

Solutions for Innovation

# JMS-S3000 SpiralTOF™ series

## Polymer Mass Spectra Collection

Edition November 2021





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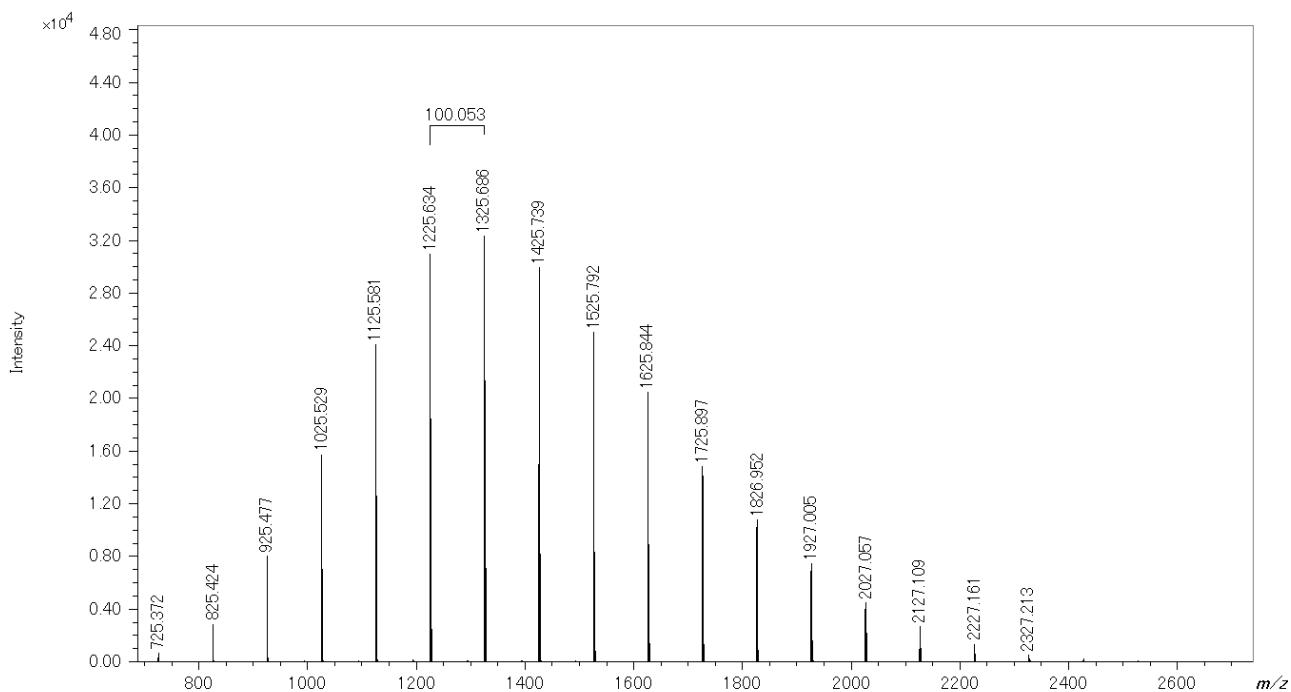
## Abbreviations

Abbreviation	Description
DCTB	<i>trans</i> -2-[3-(4- <i>tert</i> -Butylphenyl)-2-methyl-2-propenylidene] malononitrile
CHCA	α-cyano-4-hydroxy cinnamic acid
DHB	2,5-Dihydroxybenzoic Acid
HABA	2-(4-Hydroxyphenylazo)benzoic acid
THAP	2',4',6'-Trihydroxyacetophenone monohydrate
NaTFA	Sodium trifluoroacetate
AgTFA	Silver trifluoroacetate

This booklet is a collection of polymer mass spectra acquired by using JMS-S3000 SpiralTOF™-plus matrix-assisted laser desorption/ionization time-of-flight mass spectrometers (MALDI-TOFMS). Please refer the legend on page iii regarding the information attached to each mass spectrum. Note that none of the mass spectra in this booklet shows a guaranteed performance specification of the mass spectrometer.

## Legend

### PMMA (poly(methyl methacrylate))

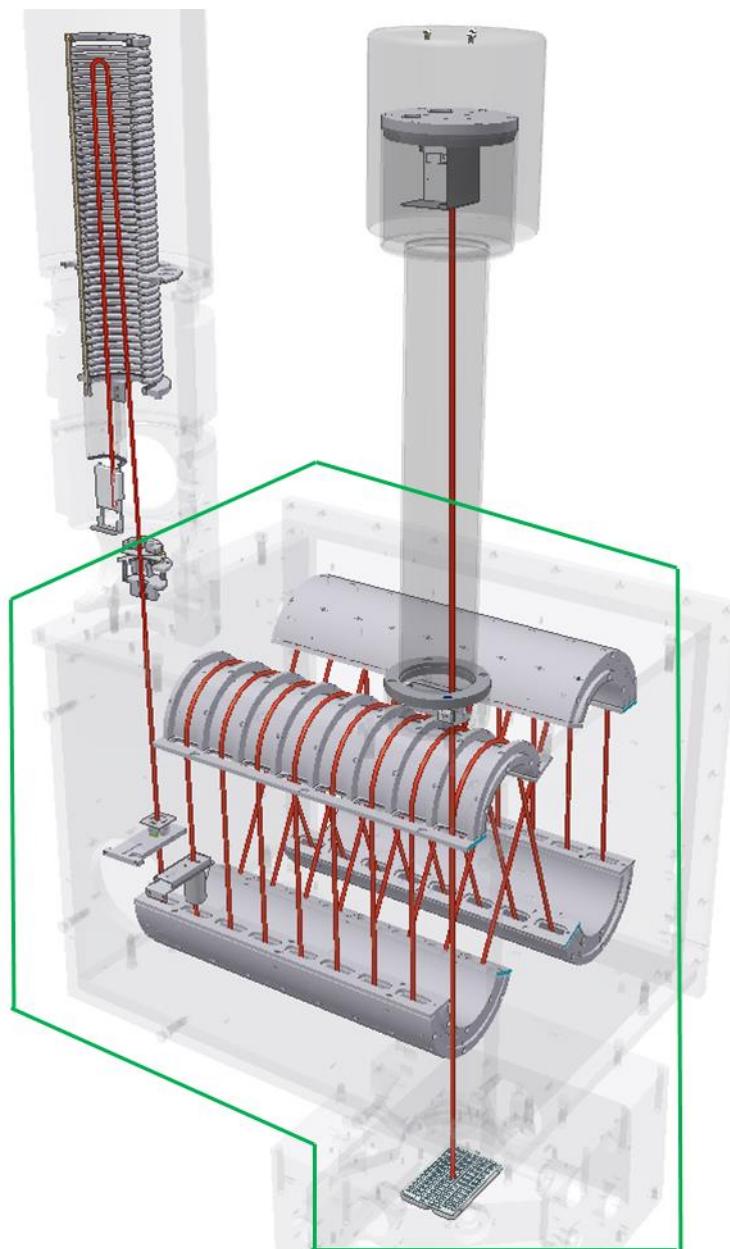


Structure	$H[CH_2C(CH_3)(CO_2CH_3)]_nH$ ; $H(C_5H_8O)_nH$	Two formulae, one showing structural information and the other showing elemental composition(s) of the repeating unit(s). If the end groups are unknown, the formula is shown as, for example: $-(C_5H_8O)_n-$
Ion Species	$[M+Na]^+$	Detected ion species
Molecular weight Information	$M_n$ 1498; $M_w$ 1559; D 1.041	Molar mass information calculated from the observed mass spectrum, NOT the values declared by the manufacturer.
Matrix	DCTB	Matrix used for the measurement
Cationization agent	NaTFA	Cationization agent used for the measurement
Mode	SpiralTOF:Positive ion	Mass analyzer mode and polarity of ions detected.

KMD (Kendrick Mass Defect) plots generated by the msRepeatFinder software are shown for the polymers other than homopolymers with only one kind of end groups.

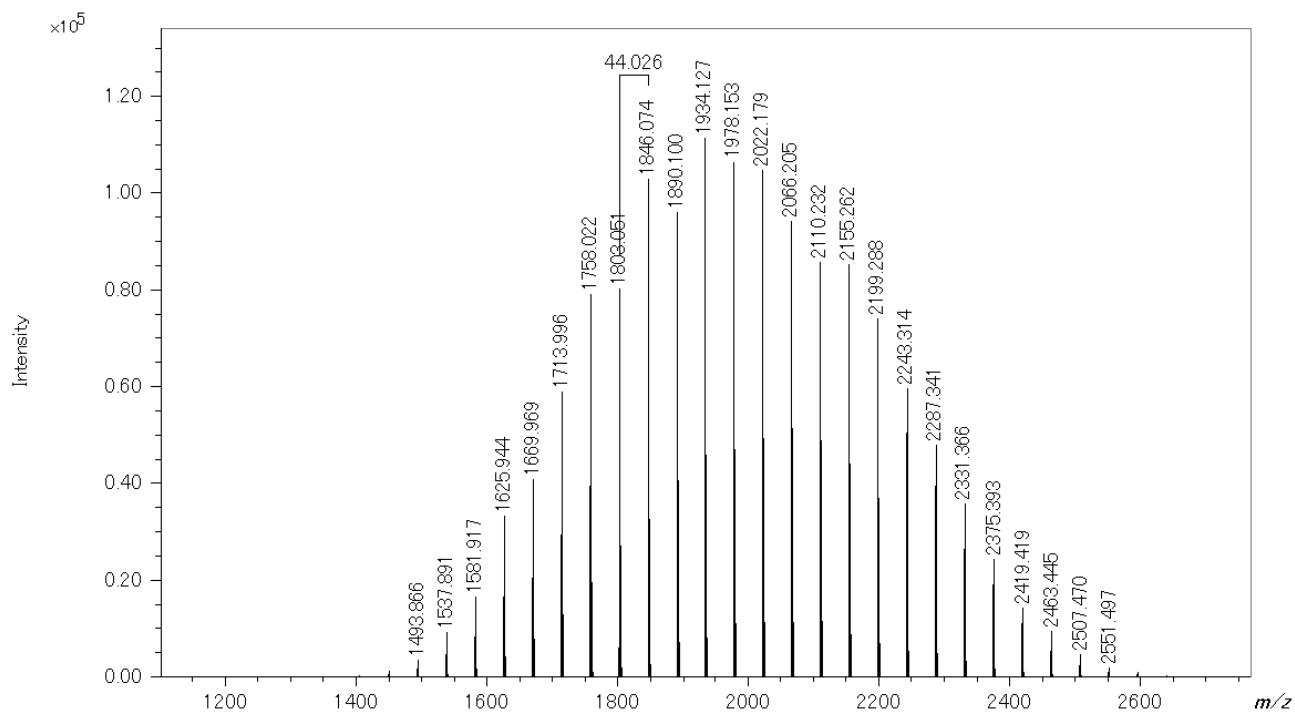
RKM (Remainder of Kendrick Mass) plots generated by the msRepeatFinder software and estimated product ion structures are shown for TOF-TOF (tandem MS) product ion mass spectra.

## SpiralTOF Mode

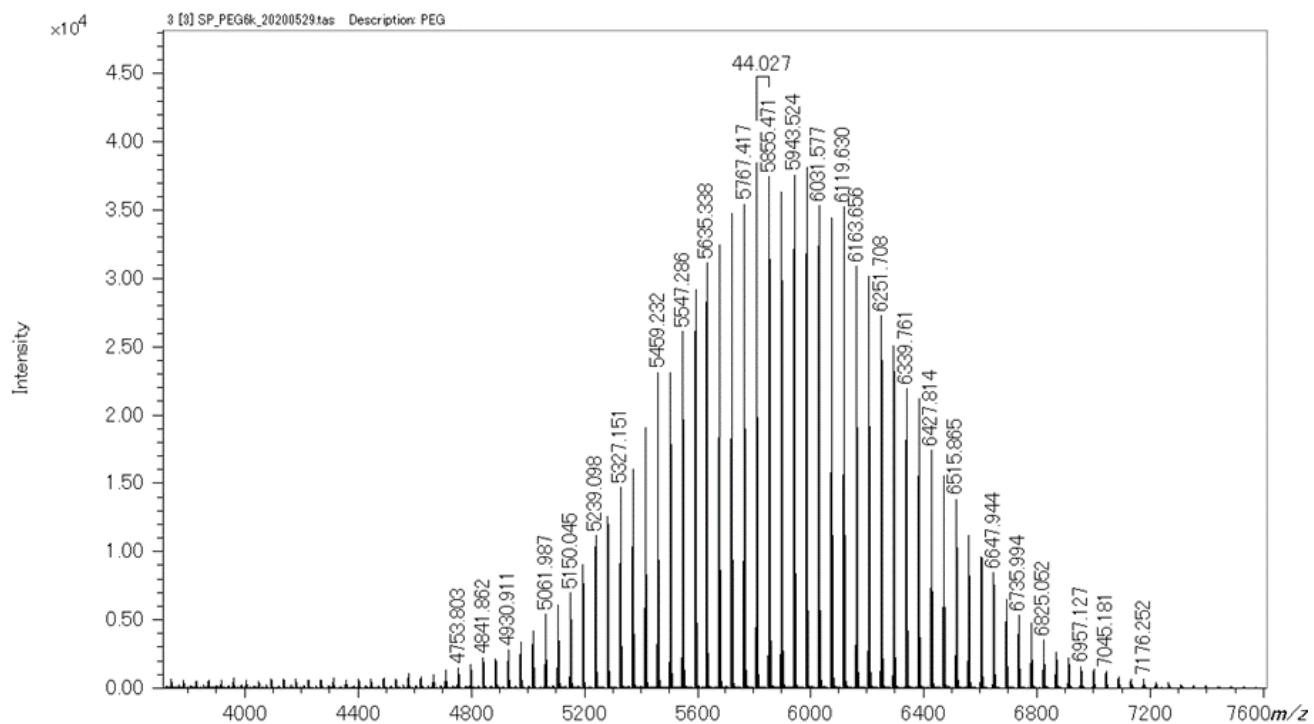


- Ultra-high mass-resolution achieved with JEOL's patented Spiral Ion Trajectory TOFMS.
- High mass-resolution and high mass-accuracy over a wide mass range – essential for polymer analysis.
- Chemical noise is reduced by eliminating post source decay (PSD) ions with electrostatic sectors – minor components can be clearly detected.

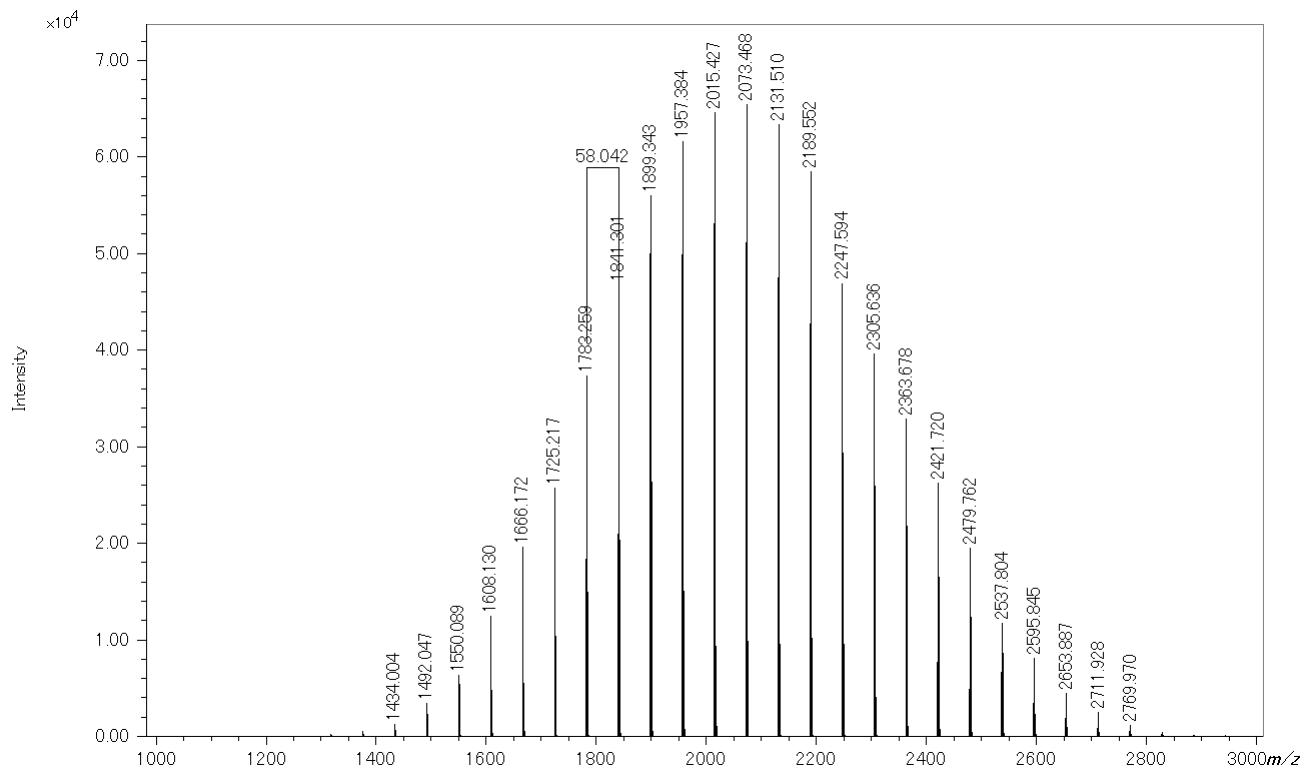


**PEO (poly(ethylene oxide)) MW 2000**

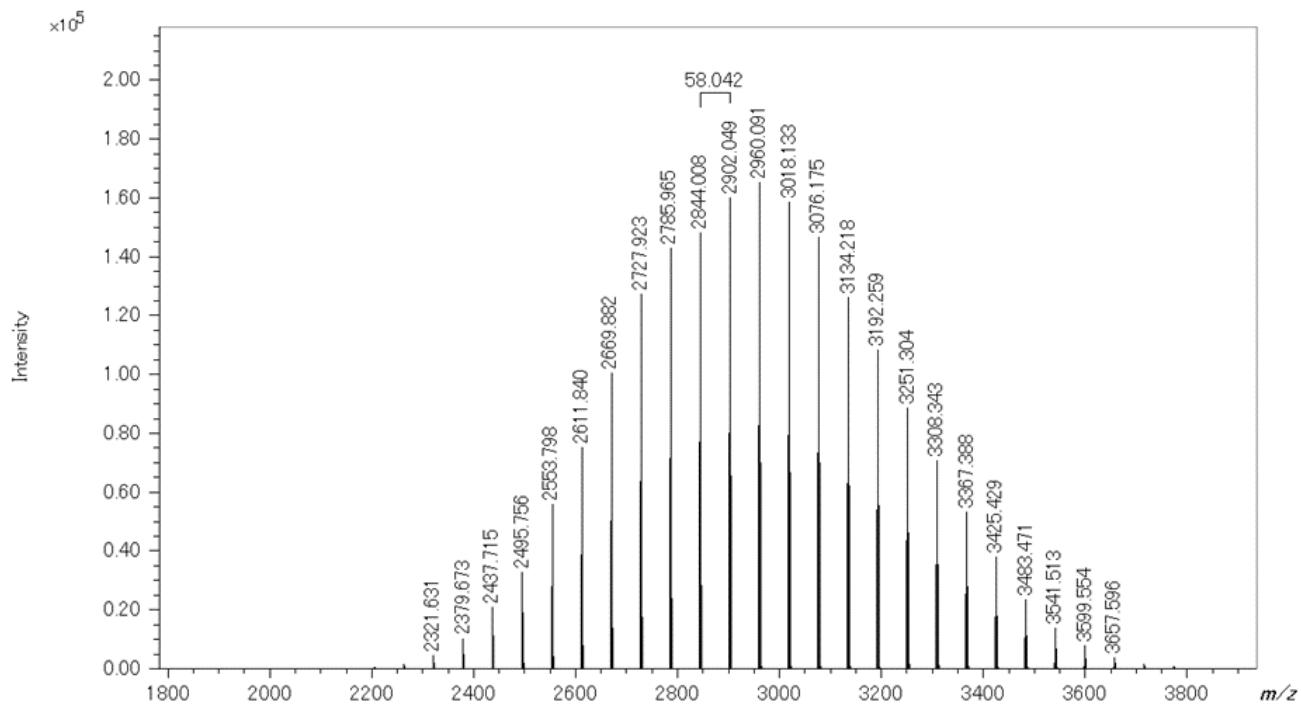
Structure	$\text{HO}(\text{CH}_2\text{CH}_2\text{O})_n\text{H}$ ; $\text{HO}(\text{C}_2\text{H}_4\text{O})_n\text{H}$
Ion Species	$[\text{M}+\text{Na}]^+$
Molecular weight Information	$M_n$ 1,998; $M_w$ 2,016; D 1.009
Matrix	CHCA
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion

**PEO (poly(ethylene oxide)) MW 6000**

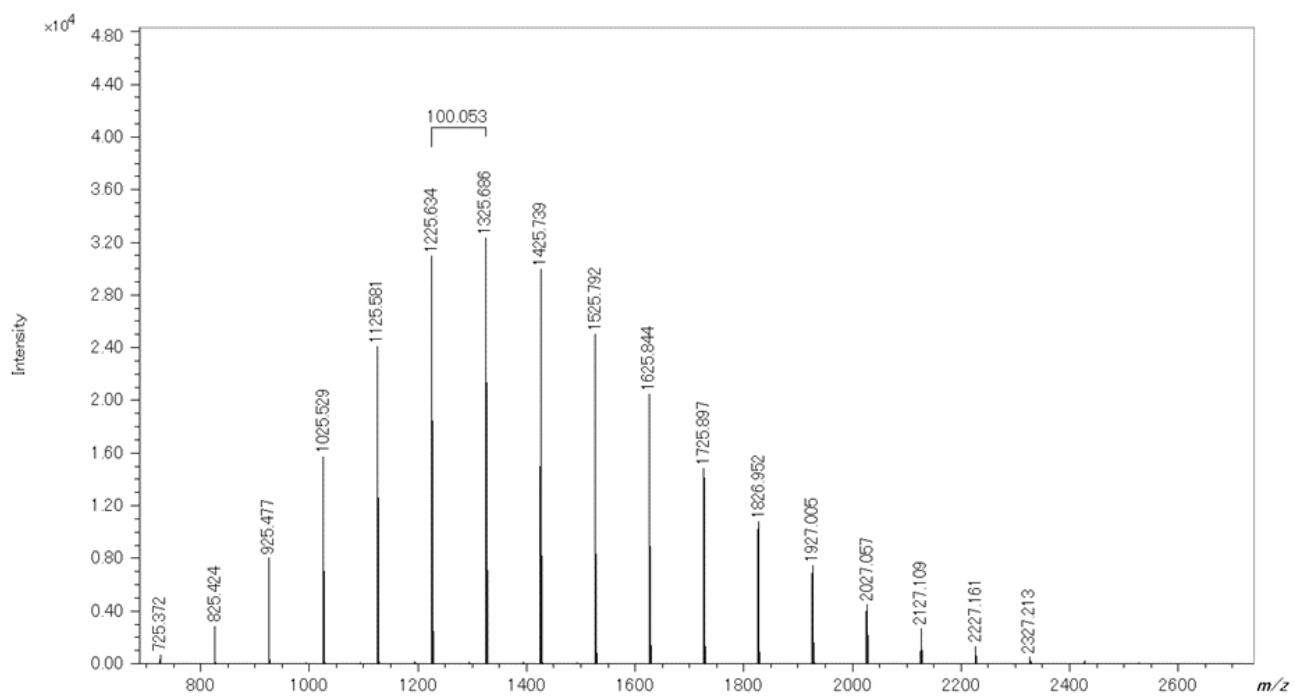
Structure	$\text{HO}(\text{CH}_2\text{CH}_2\text{O})_n\text{H}$ ; $\text{HO}(\text{C}_2\text{H}_4\text{O})_n\text{H}$
Ion Species	$[\text{M}+\text{Na}]^+$
Molecular weight Information	$M_n$ 5,878; $M_w$ 5,914; D 1.006
Matrix	DCTB
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion

**PPO (poly(propylene oxide))**

Structure	$\text{HO}[\text{CH}_2\text{CH}(\text{CH}_3)\text{O}]_n\text{H}$ ; $\text{HO}(\text{C}_3\text{H}_6\text{O})_n\text{H}$
Ion Species	$[\text{M}+\text{Na}]^+$
Molecular weight Information	$M_n$ 2,093; $M_w$ 2,119; D 1.013
Matrix	DCTB
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion

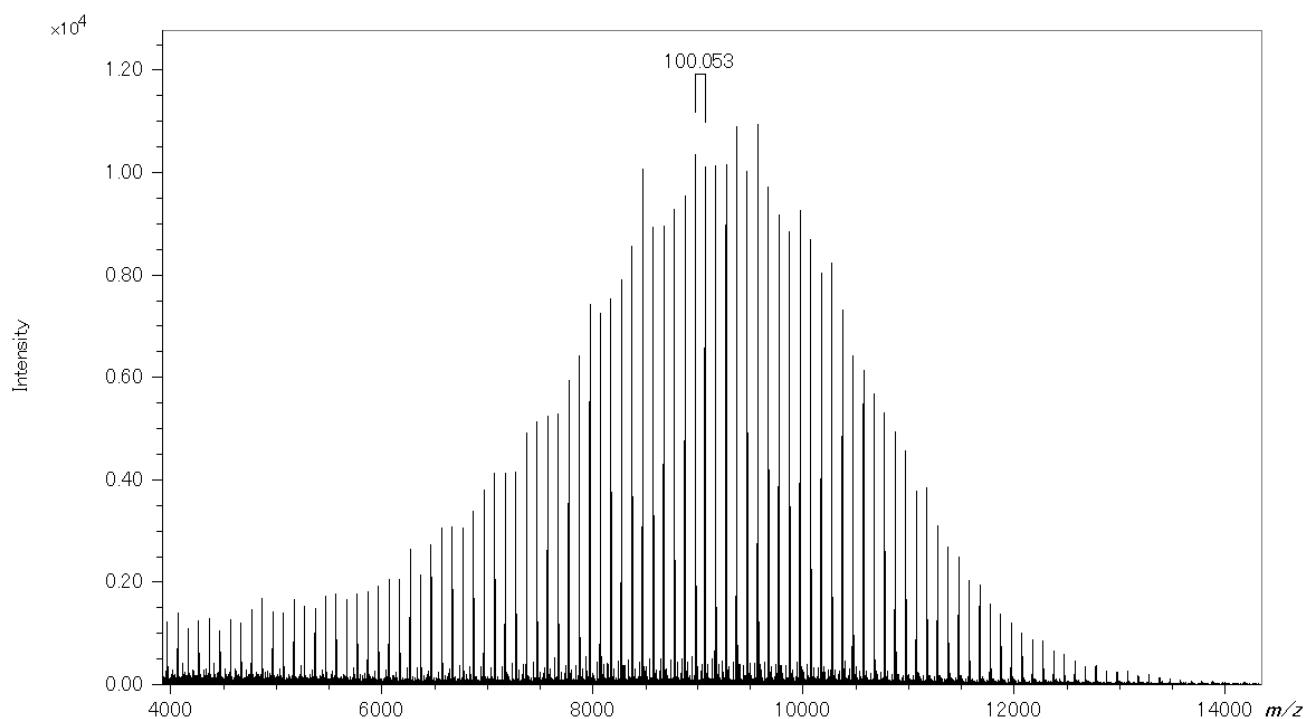
**PPO Triol (tris(polypropylene oxide) glyceryl ether)**

