

Electronic Supplementary Information (ESI)

Extractable Impurities from Fluoropolymer-Based Membrane Filters – Interference in High-throughput, Untargeted Analysis

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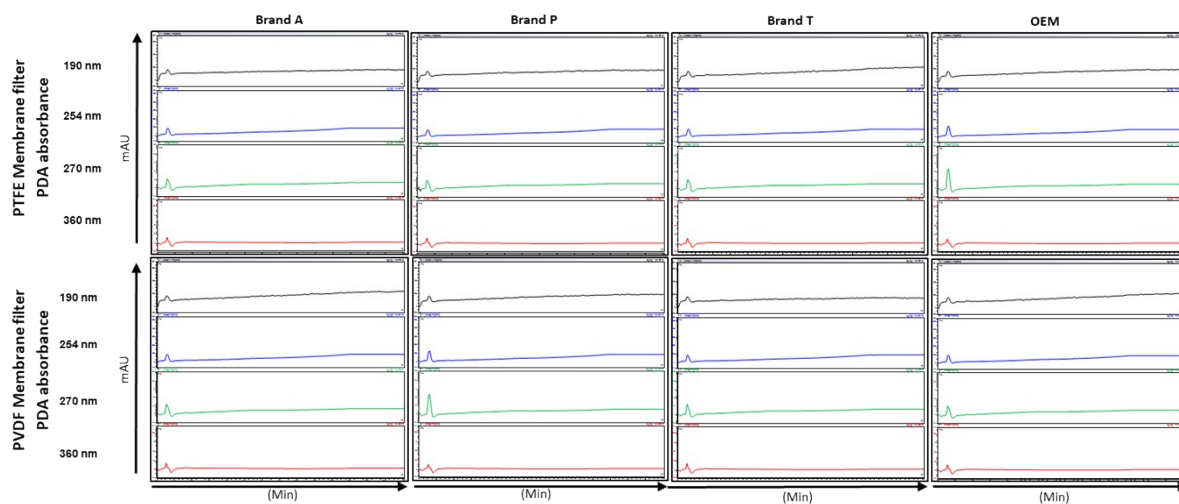


Figure S1: 1st purge of MeOH filtrates eluted from different brands scanned using different ultraviolet wavelengths ranging from 190-380 nm.

Table S 1-1: Detected extractable impurities from **Brand A, PTFE** membrane filter using Quadrupole time of flight coupled with ultra-high performance

Detected m/z^*	RT (min)	Tentative adducts	Molecular formulae	Mass error (ppm)	Signals intensities			Kruskal-Wallis (p -value) ^a	Fold-changes ^b
					1 st purge	3 rd purge	Reduction of impurities, %		
387.1788	4.9	[M+H] ⁺	C ₂₂ H ₂₆ O ₆	3.56	24992	2837	88.7	●	-1.35
409.1603	4.9	[M+Na] ⁺	C ₂₂ H ₂₆ O ₆	4.84	7704	1139	85.2	●●●	-1.28
445.2313	4.9	[M+Na] ⁺	C ₂₂ H ₃₄ N ₂ O ₆	-0.83	17659	2590	85.3	●	-1.14
415.2096	5.3	[M+H] ⁺	C ₁₇ H ₃₅ O ₉ P	-1.19	9760	1046	89.3	●●●	-1.55
473.2624	5.3	[M+H] ⁺	C ₂₆ H ₃₆ N ₂ O ₆	4.62	11381	1602	85.9	●●●	-1.71
274.2738	5.5	[M+H] ⁺	C ₁₆ H ₃₅ N O ₂	0.83	107560	2757	97.4	●●●	-2.41
338.3408	8.2	[M+H] ⁺	C ₂₂ H ₄₃ N O	2.73	18883	2042	89.2	●	-1.48
360.3222	8.2	[M+Na] ⁺	C ₂₂ H ₄₃ N O	4.34	8993	355	96.1	●●●	-2.40
721.5027	9.7	[M+H] ⁺	C ₃₈ H ₇₃ O ₁₀ P	-1.84	57613	7185	87.5	●●●	-1.74
663.4501	9.7	[M+H] ⁺	C ₃₈ H ₆₂ O ₉	-5.15	22969	2442	89.4	●●●	-1.25
701.4053	9.7	[M+Na] ⁺	C ₃₄ H ₆₂ O ₁₃	4.31	6471	1459	77.5	●●	-0.94
685.4315	9.7	[M+Na] ⁺	C ₃₈ H ₆₂ O ₉	-4.43	7793	1054	86.5	●●	-1.65
718.2898	10.0	[M+NH ₄] ⁺	C ₃₂ H ₄₄ O ₁₇	2.66	10226	1327	87.0	●●●	-1.63
702.3159	10.1	[M+NH ₄] ⁺	C ₃₆ H ₄₄ O ₁₃	-5.62	12213	1046	91.4	●	-2.02
697.3613	10.1	[M+H] ⁺	C ₃₉ H ₅₂ O ₁₁	-4.44	14749	796	94.6	●●	-2.21
746.3207	10.2	[M+NH ₄] ⁺	C ₄₁ H ₄₄ O ₁₂	-4.88	10017	2380	76.2	●●	-0.83
725.3921	10.3	[M+Na] ⁺	C ₃₉ H ₅₈ O ₁₁	-7.03	13775	1148	91.7	●	-2.01
730.3476	10.3	[M+Na] ⁺	C ₃₆ H ₅₃ N O ₁₃	-9.50	11600	1765	84.8	●●●	-1.61
753.4235	10.4	[M+H] ⁺	C ₃₆ H ₆₄ O ₁₆	4.33	7846	926	88.2	●●	-1.69
774.3522	10.4	[M+H] ⁺	C ₃₁ H ₅₅ N ₃ O ₁₉	-2.48	8024	1997	75.1	●●	-0.81

