

Closing the gap between small and smaller: Towards a framework to analyse nano- and microplastics in aqueous environmental samples

Svenja M. Mintenig ^{a,b}, Patrick S. Bäuerlein ^b, Albert A. Koelmans ^{c,d}, Stefan C. Dekker ^{a,e}, Annemarie P. van Wezel ^{a,b}

^a Copernicus Institute of Sustainable Development, Utrecht University, The Netherlands.

^b KWR Watercycle Research Institute, Nieuwegein, The Netherlands.

^c Aquatic Ecology and Water Quality Management Group, Wageningen University, The Netherlands.

^d Wageningen Marine Research, IJmuiden, The Netherlands.

^e Faculty of Management, Science and Technology, Open University, Heerlen, The Netherlands

Supporting Information

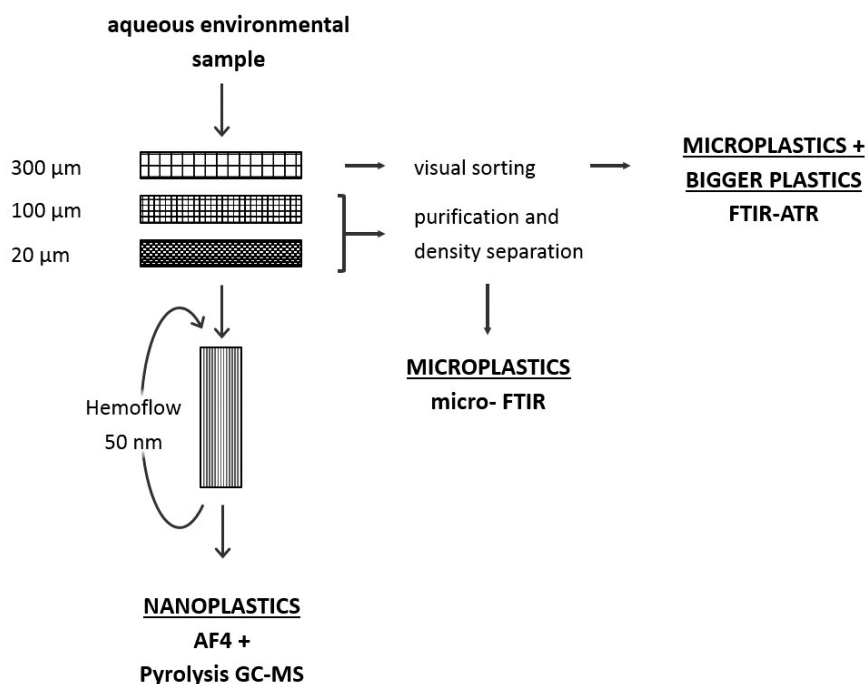


Figure S 1: Scheme of combined techniques to sample and analyse nano- and microplastics.