Structure	$\text{HO}(\text{C}_3\text{H}_6\text{O})_n\text{CH}[\text{CH}_2(\text{OC}_3\text{H}_6)_n\text{OH}]_2$ ; $(\text{C}_3\text{H}_6\text{O})_n\text{C}_3\text{H}_8\text{O}_3$
Ion Species	$[\text{M}+\text{Na}]^+$
Molecular weight Information	$M_n$ 2,982; $M_w$ 3,001; D 1.006
Matrix	DCTB
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion

**PMMA (poly(methyl methacrylate))**

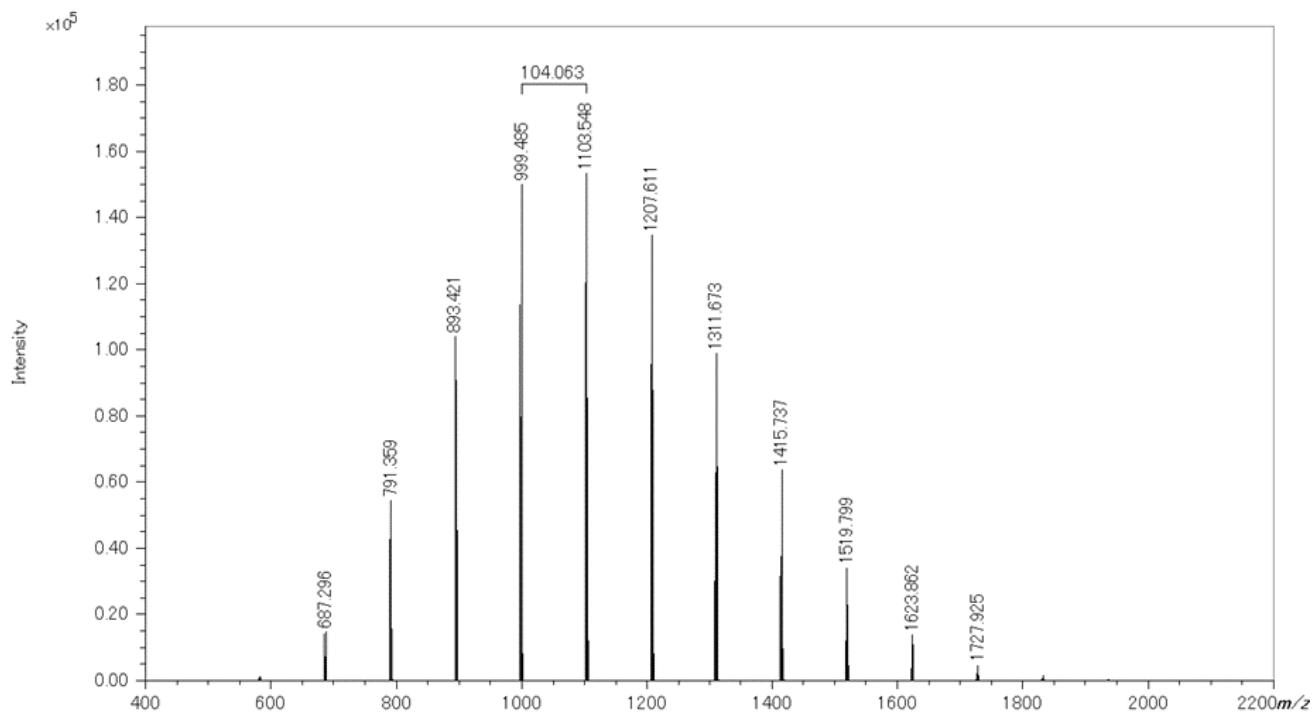
Structure	$\text{H}[\text{CH}_2\text{C}(\text{CH}_3)(\text{CO}_2\text{CH}_3)]_n\text{H}$ ; $\text{H}(\text{C}_5\text{H}_8\text{O})_n\text{H}$
Ion Species	$[\text{M}+\text{Na}]^+$
Molecular weight Information	$M_n$ 1,498; $M_w$ 1,559; D 1.041
Matrix	DCTB
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion

## PMMA (poly(methyl methacrylate))

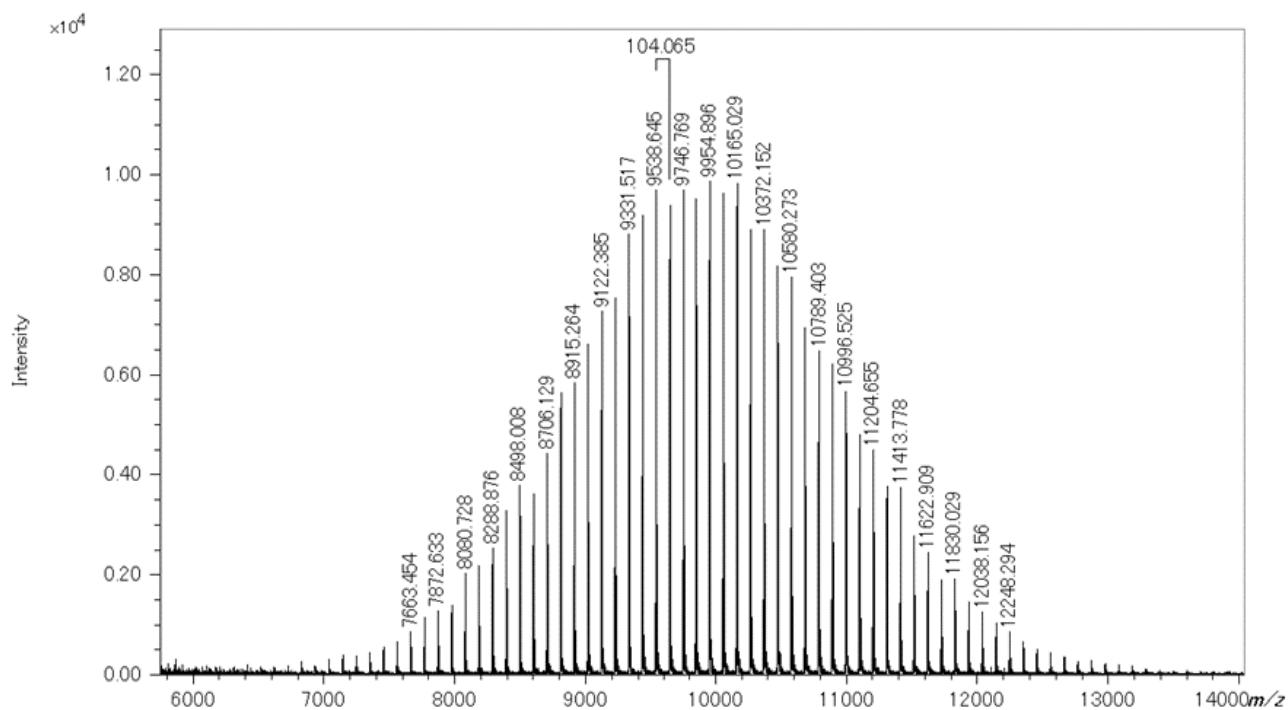


Structure	-[CH <sub>2</sub> C(CH <sub>3</sub> )(CO <sub>2</sub> CH <sub>3</sub> )] <sub>n</sub> -; -(C <sub>5</sub> H <sub>8</sub> O) <sub>n</sub> -
Ion Species	[M+Na] <sup>+</sup>
Molecular weight Information	M <sub>n</sub> 8,928; M <sub>w</sub> 9,257; D 1.037
Matrix	DCTB
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion

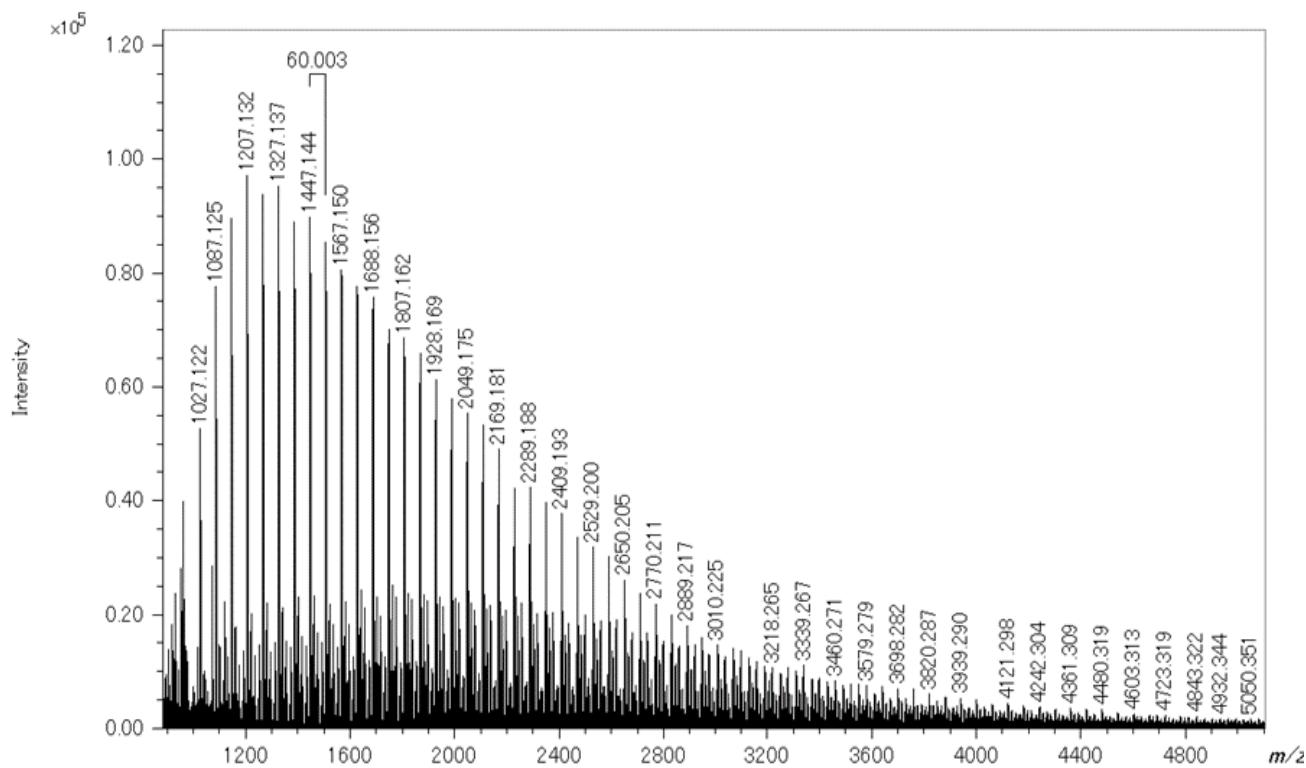
## PS (poly(styrene)) MW 1000



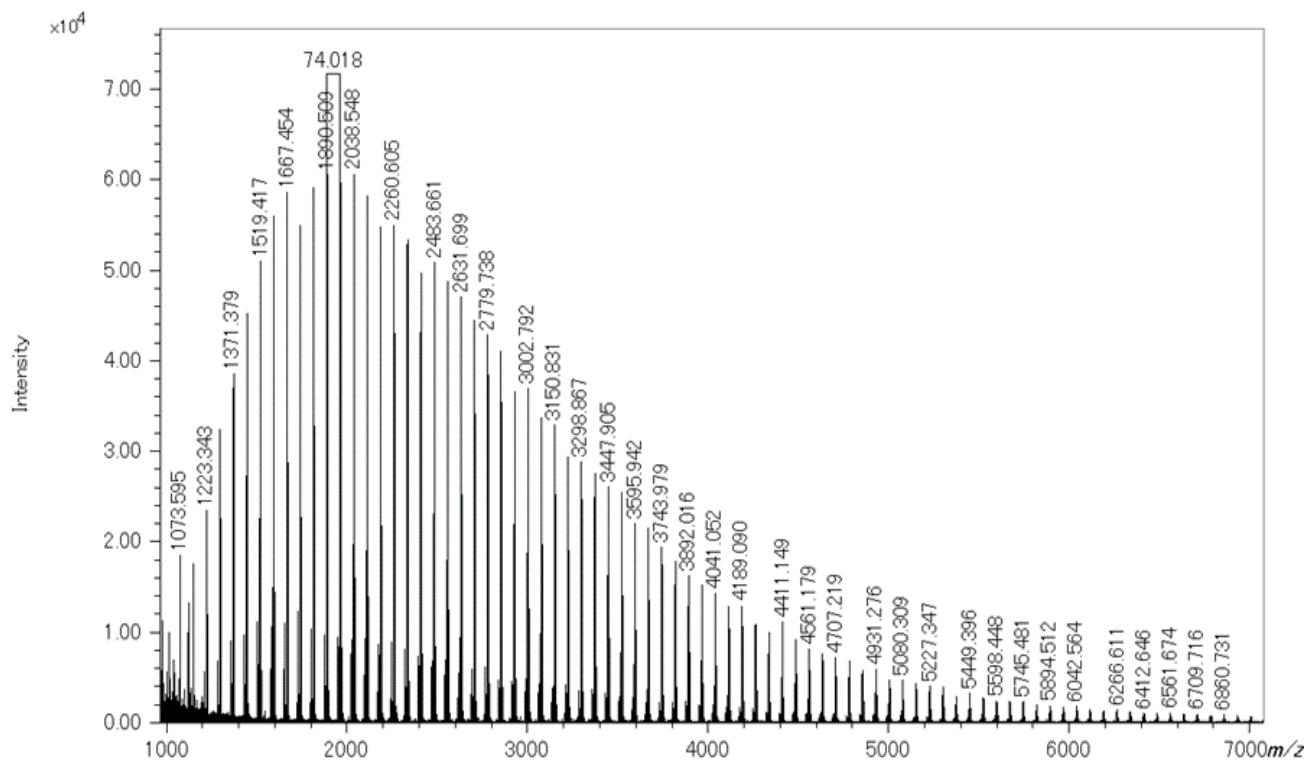
Structure	$\text{H}[\text{CH}_2\text{CH}(\text{C}_6\text{H}_5)]_n\text{C}_4\text{H}_9$ ; $\text{H}(\text{C}_8\text{H}_8)_n\text{C}_4\text{H}_9$
Ion Species	$[\text{M}+\text{Ag}]^+$
Molecular weight Information	$M_n$ 1,157; $M_w$ 1,196; D 1.033
Matrix	DCTB
Cationization agent	AgTFA
Mode	SpiralTOF:Positive ion

**PS (poly(styrene)) MW 10000**

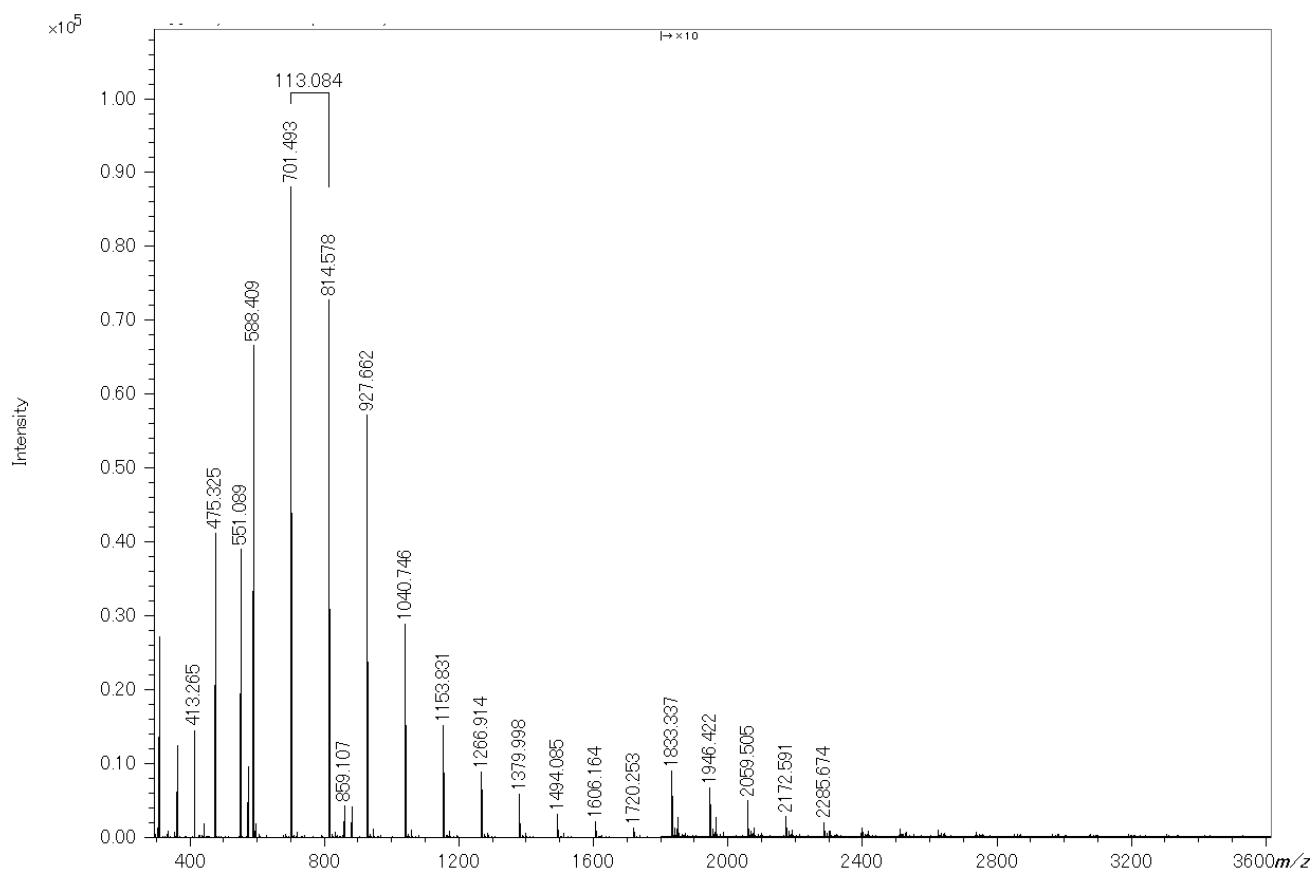
Structure	$H[CH_2CH(C_6H_5)]_nC_4H_9$ ; $H(C_8H_8)_nC_4H_9$
Ion Species	$[M+Ag]^+$
Molecular weight Information	$M_n$ 9,998; $M_w$ 10,119; D 1.012
Matrix	DCTB
Cationization agent	AgTFA
Mode	SpiralTOF:Positive ion

**PMHS (poly(methylhydrosiloxane))**

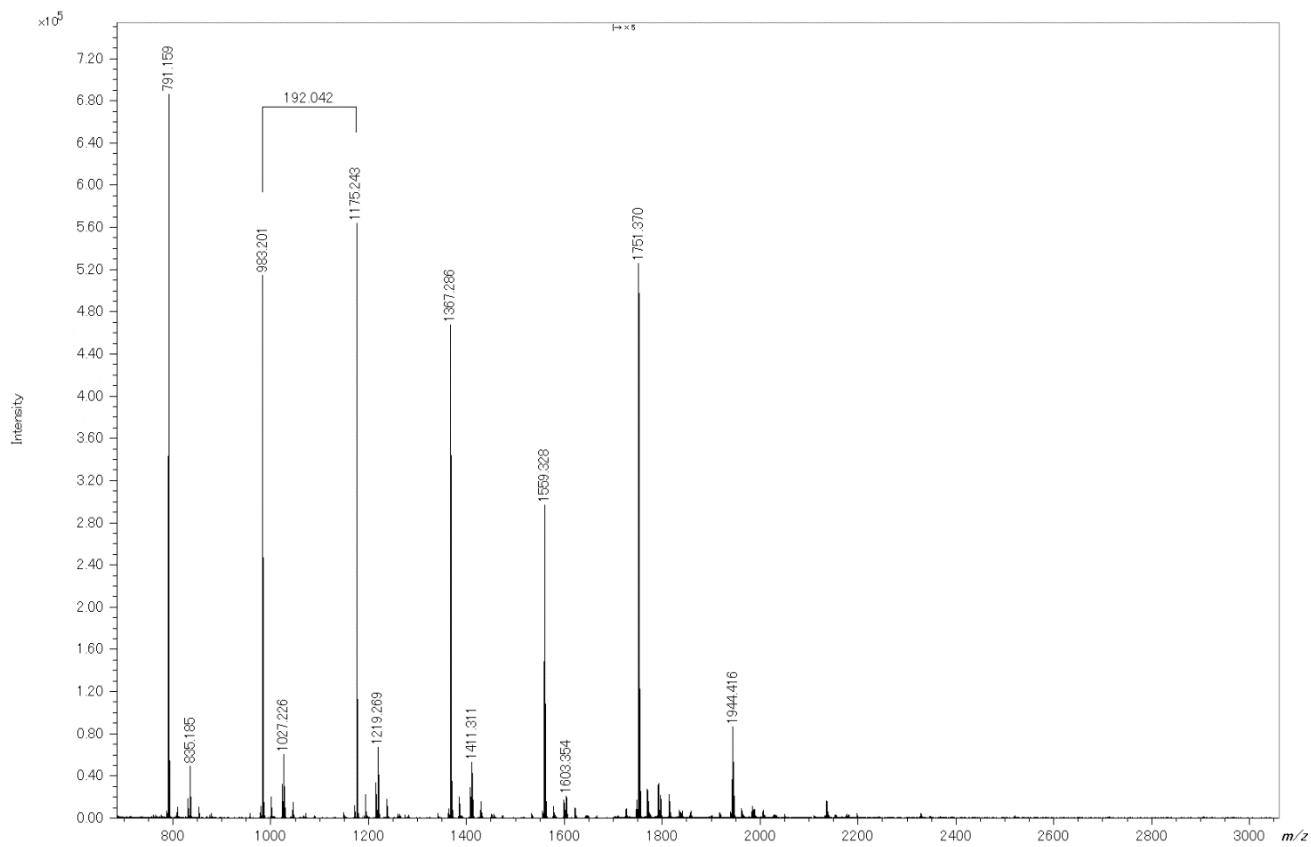
Structure	$(CH_3)_3SiO[(CH_3)HSiO]_nSi(CH_3)_3$ ; $(CH_4OSi)_nC_6H_{18}OSi_2$
Ion Species	$[M+Na]^+$
Molecular weight Information	$M_n$ 2,123; $M_w$ 2,602; D 1.226
Matrix	DCTB
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion

**PDMS (poly(dimethylsiloxane))**

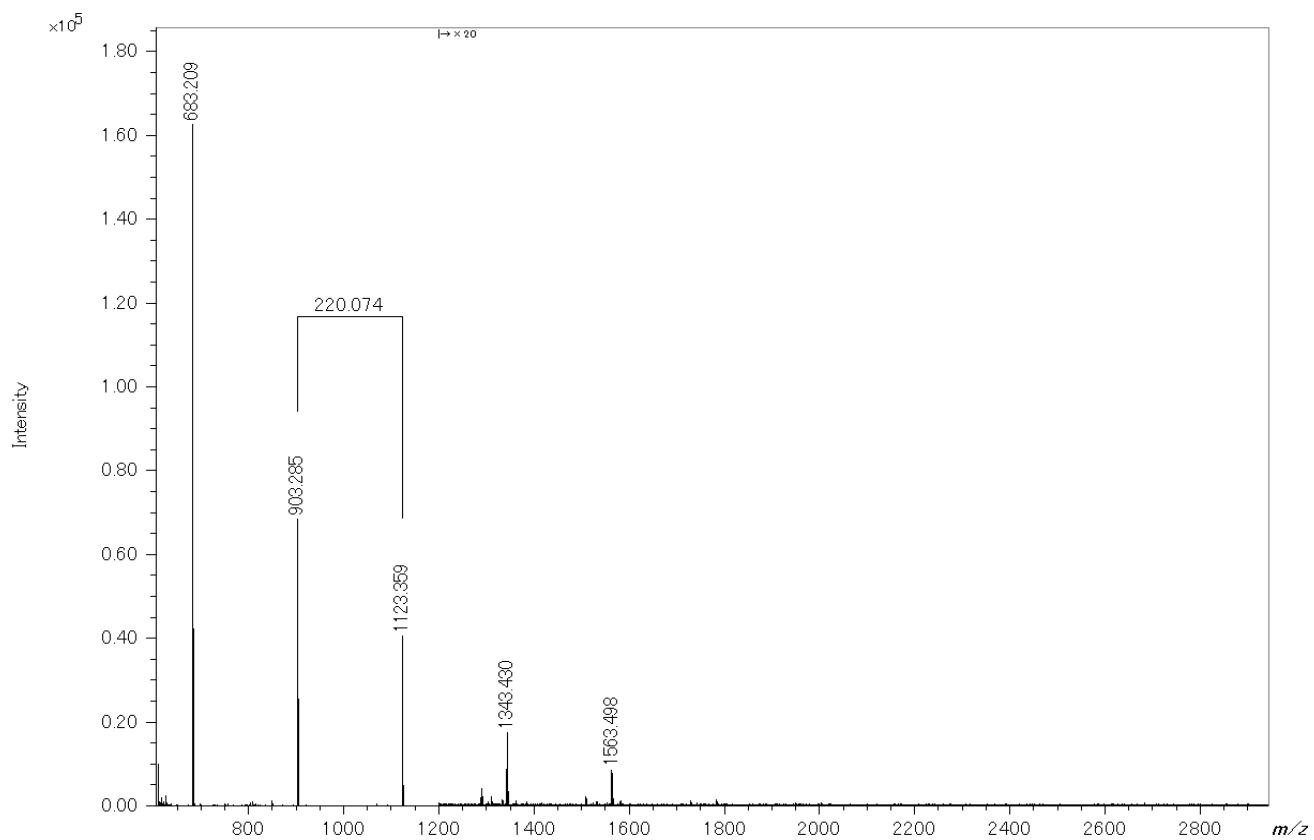
Structure	$-[(CH_3)_2SiO]_n^-$ ; $-(C_2H_6SiO)_n^-$
Ion Species	$[M+Na]^+$
Molecular weight Information	$M_n$ 2,651; $M_w$ 3,306; D 1.247
Matrix	DCTB
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion

