

N° échantillon	Référence pot	Tare (en g)	Poids brut (en g)	Poids net (en g)	Référence secteur	Quantité prélevé composites (g)	Quantité totale (g)	Quantité analyse globale (g)	Quantité analyse PFAS (g)	Référence pot global	Référence pot PFAS	Nom échantillon	Nom échantillon labo
Courcelles													
COU001	4490007163PW	16.3	22.87	6.57	CCOU5	3	21	11	10	C6800142	H0743642	CompCou 5	C5
COU003	0890721912	262.22	268.65	6.43	CCOU5	3							
COU005	0890721911	261.81	281.21	19.4	CCOU5	3							
COU006	0890721910	261.91	306	44.09	CCOU5	3							
COU013	0890721907	261.94	334.03	72.09	CCOU5	3							
COU014	0890721908	262.41	265.91	3.5	CCOU5	3							
COU015	0890721909	262	269.96	7.96	CCOU5	3							
COU009	0890721904	261.86	268.72	6.86	CCOU3	5	15	8	7	C6800133	H0743635	CompCou 3	C3
COU010	0890721905	262.7	275.91	13.21	CCOU3	5							
COU012	0890721906	262.62	299.23	36.61	CCOU3	5							
COU016	0890721903	262.43	444.18	181.75	CCOU3	→ rejeté							
COU002	0890721900	261.95	282.67	20.72	CCOU6	5	15	8	7	C6800136	H0743636	CompCou 6	C6
COU004	0890721899	261.58	275.14	13.56	CCOU6	5							
COU011	0890721897	262.24	268.63	6.39	CCOU6	5							
COU007	0890721902	262.14	274.83	12.69	CCOU8	12.69	12.69	6.69	6	C6800149	H0743637	CompCou 8	C8

COU008	0890721896	261.79	277.72	15.93	TCOU	15.93	15.93	8.93	7	C6800146	H0743638	CompCouT3	C11	
Aubange														
AUB009	0890721895	261.76	361.77	100.01	CAUB2	15	15	8	7	C6800135	H0743643	CompAub 2	A2	
AUB001	0890721898	262.16	263.93	1.77	CAUB3	1.77	1.77	1.77	0	C6800141		CompAub 3	A3	ML
AUB003	0890721901	262.42	487.57	225.15	CAUB5	3	18	10	8	C6800140	H0743639	CompAub 5	A5	
AUB007	0890721852	262.2	308.07	45.87	CAUB5	3								
AUB008	0890721851	262.33	344.66	82.33	CAUB5	3								
AUB011	0890721855	262.33	474.68	212.35	CAUB5	3								
AUB014	0890721854	262.07	374.77	112.7	CAUB5	3								
AUB012	0890721850	262.3	286.66	24.36	CAUB5	3								
AUB002	0890721853	261.88	271.61	9.73	CAUB6	3	11.41	6	5.41	C6800153	H0743640	CompAub 6	A6	
AUB010	0890721858	261.85	262.79	0.94	CAUB6	3								
AUB013	0890721857	262.25	264.66	2.41	CAUB6	2.41								
AUB015	0890721856	261.74	281.78	20.04	CAUB6	3								
AUB016	0890721838	262.15	300.14	37.99	CAUB6	3								
AUB004	0890721844	261.61	315.08	53.47	CAUB7	4	16	9	7	C6800144	H0743641	CompAub 7	A7	
AUB005	0890721849	262.11	617.07	354.96	CAUB7	4								
AUB017	0890721847	262.27	320.11	57.84	CAUB7	4								

AUB006	0890721842	261.98	435.95	173.97	CAUB7	4								
Châtelet														
CHAT017	08890721846	261.81	267.48	5.67	CCHA1	4	16	9	7	C6800089	H0745630	CompCha 1	CT1	
CHAT003	0890721848	261.78	275.65	13.87	CCHA1	4								
CHAT007	0890721845	262.2	271	8.8	CCHA1	4								
CHAT013	0890721843	261.9	268.23	6.33	CCHA1	4								
CHAT004	0890334615	264.92	309.03	44.11	CCHA2	2	18	11	8	C6800096	H0745629	CompCha 2	CT2	
CHAT005	0890334613	265.04	296.2	31.16	CCHA2	2								
CHAT009	0890334614	263.35	306.51	43.16	CCHA2	2								
CHAT026	0890334612	265.16	271.7	6.54	CCHA2	2								
CHAT016	0890334594	265.76	271.75	5.99	CCHA2	2								
CHAT018	0890334586	266.24	408.09	141.85	CCHA2	2								
CHAT020	0890334610	264.01	300.62	36.61	CCHA2	2								
CHAT033	0890334611	264.95	287.53	22.58	CCHA2	2								
CHAT025	0890334601	266.05	432.38	166.33	CCHA2	2								
CHAT019	0890334592	263.93	276.69	12.76	CCHA3	7.5								
CHAT032	0890334583	266.00	329.91	63.91	CCHA3	7.5								
CHAT008	0890334590	266.51	270.24	3.73	CCHA5	→ rejeté								