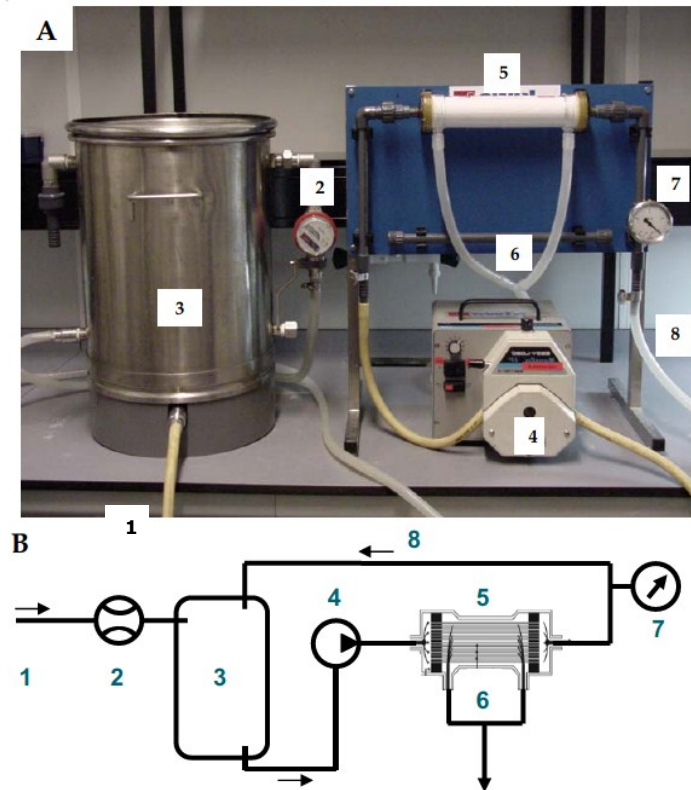


Figure S 2: Schematic presentation of the Hemoflow crossflow ultrafiltration. 1 = water inflow, 2 = water meter, 3 = tank with float valve, 4 = pump, 5 = Hemoflow filter, 6 = permeate, 7 = pressure gauge, 8 = concentrate, circulation back into (3) tank (Source: Veenendaal and Brouwer-Hanzens ¹).

Table S 1: Settings used at the AF4. For the separation of a NP mixture different crossflows were tested.

	Tested Settings (membrane & carrier liquid)	Applied Settings
Membrane	Reg. cellulose (RC) 10 kDa	RC
Carrier liquid	Polyethersulfone (PES) 10 kDa Milli-Q 0.01% SDS 0.01% TWEEN	0.01% SDS
Spacer thickness	250 μm	250 μm
Detector flow	1.0 ml min ⁻¹	1.0 ml min ⁻¹
Split flow	0 ml min ⁻¹	0 ml min ⁻¹