**Nylon 6 (polycaprolactam; poly(hexano-6-lactam))**

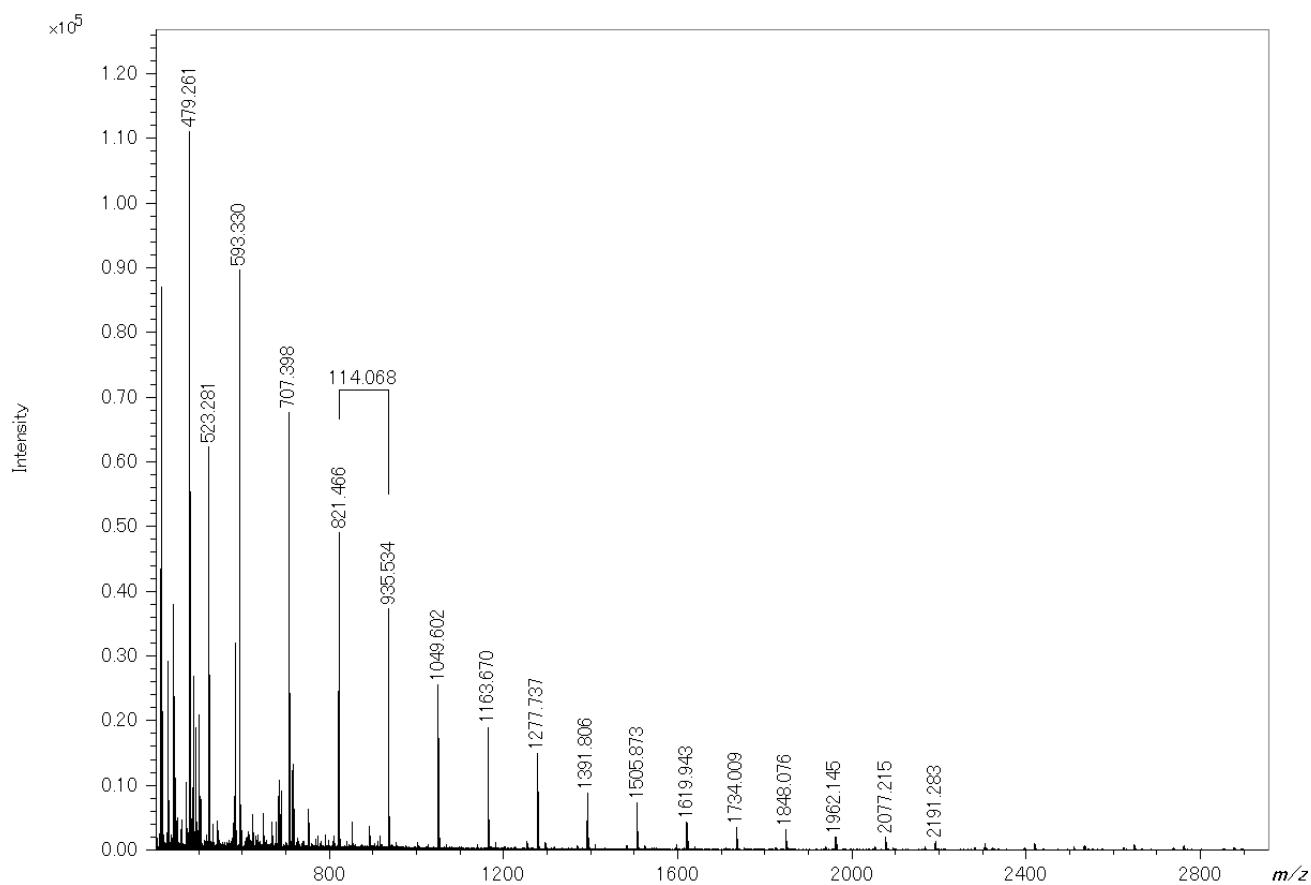
Structure	$-[NH(CH_2)_5CO]_n^-$ ; $-(C_6H_{11}NO)_n^-$
Ion Species	$[M+Na]^+$
Molecular weight Information	$M_n$ 761; $M_w$ 900; D 1.182
Matrix	HABA
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion

**PET (poly(ethylene terephthalate))**

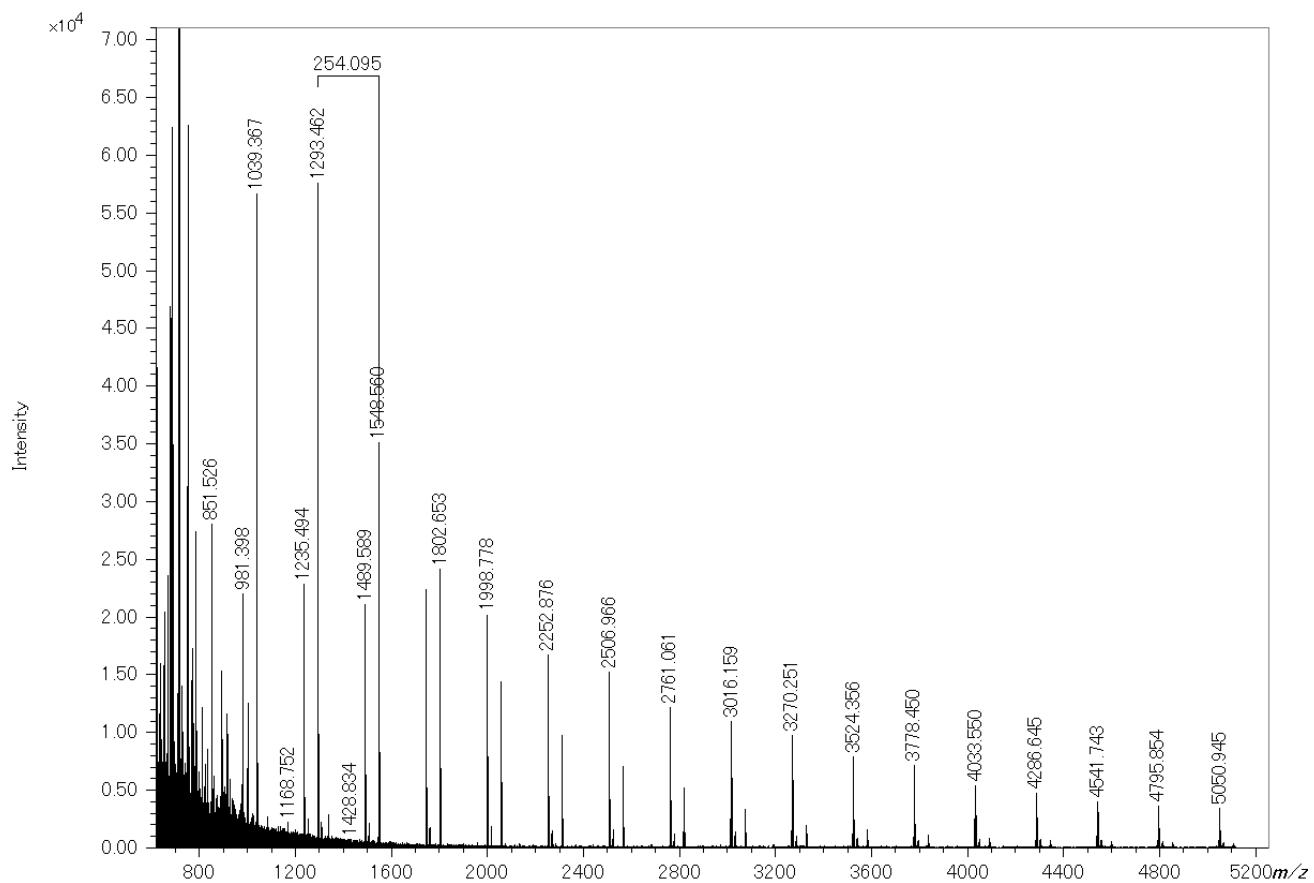
Structure	$-[COO(C_6H_4)COO(CH_2)_2]_n-$ ; $-(C_{10}H_8O_4)_n-$
Ion Species	$[M+Na]^+$
Molecular weight Information	$M_n$ 1203, $M_w$ 1281, D 1.065
Matrix	THAP
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion

**PBT (poly(butylene terephthalate))**

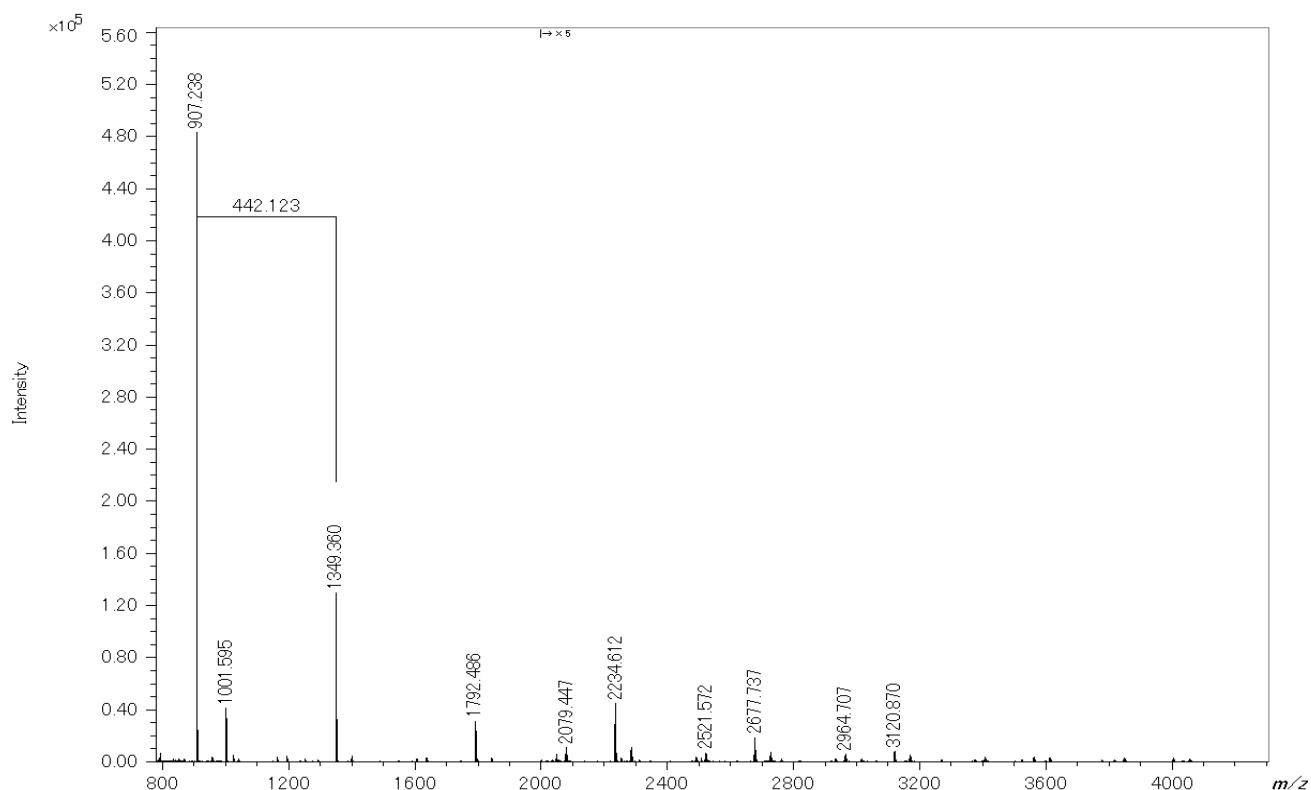
Structure	$-[CO(C_6H_4)COO(CH_2)_4O]_n^-$ ; $-(C_{12}H_{14}O_4)_n^-$
Ion Species	$[M+Na]^+$
Molecular weight Information	$M_n$ 790; $M_w$ 852; D 1.079
Matrix	DHB
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion

**PCL (poly(caprolactone))**

Structure	$-\text{O}(\text{CH}_2)_5\text{CO}]_n^-$ ; $-(\text{C}_6\text{H}_{10}\text{O}_2)_n^-$
Ion Species	$[\text{M}+\text{Na}]^+$
Molecular weight Information	$\text{M}_n$ 915; $\text{M}_w$ 1,268; D 1.386
Matrix	DCTB
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion

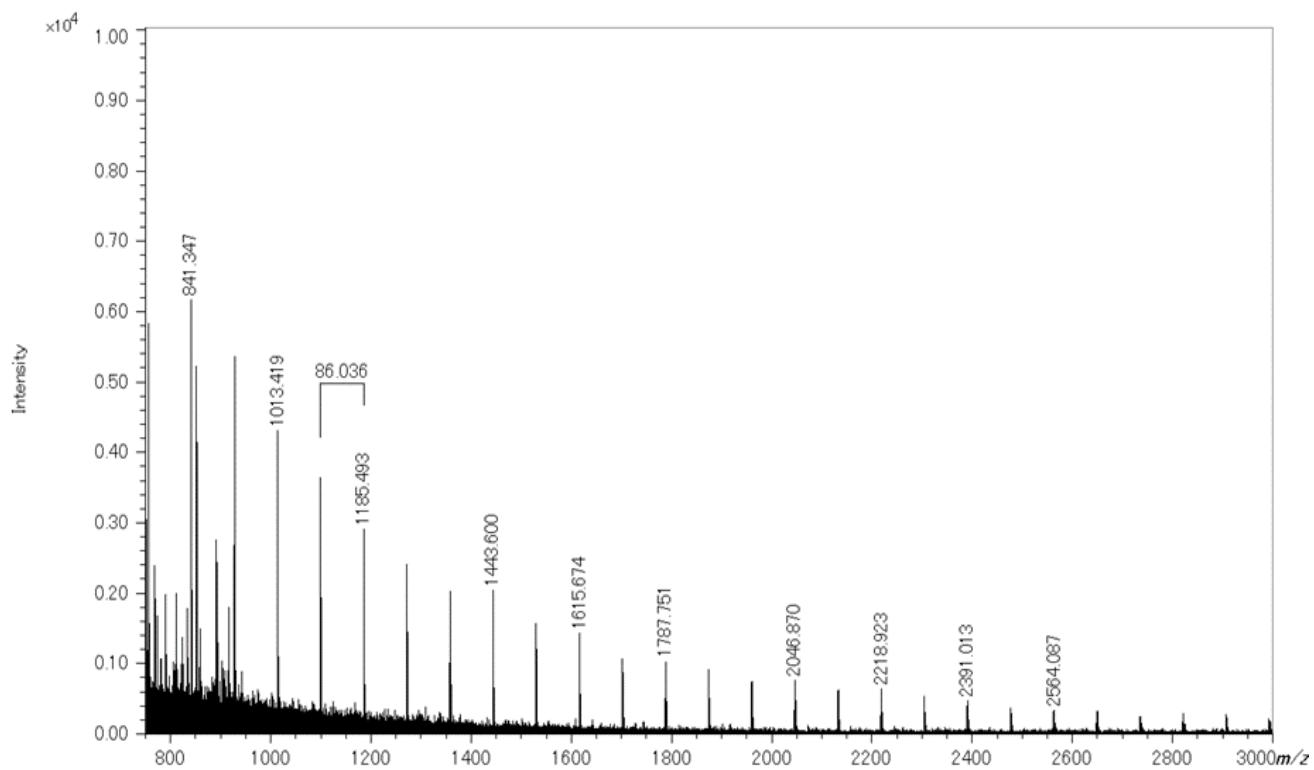
**PC (polycarbonate)**

Structure	$-[O(C_6H_4)C(CH_3)_2(C_6H_4)OCO]_n-$ ; $-(C_{16}H_{14}O_3)_n-$
Ion Species	$[M+Na]^+$
Molecular weight Information	$M_n$ 2,522; $M_w$ 3,113; D 1.236
Matrix	DCTB
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion

**PSF (polysulfone)**

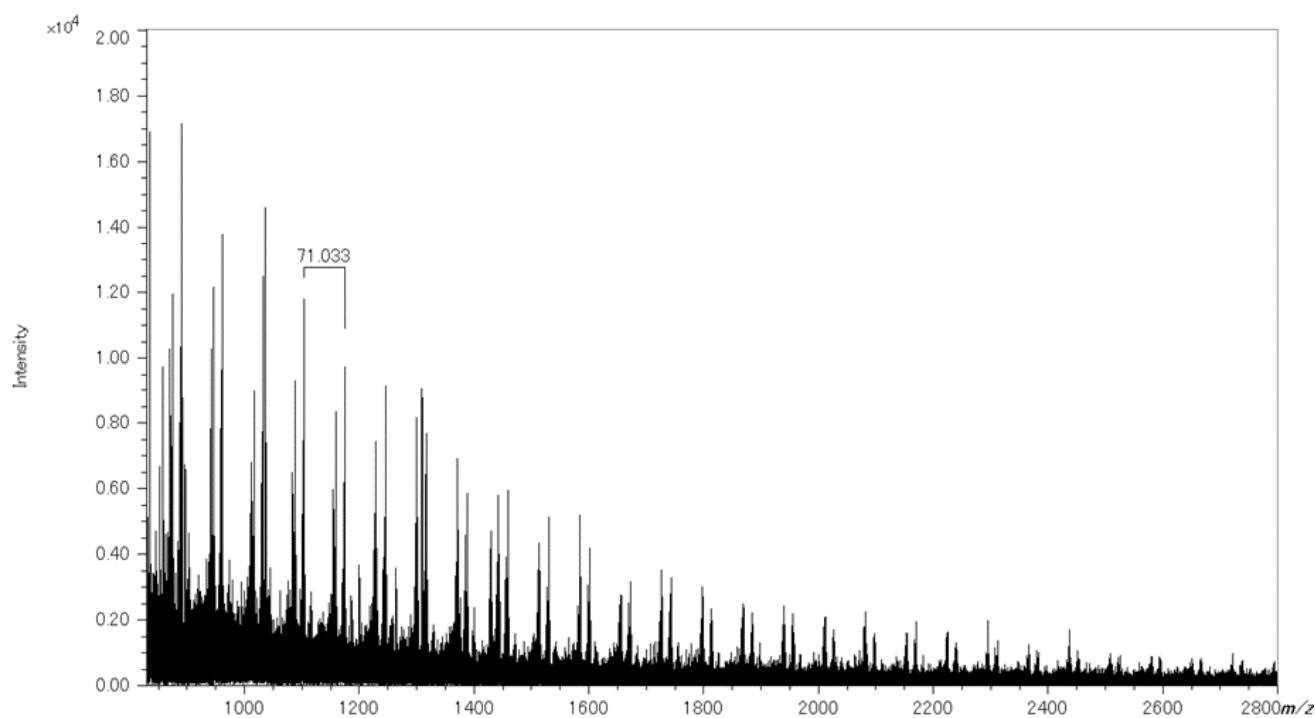
Structure	$-\left[(C_6H_4)C(CH_3)_2(C_6H_4)O(C_6H_4)SO_2(C_6H_4)O\right]_n;$ $-(C_{27}H_{22}SO_4)_n-$
Ion Species	$[M+Na]^+$
Molecular weight Information	$M_n$ 1,294; $M_w$ 1,559; D 1.205
Matrix	DCTB
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion

## PVAc (polyvinyl acetate; poly(ethenyl ethanoate))

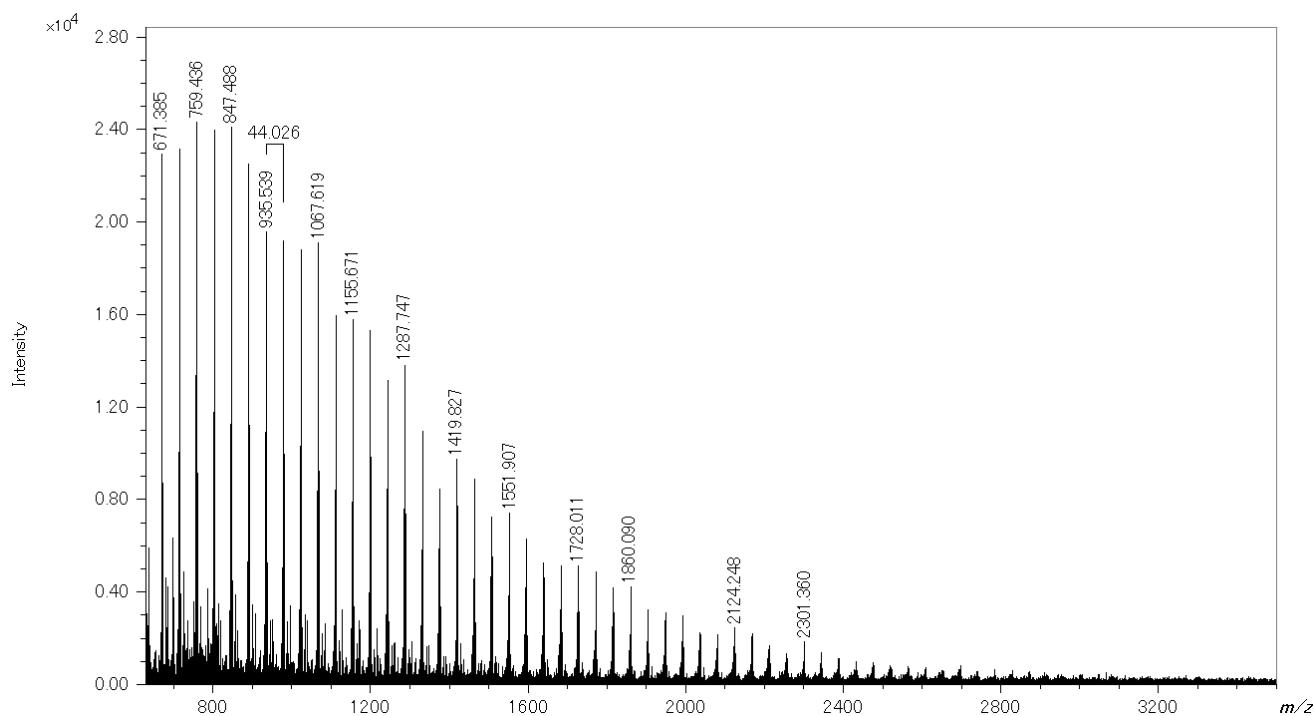


Structure	-[CH <sub>2</sub> CH(O <sub>2</sub> CCH <sub>3</sub> )] <sub>n</sub> -; -(C <sub>4</sub> H <sub>6</sub> O <sub>2</sub> ) <sub>n</sub> -
Ion Species	[M+Na] <sup>+</sup>
Molecular weight Information	$M_n$ 1,612; $M_w$ 1,818; D 1.128
Matrix	DCTB
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion

## PPA (polyacrylamide; poly(2-propenamide))

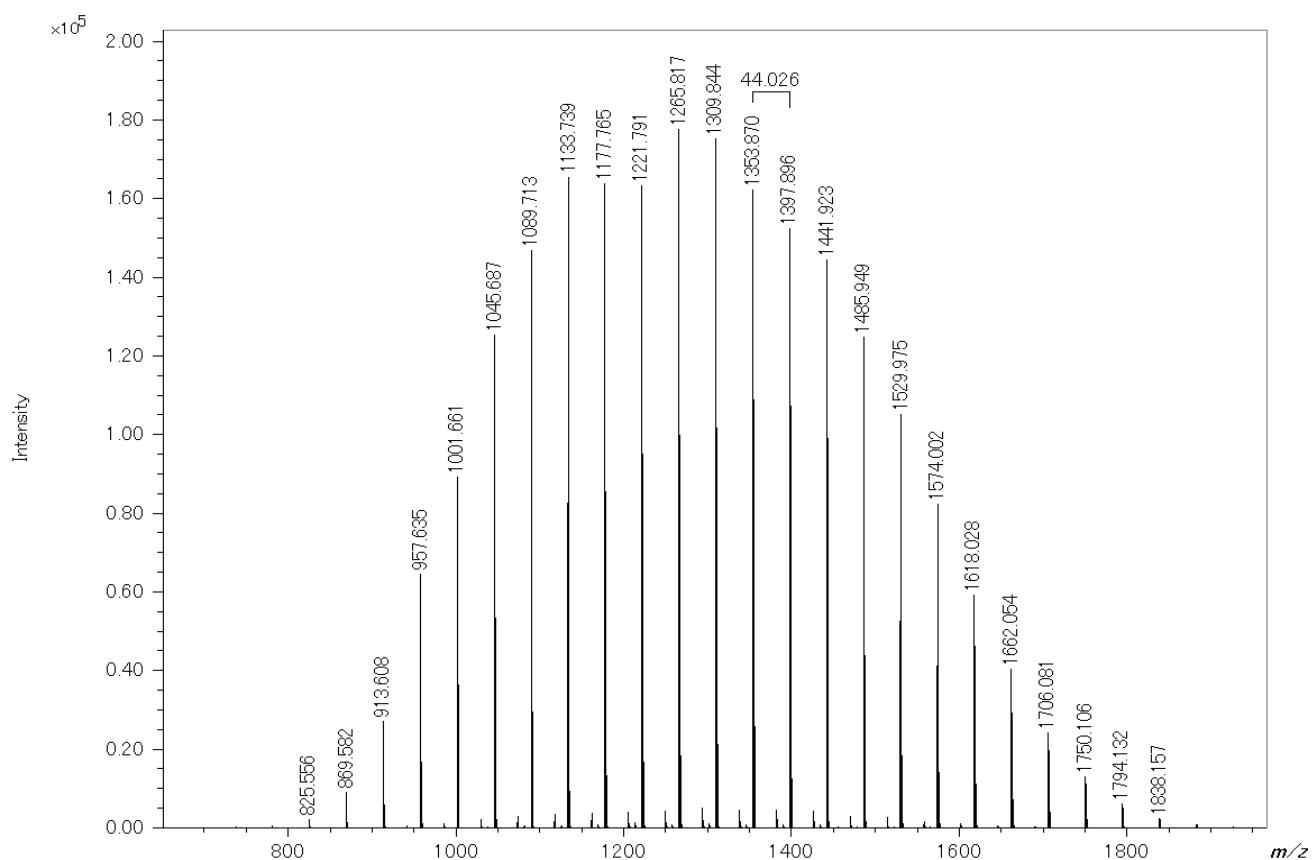


Structure	-[CH <sub>2</sub> CH(CONH <sub>2</sub> )] <sub>n</sub> -; -(C <sub>3</sub> H <sub>5</sub> NO) <sub>n</sub> -
Ion Species	[M+Na] <sup>+</sup>
Molecular weight Information	
Matrix	DHB
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion

**PVA (poly(vinyl alcohol))**

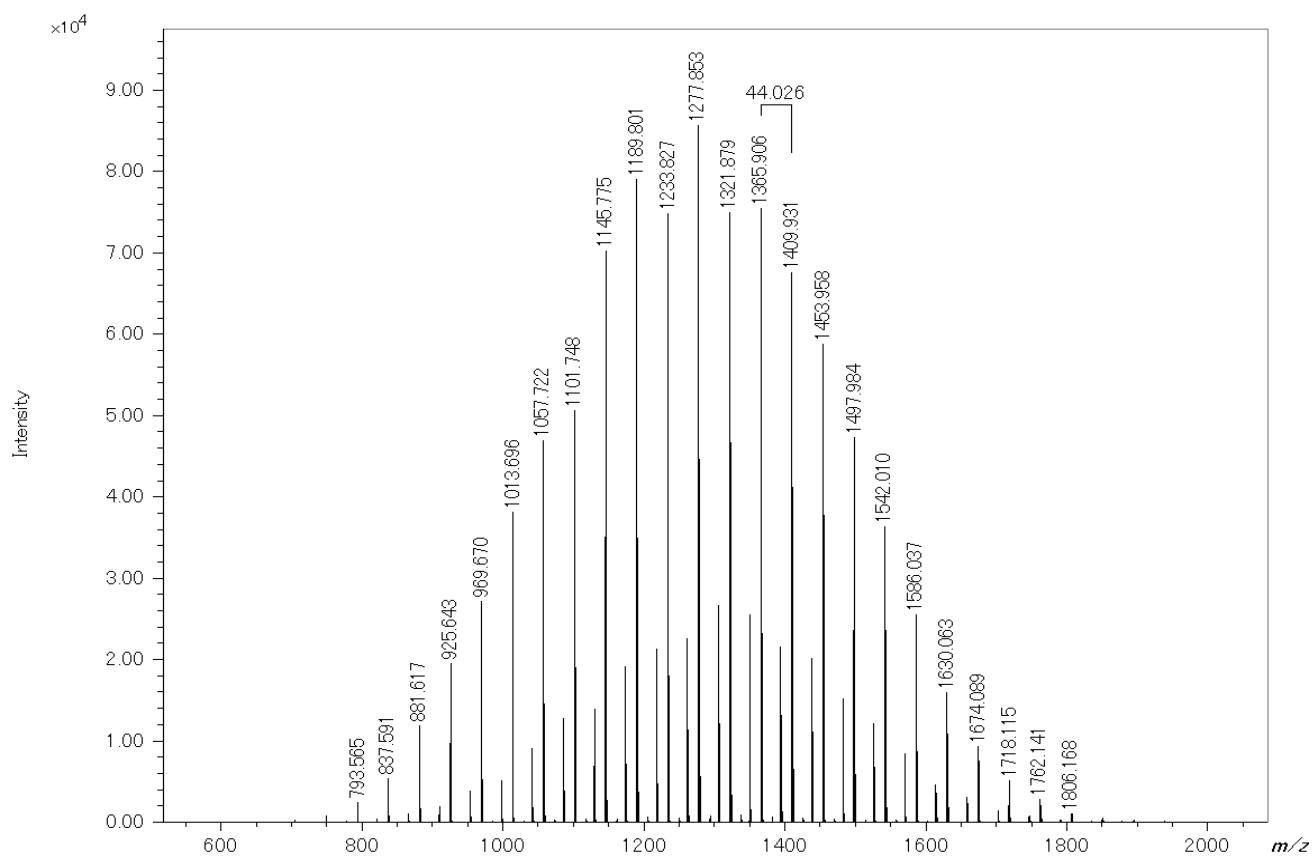
Structure	$-[\text{CH}_2\text{CH}(\text{OH})]_n-$ ; $-(\text{C}_2\text{H}_4\text{O})_n-$
Ion Species	$[\text{M}+\text{Na}]^+$
Molecular weight Information	$M_n$ 1,222; $M_w$ 1,526; D 1.249
Matrix	DHB
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion

## Polyethylene glycol monododecyl ether



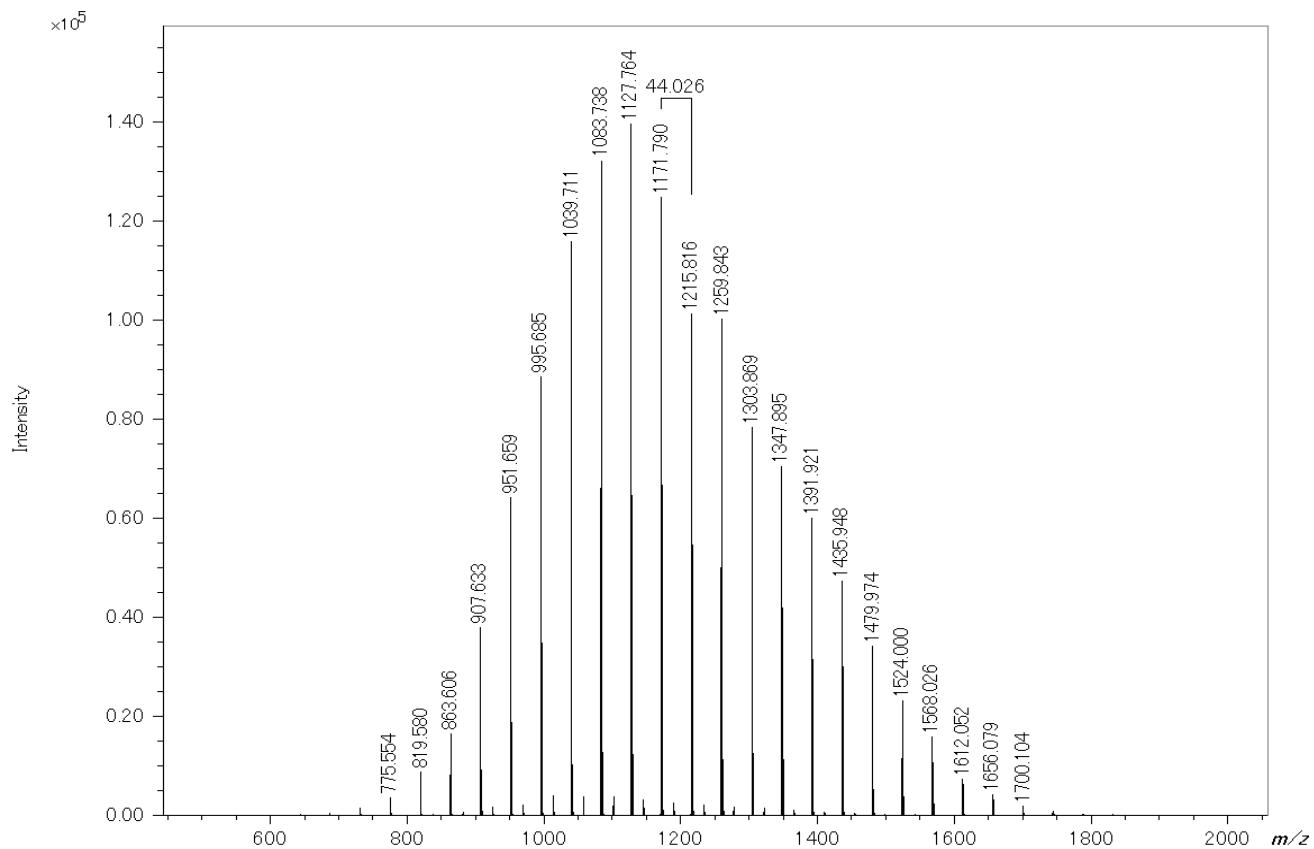
Structure	$\text{HO}[(\text{CH}_2)_2\text{O}]_n(\text{CH}_2)_{11}\text{CH}_3$ ; $\text{HO}(\text{C}_2\text{H}_4\text{O})_n\text{C}_{12}\text{H}_{25}$
Ion Species	$[\text{M}+\text{Na}]^+$
Molecular weight Information	$M_n$ 1,331; $M_w$ 1,359; D 1.021
Matrix	DCTB
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion

## Polyethylene glycol monocetyl ether



Structure	$\text{HO}[(\text{CH}_2)_2\text{O}]_n(\text{CH}_2)_{15}\text{CH}_3$ ; $\text{HO}(\text{C}_2\text{H}_4\text{O})_n\text{C}_{16}\text{H}_{33}$
Ion Species	$[\text{M}+\text{Na}]^+$
Molecular weight Information	$M_n$ 1,312; $M_w$ 1,332; D 1.016
Matrix	DCTB
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion

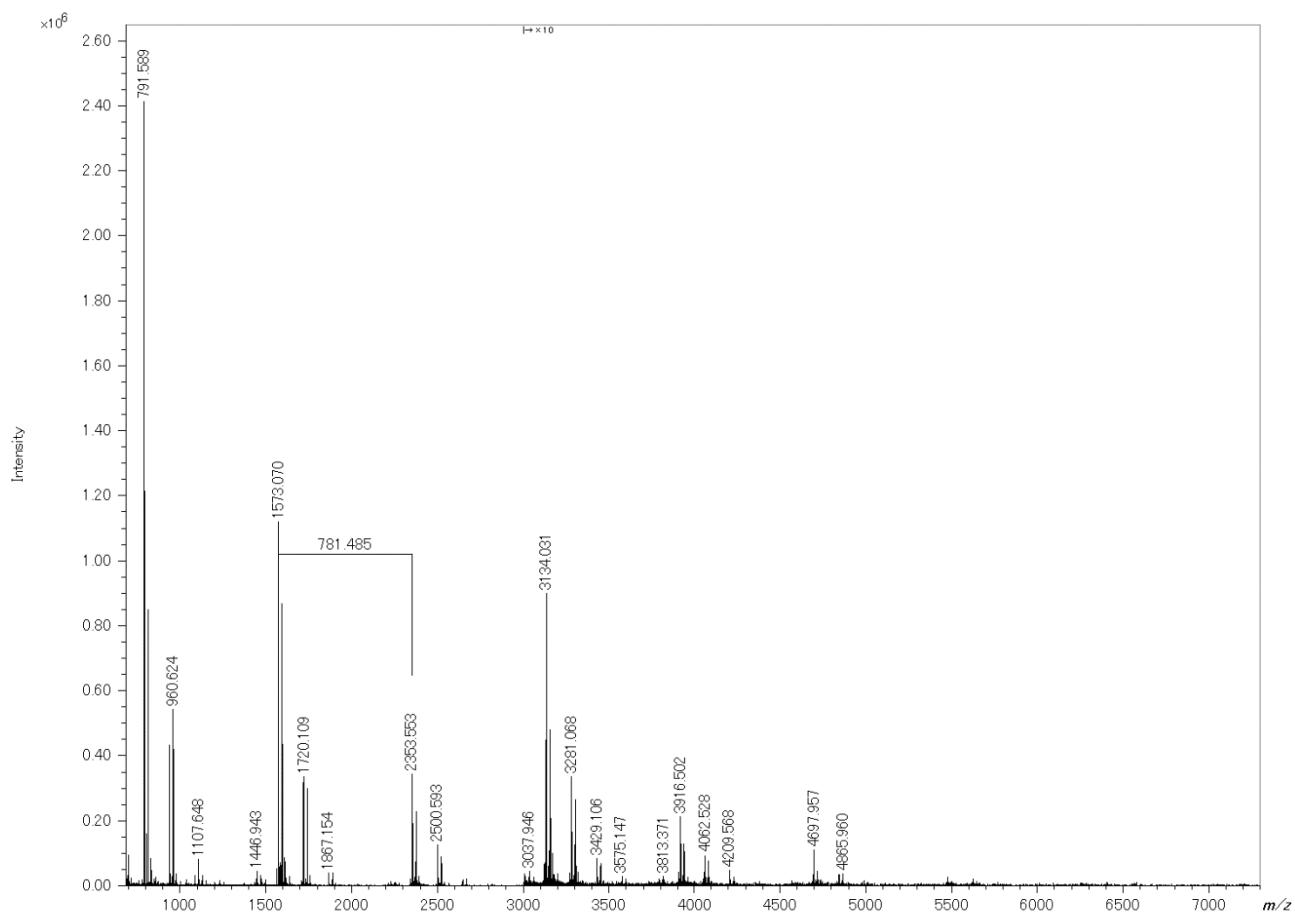
## Polyethylene glycol monooleyl ether



Structure	$\text{HO}[(\text{CH}_2)_2\text{O}]_n(\text{CH}_2)_{17}\text{CH}_3$ ; $\text{HO}(\text{C}_2\text{H}_4\text{O})\text{C}_{18}\text{H}_{35}$
Ion Species	$[\text{M}+\text{Na}]^+$
Molecular weight Information	$M_n$ 1,172; $M_w$ 1,196; D 1.021
Matrix	DCTB
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion

**ADK STAB LA-68**

(High molecular weight HALS (hindered amine light stabilizer))

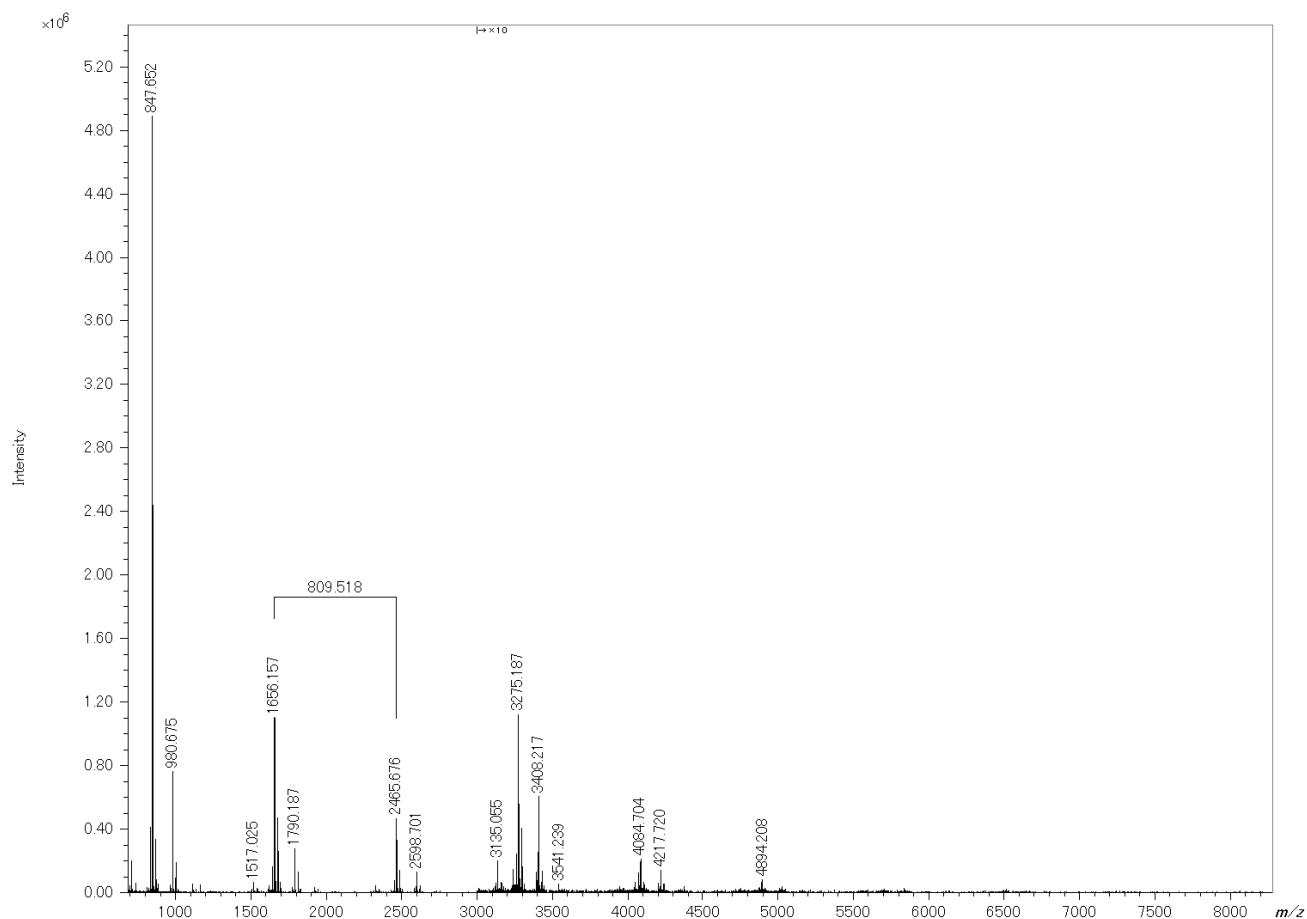


Structure	$C_9H_{18}N[C_{41}H_{68}N_2O_{12}]_n-C_{35}H_{60}O_8N_3$
Ion Species	$[M+H]^+$ , $[M+Na]^+$
Molecular weight Information	$M_n$ 1298, $M_w$ 1538, D 1.202
Matrix	DHB
Cationization agent	None
Mode	SpiralTOF:Positive ion

\* ADK STAB is either a registered trademark or a trademark of ADEKA Corporation in Japan and/or other countries.

**ADK STAB LA-63P**

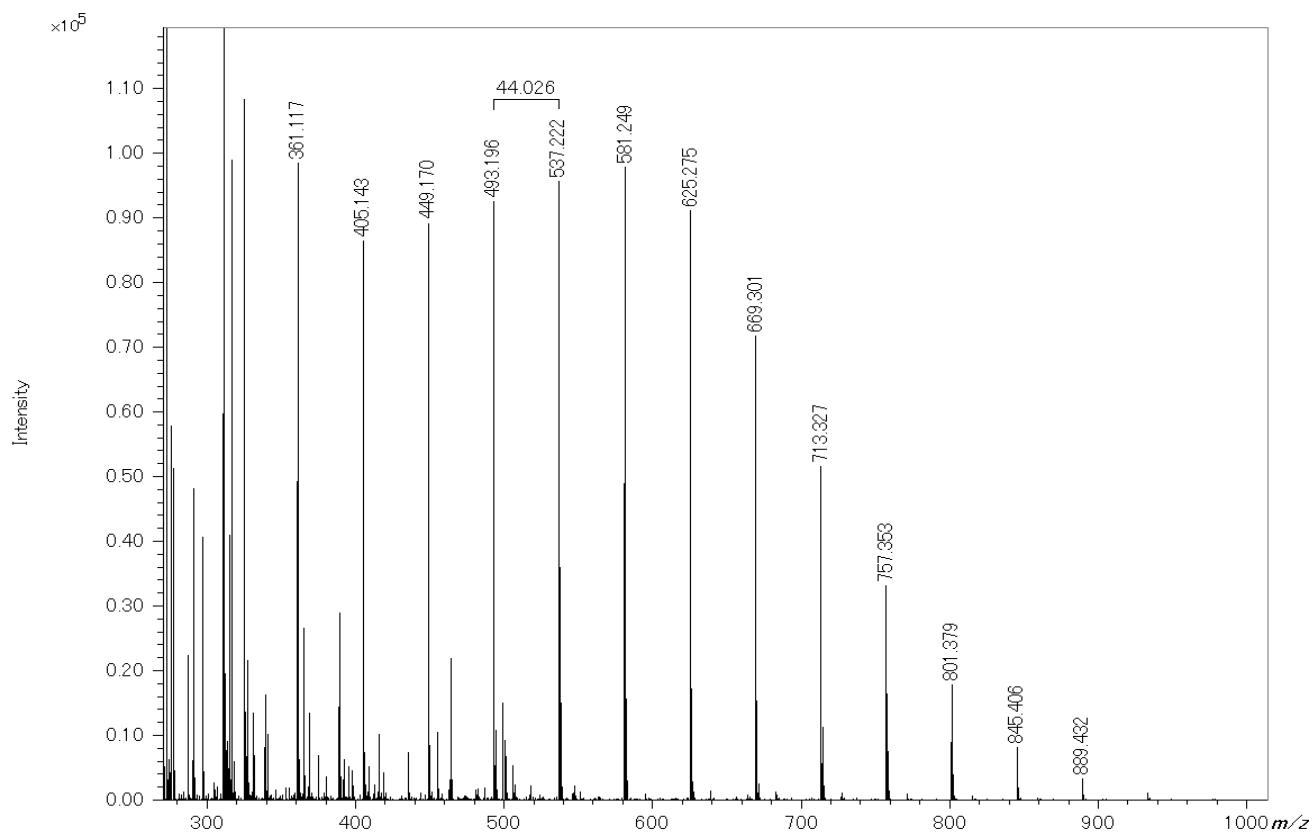
(High molecular weight HALS (hindered amine light stabilizer))



Structure	$C_{10}H_{20}N[C_{43}H_{72}N_2O_{12}]_n-C_{38}H_{66}O_8N_3$
Ion Species	$[M+H]^+$ , $[M+Na]^+$
Molecular weight Information	$M_n$ 1157, $M_w$ 1424, D 1.231
Matrix	DHB
Cationization agent	None
Mode	SpiralTOF:Positive ion

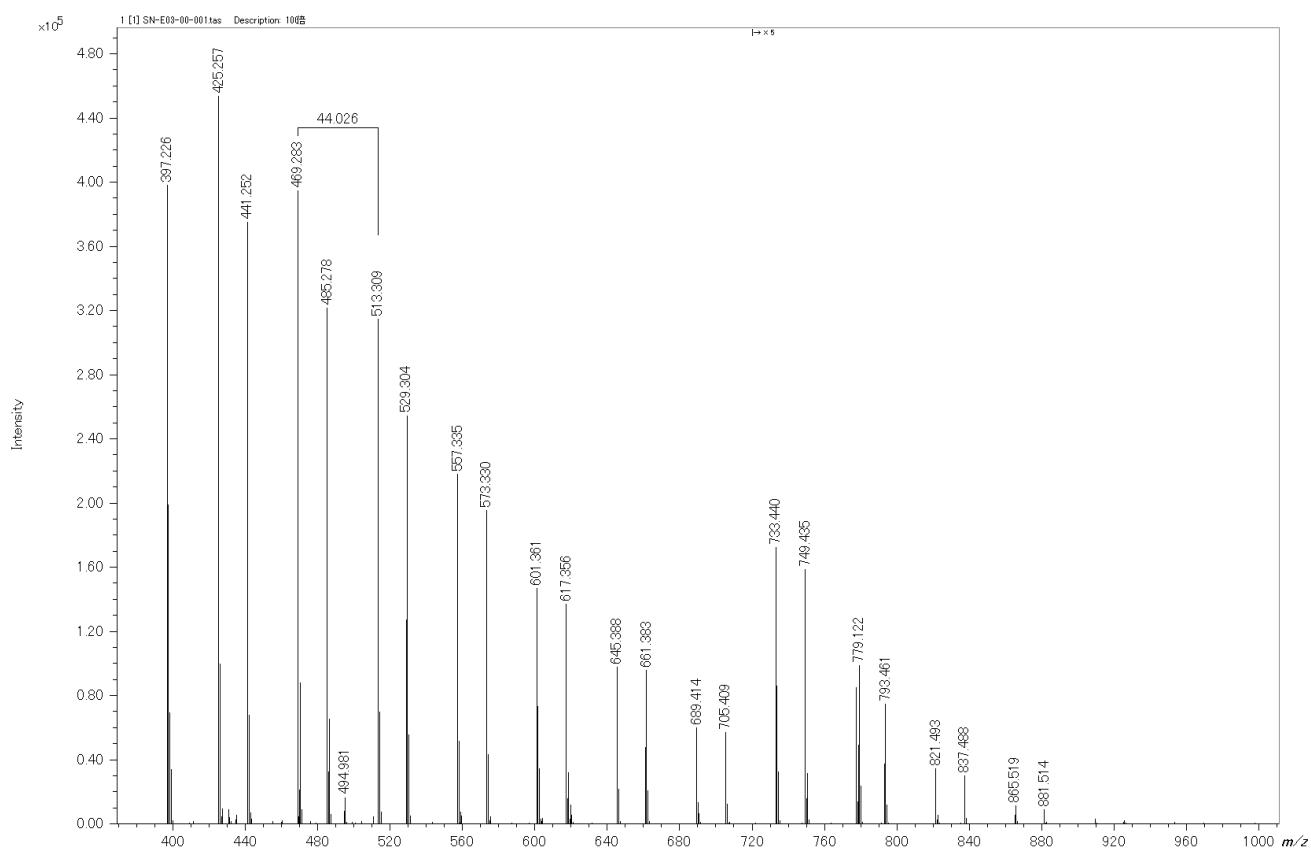
\* ADK STAB is either a registered trademark or a trademark of ADEKA Corporation in Japan and/or other countries.

## Polyethylene glycol sulfate

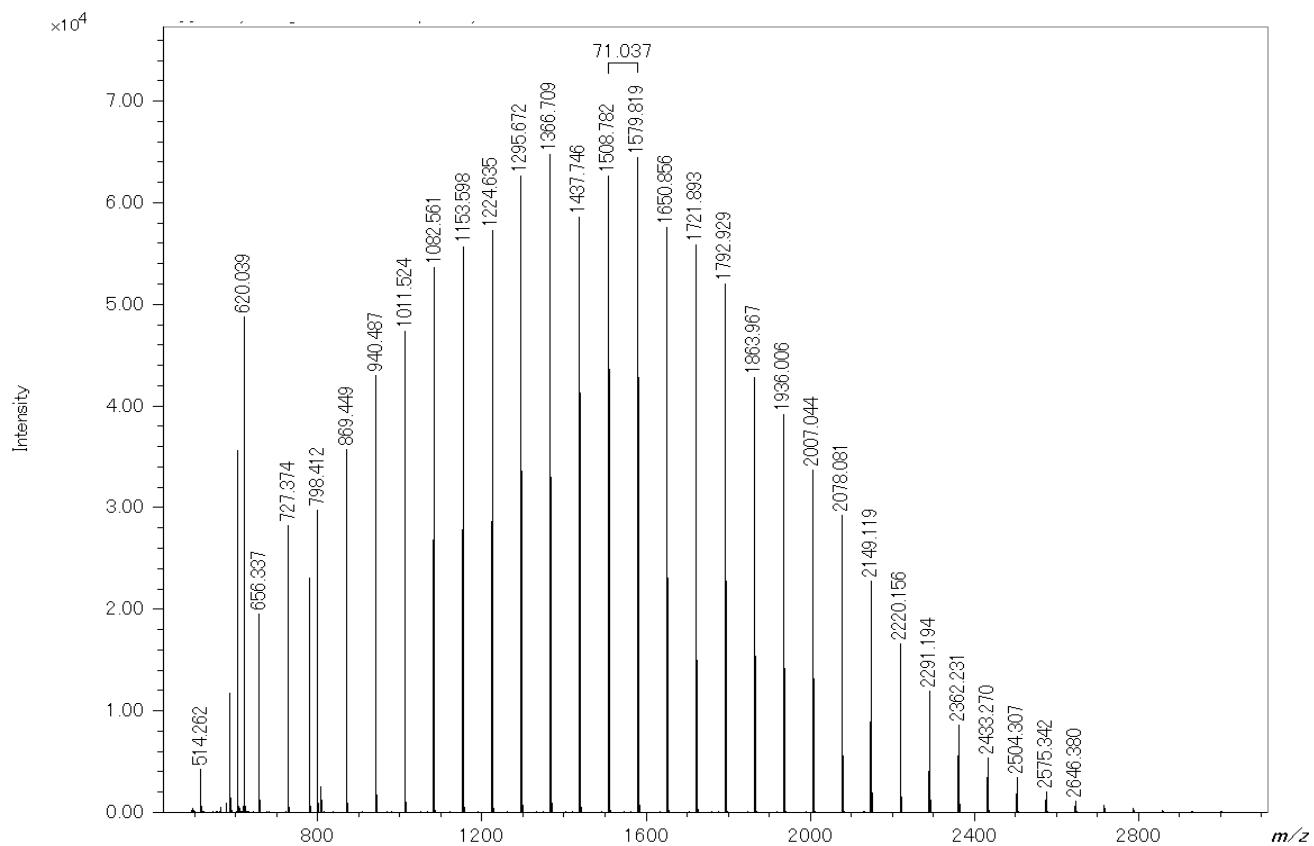


Structure	$\text{HO}[(\text{CH}_2)_2\text{O}]_n\text{SO}_3\text{H}$ ; $\text{HO}(\text{C}_2\text{H}_4\text{O})_n\text{SO}_3\text{H}$
Ion Species	$[\text{M}-\text{H}]^-$
Molecular weight Information	$M_n$ 535; $M_w$ 573; D 1.070
Matrix	DCTB
Cationization agent	None
Mode	SpiralTOF:Negative ion