liquid chromatography.

* Listed molecules are single-charged

^a●, p -value < 0.05; ●●, p -value < 0.01; ●●●, p -value < 0.001 determined by parametric Kruskal-Wallis test

^bFold changes of peak area relative to mean control. Log₂ transformed. Values > 0 represent increase relative to 1st purge and values < 0 represent decrease relative to 1st purge

Table S 1-2: Detected extractable impurities from **Brand P, PTFE** membrane filter using Quadrupole-time-of flight coupled with ultra-high performance

Detected m/z^*	RT (min)	Tentative adducts	Molecular formulae	Mass error (ppm)	Signals intensities			Kruskal-Wallis (p -value) ^a	Fold-changes ^b
					1 st purge	3 rd purge	Reduction of impurities, %		
172.09465	0.5	[M+Na] ⁺	C ₆ H ₁₅ N O ₃	-1.58	6244	267	95.7	●●●	-4.99
150.11279	0.5	[M+H] ⁺	C ₆ H ₁₅ N O ₃	-2.15	33442	1086	96.8	●●●	-4.95
132.10245	0.6	[M+H] ⁺	C ₆ H ₁₃ N O ₂	-4.16	7018	204	97.1	●●●	-3.21
166.10751	0.6	[M+NH ₄] ⁺	C ₆ H ₁₂ O ₄	-0.85	5501	131	97.6	●●●	-6.20
387.17871	4.9	[M+H] ⁺	C ₂₂ H ₂₆ O ₆	3.90	5836	2193	62.4	●	-1.21
445.23150	4.9	[M+Na] ⁺	C ₂₂ H ₃₄ N ₂ O ₆	-1.40	5697	2044	64.1	●●	-1.39
547.32206	5.3	[M+NH ₄] ⁺	C ₂₇ H ₃₉ N ₅ O ₆	3.40	7913	158	98.0	●●●	-5.59
591.34787	5.4	[M+H] ⁺	C ₃₃ H ₅₀ O ₉	8.28	7715	157	98.0	●●●	-5.80
635.37277	5.5	[M+Na] ⁺	C ₃₃ H ₅₆ O ₁₀	6.20	5585	116	97.9	●●●	-6.88
338.34141	8.2	[M+H] ⁺	C ₂₂ H ₄₃ N O	0.98	17356	5924	65.9	●	-1.42
360.32276	8.2	[M+Na] ⁺	C ₂₂ H ₄₃ N O	2.74	7938	2363	70.2	●●●	-1.56
721.50319	9.7	[M+H] ⁺	C ₃₈ H ₇₃ O ₁₀ P	-2.47	55072	34085	38.1	●●	-1.34
663.45047	9.7	[M+H] ⁺	C ₃₈ H ₆₂ O ₉	-5.75	22178	12156	45.2	●	-1.44
701.40561	9.7	[M+Na] ⁺	C ₃₄ H ₆₂ O ₁₃	3.91	7140	4832	32.3	●●●	-1.40
685.43188	9.7	[M+Na] ⁺	C ₃₈ H ₆₂ O ₉	-4.94	7627	4424	42.0	●●●	-2.02

liquid chromatography.