Cross flow	1.5 ml min ⁻¹ (0-11 min) 1.5-0 ml min ⁻¹ (11-50 min, exp. 0.2)	1 ml min ⁻¹ (0-8 min) 1-0 ml min ⁻¹ (8-28 min, exp. 0.2)
Focusing flow	0 ml ml min ⁻¹ (50-65 min)	0 ml min ⁻¹ (28-33 min)
Injection flow	2.3 ml min ⁻¹	1.8 ml min ⁻¹
Injection time	0.2 ml min ⁻¹	0.2 ml min ⁻¹
Injection volume	6 min 30 μl (monodispersed) (10 μl PS polydispersion)	4 min 50 μl

Table S 2: Settings to run the pyrolysis of the samples.

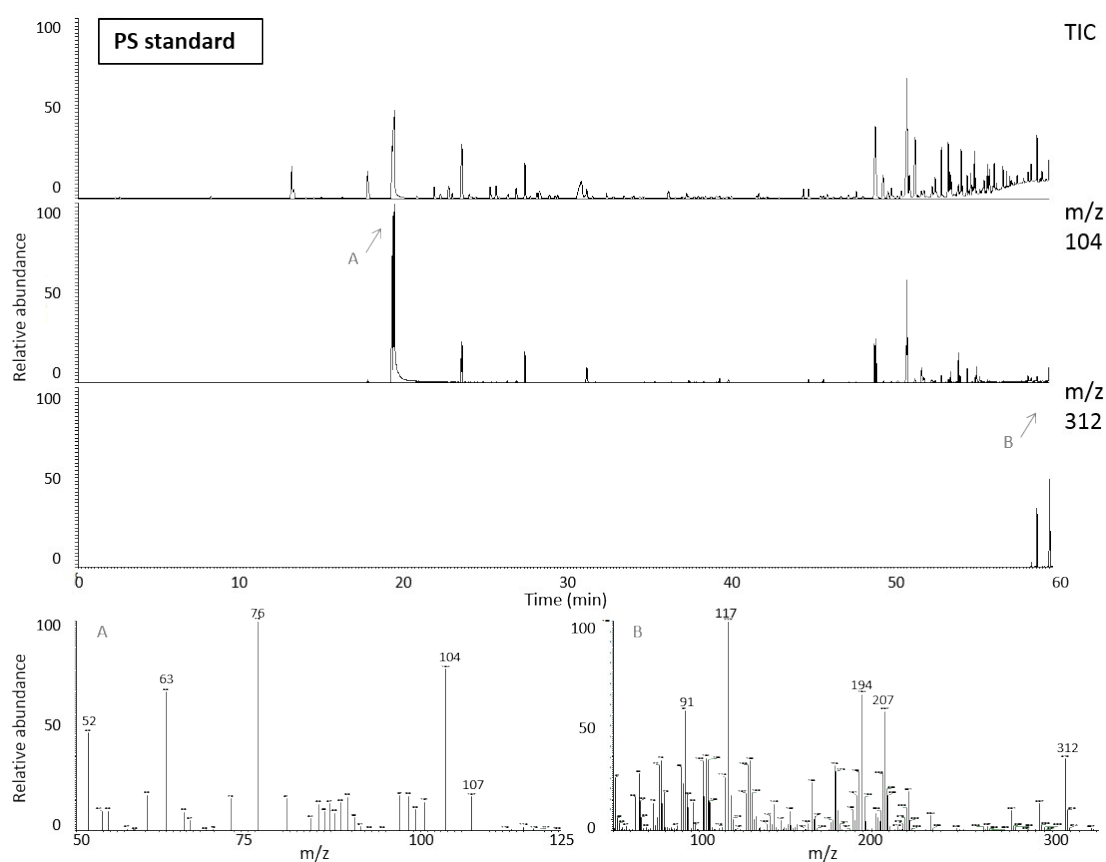
General timing:	
Clean time	20.0 s
Clean time #2	60.0 s
Delay time	0.0 min
Equilibration time	20.0 s
Standby Temperature	
Head temperature	150.0 °C
Offset AS	50.0 °C
Default Parameters	
Temperature	150.0 °C
Pyro Time	10.0 s
Table	single
Pyrolysis Cup	560 °C

