



Supplementary Materials

Table S1. The characteristics of oilfield wastewater

Parameters	Mean value*
Total soluble salt (mg/L)	12969.3±1058.6
pH	7.8±0.2
Chemical oxygen demand (mg/L)	249.3±45.6
Biochemical oxygen demand	21.1±5.7
Oil content (mg/L)	4.9±1.8
Ammonia nitrogen (mg/L)	5.6±1.9
Total suspended solid (mg/L)	58.5±15.1
Temperature (°C)	47.9±1.6
Chloride ions (mg/L)	2535.2±595.3
Sulfide (mg/L)	2.6±0.7

* The mean value for each parameter was calculated from 90 samples.

Table S2. Alpha diversity indices of fungal community in different biofilm samples

Sample name	OTUs	Richness indices		Good's coverage	Diversity indices	
		ACE	Chao1		Simpson	Shannon
9.1A	17	297.83	291.62	0.995	0.86	4.21
9.1B	22	279.00	279.00	0.994	0.96	5.61
9.1C	22	318.21	312.43	0.992	0.96	5.85
9.2A	27	282.00	282.00	0.994	0.48	2.15
9.2B	22	265.00	265.00	0.993	0.52	2.29
9.2C	11	145.00	145.00	0.994	0.39	1.47
9.3A	30	273.93	268.60	0.994	0.66	2.91
9.3B	14	208.33	203.62	0.994	0.60	2.50
9.3C	33	417.86	401.94	0.993	0.84	4.24
11.1A	44	193.00	193.00	0.997	0.88	4.61
11.1B	45	239.41	235.79	0.996	0.85	4.20
11.1C	48	184.68	181.88	0.998	0.91	5.23
11.2A	22	192.00	192.00	0.996	0.61	2.36
11.2B	14	144.00	144.00	0.995	0.55	1.87
11.2C	19	190.52	189.97	0.995	0.54	1.93
11.3A	27	204.00	204.00	0.993	0.51	2.01
11.3B	12	141.00	141.00	0.995	0.48	1.70
11.3C	17	235.40	228.57	0.993	0.53	2.20

Table S3. Topological properties of fungal networks and Erdős–Rényi random networks

Parameters	Sept. network	Random network	Nov. network	Random network
Node	74	74	100	100
Edge	230	230	602	602
Average degree (AD)	6.216	6.216	12.04	12.04
Clustering coefficient (CC)	0.302	0.085	0.269	0.121
Graph Density (GD)	0.085	0.085	0.122	0.122
Modularity (MD)	0.536	0.304	0.401	0.206
Average path length (APL)	3.388	2.526	2.615	2.081
Network diameter (ND)	9.00	4.79	6.00	3.31

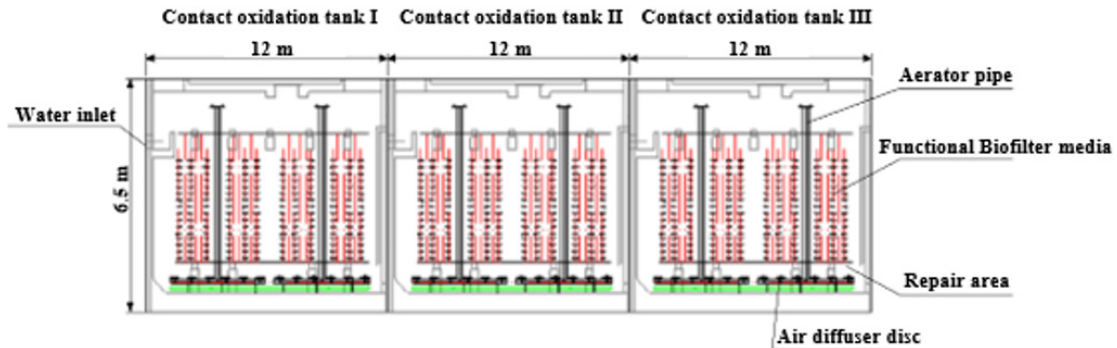


Fig. S1. Schematic diagram of the multistage bio-contact oxidation reactor used in this study

