

Supplementary Materials

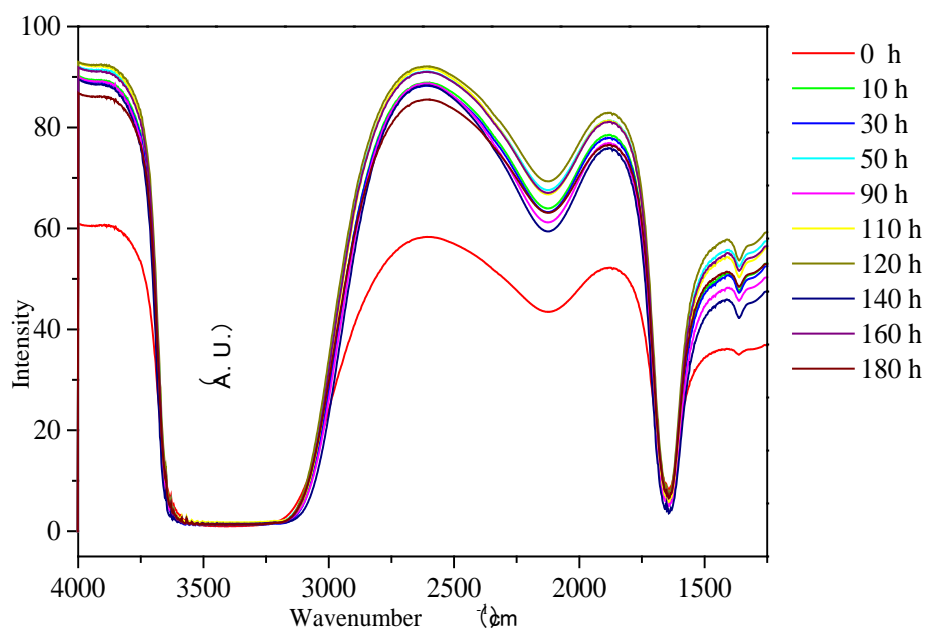


Fig. S1 Time-dependent FTIR spectra during the humification process

Table S1 Assignments of FTIR absorption bands in the humification process

Wavenumber (cm^{-1})	Assignment	Reference
3400	stretching vibration of OH	(Chen et al., 2015; Silva et al., 2020)
2130–2120	stretching vibration of unsaturated groups such as triple bonds and cumulative double bonds	(Abdulla et al., 2010; Li et al., 2014)
1720	C = O stretching of carboxyl and ketonic carbonyls	(Yang et al., 2015; Lee and Hur, 2020)
1660	C = O stretching in conjugation to the aromatic ring	(Amir et al., 2010; Lee and Hur, 2020)
1640	C = O stretching of amide groups in proteins or protein-like compounds, i.e., amide I	(Li et al., 2017)
1606	-COO- stretching of carboxylic acid	(Amir et al., 2010; Li et al., 2014)
1580	C-N stretching and N-H bending vibrations of amide groups in proteins or protein-like compounds, i.e., amide II	(Abdulla et al., 2010; Amir et al., 2010)
1515	Aromatic ring C = C vibrations	(Amir et al., 2010; Li et al., 2014; Yang et al., 2015)
1420	C = N stretching of primary amides, amide III band	(Li et al., 2014; Li et al., 2017; Xu et al., 2018)
1380	OH deformation of phenol-OH and C-O stretching of phenols	(Abdulla et al., 2010)
1350	H deformation of ester C-H bonds (CH_2 and CH_3 groups)	(Chen et al., 2015; Yang et al., 2015)
1190–1180	C-O stretching of phenols	(Amir et al., 2010; Li et al., 2014)

Table S2 DOM fluorescence results for the digestates

Time (h)	Mean component score n ^{a)} (a.u.)						Relative intensity of fluorescence ^{b)}					
	C1	C2	C3	C4	C5	R ^{c)}	Total	C1	C2	C3	C4	C5
0	6.17	10.14	14.78	1.69	2.18	0.76	34.95	0.18	0.29	0.42	0.05	0.06
5	6.42	9.40	14.85	4.97	2.80	0.76	38.44	0.17	0.24	0.39	0.13	0.07
10	6.29	8.42	14.12	5.12	2.91	0.75	36.86	0.17	0.23	0.38	0.14	0.08
20	6.47	8.53	14.23	4.29	3.53	0.73	37.04	0.17	0.23	0.38	0.12	0.10
30	7.09	9.30	14.66	3.45	4.26	0.71	38.77	0.18	0.24	0.38	0.09	0.11
40	6.55	8.76	12.79	3.79	3.96	0.71	35.84	0.18	0.24	0.36	0.11	0.11
50	6.97	9.19	13.64	3.16	4.55	0.69	37.51	0.19	0.25	0.36	0.08	0.12
60	6.60	8.97	11.35	2.61	4.49	0.67	34.03	0.19	0.26	0.33	0.08	0.13
80	6.45	8.65	11.82	2.85	4.60	0.67	34.38	0.19	0.25	0.34	0.08	0.13
100	6.95	8.90	11.50	3.75	4.39	0.68	35.50	0.20	0.25	0.32	0.11	0.12
110	6.94	8.77	12.91	3.85	4.72	0.69	37.18	0.19	0.24	0.35	0.10	0.13
120	6.99	8.83	11.53	4.77	4.84	0.68	36.95	0.19	0.24	0.31	0.13	0.13
125	7.39	8.49	10.47	5.02	4.79	0.66	36.16	0.20	0.23	0.29	0.14	0.13
130	7.08	8.32	9.35	5.66	4.62	0.67	35.03	0.20	0.24	0.27	0.16	0.13
135	7.79	8.95	10.24	4.08	4.94	0.64	36.00	0.22	0.25	0.28	0.11	0.14
140	7.19	8.18	8.04	2.72	4.82	0.61	30.95	0.23	0.26	0.26	0.09	0.16
150	6.21	6.51	4.79	2.79	4.43	0.56	24.73	0.25	0.26	0.19	0.11	0.18
160	6.29	6.99	3.03	3.85	4.74	0.55	24.90	0.25	0.28	0.12	0.15	0.19
170	5.71	5.07	2.99	2.57	4.79	0.5	21.13	0.27	0.24	0.14	0.12	0.23
180	5.64	3.44	1.78	2.16	4.62	0.42	17.65	0.32	0.20	0.10	0.12	0.26

Notes: a) SD-standard deviation. b) Calculated according to Carstea et al. (2014) as a percentage of the total fluorescence. c) Proportion of non-HLAs in the total fluorescence of DOM

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