* Listed molecules are single-charged

^a●, p -value < 0.05; ●●, p -value < 0.01; ●●●, p -value < 0.001 determined by parametric Kruskal-Wallis test

^bFold changes of peak area relative to mean control. Log₂ transformed. Values > 0 represent increase relative to 1st purge and values < 0 represent decrease relative to 1st purge

Table S 1-3: Detected extractable impurities from **Brand T, PTFE** membrane filter using Quadrupole time of flight coupled with ultra-high performance

Detected m/z^*	RT (min)	Tentative adducts	Molecular formulae	Mass error (ppm)	Signals intensities			Kruskal-Wallis (p -value) ^a	Fold-changes ^b
					1 st purge	3 rd purge	Reduction of impurities, %		
445.2324	4.9	[M+Na] ⁺	C ₂₂ H ₃₄ N ₂ O ₆	-3.42	27139	16528	39.1	●●	-1.56
387.1803	4.9	[M+H] ⁺	C ₂₂ H ₂₆ O ₆	-0.19	42488	20175	52.5	●●●	-1.75
409.1615	4.9	[M+Na] ⁺	C ₂₂ H ₂₆ O ₆	1.63	12694	7525	40.7	●●●	-1.82
425.1353	5.0	[M+H] ⁺	C ₂₂ H ₂₀ N ₂ O ₇	-2.36	6729	5267	21.7	●	-0.38
415.2113	5.4	[M+H] ⁺	C ₁₇ H ₃₅ O ₉ P	-5.30	9428	409	95.7	●●●	-1.79
473.2638	5.5	[M+H] ⁺	C ₂₆ H ₃₆ N ₂ O ₆	1.70	11429	594	94.8	●●●	-1.69
274.2751	5.5	[M+H] ⁺	C ₁₆ H ₃₅ N O ₂	-3.75	71064	2439	96.6	●●●	-3.19
338.3425	8.2	[M+H] ⁺	C ₂₂ H ₄₃ N O	-2.37	10116	1907	81.2	●	-1.01
360.3243	8.2	[M+Na] ⁺	C ₂₂ H ₄₃ N O	-1.67	5105	314	93.9	●●	-0.64
721.5036	9.7	[M+H] ⁺	C ₃₈ H ₇₃ O ₁₀ P	-2.98	46259	13268	71.3	●	-1.17
663.4506	9.7	[M+H] ⁺	C ₃₈ H ₆₂ O ₉	-5.95	18813	4420	76.5	●●	-1.25
701.4069	9.7	[M+Na] ⁺	C ₃₄ H ₆₂ O ₁₃	2.04	6417	2520	60.7	●	-0.67
685.4326	9.7	[M+Na] ⁺	C ₃₈ H ₆₂ O ₉	-6.06	6515	1915	70.6	●●	-0.87

liquid chromatography.

* Listed molecules are single-charged

^a●, p -value < 0.05; ●●, p -value < 0.01; ●●●, p -value < 0.001 determined by parametric Kruskal-Wallis test

^bFold changes of peak area relative to mean control. Log₂ transformed. Values > 0 represent increase relative to 1st purge and values < 0 represent decrease relative to 1st purge

Table S 1-4: Detected extractable impurities from **OEM, PTFE** membrane filter using Quadrupole time of flight coupled with ultra-high performance liquid