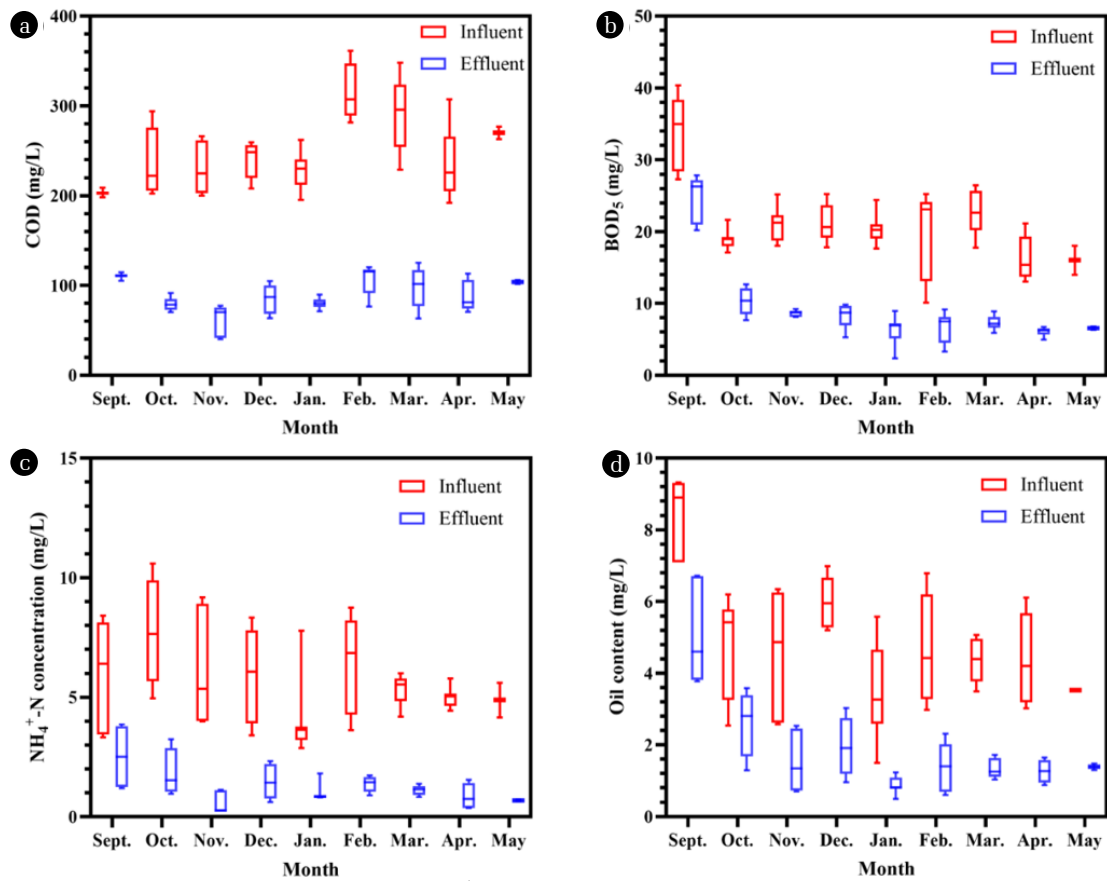


Fig. S2. Monthly average content of COD, BOD₅, NH₄⁺-N, and oil pollutants in influent and effluent over the operation period.