CHAT014	0890334593	263.34	268.32	4.98	CCHA5	5	15	8	7	C6800088	H0745635	CompCha 5	CT5	
CHAT022	0890334591	265.66	283.35	17.69	CCHA5	5								
CHAT027	0890334609	263.99	276.63	12.64	CCHA5	5								
CHAT021	0890334608	266.35	271.34	4.99	CCHA7	5	5	5	0	C6800086		CompCha 7	CT7	ML
CHAT023	0890334607	264.00	275.36	11.36	CCHA8	7.5	15	8	7	C6800080	H0745634	CompCha 8	CT8	
CHAT029	0890334606	264.17	274.43	10.26	CCHA8	7.5								
CHAT012	0890722265	263.55	276.61	13.06	TCHA2	13	13	7	6	C6800084	H0745633	CompCha T2	CT10	
CHAT006	0890722264	263.29	290.26	26.97	TCHA1	15	15	8	7	C6800260	H0745640	CompCha T1-	CT9-1	
CHAT024	0890722269	263.37	361.46	98.09	TCHA1	15	15	8	7	C6800277	H0745639	CompCha T1-	CT9-2	
CHAT028	0890722268	263.12	278.53	15.41	CCHA6	2	20	12	8	C6800276	H0745638	CompCha 6	CT6	
CHAT015	0890722267	263.25	291.07	27.82	CCHA6	2								
CHAT011	0890722270	263.47	270.54	7.07	CCHA6	2								
CHAT034	0890722271	263.08	270.31	7.23	CCHA6	2								
CHAT031	0890722266	263.33	274.05	10.72	CCHA6	2								
CHAT030	0890722272	263.28	283.66	20.38	CCHA6	2								
CHAT001	0890722255	263.24	270.66	7.42	CCHA6	2								
CHAT050	0890722261	263.39	275.48	12.09	CCHA6	2								
CHAT010	0890722257	263.32	276.76	13.44	CCHA6	2								

CHAT002	089722260	263.38	346.83	83.45	CCHA6	2								
Namur														
NAM002	0890722258	263.37	339.93	76.56	CDU2	5	15	8	7	C6800444	H0743650	CompNam 2	N2	
NAM007	0890722263	263.17	333.37	70.2	CDU2	5								
NAM004	C6800344	265.64	326.52	60.88	CDU2	5								
NAM001	C6800335	265.4	414.33	148.93	CDU4	15	15	8	7	C6800441	H0743645	CompNam 4	N4	
NAM006	0890722256	263.34	495.03	231.69	CDU6	7.5	15	8	7	C6800434	H0743646	CompNam 6	N6	
NAM008	C6800347	265	401.6	136.6	CDU6	7.5								
NAM003	0890722262	263.09	314.66	51.57	CDU8	15	15	8	7	C6800432	H0743644	CompNam 8	N8	
NAM005	0890722259	263.12	307.27	44.15	TDU2	15	15	8	7	C6800429	H0743647	CompNam T2	N10	
Charleroi														
CRL004	C6800334	265.1	284.17	19.07	CDe2	5	15	8	7	C6800448	H0743649	CompChl2	CR2	
CRL006	C6800343	265.42	322.27	56.85	CDe2	5								
CRL007	C6800333	264.83	354.15	89.32	CDe2	5								
CRL003	C6800338	264.73	286.84	22.11	CDe8	15	15	8	7	C6800450	H0743648	CompChl 8	CR8	
CRL008	C6800340	264.75	273.45	8.7	CDe3	7.5	15	8	7	C6800438	H0743654	CompChl 3	CR3	
CRL009	C6800326	264.93	276.65	11.72	CDe3	7.5								

