



*Supplement of*

**Controlled chamber formation of per- and polyfluoroalkyl substances (PFAS) aerosols with *Pseudomonas fluorescens*: size distributions, effects, and inhalation deposition potential**

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## Content

<b>Table S1.</b> <sup>13</sup> C labelled PFAS in the EPA 533 ES isotope dilution standard mix (Wellington laboratories Inc, Canada).....	<b>3</b>
<b>Table S2.</b> Breathing parameters used in MPPD model (version 2.11, Applied Research Associates, Inc.) to estimate deposition fraction as a function of particle size.....	<b>4</b>
<b>Table S3.</b> Regional (head, tracheobronchial region – TB, and Pulmonary/Alveolar Region – Pul) and total deposition efficiencies calculated using MPPD (version 2.11, Applied Research Associates, Inc.) for each nanoMOUDI size bin.....	<b>5</b>
<b>Table S4.</b> Size segregated deposition flux (pg/h) estimates in the human respiratory tract.....	<b>6-8</b>
<b>Figure S1.</b> Schematic representation of the experimental setup in the ChAMBRé atmospheric simulation facility. (1) ChAMBRé; (2) three-jet Collison nebuliser used for aerosolisation of the PFAS-containing solution; (3) Sparging Liquid Aerosol Generator (SLAG) used for nebulisation of Pseudomonas fluorescens; (4) Optical Particle Sizer (OPS); (5) Scanning Mobility Particle Sizer (SMPS); (6) Waveband Integrated Bioaerosol Sensor (WIBS-NEO); and (7) Nano Micro-Orifice Uniform Deposit Impactor (NanoMOUDI) used for size-resolved aerosol sampling.....	<b>9</b>
<b>Figure S2.</b> MPPD modelled deposition efficiency in the head, tracheobronchial region, pulmonary region and total human respiratory tract as a function of particle size. MPPD model parameters are given in Table S2.....	<b>10</b>
<b>Chamber dilution correction</b> .....	<b>11</b>

**Table S1** <sup>13</sup>C labelled PFAS in the EPA 533 ES isotope dilution standard mix (Wellington laboratories Inc, Canada).

<sup>13</sup> C labelled PFAS	Abbreviation
Perfluorobutanoic acid, <sup>13</sup> C <sub>4</sub>	MPFBA
Perfluorobutane sulfonic acid, <sup>13</sup> C <sub>3</sub>	M3PFBS
Perfluoropentanoic acid, <sup>13</sup> C <sub>5</sub>	M5PFPeA
4:2 fluorotelomer sulfonate, <sup>13</sup> C <sub>2</sub>	M2-4:2FTS
Perfluorohexanoic acid, <sup>13</sup> C <sub>5</sub>	M5PFHxA
Perfluoroheptanoic acid, <sup>13</sup> C <sub>4</sub>	M4PFHpA
Perfluorohexane sulfonic acid, <sup>13</sup> C <sub>3</sub>	M3PFHxS
Perfluorooctanoic acid, <sup>13</sup> C <sub>8</sub>	M8PFOA
6:2 fluorotelomer sulfonate, <sup>13</sup> C <sub>2</sub>	M2-6:2FTS
Perfluorononanoic acid, <sup>13</sup> C <sub>9</sub>	M9PFNA
Perfluorooctane sulfonic acid, <sup>13</sup> C <sub>8</sub>	M8PFOS
Perfluorodecanoic acid, <sup>13</sup> C <sub>6</sub>	M6PFDA
8:2 fluorotelomer sulfonate, <sup>13</sup> C <sub>2</sub>	M2-8:2-FTS
Perfluoroundecanoic acid, <sup>13</sup> C <sub>7</sub>	M7PFUdA
Hexafluoropropylene oxide, <sup>13</sup> C <sub>3</sub>	M3HFPO-DA
Perfluorododecanoic acid, <sup>13</sup> C <sub>2</sub>	MPFDoA

**Table S2.** Breathing parameters used in MPPD model (version 2.11, Applied Research Associates, Inc.) to estimate deposition fraction as a function of particle size (Anjilvel and Asgharian, 1995, <https://doi.org/10.1006/faat.1995.1144>).

