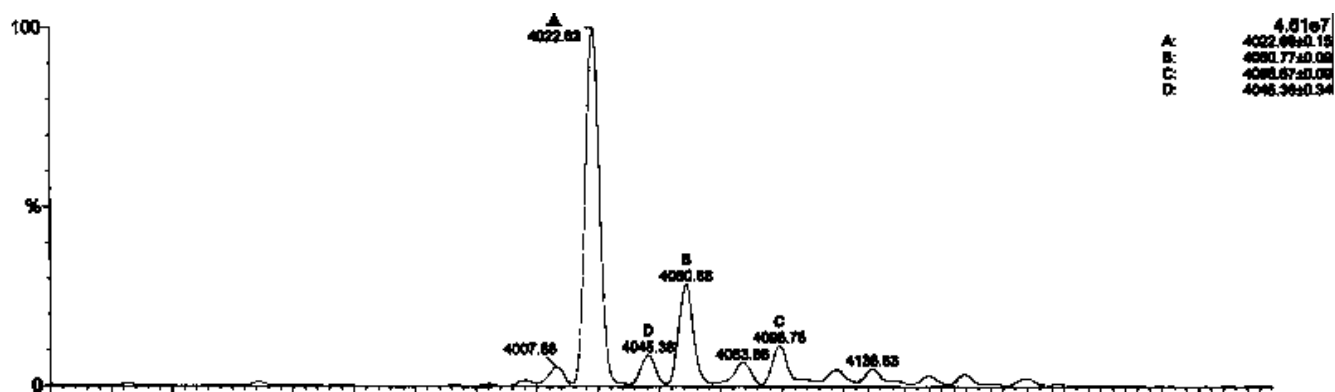
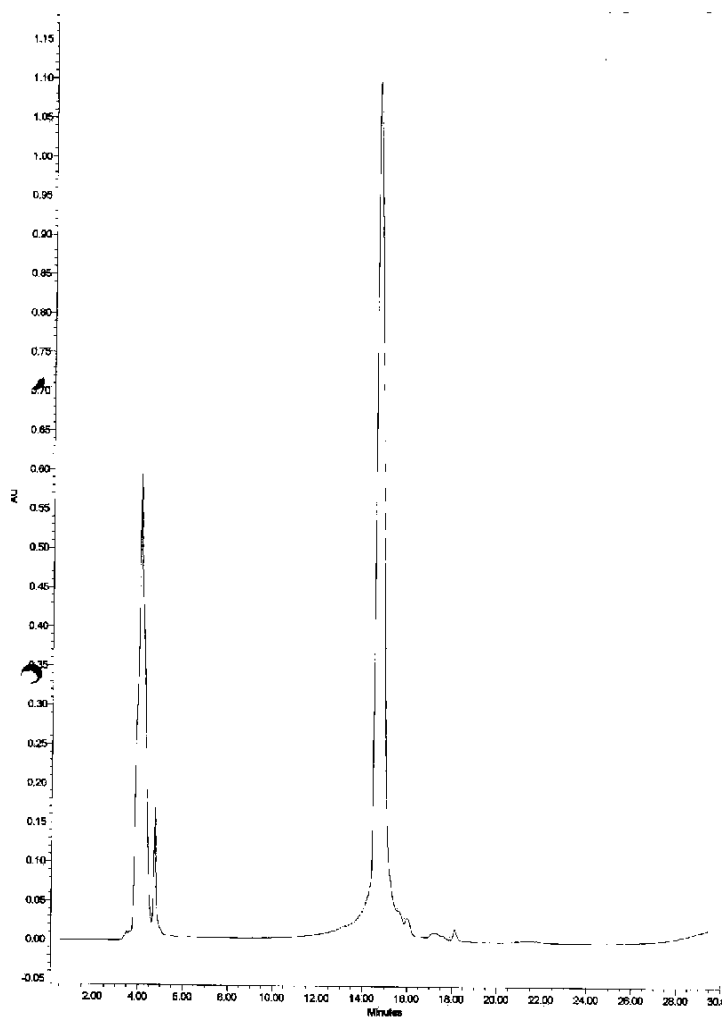
Dendrimer F<sub>1</sub>

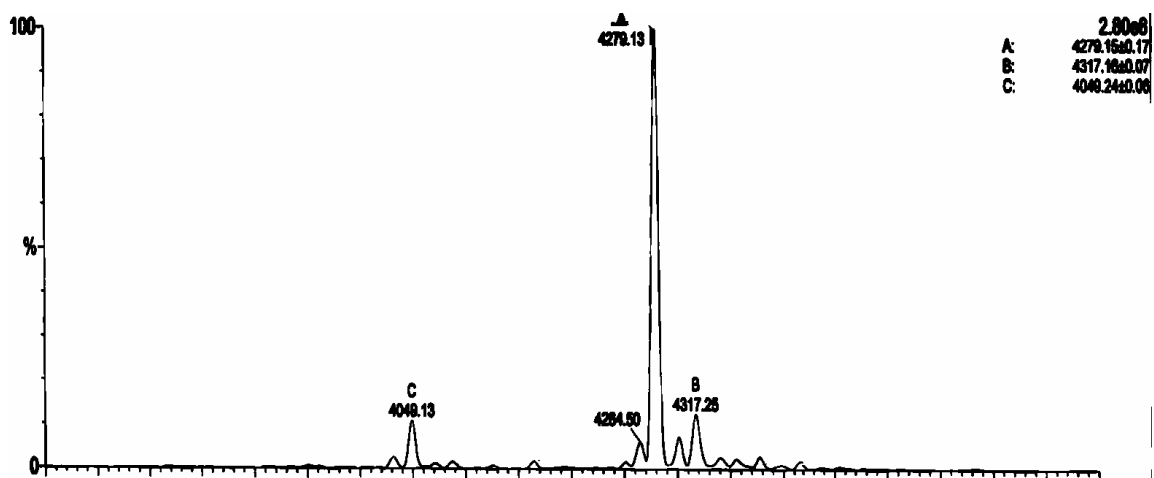
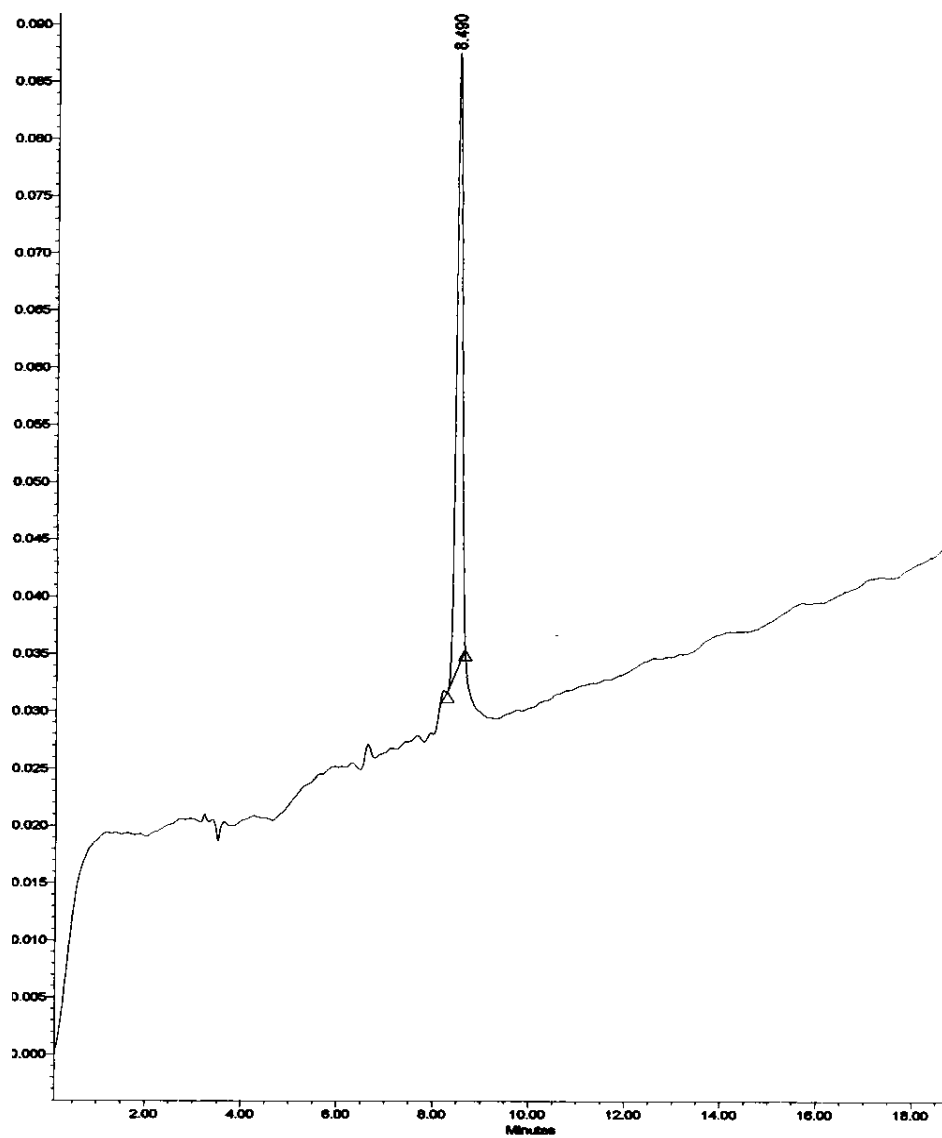
Dendrimer A<sub>1</sub>-A<sub>1</sub>

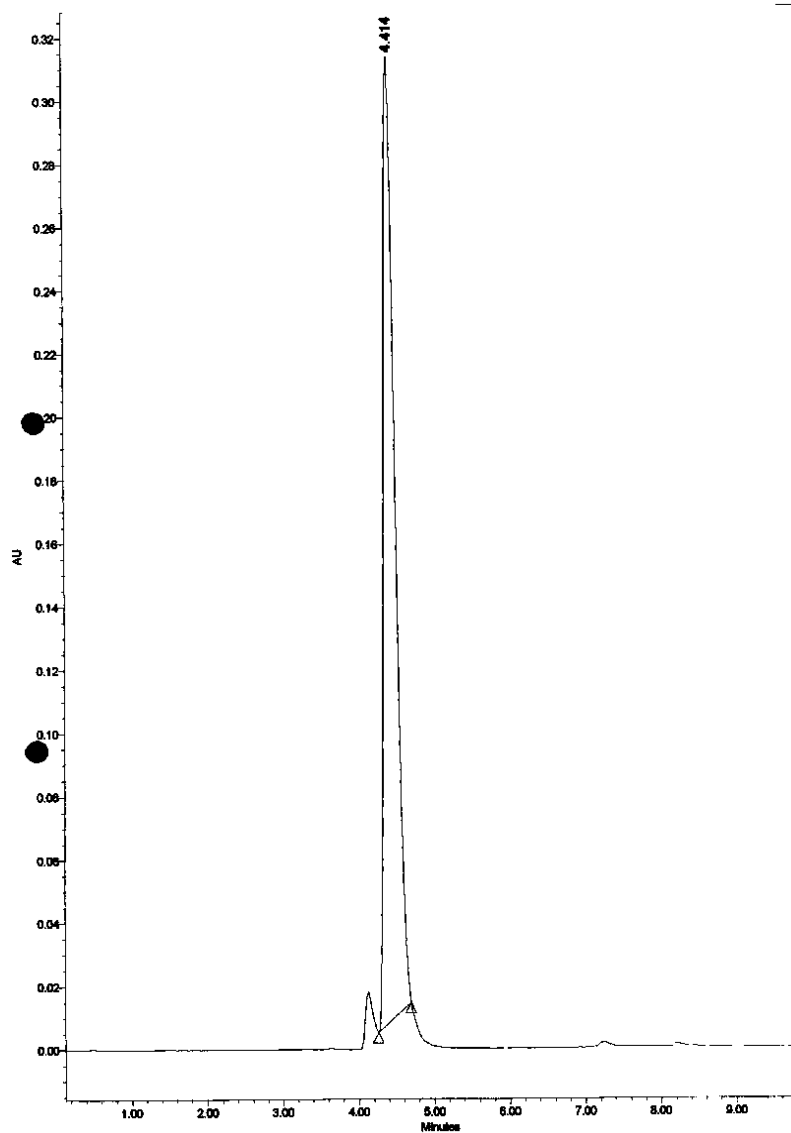


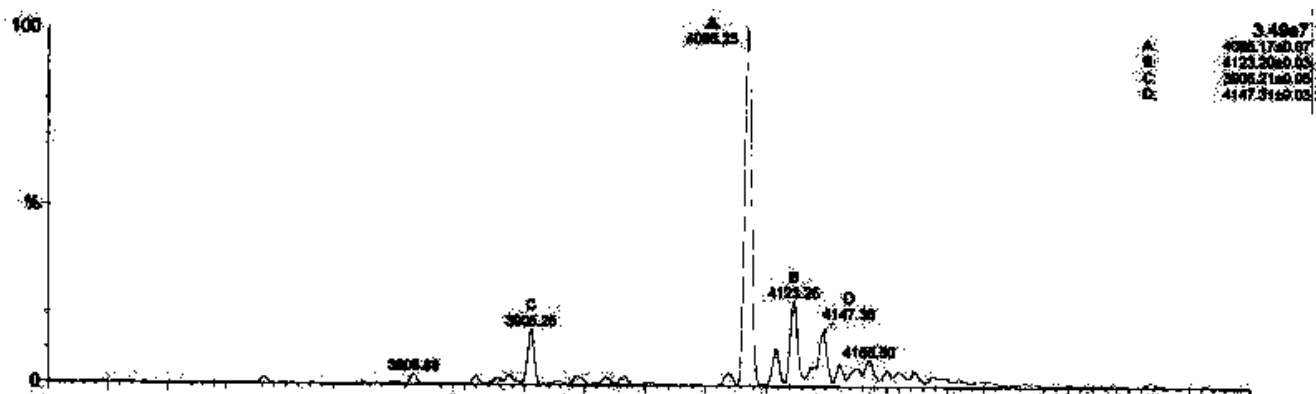
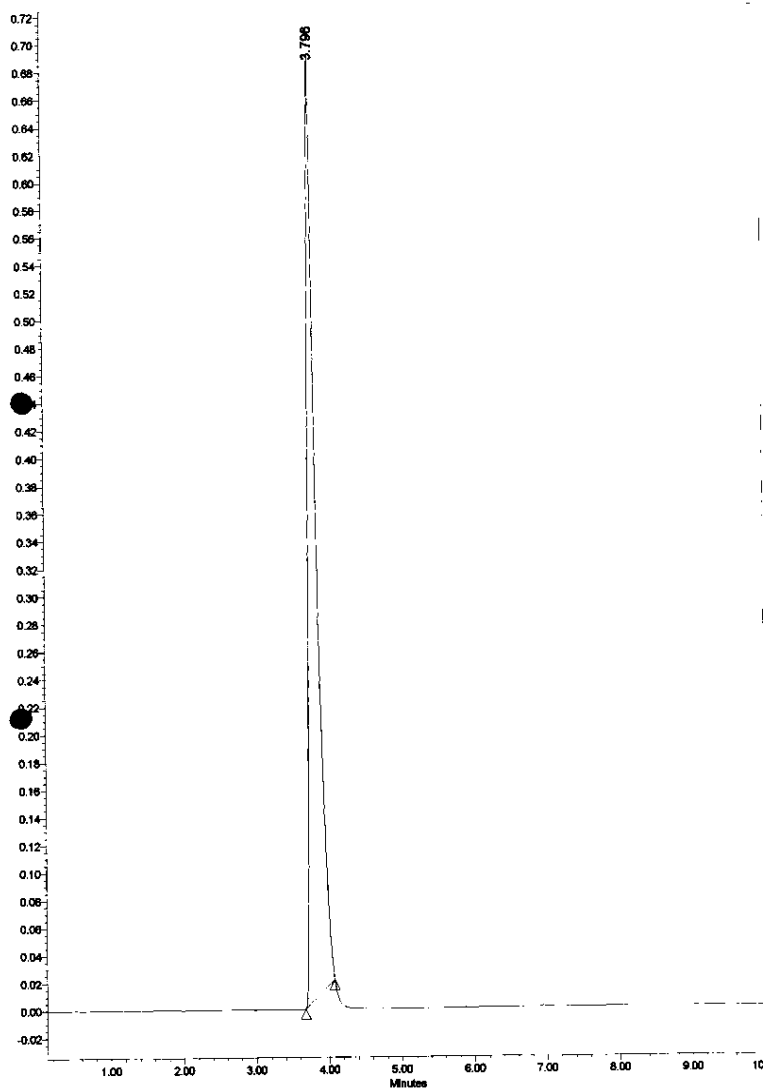
Dendrimer B<sub>1</sub>-B<sub>1</sub>

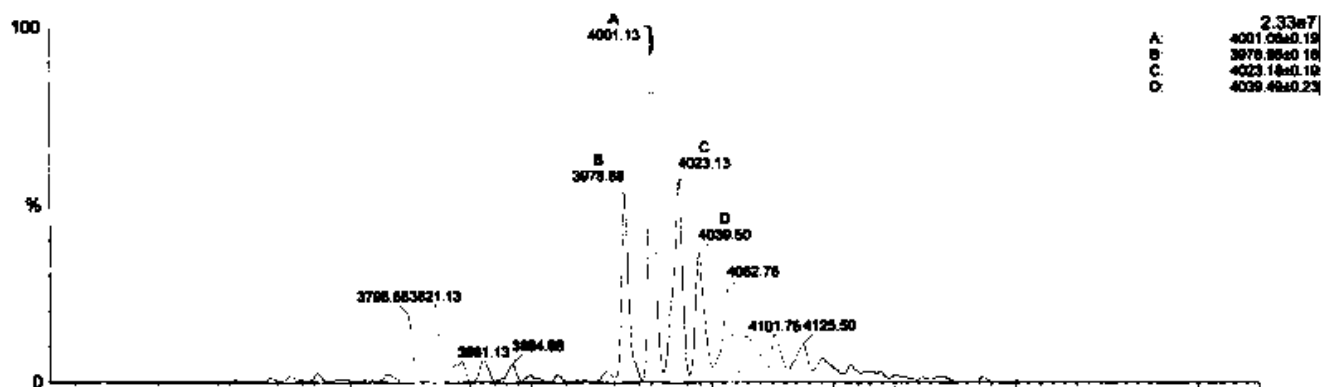
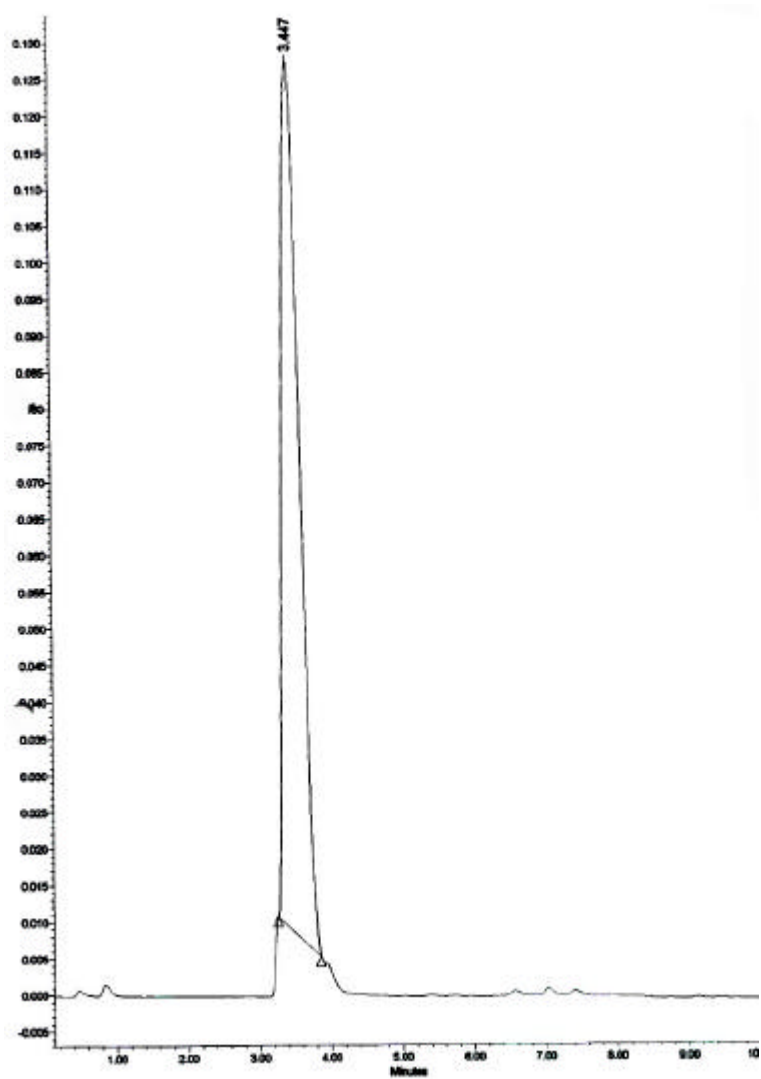
Dendrimer C<sub>1</sub>-C<sub>1</sub>

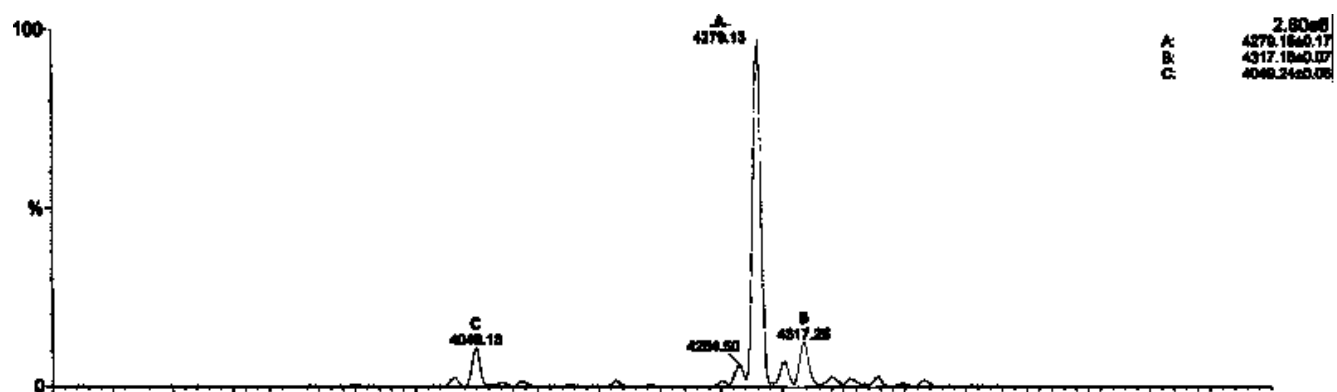
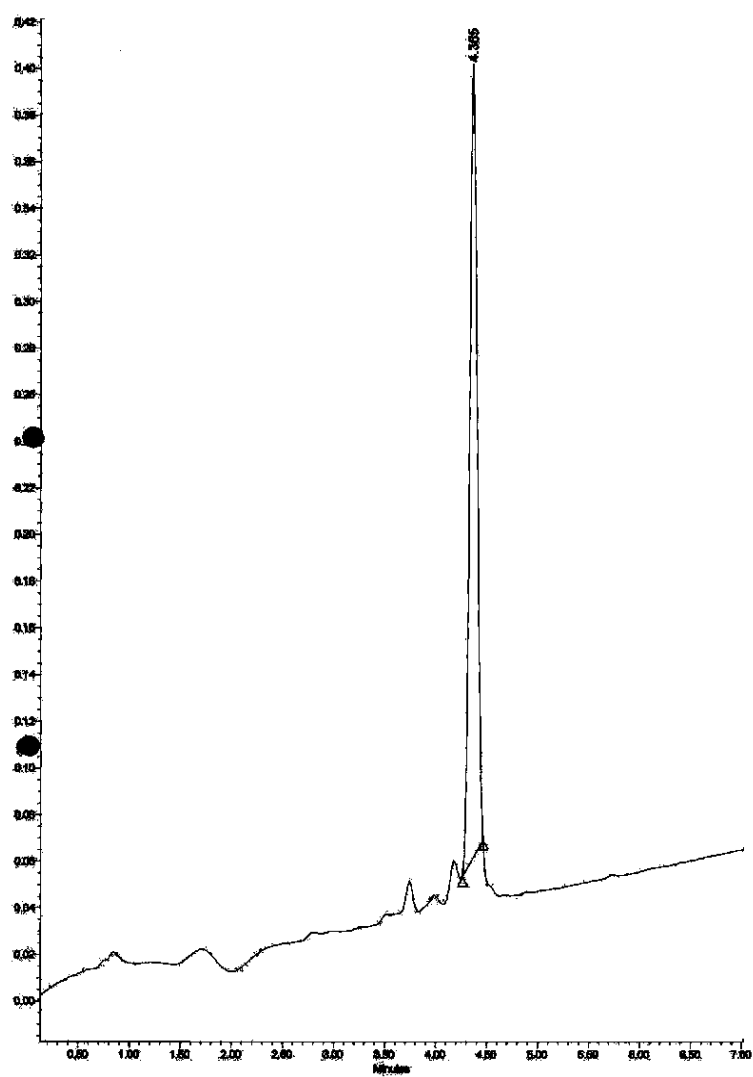
Dendrimer D<sub>1</sub>-D<sub>1</sub>

Dendrimer E<sub>1</sub>-E<sub>1</sub>

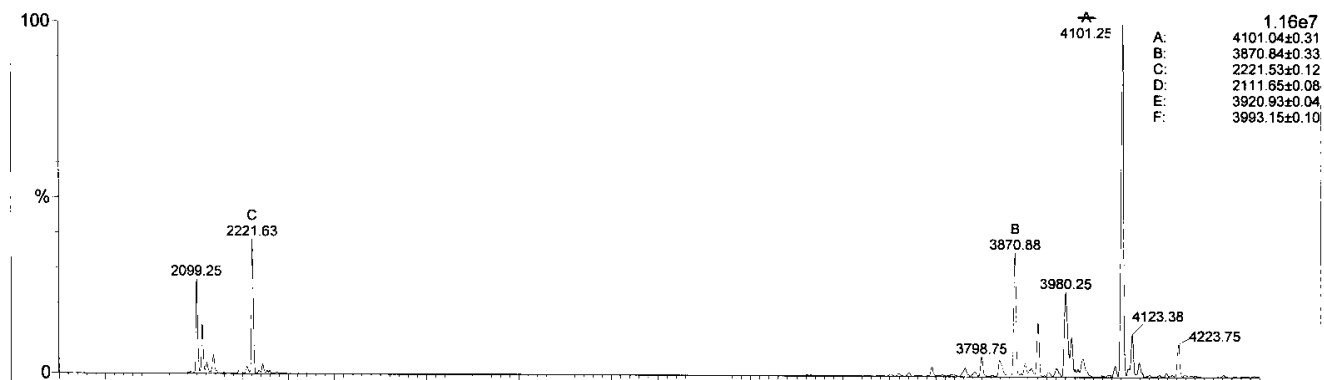
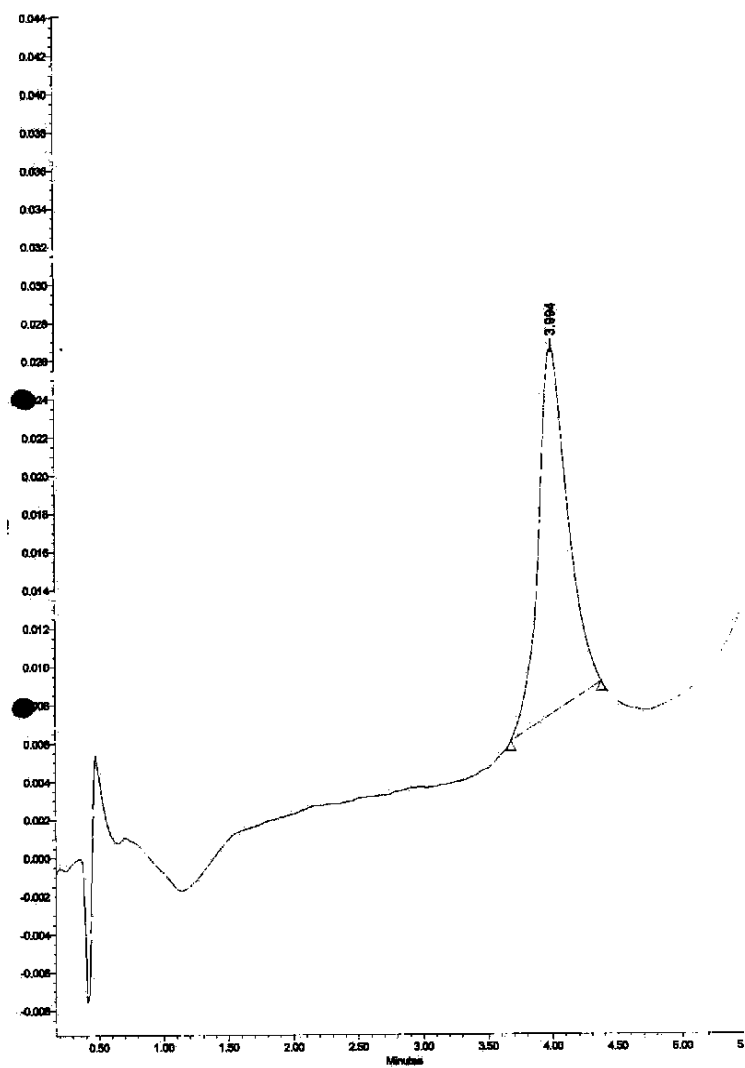
Dendrimer A<sub>1</sub>-B<sub>1</sub>

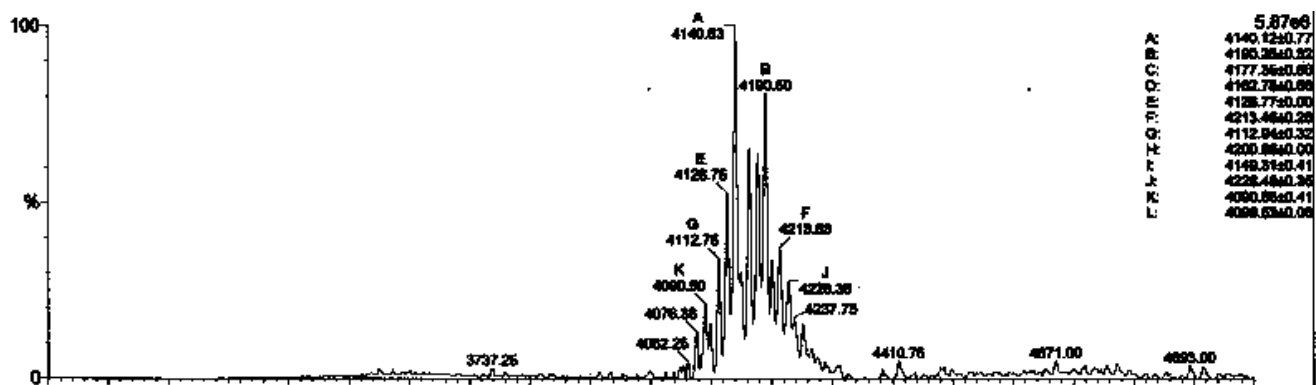
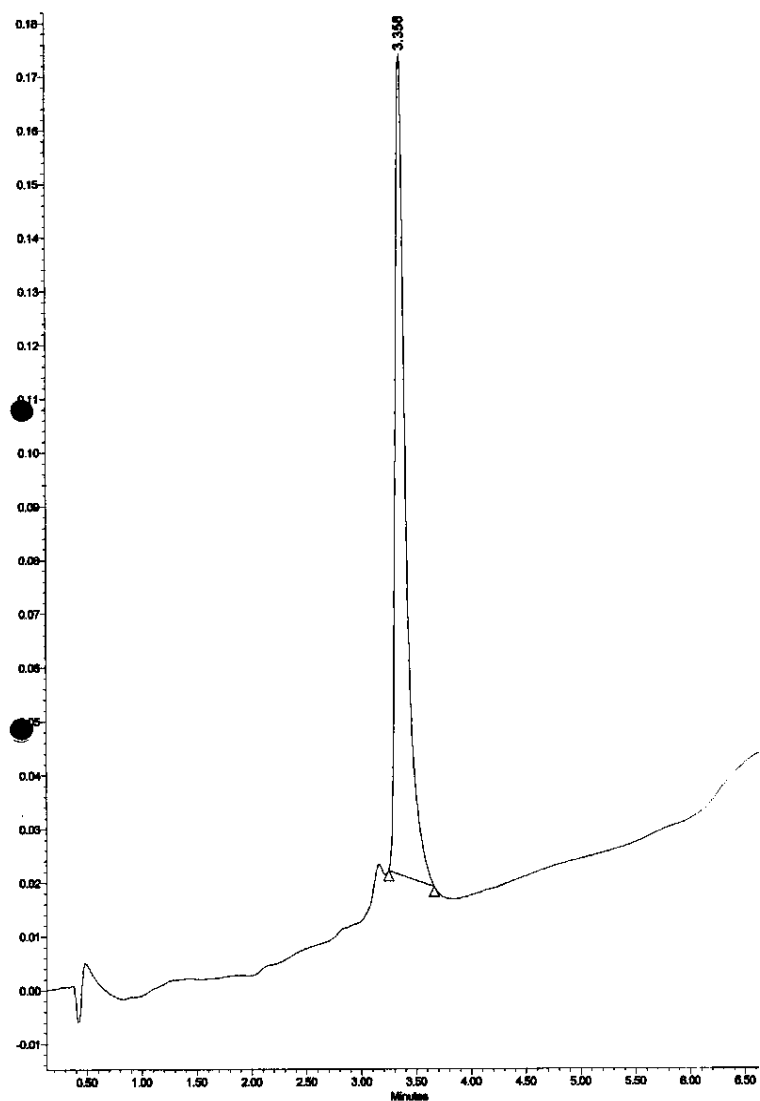
Dendrimer A<sub>1</sub>-C<sub>1</sub>

Dendrimer A<sub>1</sub>-D<sub>1</sub>

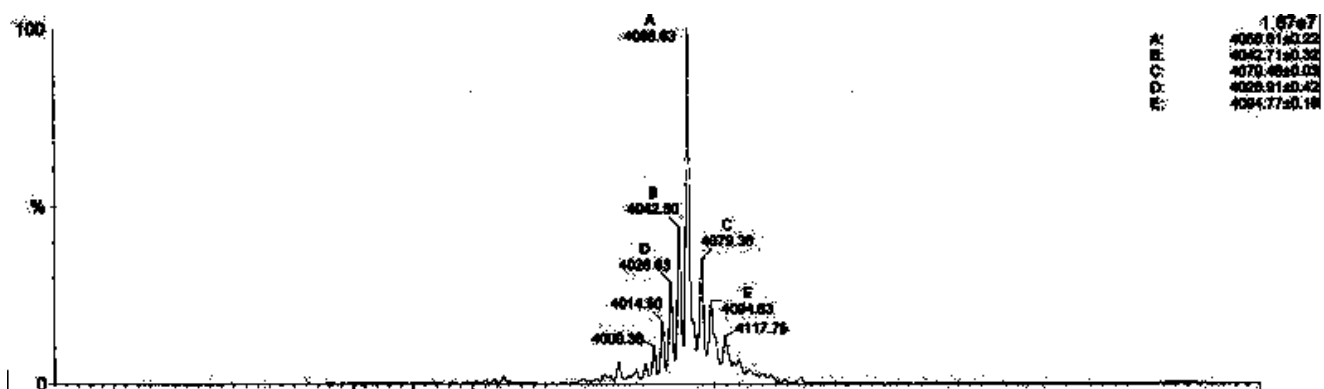
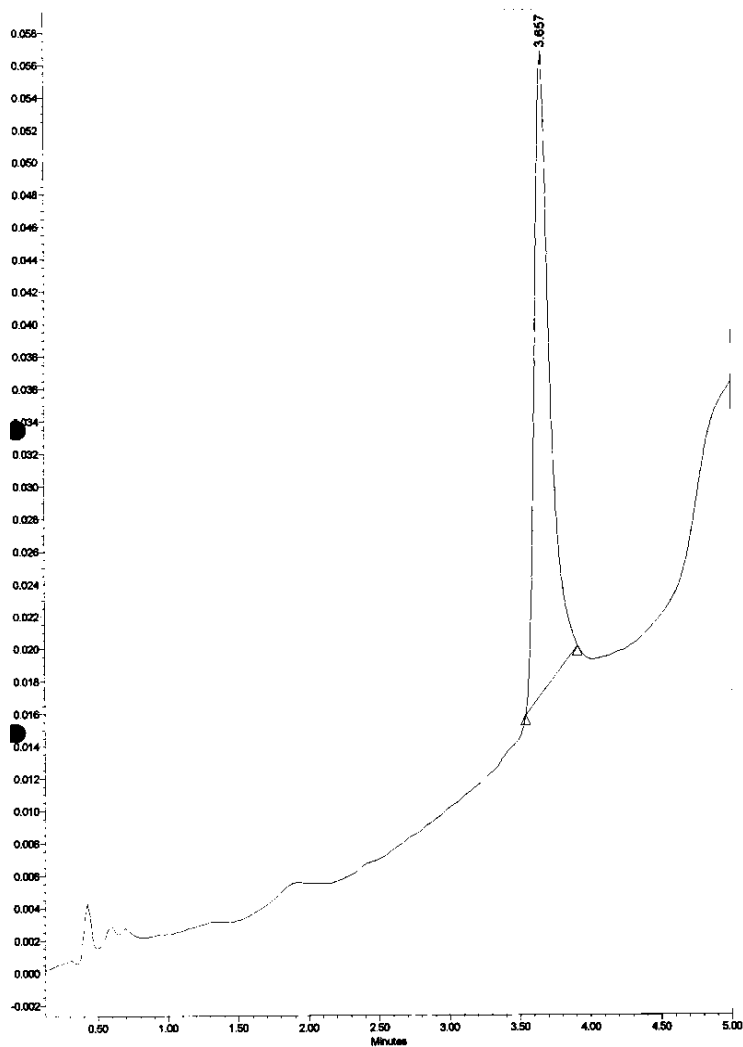
Dendrimer A<sub>1</sub>-E<sub>1</sub>

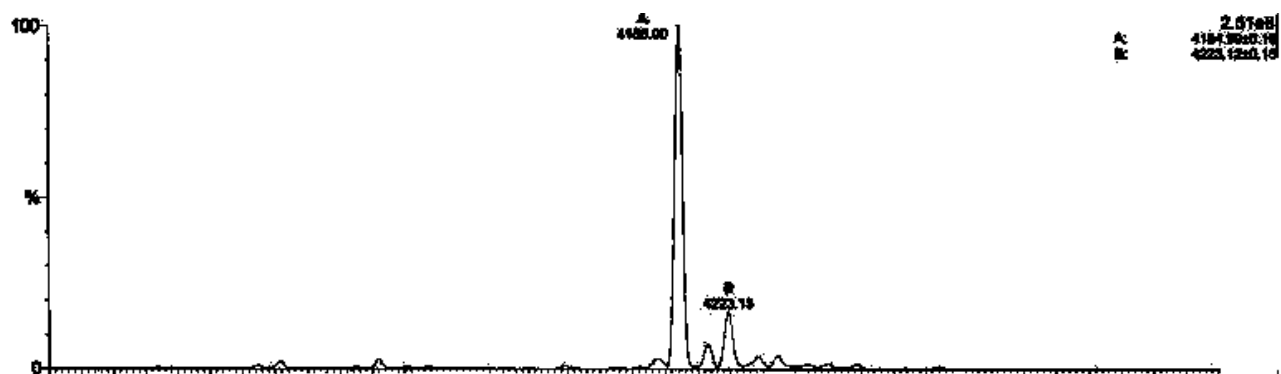
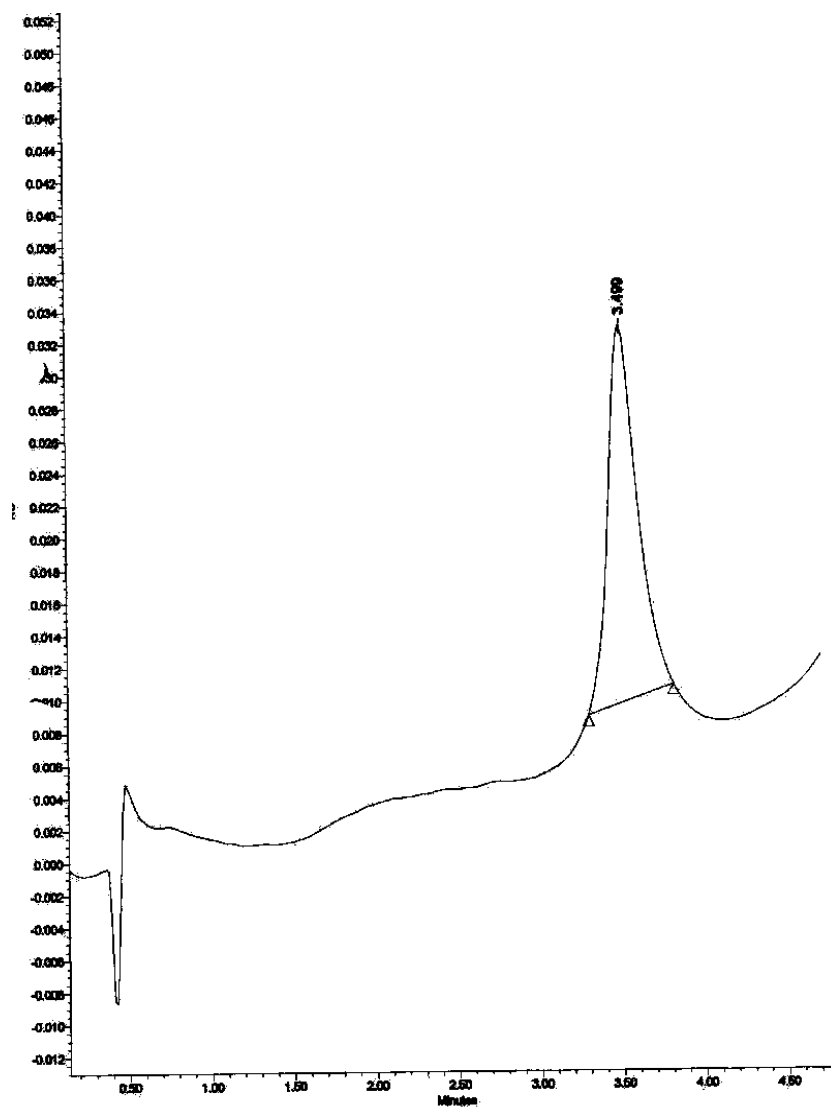


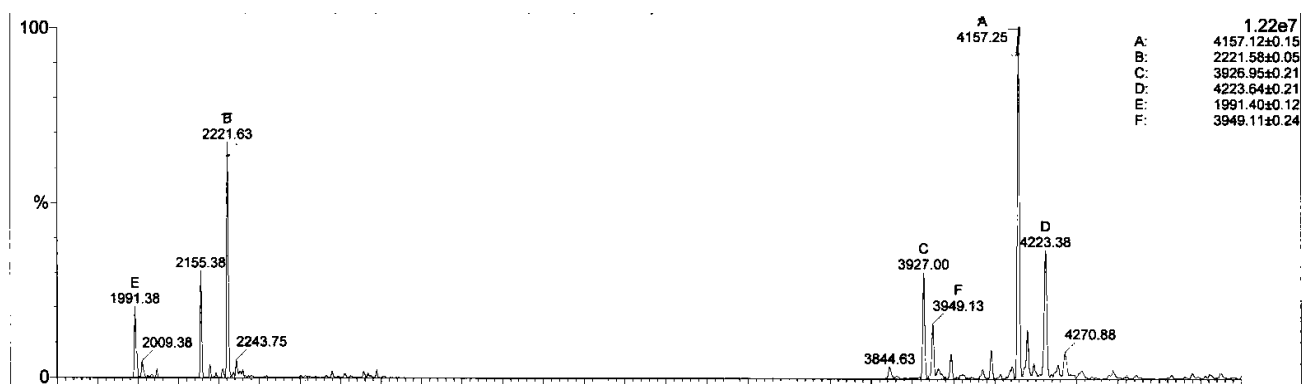
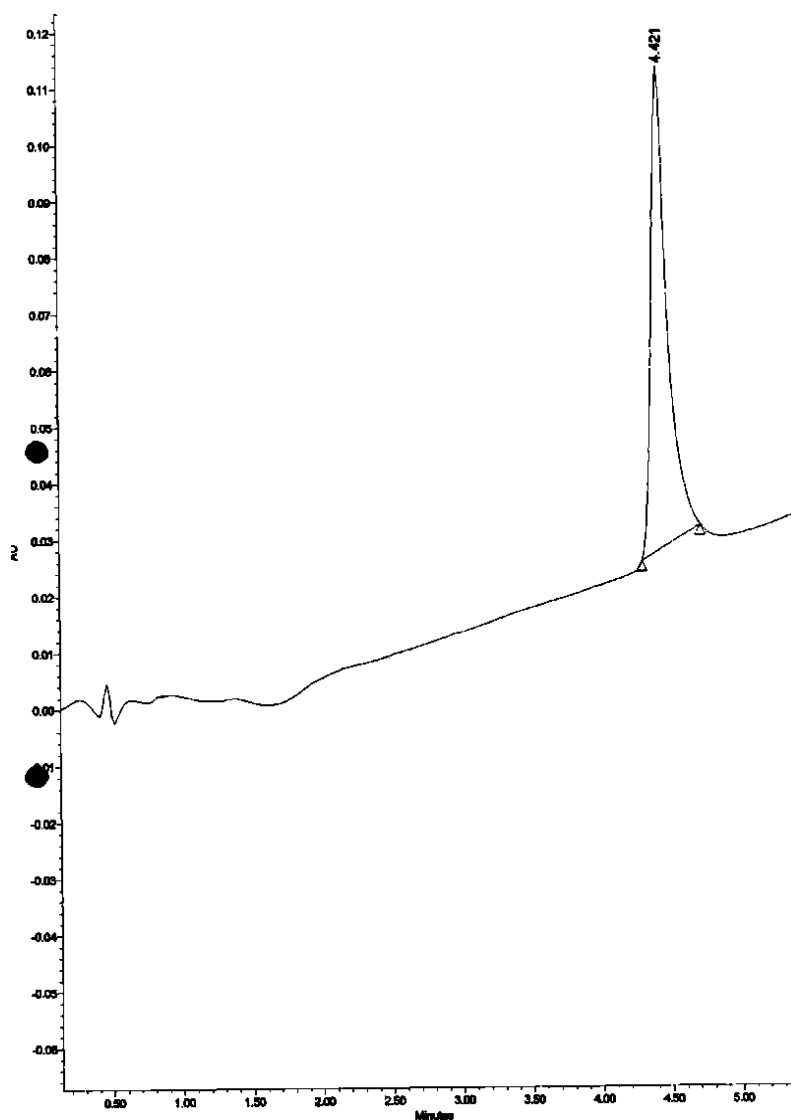
Dendrimer A<sub>1</sub>-F<sub>1</sub>

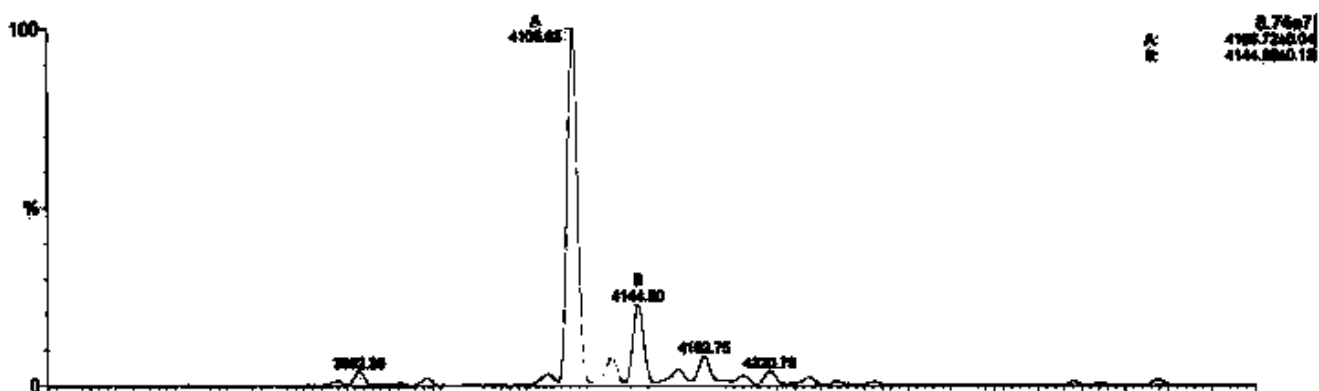
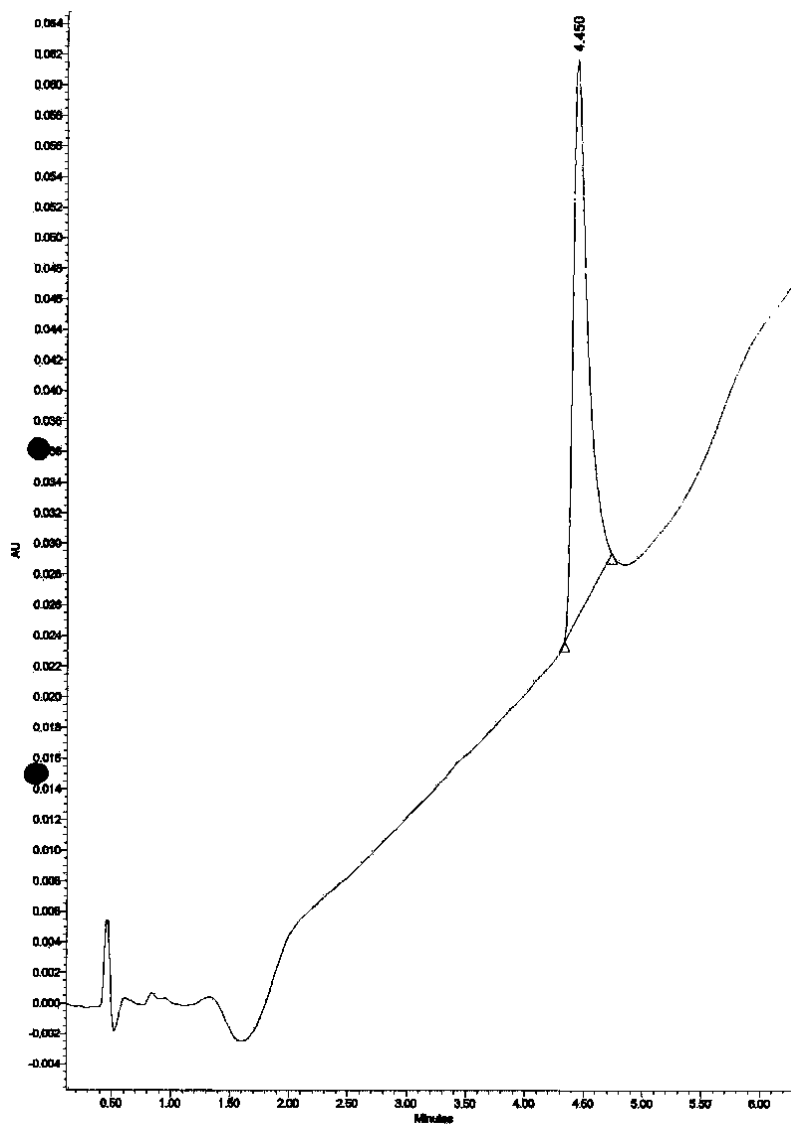
Dendrimer B<sub>1</sub>-C<sub>1</sub>