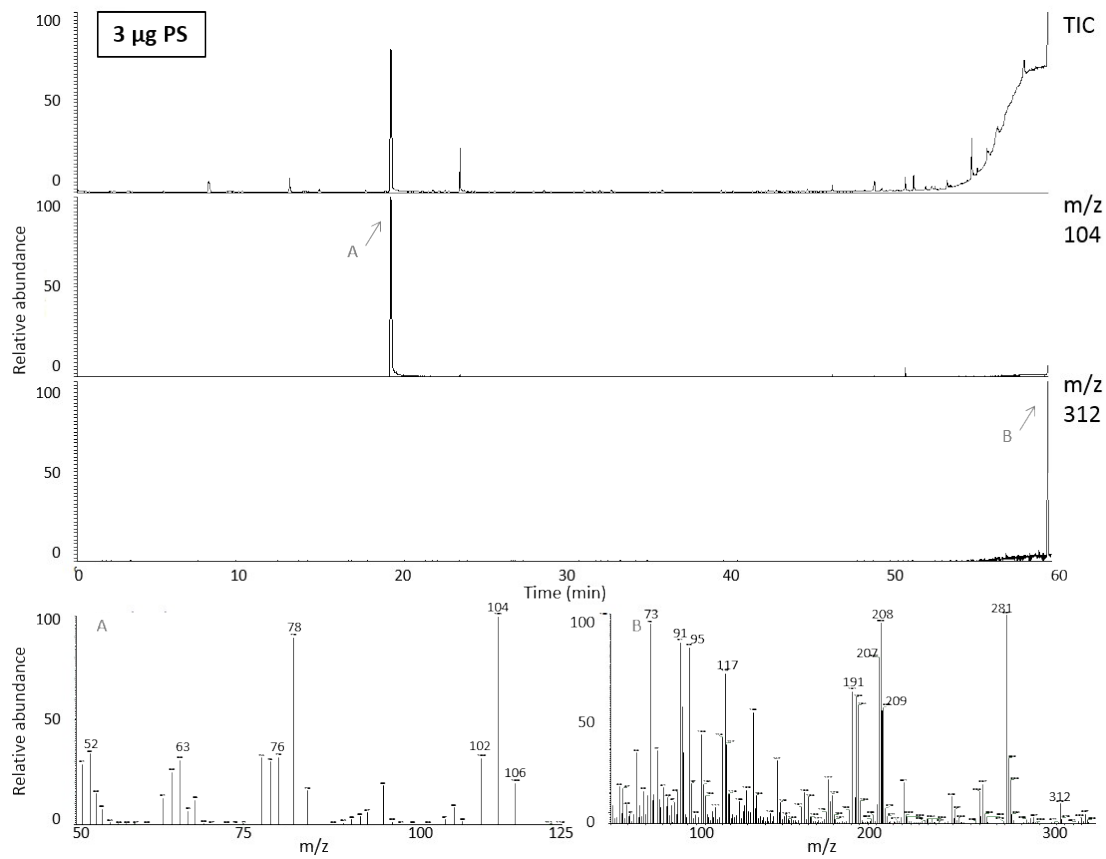
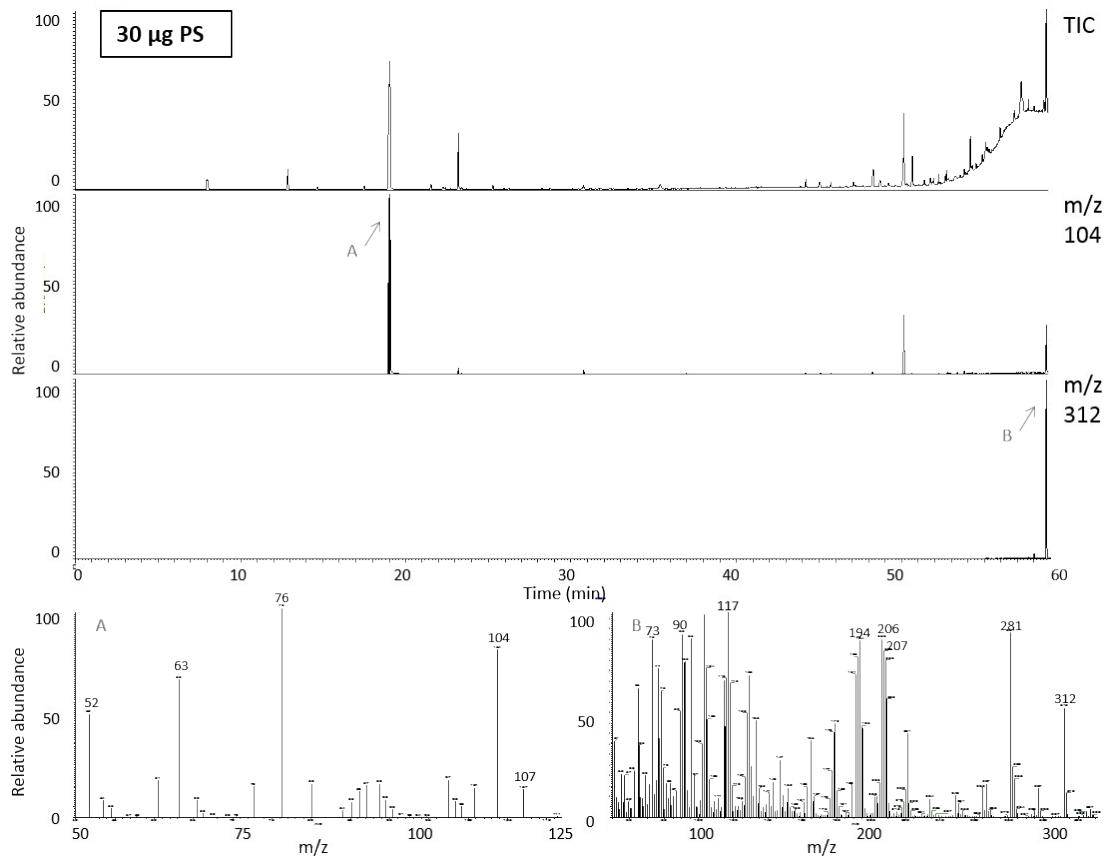
Table S 3: Settings to run the GC-MS.