CRL002	C6800323	264.31	325.77	61.46	CDe4	15	15	8	7	C6800447	H0743653	CompChl 4	CR4	
CRL001	C6800329	265.07	364.37	99.3	CDe7	15	15	8	7	C6800132	H0743652	CompChl 7	CR7	
CRL005	C6800341	264.75	305.92	41.17	TDe2	15	15	8	7	C6800143	H0743651	CompChl T2	CR10	
Engis														
ENG012	C6801006	264.95	267.51	2.56	CS2	2.56	2.56	2.56	0	C6800137		CompEng 2	E2	ML
ENG002	C6801004	265.16	273.3	8.14	CS3	3	15	8	7	C6800102	H0745628	CompEng 3	E3	
ENG005	C6801017	264.77	312.79	48.02	CS3	3								
ENG007	C6801012	264.96	271.22	6.26	CS3	3								
ENG010	C6801011	264.53	267.61	3.08	CS3	3								
ENG015	C6801014	264.98	429.36	164.38	CS3	3								
ENG003	C6801010	265.5	355.33	89.83	CS5	3	15	8	7	C6800100	H0745627	CompEng 5	E5	
ENG004	C6801016	265.46	387.13	121.67	CS5	3								
ENG008	C6801015	265.09	320.97	55.88	CS5	3								
ENG009	C6801018	264.88	285.33	20.45	CS5	3								
ENG011	C6801007	265.38	417.26	151.88	CS5	3								

ENG001	C6801008	264.98	270.14	5.16	CS6	5.16	11	6	5	C6800095	H0745626	CompEng 6	E6	
ENG016	C6789659	267.4	702.92	435.52	CS6	6								
ENG006	C6789665	266.93	279.58	12.65	CS7	12.65	12.65	6.65	6	C6800098	H0745625	CompEng 7	E7	
ENG013	C6800337	264.83	285.79	20.96	TS	15	15	8	7	C6800092	H0745632	CompEng T1	E9-1	
ENG014	C6801009	265.26	323.8	58.54	TS	15	15	8	7	C6800087	H0745631	CompEng T	E9-2	
Mons														
MONS-066	C6800206	266.02	284.02	18	C01	7.5	15	8	7	C6800268	H0745141	CompMon 2	M2	
MONS-033	C6800431	264.54	277.96	13.42	C01	7.5								
MONS-058	C6800328	265.37	300.13	34.76	C02	3	15	8	7	C6800271	H0745137	CompMon 3	M3	
MONS-071	C6800212	264.73	307.17	42.44	C02	3								
MONS-072	C6800211	264.48	304.03	39.55	C02	3								
MONS-048	C6800317	264.98	278.27	13.29	C02	3								
MONS-043	C6800330	264.5	332.65	68.15	C02	3								
MONS-023	C6800391	264.72	289.92	25.2	C03	4	16	9	7	C6800273	H0745139	CompMon 4	M4	
MONS-038	C6800421	265.09	366.72	101.63	C03	4								
MONS-054	C6800439	265.08	360.32	95.24	C03	4								
MONS-073	C6789661	266.61	293.92	27.31	C03	4								
MONS-018	C6800437	264.61	468.55	203.94	C04	1	15	8	7	C6800275	H0745637	CompMon 1	M1	

MONS-060	C6800199	265.86	341.45	75.59	C05	1	21	12	9	C6800270	H0745643	CompMon 6	M6
MONS-063	C6769842	256.81	314.69	57.88	C05	1							
MONS-052	C6800327	265	272.4	7.4	C05	1							
MONS-031	C6769841	263.03	274.67	11.64	C05	1							
MONS-055	C6769838	255.77	307.44	51.67	C05	1							
MONS-028	C6769836	256.78	295.15	38.37	C05	1							
MONS-032	C6769835	257.37	290.61	33.24	C05	1							
MONS-035	C6800208	262.42	327.18	64.76	C05	1							
MONS-036	C6769839	255.38	263.74	8.36	C05	1							
MONS-040	C6800209	265.15	285.72	20.57	C05	1							
MONS-046	C6800198	265.9	290	24.1	C05	1							
MONS-042	C6800284	264.41	266.5	2.09	C05	1							
MONS-049	C6800332	264.9	278.53	13.63	C05	1							
MONS-047	C6800309	264.6	269.74	5.14	C05	1							
MONS-009	C6800394	264.81	334.32	69.51	C05	1							
MONS-014	C6800390	265.2	273.98	8.78	C05	1							
MONS-019	C6800377	265.4	277.03	11.63	C05	1							
MONS-027	C6800425	264.75	265.82	1.07	C05	1							