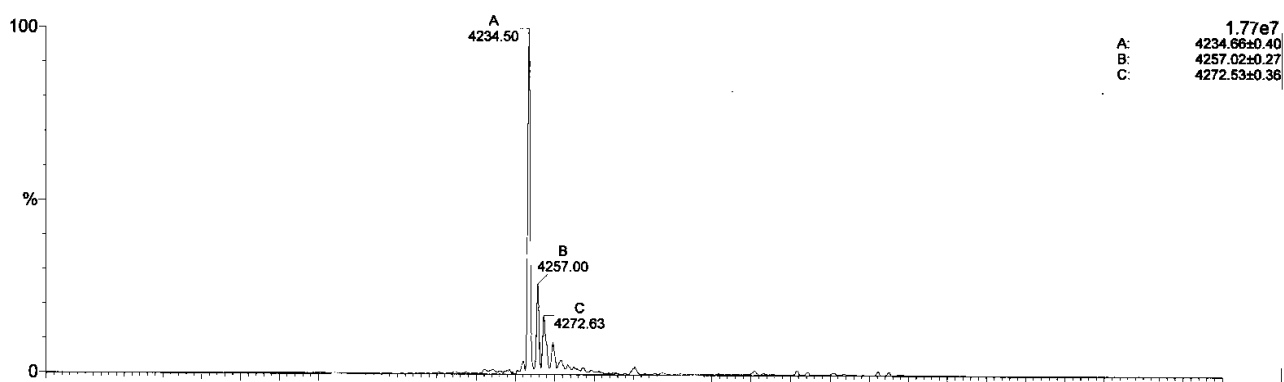
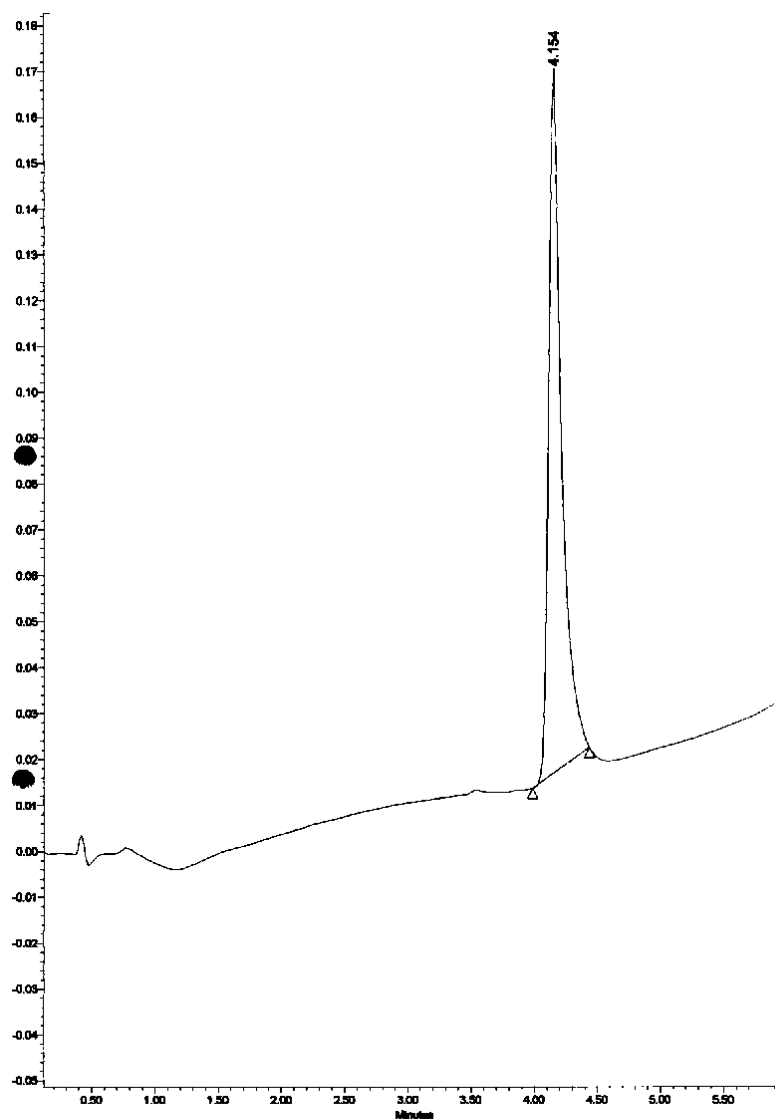
Dendrimer B<sub>1</sub>-D<sub>1</sub>

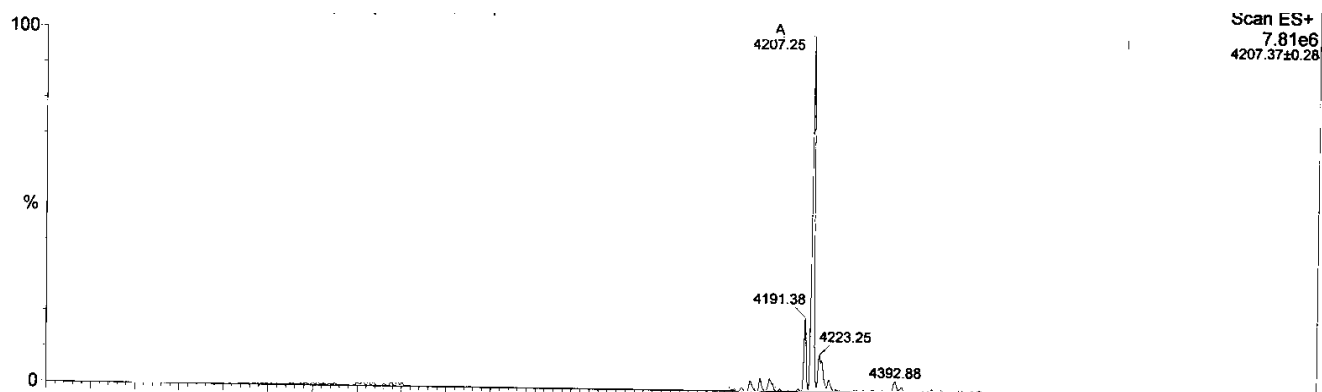
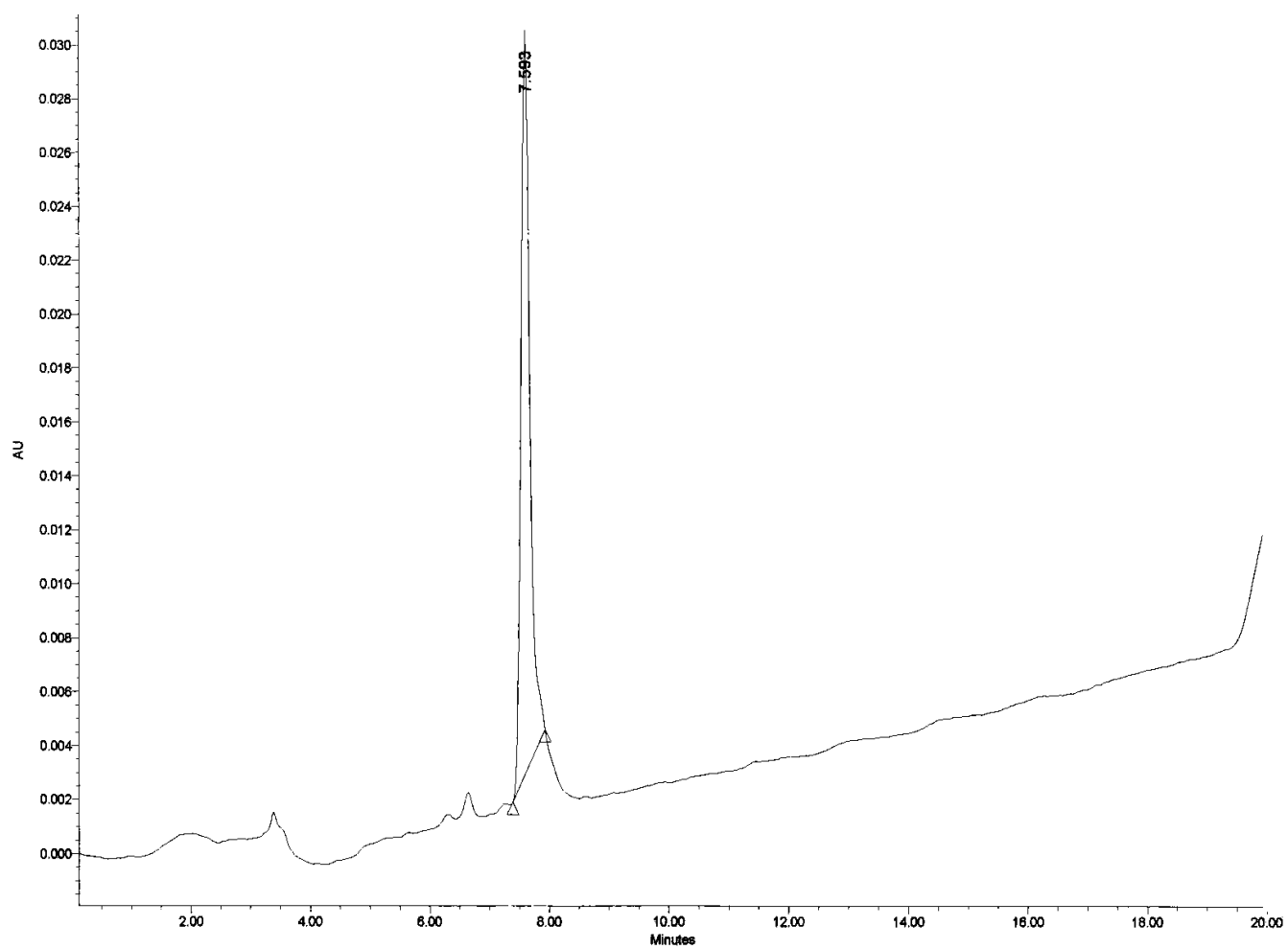


Dendrimer B<sub>1</sub>-E<sub>1</sub>

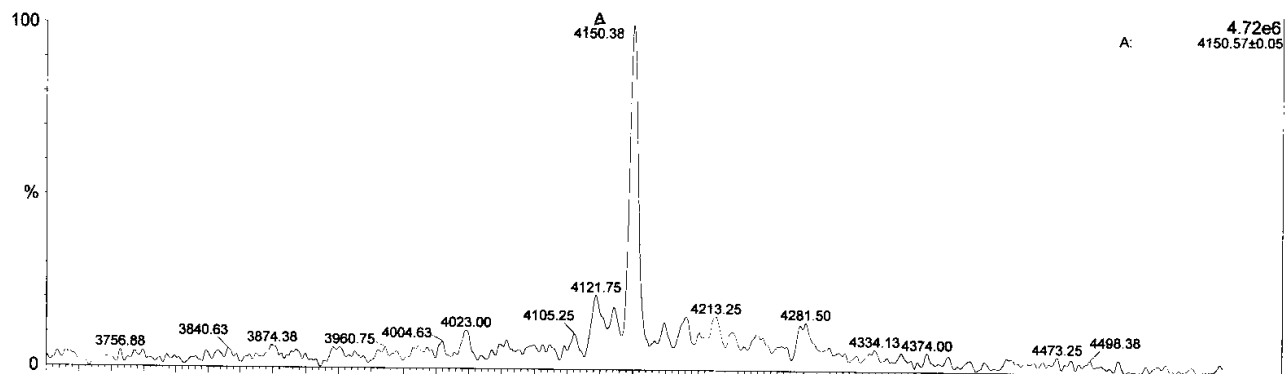
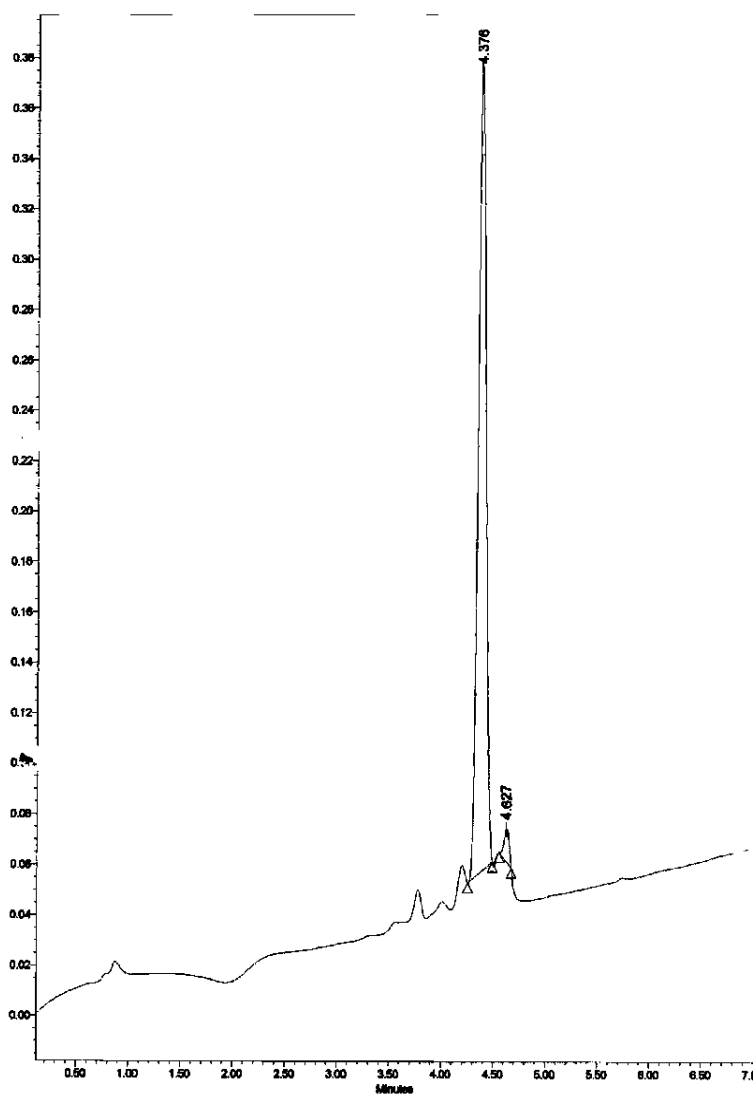
Dendrimer B<sub>1</sub>-F<sub>1</sub>

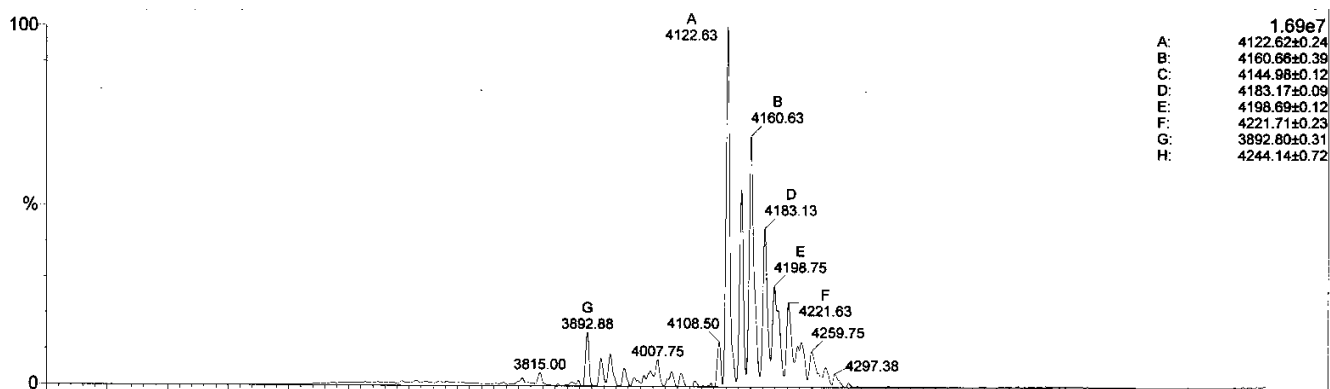
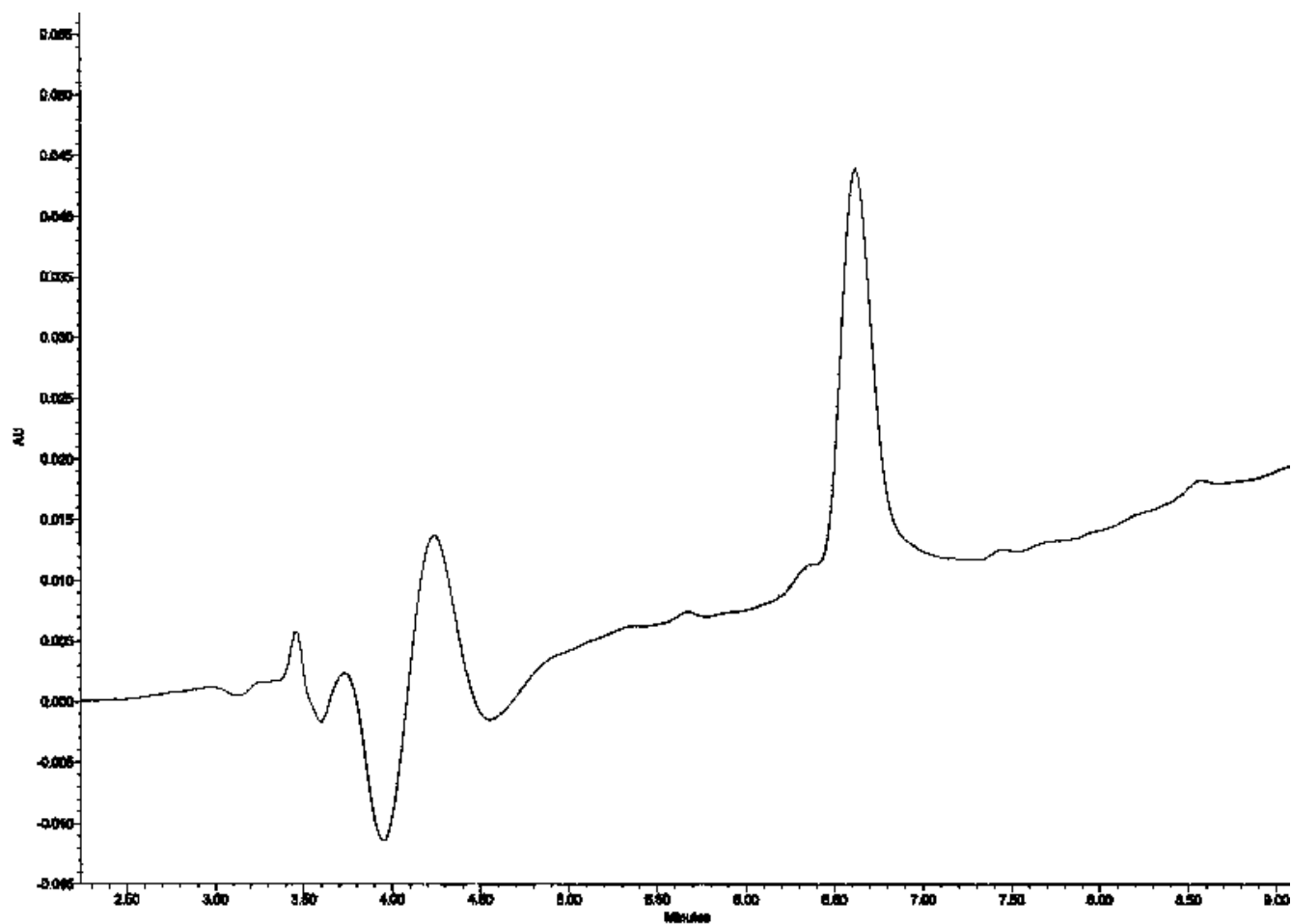
Dendrimer C<sub>1</sub>-D<sub>1</sub>

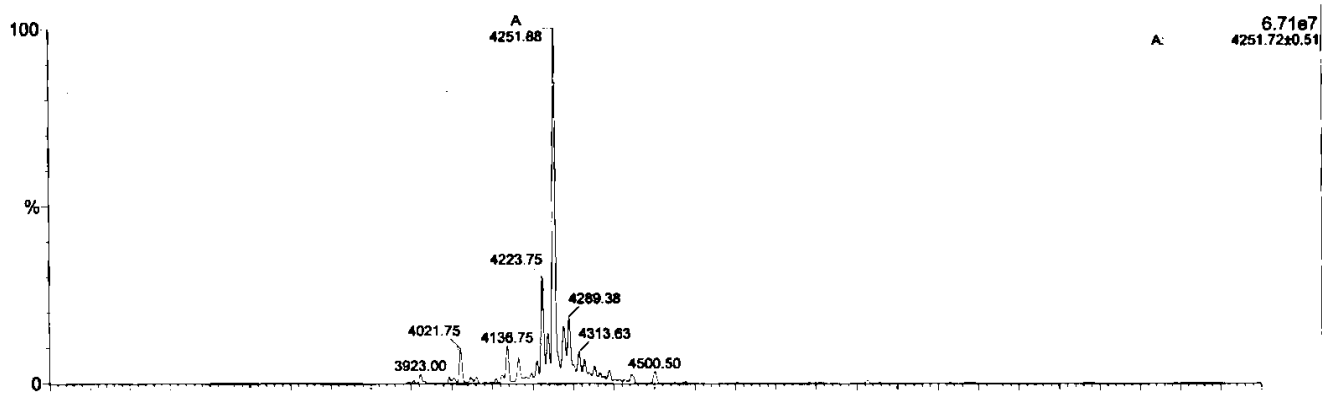
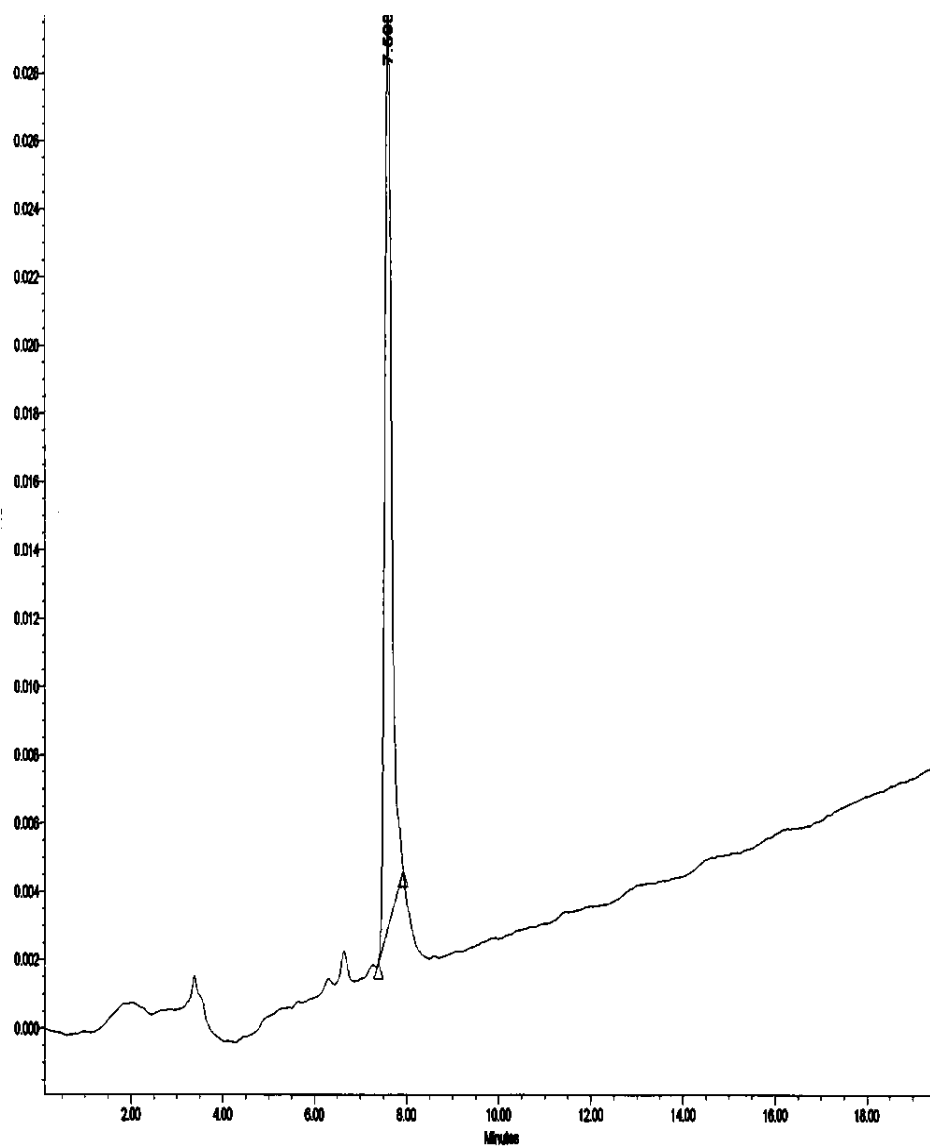
Dendrimer C<sub>1</sub>-E<sub>1</sub>

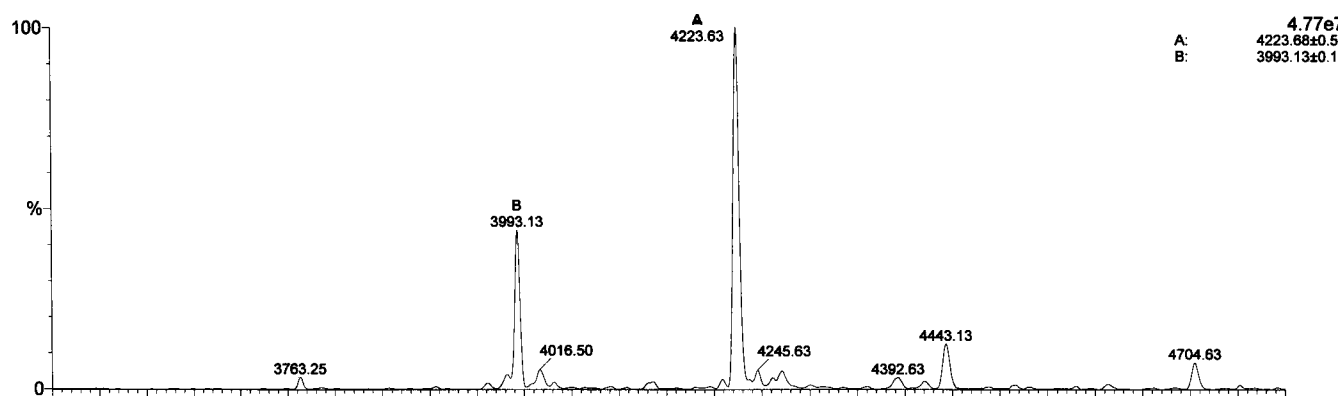
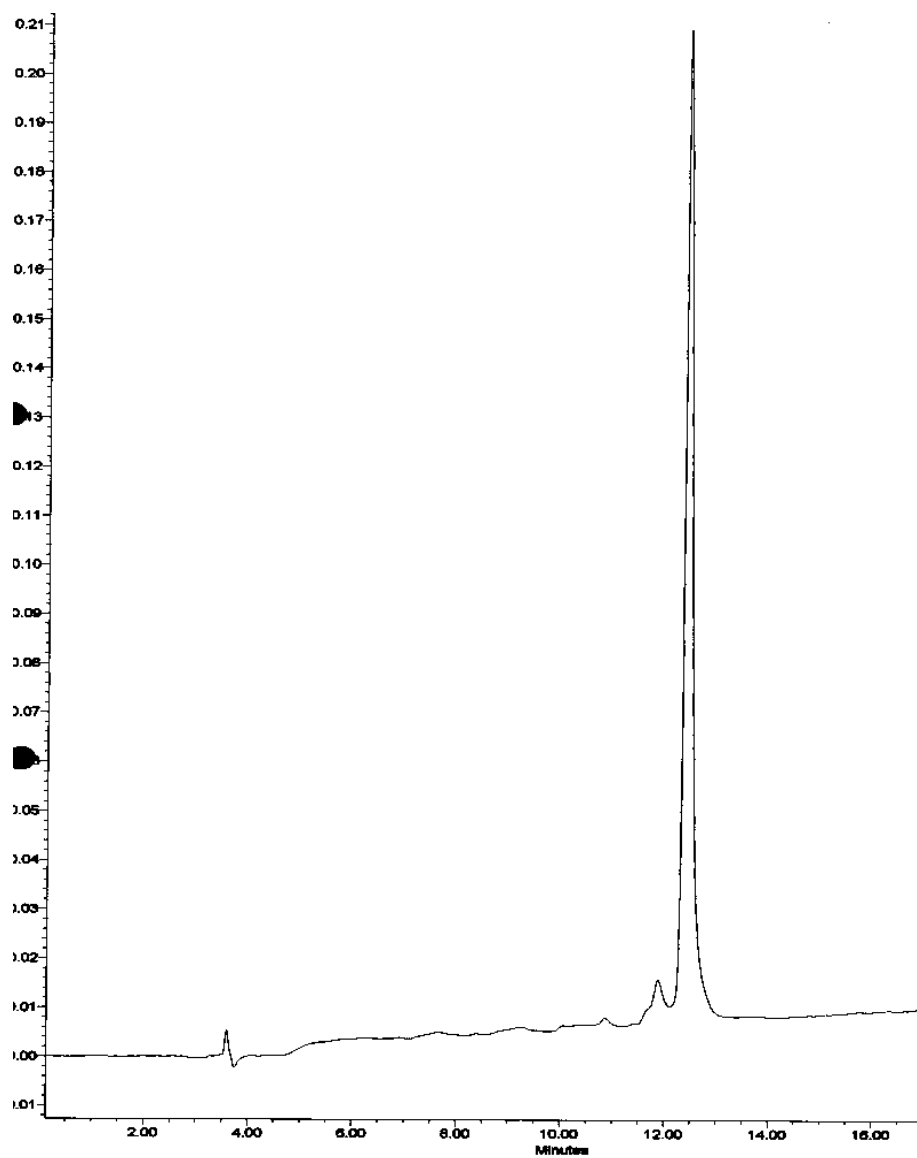
Dendrimer C<sub>1</sub>-F<sub>1</sub>

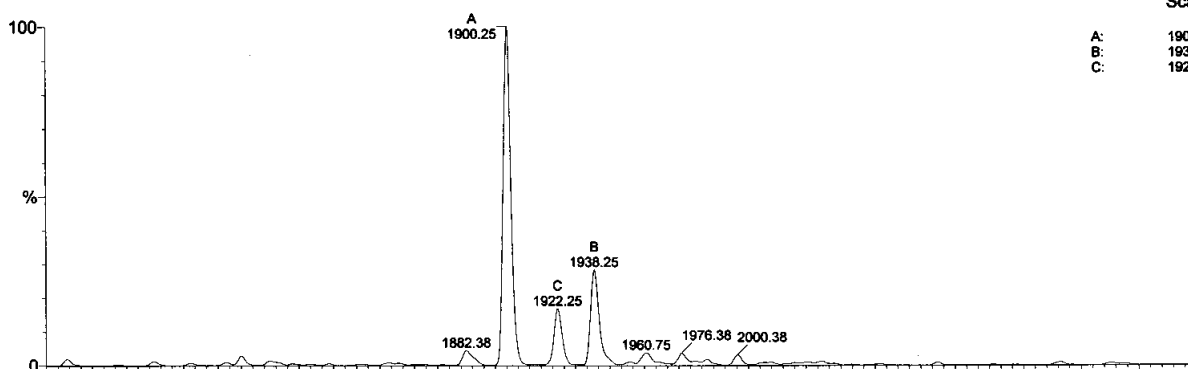
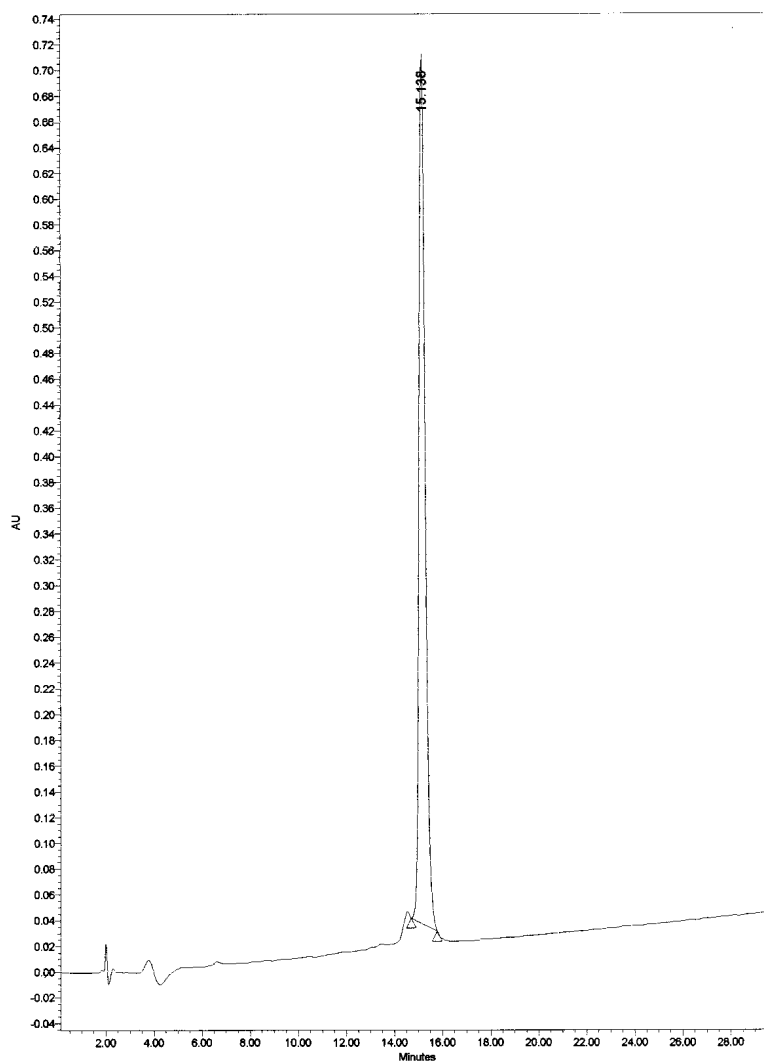


Dendrimer D<sub>1</sub>-E<sub>1</sub>

Dendrimer D<sub>1</sub>-F<sub>1</sub>

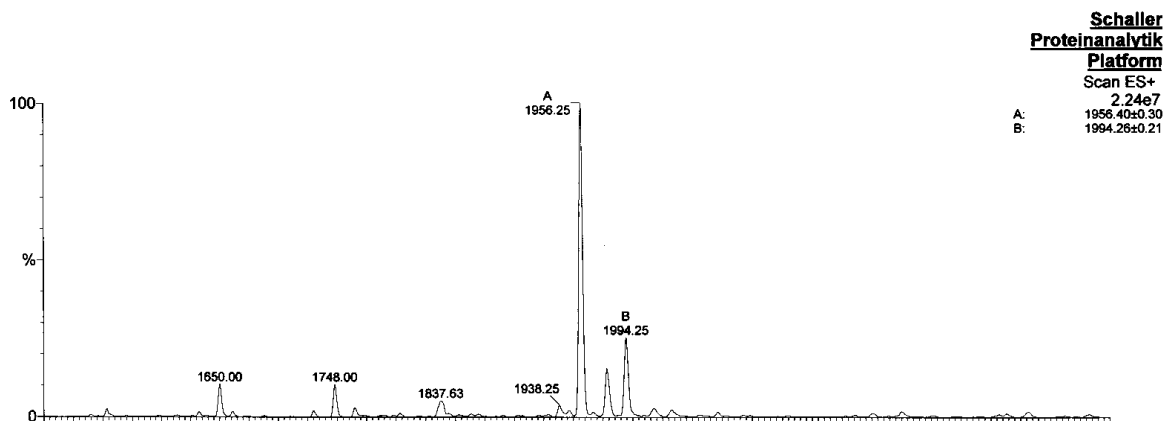
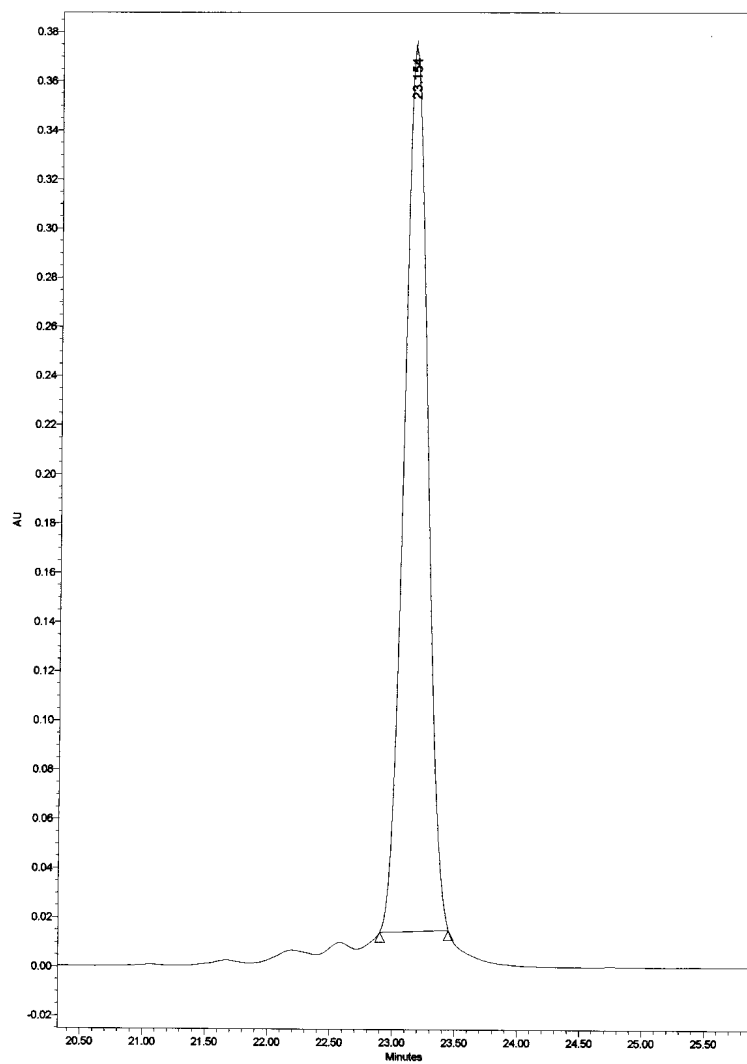
Dendrimer E<sub>1</sub>-F<sub>1</sub>

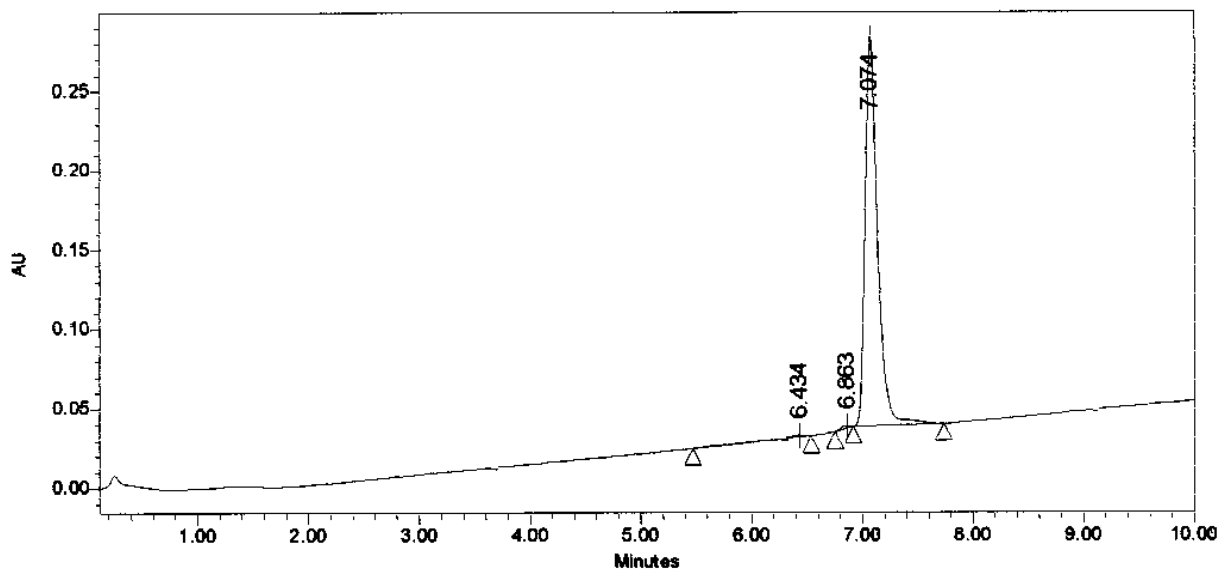
Dendrimer F<sub>1</sub>-F<sub>1</sub>

Dendrimer A<sub>2</sub>

**Schaller**  
**Proteinanalytik**  
**Platform**  
Scan ES+  
7.98e7

Label	m/z
A:	1900.29±0.09
B:	1938.25±0.22
C:	1922.44±0.40

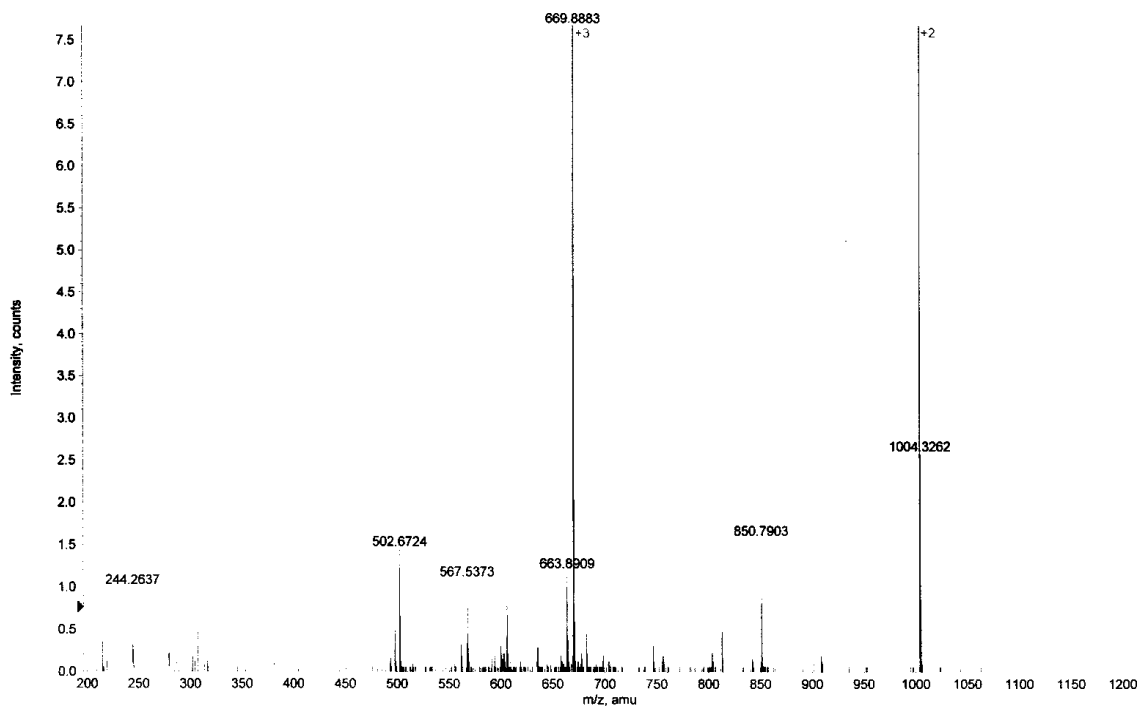
**DendrimerB<sub>2</sub>**

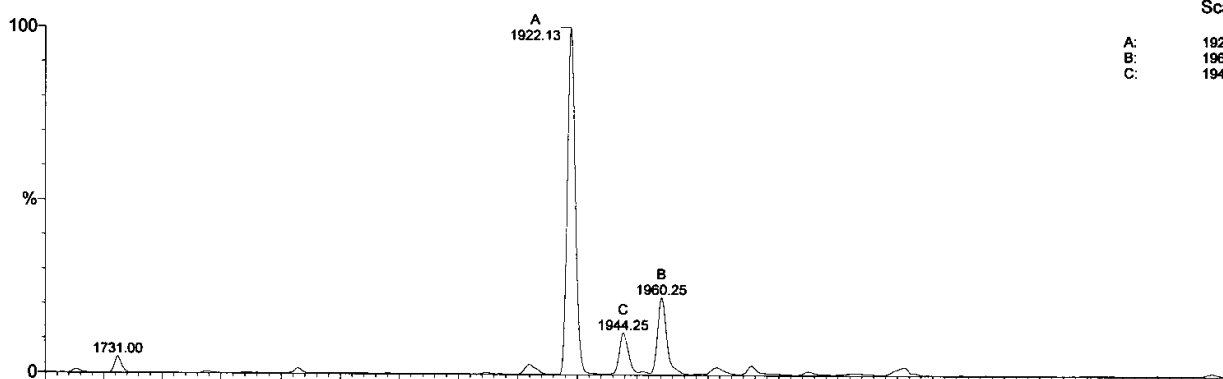
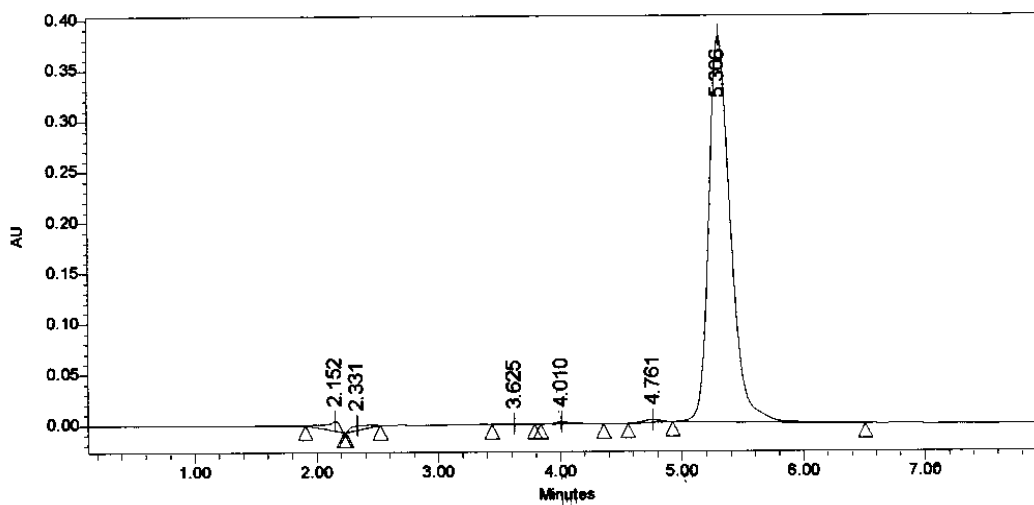
Dendrimer C<sub>2</sub>

+TOF MS: 1.534 to 2.034 min from A28\_A.wiff  
a=3.55688854675700220e-004, t0=3.78043834838899780e+000

ESI-MS positive mode

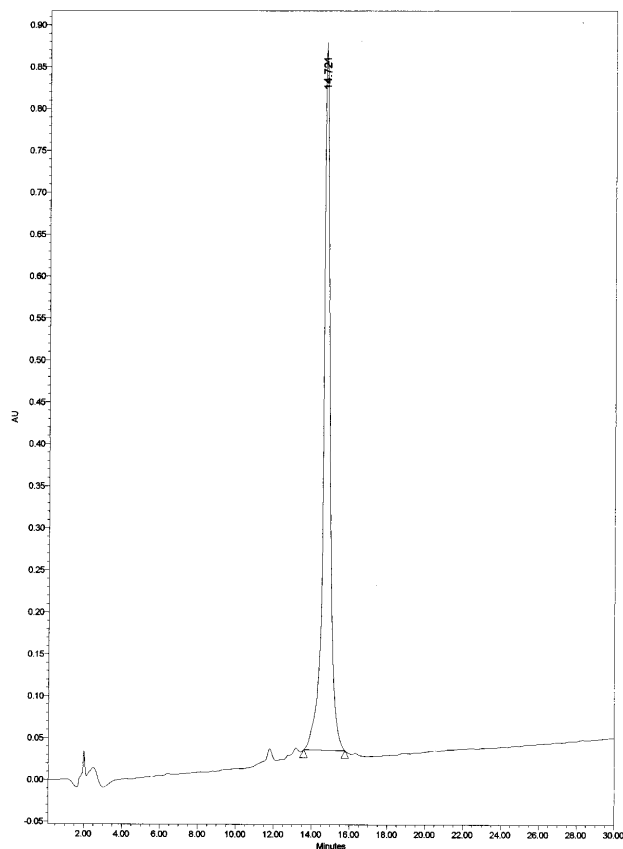
Sample dissolved in MeOH/H<sub>2</sub>O/HFo (74+25+1)