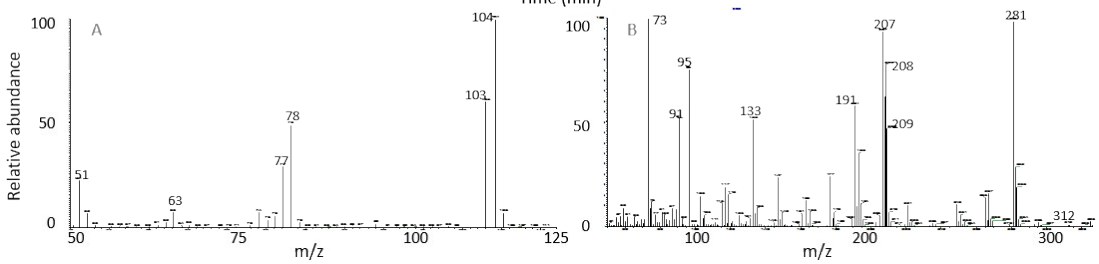
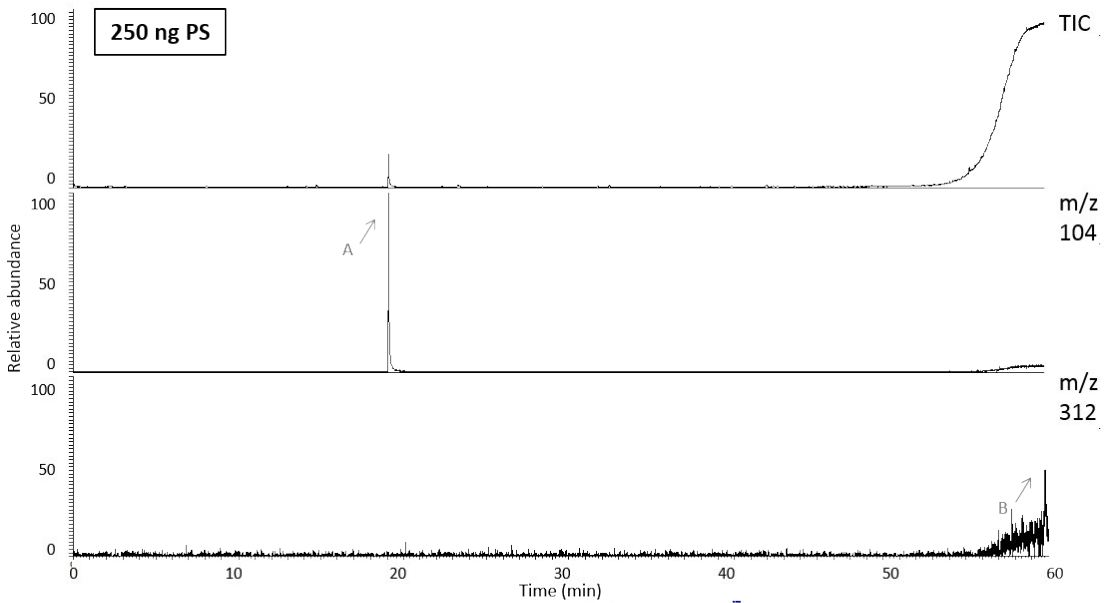
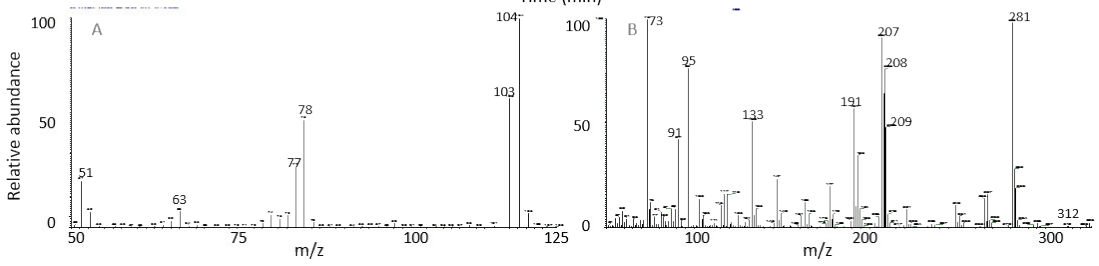
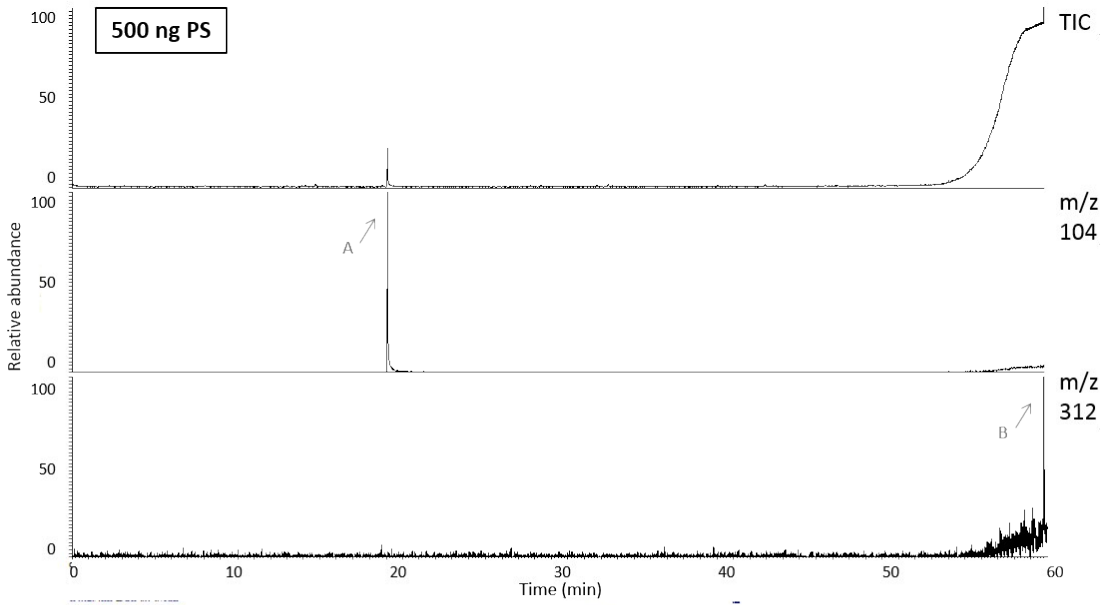
Oven	
Initial Temperature	40 °C
Initial Time	4.00 min
Number of Ramps	2
Rate #1	4.0 °C/min
Final Temperature #1	230 °C
Hold Time #1	0.00 min
Rate #2	20.0 °C/min
Final Temperature #2	325 °C
Hold Time #2	5.00 min
Maximum Temperature	350 °C
Prep Run Timeout	10.00 min
Equilibration Time	0.50 min
Inlet	
Mode	split
Base Temperature	200 °C
Split Flow	40 ml/min
Split ratio	10
Carrier	
Mode	Constant flow
Initial Value	4.00 ml/min
Detector	
Mode	Full scan
Mass Range	50 – 1000 amu
Time Range	0 – 59 min
Peak Format	Centriod
Scan Time	0.40 s
Multiplier	600 V
Ionisation Mode	EI+
Source Temperature	200 °C
Interface Temperature	280 °C