Detected m/z^*	RT (min)	Tentative adducts	Molecular formulae	Mass error (ppm)	Signals intensities			Kruskal-Wallis (p -value) ^a	Fold-changes ^b
					1 st purge	3 rd purge	Reduction of impurities, %		
387.1779	4.9	[M+H] ⁺	C ₂₂ H ₂₆ O ₆	5.89	23384	745	96.8	●●●	-1.07
445.2302	4.9	[M+Na] ⁺	C ₂₂ H ₃₄ N ₂ O ₆	1.58	15939	726	95.5	●●	-0.80
409.1597	4.9	[M+Na] ⁺	C ₂₂ H ₂₆ O ₆	6.26	7330	331	95.5	●●	-1.45
437.1907	5.4	[M+NH ₄] ⁺	C ₂₁ H ₂₅ N O ₈	2.84	7154	139	98.1	●●●	-2.83
415.2090	5.4	[M+H] ⁺	C ₁₇ H ₃₅ O ₉ P	0.30	10119	223	97.8	●●●	-1.56
473.2615	5.5	[M+H] ⁺	C ₂₆ H ₃₆ N ₂ O ₆	6.51	12350	345	97.2	●●	-1.41
274.2731	5.5	[M+H] ⁺	C ₁₆ H ₃₅ N O ₂	3.57	618291	6833	98.9	●●●	-2.76
296.2547	5.5	[M+Na] ⁺	C ₁₆ H ₃₅ N O ₂	4.94	11874	312	97.4	●	-1.73
230.2471	5.7	[M+H] ⁺	C ₁₄ H ₃₁ N O	3.10	7311	153	97.9	●●●	-2.41
318.2988	5.7	[M+H] ⁺	C ₁₈ H ₃₉ N O ₃	4.73	7432	160	97.9	●●	-2.01
492.4018	8.2	[M+NH ₄] ⁺	C ₃₀ H ₅₀ O ₄	6.21	7397	NA	100.0	-	-
520.4332	8.8	[M+NH ₄] ⁺	C ₃₂ H ₅₄ O ₄	5.72	14659	179	98.8	●●●	-1.36
721.5020	9.7	[M+H] ⁺	C ₃₈ H ₇₃ O ₁₀ P	-0.75	88231	10157	88.5	●●	-1.11
663.4489	9.7	[M+H] ⁺	C ₃₈ H ₆₂ O ₉	-3.44	40390	3520	91.3	●●●	-0.98
701.4048	9.7	[M+Na] ⁺	C ₃₄ H ₆₂ O ₁₃	5.18	8915	1944	78.2	●●	-0.87
685.4310	9.7	[M+Na] ⁺	C ₃₈ H ₆₂ O ₉	-3.60	11047	1495	86.5	●●●	-1.02

chromatography.

* Listed molecules are single-charged

^a●, p -value < 0.05; ●●, p -value < 0.01; ●●●, p -value < 0.001 determined by parametric Kruskal-Wallis test

^bFold changes of peak area relative to mean control. Log₂ transformed. Values > 0 represent increase relative to 1st purge and values < 0 represent decrease relative to 1st purge

Table S 2-1: Detected extractable impurities from **Brand A, PVDF** membrane filter using Quadrupole-time-of flight coupled with ultra-high performance

Detected m/z^*	RT (min)	Tentative adducts	Molecular formulae	Mass error (ppm)	Signals intensities			Kruskal-Wallis (p -value) ^a	Fold-changes ^b
					1 st purge	3 rd purge	Reduction of impurities, %		
568.29327	4.5	[M+H] ⁺	C ₂₁ H ₄₁ N ₇ O ₁₁	0.73	16384	2957	81.95	●●●	-1.21
445.23136	4.9	[M+Na] ⁺	C ₂₂ H ₃₄ N ₂ O ₆	-1.07	13803	1066	92.28	●●	-1.21
387.17866	4.9	[M+H] ⁺	C ₂₂ H ₂₆ O ₆	4.03	19728	1131	94.27	●●●	-1.86
296.25550	5.5	[M+Na] ⁺	C ₁₆ H ₃₅ N O ₂	1.83	26579	3221	87.88	●●●	-1.31
274.27417	5.5	[M+H] ⁺	C ₁₆ H ₃₅ N O ₂	-0.42	1428802	85918	93.99	●●	-1.81
230.24798	5.6	[M+H] ⁺	C ₁₄ H ₃₁ N O	-0.61	99261	2580	97.40	●●●	-1.42
318.29969	5.7	[M+H] ⁺	C ₁₈ H ₃₉ N O ₃	1.83	22997	4727	79.45	●●●	-1.11
338.34181	8.2	[M+H] ⁺	C ₂₂ H ₄₃ N O	-0.20	19671	1671	91.51	●●●	-1.72
360.32239	8.2	[M+Na] ⁺	C ₂₂ H ₄₃ N O	3.84	9364	285	96.96	●●●	-1.74
665.44103	9.5	[M+H] ⁺	C ₃₄ H ₆₅ O ₁₀ P	-3.34	7563	343	95.46	●●	-1.53
721.50284	9.7	[M+H] ⁺	C ₃₈ H ₇₃ O ₁₀ P	-1.98	152351	8306	94.55	●●	-1.44
663.45021	9.7	[M+H] ⁺	C ₃₈ H ₆₂ O ₉	-5.36	75075	2802	96.27	●●	-1.71
685.43190	9.7	[M+Na] ⁺	C ₃₈ H ₆₂ O ₉	-4.98	17092	1231	92.80	●●	-1.06