DendrimerD<sub>2</sub>

**Schaller**  
**Proteinanalytik**  
**Platform**  
Scan ES+  
1.69e8  
A: 1922.33±0.34  
B: 1960.18±0.24  
C: 1944.46±0.52

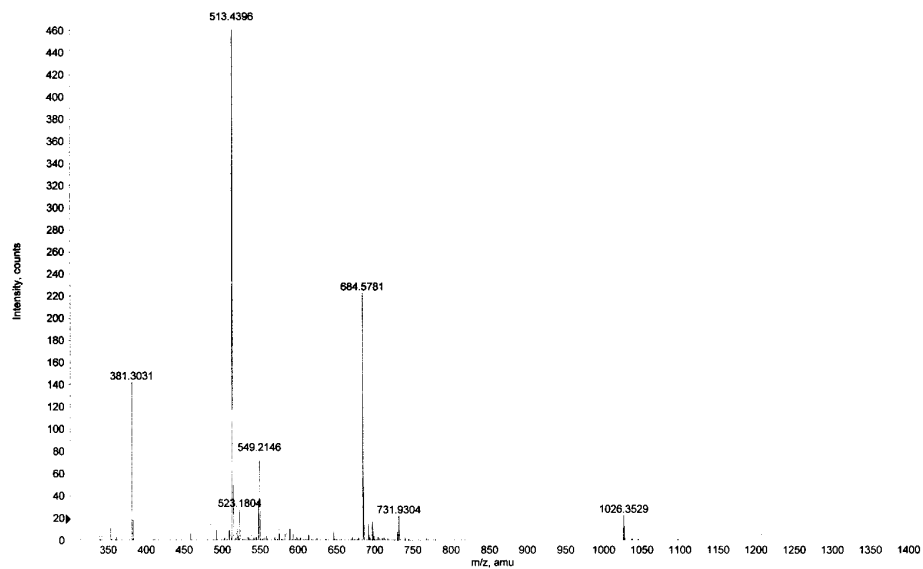


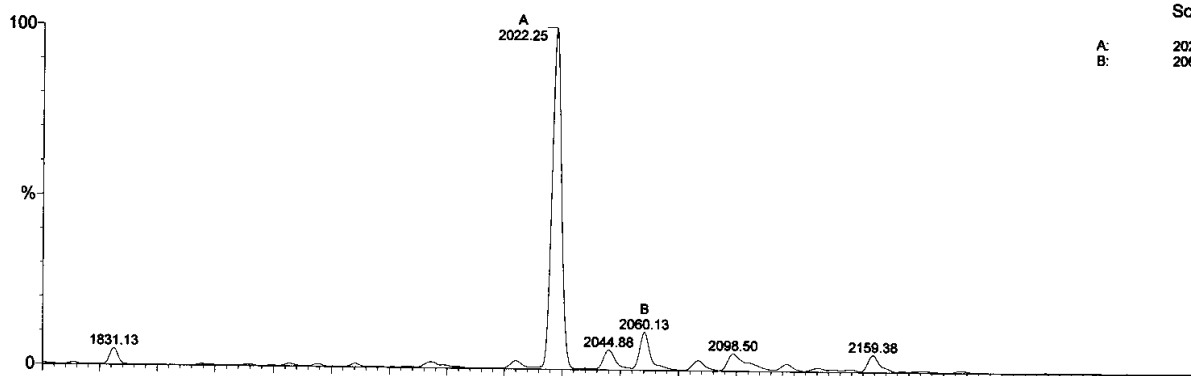
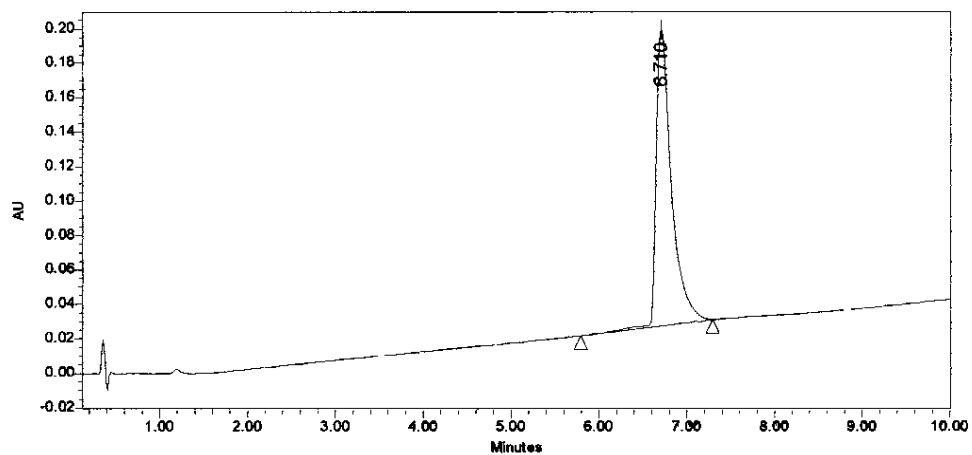
Dendrimer E<sub>2</sub>

\*TOF MS: 1.634 to 2.951 min from A29\_42\_45\_A.wiff  
a=3.55671773310228740e-004, k0=3.47708225918177050e+000

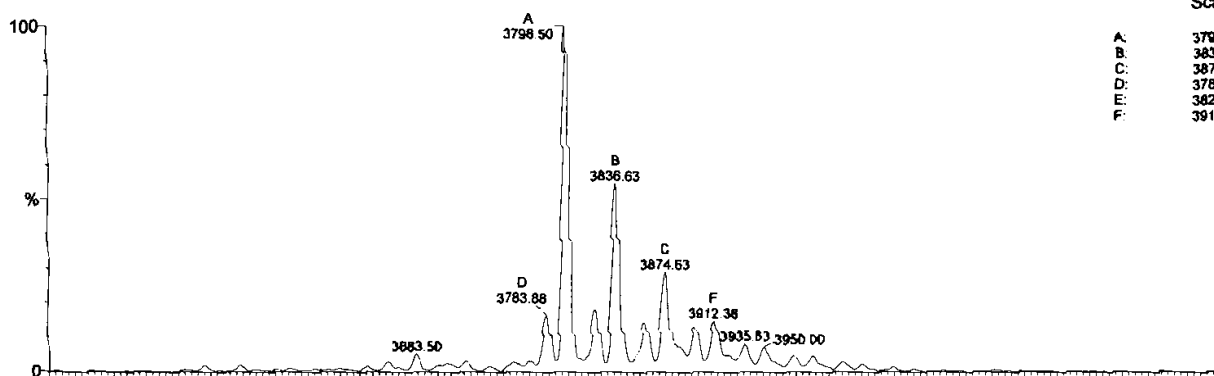
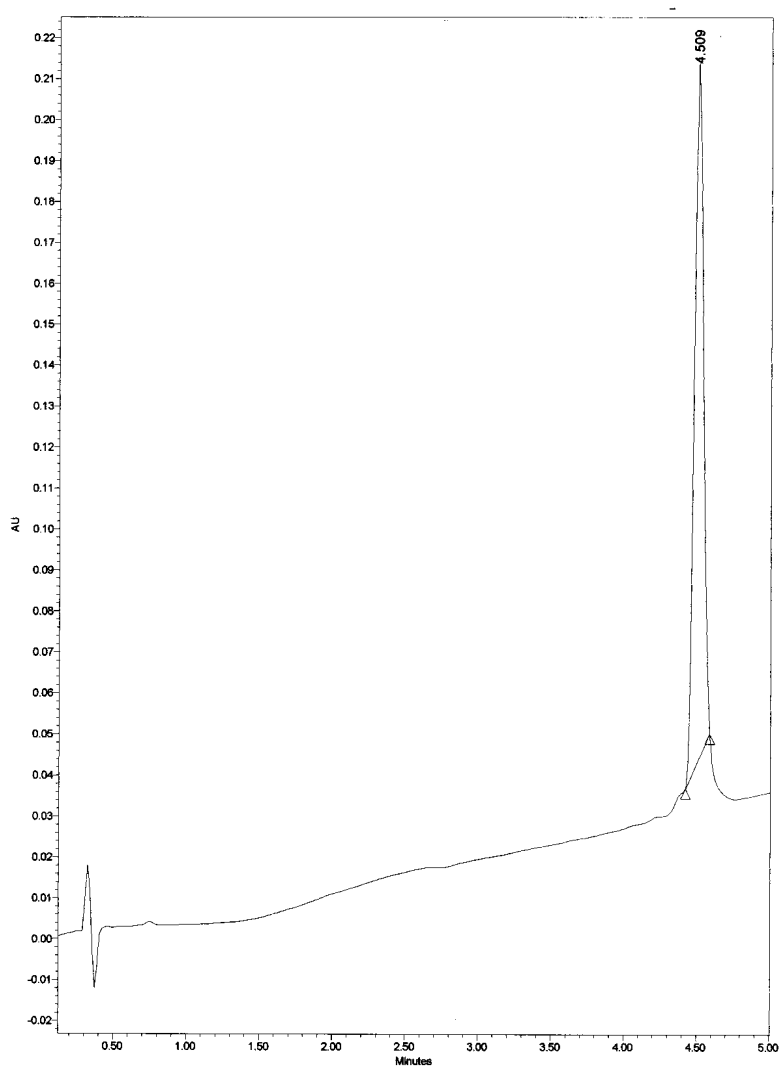
ESI-MS positive mode

Sample dissolved in MeOH/H<sub>2</sub>O/HFo (74+25+1)



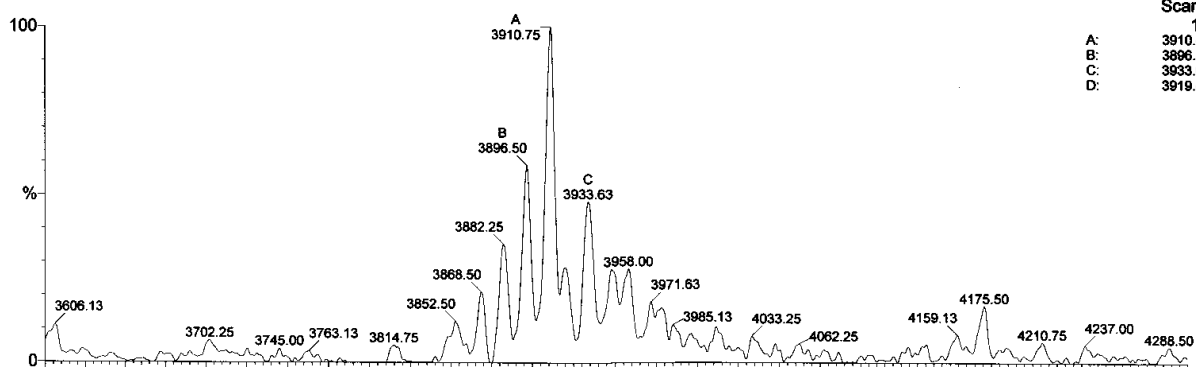
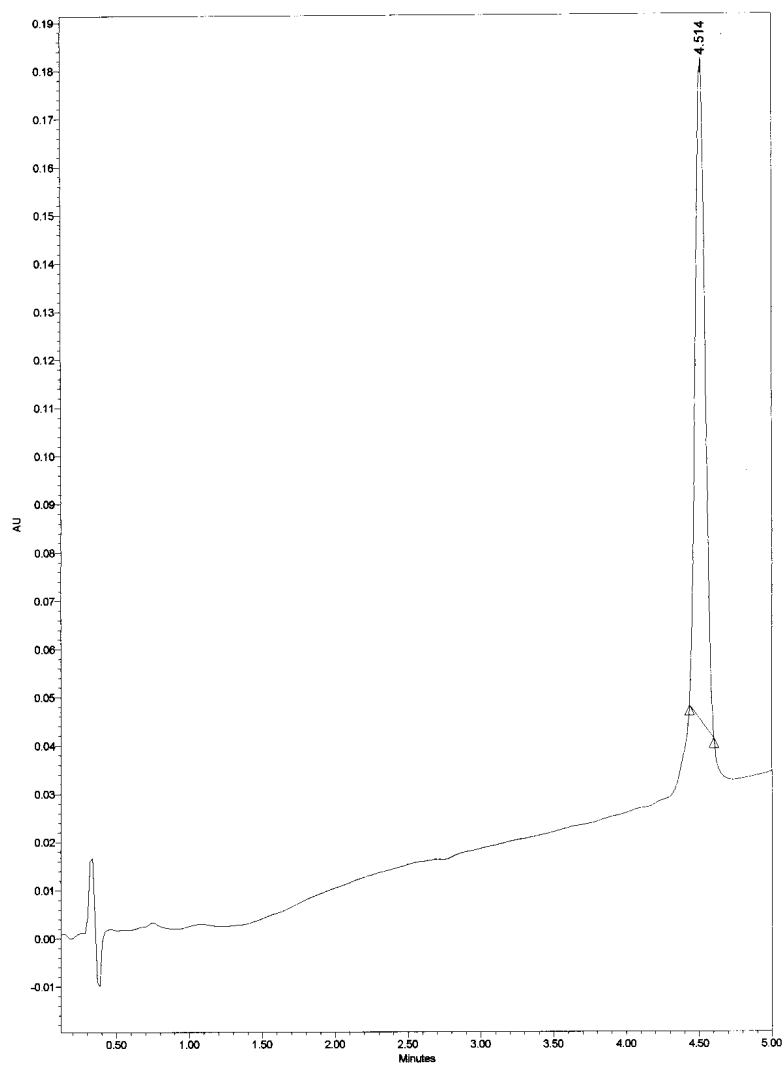
DendrimerF<sub>2</sub>

**Schaller**  
**Proteinanalytik**  
**Plattform**  
Scan ES+  
2.38e8  
A: 2022.27±0.14  
B: 2060.29±0.14

Dendrimer A<sub>2</sub>-A<sub>2</sub>

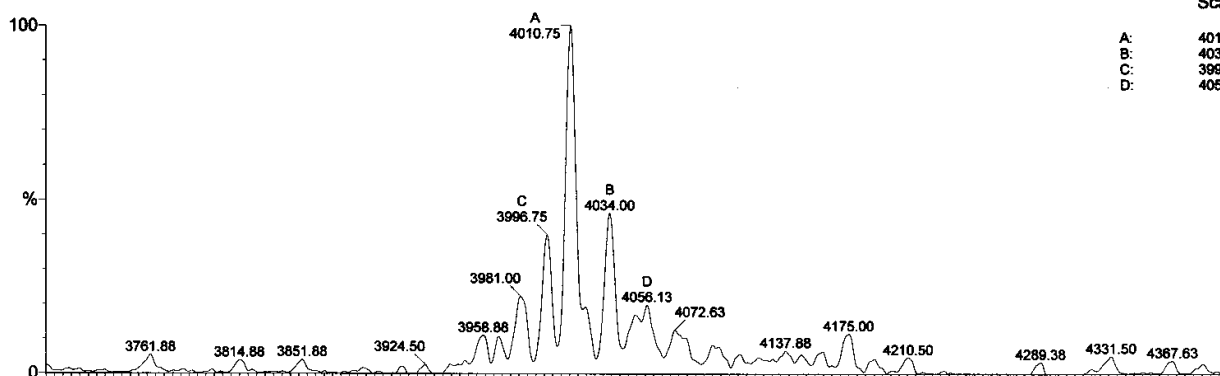
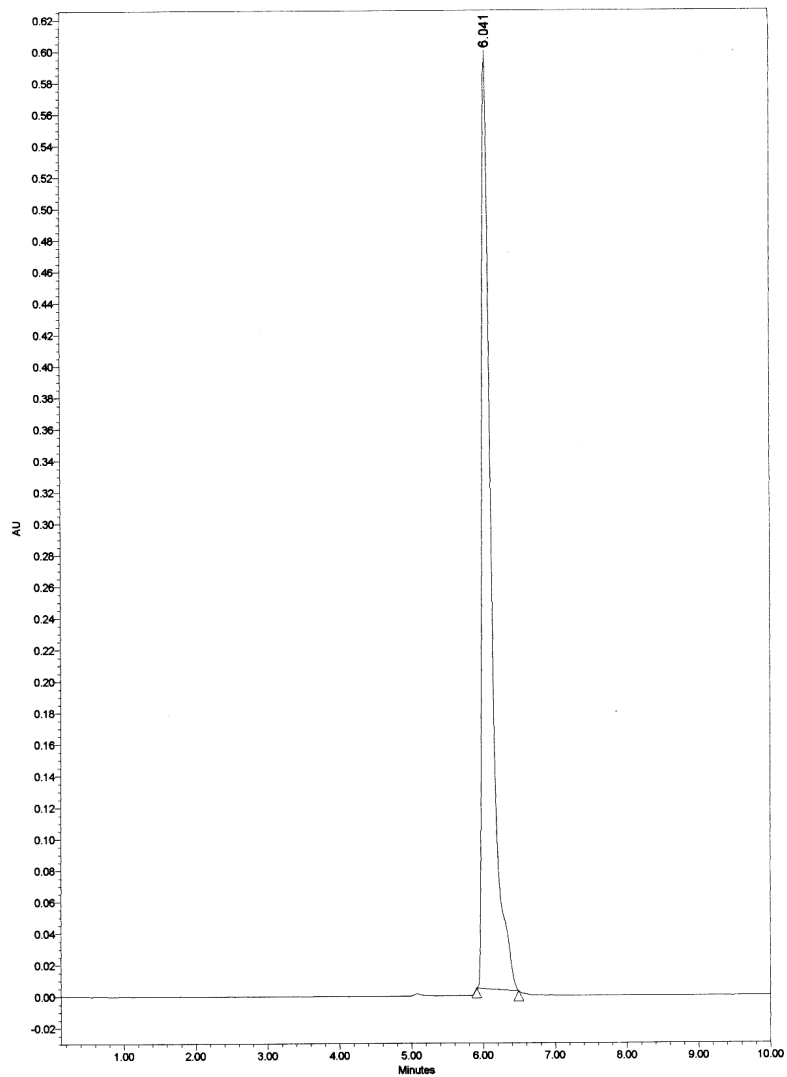
**Schaller**  
**Proteinanalytik**  
**Platform**  
Scan ES+  
2.45e7

A:	3798.47±0.24
B:	3836.06±0.15
C:	3874.76±0.07
D:	3783.50±0.20
E:	3820.86±0.13
F:	3912.30±0.16

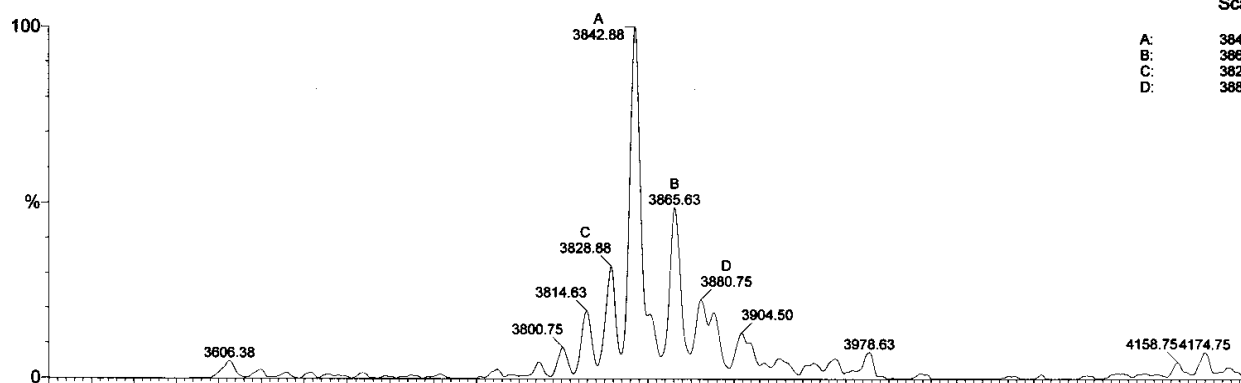
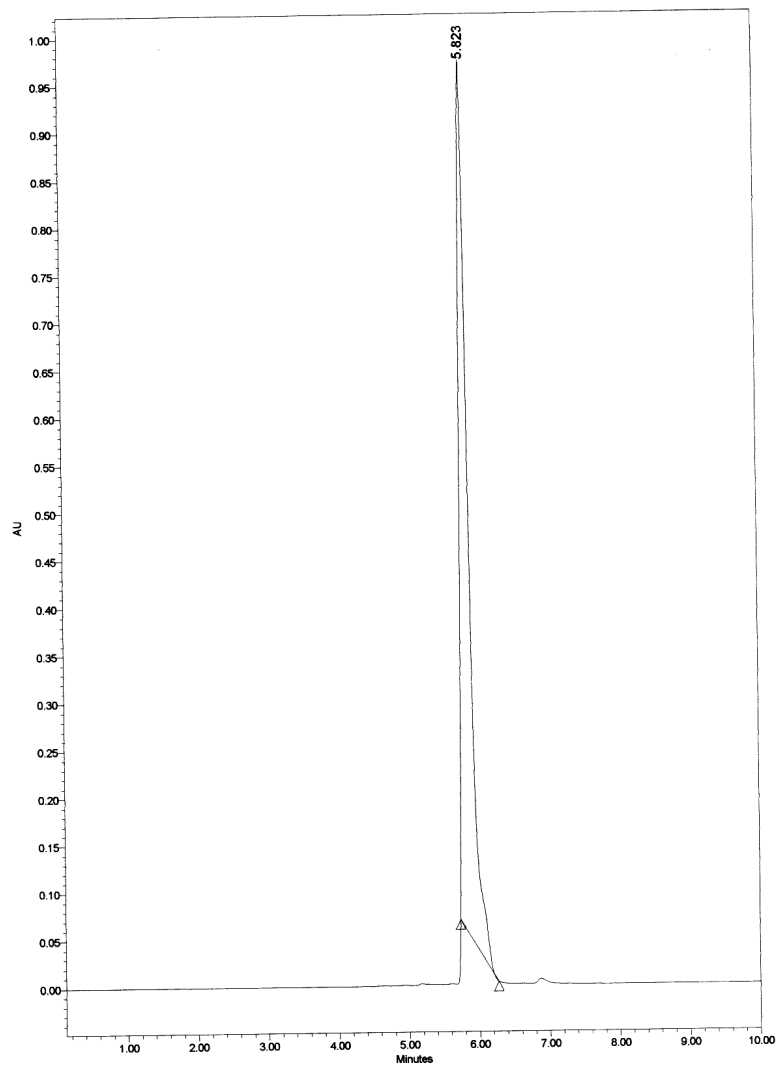
Dendrimer B<sub>2</sub>-B<sub>2</sub>

**Schaller**  
**Proteinanalytik**  
**Platform**  
 Scan ES+

1.81e6  
 A: 3910.46±0.60  
 B: 3896.50±0.04  
 C: 3933.67±0.26  
 D: 3919.46±0.63

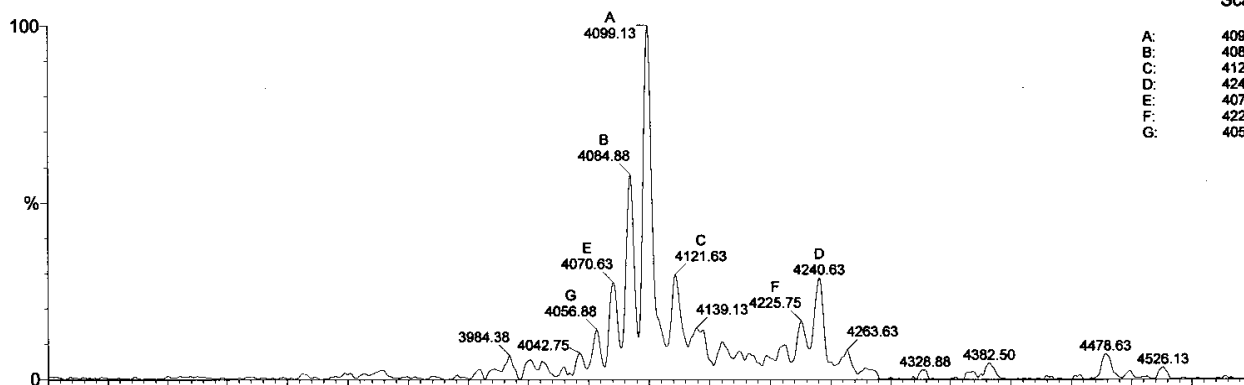
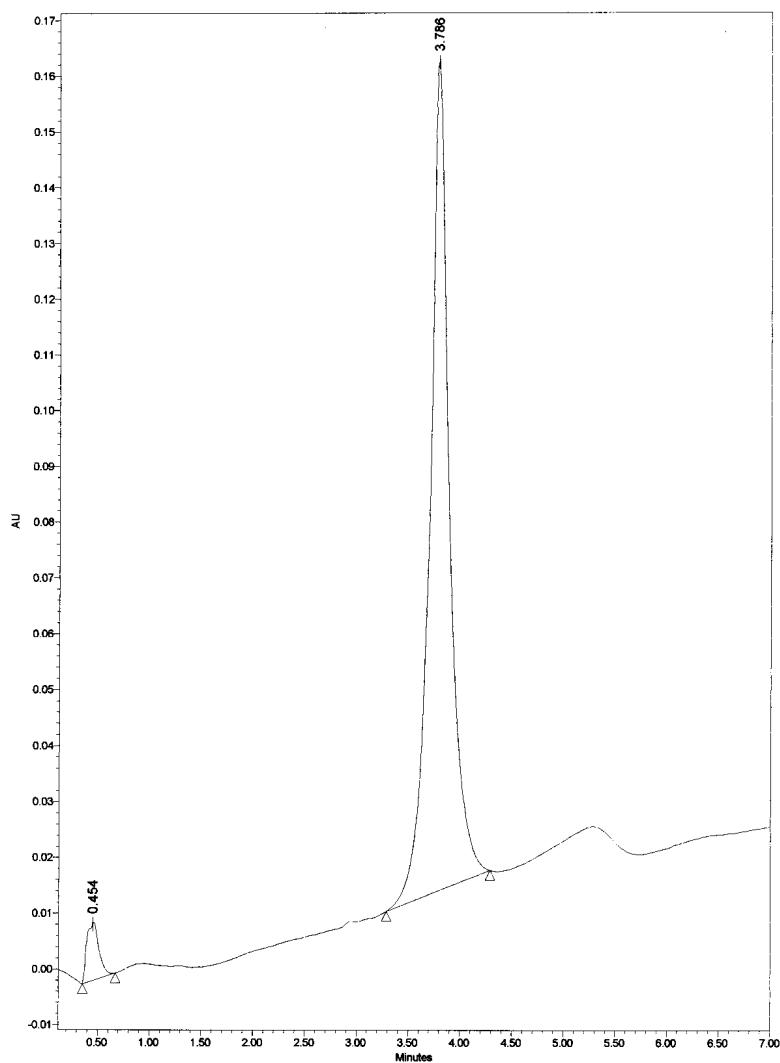
Dendrimer C<sub>2</sub>-C<sub>2</sub>

**Schaller**  
**Proteinanalytik**  
**Platform**  
Scan ES+  
3.43e6  
A: 4011.12±0.39  
B: 4034.07±0.37  
C: 3996.47±0.42  
D: 4056.01±0.01

**Dendrimer D<sub>2</sub>-D<sub>2</sub>**

**Schaller**  
**Proteinanalytik**  
**Platform**  
Scan ES+  
7.56e6

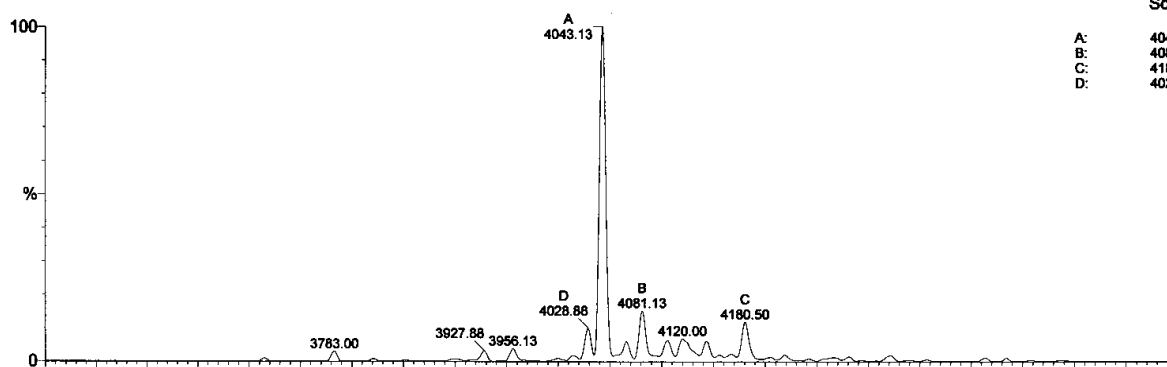
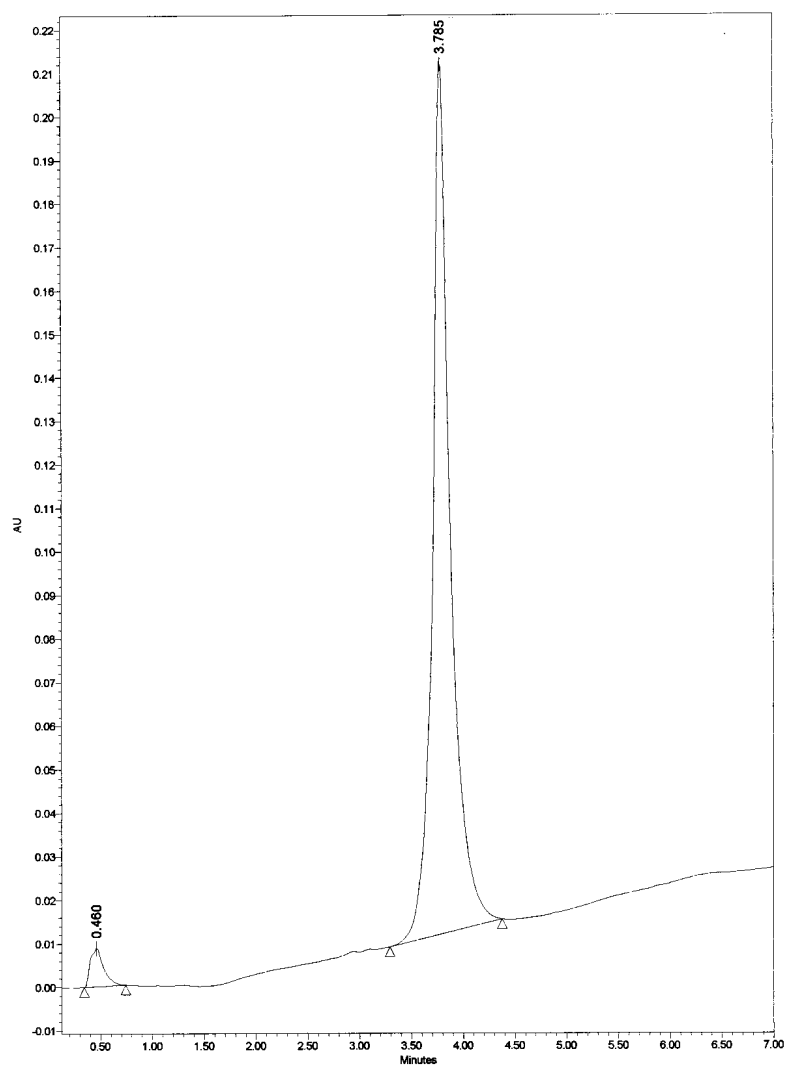
Label	m/z
A:	3842.88±0.20
B:	3865.70±0.25
C:	3828.65±0.18
D:	3880.57±0.30

Dendrimer E<sub>2</sub>-E<sub>2</sub>

**Schaller**  
**Proteinanalytik**  
**Platform**

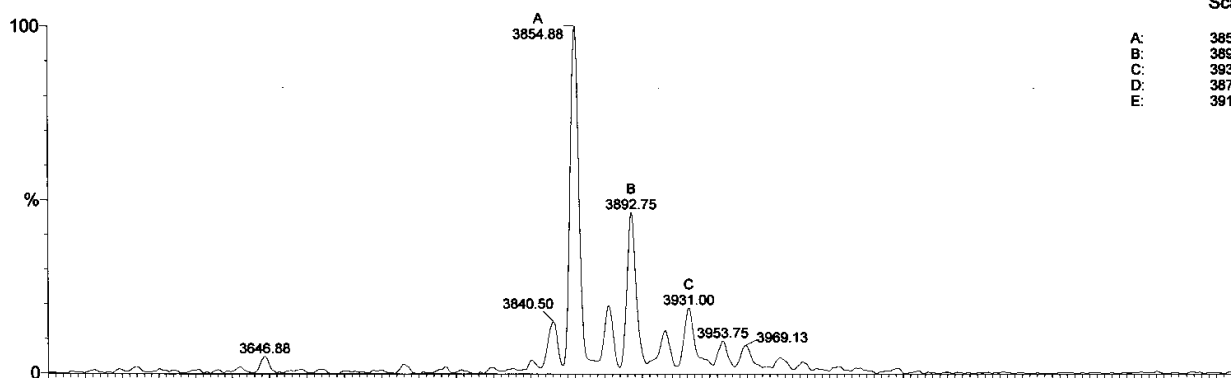
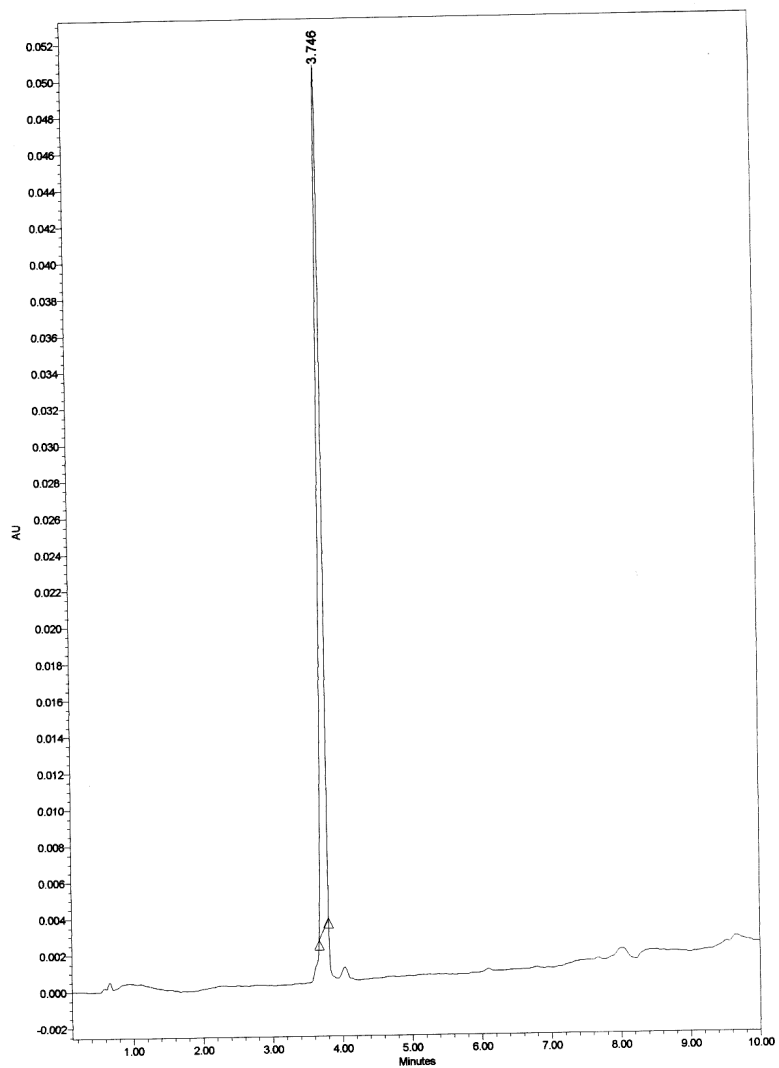
Scan ES+  
8.01e6

A: 4099.05±0.16  
B: 4084.77±0.08  
C: 4122.10±0.73  
D: 4240.41±0.19  
E: 4070.54±0.38  
F: 4225.09±0.87  
G: 4056.74±0.40

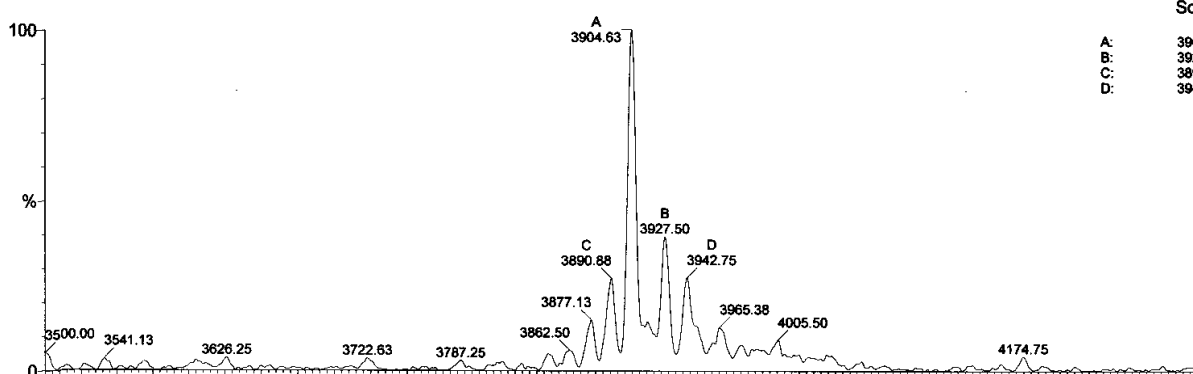
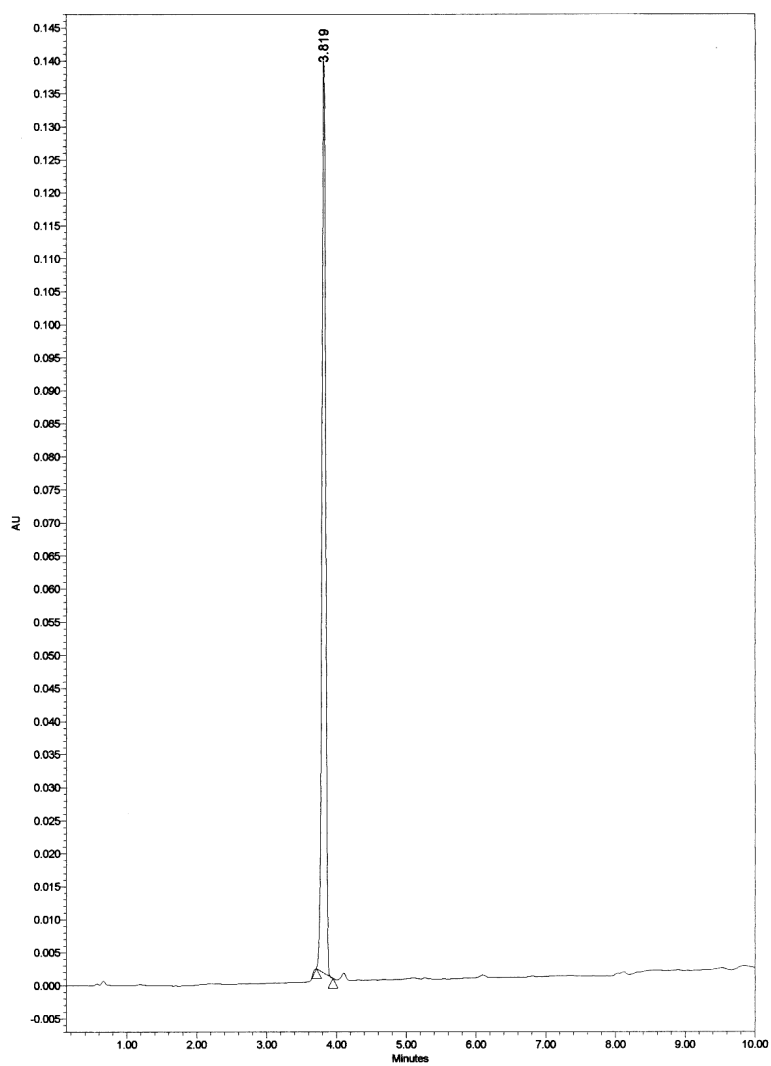
Dendrimer F<sub>2</sub>-F<sub>2</sub>

**Schaller**  
**Proteinanalytik**  
**Platform**  
Scan ES+  
1.55e8  
A: 4043.09±0.22  
B: 4081.02±0.03  
C: 4180.38±0.07  
D: 4028.90±0.04

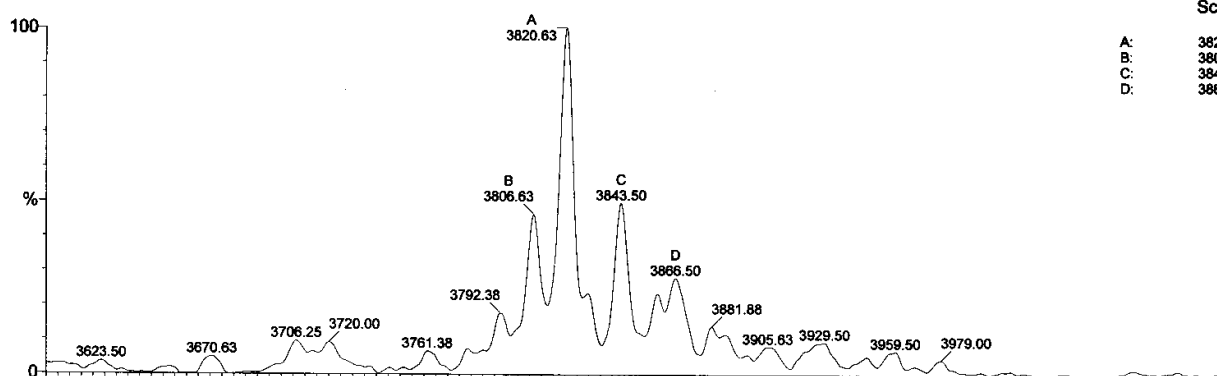
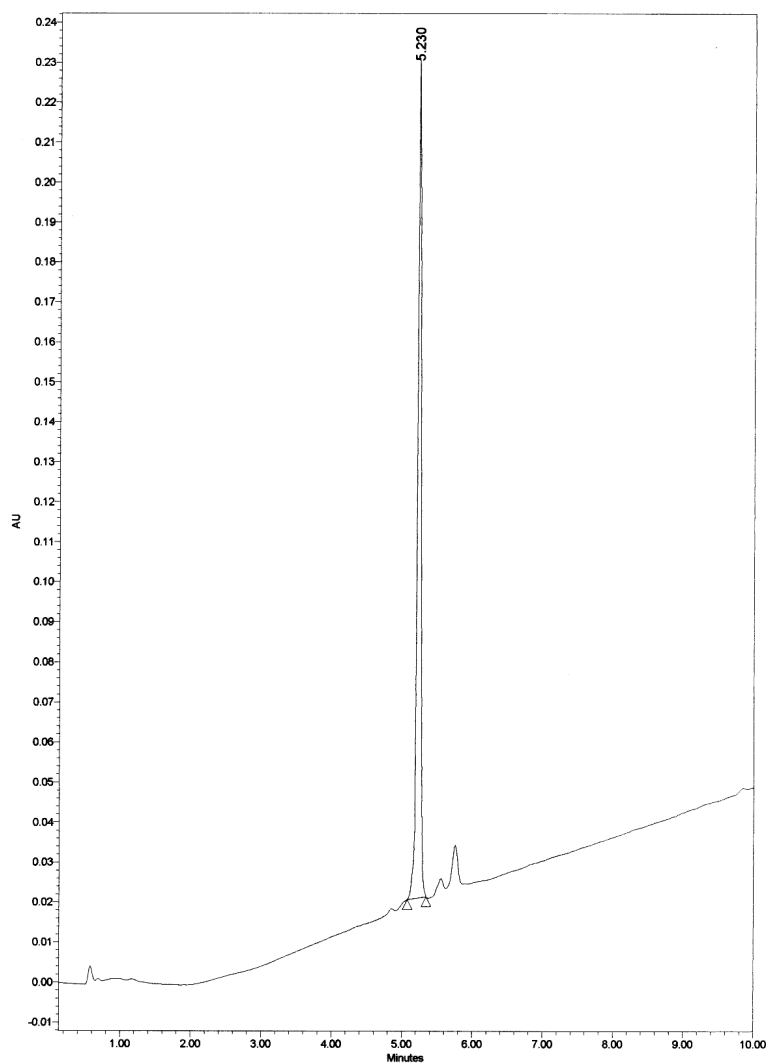


Dendrimer A<sub>2</sub>-B<sub>2</sub>

**Schaller**  
**Proteinanalytik**  
**Platform**  
Scan ES+  
8.35e6  
A: 3854.83±0.20  
B: 3892.68±0.10  
C: 3931.02±0.29  
D: 3877.26±0.35  
E: 3915.26±0.06

Dendrimer A<sub>2</sub>-C<sub>2</sub>

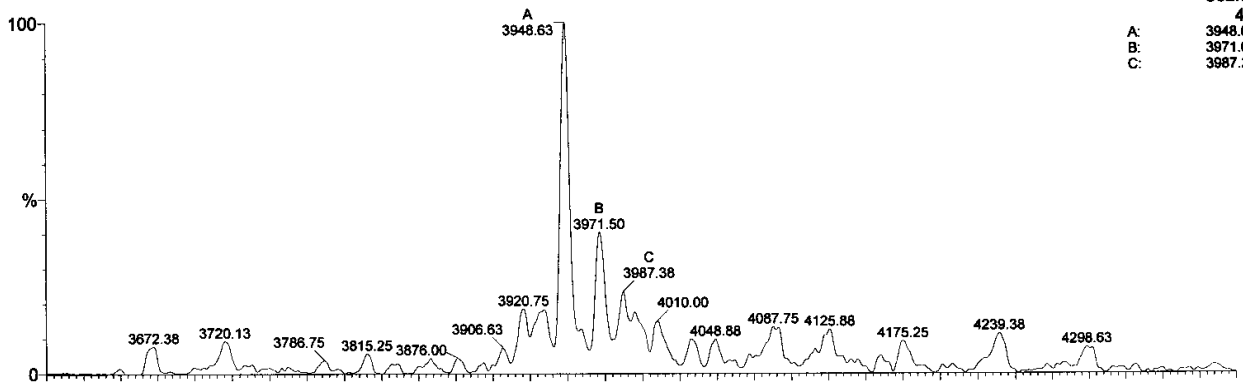
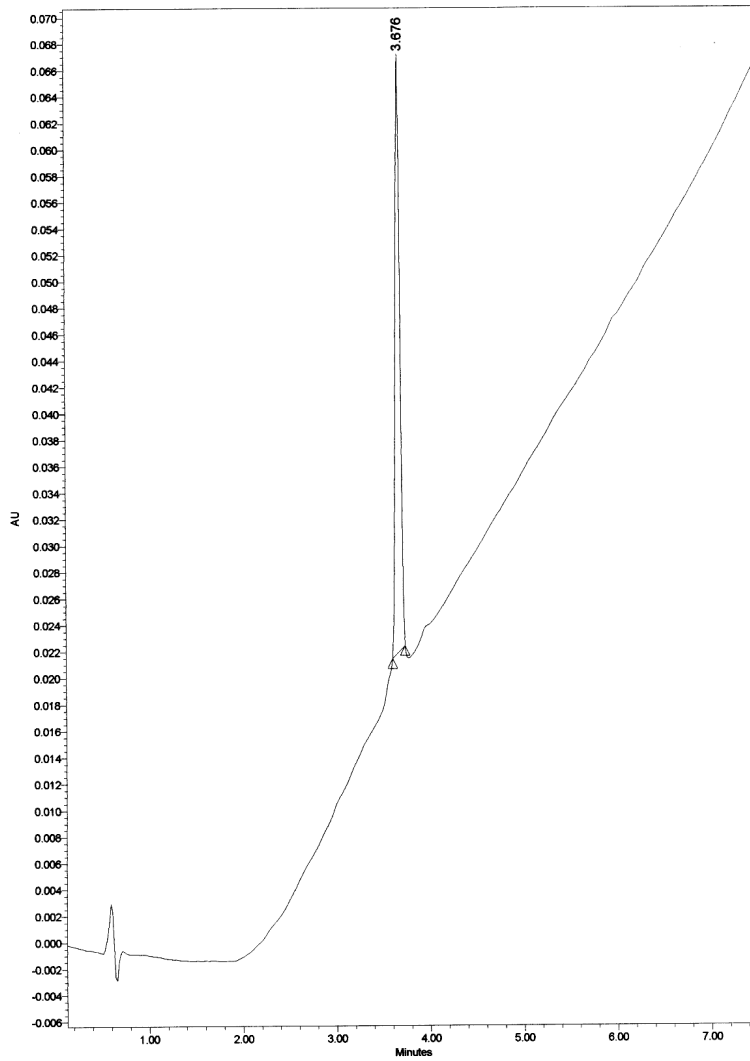
**Schaller**  
**Proteinanalytik**  
**Plattform**  
Scan ES+  
3.87e6  
A: 3904.44±0.58  
B: 3927.82±0.15  
C: 3890.82±0.17  
D: 3942.97±0.05