## Polyoxyethylene alkyl ether sulfuric acid

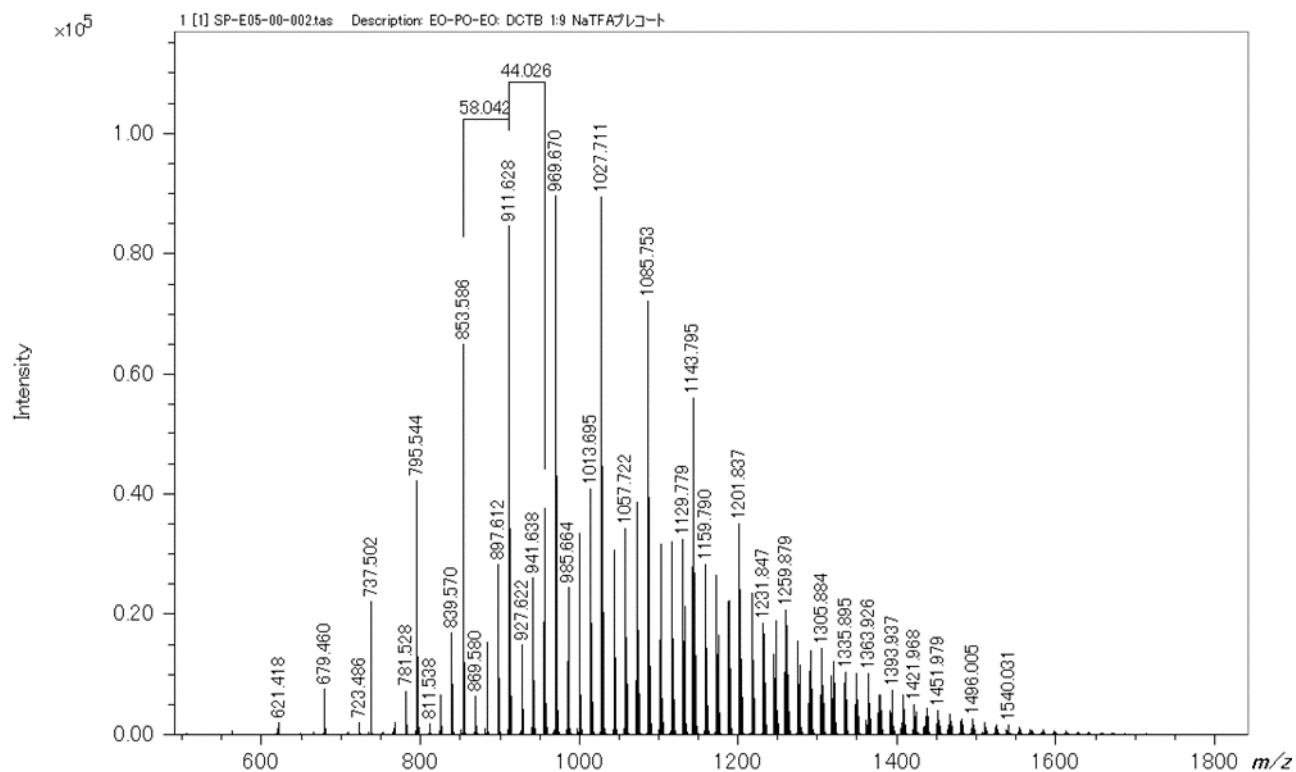


Structure	$\text{NaSO}_3[(\text{CH}_2)_2\text{O}]_n\text{O}(\text{CH}_2)_{11}\text{CH}_3$ ; $\text{NaSO}_3(\text{C}_2\text{H}_4\text{O})_n\text{OC}_{12}\text{H}_{25}$ $\text{NaSO}_3[(\text{CH}_2)_2\text{O}]_n\text{O}(\text{CH}_2)_{13}\text{CH}_3$ ; $\text{NaSO}_3(\text{C}_2\text{H}_4\text{O})_n\text{OC}_{14}\text{H}_{29}$
Ion Species	$[\text{M}-\text{Na}]^-$
Molecular weight Information	$M_n$ 511, $M_w$ 527, D 1.032
Matrix	CHCA
Cationization agent	None
Mode	SpiralTOF:Negative ion

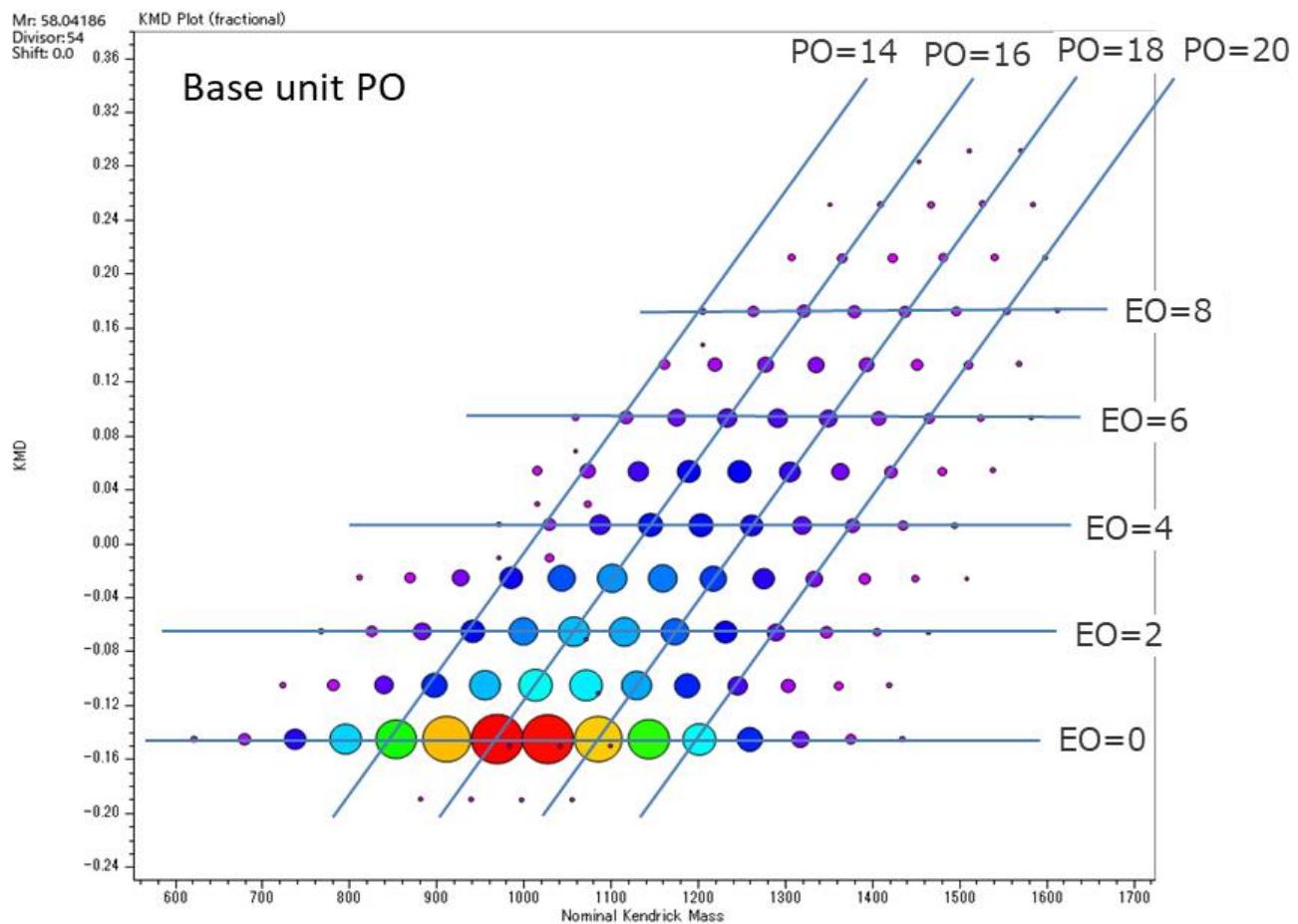
**Poly(alanine)**

Structure	$\text{H}[\text{NHCH}(\text{CH}_3)\text{CO}]_n\text{OH}$ ; $\text{H}(\text{C}_3\text{H}_5\text{NO})_n\text{OH}$
Ion Species	$[\text{M}-\text{H}]^-$
Molecular weight Information	$M_n$ 1,652; $M_w$ 1,765; D 1.069
Matrix	DCTB
Cationization agent	None
Mode	SpiralTOF:Negative ion

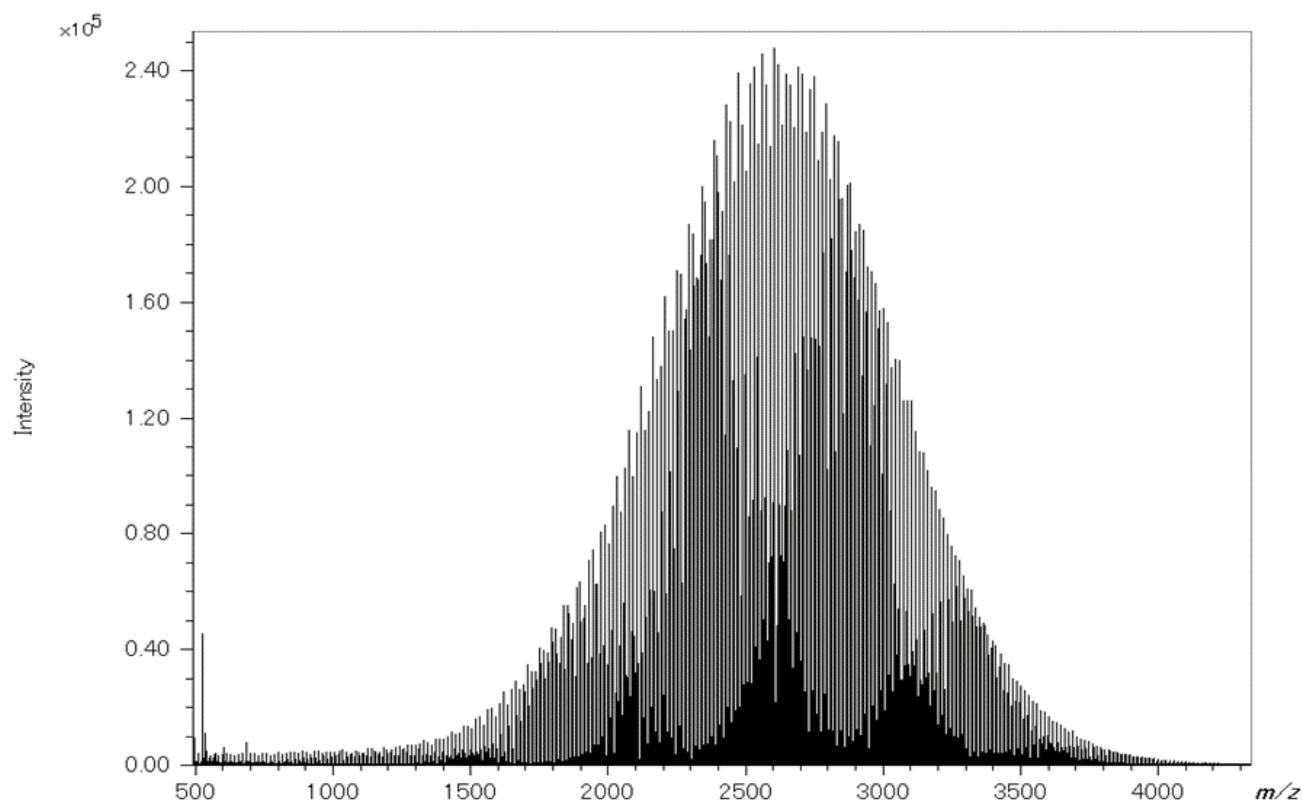
## Poly(ethylene glycol)-*b*-block-poly(propylene glycol)-*b*-block-poly(ethylene glycol)



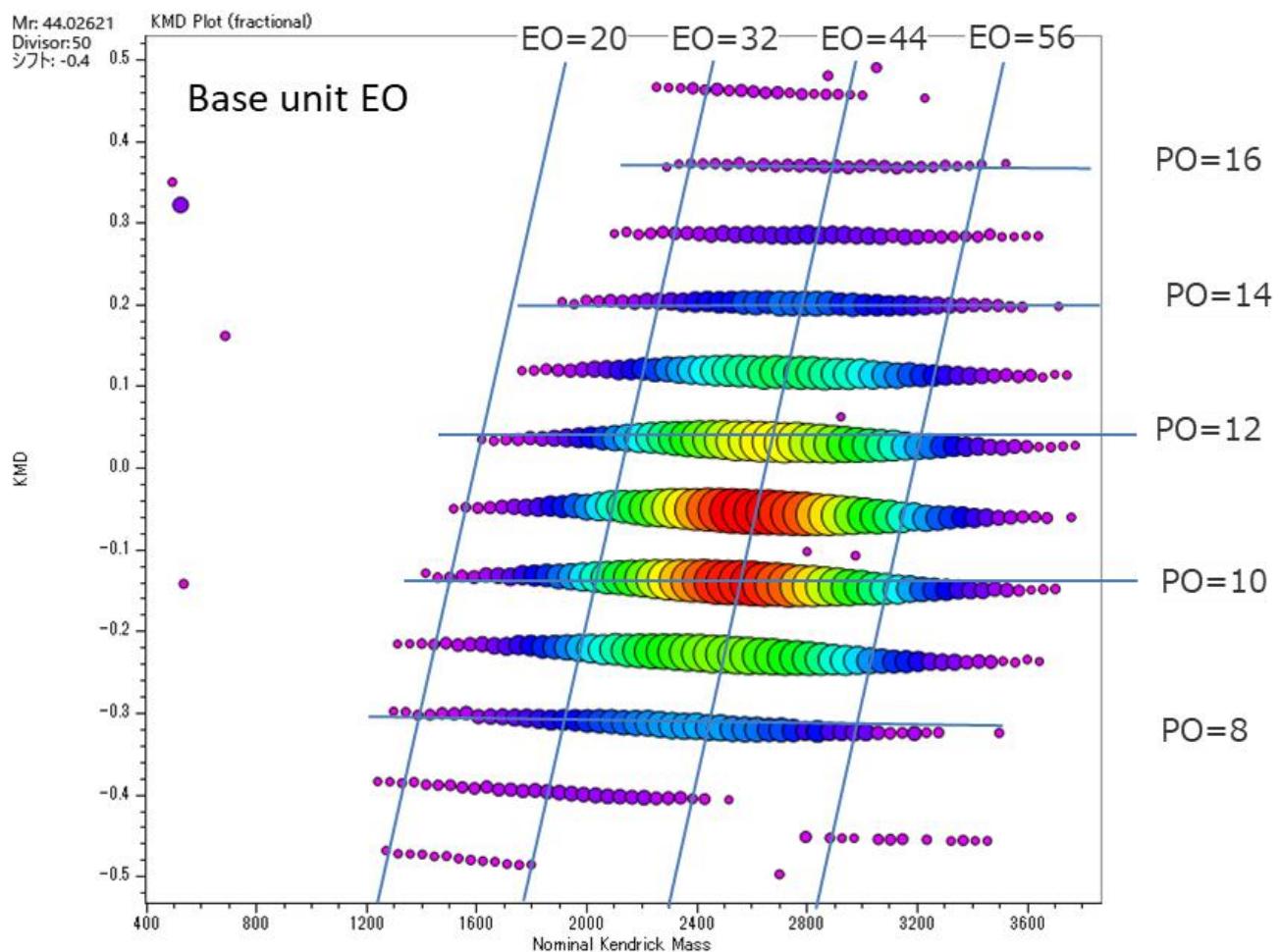
Structure	$O[O(CH_2)_2]_x-b-[OCH(CH_3)CH_2]_y-b-[O(CH_2)_2]_zOH$ ; $H(C_2H_4O)_{x+z}(C_3H_6O)_yOH$ ; PEG, 10 wt. %
Ion Species	$[M+Na]^+$
Molecular weight Information	$M_n$ 1,101; $M_w$ 1,128; D 1.025
Matrix	DCTB
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion



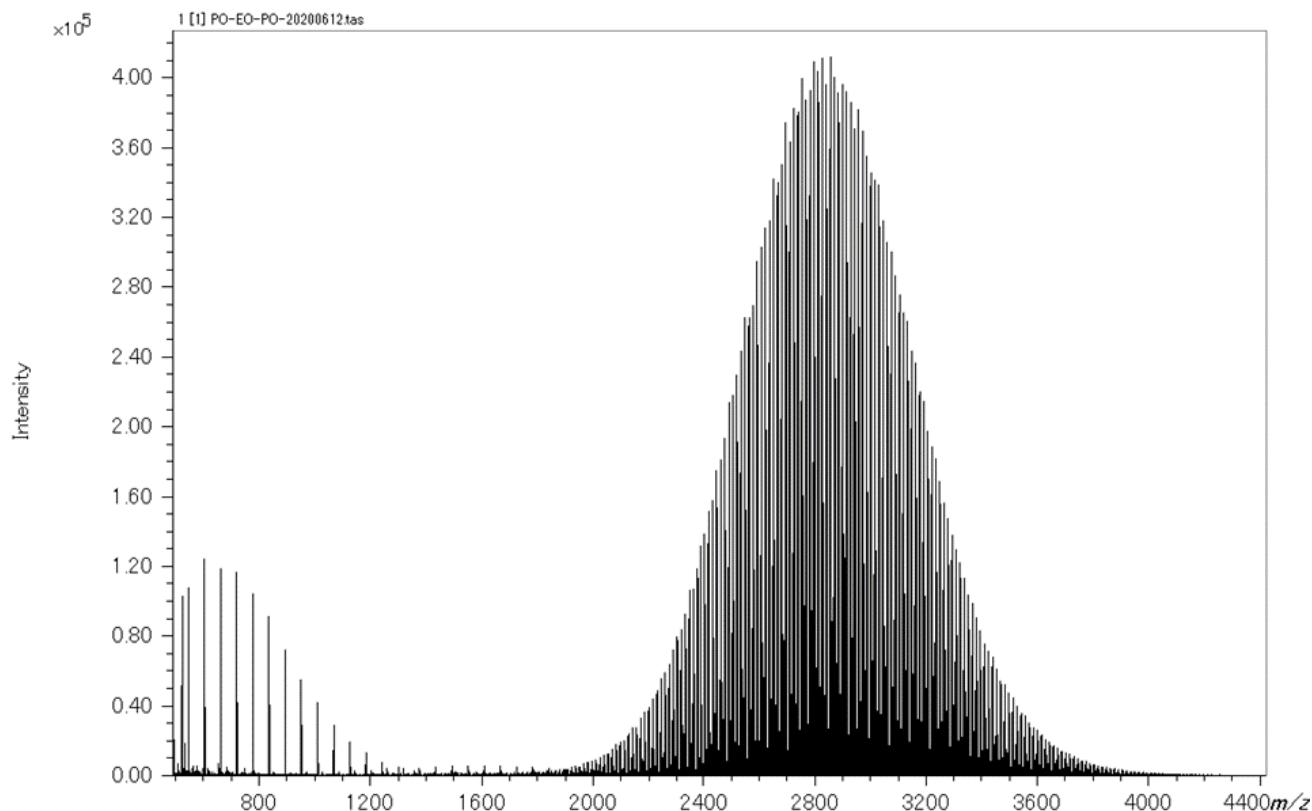
## Poly(ethylene glycol-*ran*-propylene glycol)



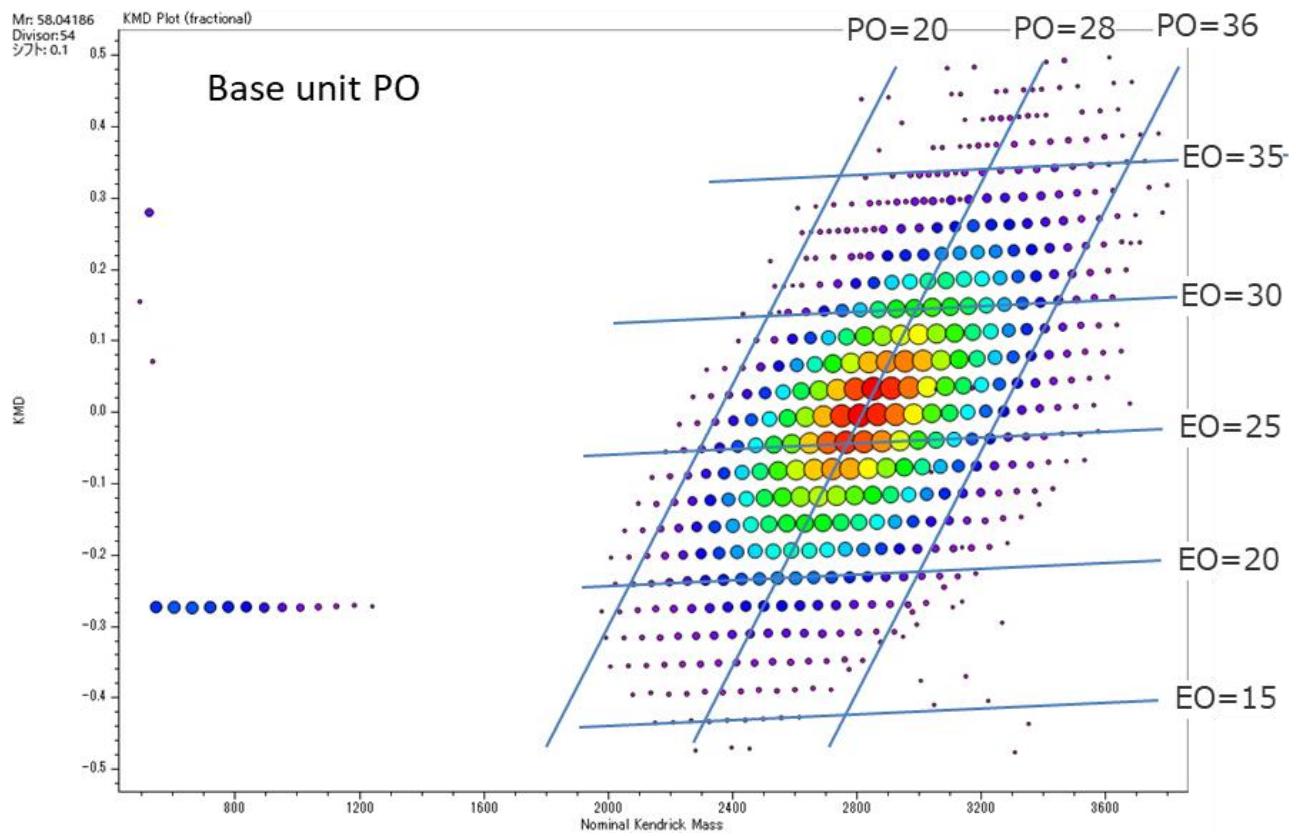
Structure	$H[O(CH_2)_2]_x\text{-}ran\text{-}[OCH(CH_3)CH_2]_yOH$ ; $O(C_2H_4O)_x(C_3H_6O)_yOH$
Ion Species	$[M+Na]^+$
Molecular weight Information	$M_n$ 2,554; $M_w$ 2,674; D 1.047
Matrix	DCTB
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion



## Poly(propylene glycol)-*b*-block-poly(ethylene glycol)-*b*-block-poly(propylene glycol)

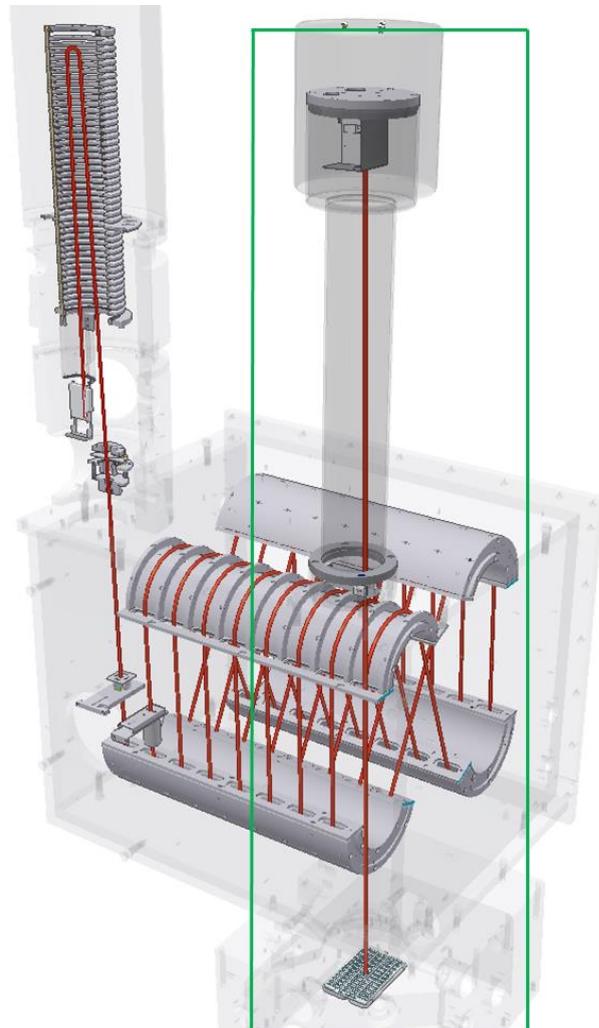


Structure	O[OCH(CH <sub>3</sub> )CH <sub>2</sub> ] <sub>x</sub> - <i>b</i> -[O(CH <sub>2</sub> ) <sub>2</sub> ] <sub>y</sub> - <i>b</i> -[OCH(CH <sub>3</sub> )CH <sub>2</sub> ] <sub>z</sub> OH; H(C <sub>3</sub> H <sub>6</sub> O) <sub>x+z</sub> (C <sub>2</sub> H <sub>4</sub> O) <sub>y</sub> OH; PEG, 40 wt. %
Ion Species	[M+Na] <sup>+</sup>
Molecular weight Information	$M_n$ 2,554; $M_w$ 2,674; D 1.047
Matrix	DCTB
Cationization agent	NaTFA
Mode	SpiralTOF:Positive ion





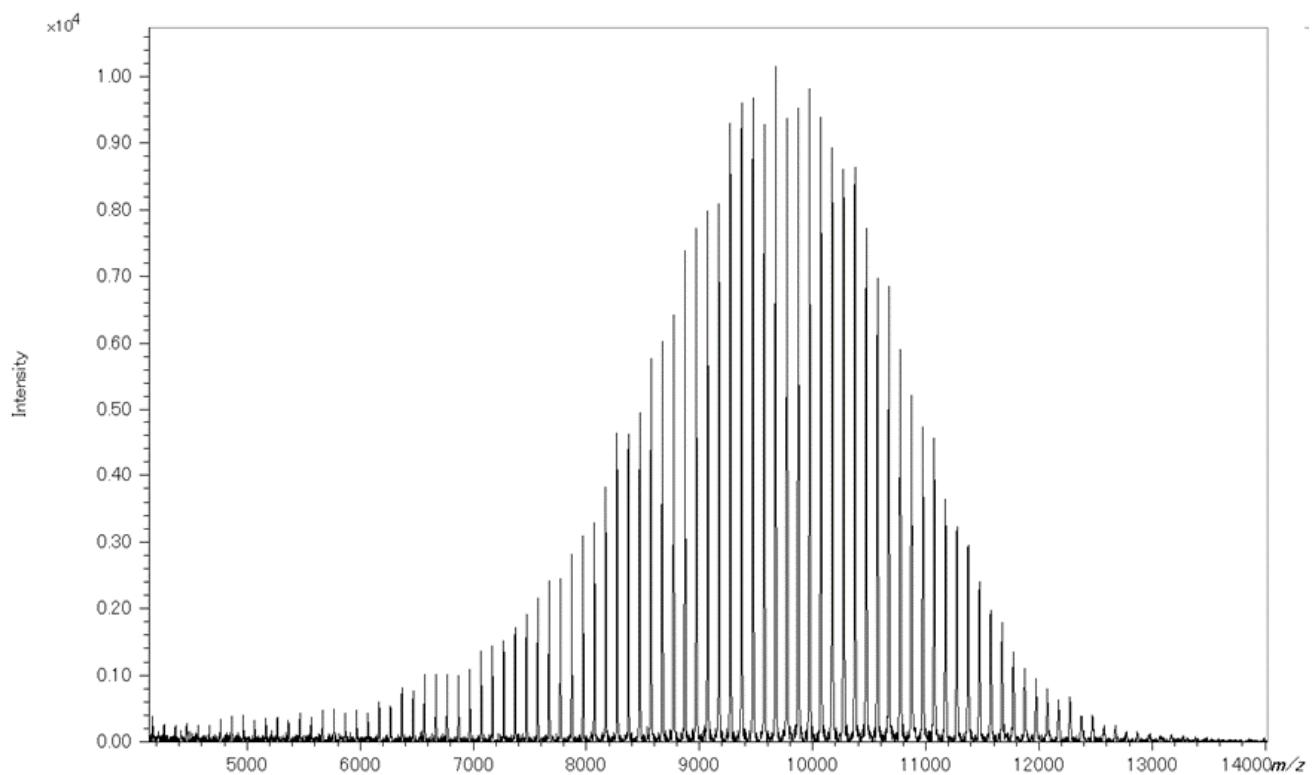
## Linear Mode



- Ions fly straight to the detector after being accelerated.
- High sensitivity can be achieved for high mass samples – suitable for polymers with the molecular weight of tens of thousands.
- The mass spectrum reflects ion population right after being accelerated – suitable for the determination of molecular weight distribution and for polymers that are easily fragmented.