<b>Species</b>	Human
<b>Model</b>	Yeh/Schum Symmetric
<b>FRC</b>	3300 ml
<b>URT Volume</b>	50 ml
<b>Breathing frequency</b>	12 /min
<b>Tidal volume</b>	625 ml
<b>Inspiration Fraction</b>	0.5
<b>Pause fraction</b>	0
<b>Body position</b>	Upright
<b>Breathing Scenario</b>	Nasal

**Table S3.** Regional (head, tracheobronchial region – TB, and Pulmonary/Alveolar Region – Pul) and total deposition efficiencies calculated using MPPD (version 2.11, Applied Research Associates, Inc.) for each nanoMOUDI size bin.

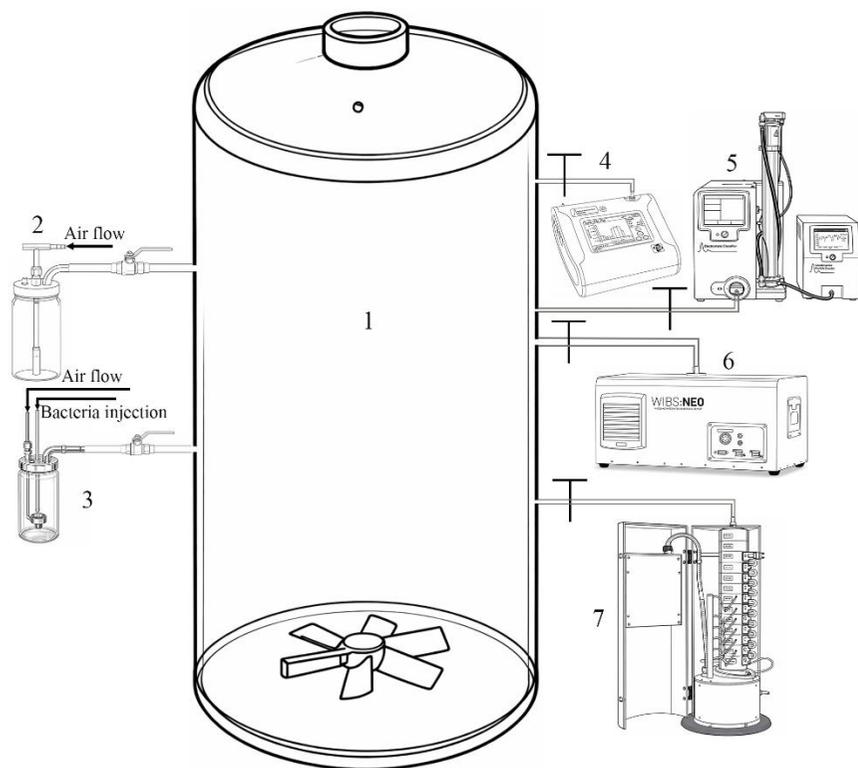
<b>MOUDI stage</b>	13	12	11	10	9	8	7	6	5	4	3	2	1
<b>Particle Size (µm)</b>	0.01–0.018	0.018–0.032	0.032–0.056	0.056–0.1	0.1–0.18	0.18–0.32	0.32–0.56	0.56–1	1–1.8	1.8–3.2	3.2–5.6	5.6–10	10-30
<b>Head</b>	0.1333	0.0825	0.045	0.025	0.0315	0.0235	0.029	0.0537	0.1437	0.2775	0.578	0.886	0.9884
<b>TB</b>	0.406	0.313	0.2105	0.1395	0.092	0.066	0.052	0.0507	0.0555	0.0755	0.0875	0.0605	0.0085
<b>Pul</b>	0.1933	0.2745	0.2855	0.219	0.1415	0.094	0.0755	0.091	0.119	0.1845	0.15	0.037	0.0002
<b>Total</b>	0.7785	0.7066	0.5699	0.4058	0.2825	0.1993	0.179	0.2312	0.3908	0.6557	0.9122	0.9902	0.9998