Dendrimer A<sub>2</sub>-D<sub>2</sub>

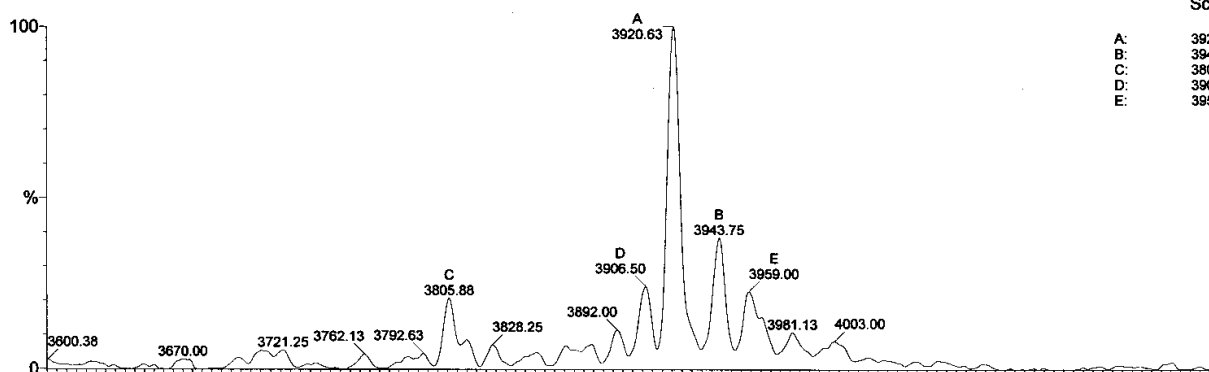
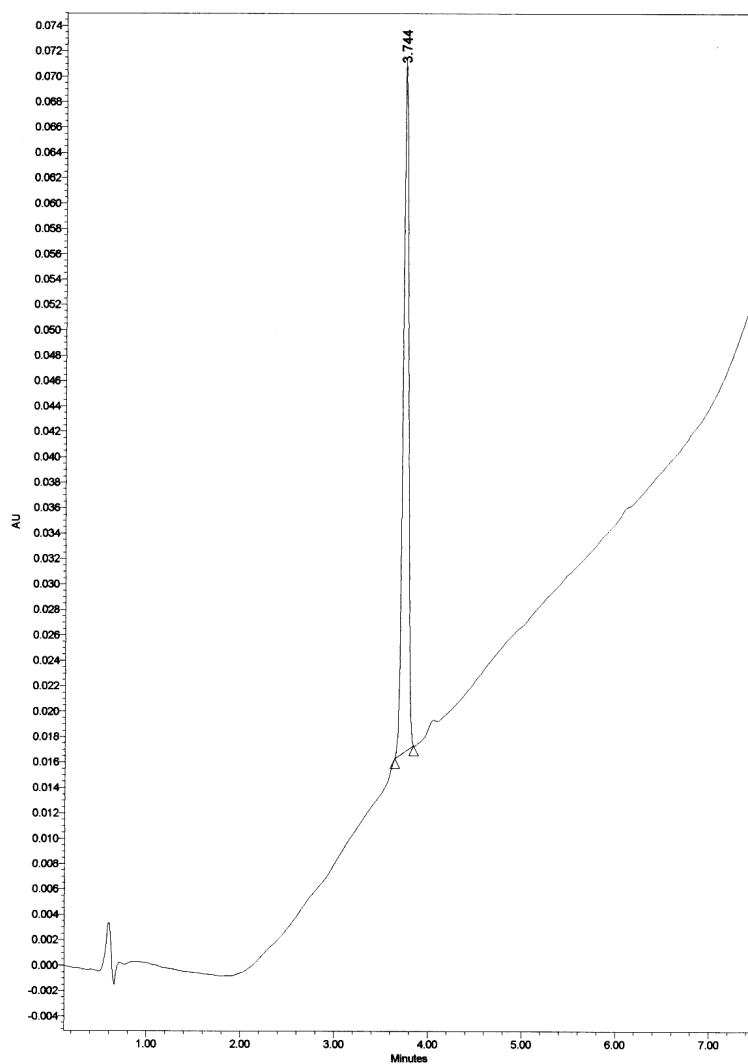
**Schaller**  
**Proteinanalytik**  
**Platform**  
Scan ES+  
5.84e6

A:	3820.93±0.30
B:	3806.51±0.15
C:	3843.31±0.01
D:	3866.45±0.44

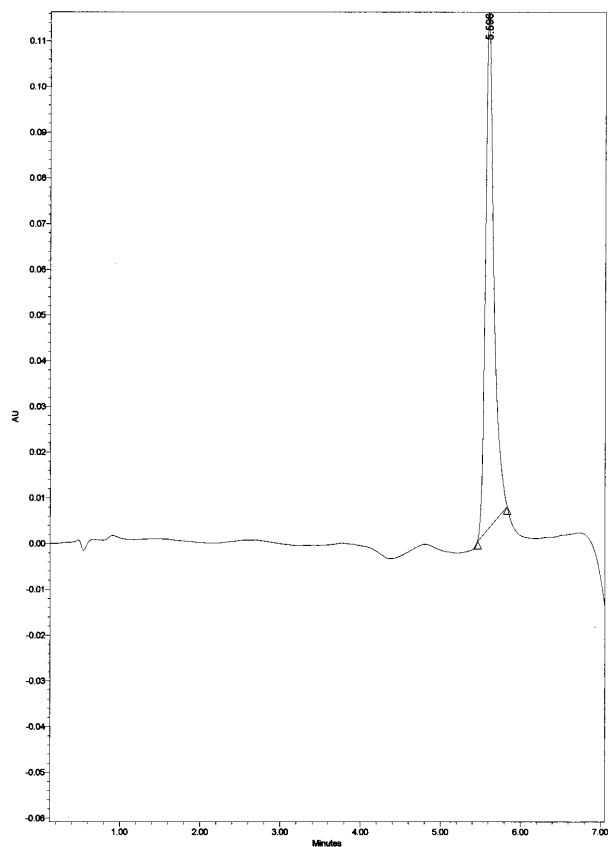
Dendrimer A<sub>2</sub>-E<sub>2</sub>



**Schaller**  
**Proteinanalytik**  
**Platform**  
 Scan ES+  
 4.52e6  
 A: 3948.63±0.28  
 B: 3971.65±0.35  
 C: 3987.25±0.24

Dendrimer A<sub>2</sub>-F<sub>2</sub>

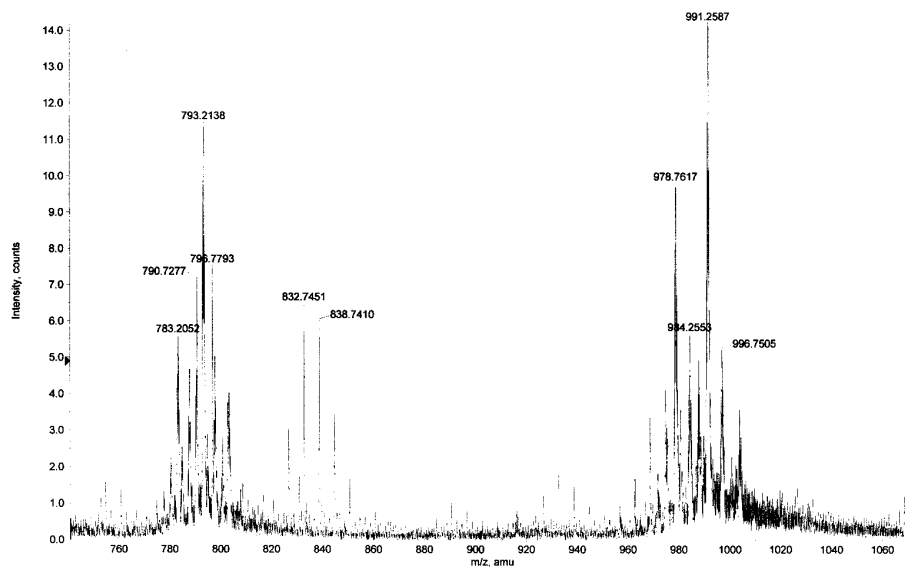
**Schaller**  
**Proteinanalytik**  
**Platform**  
Scan ES+  
7.53e6  
A: 3920.77±0.35  
B: 3943.84±0.29  
C: 3805.86±0.25  
D: 3906.13±0.46  
E: 3958.57±0.13

Dendrimer B<sub>2</sub>-C<sub>2</sub>

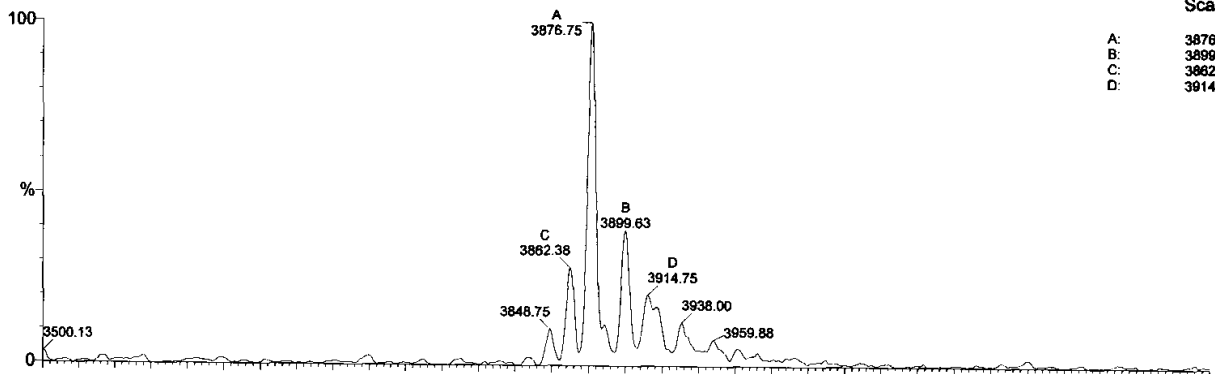
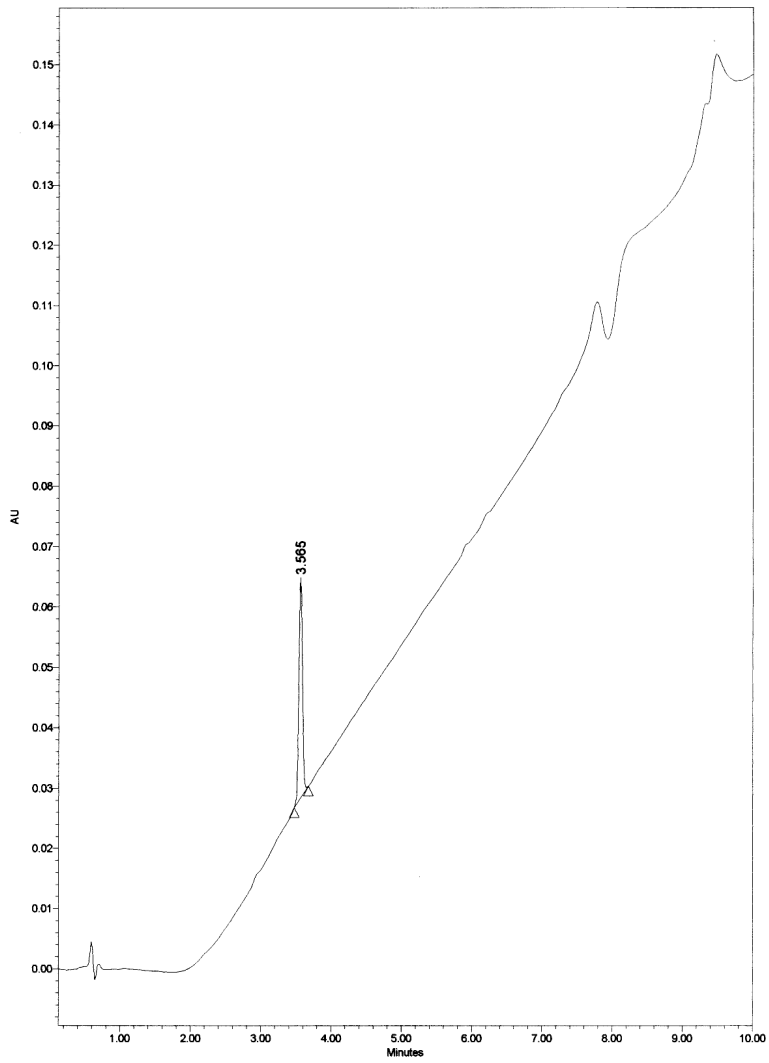
+TOF MS: 1.784 to 3.868 min from A28\_A18\_A.wiff  
a=3.55634542703044630e-004, 10=e-63729501778552780e+000

ESI-MS positive mode

Sample dissolved in MeOH/H<sub>2</sub>O/HFo (74\*25\*1)

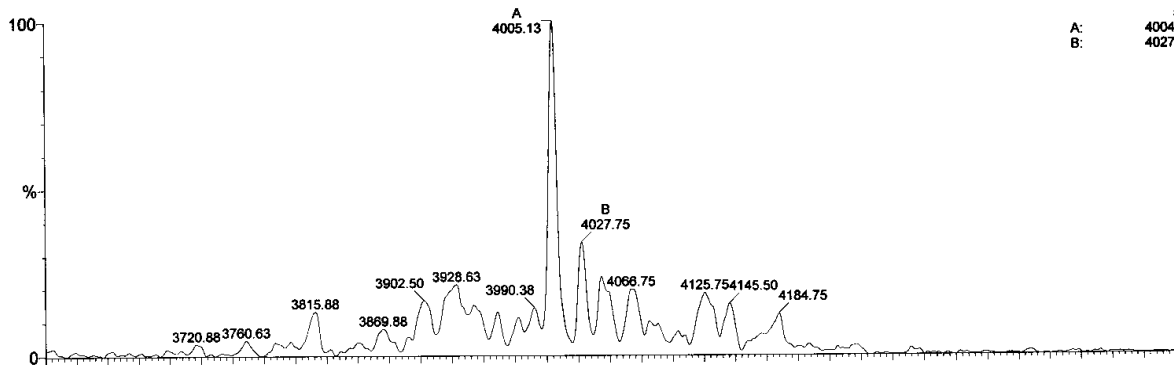
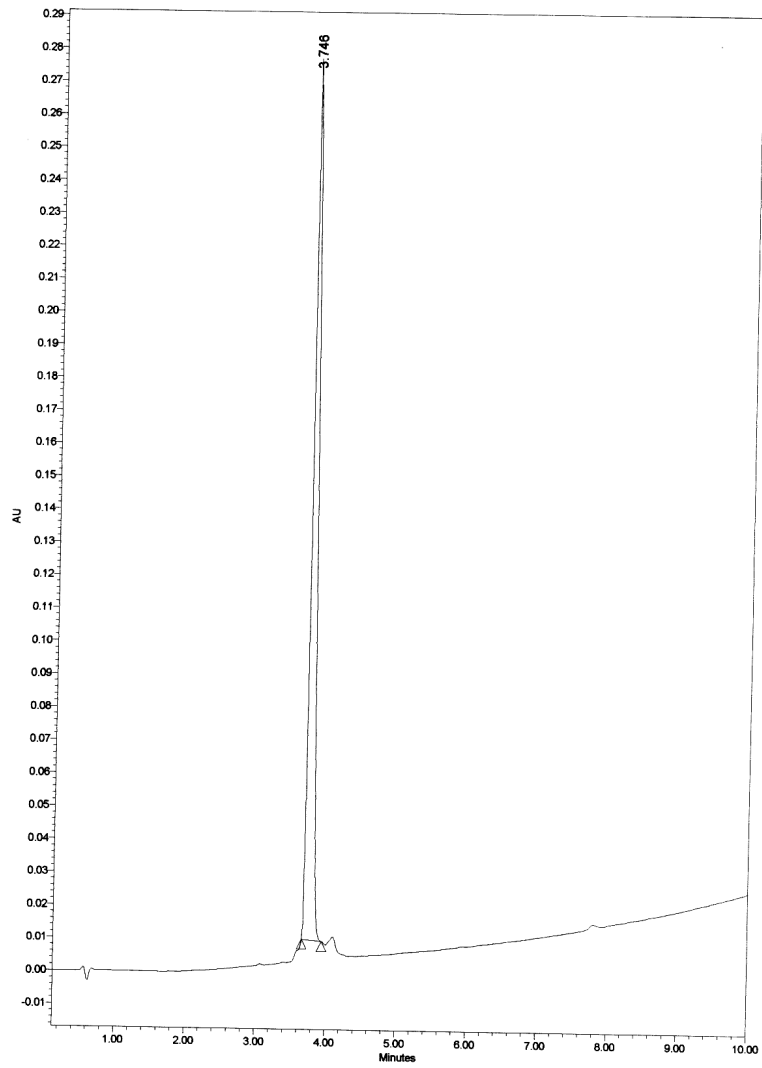


**Dendrimer B<sub>2</sub>-D<sub>2</sub>**



**Schaller**  
**Proteinanalytik**  
**Platform**  
 Scan ES+  
 5.97e6  
 A: 3876.63±0.22  
 B: 3889.69±0.22  
 C: 3882.55±0.53  
 D: 3914.47±0.33

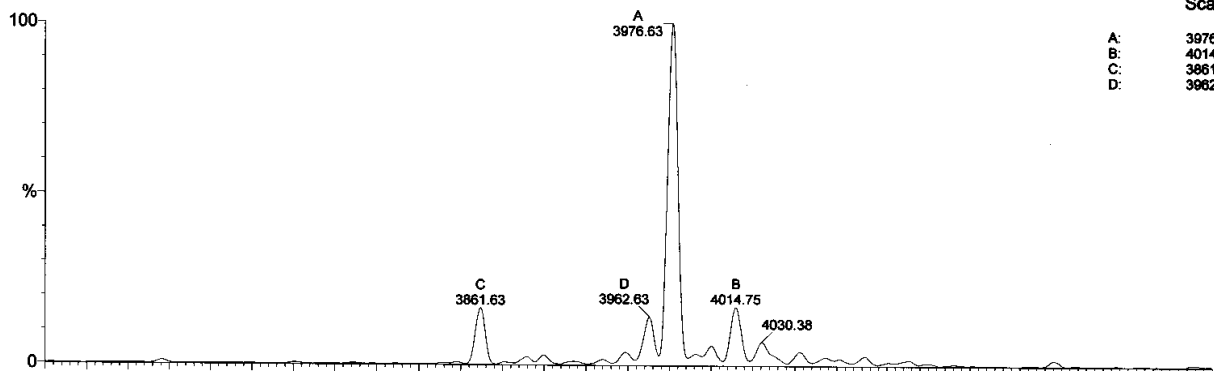
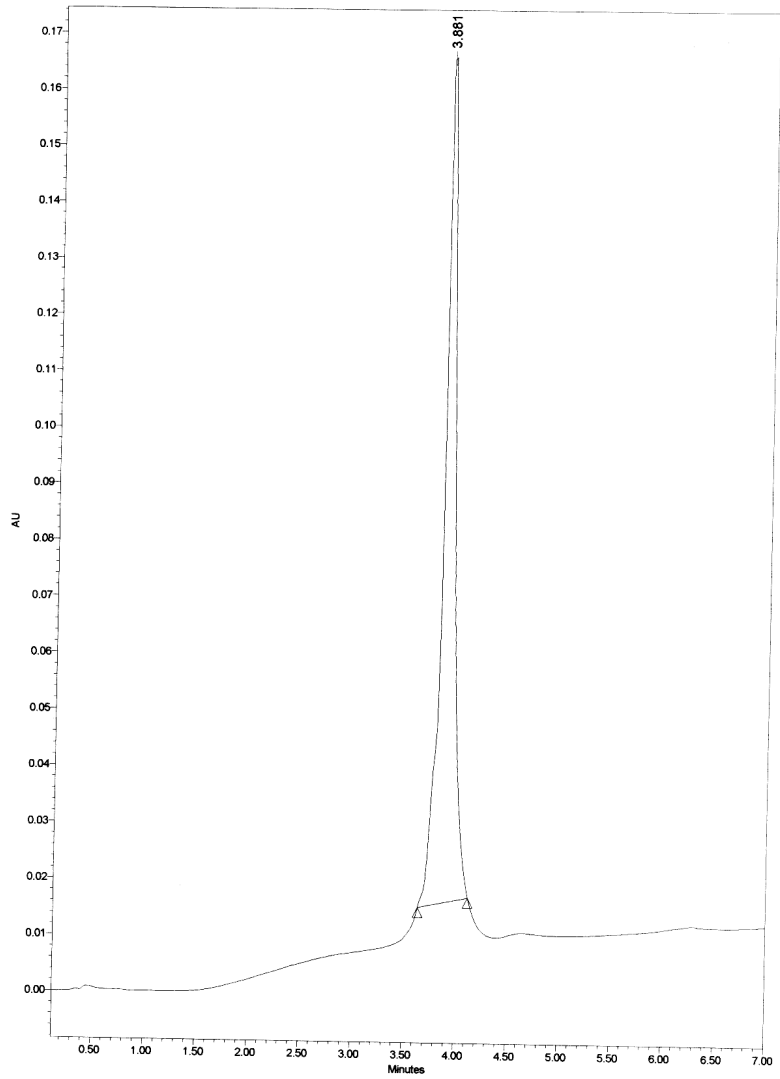
Dendrimer B<sub>2</sub>-E<sub>2</sub>



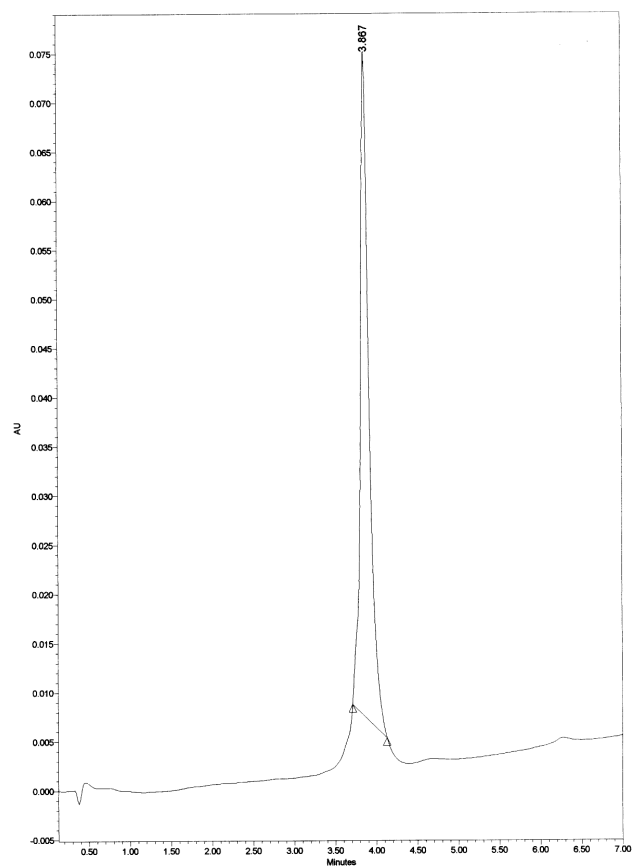
**Schaller**  
**Proteinanalytik**  
**Plattform**  
Scan ES+  
5.48e6  
A: 4004.93±0.16  
B: 4027.99±0.26



**Dendrimer B<sub>2</sub>-F<sub>2</sub>**



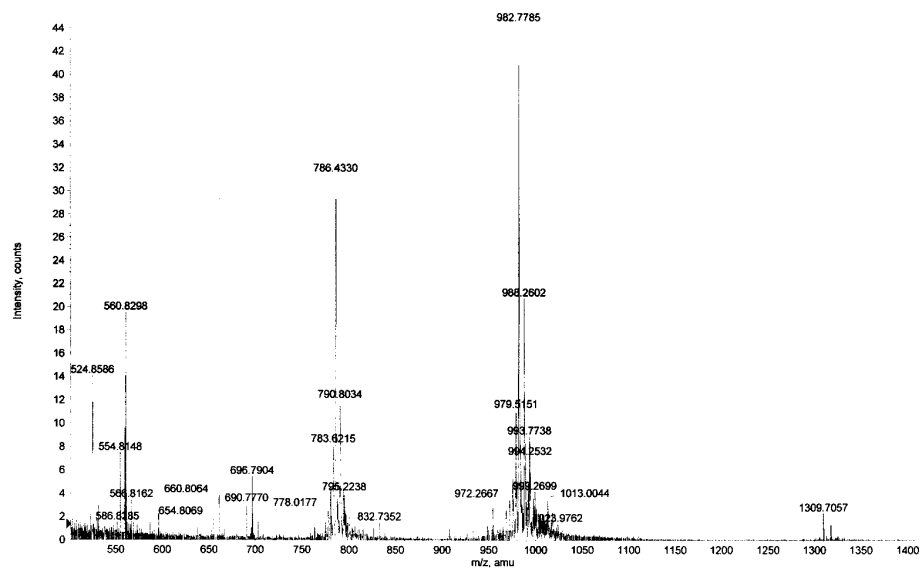
**Schaller**  
**Proteinanalytik**  
**Platform**  
Scan ES+  
4.00e7  
A: 3976.81±0.31  
B: 4014.80±0.08  
C: 3861.75±0.25  
D: 3962.83±0.24

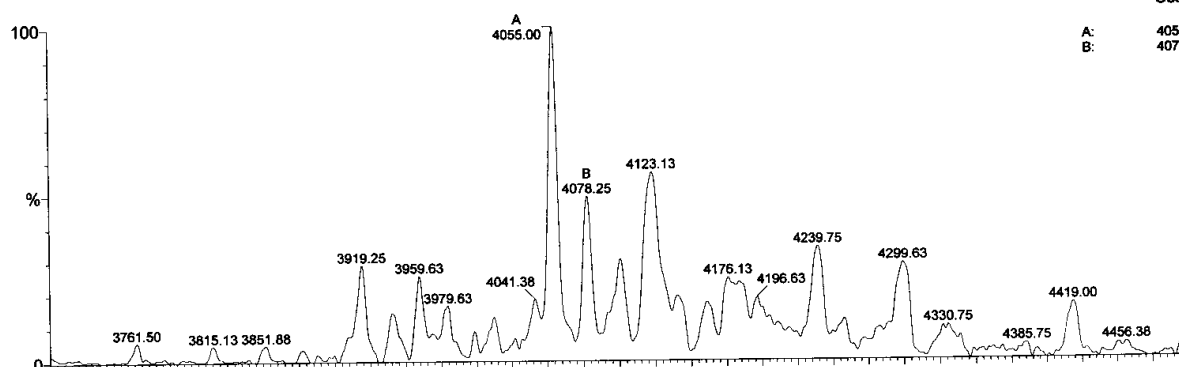
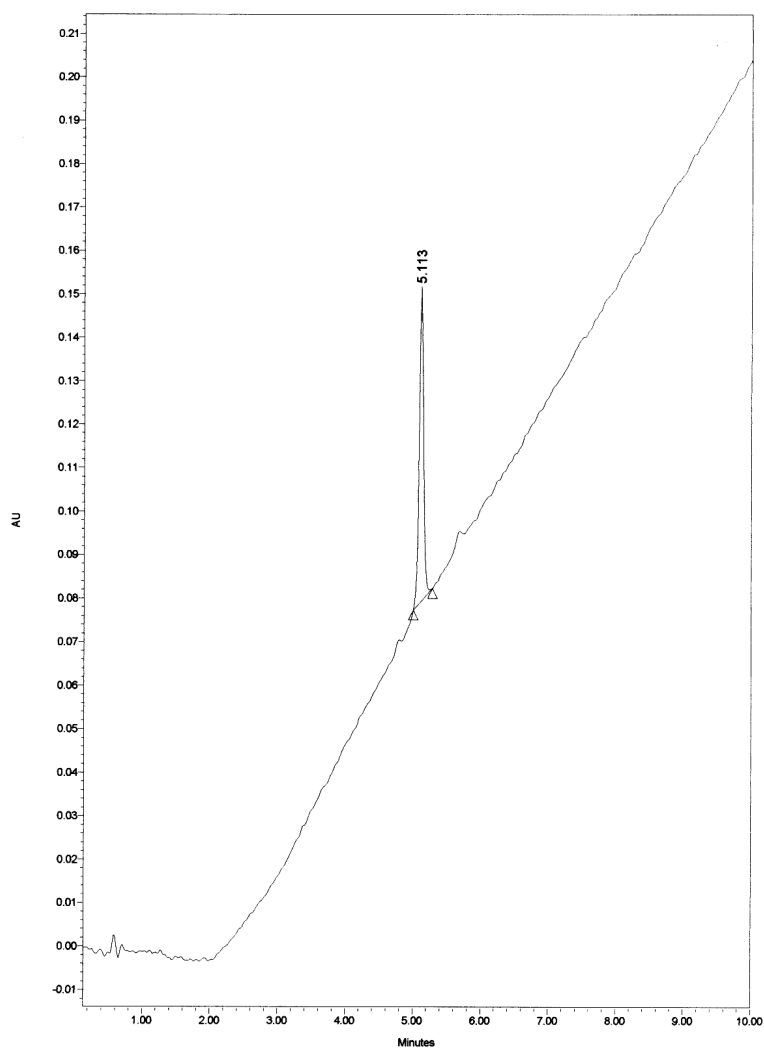
Dendrimer C<sub>2</sub>-D<sub>2</sub>

\*TOF MS: 2.451 to 3.151 min from Sample 2 (Lagnoux) of A28\_A10\_12\_A.wiff  
a=3.55534542703044650e-004, b=6.63729501778652780e+000

ESI-MS positive mode

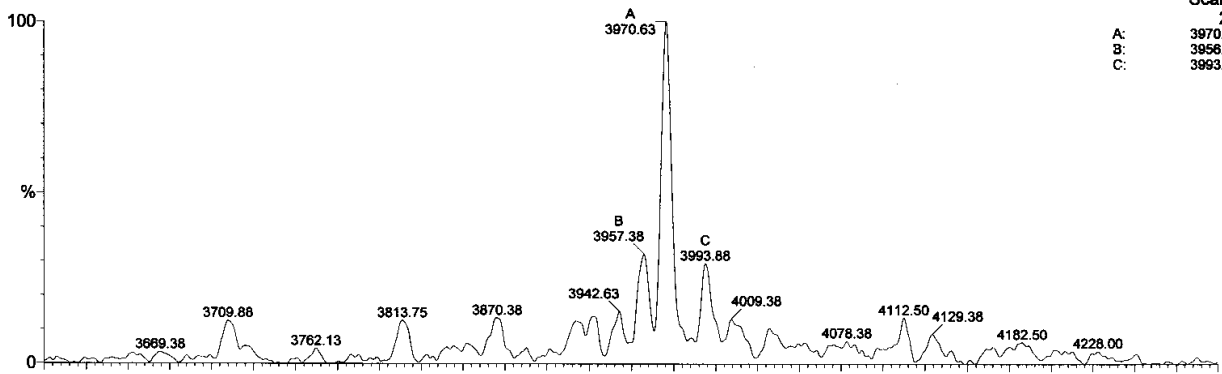
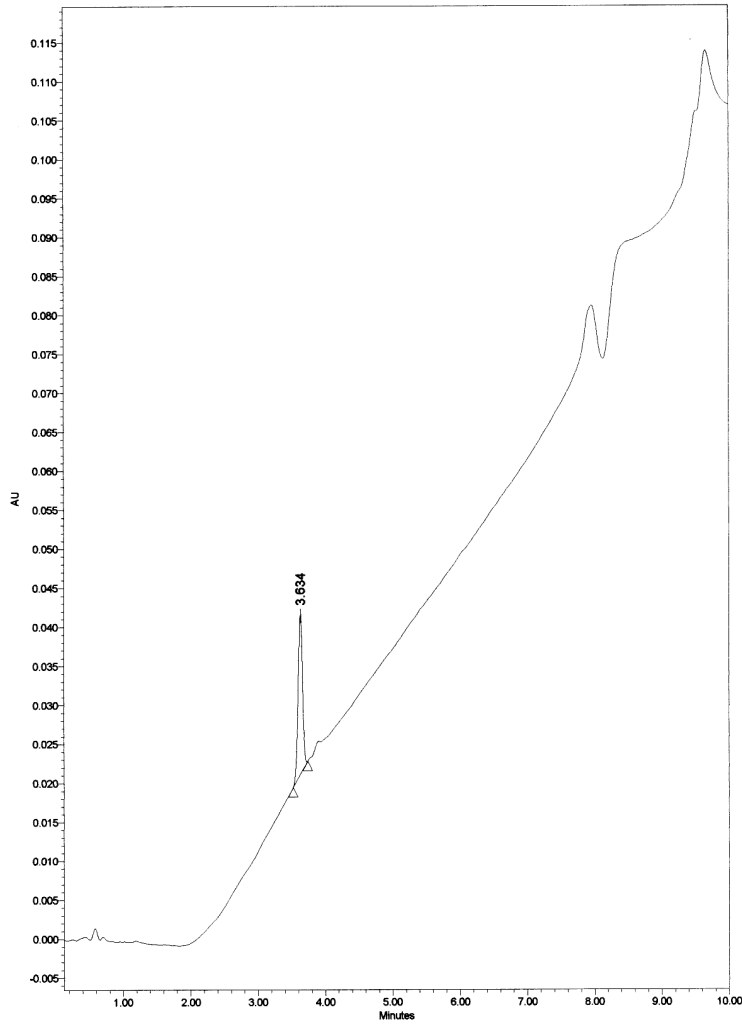
Sample dissolved in MeOH/H<sub>2</sub>O/HFo (74+25+1)



Dendrimer C<sub>2</sub>-E<sub>2</sub>

**Schaller**  
**Proteinanalytik**  
**Platform**  
Scan ES+  
2.57e<sup>1</sup>  
A: 4055.10±0.17  
B: 4077.05±0.17

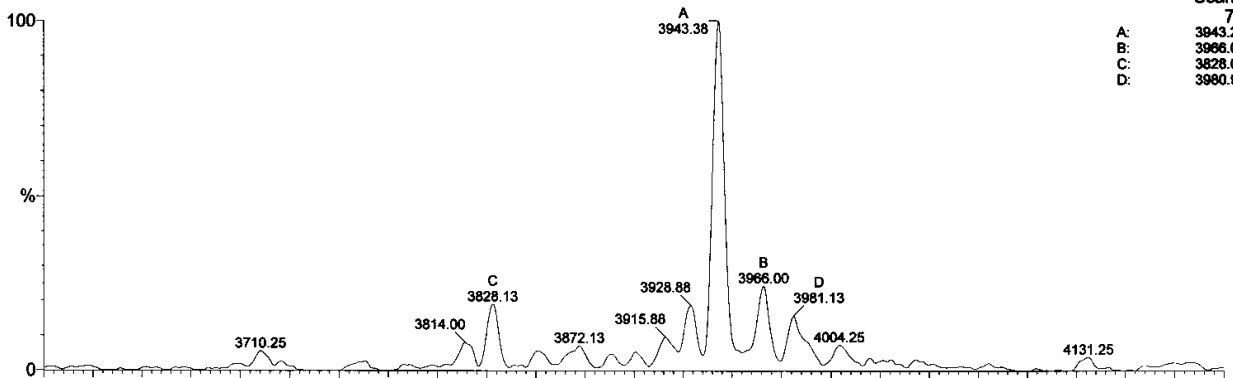
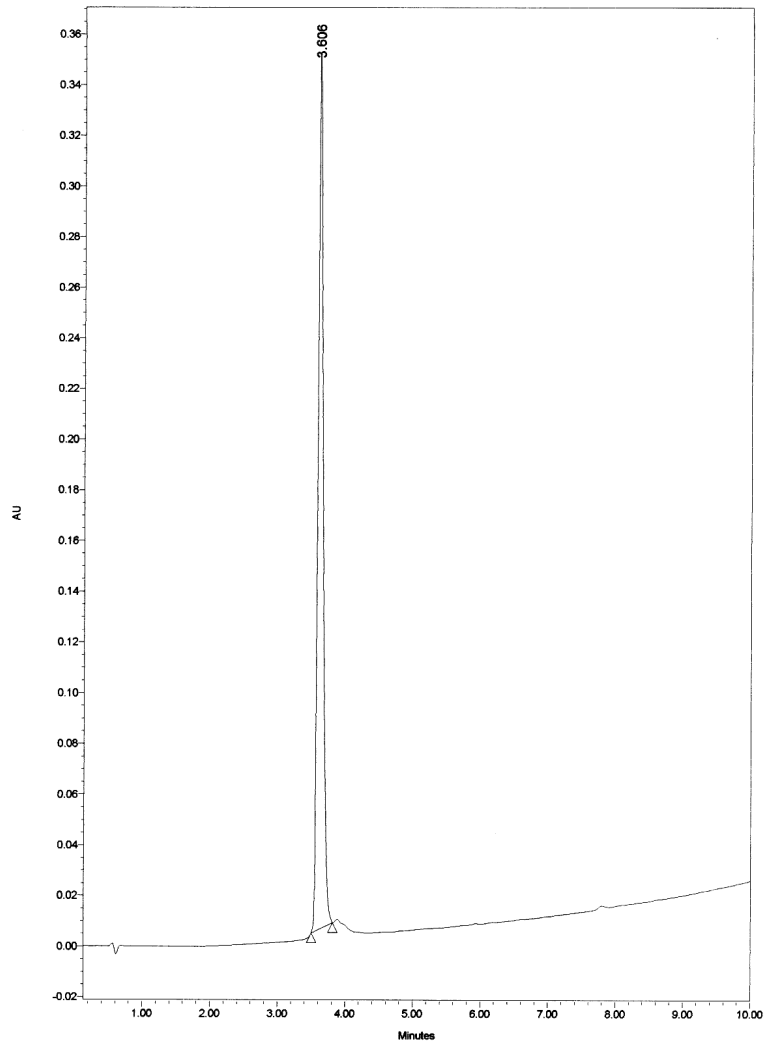
**Dendrimer D<sub>2</sub>-E<sub>2</sub>**



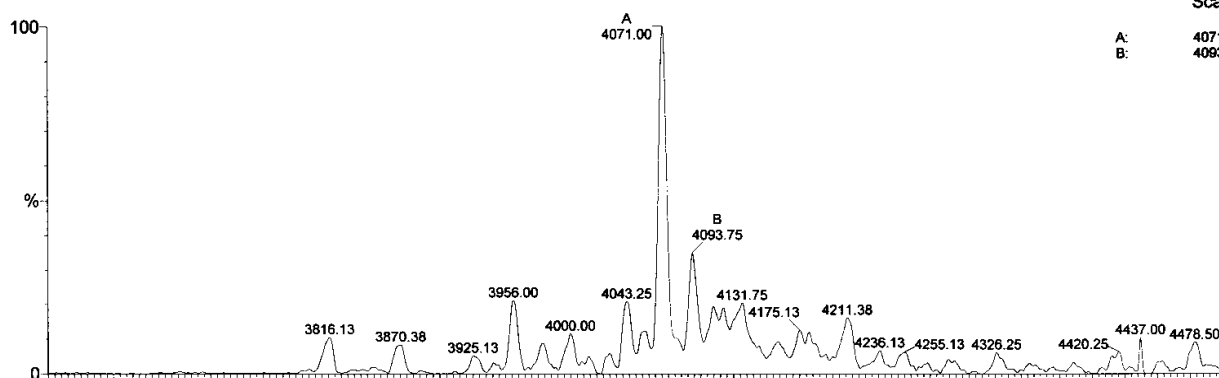
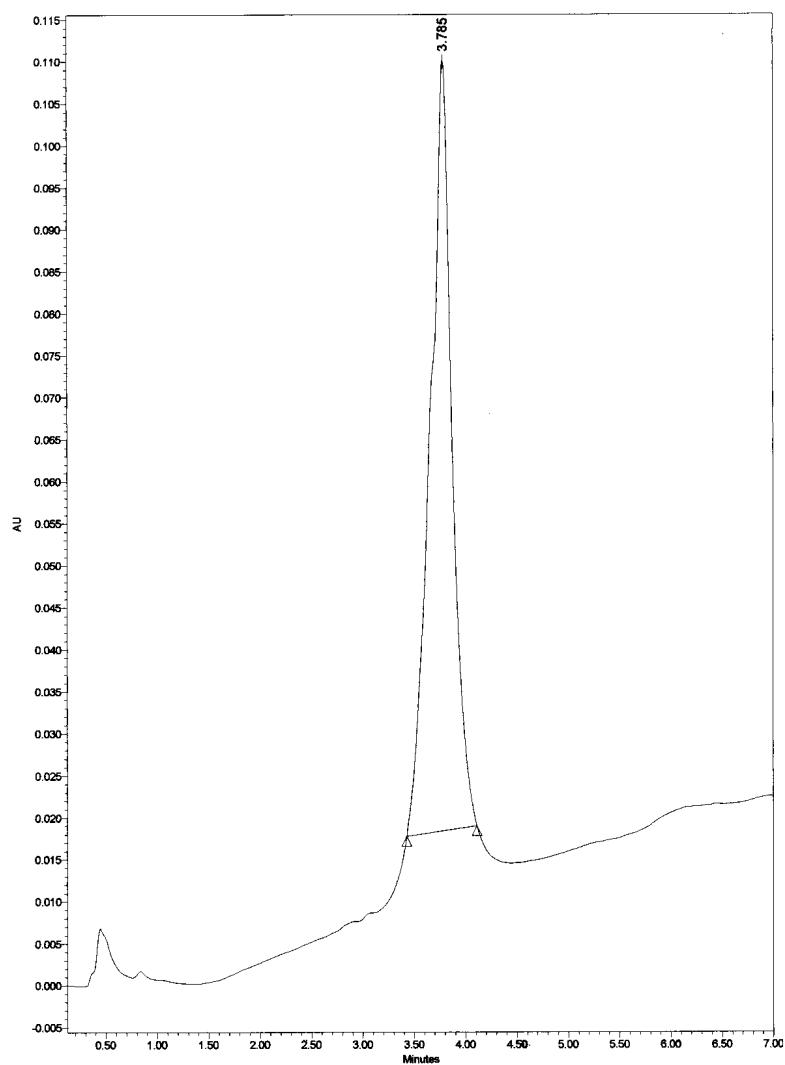
**Schaller**  
**Proteinanalytik**  
**Platform**  
 Scan ES+  
 2.75e6  
 A: 3970.58±0.32  
 B: 3956.95±0.08  
 C: 3993.95±0.36

:

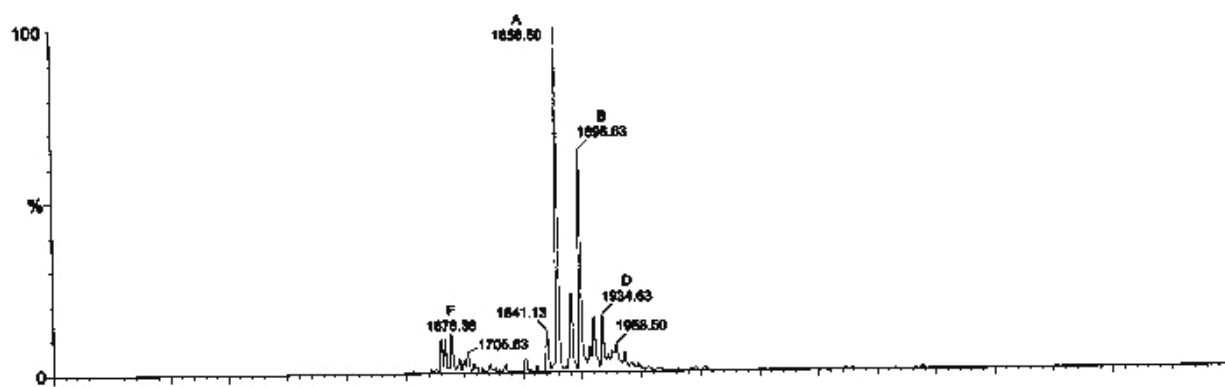
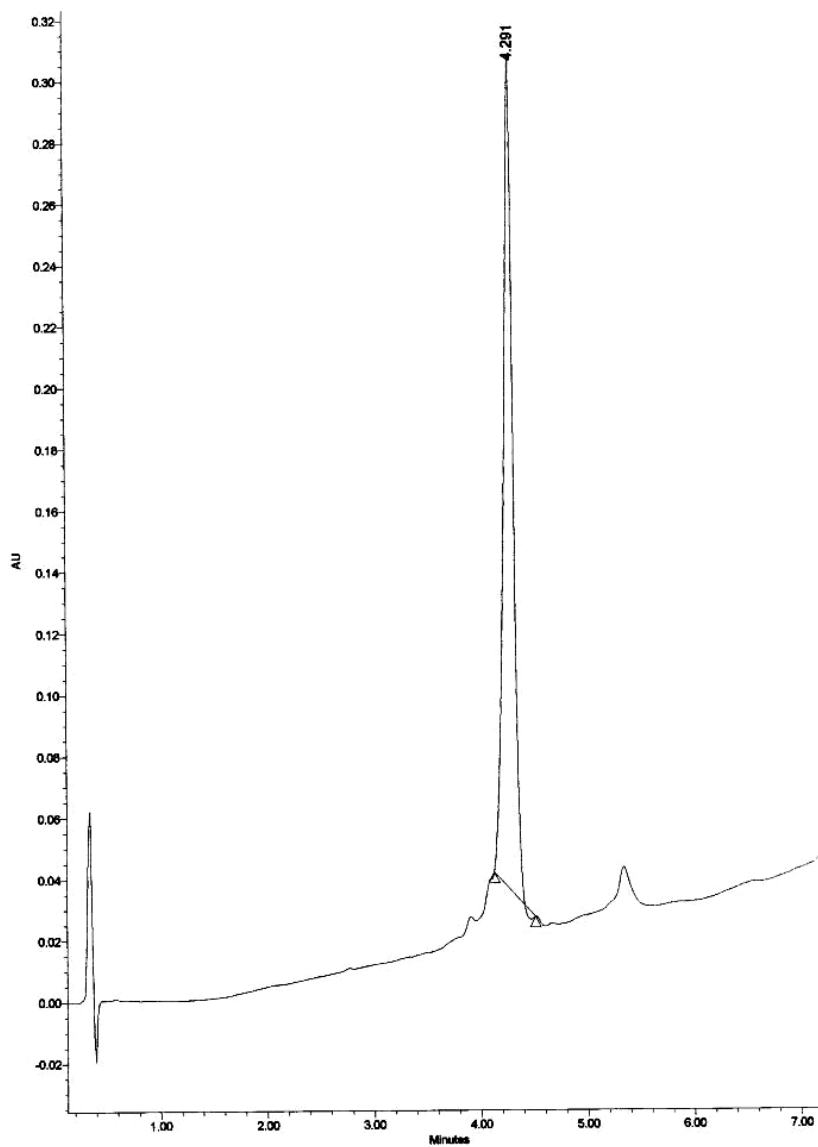
DendrimerD<sub>2</sub>-F<sub>2</sub>

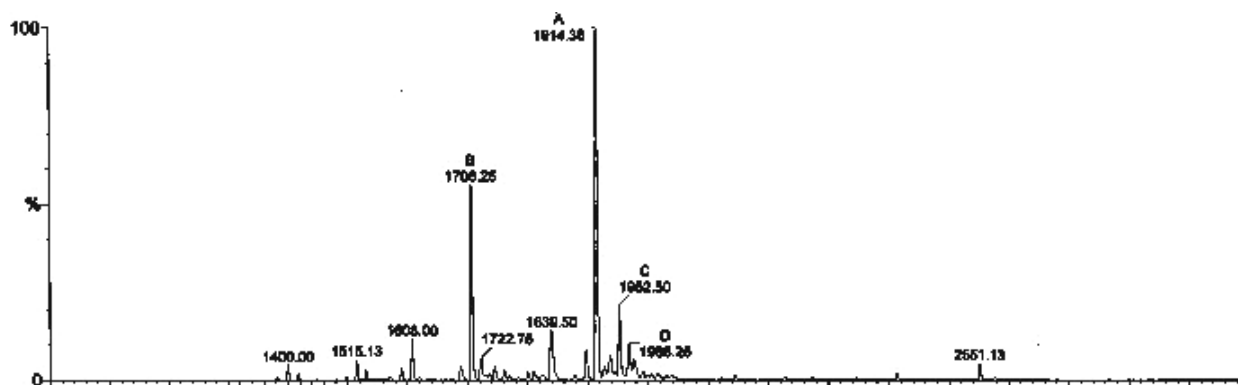
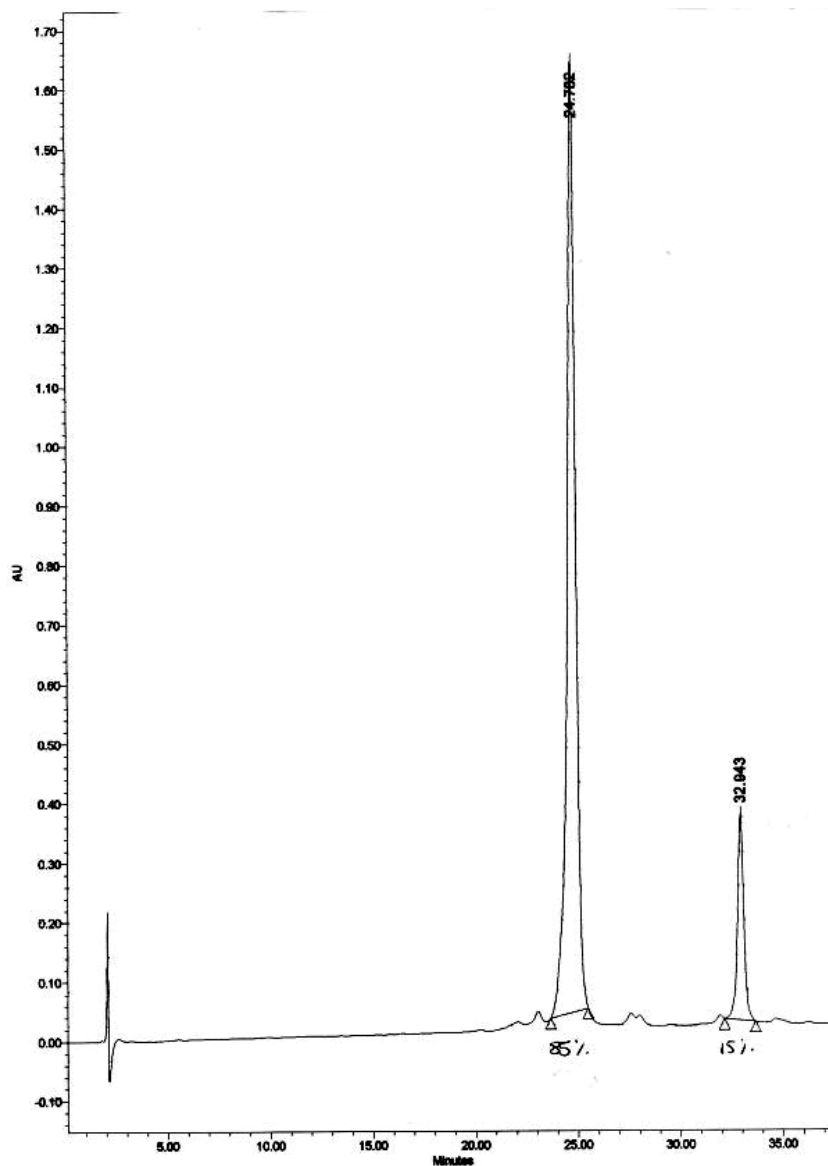