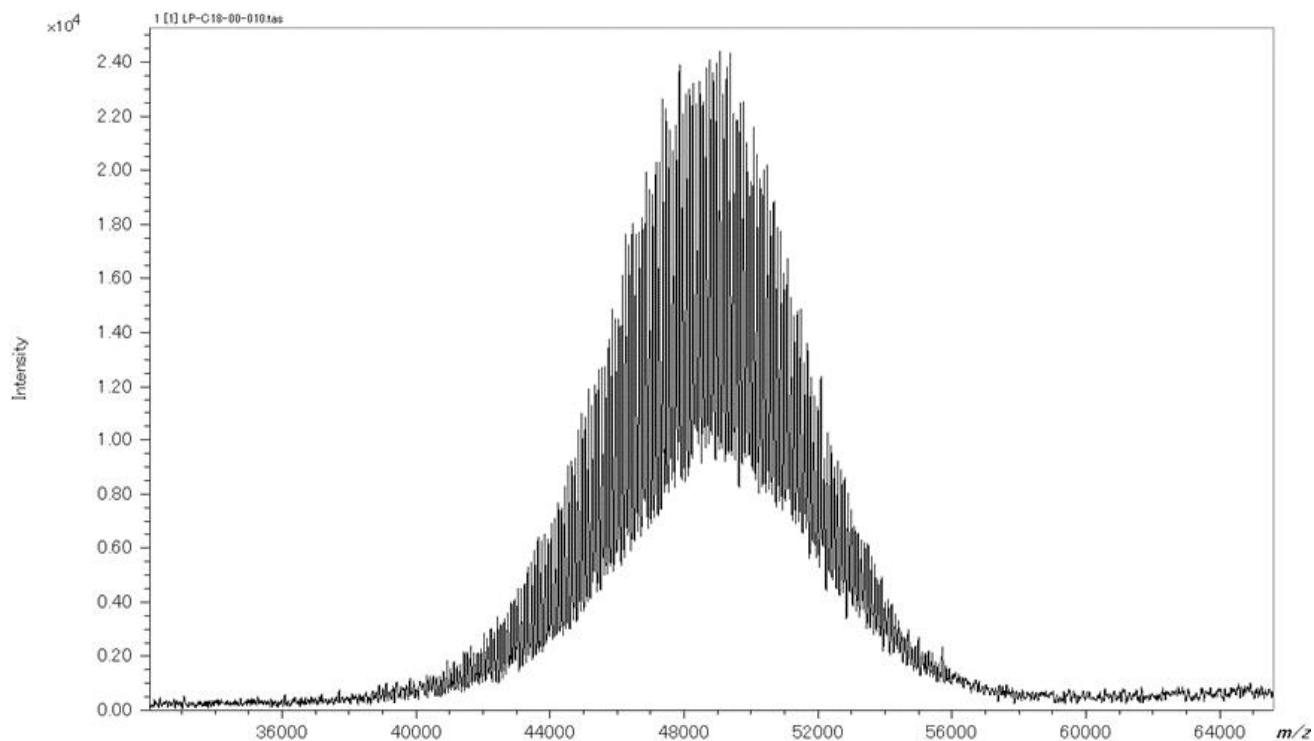


## PMMA (poly(methyl methacrylate)) MW 10 kDa



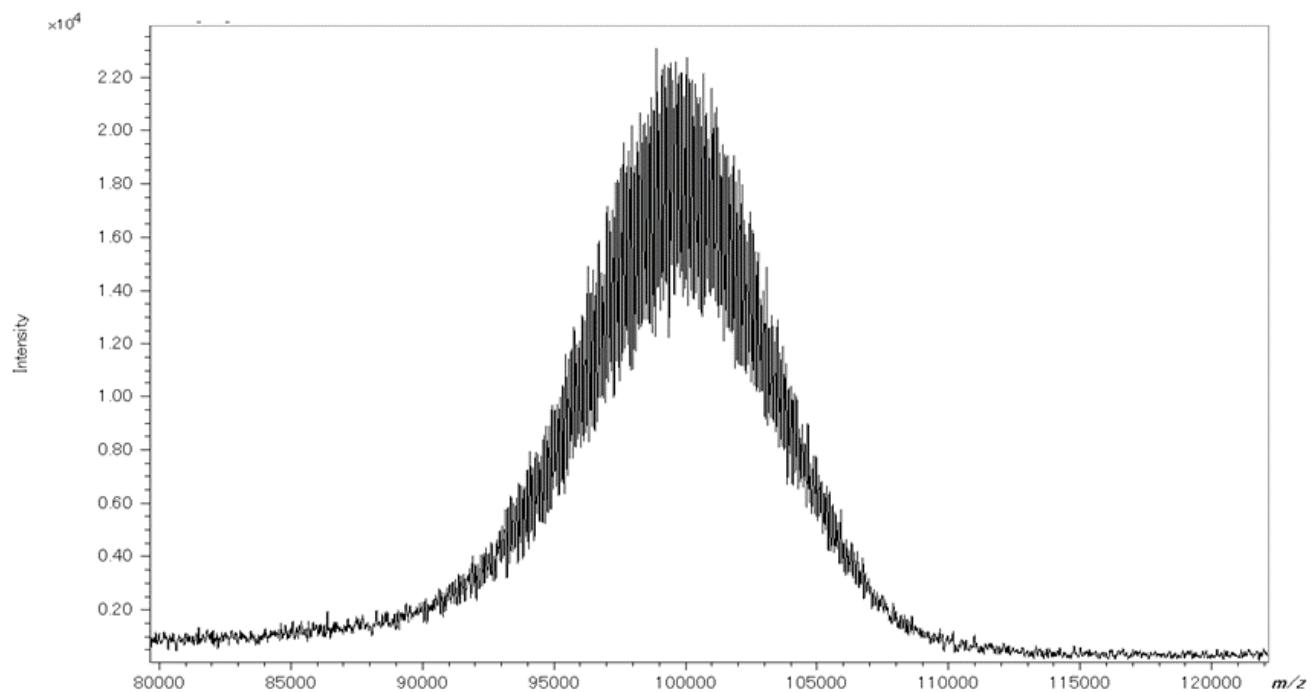
Structure	$H[CH_2C(CH_3)(CO_2CH_3)]_nH$ ; $H(C_5H_8O)_nH$
Ion Species	$[M+Na]^+$
Molecular weight Information	$M_n$ 9,510; $M_w$ 9,779; D 1.028
Matrix	DCTB
Cationization agent	NaTFA
Mode	Linear :Positive ion

## PMMA (poly(methyl methacrylate)) MW 50 kDa



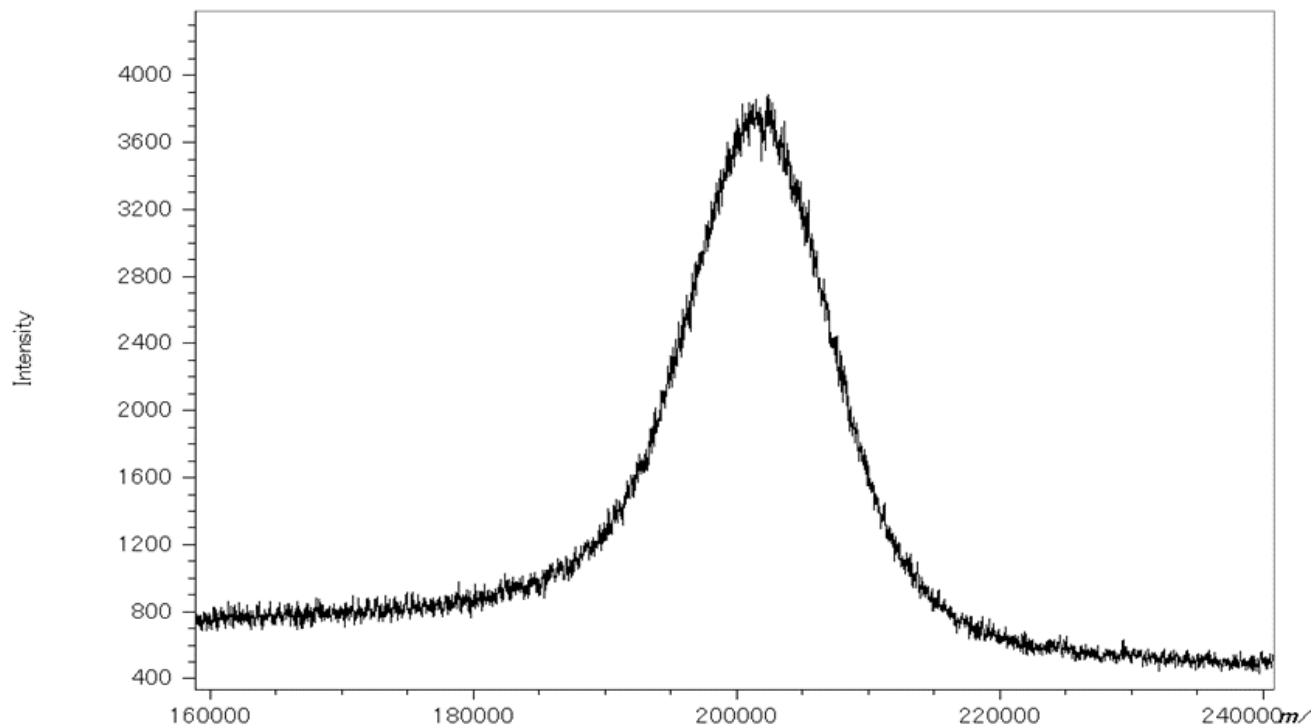
Structure	$H[CH_2C(CH_3)(CO_2CH_3)]_nH$ ; $H(C_5H_8O)_nH$
Ion Species	$[M+Na]^+$
Molecular weight Information	$M_n$ 48,550; $M_w$ 48,821; D 1.006
Matrix	DCTB
Cationization agent	NaTFA
Mode	Linear :Positive ion

## PS (poly(styrene)) MW 100 kDa

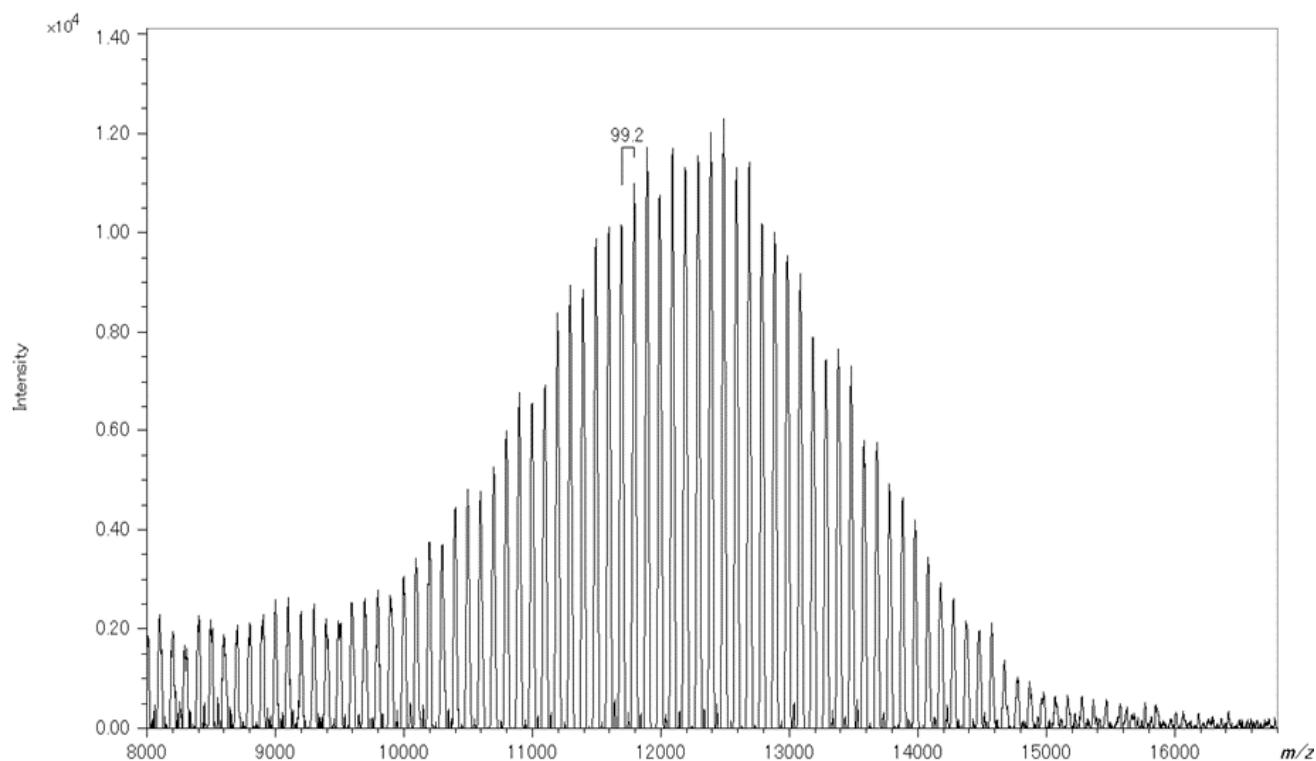


Structure	-(C <sub>8</sub> H <sub>8</sub> )-
Ion Species	[M+Ag] <sup>+</sup>
Molecular weight Information	M <sub>n</sub> 97,059; M <sub>w</sub> 97,861; D 1.008
Matrix	DCTB
Cationization agent	AgTFA
Mode	Linear :Positive ion

**PS (poly(styrene)) MW 200 kDa**



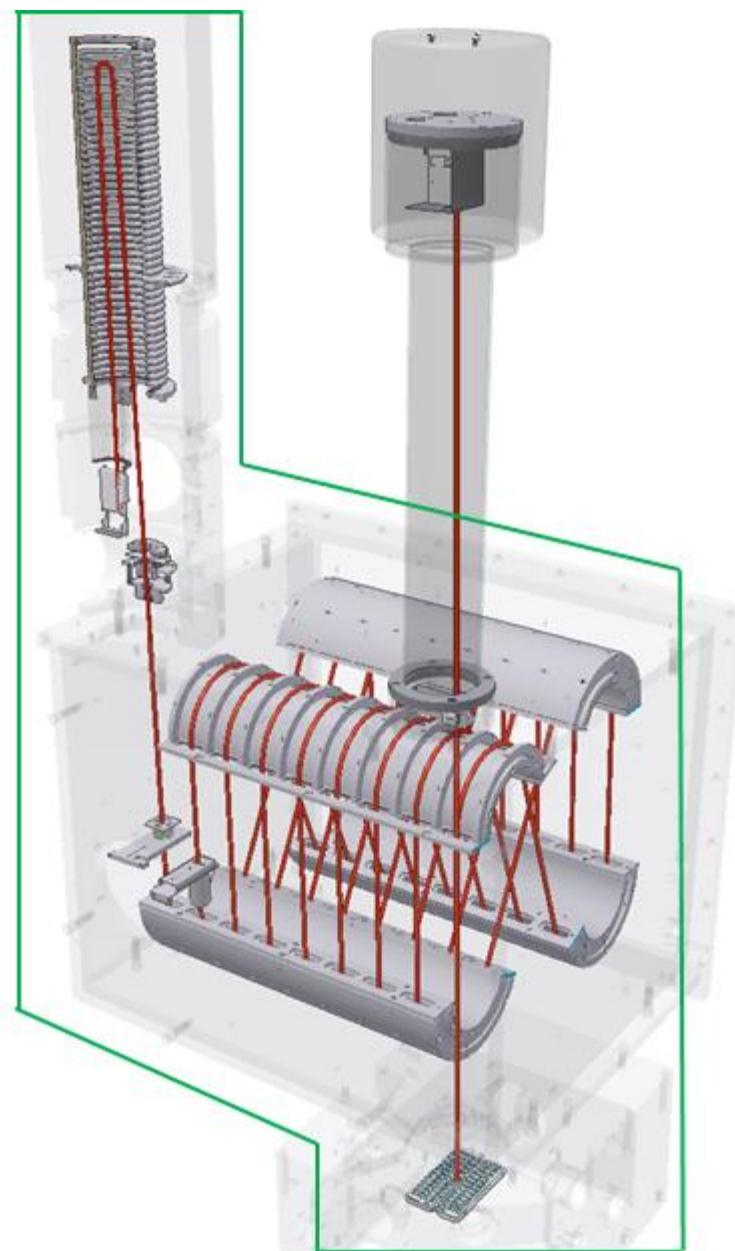
Structure	-[CH <sub>2</sub> CH(C <sub>6</sub> H <sub>5</sub> )] <sub>n</sub> -; -(C <sub>8</sub> H <sub>8</sub> ) <sub>n</sub> -
Ion Species	[M+Ag] <sup>+</sup>
Molecular weight Information	$M_n$ 197,860; $M_w$ 199,401; D 1.008
Matrix	DCTB
Cationization agent	AgTFA
Mode	Linear :Positive ion

**POZ (poly(2-ethyl-2-oxazoline)) MW 10 kDa**

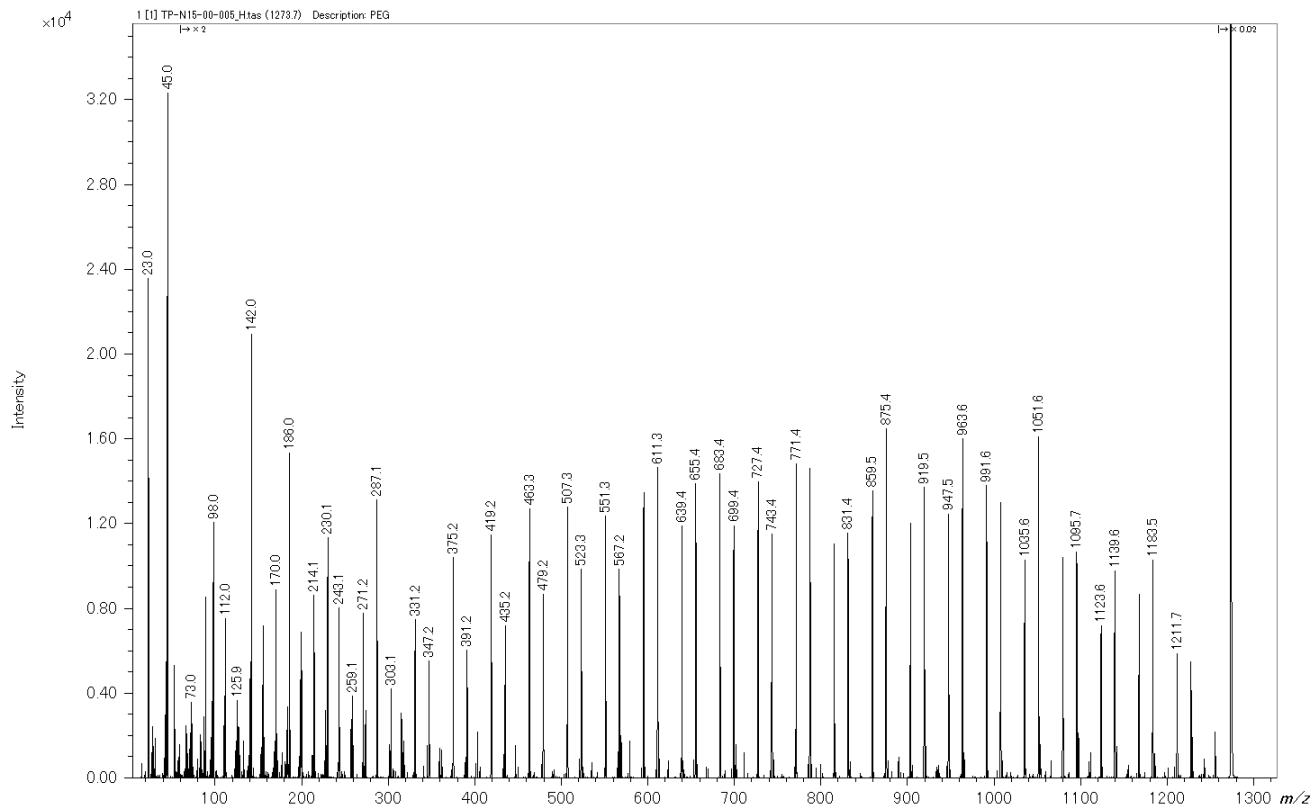
Structure	$\text{CH}_3[\text{N}(\text{COC}_2\text{H}_5)\text{CH}_2\text{CH}_2]_n\text{OH}$ ; $\text{CH}_3(\text{C}_5\text{H}_9\text{NO})\text{OH}$
Ion Species	$[\text{M}+\text{Na}]^+$
Molecular weight Information	$M_n$ 11,307; $M_w$ 11,,596; D 1.026
Matrix	DCTB
Cationization agent	NaTFA
Mode	Linear :Positive ion



## TOF-TOF Mode

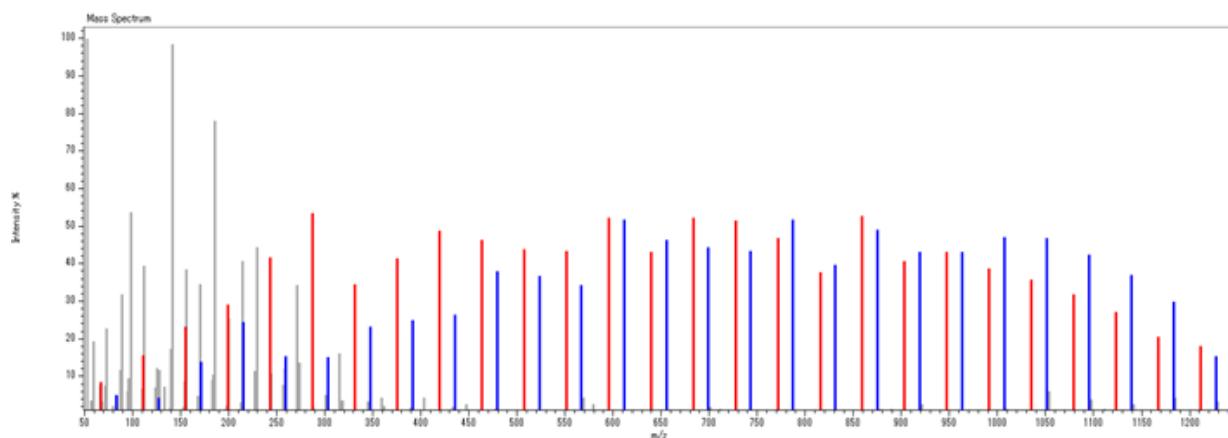


- Information-rich product ion spectra with high-energy collision induced dissociation (HE-CID).
- High precursor ion selectivity due to long flight distance of 15 m from the ion source to the ion gate.
- The Offset Parabolic Reflectron used in the 2nd TOFMS can detect product ions over a wide mass range from alkali metal adduct ions to the precursor ions.

