Changes in BNR microbial community in response to different selection pressure

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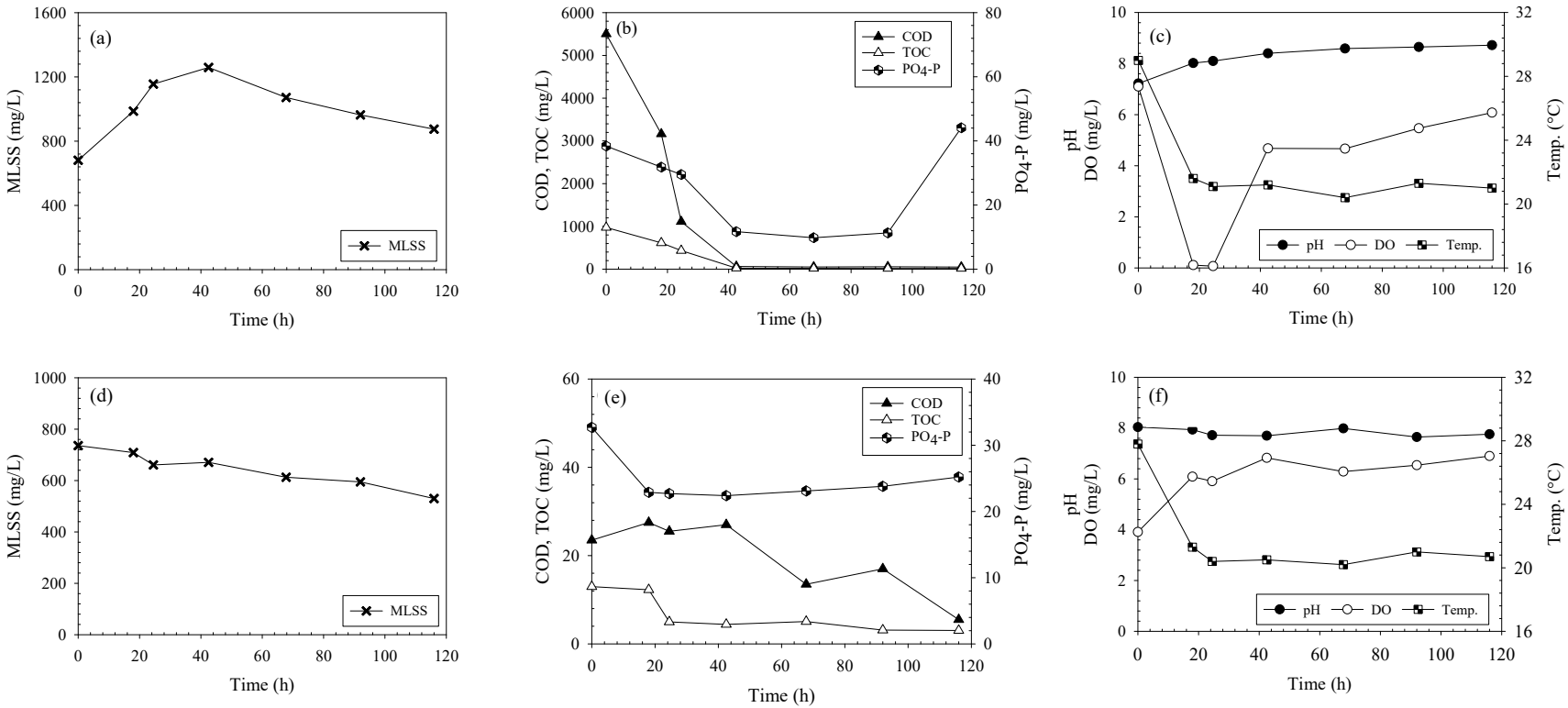

Fig. S1. Concentration profiles of soluble constituents, biomass, and environmental factors in (a – c) heterotrophic nitrification (HN) and (d – f) autotrophic nitrification (AN) suspended-biomass batch bioreactors