**Schaller**  
**Proteinanalytik**  
**Platform**  
Scan ES+  
7.39e6  
A: 3943.21±0.07  
B: 3966.02±0.13  
C: 3828.00±0.18  
D: 3980.90±0.13

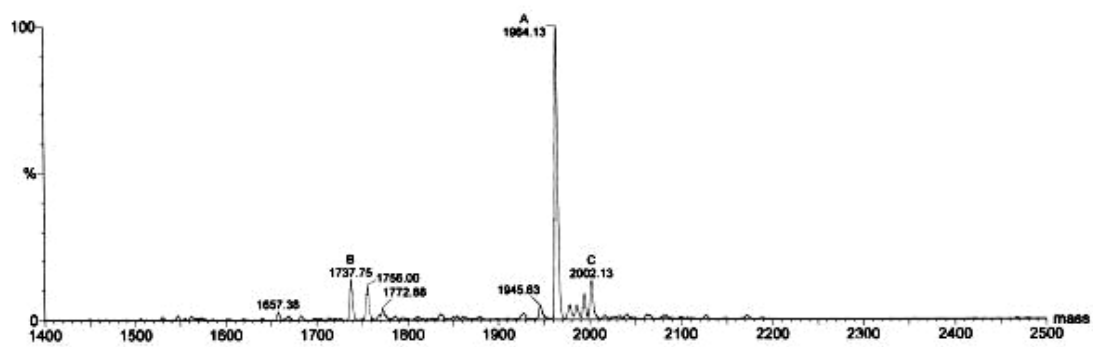
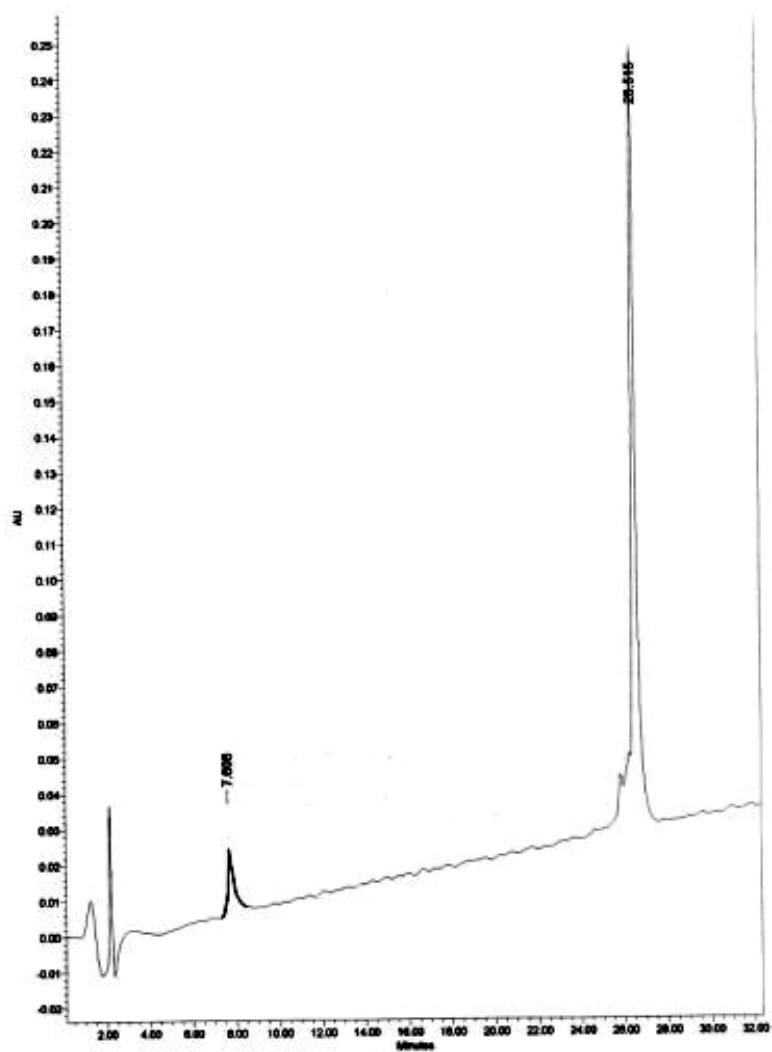
DendrimerE<sub>2</sub>-F<sub>2</sub>

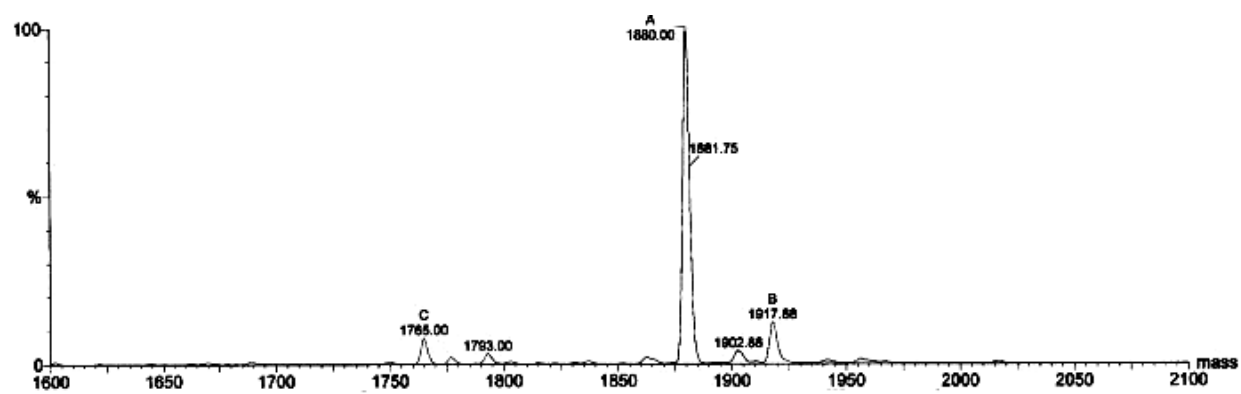
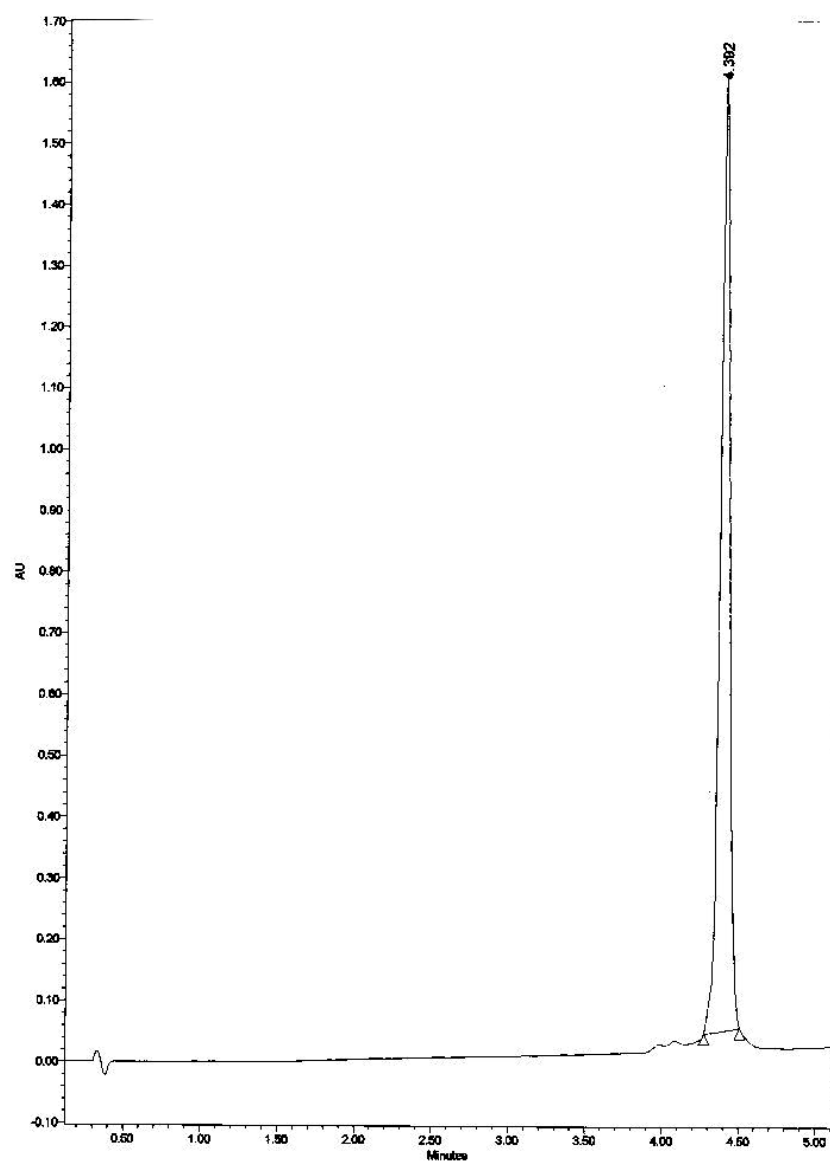
**Schaller**  
**Proteinanalytik**  
**Platform**  
Scan ES+  
5.17e6  
A: 4071.11±0.30  
B: 4093.62±0.02

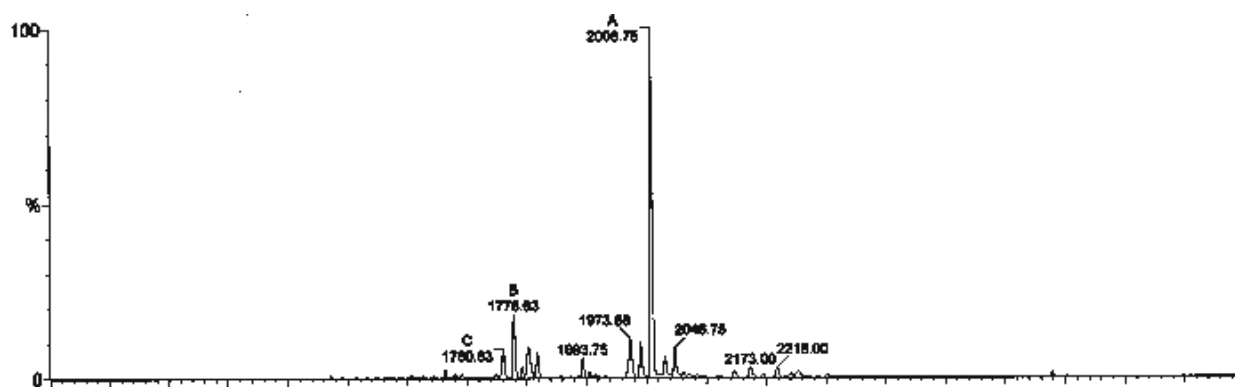
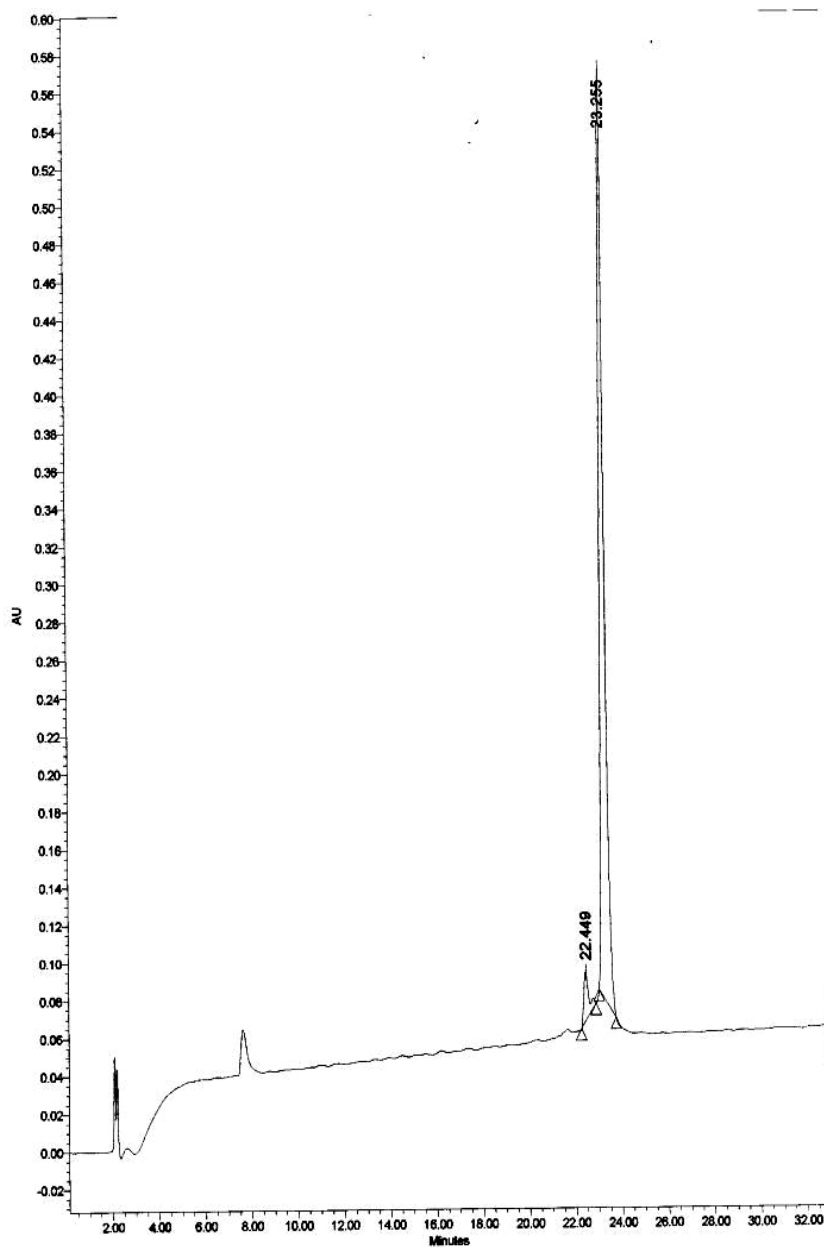
Dendrimer A<sub>3</sub>

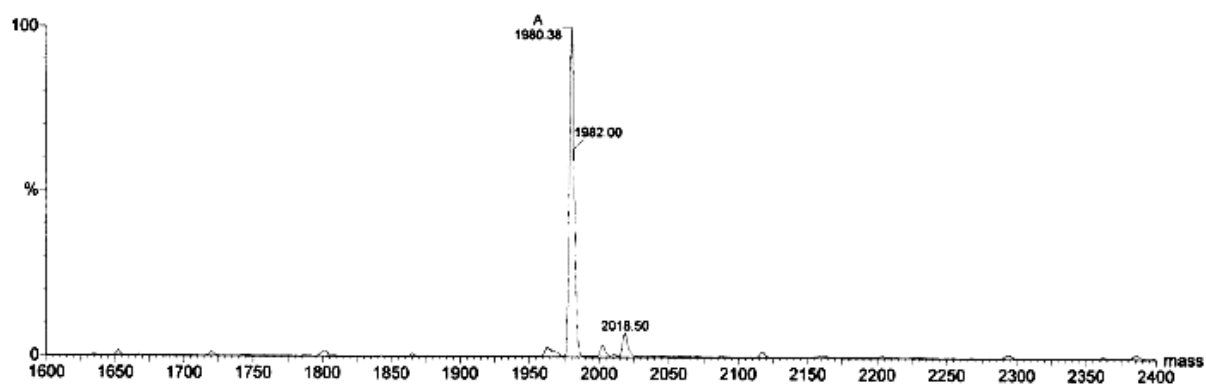
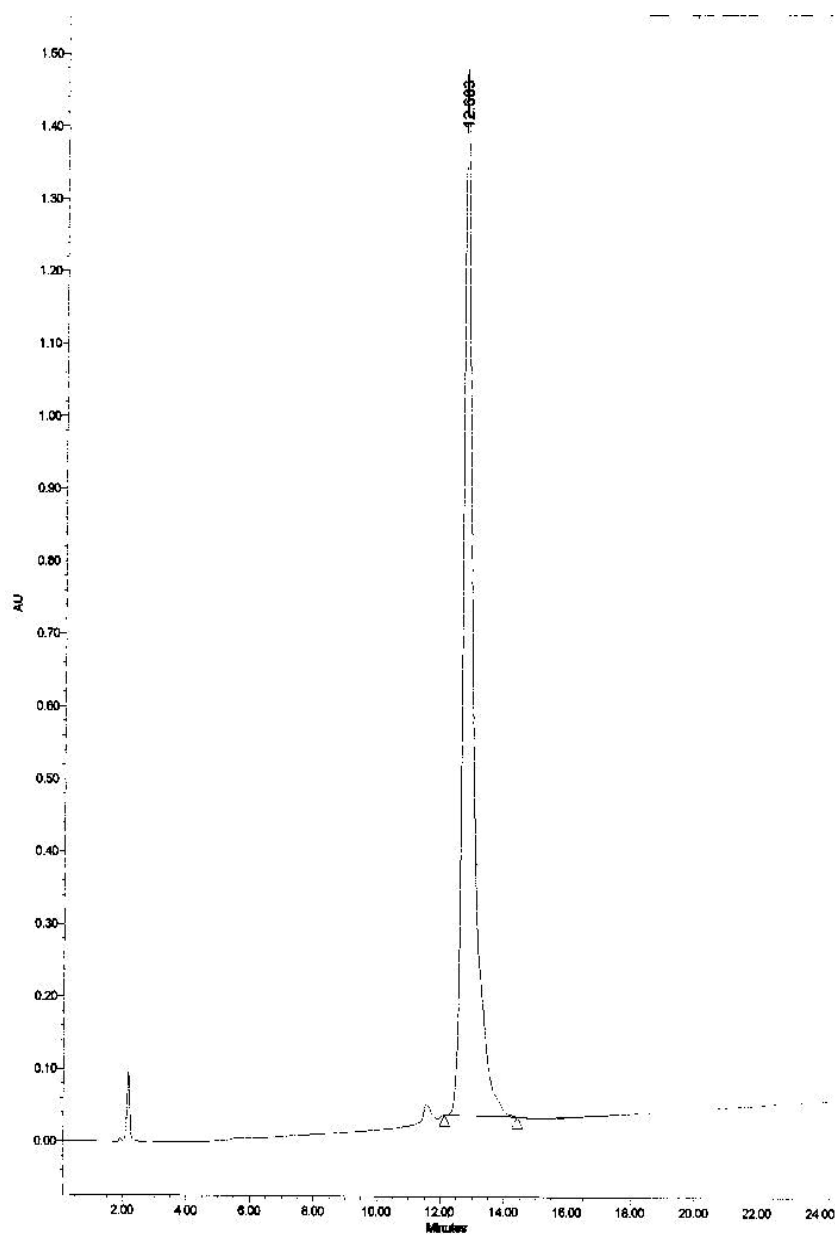
Dendrimer B<sub>3</sub>

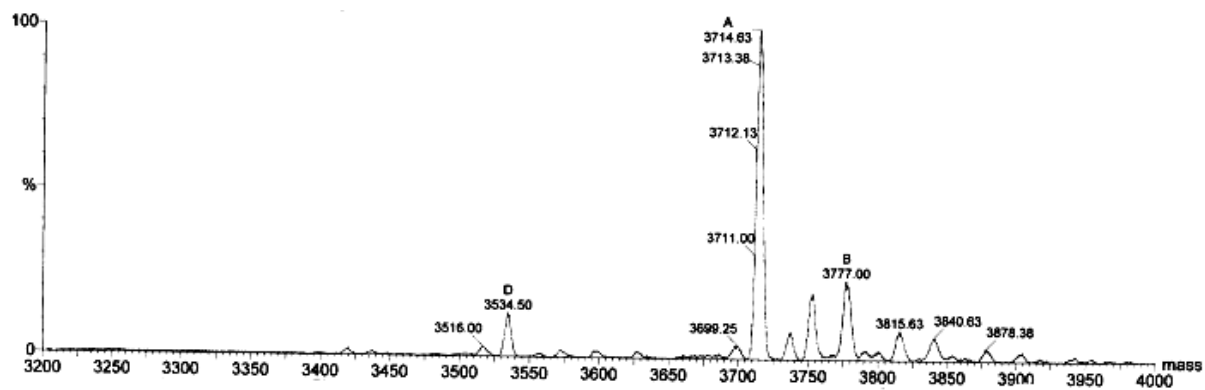
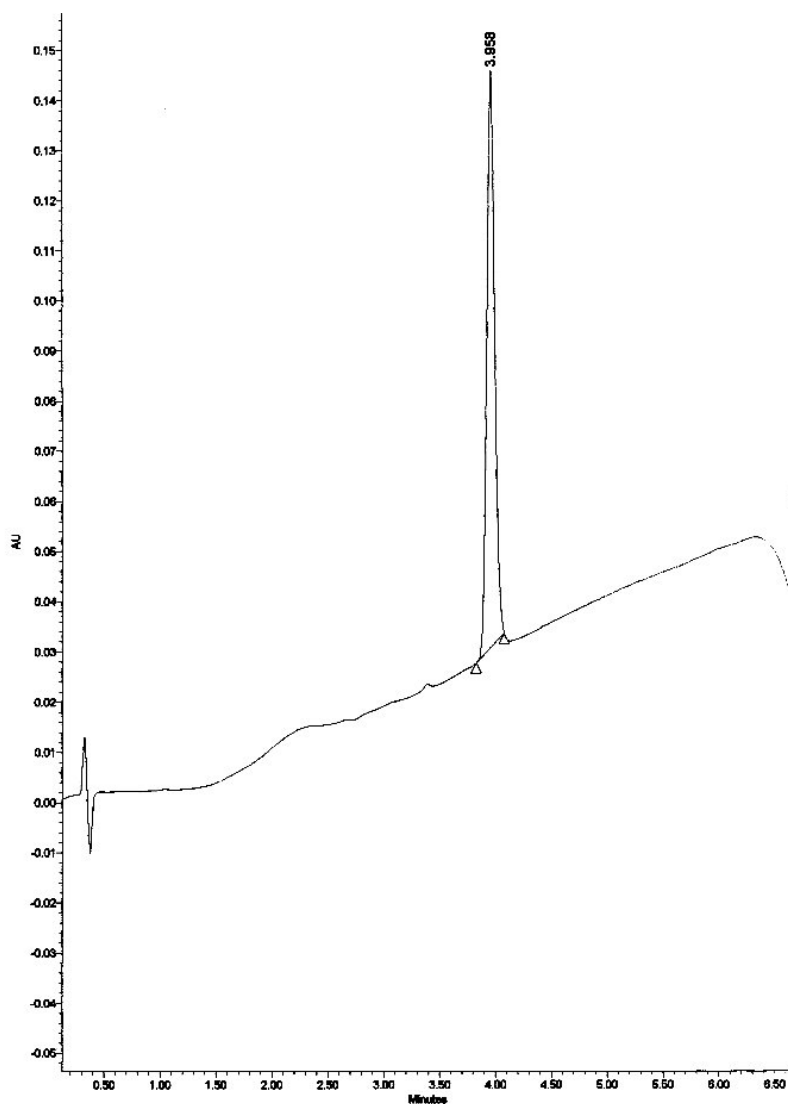


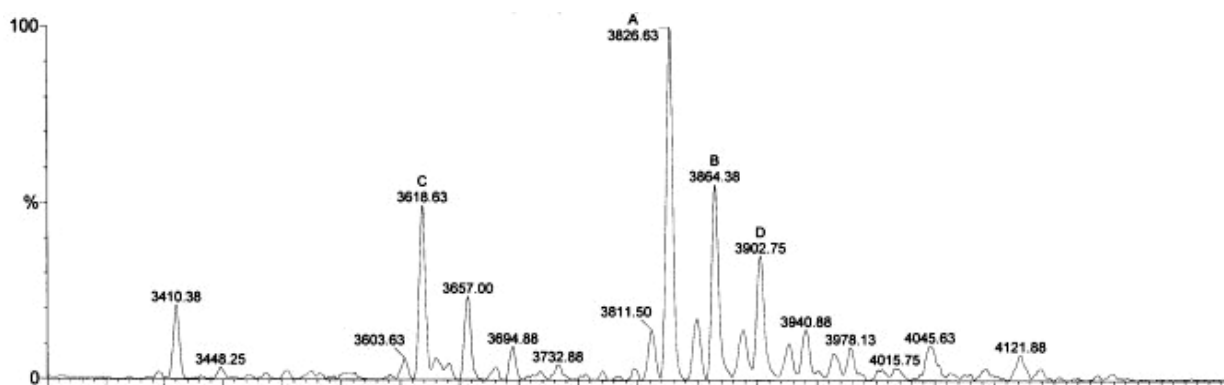
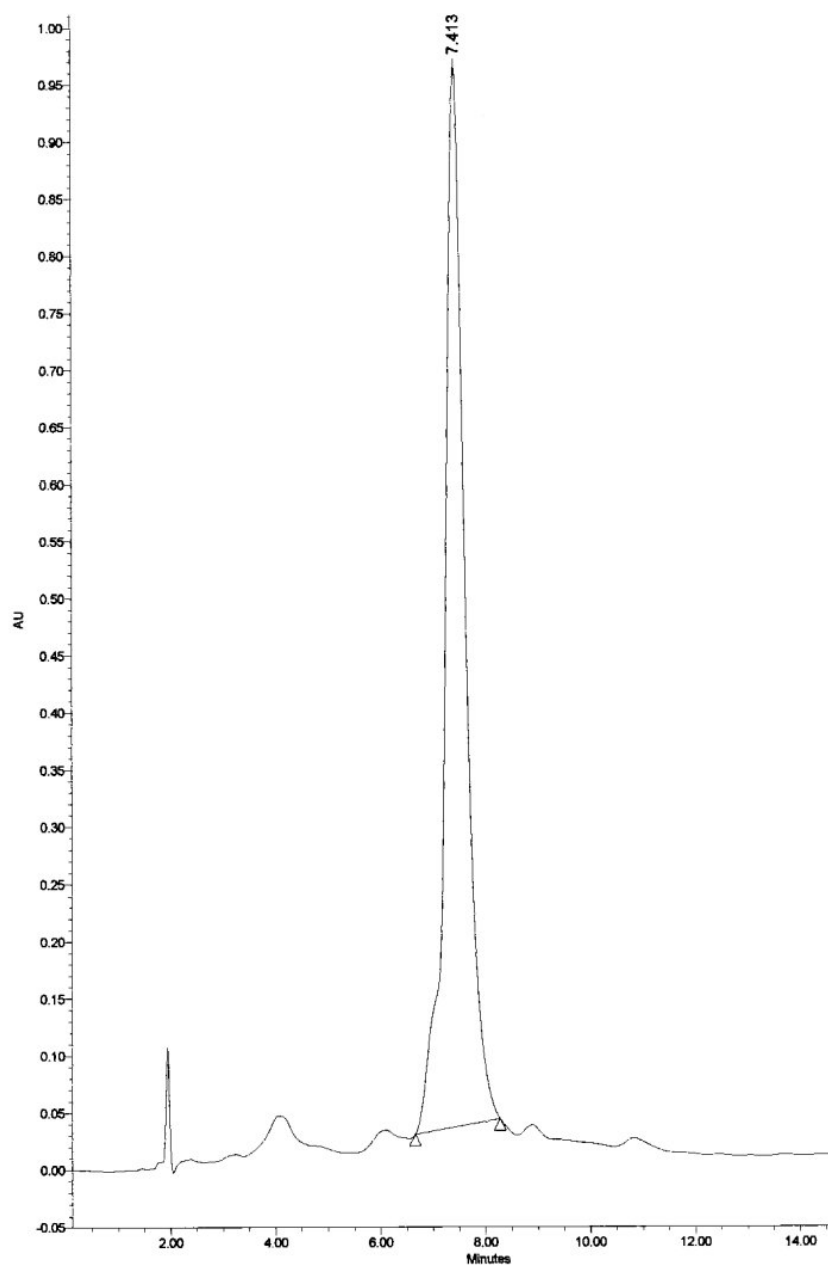
Dendrimer C<sub>3</sub>

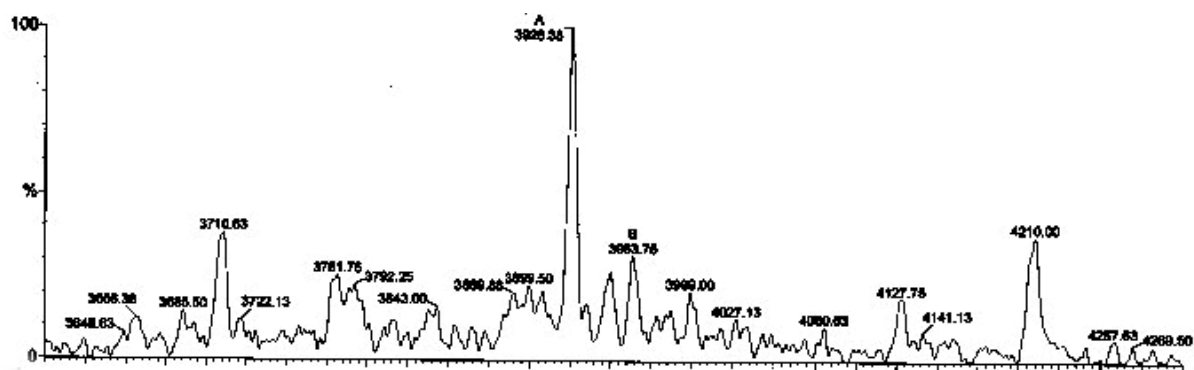
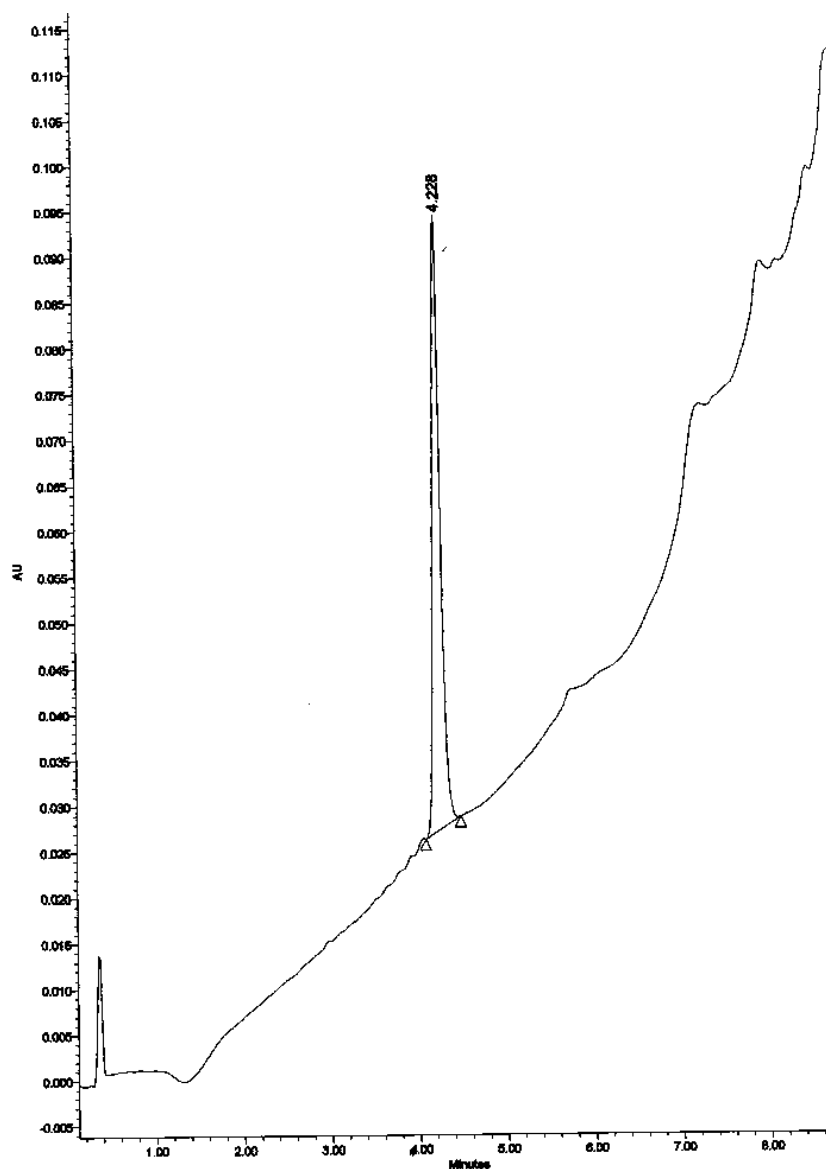
DendrimerD<sub>3</sub>

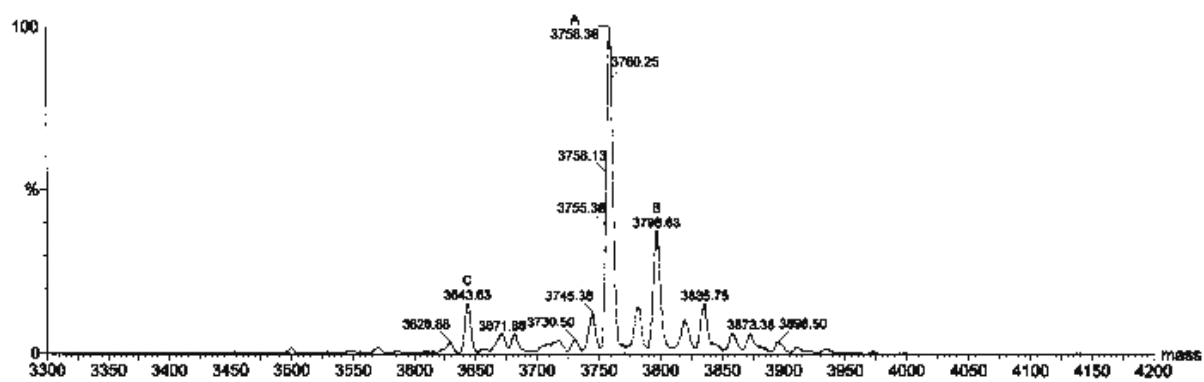
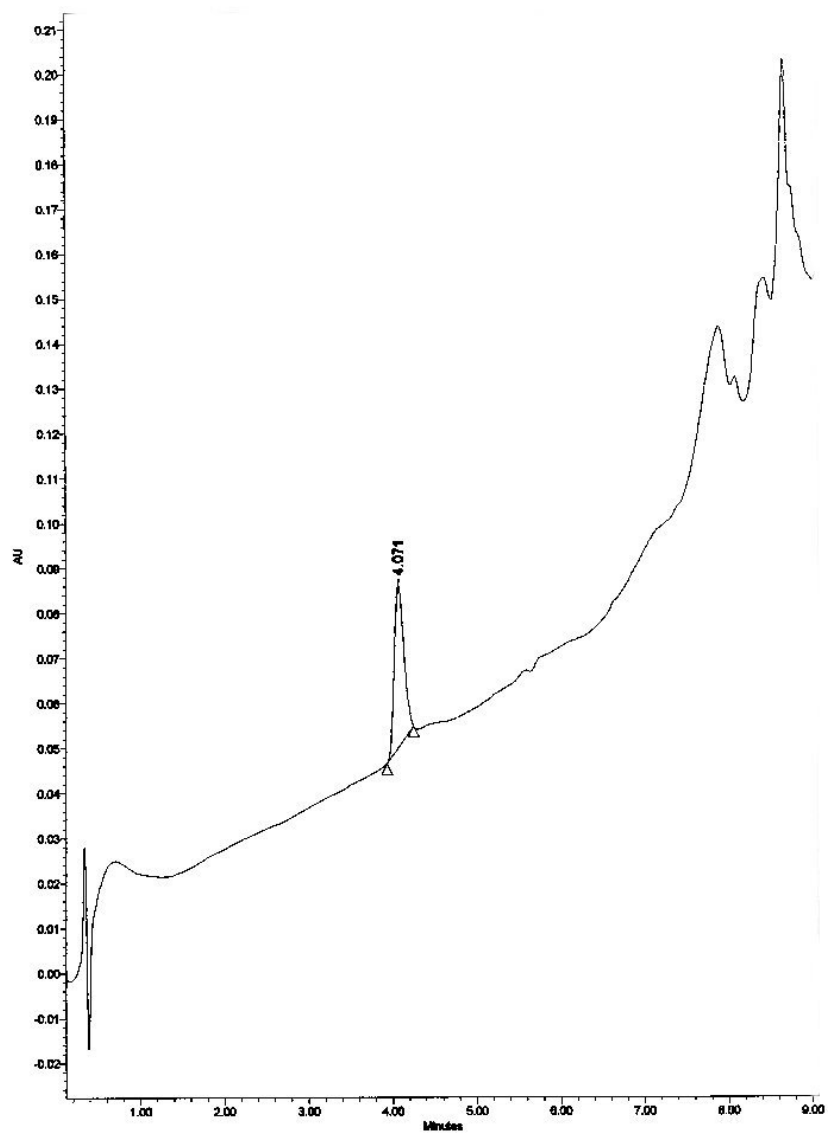
DendrimerE<sub>3</sub>

DendrimerF<sub>3</sub>

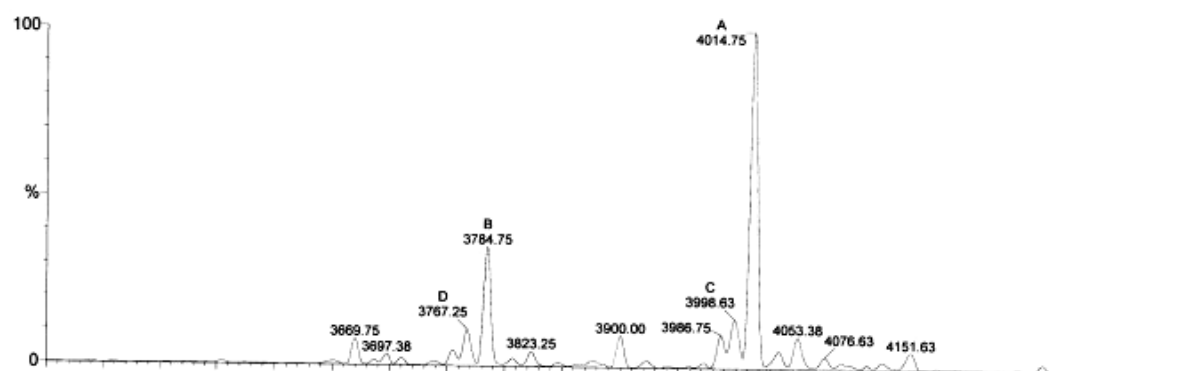
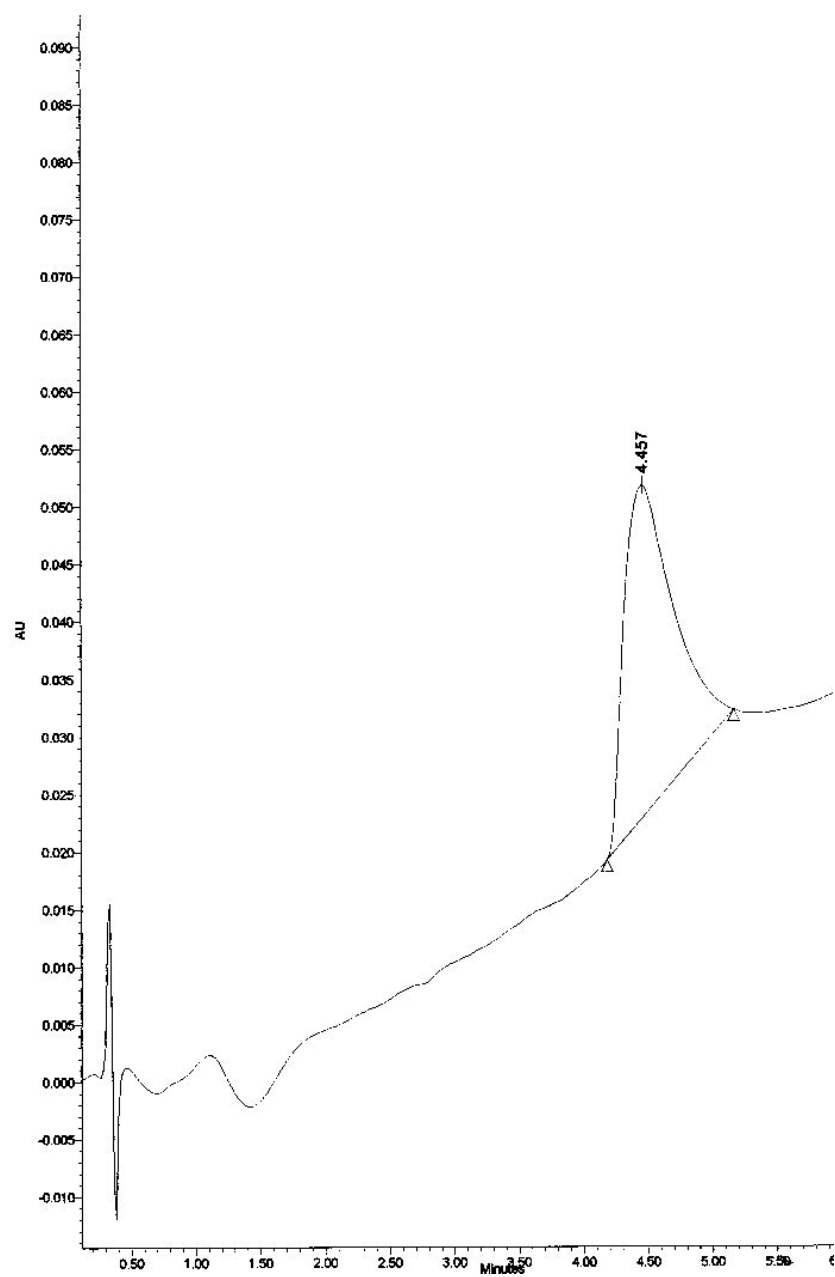
Dendrimer A<sub>3</sub>A<sub>3</sub>

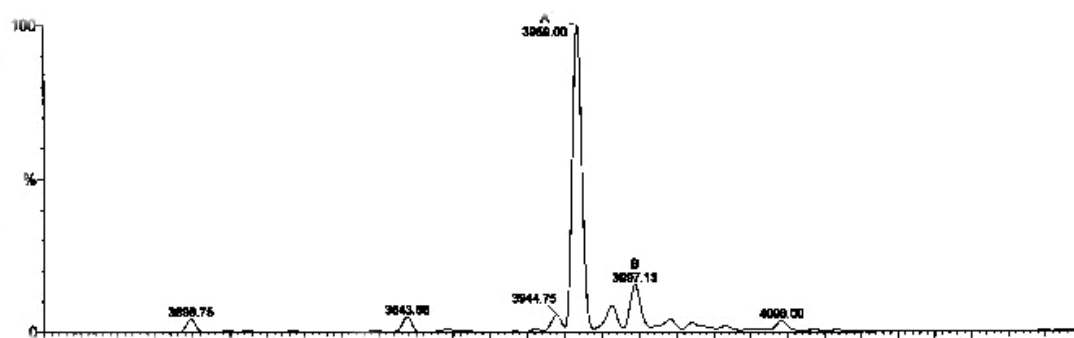
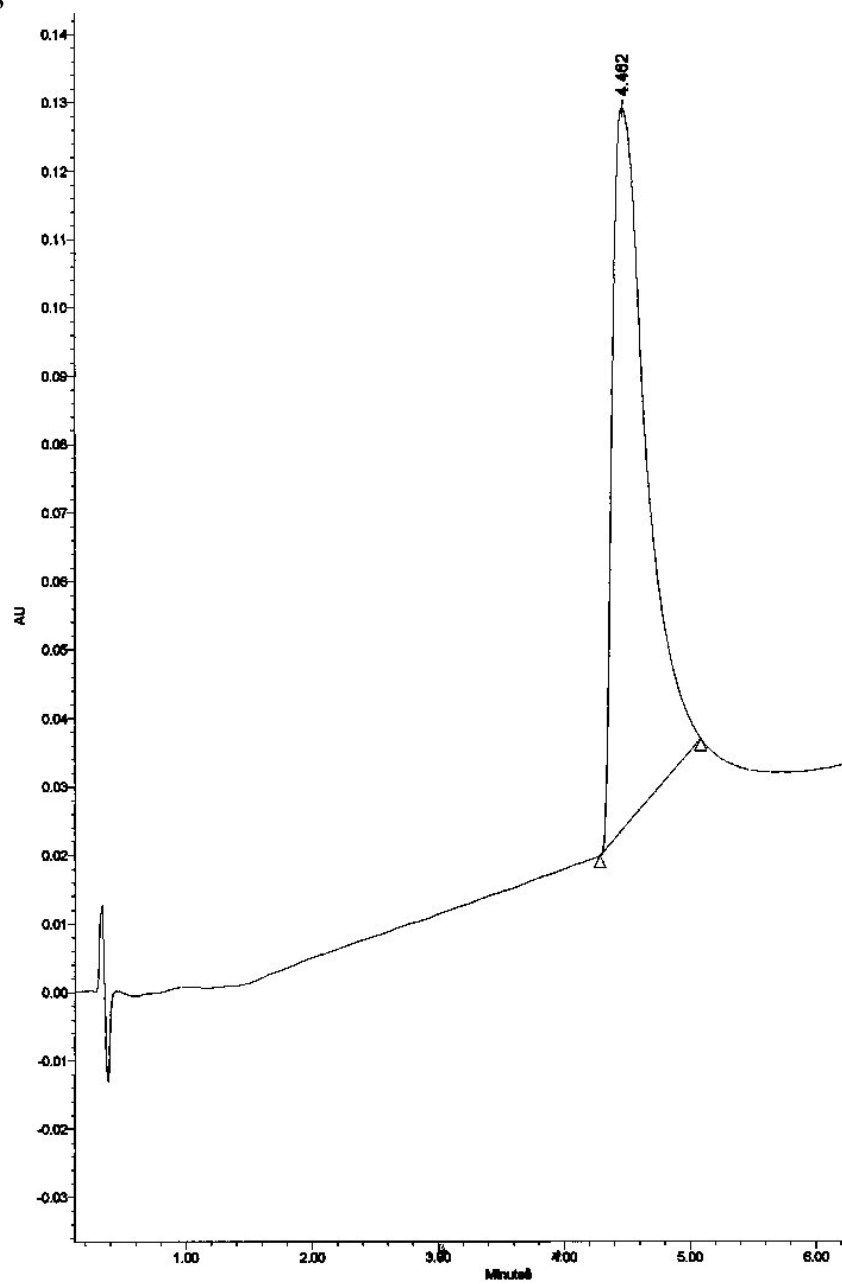
Dendrimer B<sub>3</sub>B<sub>3</sub>