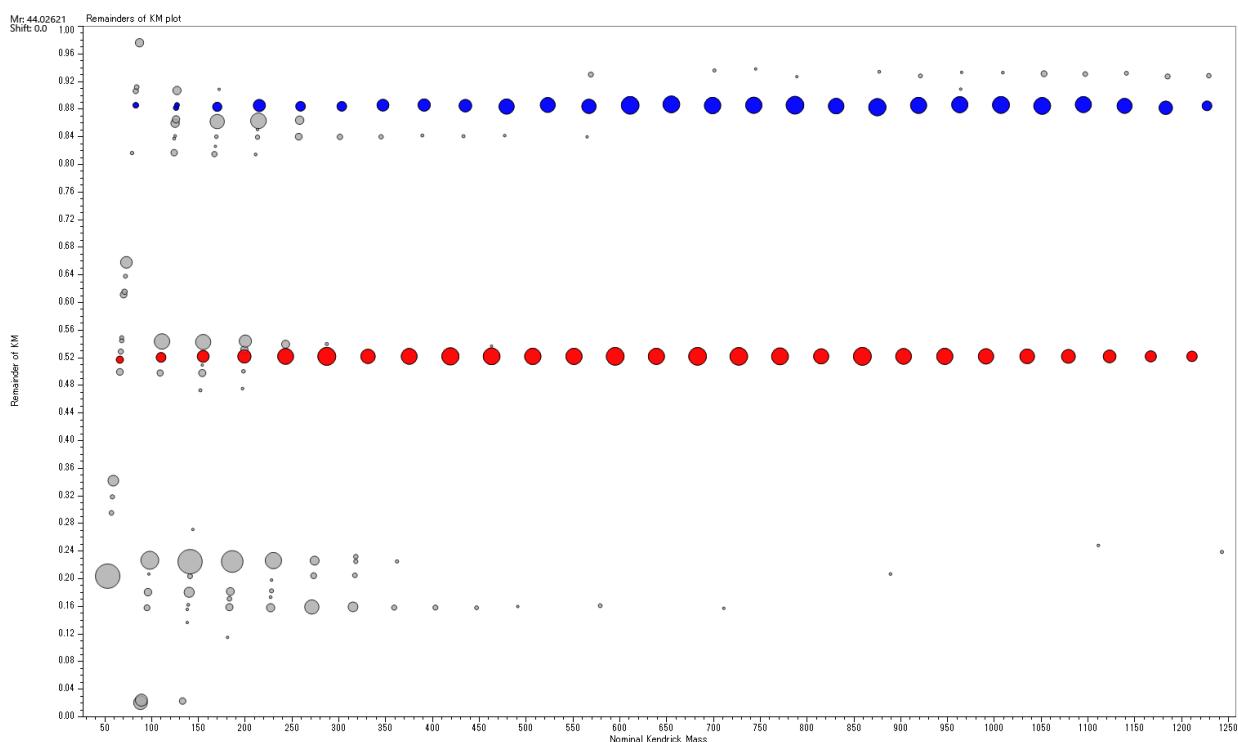
**PEO (poly(ethylene oxide))**

Structure	$\text{HO}(\text{CH}_2\text{CH}_2\text{O})_n\text{H}$
Ion Species	$[\text{M}+\text{Na}]^+$
Precursor ion	$m/z$ 1,273.73380 ( $n = 28$ )
Matrix	DCTB
Cationization agent	NaTFA
Mode	TOF-TOF :Positive ion

## Peak list



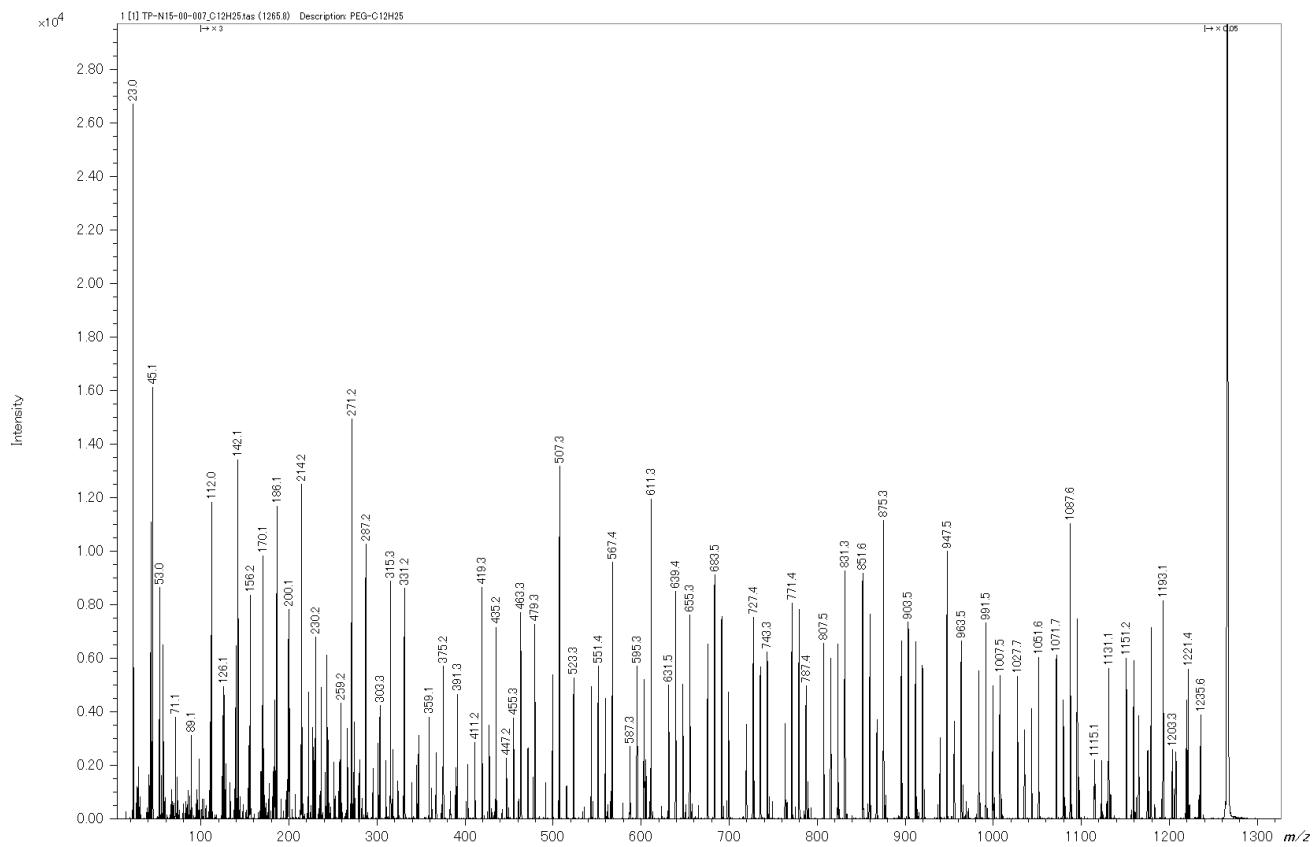
## RKM Plot



## Estimated Structure

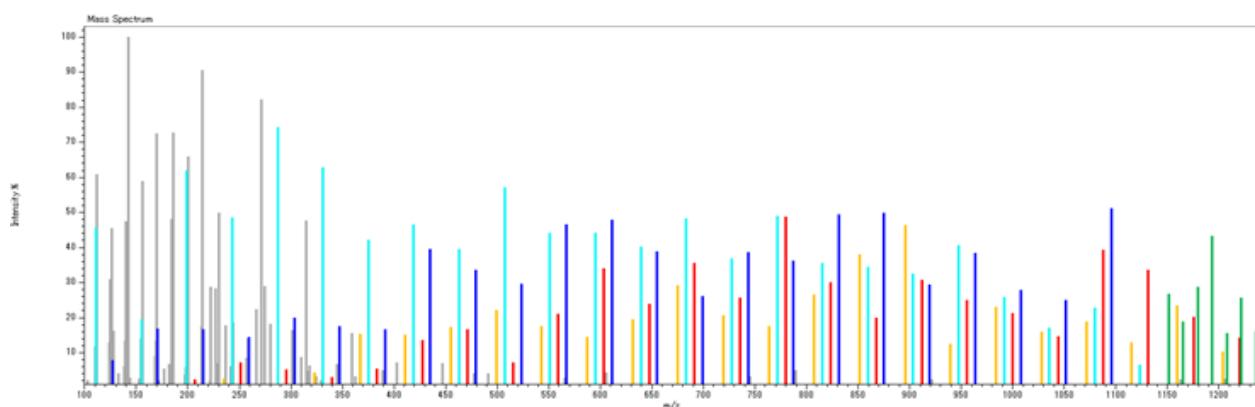


## Polyethylene glycol monododecyl ether

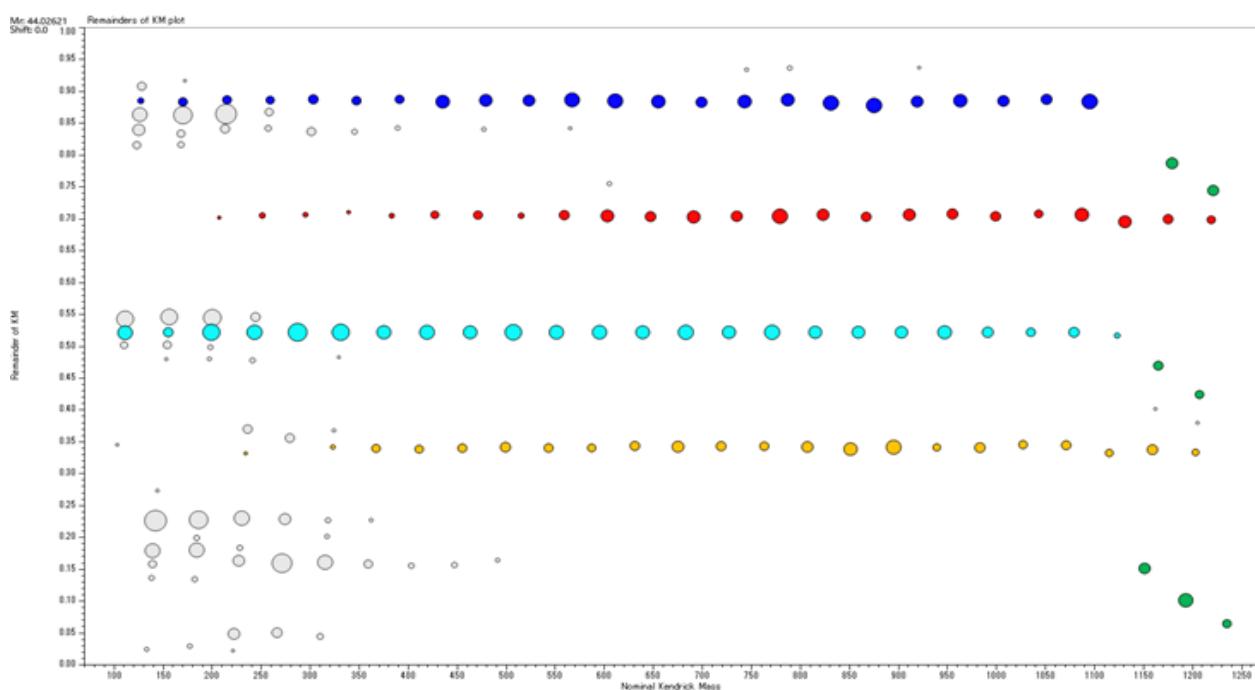


Structure	$\text{HO}[(\text{CH}_2)_2\text{O}]_n(\text{CH}_2)_{11}\text{CH}_3, \text{HO}(\text{C}_2\text{H}_4\text{O})_n\text{C}_{12}\text{H}_{25}$
Ion Species	$[\text{M}+\text{Na}]^+$
Precursor ion	$m/z$ 1,265.817 ( $n = 24$ )
Matrix	DCTB
Cationization agent	NaTFA
Mode	TOF-TOF:Positive ion

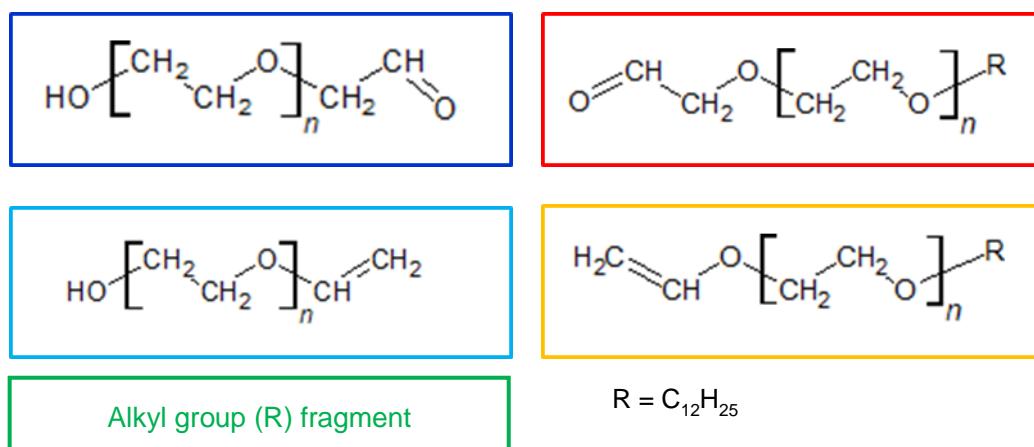
## Peak list



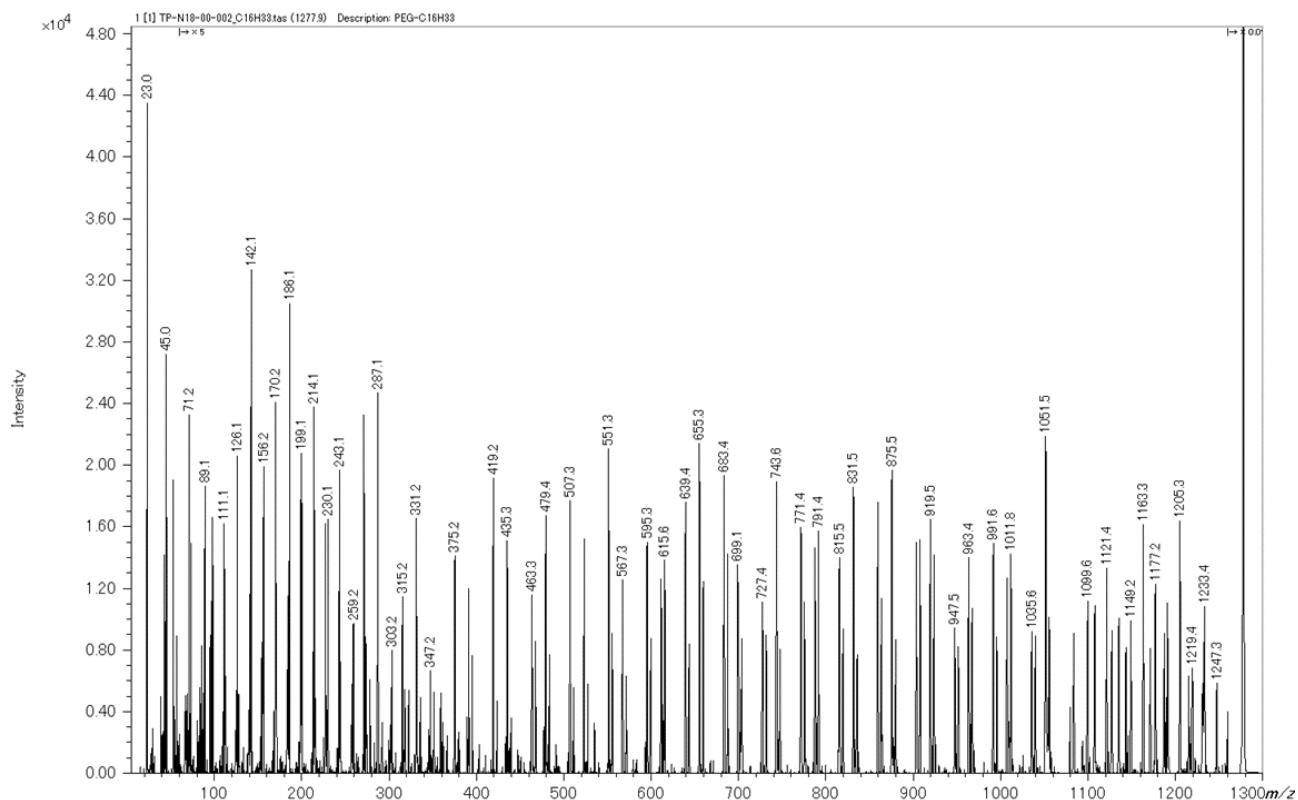
## RKM Plot



## Estimated Structure

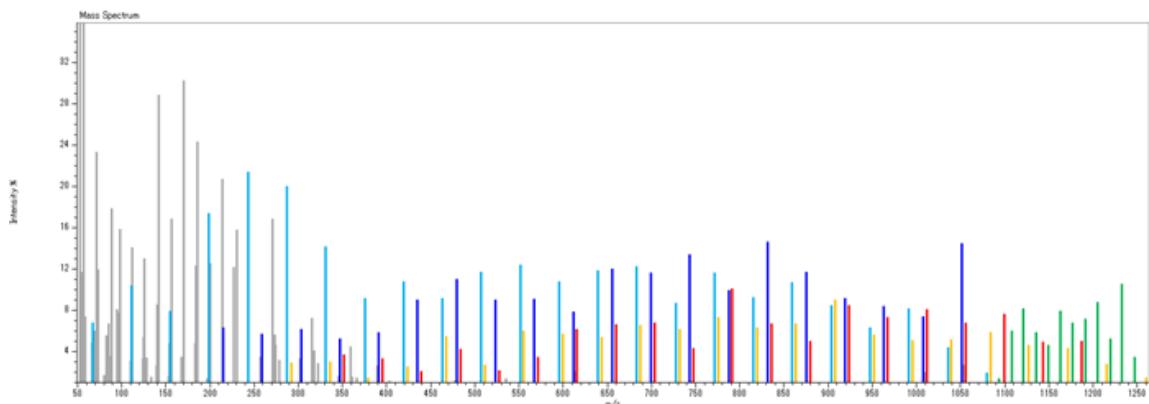


## Polyethylene glycol monocetyl ether

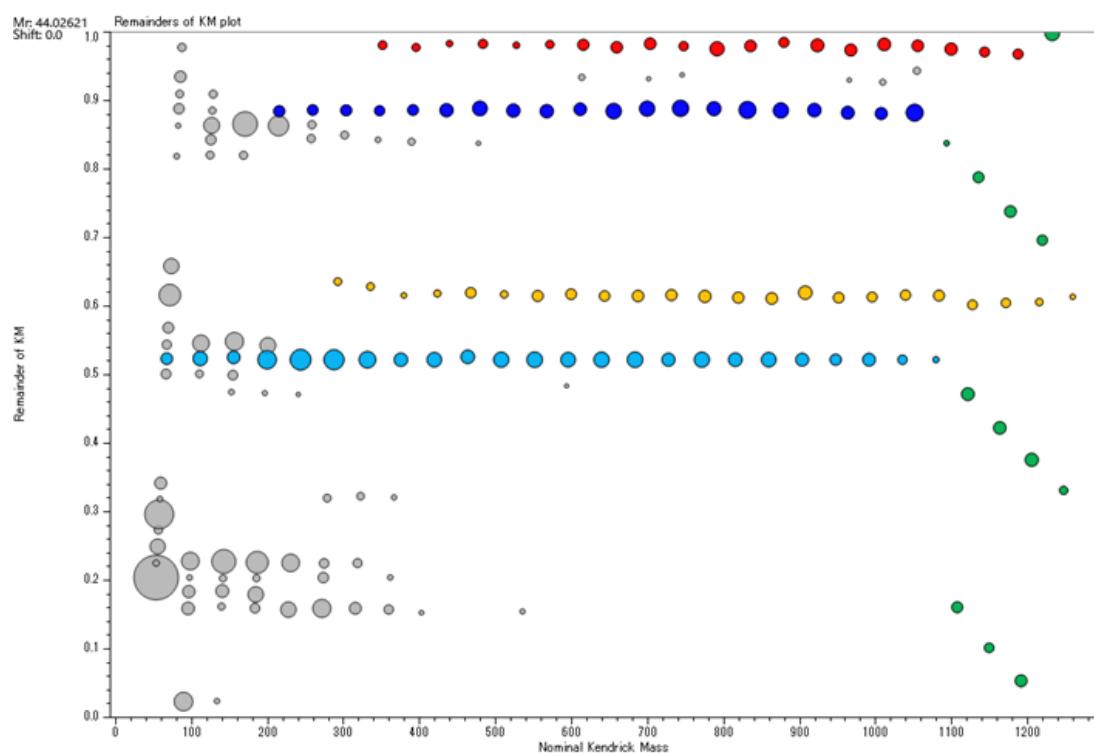


Structure	$\text{HO}[(\text{CH}_2)_2\text{O}]_n(\text{CH}_2)_{15}\text{CH}_3$ ; $\text{HO}(\text{C}_2\text{H}_4\text{O})_n\text{C}_{16}\text{H}_{33}$
Ion Species	$[\text{M}+\text{Na}]^+$
Precursor ion	$m/z$ 1,277.853 ( $n = 23$ )
Matrix	DCTB
Cationization agent	NaTFA
Mode	TOF-TOF:Positive ion

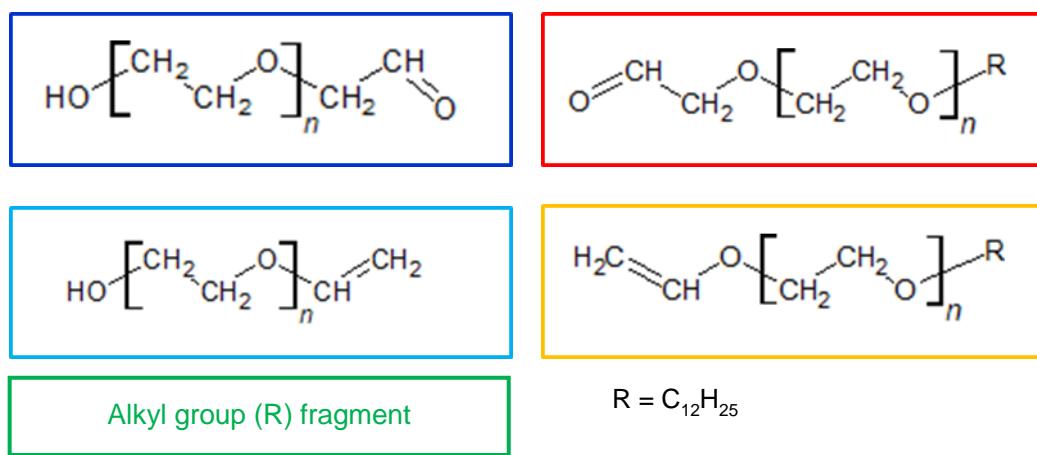
## Peak list

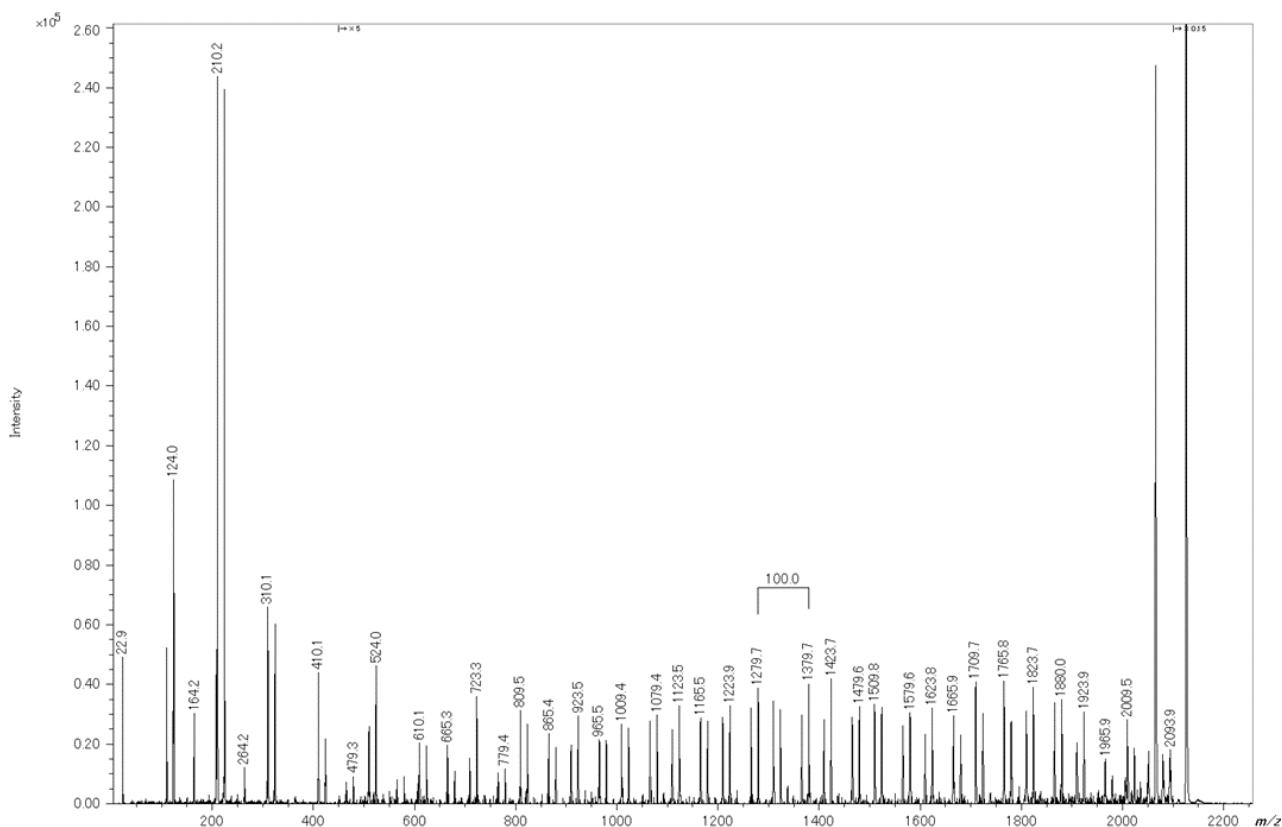


## RKM Plot



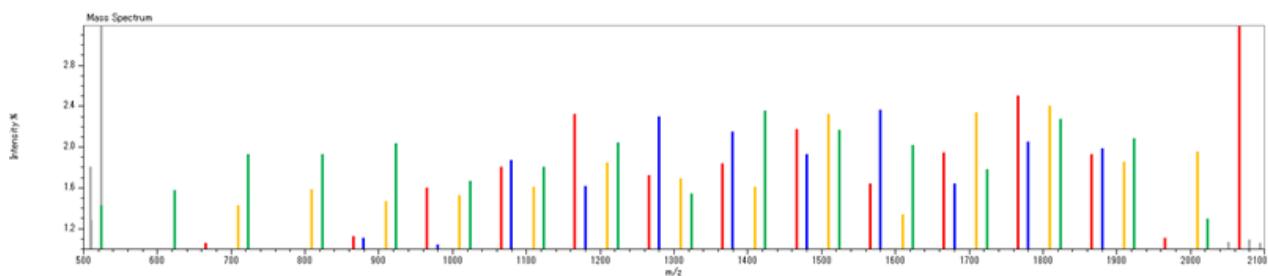
## Estimated Structure



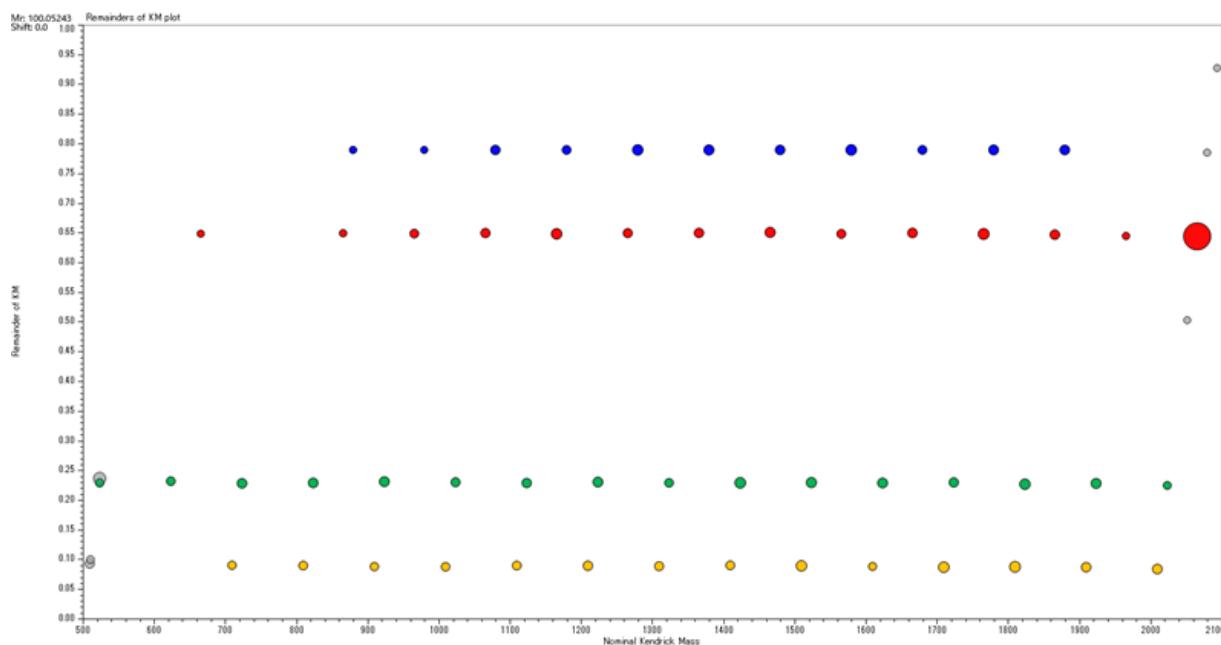
**PMMA (poly(methyl methacrylate))**

Structure	$\text{HO}(\text{CH}_2\text{CH}_2\text{O})_n\text{H}$ ; $\text{HO}(\text{C}_2\text{H}_4\text{O})_n\text{H}$
Ion Species	$[\text{M}+\text{Na}]^+$
Precursor ion	$m/z$ 2,125 ( $n = 25$ )
Matrix	DCTB
Cationization agent	NaTFA
Mode	TOF-TOF:Positive ion