**Table S4.** Size segregated deposition flux (pg/h) estimates in the human respiratory tract.

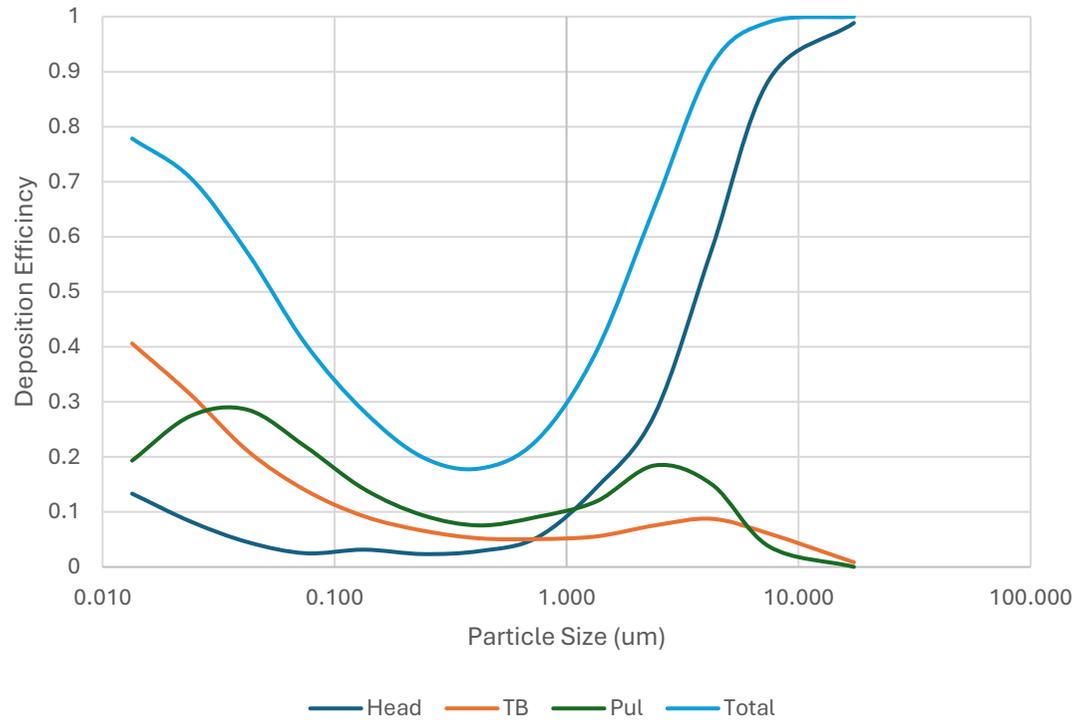
PFAS	Tract region	0.01–0.018	0.018–0.032	0.032–0.056	0.056–0.1	0.1–0.18	0.18–0.32	0.32–0.56	0.56–1	1–1.8	1.8–3.2	3.2–5.6	5.6–10	10–30	Sum	Course (>1.8 um)	Fine (0.1-1.8 um)	UF (<0.1 um)
<b>11 Cl-PF2OUdS</b>	<b>Head</b>	0.000	0.000	0.000	0.009	0.082	0.132	0.373	0.143	0.521	0.081	0.029	0.015	0.000	1.384	0.125	1.251	0.009
	<b>TB</b>	0.000	0.000	0.000	0.049	0.238	0.371	0.669	0.135	0.201	0.022	0.004	0.001	0.000	1.691	0.027	1.615	0.049
	<b>AL</b>	0.000	0.000	0.000	0.077	0.366	0.528	0.972	0.243	0.431	0.054	0.007	0.001	0.000	2.679	0.062	2.540	0.077
	<b>Total</b>	0.000	0.000	0.000	0.142	0.731	1.120	2.304	0.616	1.417	0.190	0.045	0.017	0.000	6.583	0.253	6.189	0.142
<b>L-PFBS</b>	<b>Head</b>	0.000	0.000	0.000	0.000	0.042	0.082	0.262	0.051	0.146	0.000	0.000	0.000	0.000	0.584	0.000	0.584	0.000
	<b>TB</b>	0.000	0.000	0.000	0.000	0.124	0.232	0.469	0.048	0.056	0.000	0.000	0.000	0.000	0.930	0.000	0.930	0.000
	<b>AL</b>	0.000	0.000	0.000	0.000	0.191	0.330	0.681	0.087	0.121	0.000	0.000	0.000	0.000	1.409	0.000	1.409	0.000
	<b>Total</b>	0.000	0.000	0.000	0.000	0.381	0.700	1.615	0.220	0.397	0.000	0.000	0.000	0.000	3.313	0.000	3.313	0.000
<b>82 FTS</b>	<b>Head</b>	0.000	0.000	0.003	0.005	0.064	0.084	0.235	0.076	0.238	0.047	0.103	0.170	0.015	1.039	0.334	0.697	0.008
	<b>TB</b>	0.000	0.000	0.014	0.026	0.188	0.236	0.421	0.071	0.092	0.013	0.016	0.012	0.000	1.088	0.040	1.008	0.040
	<b>AL</b>	0.000	0.000	0.018	0.041	0.288	0.337	0.611	0.128	0.197	0.031	0.027	0.007	0.000	1.687	0.065	1.562	0.060
	<b>Total</b>	0.000	0.000	0.037	0.077	0.576	0.714	1.450	0.325	0.648	0.111	0.162	0.190	0.015	4.304	0.478	3.713	0.113