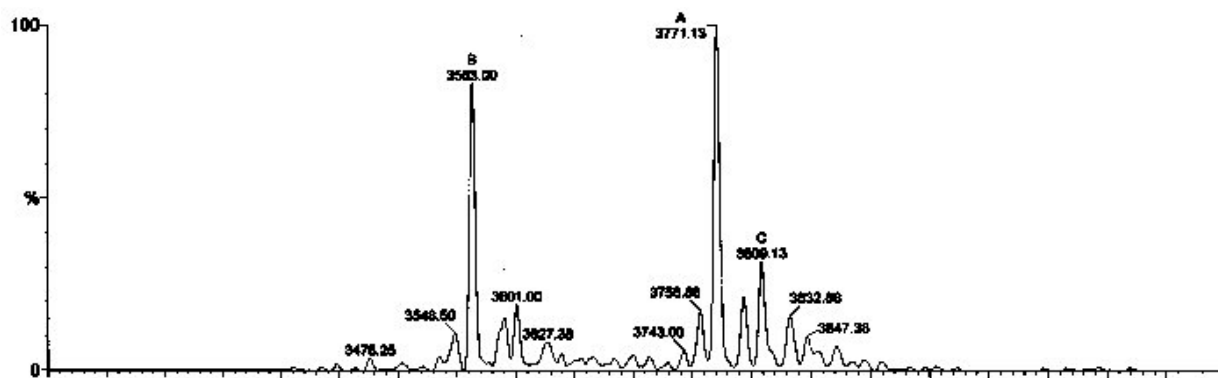
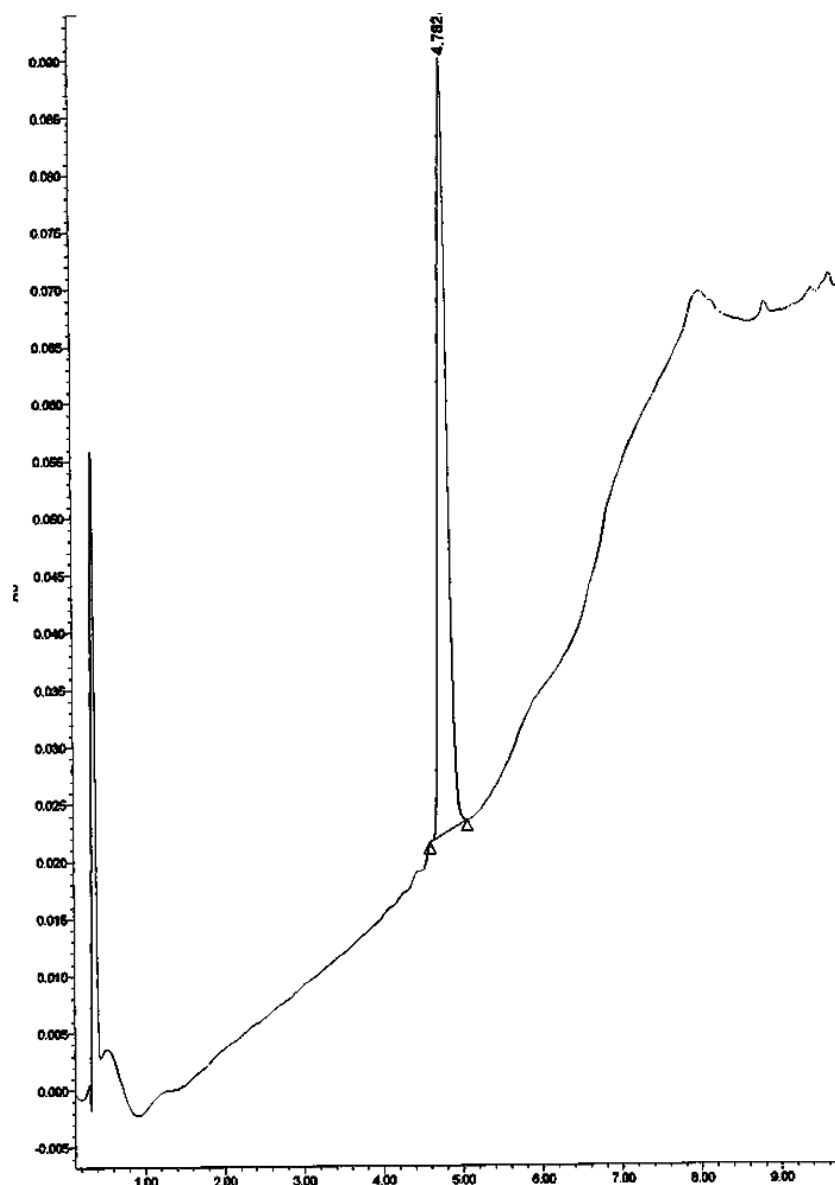
Dendrimer C<sub>3</sub>C<sub>3</sub>

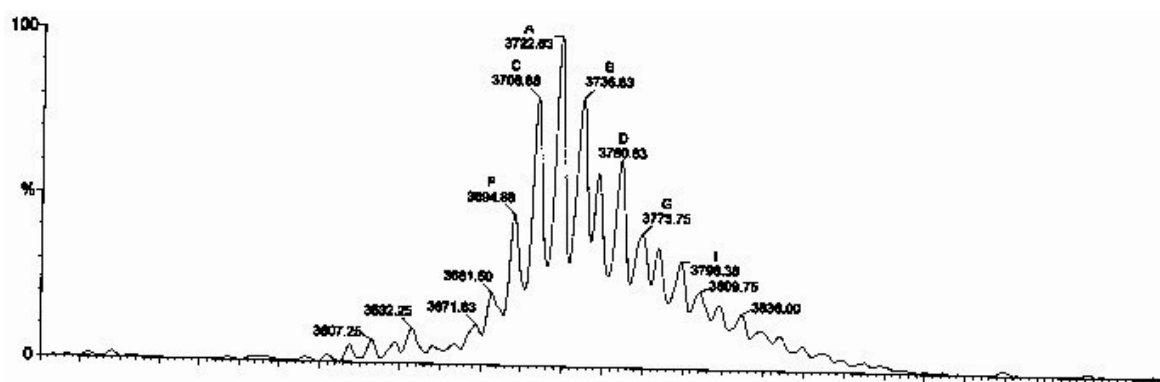
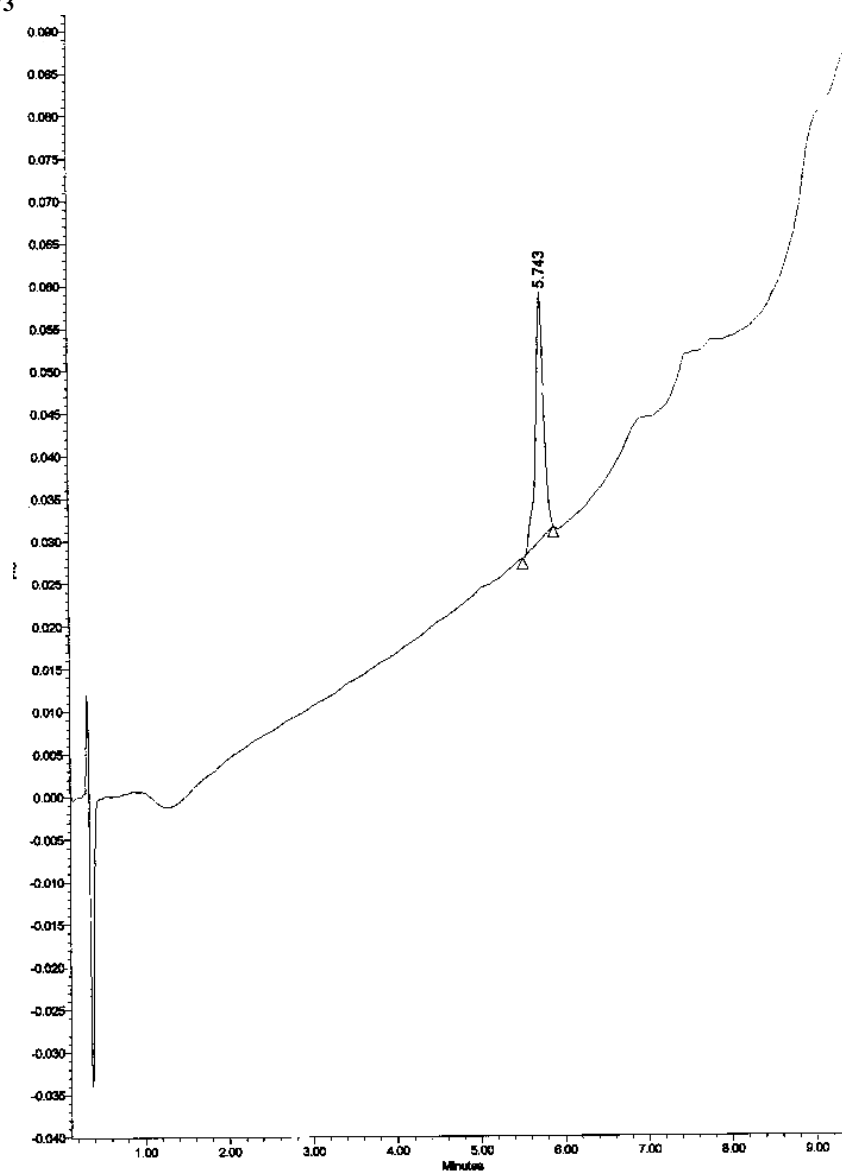
Dendrimer D<sub>3</sub>D<sub>3</sub>

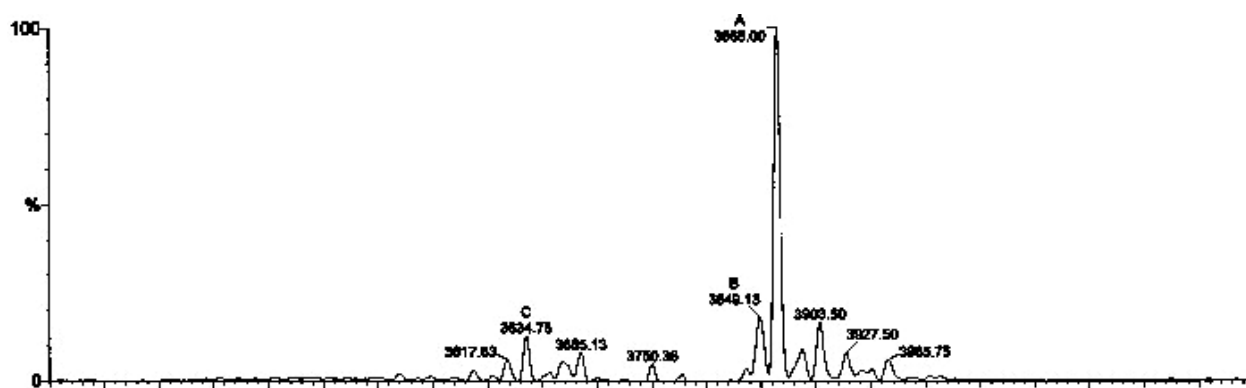
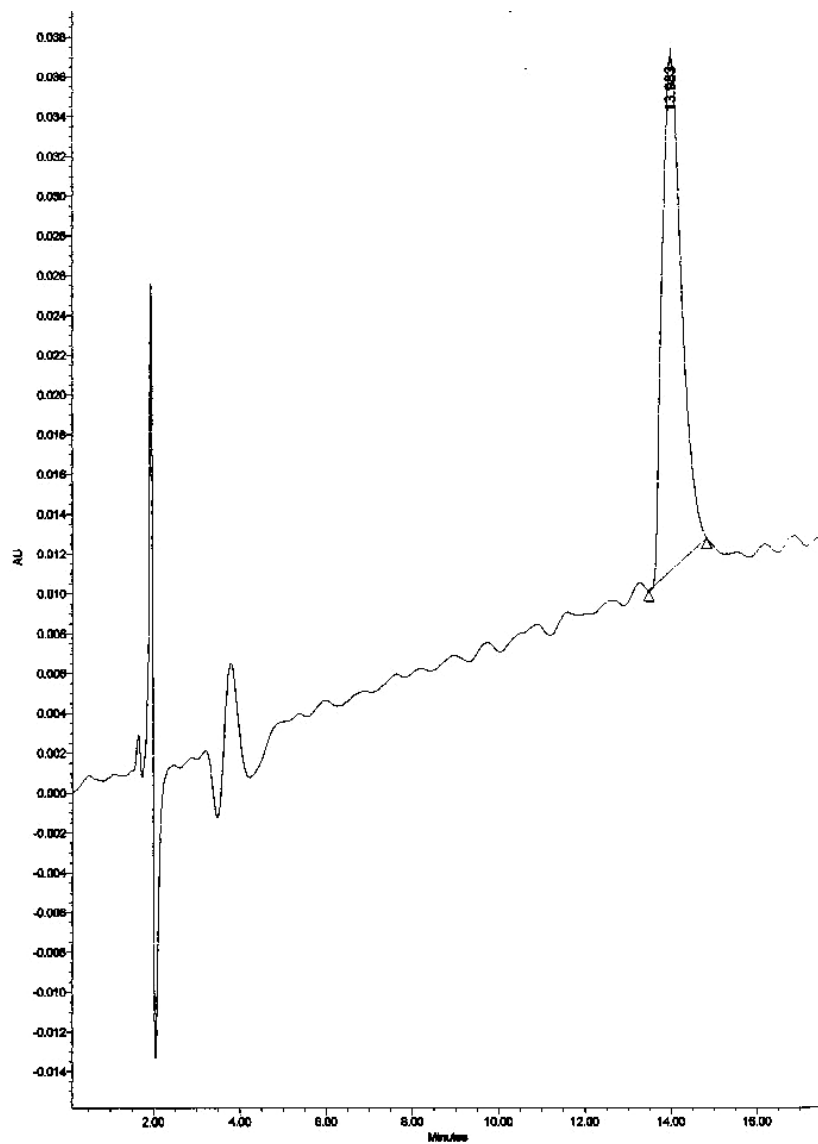


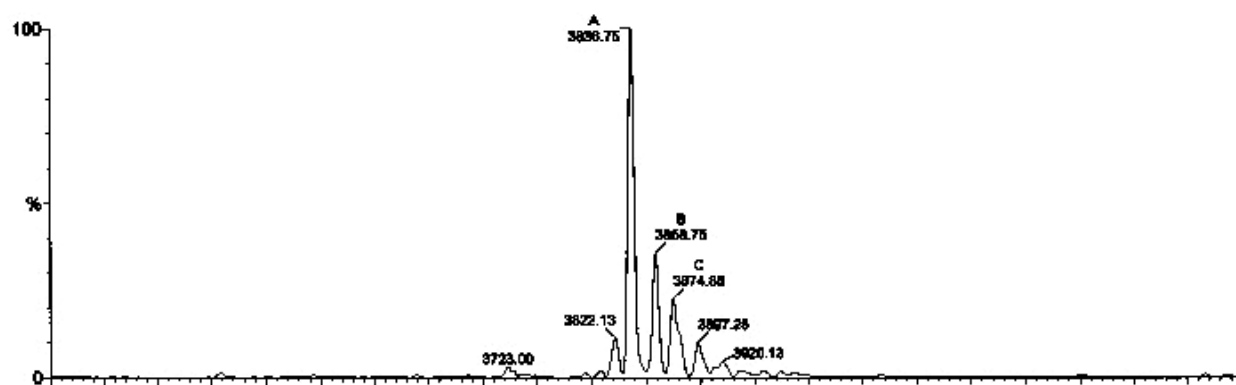
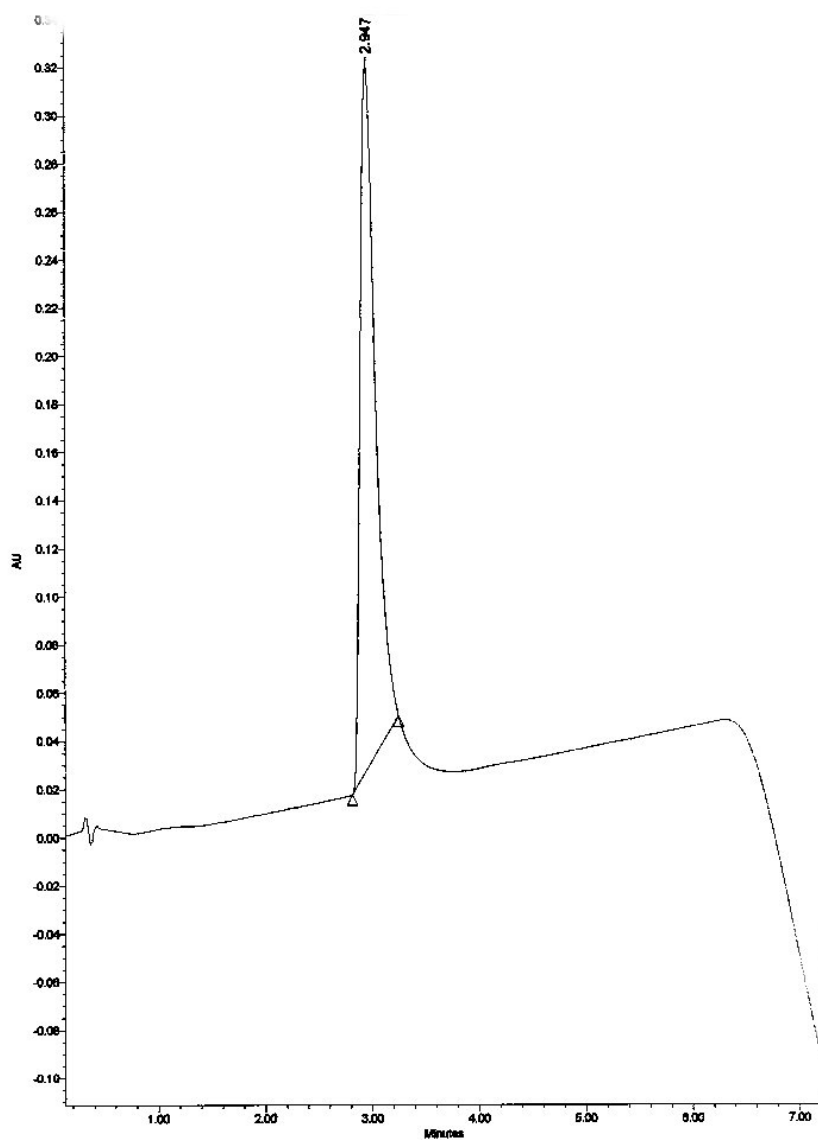
DendrimerE<sub>3</sub>E<sub>3</sub>

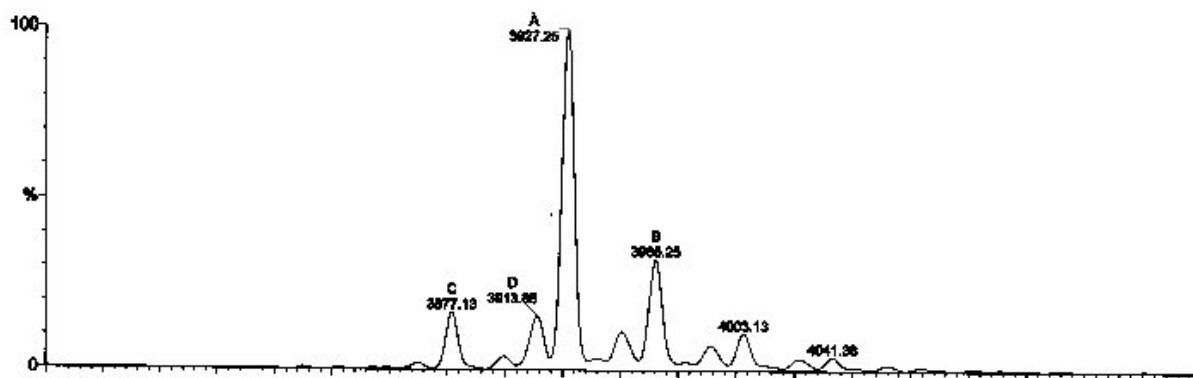
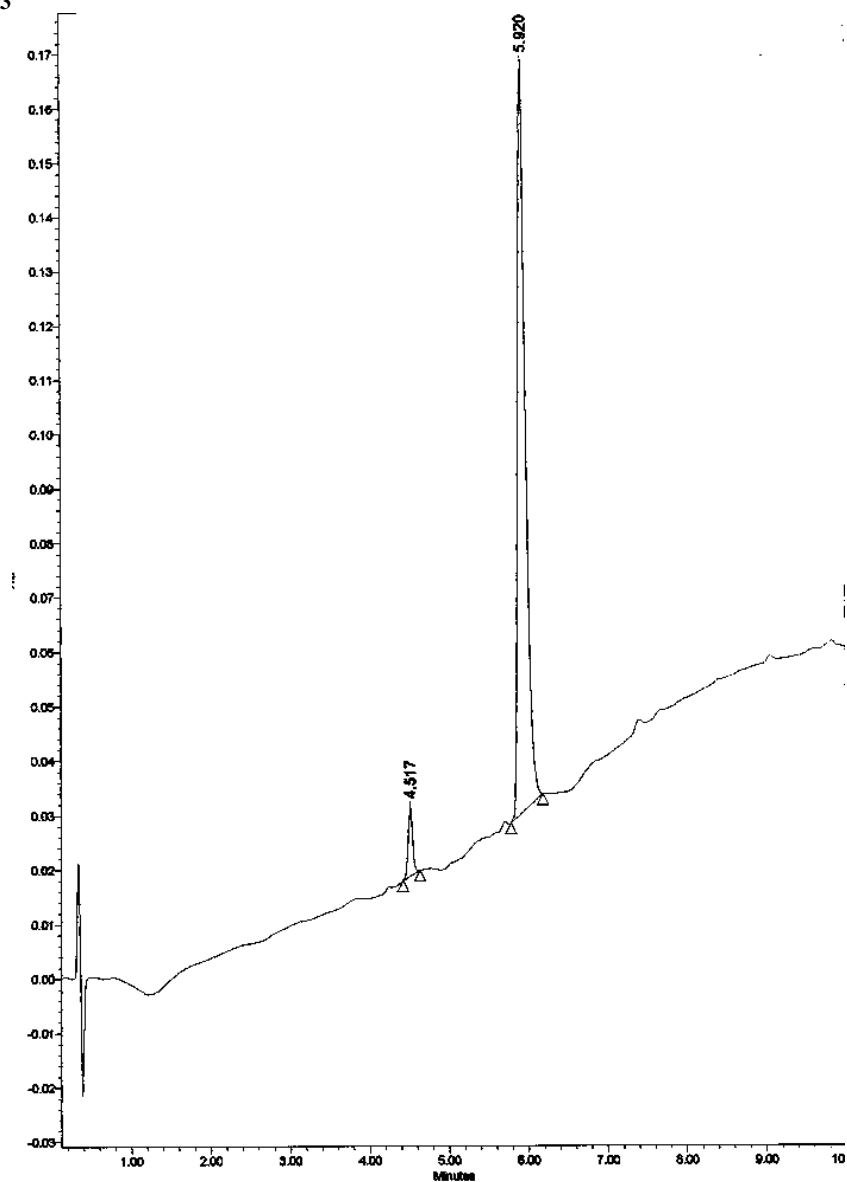
DendrimerF<sub>3</sub>F<sub>3</sub>

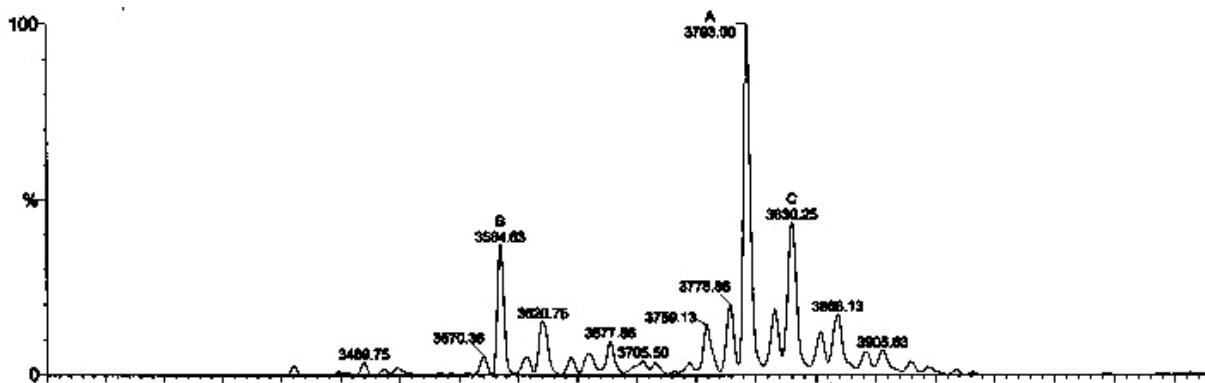
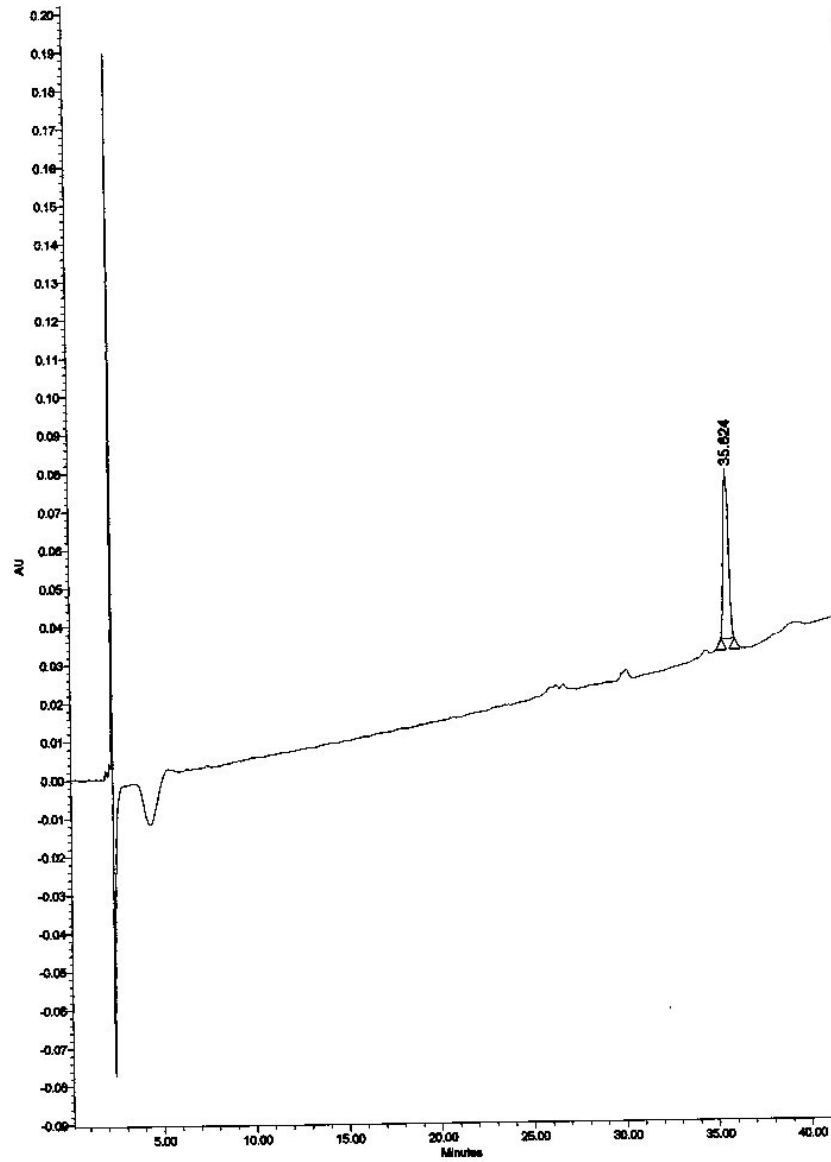
Dendrimer A<sub>3</sub>B<sub>3</sub>

Dendrimer A<sub>3</sub>D<sub>3</sub>

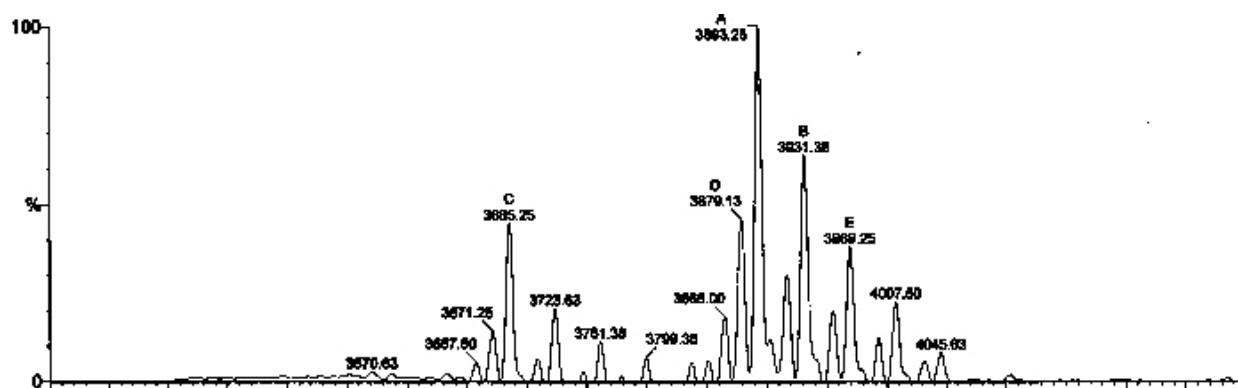
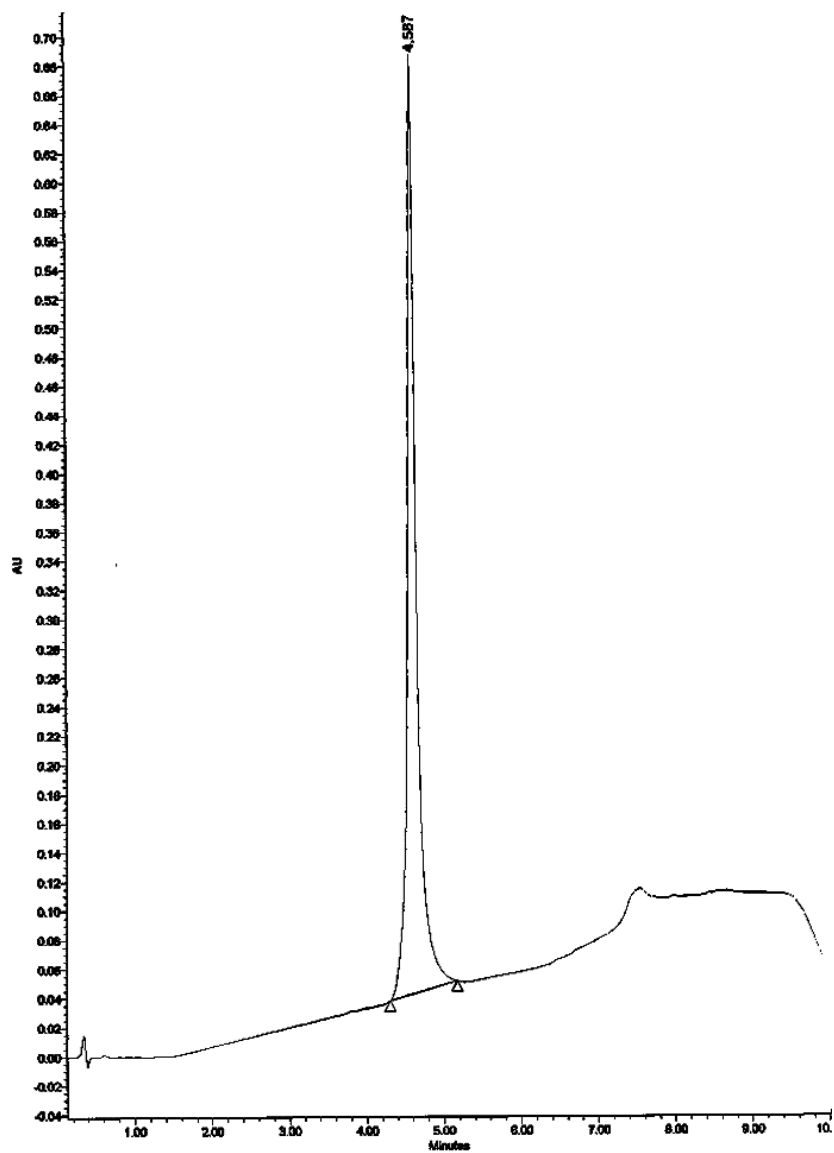
Dendrimer A<sub>3</sub>E<sub>3</sub>

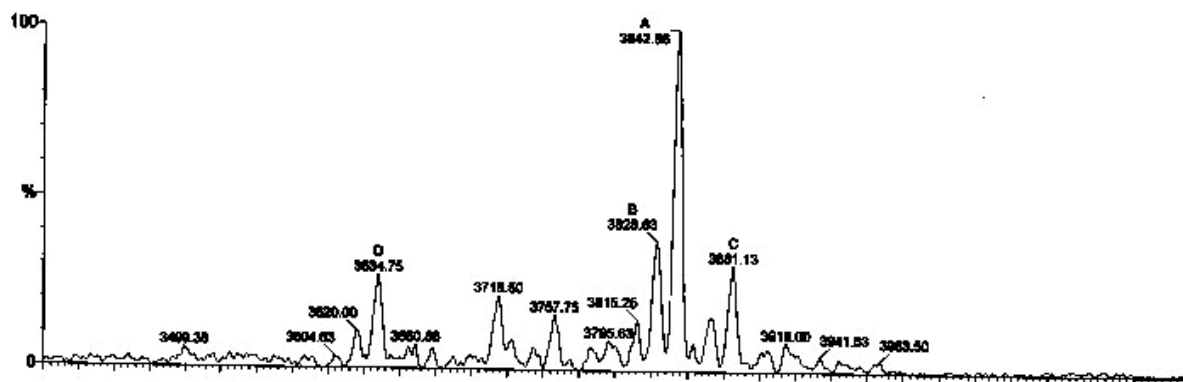
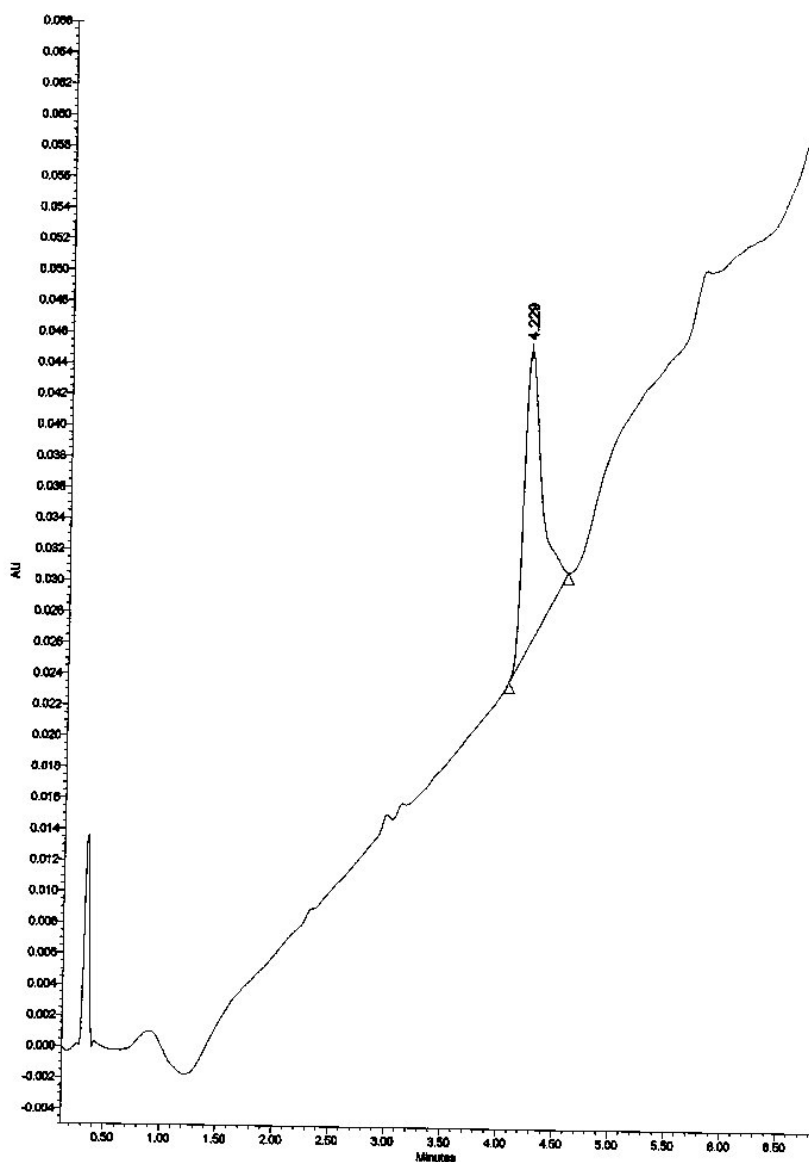
Dendrimer A<sub>3</sub>F<sub>3</sub>

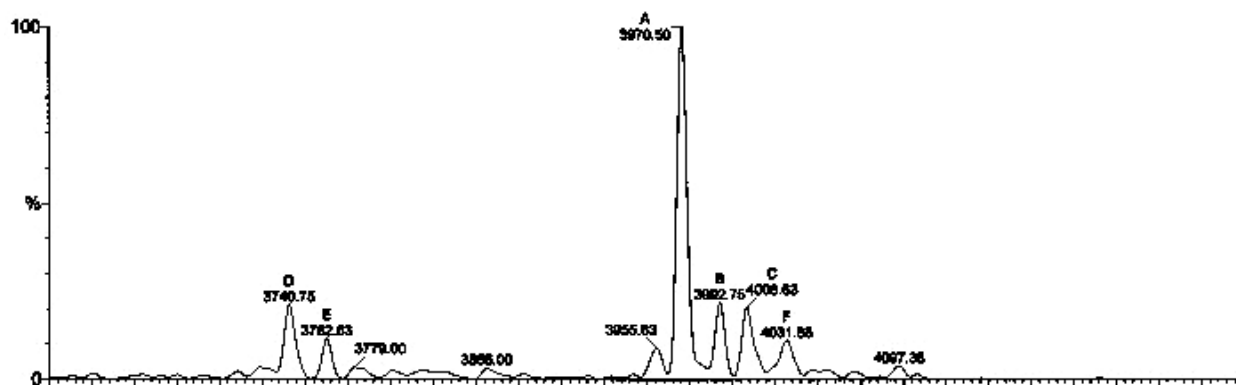
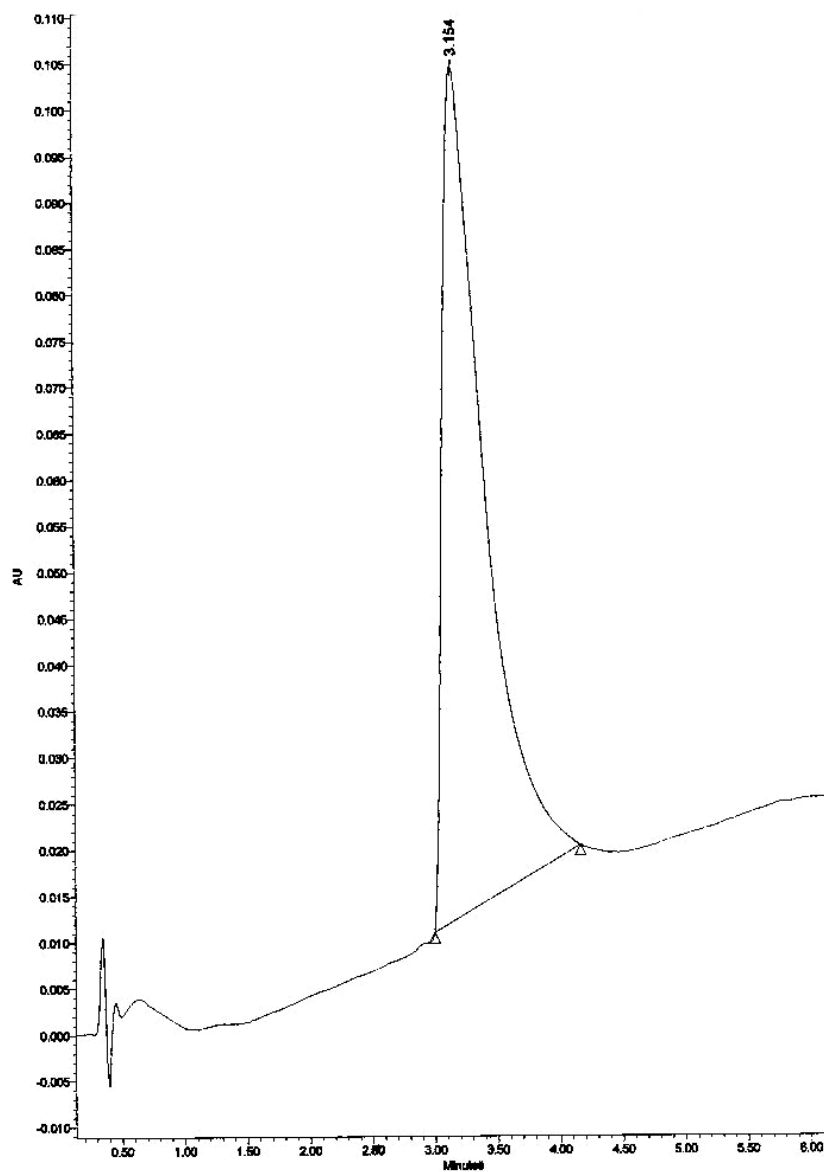
Dendrimer B<sub>3</sub>C<sub>3</sub>

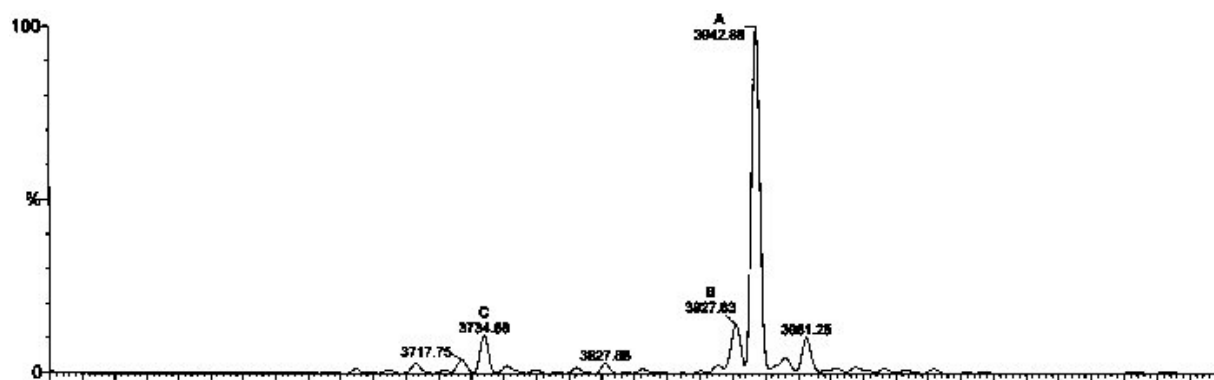
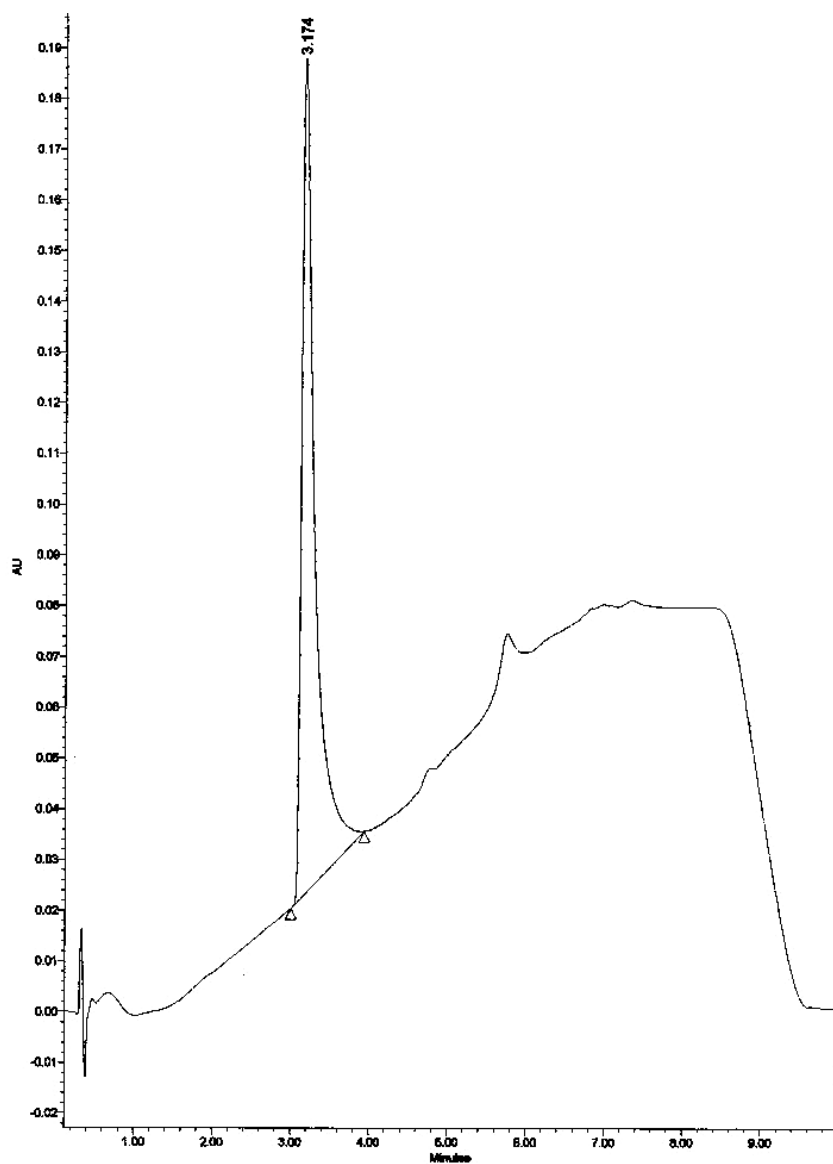
Dendrimer B<sub>3</sub>D<sub>3</sub>

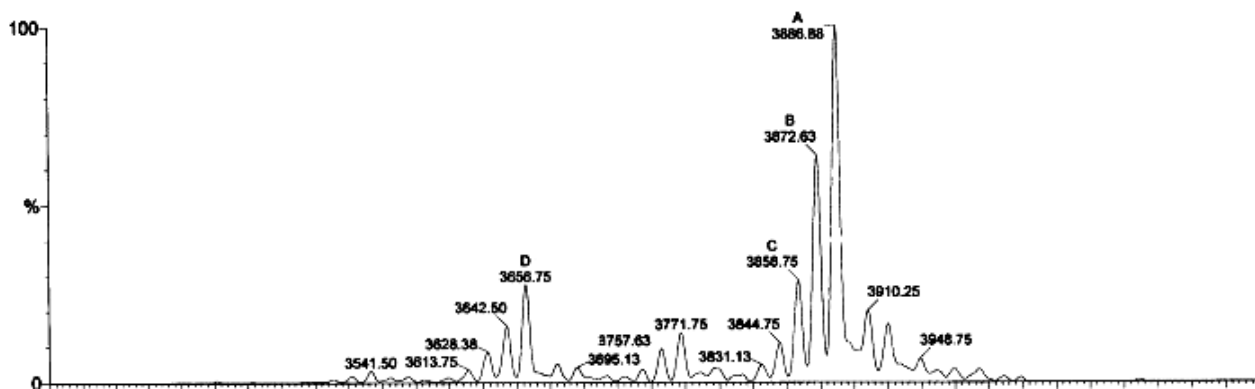
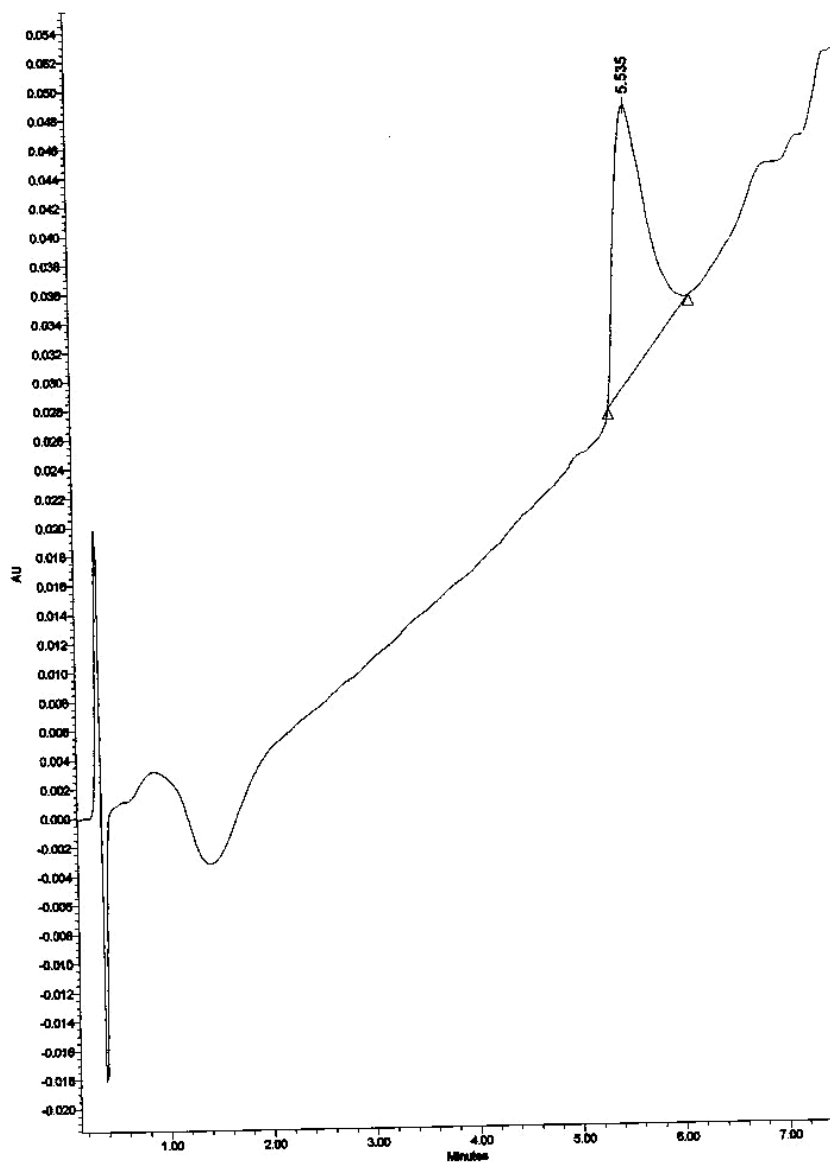