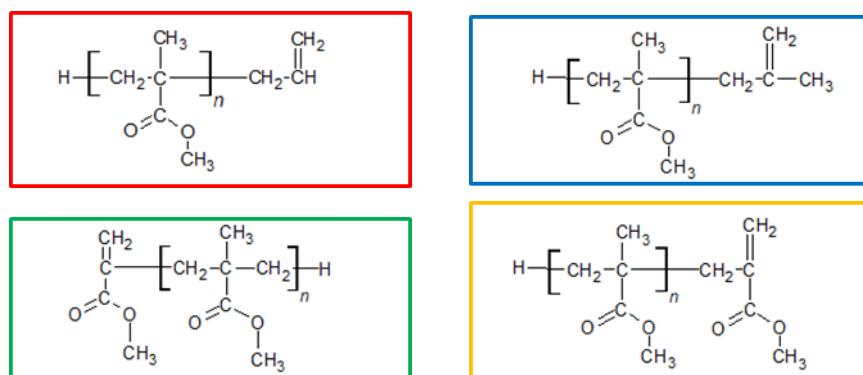
## Peak list

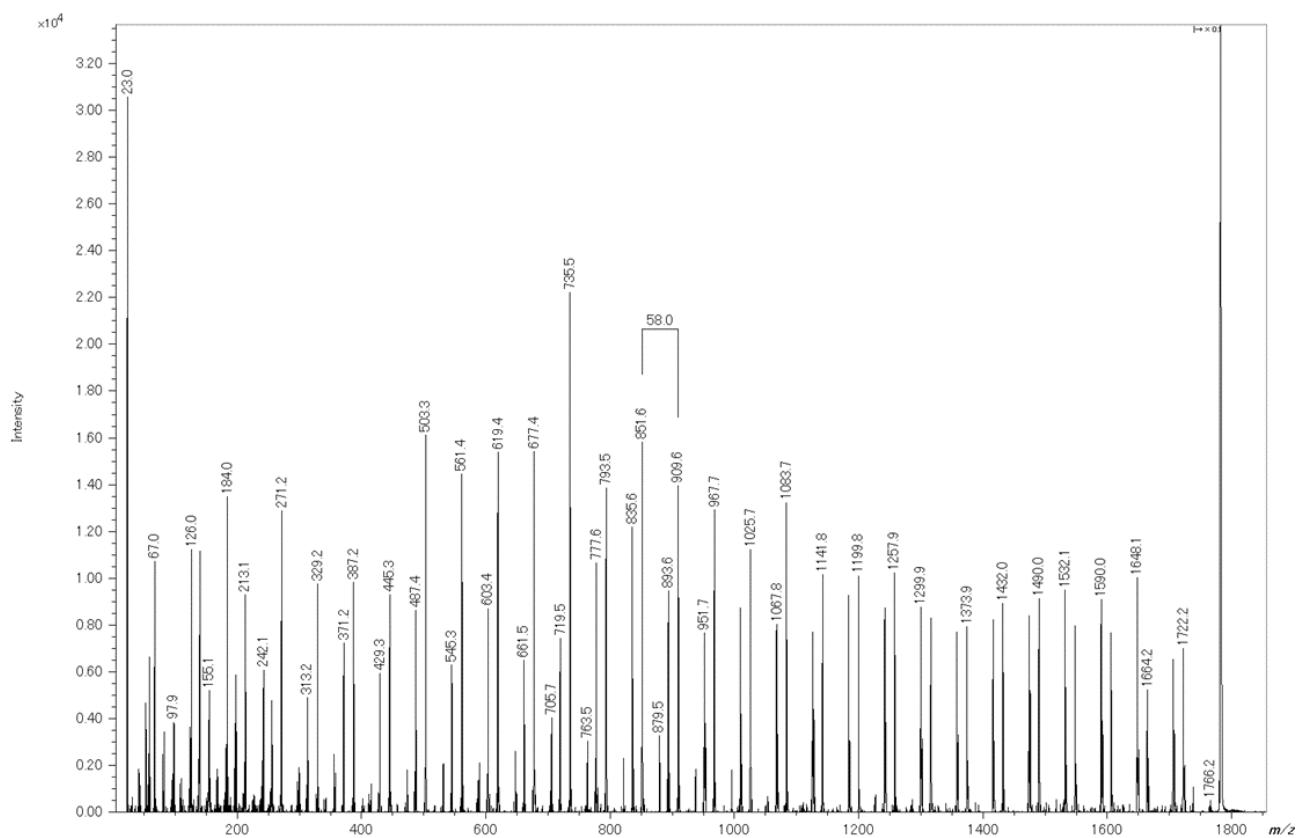


## RKM Plot



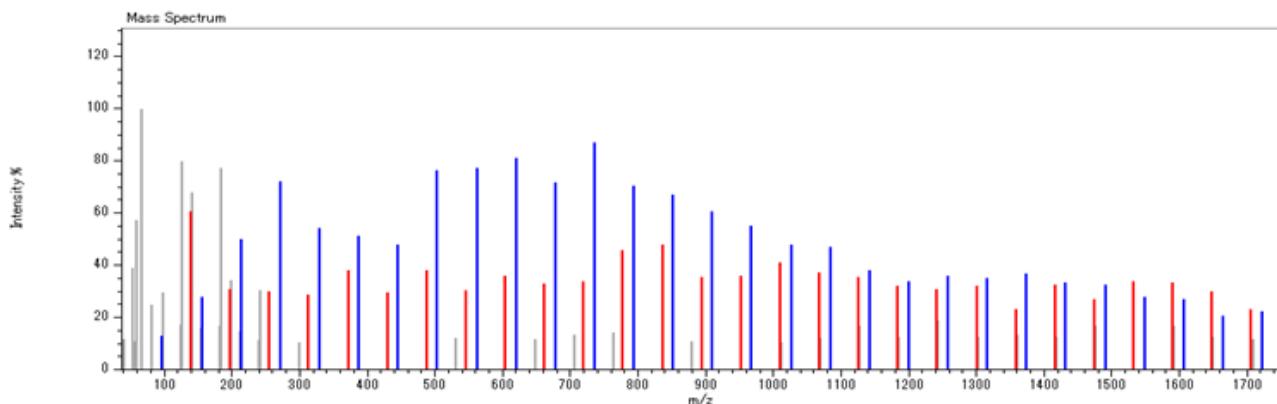
## Estimated Structure



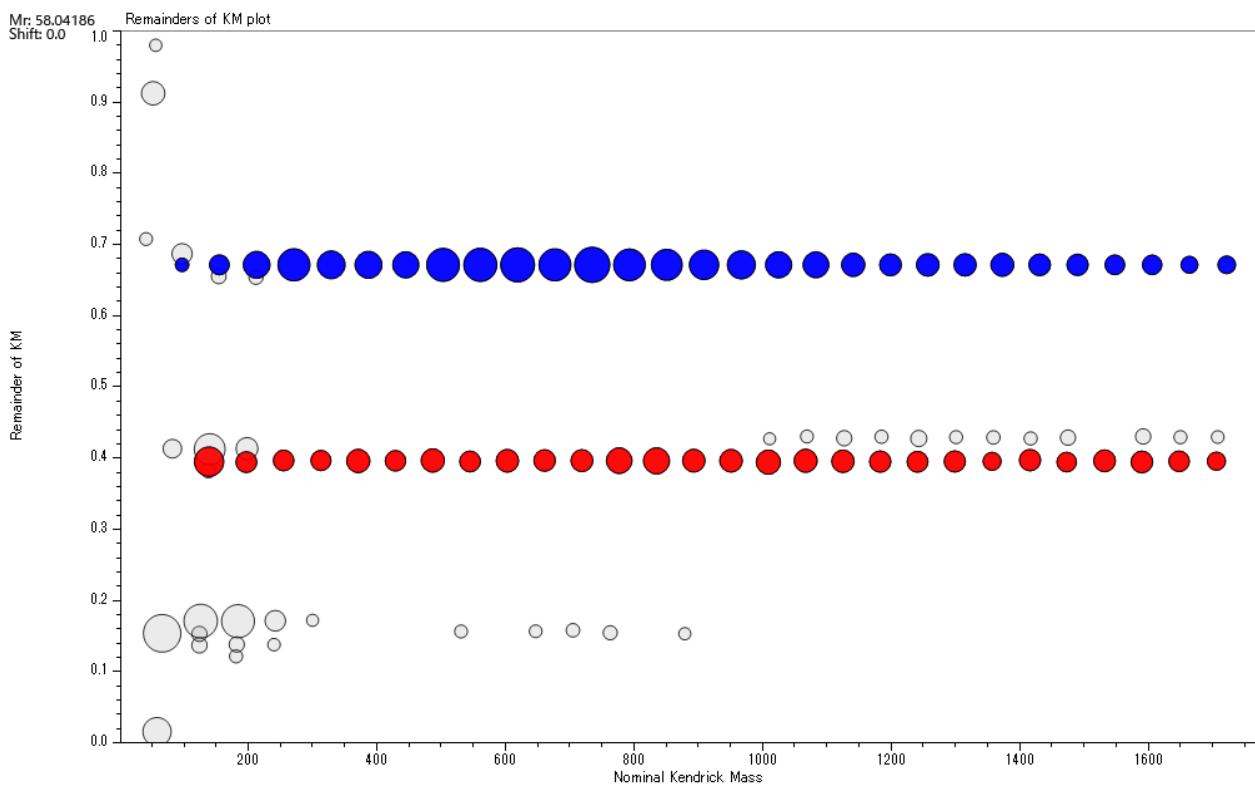
**PPO (poly(propylene oxide))**

Structure	$\text{HO}[\text{CH}_2\text{CH}(\text{CH}_3)\text{O}]_n\text{H}$ ; $\text{HO}(\text{C}_3\text{H}_6\text{O})_n\text{H}$
Ion Species	$[\text{M}+\text{Na}]^+$
Precursor ion	$m/z$ 1,782.25573 ( $n = 30$ )
Matrix	DCTB
Cationization agent	NaTFA
Mode	TOF-TOF:Positive ion

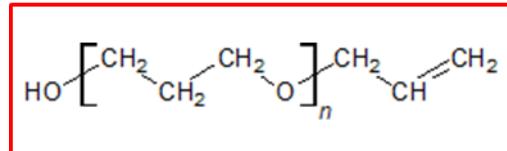
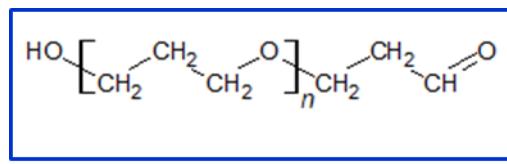
### Peak list

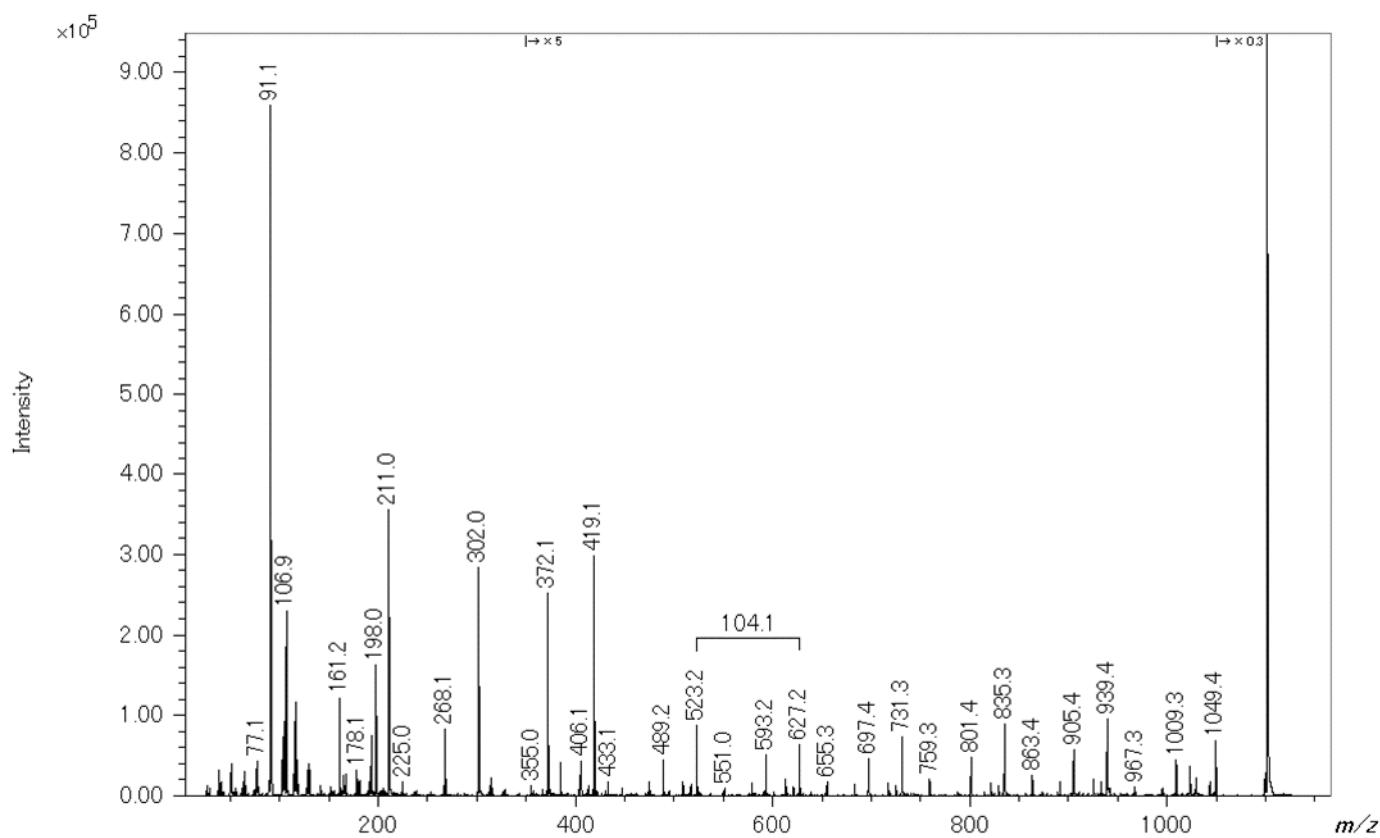


### RKM Plot



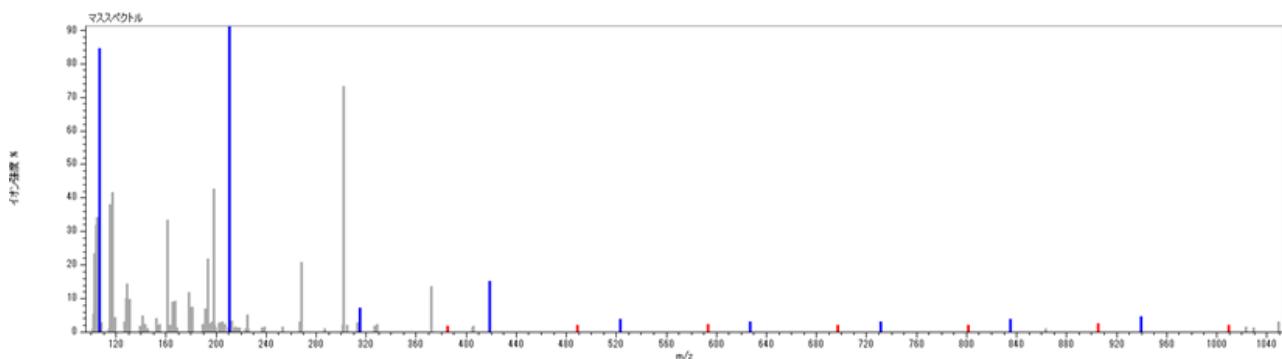
### Estimated Structure



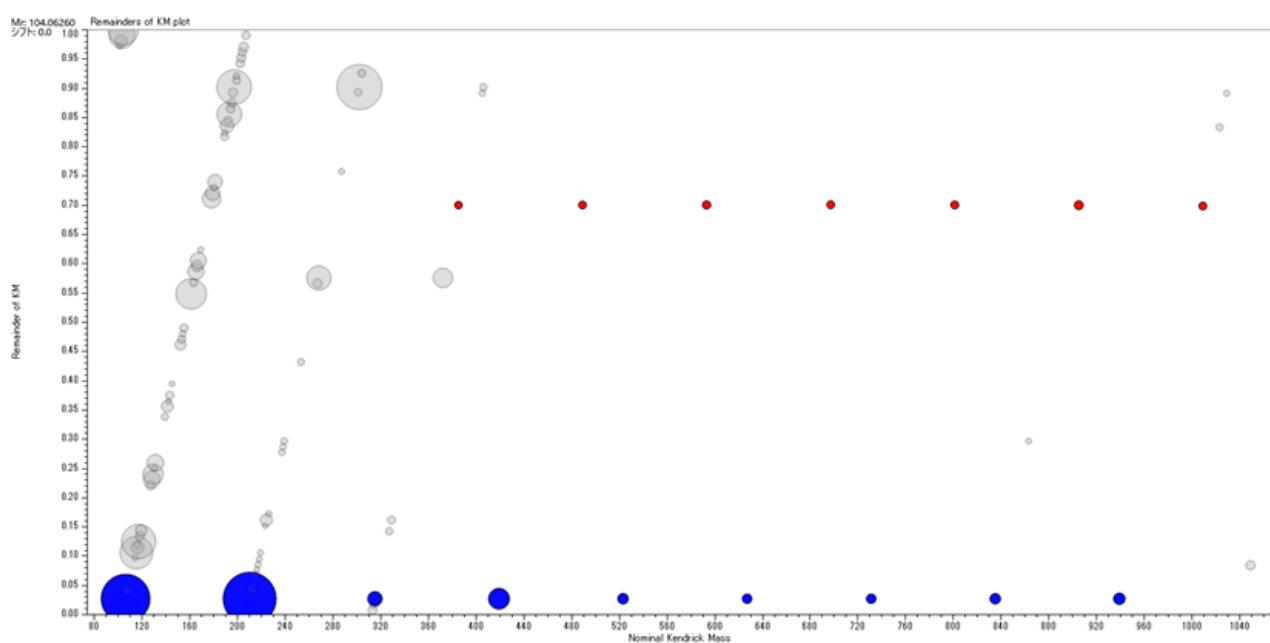
**PS (poly(styrene))**

Structure	$H[CH_2CH(C_6H_5)]_nC_4H_9$ ; $H(C_8H_8)_nC_4H_9$
Ion Species	$[M+Na]^+$
Precursor ion	$m/z$ 1,101.5462 ( $n = 9$ )
Matrix	DCTB
Cationization agent	AgTFA
Mode	TOF-TOF:Positive ion

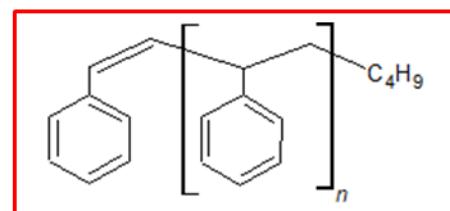
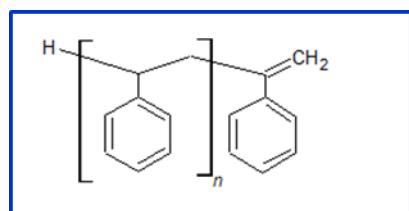
## Peak list



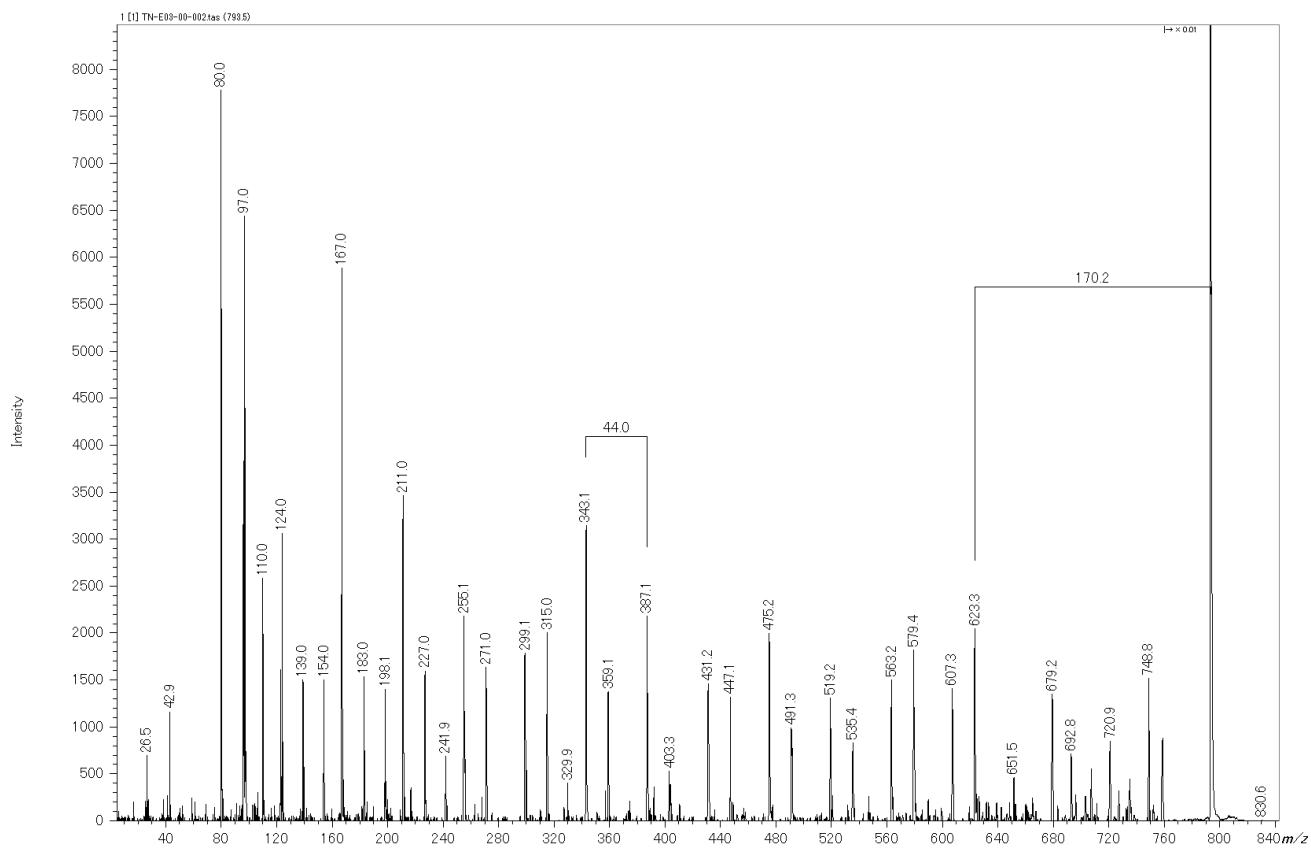
## RKM Plot



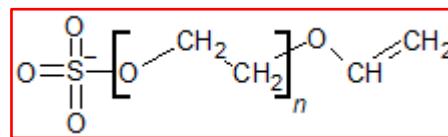
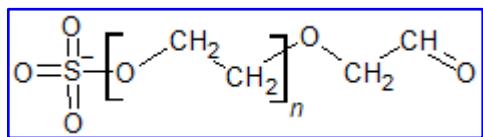
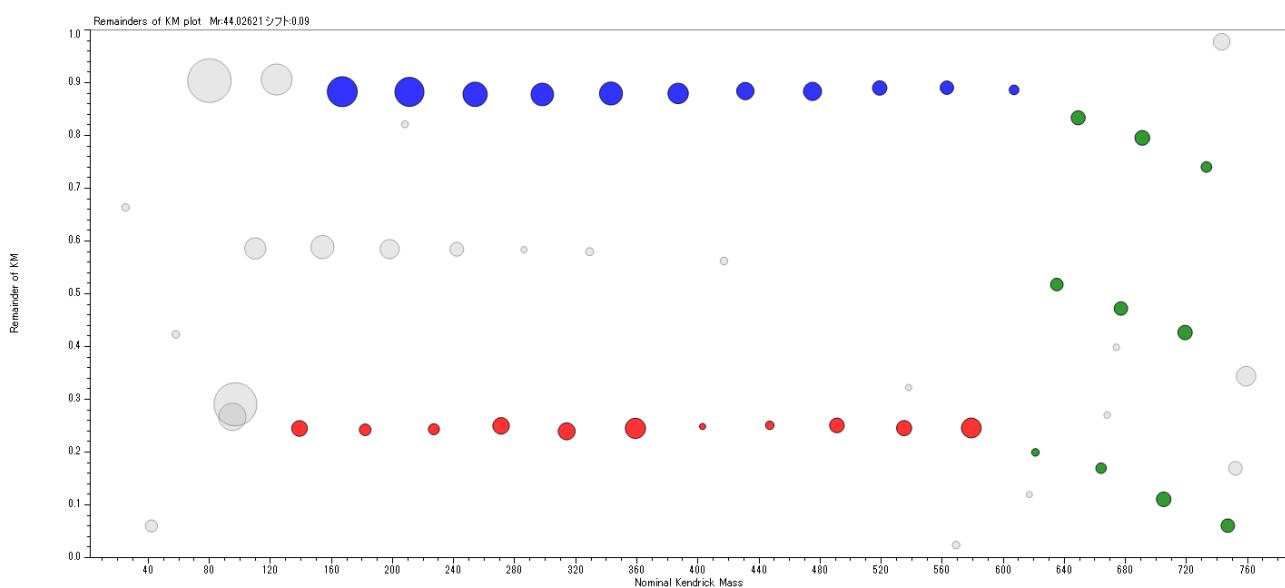
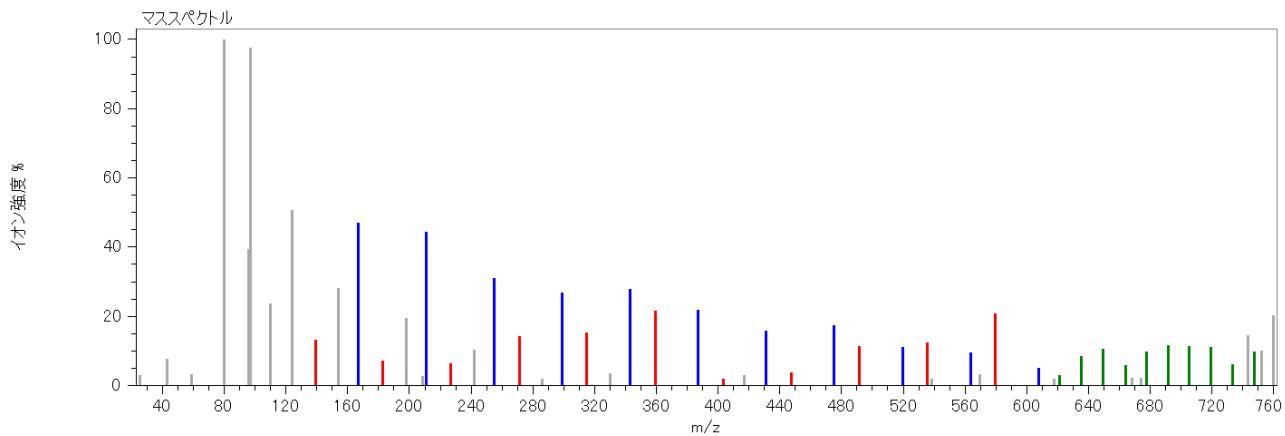
## Estimated Structure



## Polyoxyethylene alkyl ether sulfuric acid



Structure	$\text{NaSO}_3[(\text{CH}_2)_2\text{O}]_n\text{O}(\text{CH}_2)_{11}\text{CH}_3$ ; $\text{NaSO}_3(\text{C}_2\text{H}_4\text{O})_n\text{OC}_{12}\text{H}_{25}$
Ion Species	$[\text{M}-\text{Na}]^-$
Precursor ion	$m/z$ 793.46248 ( $n = 12$ )
Matrix	CHCA
Cationization agent	None
Mode	TOF-TOF:Negative ion



Alkyl group (R) fragment



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