liquid chromatography.

* Listed molecules are single-charged

^a●, p -value < 0.05; ●●, p -value < 0.01; ●●●, p -value < 0.001 determined by parametric Kruskal-Wallis test

^bFold changes of peak area relative to mean control. Log₂ transformed. Values > 0 represent increase relative to 1st purge and values < 0 represent decrease relative to 1st purge

Table S 2-2: Detected extractable impurities from **Brand P, PVDF** membrane filter using Quadrupole-time-of flight coupled with ultra-high performance liquid chromatography.

* Listed molecules are single-charged

^a●, *p*-value < 0.05; ●●, *p*-value < 0.01; ●●●, *p*-value < 0.001 determined by parametric Kruskal-Wallis test

Detected <i>m/z</i> *	RT (min)	Tentative adducts	Molecular formulae	Mass error (ppm)	Signals intensities			Kruskal-Wallis (<i>p</i> -value) ^a	Fold-changes ^b
					1 st purge	3 rd purge	Reduction of impurities, %		
164.12852	0.5	[M+H] ⁺	C ₇ H ₁₇ N O ₃	-2.45	8417	306	96.36	●●●	-2.21
178.14397	0.5	[M+NH ₄] ⁺	C ₈ H ¹⁶ O ₃	-1.25	10712	391	96.35	●●●	-1.31
162.14914	0.7	[M+H] ⁺	C ₈ H ₁₉ N O ₂	-1.77	5089	195	96.17	●	-2.14
192.15975	0.8	[M+H] ⁺	C ₉ H ₂₁ N O ₃	-1.73	10088	1298	87.13	●	-1.63
304.28455	4.9	[M+NH ₄] ⁺	C ₁₇ H ₃₄ O ₃	0.25	183484	10164	94.46	●●●	-1.26
316.28423	5.0	[M+H] ⁺	C ₁₈ H ₃₇ N O ₃	1.24	14408	847	94.12	●●●	-1.21
387.17928	5.1	[M+NH ₄] ⁺	C ₂₃ H ₁₉ N ₃ O ₂	6.16	6590	1757	73.34	●	-0.84
635.59063	5.3	[M+H] ⁺	C ₃₂ H ₇₄ N ₈ O ₄	-0.08	47044	N/A	100.00	-	-
362.32605	5.4	[M+H] ⁺	C ₂₀ H ₄₃ N O ₄	1.20	87885	11691	86.70	●●●	-0.69
288.28976	5.5	[M+H] ⁺	C ₁₇ H ₃₇ N O ₂	-0.19	11067	1450	86.90	●	-0.93
332.31577	5.5	[M+NH ₄] ⁺	C ₁₉ H ₃₈ O ₃	0.48	546457	35512	93.50	●●●	-1.58
318.30074	5.7	[M+H] ⁺	C ₁₈ H ₃₉ N O ₃	-1.48	2724822	434688	84.05	●●●	-1.20
321.26317	5.7	[M+Na] ⁺	C ₁₆ H ₃₄ N ₄ O	-2.31	48521	1052	97.83	●●●	-1.57
470.36726	5.7	[M+Na] ⁺	C ₂₂ H ₄₉ N ₅ O ₄	0.93	72748	3799	94.78	●●	-1.95
514.39298	5.8	[M+Na] ⁺	C ₂₄ H ₅₃ N ₅ O ₅	1.85	53396	3362	93.70	●●●	-1.62
484.38262	6.0	[M+Na] ⁺	C ₂₃ H ₅₁ N ₅ O ₄	1.53	157977	6307	96.01	●●●	-1.79
528.40850	6.0	[M+Na] ⁺	C ₂₅ H ₅₅ N ₅ O ₅	2.06	131285	7377	94.38	●●	-1.88
335.27886	6.1	[M+Na] ⁺	C ₁₇ H ₃₆ N ₄ O	-2.33	63868	972	98.48	●●●	-1.82
572.43424	6.1	[M+Na] ⁺	C ₂₇ H ₅₉ N ₅ O ₆	2.76	103108	7355	92.87	●	-1.59
616.45990	6.2	[M+NH ₄] ⁺	C ₃₇ H ₅₈ O ₆	-4.57	76127	6710	91.19	●	-1.48
349.29432	6.4	[M+H] ⁺	C ₁₉ H ₄₀ O ₅	1.52	66578	1369	97.94	●●●	-1.81
377.32540	6.5	[M+Na] ⁺	C ₂₀ H ₄₂ N ₄ O	-0.90	271253	6976	97.43	●●●	-1.89
586.44980	6.6	[M+NH ₄] ⁺	C ₃₆ H ₅₆ O ₅	-5.63	80630	5337	93.38	●●	-1.76
630.47551	6.6	[M+NH ₄] ⁺	C ₃₈ H ₆₀ O ₆	-4.40	57057	5105	91.05	●●	-1.11
333.29962	6.9	[M+H] ⁺	C ₁₉ H ₄₀ O ₄	0.95	65947	1539	97.67	●●	-1.53