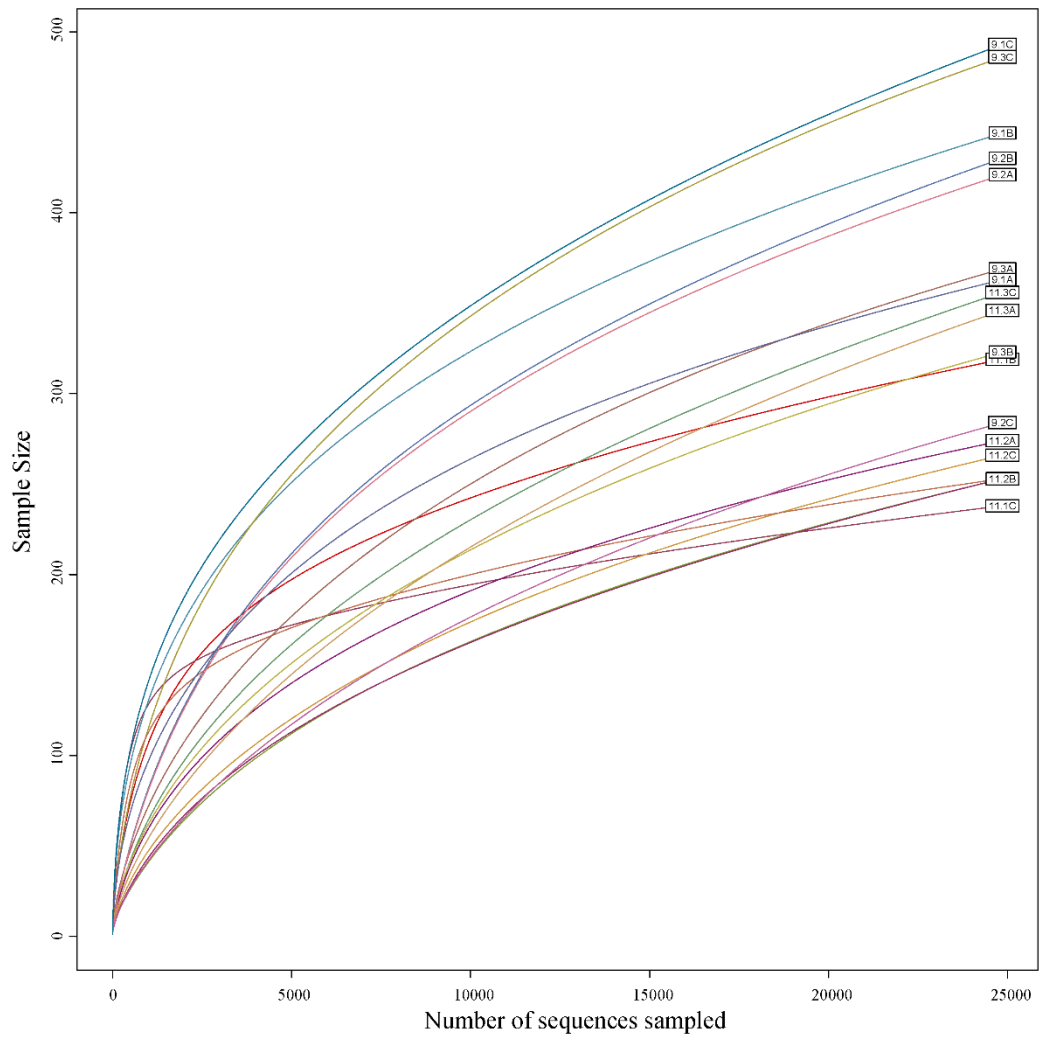


Fig. S3. The rarefaction curves of different biofilm samples

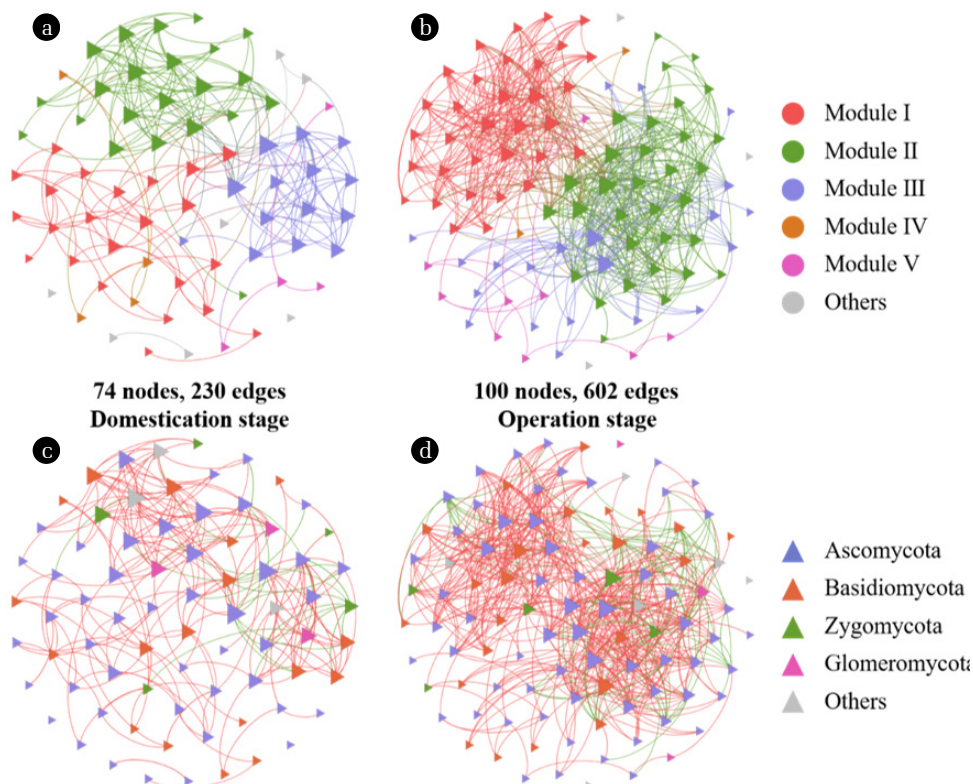


Fig. S4. Co-occurring network of fungal communities in biofilm samples during domestication stage (a, c) and operation stage (b, d) based on correlation analysis. a, b: nodes were colored by