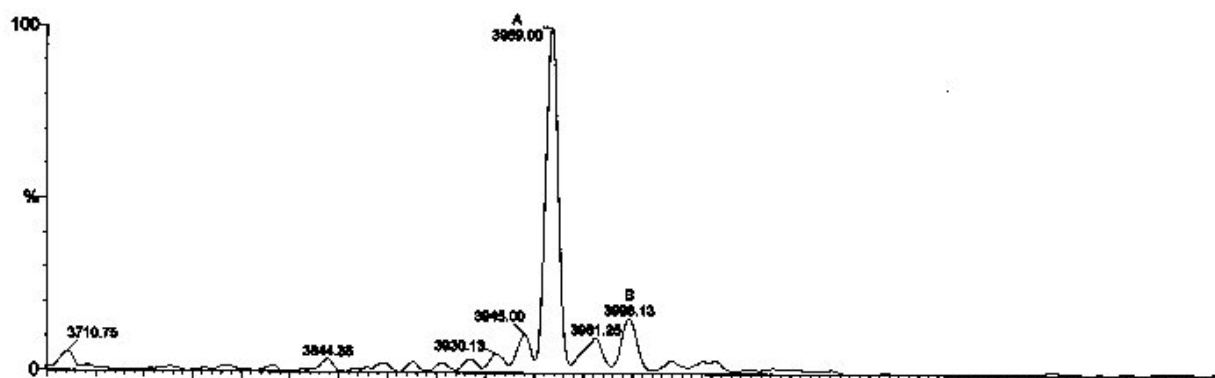
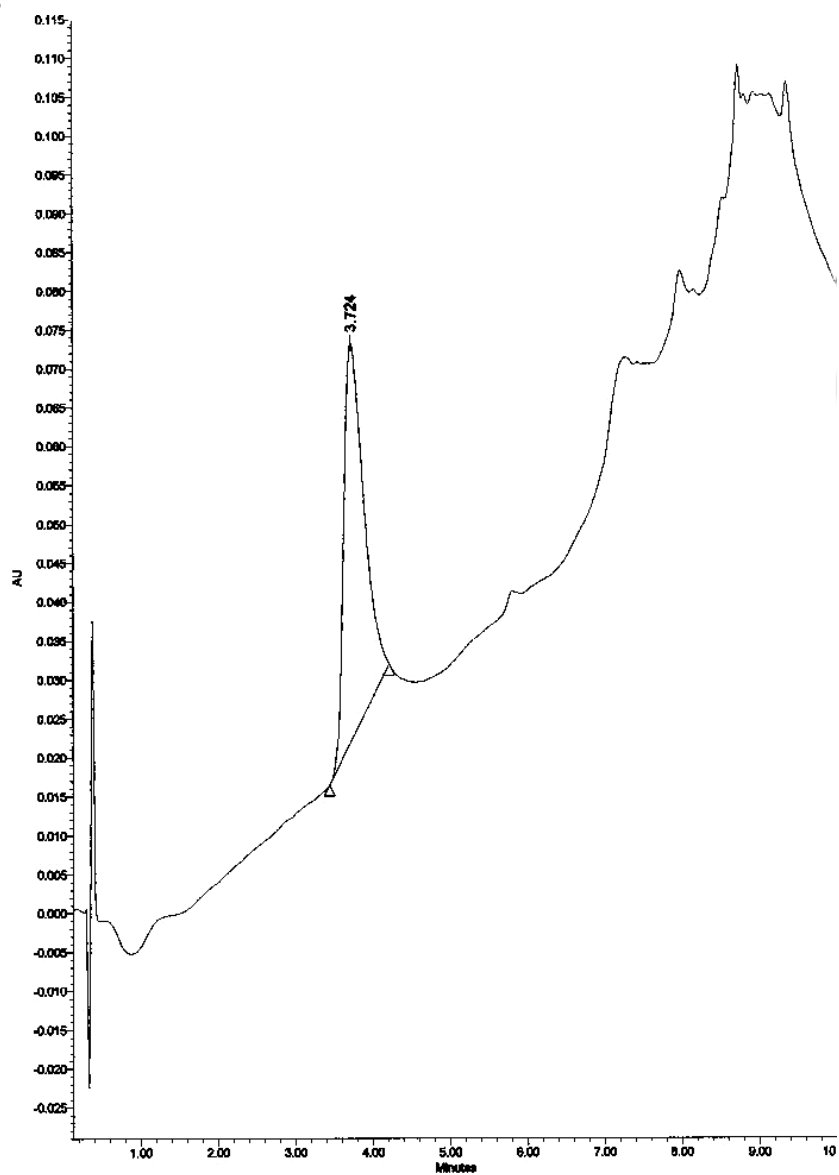
Dendrimer B<sub>3</sub>F<sub>3</sub>

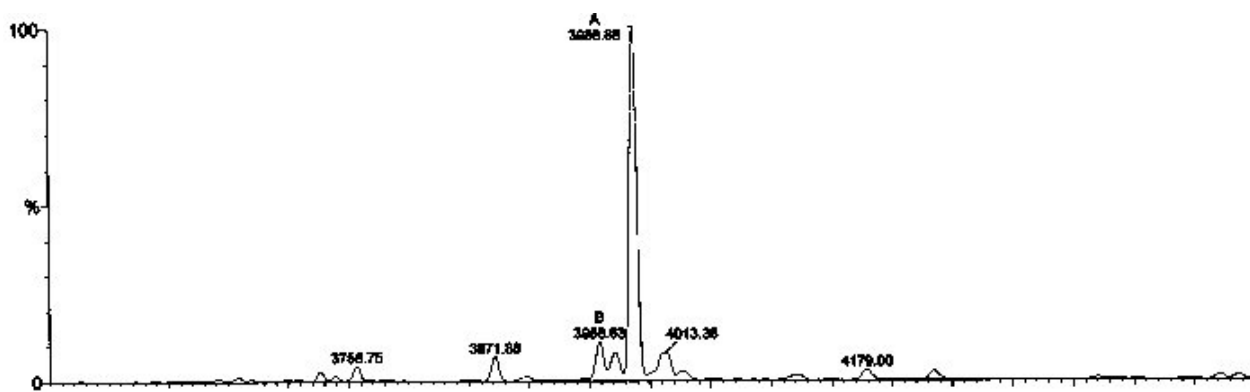
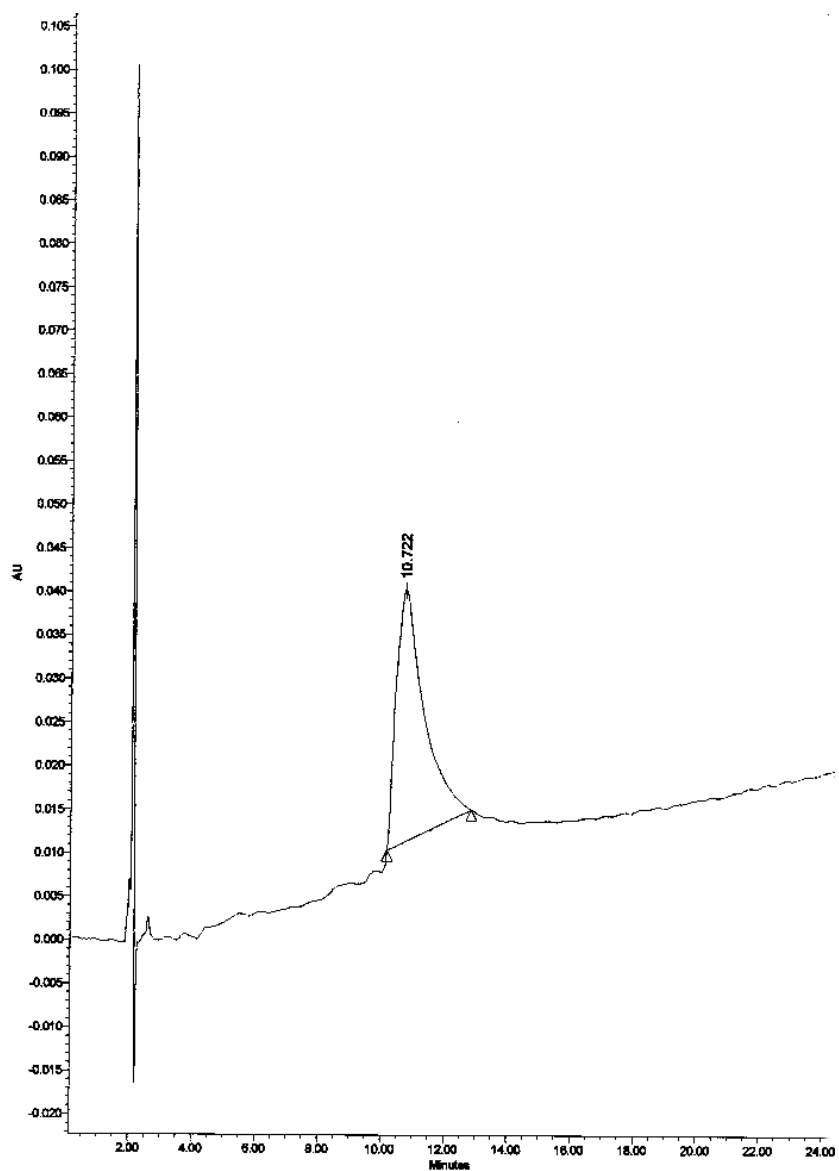
Dendrimer C<sub>3</sub>D<sub>3</sub>

Dendrimer C<sub>3</sub>E<sub>3</sub>

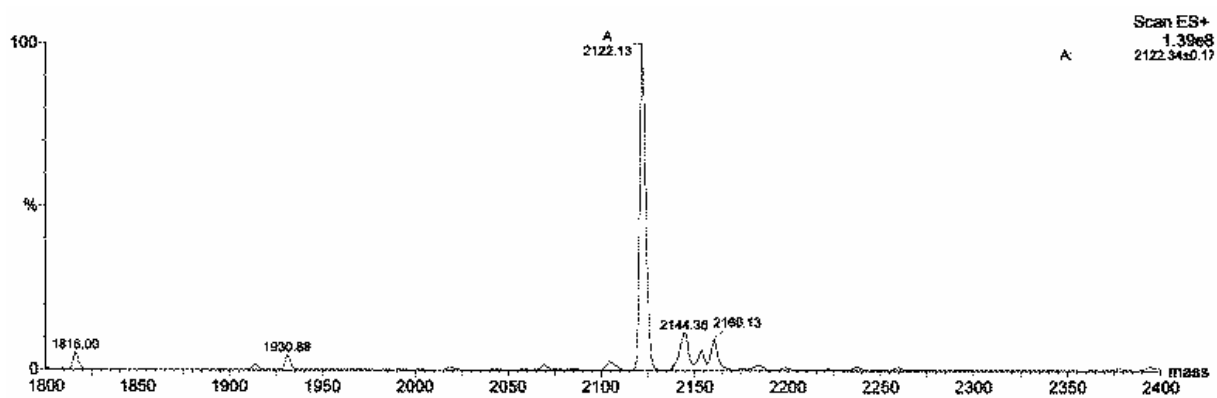
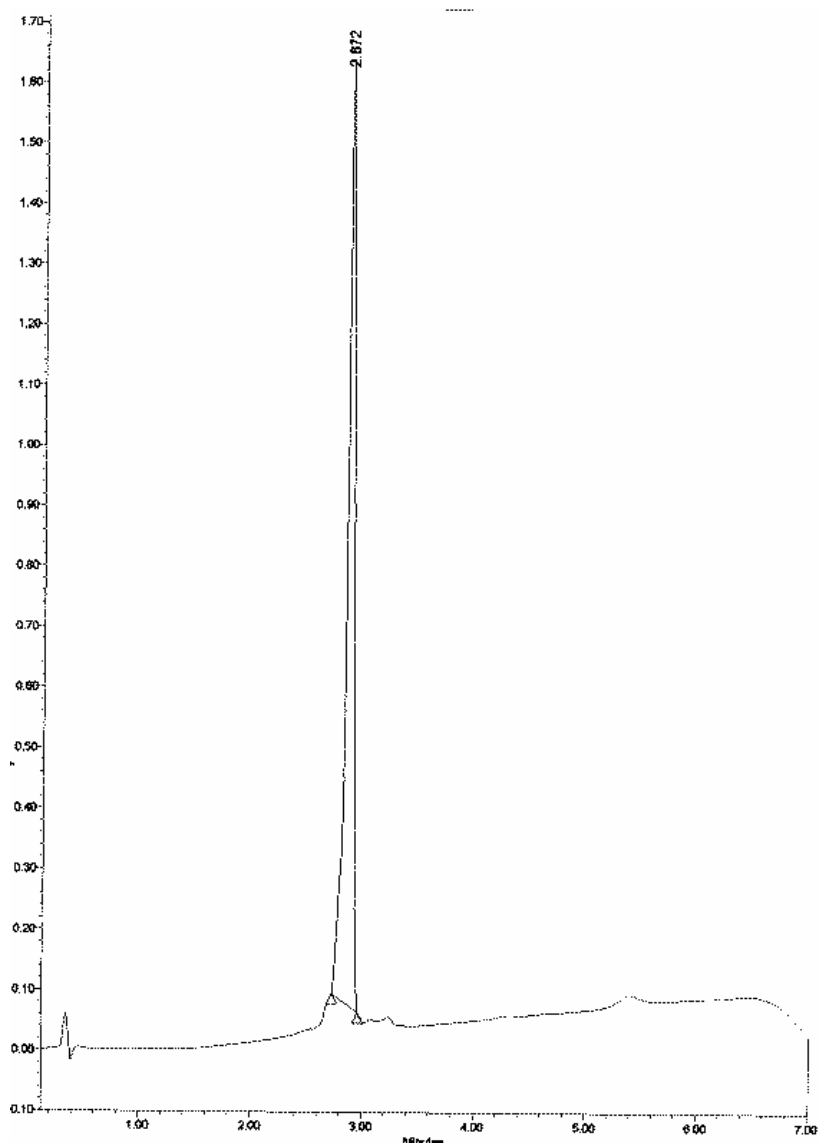
Dendrimer C<sub>3</sub>F<sub>3</sub>

Dendrimer D<sub>3</sub>E<sub>3</sub>

DendrimerD<sub>3</sub>F<sub>3</sub>

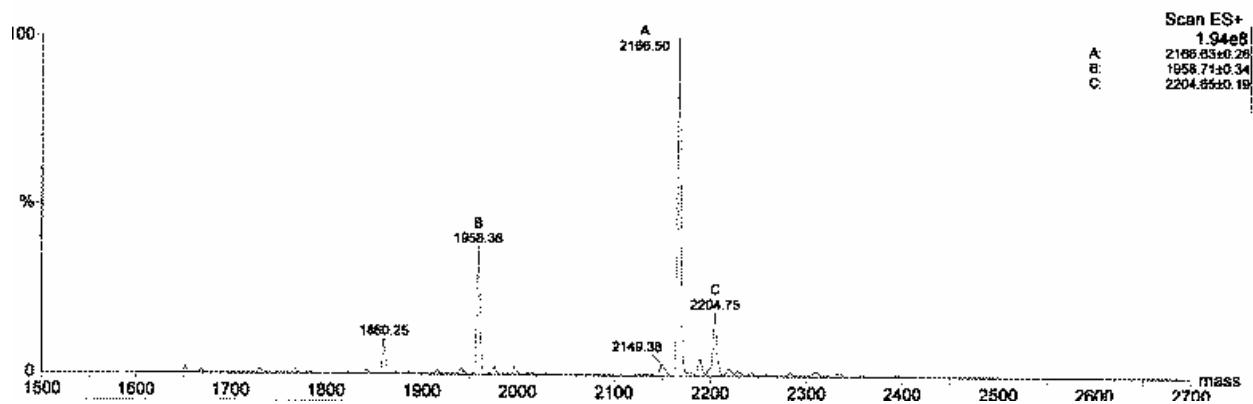
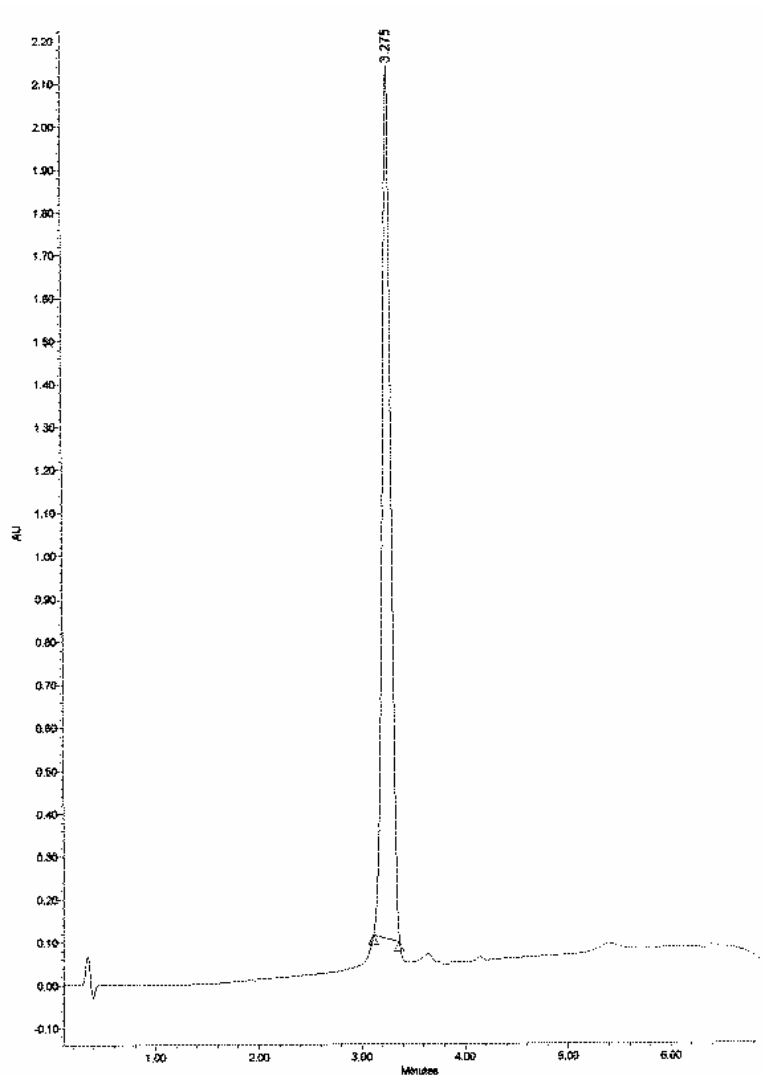
DendrimerE<sub>3</sub>F<sub>3</sub>

## Monomer DHL

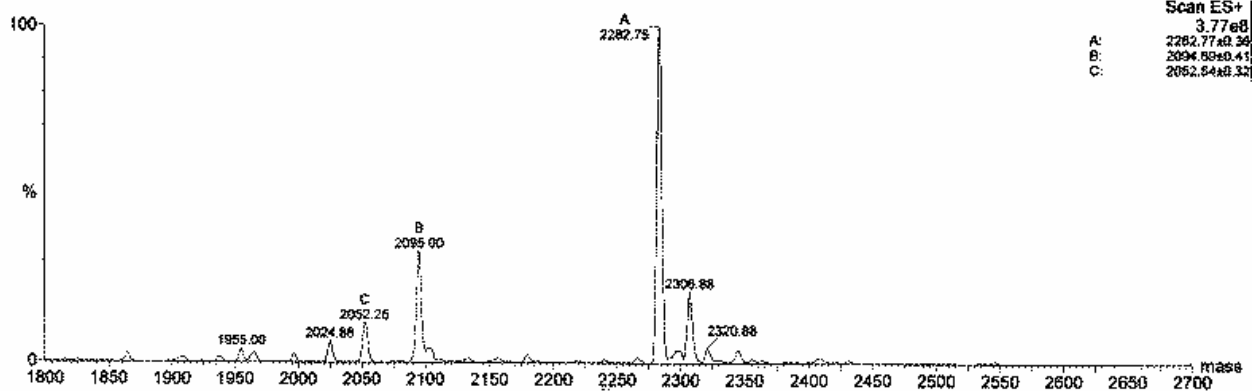
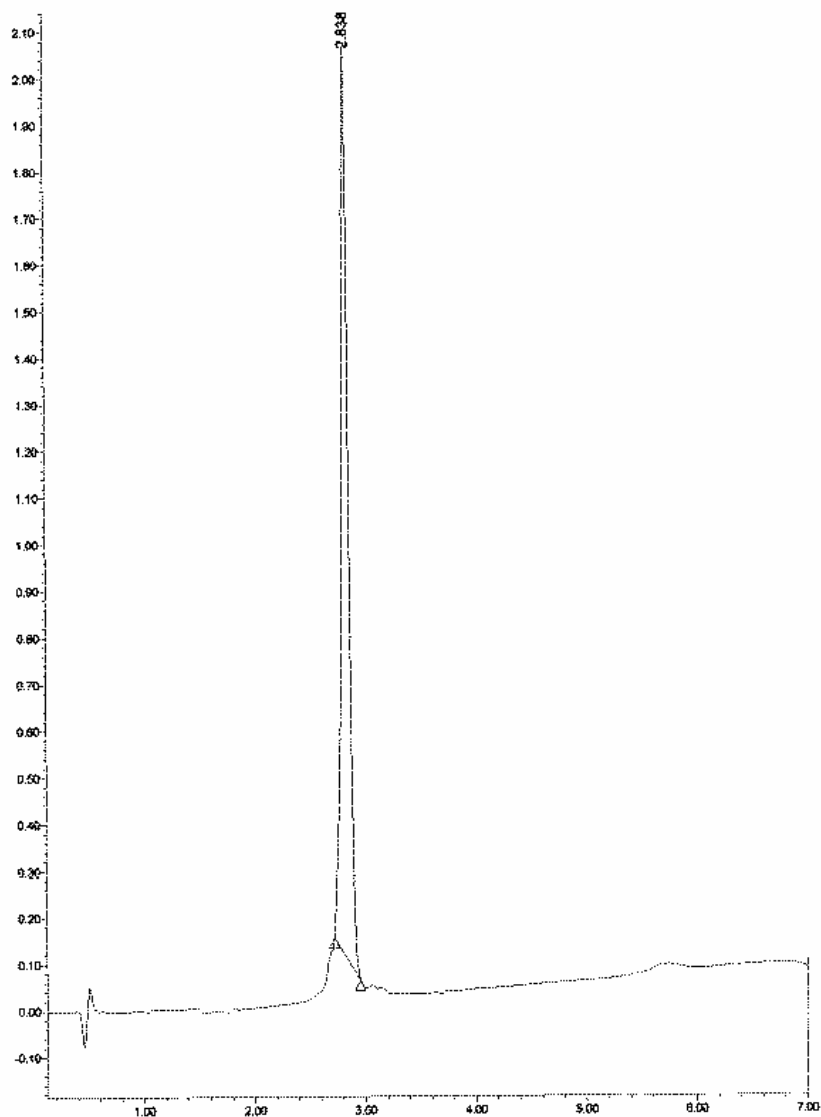




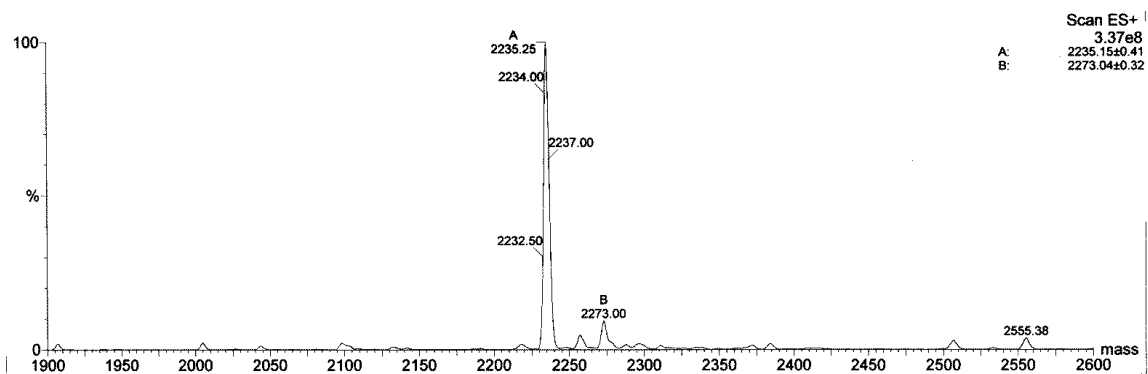
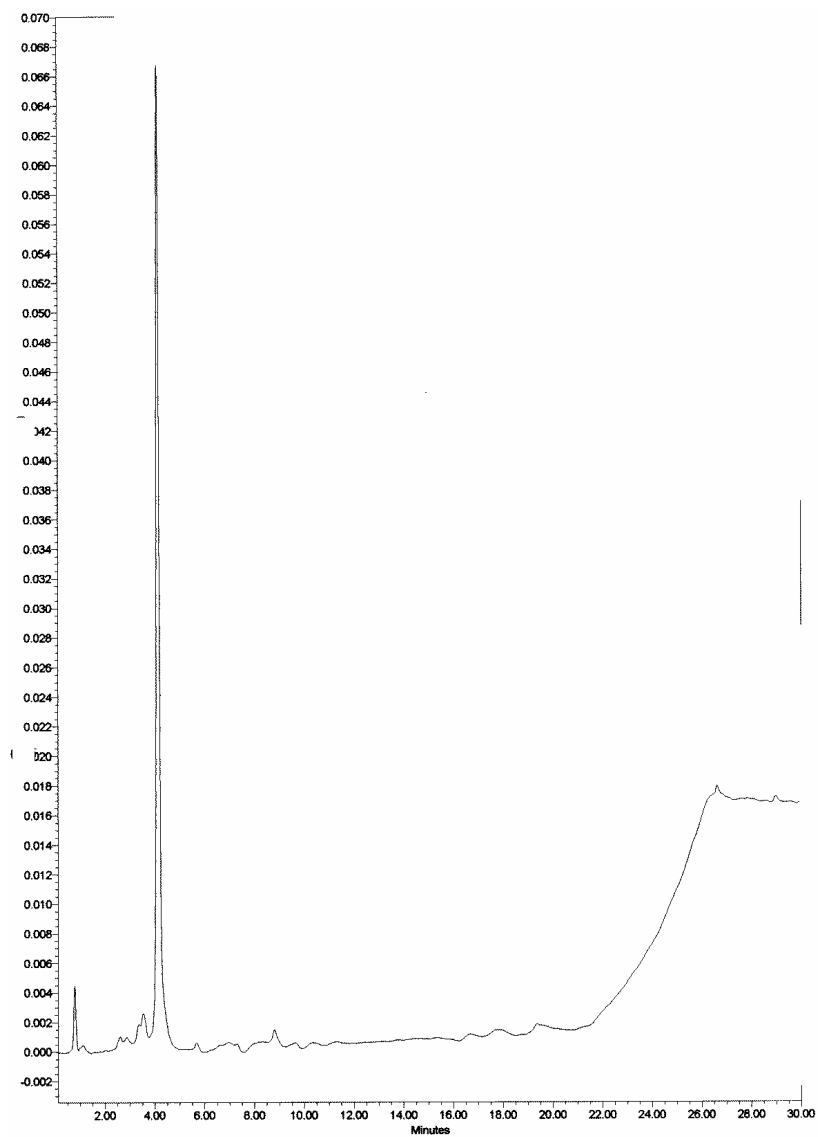
## Monomer DFH



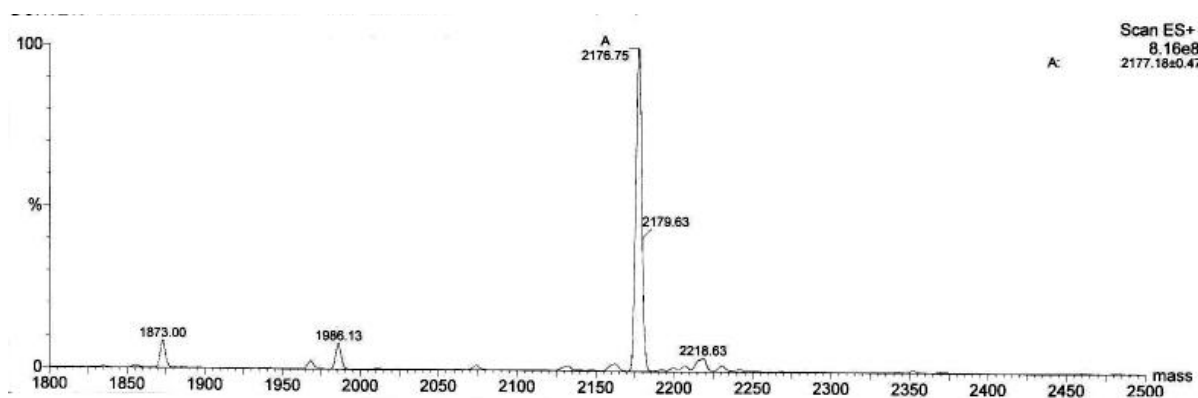
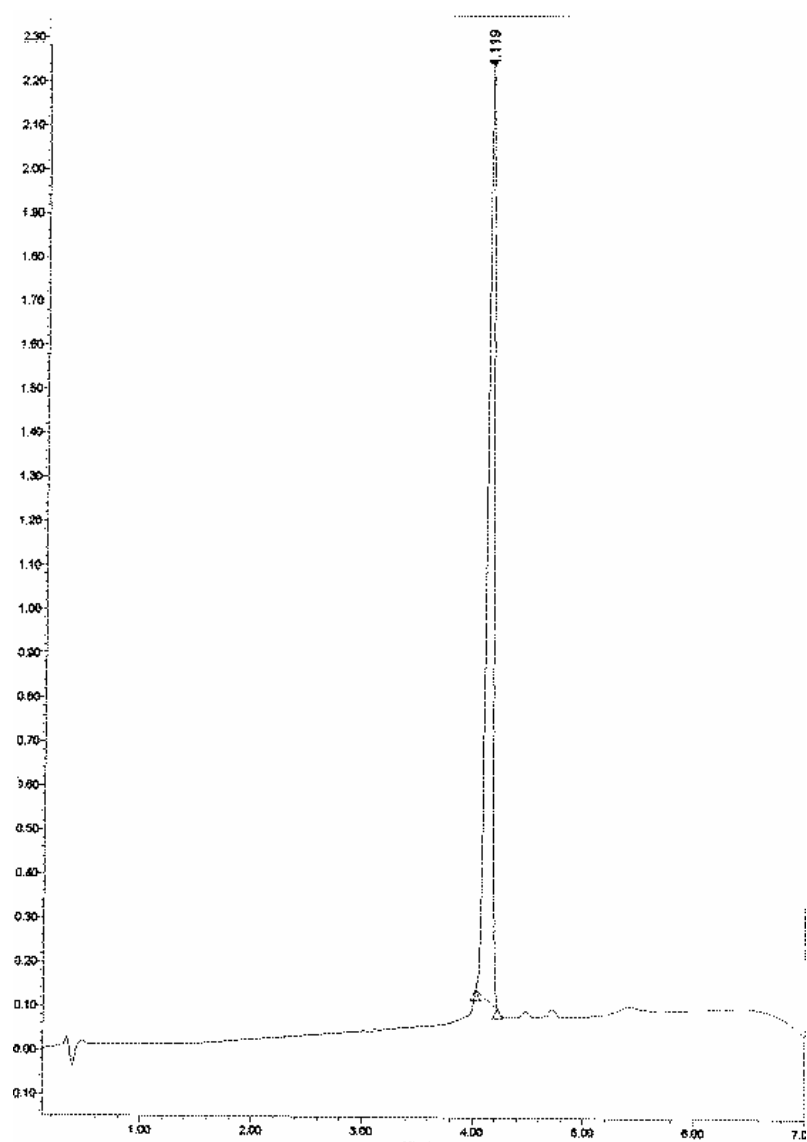
## Monomer HWS



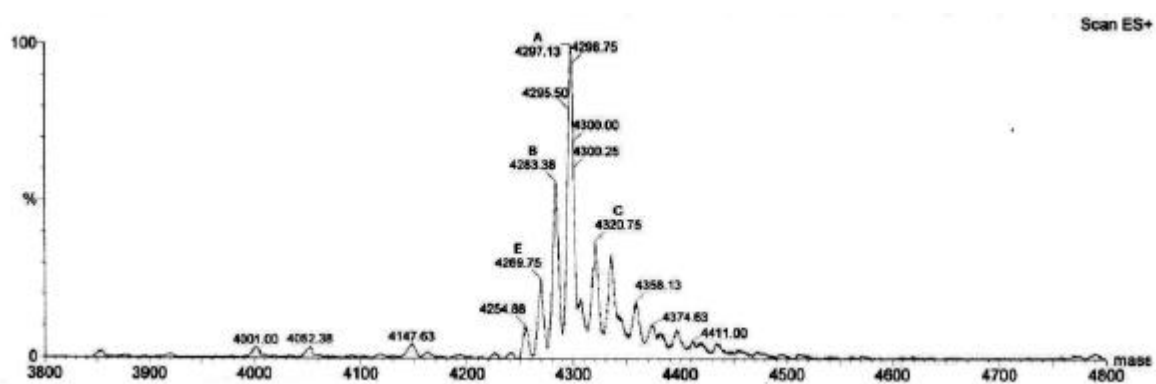
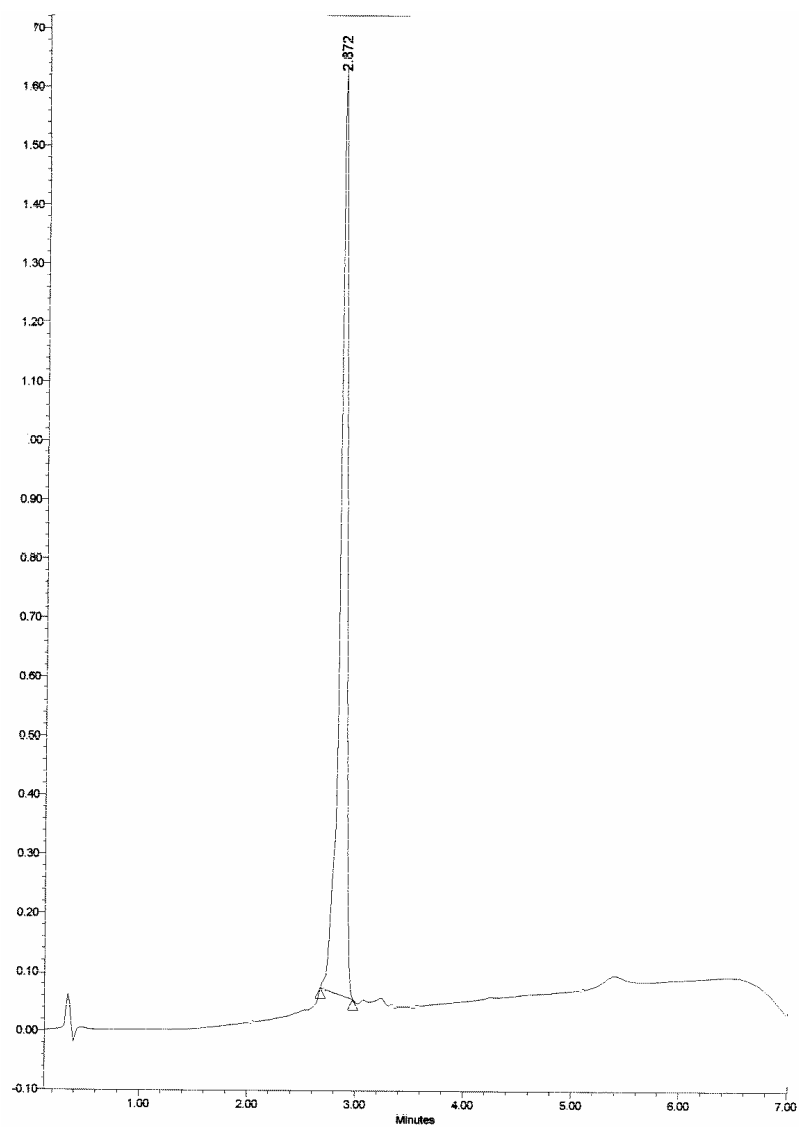
## Monomer HHH



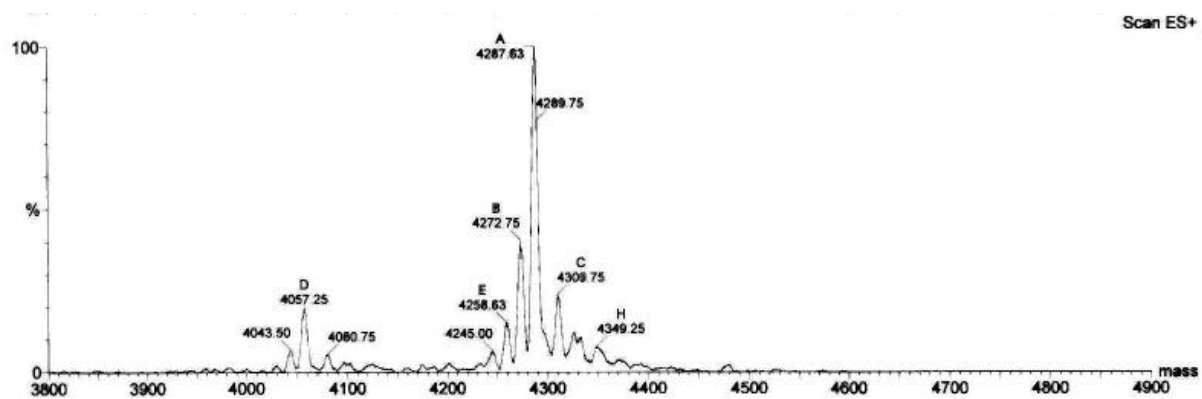
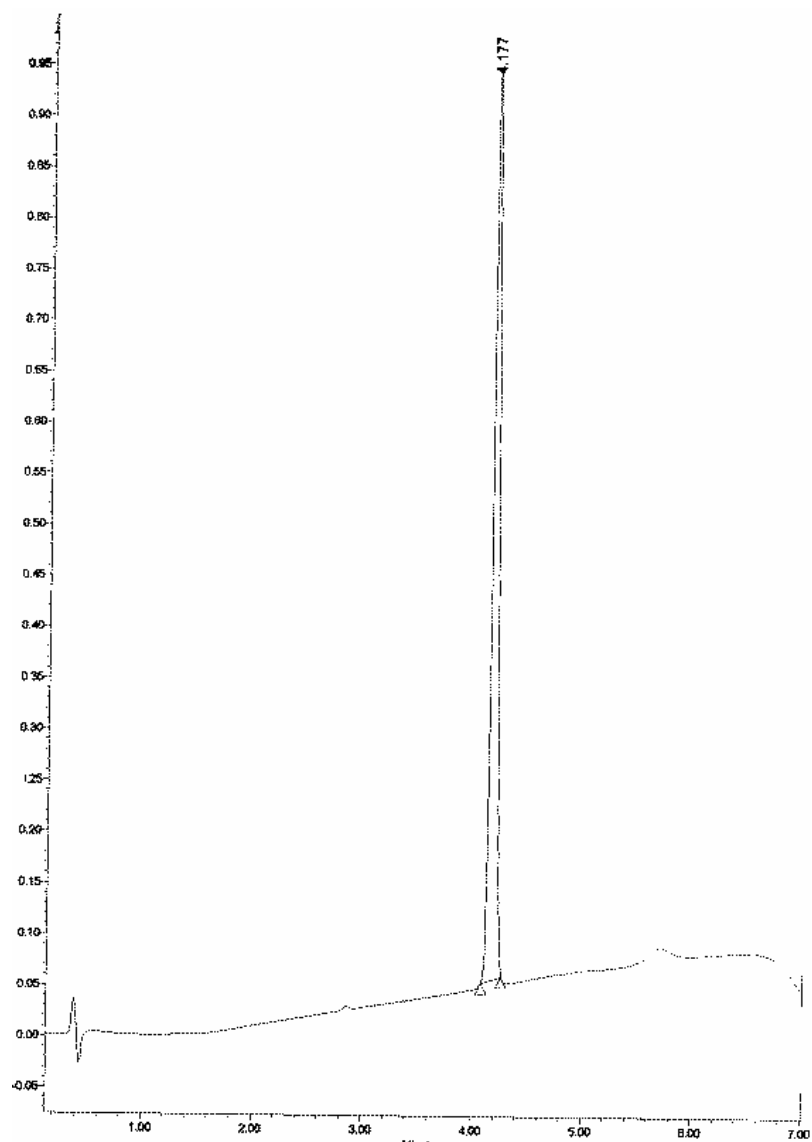
## MonomerLRH



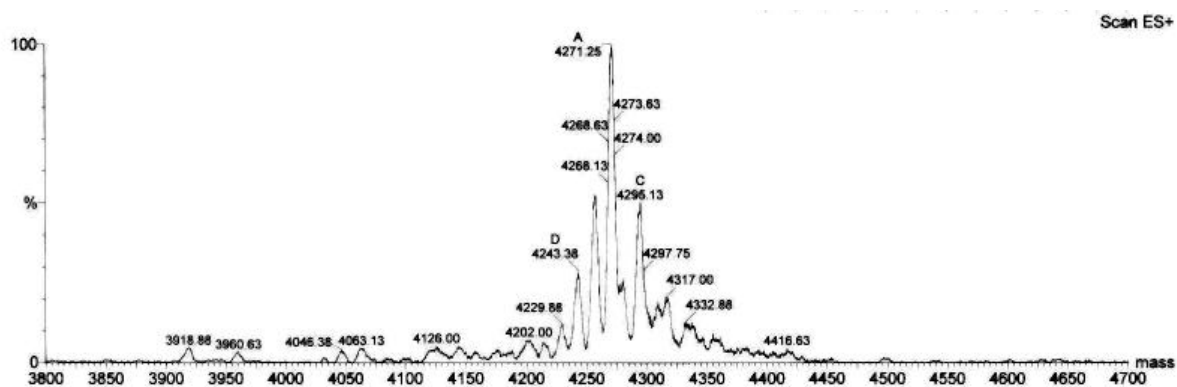
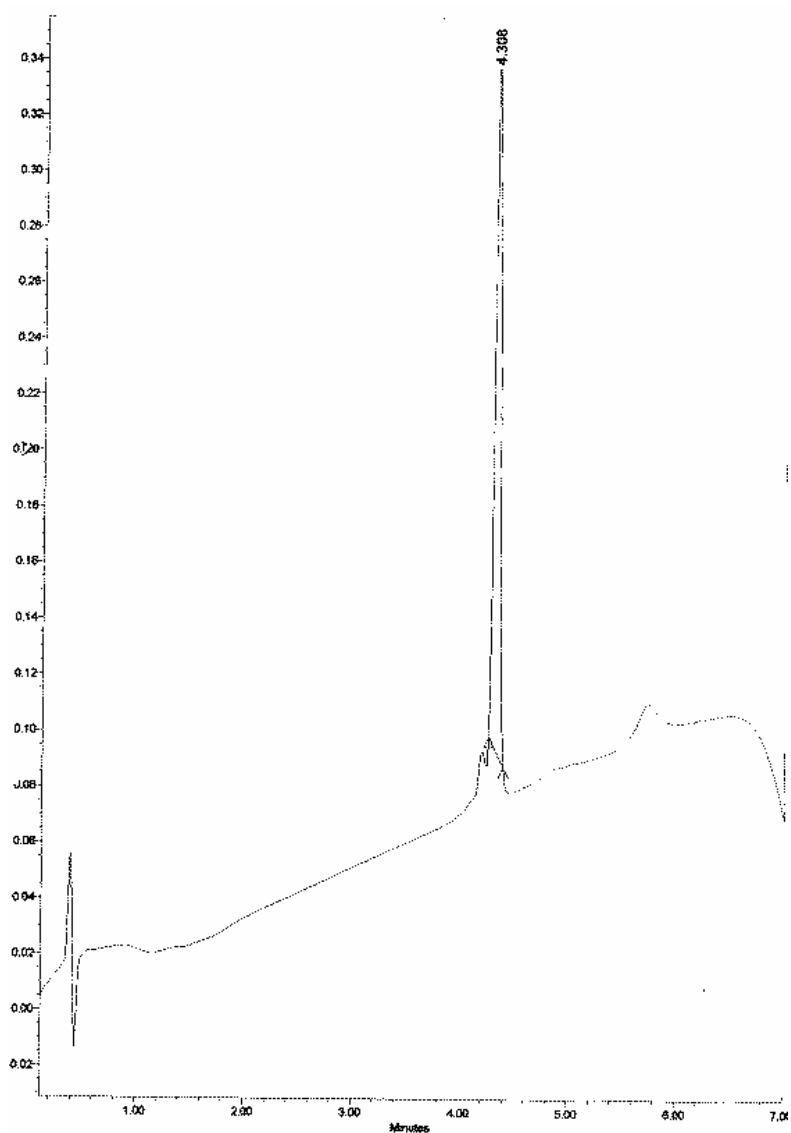
## Dendrimer LRH-DHL



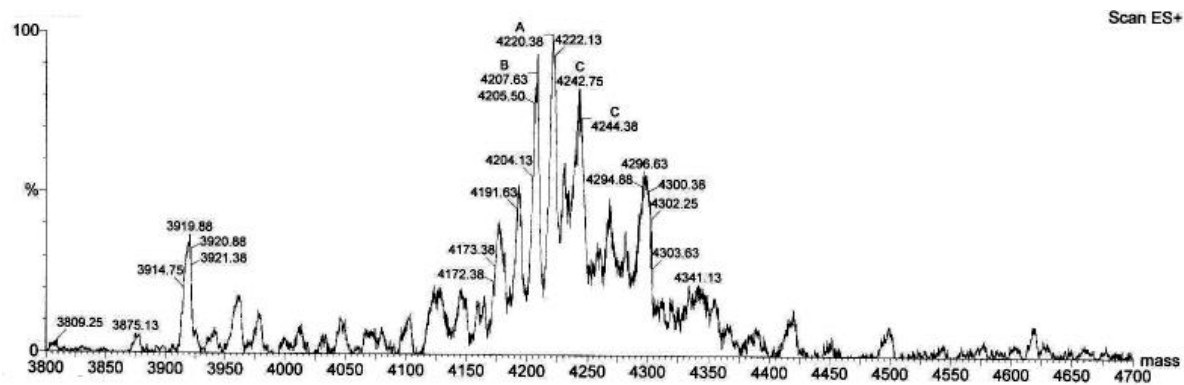
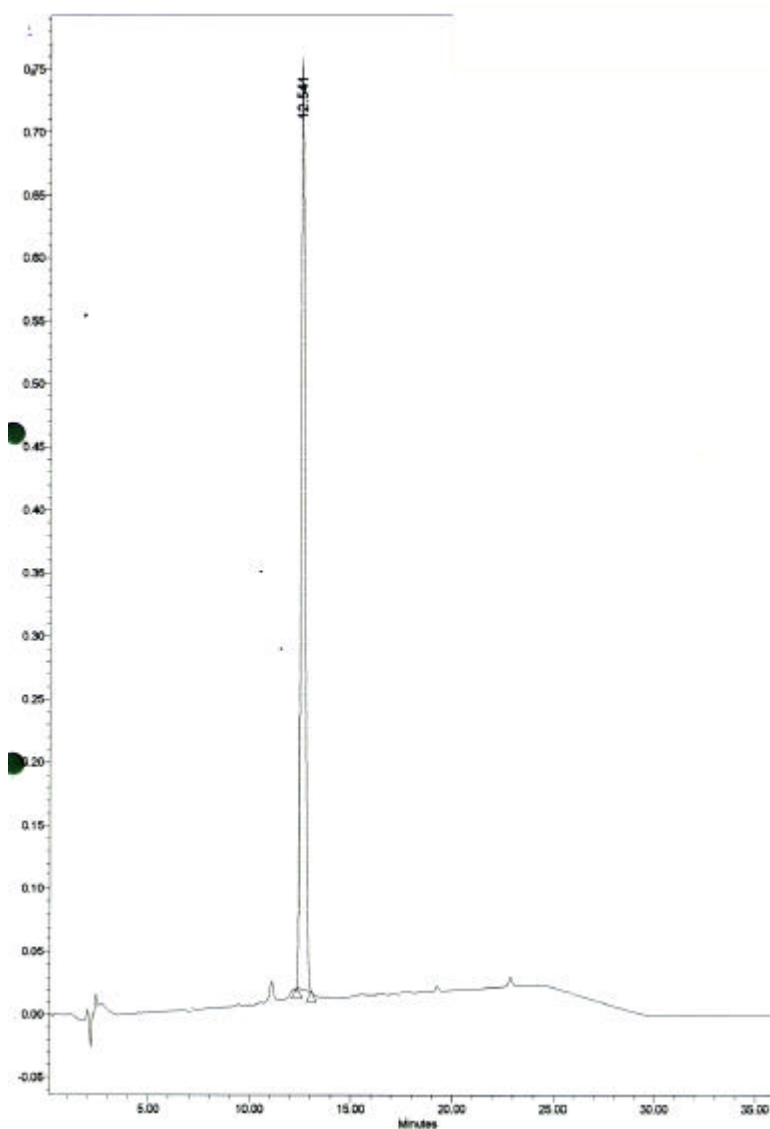
## Dendrimer LRH-HSD