614.48077	6.8	[M+NH ₄] ⁺	C ₃₈ H ₆₀ O ₅	-4.81	142766	12653	91.14	●●●	-1.93
658.50671	6.8	[M+NH ₄] ⁺	C ₄₁ H ₆₀ N ₄ O ₂	-1.96	98583	10203	89.65	●●●	-0.71
702.53250	7.0	[M+NH ₄] ⁺	C ₄₂ H ₆₈ O ₇	-3.17	64635	7678	88.12	●●●	-1.02
353.26556	7.1	[M+Na] ⁺	C ₁₉ H ₃₈ O ₄	2.03	34589	3979	88.50	●	-0.63
359.31493	7.8	[M+H] ⁺	C ₂₁ H ₄₂ O ₄	1.83	11045	485	95.61	●●●	-2.35
341.30453	7.9	[M+H] ⁺	C ₂₁ H ₄₀ O ₃	1.44	17312	800	95.38	●●	-2.26
381.29675	7.5	[M+Na] ⁺	C ₂₁ H ₄₂ O ₄	2.18	85952	10250	88.07	●	-0.97
267.26817	7.6	[M+H] ⁺	C ₁₈ H ₃₄ O	0.27	8024	431	94.63	●●	-1.93
695.35529	7.8	[M+H] ⁺	C ₃₈ H ₅₀ N ₂ O ₁₀	-2.11	10634	1717	83.85	●	-2.79
338.34161	8.2	[M+H] ⁺	C ₂₂ H ₄₃ NO	0.39	6973	5128	26.46	●	-0.32
738.38508	8.3	[M+Na] ⁺	C ₂₈ H ₅₇ N ₇ O ₁₄	0.69	6924	2554	63.11	●	-0.72
458.45521	8.4	[M+NH ₄] ⁺	C ₂₈ H ₅₆ O ₃	3.54	7113	368	94.83	●●	-2.03
472.47111	8.6	[M+NH ₄] ⁺	C ₂₉ H ₅₈ O ₃	2.89	225560	9729	95.69	●●●	-2.68
679.36023	8.6	[M+Na] ⁺	C ₃₇ H ₄₈ N ₆ O ₅	-3.64	5801	1443	75.12	●●	-1.42
486.48655	8.8	[M+NH ₄] ⁺	C ₃₀ H ₆₀ O ₃	3.25	36366	427	98.83	●●●	-1.82
663.45062	9.7	[M+H] ⁺	C ₃₈ H ₆₂ O ₉	-5.98	30511	4946	83.79	●●●	-1.68
721.50318	9.7	[M+H] ⁺	C ₃₈ H ₇₃ O ₁₀ P	-2.45	68094	14886	78.14	●●	-1.64
685.43224	9.7	[M+Na] ⁺	C ₃₈ H ₆₂ O ₉	-5.49	9095	2113	76.77	●●	-1.62
579.53232	10.1	[M+H] ⁺	C ₃₇ H ₇₀ O ₄	4.09	27367	3934	85.63	●●●	-1.98
619.52456	10.1	[M+Na] ⁺	C ₃₇ H ₇₂ O ₅	4.42	30693	13111	57.28	●	-0.47
607.56350	10.3	[M+H] ⁺	C ₃₉ H ₇₄ O ₄	4.10	219735	24254	88.96	●●	-0.85
647.55565	10.3	[M+Na] ⁺	C ₃₉ H ₇₆ O ₅	4.56	54029	25821	52.21	●	-1.13
642.60004	10.4	[M+NH ₄] ⁺	C ₃₉ H ₇₆ O ₅	4.90	16579	547	96.70	●●●	-1.84