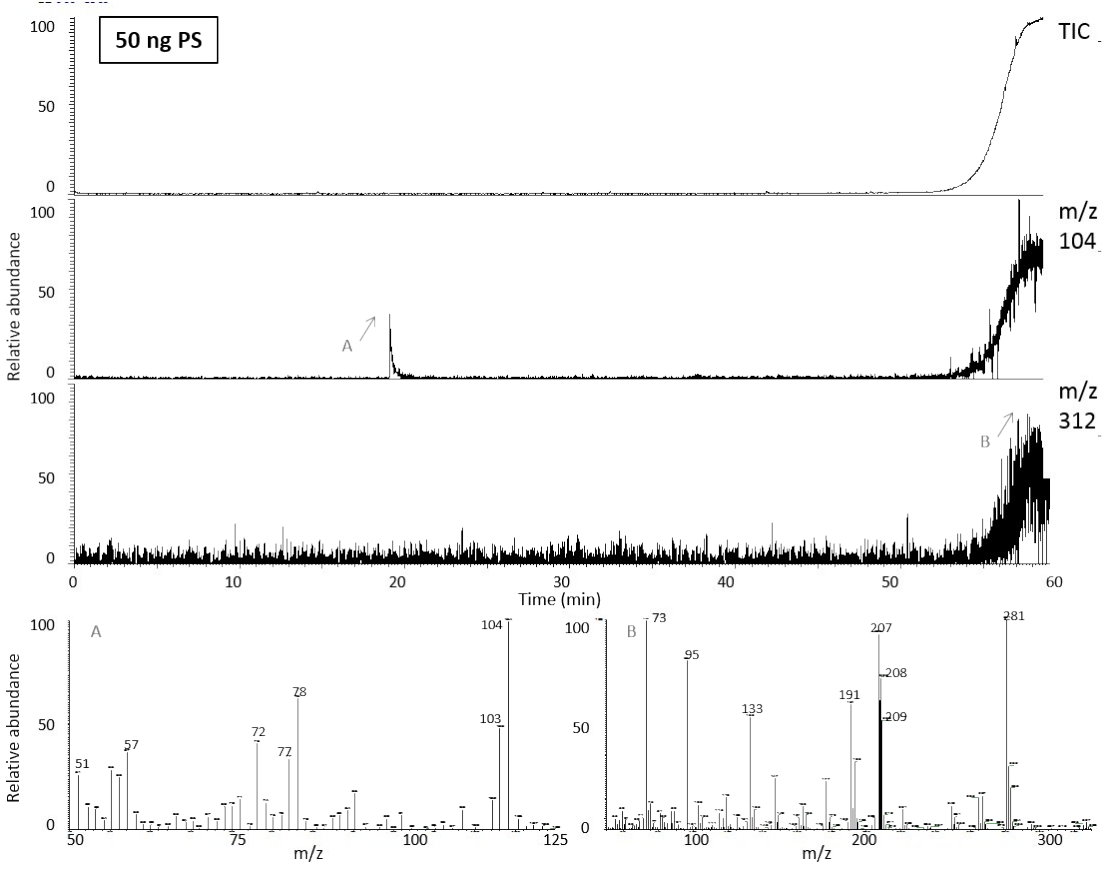
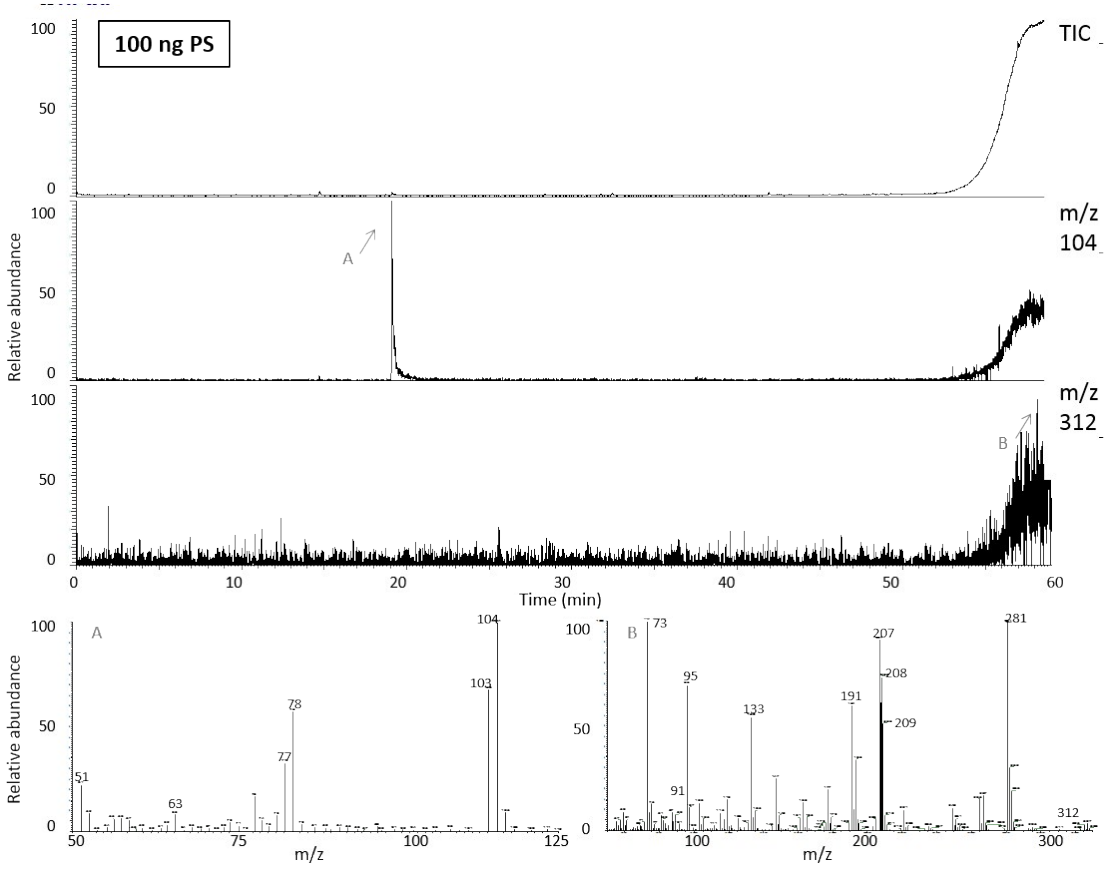
Table S 4: Separation efficiency for various AF4 membrane/ carrier liquid combinations. The fractionation of mono- and polydispersed solutions was concerned successful (marked with an “Y”) when resulting in clear distinct peaks.