modules; c, d: nodes were colored by fungal phyla. A connection stands for a strong (Spearman's $\rho > 0.65$) and significant ($P < 0.05$) correlation. The size of each node is proportional to its degree.

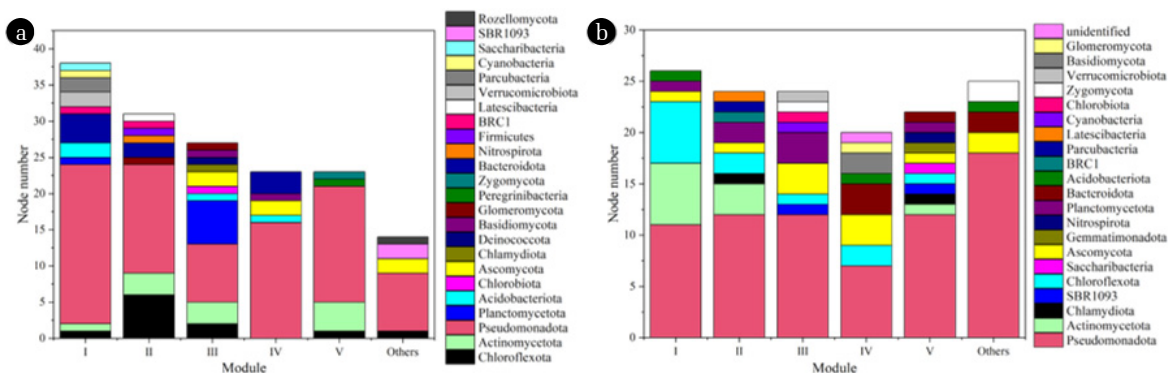


Fig. S5. Number of nodes belonging to different phyla in each module in D-network (a) / O-network (b).