^bFold changes of peak area relative to mean control. Log₂ transformed. Values > 0 represent increase relative to 1st purge and values < 0 represent decrease relative to 1st purge

Table S2-3: Detected extractable impurities from **Brand T, PVDF** membrane filter using Quadrupole-time-of flight coupled with ultra-high performance

Detected m/z^*	RT (min)	Tentative adducts	Molecular formulae	Mass error (ppm)	Signals intensities			Kruskal-Wallis (p -value) ^a	Fold-changes ^b
					1 st purge	3 rd purge	Reduction of impurities, %		
409.16034	4.9	[M+Na] ⁺	C ₂₂ H ₂₆ O ₆	4.71	7487	380	94.92	●●	-1.69
445.23093	4.9	[M+Na] ⁺	C ₂₂ H ₃₄ N ₂ O ₆	-0.05	15085	854	94.34	●●	-1.72
387.17860	4.9	[M+H] ⁺	C ₂₂ H ₂₆ O ₆	4.18	22440	939	95.82	●●●	-1.80
415.20965	5.4	[M+H] ⁺	C ₁₇ H ₃₅ O ₉ P	-1.22	7862	198	97.48	●●●	-1.81
473.26251	5.3	[M+Na] ⁺	C ₂₂ H ₄₃ O ₇ P	3.00	9429	257	97.27	●	-1.87
296.25495	5.5	[M+Na] ⁺	C ₁₆ H ₃₅ N O ₂	3.84	14430	2187	84.84	●●●	-1.52
437.19151	5.5	[M+NH ₄] ⁺	C ₂₁ H ₂₅ N O ₈	0.79	6263	147	97.65	●●●	-1.78
274.27362	5.5	[M+H] ⁺	C ₁₆ H ₃₅ N O ₂	1.59	806372	53297	93.39	●●	-1.55
230.24763	5.6	[M+H] ⁺	C ₁₄ H ₃₁ N O	0.92	17558	763	95.65	●●●	-1.64
318.29949	5.7	[M+H] ⁺	C ₁₈ H ₃₉ N O ₃	2.46	16126	1721	89.33	●●●	-0.89
338.34016	8.2	[M+H] ⁺	C ₂₂ H ₄₃ N O	4.69	13743	2284	83.38	●●●	-1.66
360.32230	8.2	[M+Na] ⁺	C ₂₂ H ₄₃ N O	4.11	6262	651	89.60	●	-0.92
663.44944	9.7	[M+H] ⁺	C ₃₈ H ₆₂ O ₉	-4.20	51440	3240	93.70	●●●	-1.14
701.40482	9.7	[M+Na] ⁺	C ₃₄ H ₆₂ O ₁₃	5.07	9317	1923	79.36	●●	-0.69
685.43089	9.7	[M+Na] ⁺	C ₃₈ H ₆₂ O ₉	-3.45	12762	1351	89.41	●	-1.61
721.50202	9.7	[M+H] ⁺	C ₃₈ H ₇₃ O ₁₀ P	-0.84	106323	9649	90.92	●●●	-0.99

liquid chromatography.