	50 nm	500 nm	fractionation of mixture
PES & Milli-Q	Y	Y	-
PES & SDS	-	-	-
PES & TWEEN	-	-	-
RC & Milli-Q	(y)	Y	-
RC & SDS	Y	Y	Y
RC & TWEEN	-	-	-









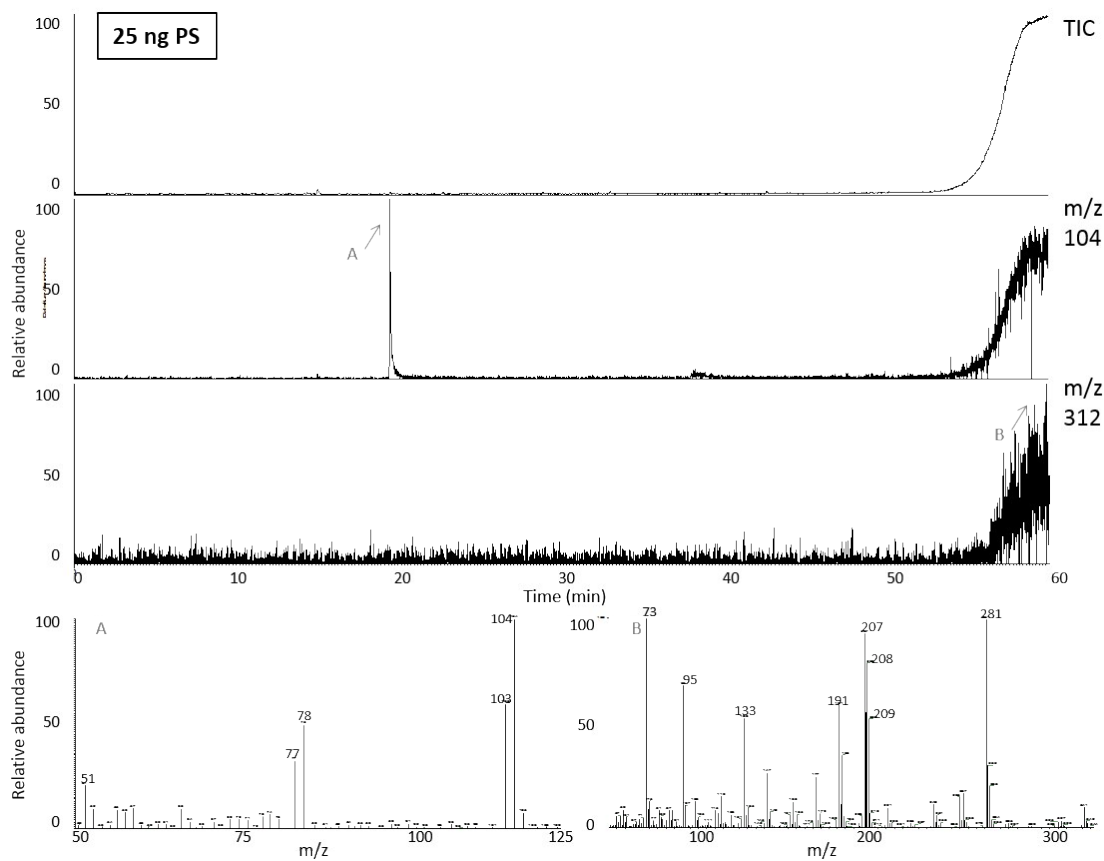


Figure S3: The pyrograms of a PS standard, and of PS (30 μg to 25 ng) that was added to surface water samples after analysis with Pyrolysis GC-MS. Each showing the total ion current (TIC), the chromatogram of selected masses (styrene m/z 104; tri-styrene m/z 312) and the mass spectra of selected peaks (A, B).

References

1. H. R. Veenendaal and A. J. Brouwer-Hanzens, *A method for the concentration of microbes in large volumes of water*, 2007.