<b>PFOS</b>	<b>Head</b>	5.236	1.234	0.036	0.040	0.076	0.112	0.401	0.098	0.289	0.068	0.213	0.064	0.000	7.867	0.345	0.976	6.546
	<b>TB</b>	15.947	4.682	0.170	0.223	0.222	0.316	0.718	0.093	0.111	0.019	0.032	0.004	0.000	22.538	0.055	1.460	21.023
	<b>AL</b>	7.593	4.106	0.231	0.351	0.341	0.450	1.043	0.167	0.239	0.045	0.055	0.003	0.000	14.623	0.103	2.239	12.280
	<b>Total</b>	30.579	10.570	0.460	0.650	0.681	0.954	2.473	0.423	0.785	0.161	0.336	0.071	0.000	48.143	0.569	5.315	42.259
<b>L-PFHpS</b>	<b>Head</b>	0.011	0.001	0.007	0.014	0.078	0.106	0.267	0.104	0.284	0.122	0.147	0.191	0.000	1.332	0.460	0.839	0.033
	<b>TB</b>	0.033	0.003	0.034	0.077	0.226	0.297	0.478	0.099	0.110	0.033	0.022	0.013	0.000	1.425	0.069	1.210	0.147
	<b>AL</b>	0.016	0.003	0.046	0.122	0.348	0.423	0.694	0.177	0.236	0.081	0.038	0.008	0.000	2.191	0.127	1.878	0.185
	<b>Total</b>	0.063	0.007	0.091	0.225	0.695	0.897	1.646	0.449	0.774	0.288	0.232	0.214	0.000	5.581	0.734	4.461	0.386
<b>62 FTS</b>	<b>Head</b>	0.071	0.106	0.656	0.213	0.136	0.123	0.738	0.132	0.884	0.235	26.555	3.833	4.687	38.369	35.310	2.013	1.046
	<b>TB</b>	0.216	0.402	3.070	1.190	0.398	0.345	1.324	0.124	0.342	0.064	4.020	0.262	0.040	11.797	4.386	2.533	4.878
	<b>AL</b>	0.103	0.353	4.164	1.868	0.612	0.492	1.922	0.223	0.732	0.156	6.891	0.160	0.001	17.677	7.209	3.981	6.487
	<b>Total</b>	0.414	0.908	8.312	3.461	1.223	1.043	4.556	0.566	2.405	0.555	41.909	4.284	4.741	74.376	51.489	9.793	13.094
<b>PFHxSK(br+Lin)</b>	<b>Head</b>	0.006	0.000	0.006	0.011	0.076	0.103	0.277	0.093	0.272	0.112	0.132	0.184	0.000	1.273	0.428	0.821	0.024
	<b>TB</b>	0.017	0.001	0.030	0.063	0.222	0.289	0.497	0.088	0.105	0.031	0.020	0.013	0.000	1.375	0.063	1.201	0.111
	<b>AL</b>	0.008	0.001	0.040	0.099	0.341	0.412	0.722	0.157	0.226	0.075	0.034	0.008	0.000	2.123	0.117	1.857	0.149