* Listed molecules are single-charged

^a●, p -value < 0.05; ●●, p -value < 0.01; ●●●, p -value < 0.001 determined by parametric Kruskal-Wallis test

^bFold changes of peak area relative to mean control. Log₂ transformed. Values > 0 represent increase relative to 1st purge and values < 0 represent decrease relative to 1st purge

Table S 2-4: Detected extractable impurities from **OEM, PVDF** membrane filter using Quadrupole-time-of flight coupled with ultra-high performance liquid

Detected <i>m/z</i> *	RT (min)	Tentative adducts	Molecular formulae	Mass error (ppm)	Signals intensities			Kruskal-Wallis (<i>p</i> -value) ^a	Fold-changes ^b
					1 st purge	3 rd purge	Reduction of impurities, %		
274.27292	5.5	[M+H] ⁺	C ₁₆ H ₃₅ N O ₂	4.16	7726	998	87.08	●●●	-1.46
288.28856	5.7	[M+NH ₄] ⁺	C ₁₇ H ₃₄ O ₂	4.24	45422	4070	91.04	●●●	-1.94
301.13964	6.0	[M+Na] ⁺	C ₁₆ H ₂₂ O ₄	4.99	11150	381	96.58	●	-0.79
205.08491	6.1	[M+H] ⁺	C ₁₂ H ₁₂ O ₃	4.95	5050	158	96.87	●	-0.85
316.31927	6.3	[M+NH ₄] ⁺	C ₁₉ H ₃₈ O ₂	5.82	7525	997	86.75	●●●	-1.97
353.26433	6.9	[M+Na] ⁺	C ₁₉ H ₃₈ O ₄	5.75	75362	1289	98.29	●●●	-1.64
313.27203	6.9	[M+H] ⁺	C ₁₉ H ₃₆ O ₃	5.42	16267	100	99.39	●●●	-1.85
331.28313	7.1	[M+H] ⁺	C ₁₉ H ₃₈ O ₄	3.50	7477	N/A	100.00	-	-
381.29515	7.5	[M+Na] ⁺	C ₂₁ H ₄₂ O ₄	6.64	58864	716	98.78	●●	-1.61
341.30298	7.5	[M+H] ⁺	C ₂₁ H ₄₀ O ₃	6.00	9670	N/A	100.00	-	-
359.31373	7.5	[M+H] ⁺	C ₂₁ H ₄₂ O ₄	5.18	6045	N/A	100.00	-	-
492.40173	8.1	[M+NH ₄] ⁺	C ₃₀ H ₅₀ O ₄	6.34	79756	432	99.46	●●●	-6.47
520.43296	8.7	[M+NH ₄] ⁺	C ₃₂ H ₅₄ O ₄	6.12	169811	632	99.63	●●●	-6.91
534.44833	8.9	[M+NH ₄] ⁺	C ₃₃ H ₅₆ O ₄	6.50	5171	143	97.23	●●●	-5.43
663.44876	9.7	[M+H] ⁺	C ₃₈ H ₆₂ O ₉	-3.17	32161	1446	95.50	●●	-1.87
721.50163	9.7	[M+H] ⁺	C ₃₈ H ₇₃ O ₁₀ P	-0.30	74355	5246	92.94	●●●	-1.92
701.40461	9.7	[M+Na] ⁺	C ₃₄ H ₆₂ O ₁₃	5.38	6687	852	87.26	●●●	-1.39
685.43059	9.7	[M+Na] ⁺	C ₃₈ H ₆₂ O ₉	-2.99	8480	622	92.67	●●●	-2.71
526.51643	9.8	[M+H] ⁺	C ₃₃ H ₆₇ N O ₃	5.60	5306	252	95.25	●●●	-3.62

chromatography.

* Listed molecules are single-charged

^a●, *p*-value < 0.05; ●●, *p*-value < 0.01; ●●●, *p*-value < 0.001 determined by parametric Kruskal-Wallis test

^bFold changes of peak area relative to mean control. Log₂ transformed. Values > 0 represent increase relative to 1st purge and values < 0 represent decrease relative to 1st purge