## Dendrimer LRH-DHS

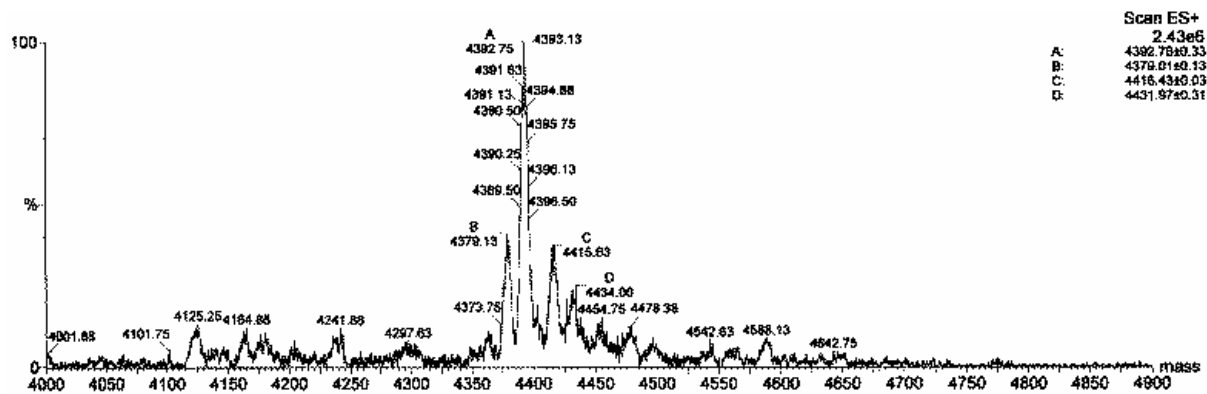
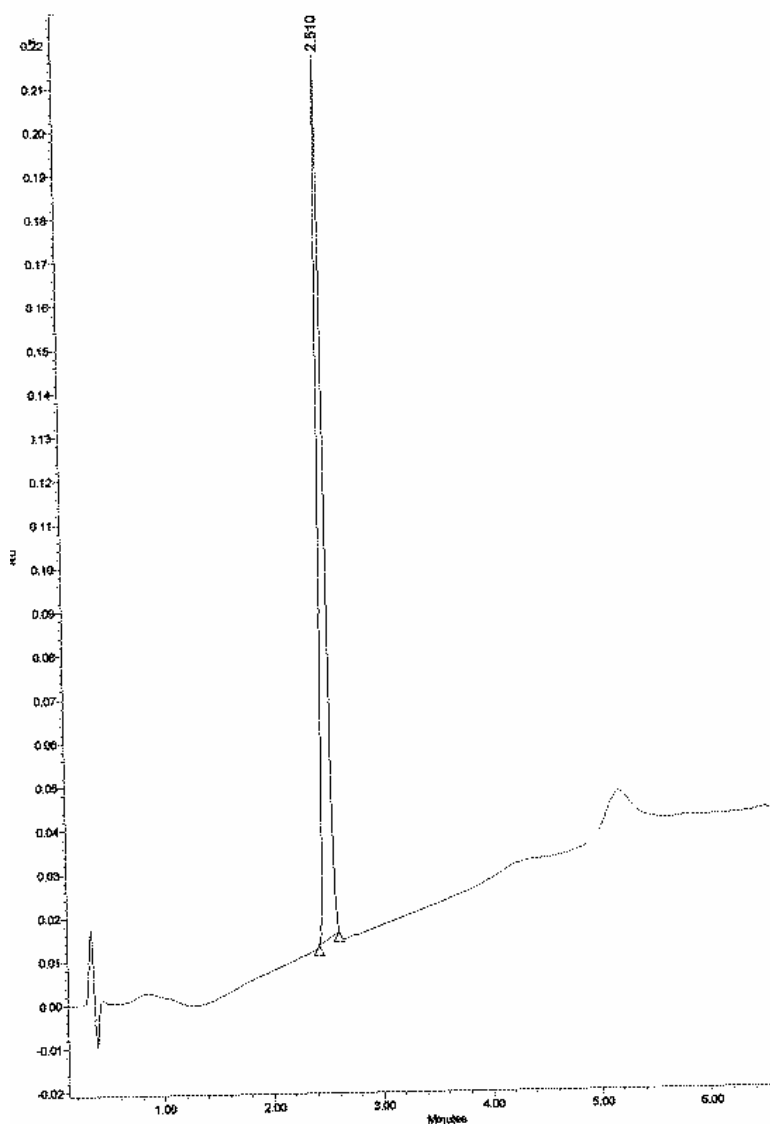


## Dendrimer LRH-DSH

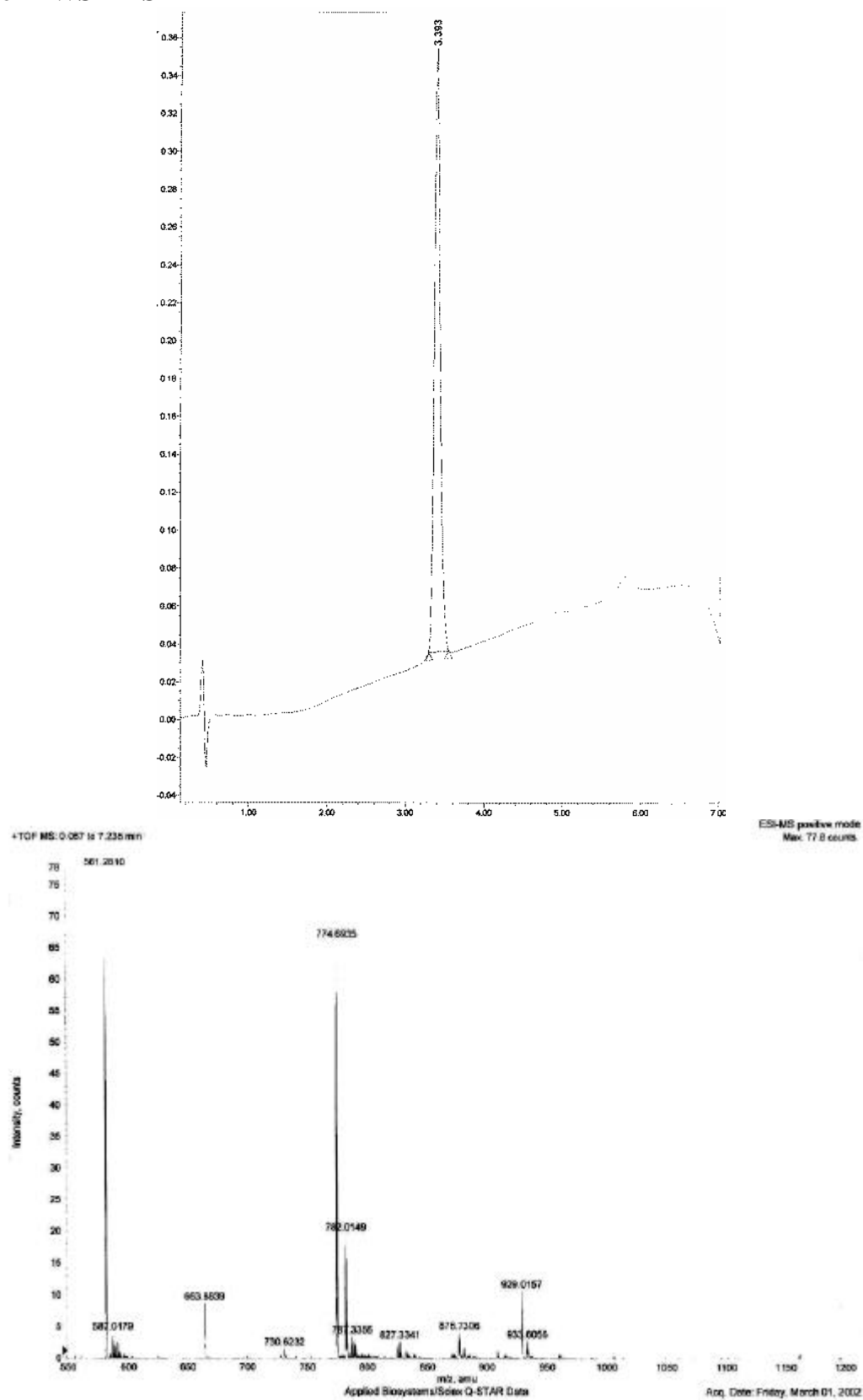




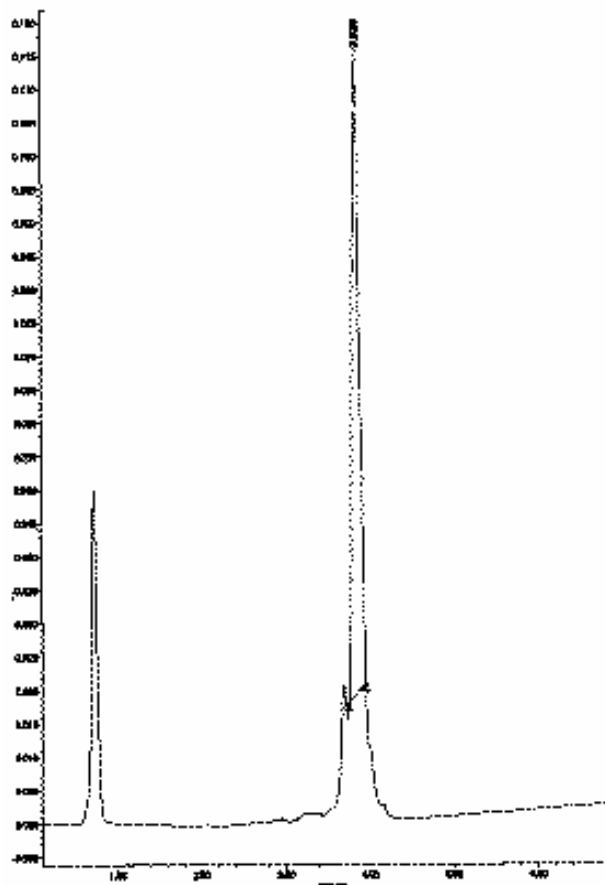
Dendrimer HWS-HSD



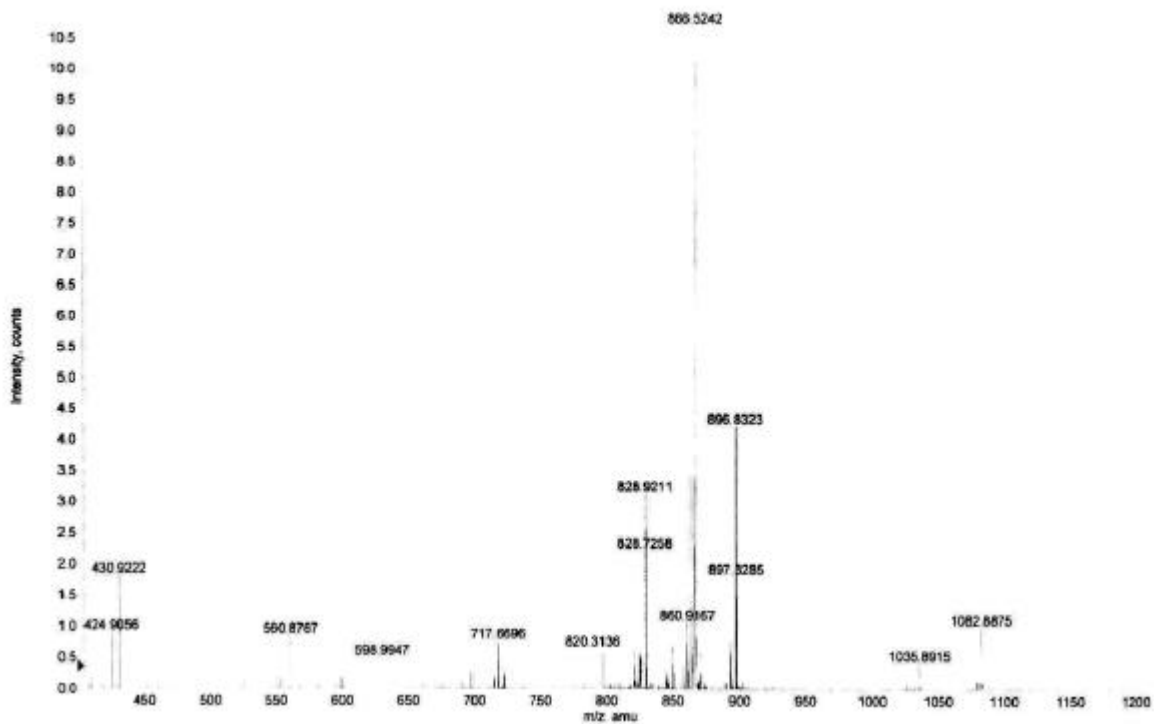
## Dendrimer HWS-DHS



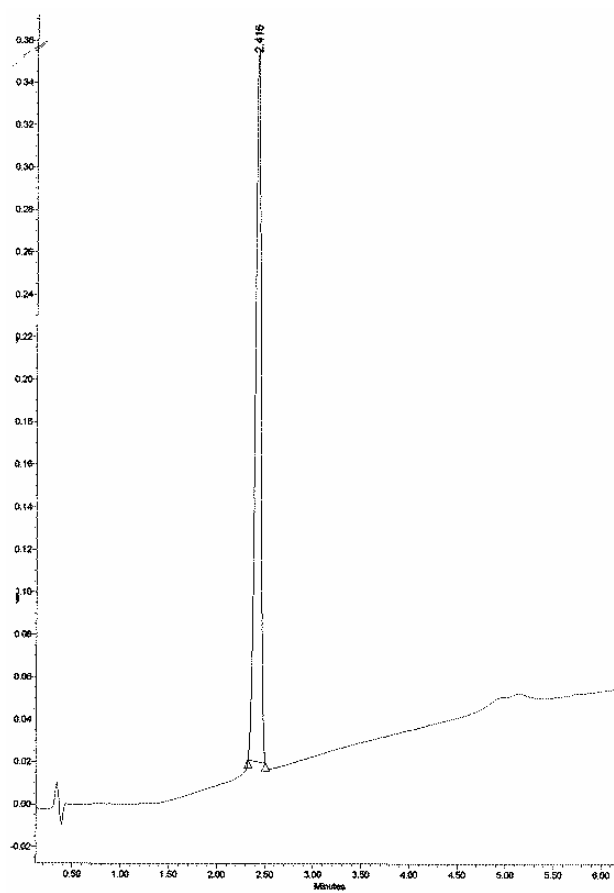
## Dendrimer HWS-DSH



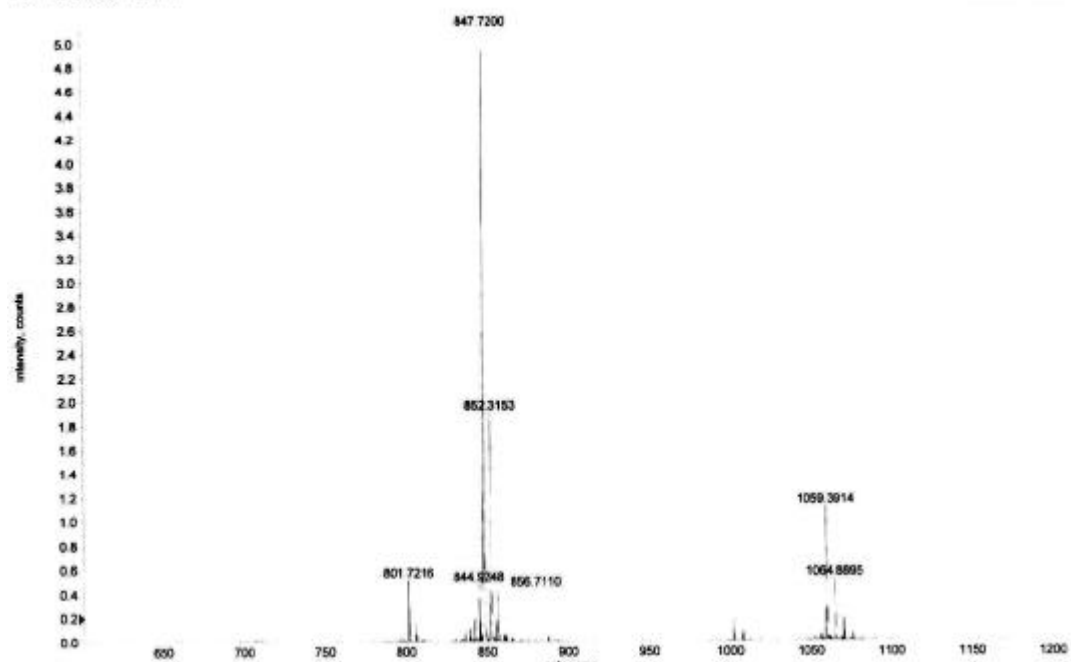
+TOF MS: 0.817 to 6.685 min

ESI-MS positive mode  
Max: 10.7 counts

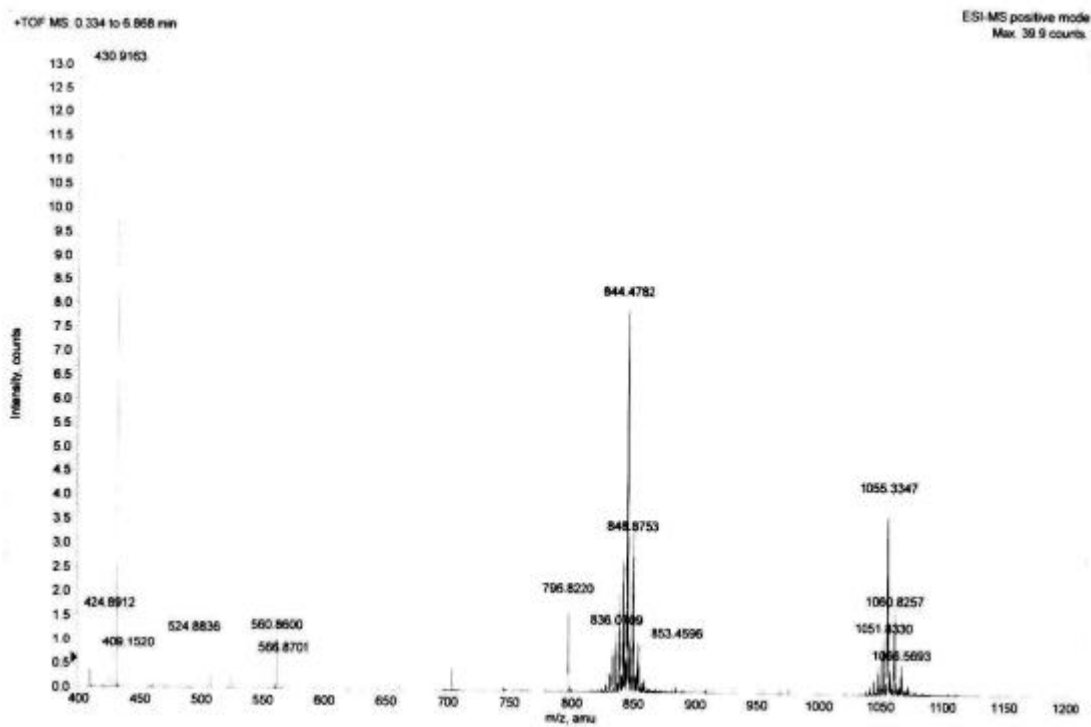
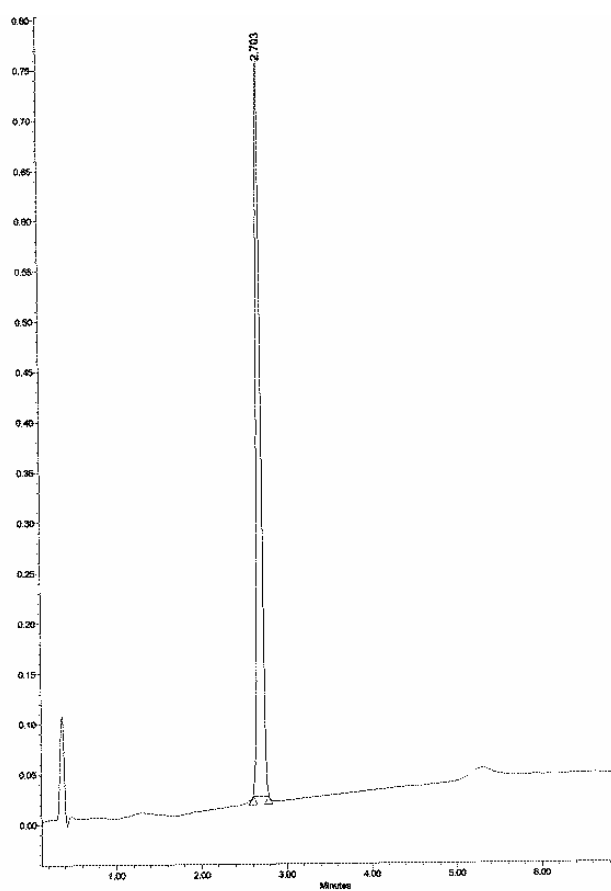
## Dendrimer DHL-HSD



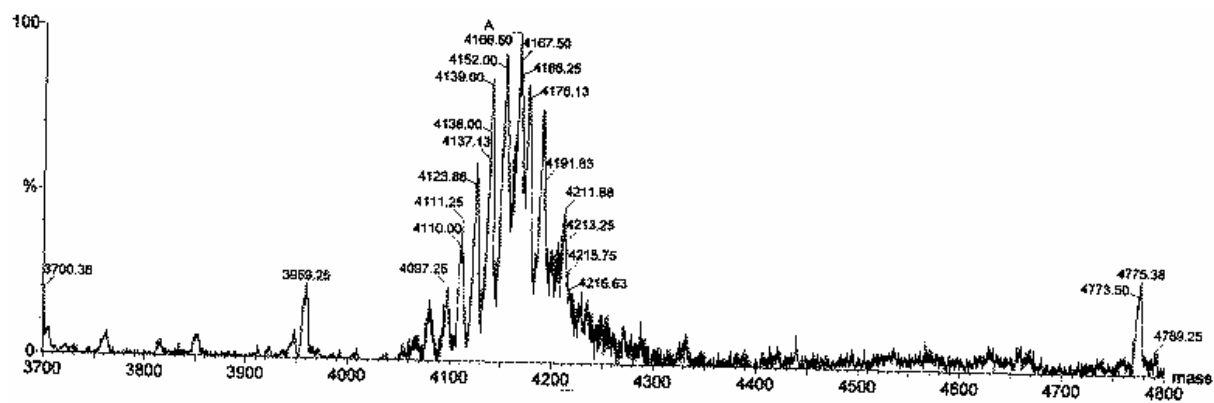
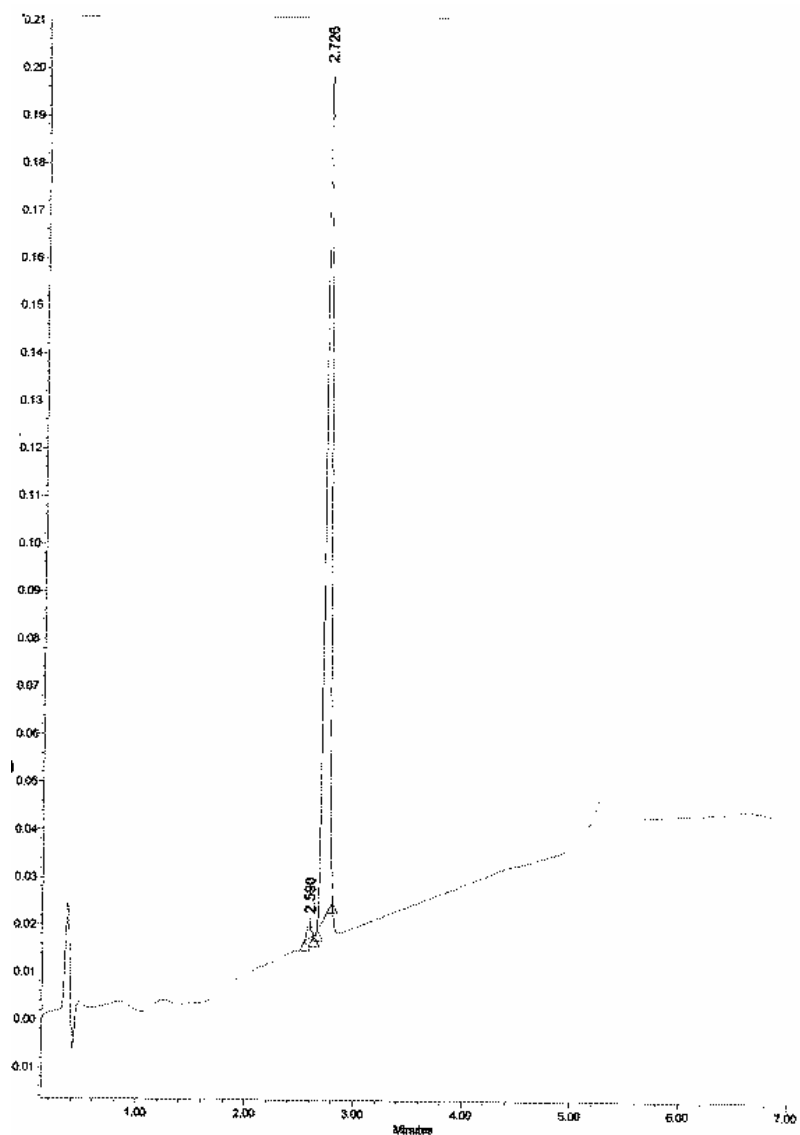
+TOF MS: 0.050 to 7.102 min



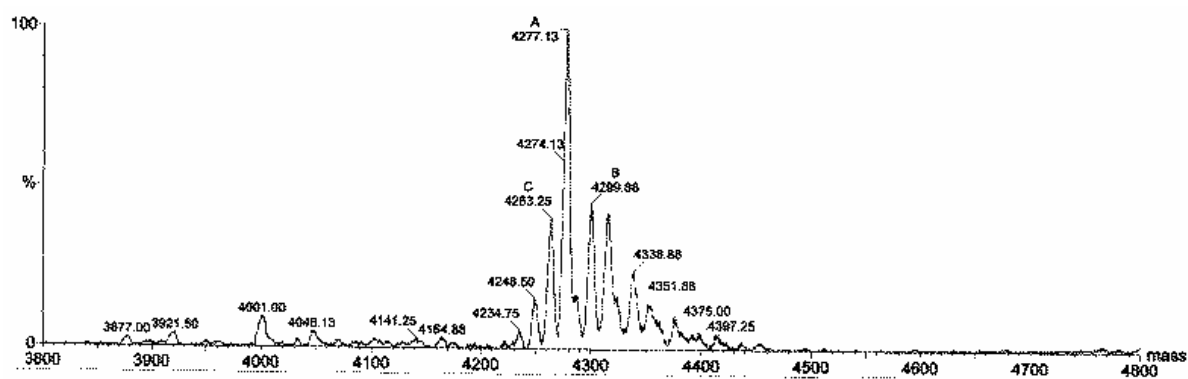
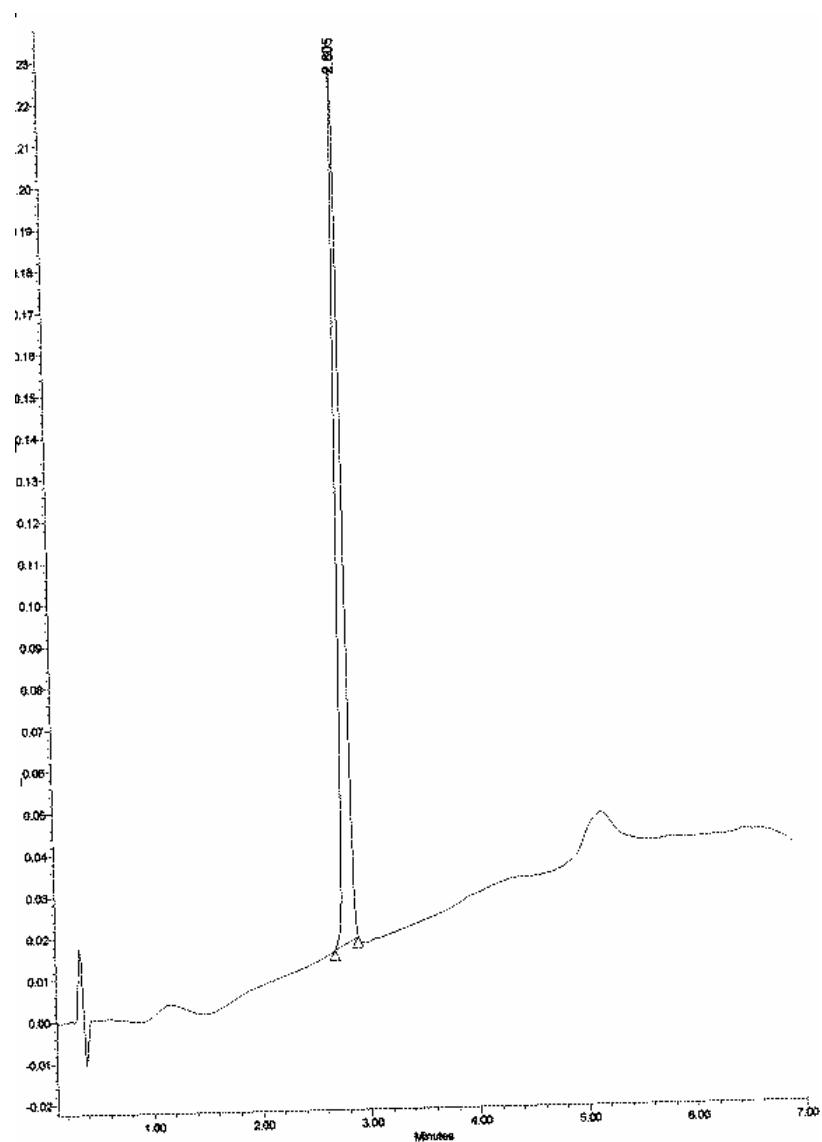
## Dendrimer DHL-DHS



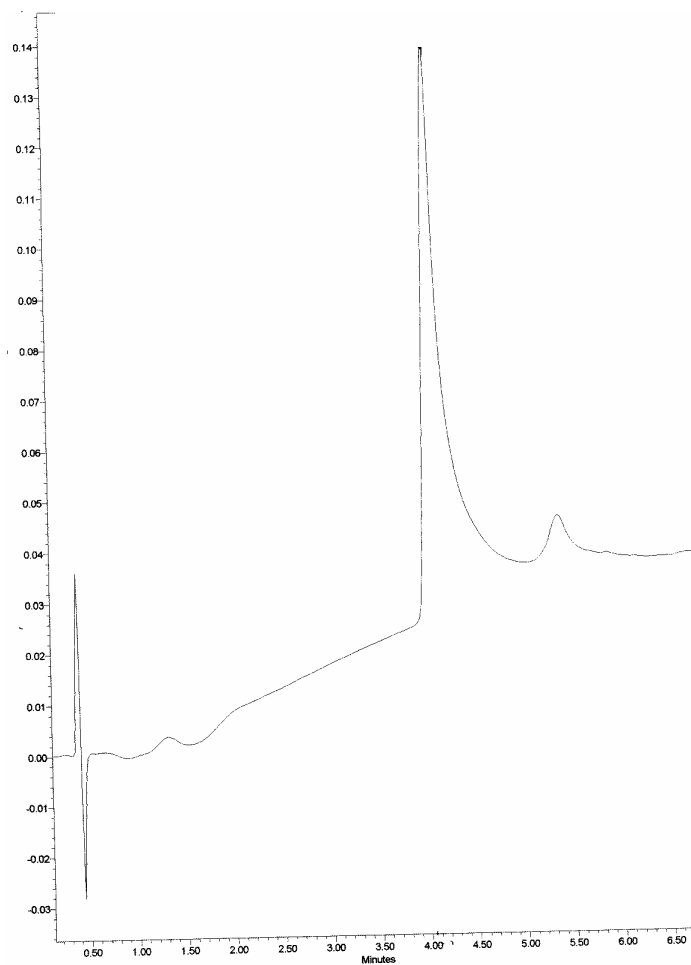
## Dendrimer DHL-DSH



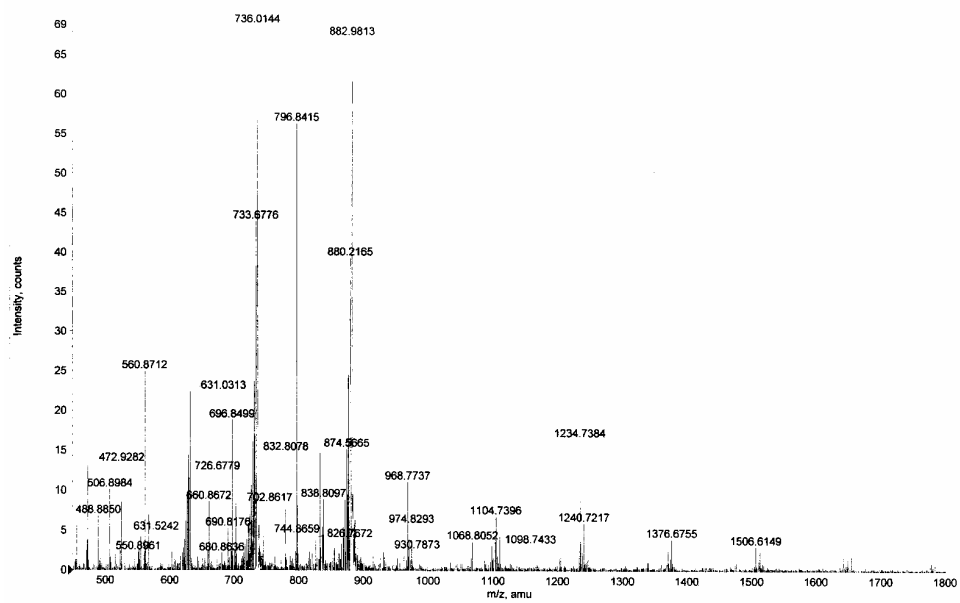
## DendrimerDFH-HSD



## Dendrimer LRH-HHH

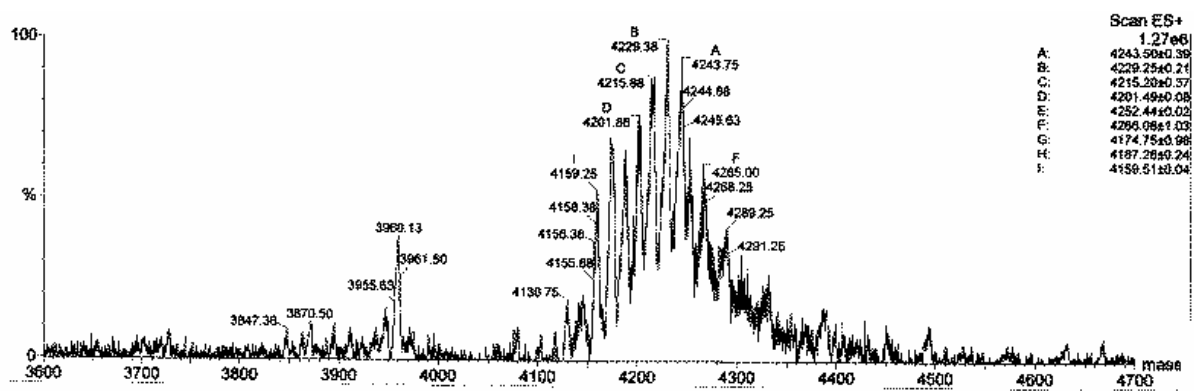
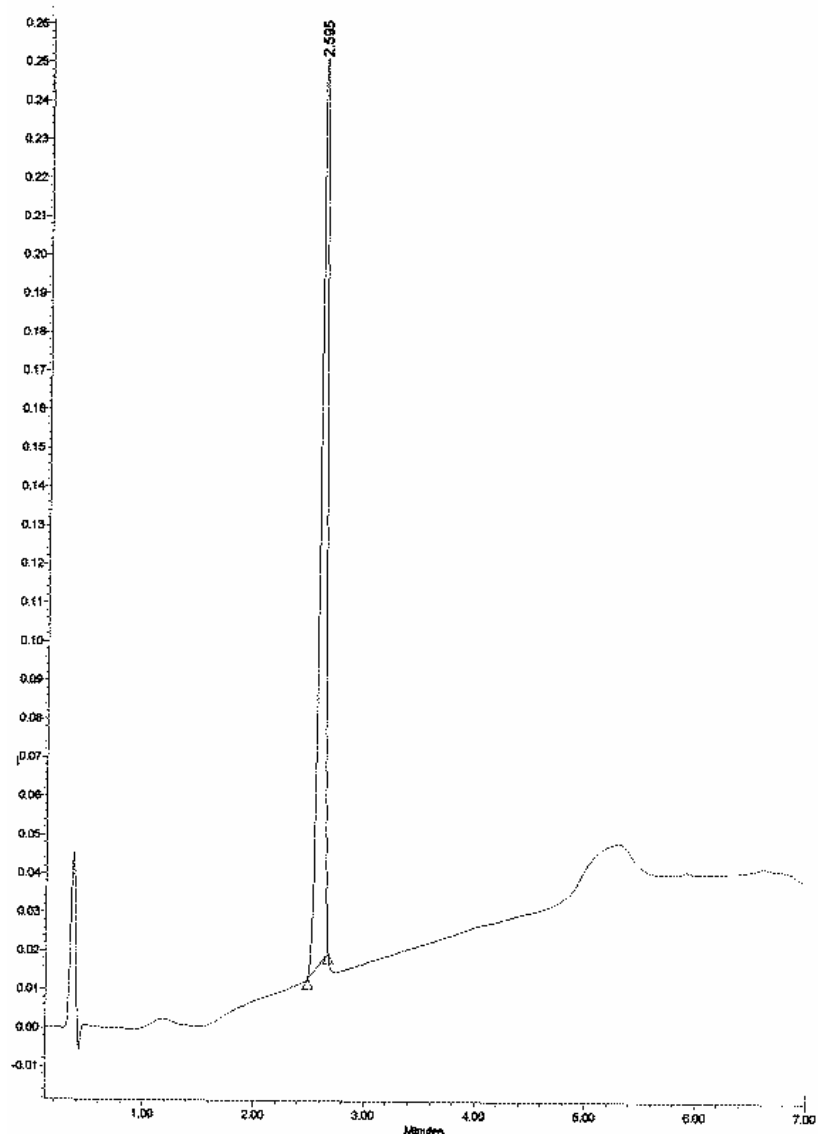


+TOF MS: 3.184 to 3.384 min

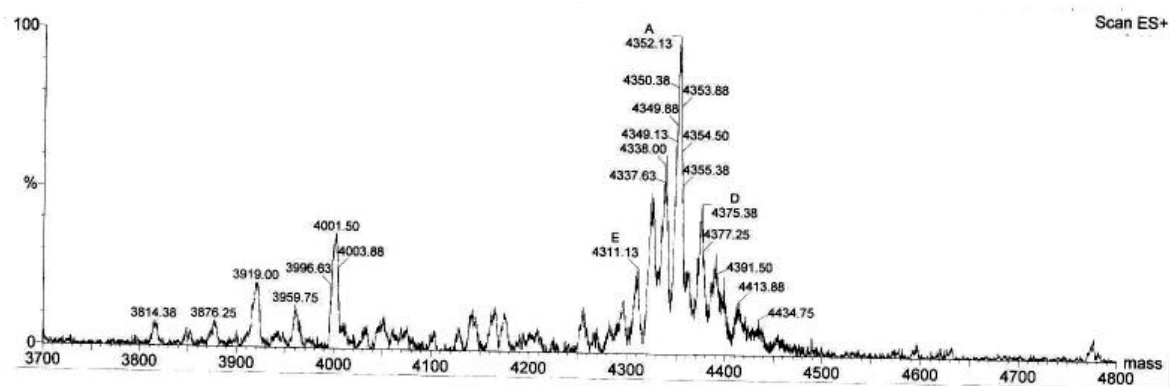
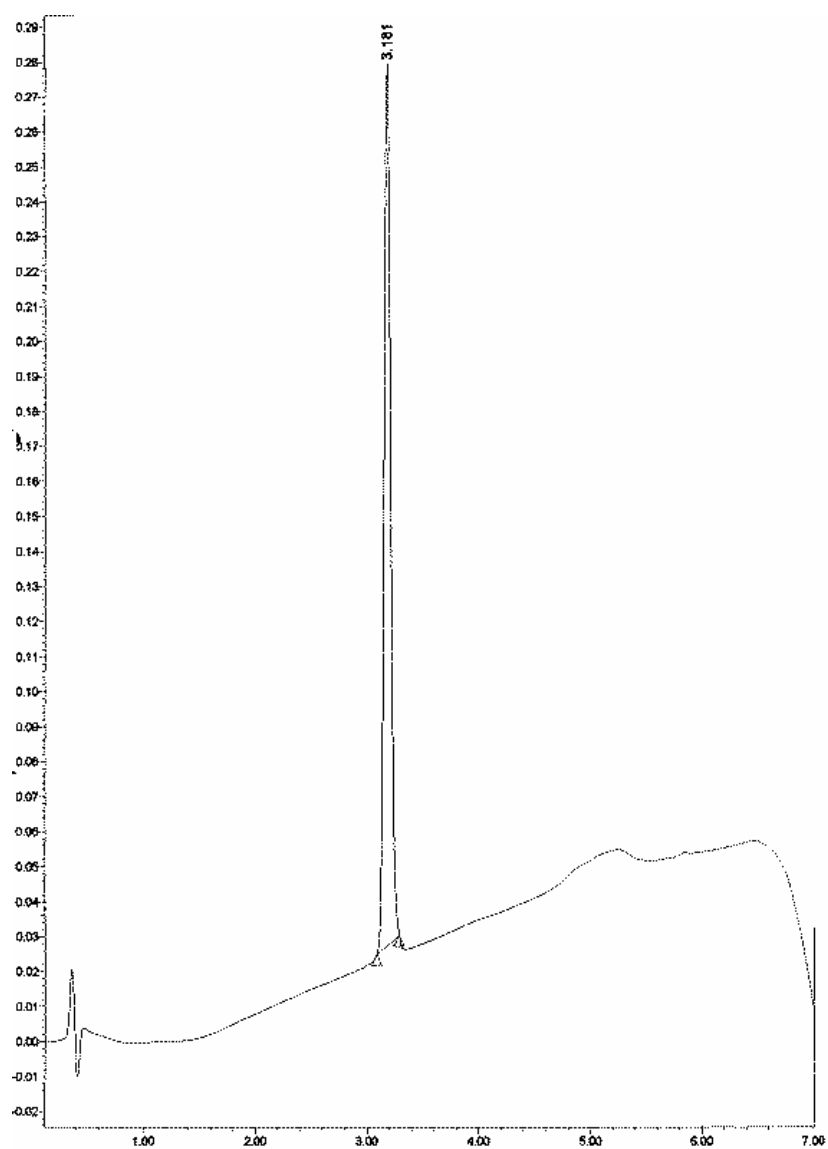
ESI-MS positive mode  
Max. 106.5 counts.



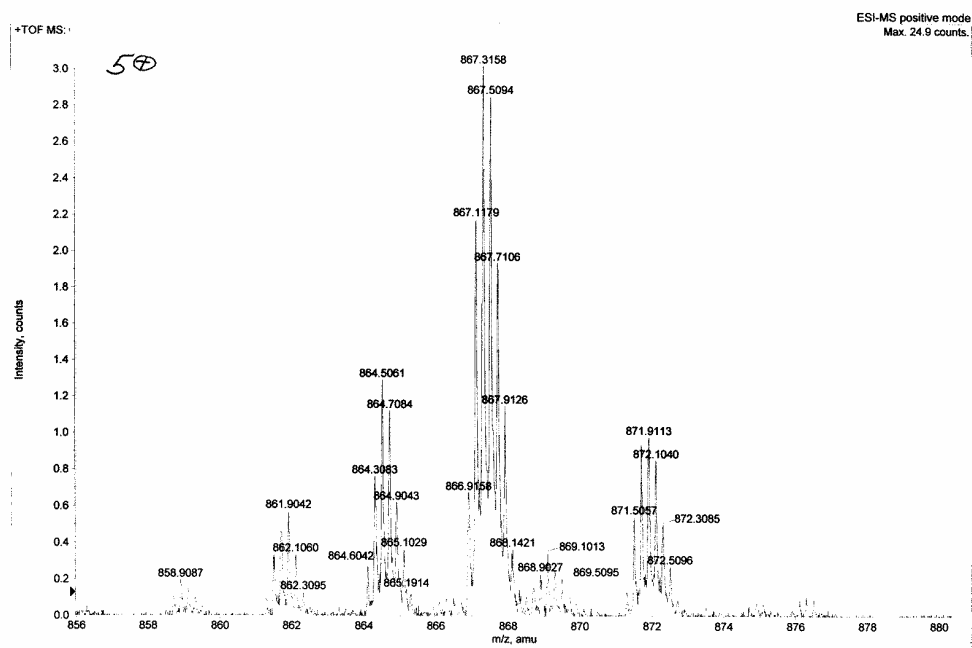
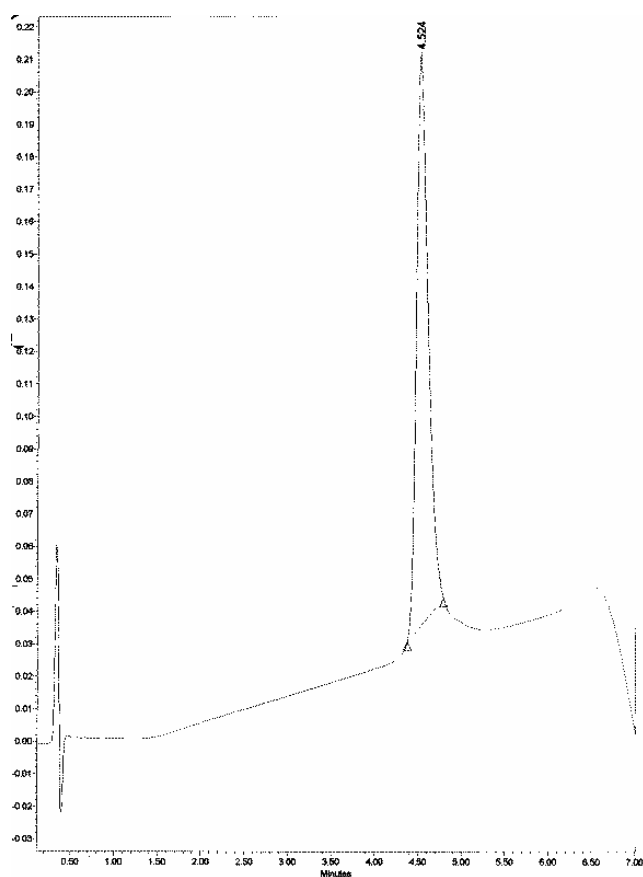
## Dendrimer DHL-DHL



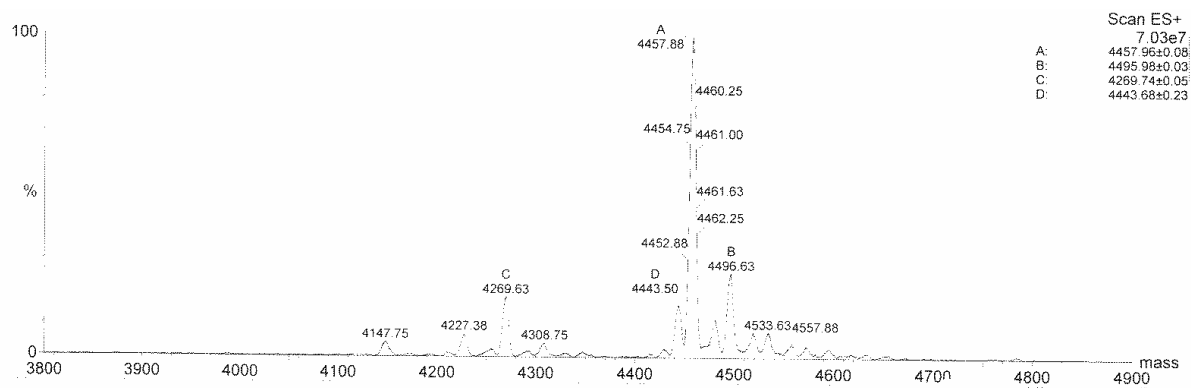
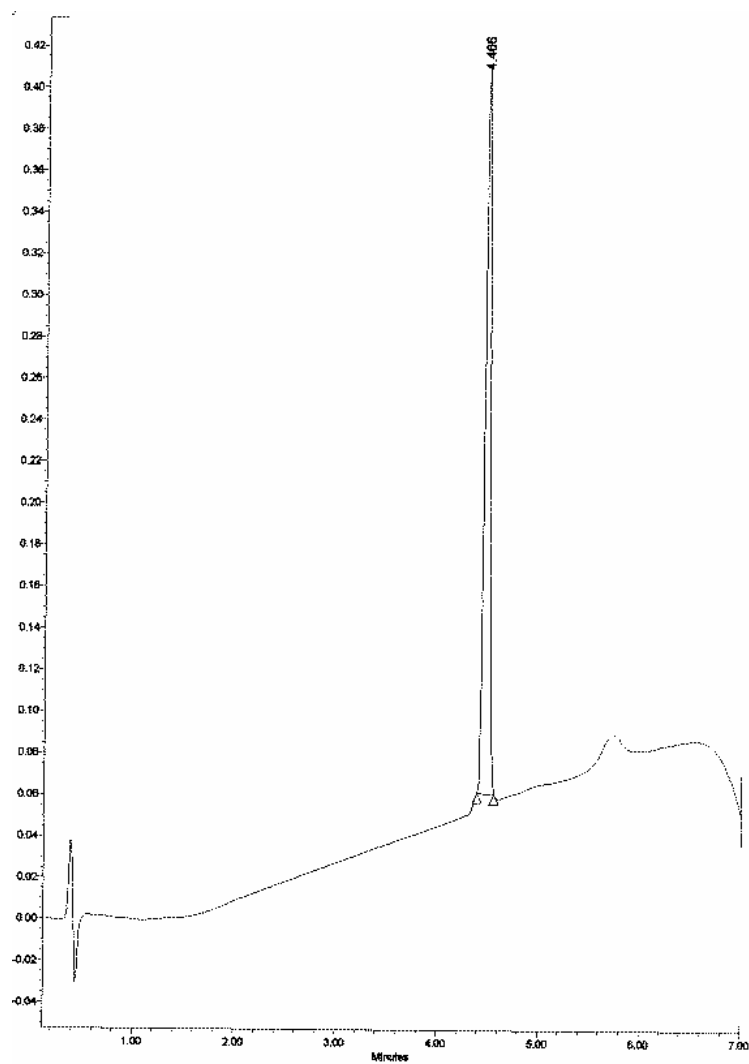
## Dendrimer LRH-LRH



## Dendrimer DFH-DFH



## Dendrimer LRH-HWS



## Dendrimer LRH-DFH