	<b>Total</b>	0.032	0.003	0.081	0.184	0.681	0.874	1.711	0.399	0.741	0.265	0.208	0.205	0.000	5.384	0.678	4.405	0.300
<b>L-PFPeS</b>	<b>Head</b>	0.000	0.001	0.008	0.014	0.085	0.106	0.275	0.103	0.313	0.156	0.163	0.236	0.000	1.459	0.554	0.882	0.023
	<b>TB</b>	0.000	0.002	0.037	0.080	0.249	0.299	0.493	0.097	0.121	0.042	0.025	0.016	0.000	1.461	0.083	1.258	0.119
	<b>AL</b>	0.000	0.002	0.051	0.125	0.383	0.426	0.715	0.175	0.259	0.103	0.042	0.010	0.000	2.290	0.156	1.957	0.178
	<b>Total</b>	0.000	0.005	0.101	0.232	0.764	0.902	1.696	0.443	0.851	0.368	0.257	0.264	0.000	5.883	0.888	4.657	0.338
<b>42 FTS</b>	<b>Head</b>	0.000	0.002	0.007	0.011	0.081	0.119	0.288	0.099	0.298	0.109	0.272	0.079	0.045	1.410	0.505	0.885	0.021
	<b>TB</b>	0.000	0.007	0.035	0.063	0.238	0.334	0.517	0.094	0.115	0.030	0.041	0.005	0.000	1.478	0.077	1.296	0.105
	<b>AL</b>	0.000	0.006	0.047	0.099	0.366	0.475	0.750	0.168	0.246	0.072	0.071	0.003	0.000	2.304	0.146	2.005	0.152
	<b>Total</b>	0.000	0.015	0.094	0.184	0.730	1.007	1.778	0.427	0.809	0.257	0.430	0.088	0.045	5.865	0.820	4.751	0.294
<b>PFBA</b>	<b>Head</b>	0.020	0.318	0.023	0.193	0.377	0.585	1.071	1.086	1.847	1.148	1.692	3.423	4.115	15.897	10.378	4.966	0.553
	<b>TB</b>	0.060	1.205	0.106	1.076	1.102	1.642	1.921	1.025	0.714	0.312	0.256	0.234	0.035	9.688	0.838	6.403	2.447
	<b>AL</b>	0.029	1.057	0.143	1.689	1.695	2.338	2.789	1.840	1.530	0.763	0.439	0.143	0.001	14.456	1.346	10.192	2.918
	<b>Total</b>	0.116	2.721	0.286	3.129	3.383	4.958	6.612	4.675	5.024	2.712	2.670	3.825	4.162	44.275	13.370	24.653	6.252
<b>PFDaA</b>	<b>Head</b>	0.000	0.003	0.000	0.003	0.010	0.010	0.045	0.036	0.111	0.032	0.015	0.073	0.000	0.340	0.120	0.213	0.007
	<b>TB</b>	0.000	0.012	0.000	0.019	0.031	0.028	0.081	0.034	0.043	0.009	0.002	0.005	0.000	0.264	0.016	0.217	0.031
	<b>AL</b>	0.000	0.011	0.000	0.030	0.047	0.039	0.118	0.062	0.092	0.021	0.004	0.003	0.000	0.427	0.028	0.358	0.041
	<b>Total</b>	0.000	0.027	0.000	0.056	0.094	0.083	0.279	0.157	0.302	0.075	0.023	0.082	0.000	1.180	0.181	0.916	0.083
<b>CI-PF3ONS</b>	<b>Head</b>	0.000	0.000	0.004	0.009	0.079	0.104	0.263	0.104	0.303	0.143	0.186	0.146	0.000	1.342	0.475	0.853	0.014
	<b>TB</b>	0.001	0.000	0.021	0.050	0.230	0.292	0.472	0.098	0.117	0.039	0.028	0.010	0.000	1.358	0.077	1.209	0.072
	<b>AL</b>	0.001	0.000	0.028	0.078	0.354	0.415	0.685	0.176	0.251	0.095	0.048	0.006	0.000	2.138	0.149	1.882	0.107
	<b>Total</b>	0.002	0.000	0.056	0.145	0.706	0.881	1.624	0.448	0.825	0.337	0.294	0.163	0.000	5.482	0.794	4.484	0.203
<b>NaDONA</b>	<b>Head</b>	0.000	0.000	0.000	0.000	0.000	0.001	0.003	0.003	0.000	0.000	0.043	0.008	0.021	0.078	0.071	0.007	0.000
	<b>TB</b>	0.000	0.000	0.000	0.000	0.000	0.004	0.005	0.003	0.000	0.000	0.007	0.001	0.000	0.019	0.007	0.011	0.000
	<b>AL</b>	0.000	0.000	0.000	0.000	0.000	0.005	0.007	0.005	0.000	0.000	0.011	0.000	0.000	0.029	0.011	0.017	0.000
	<b>Total</b>	0.000	0.000	0.000	0.000	0.000	0.011	0.017	0.012	0.000	0.000	0.068	0.009	0.021	0.138	0.097	0.040	0.000
<b>GenX</b>	<b>Head</b>	0.694	1.001	0.299	0.306	1.015	1.016	1.222	1.719	2.025	2.203	3.408	5.052	1.673	21.633	12.337	6.997	2.300
	<b>TB</b>	2.113	3.796	1.400	1.706	2.963	2.854	2.191	1.623	0.782	0.599	0.516	0.345	0.014	20.904	1.475	10.413	9.016
	<b>AL</b>	1.006	3.329	1.899	2.679	4.558	4.064	3.181	2.914	1.677	1.465	0.885	0.211	0.000	27.868	2.561	16.394	8.913
	<b>Total</b>	4.052	8.570	3.791	4.964	9.100	8.617	7.543	7.402	5.507	5.205	5.379	5.647	1.692	77.468	17.923	38.168	21.377