MONS-004	C6800424	265.27	289.26	23.99	C05	1							
MONS-068	C6789669	266.95	341.55	74.6	C05	1							
MONS-062	C6789664	266.92	267.80	0.88	C05	1							
MONS-064	C6800207	265.42	307.36	41.94	C06	1	20	12	8	C6800265	H0745143	CompMon7	M7
MONS-053	C6800331	265.06	277.39	12.33	C06	1							
MONS-034	C6769832	256.59	264.34	7.75	C06	1							
MONS-002	C6769834	256.62	258.57	1.95	C06	1							
MONS-037	C6800205	264.92	267.27	2.35	C06	1							
MONS-075	C6800200	265.65	275.34	9.69	C06	1							
MONS-051	C6800314	264.63	293.65	29.02	C06	1							
MONS-008	C6800379	265.59	269.47	3.88	C06	1							
MONS-005	C6800392	265.41	278.51	13.1	C06	1							
MONS-006	C6800388	264.67	270.95	6.28	C06	1							
MONS-017	C6800387	265.18	276.32	11.14	C06	1							
MONS-057	C6800427	264.74	296.77	32.03	C06	1							
MONS-074	C6800426	265.4	336.02	70.62	C06	1							
MONS-012	C6800428	264.46	280.71	16.25	C06	1							
MONS-015	C6800430	264.82	279.37	14.55	C06	1							

MONS-010	C6800435	265.05	285.32	20.27	C06	1								
MONS-020	C6800442	264.77	331.05	66.28	C06	1								
MONS-065	C6789670	266.7	273.00	6.3	C06	1								
MONS-059	C6789660	267.07	282.19	15.12	C06	1								
MONS-077	C6789671	267.15	321.26	54.11	C06	1								
MONS-070	C6789667	267.05	321.22	54.17	C07	2	22	13	9	C6800264	H0745644	CompMon 8	M8	
MONS-067	C6800204	265.4	656.22	390.82	C07	2								
MONS-056	C6800319	264.41	282.95	18.54	C07	2								
MONS-025	C6769843	255.87	285.83	29.96	C07	2								
MONS-041	C6789662	262.93	331.44	68.51	C07	2								
MONS-024	C6769837	257.45	292.08	34.63	C07	2								
MONS-050	C6800318	265.42	307.21	41.79	C07	2								
MONS-013	C6800385	265.5	306.38	40.88	C07	2								
MONS-016	C6800386	264.47	280.03	15.56	C07	2								
MONS-029	C6800383	264.49	271.75	7.26	C07	2								
MONS-011	C6800452	264.73	269.53	4.8	C07	2								
MONS-044	C6800324	265.22	294.65	29.43	C08	4								
MONS-003	C6800393	265.23	336.15	70.92	C08	4								

MONS-007	C6800389	264.88	348.3	83.42	C08	4	10	9	7	C6800272	H0745144	CompMonT 3	M9	
MONS-030	C6800417	264.58	396.59	132.01	C08	4								
MONS-079	C6800210	265	342.64	77.64	T01	7.5	15	8	7	C6800274	H0745142	CompMonT 2	M10	
MONS-039	C6800422	264.41	301.62	37.21	T01	7.5								
MONS-001	C6800446	264.62	366.21	101.59	T02	3	18	10	8	C6800962	H0745138	CompMonT 3	M11	
MONS-021	C6769844	263.98	267.38	3.4	T02	3								
MONS-045	C6800203	265.91	275.36	9.45	T02	3								
MONS-078	C6789666	266.89	279.98	13.09	T02	3								
MONS-061	C6789658	266.54	285.97	19.43	T02	3								
MONS-076	C6789668	267.27	310.8	43.53	T02	3								
MONS-069	C6769840	256.67	274.88	18.21	T03	7.5	15	8	7	C6800965	H0745642	CompMonT 4	M12	
MONS-026	C6800325	264.53	295.13	30.6	T03	7.5								
MONS-022	C6769833	256.79	258.6	1.81	T04	1.81	1.81	1.81	0	C6800269		CompMonT 1	M9	ML