<b>PFESA</b>	<b>Head</b>	0.000	0.000	0.005	0.011	0.074	0.093	0.229	0.090	0.276	0.115	0.116	0.175	0.000	1.184	0.406	0.761	0.017
	<b>TB</b>	0.000	0.000	0.025	0.063	0.215	0.260	0.410	0.085	0.107	0.031	0.018	0.012	0.000	1.226	0.061	1.077	0.088
	<b>AL</b>	0.000	0.000	0.034	0.098	0.331	0.370	0.596	0.153	0.228	0.077	0.030	0.007	0.000	1.925	0.114	1.679	0.133
	<b>Total</b>	0.000	0.000	0.068	0.182	0.660	0.786	1.413	0.389	0.750	0.272	0.183	0.195	0.000	4.899	0.650	3.998	0.251



**Figure S1.** Schematic representation of the experimental setup in the ChAMBRe atmospheric simulation facility. (1) ChAMBRe; (2) three-jet Collision nebuliser used for aerosolisation of the PFAS-containing solution; (3) Sparging Liquid Aerosol Generator (SLAG) used for nebulisation of *Pseudomonas fluorescens*; (4) Optical Particle Sizer (OPS); (5) Scanning Mobility Particle Sizer (SMPS); (6) Waveband Integrated Bioaerosol Sensor (WIBS-NEO); and (7) Nano Micro-Orifice Uniform Deposit Impactor (NanoMOUDI) used for size-resolved aerosol sampling.



**Figure S2.** MPPD modelled deposition efficiency in the head, tracheobronchial region, pulmonary region and total human respiratory tract as a function of particle size. MPPD model parameters are given in Table S2.

### Chamber dilution correction

Dilution due to aerosol sampling was corrected by accounting for the total sampling flow rate relative to the chamber volume, treating sampling as a first-order loss process. The total volume of ChAMBRé is 2200 L. The connected to the chamber instruments flow was:

- NanoMOUDI 10 L min<sup>-1</sup>;
- TSP: 10 L min<sup>-1</sup>;
- OPS: 1 L min<sup>-1</sup>;
- SMPS: 1 L min<sup>-1</sup>;
- WIBS: 0.3 L min<sup>-1</sup>;

In the experiments with only PFAS, the instruments used were MOUDI, TSP, OPS and SMPS. The dilution factor was  $22/2200 \text{ min}^{-1} = 0.01 \text{ min}^{-1}$ .

In the experiments with PFAS + bacteria, the instruments used were MOUDI, TSP, OPS, SMPS and WIBS. The dilution factor was  $22.3/2200 \text{ min}^{-1} = 0.0101 \text{ min}^{-1}$ .