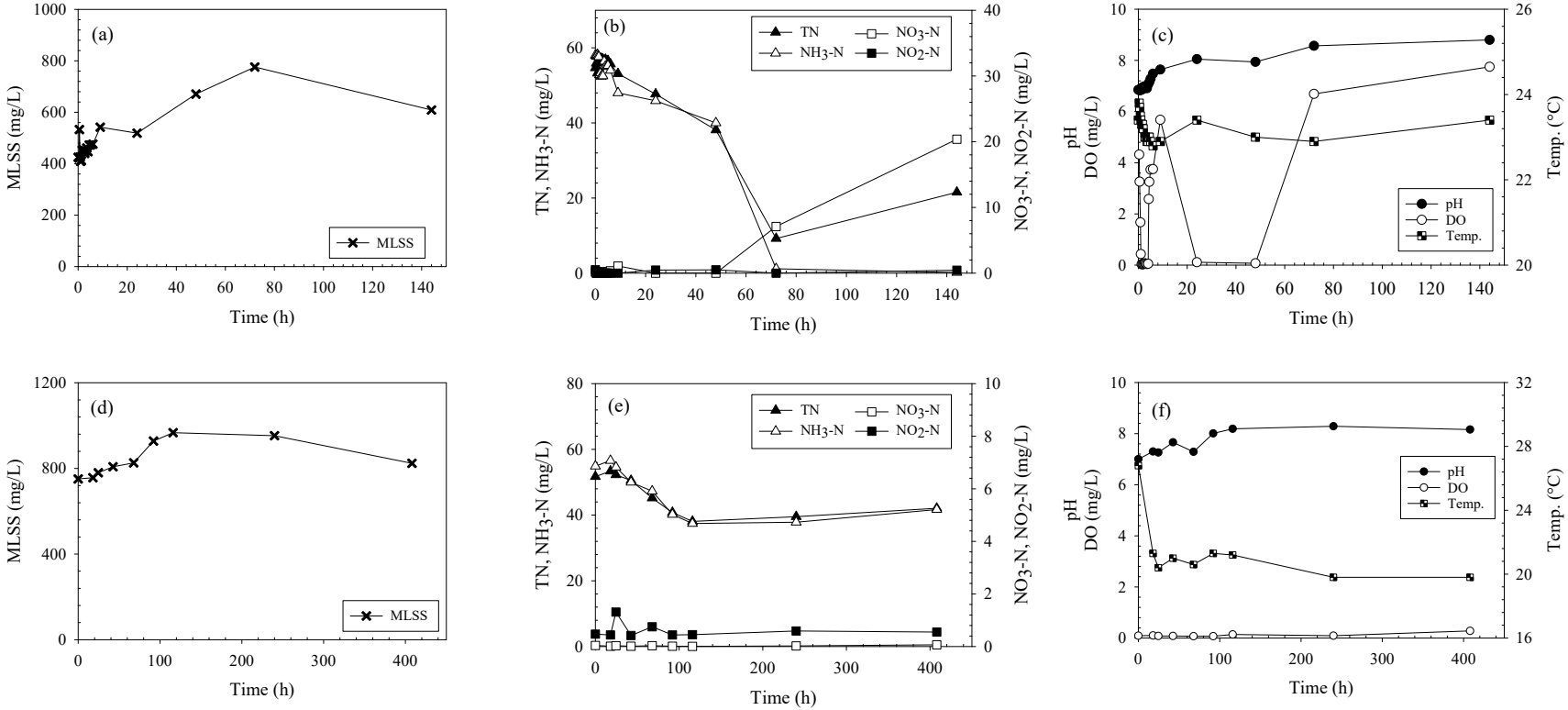

Fig. S2. Concentration profiles of soluble constituents, biomass, and environmental factors in (a – c) alternating anaerobic/aerobic (BioP_AN/O) and (d – f) extended biomass (BioP_ExAN) suspended-biomass batch bioreactors

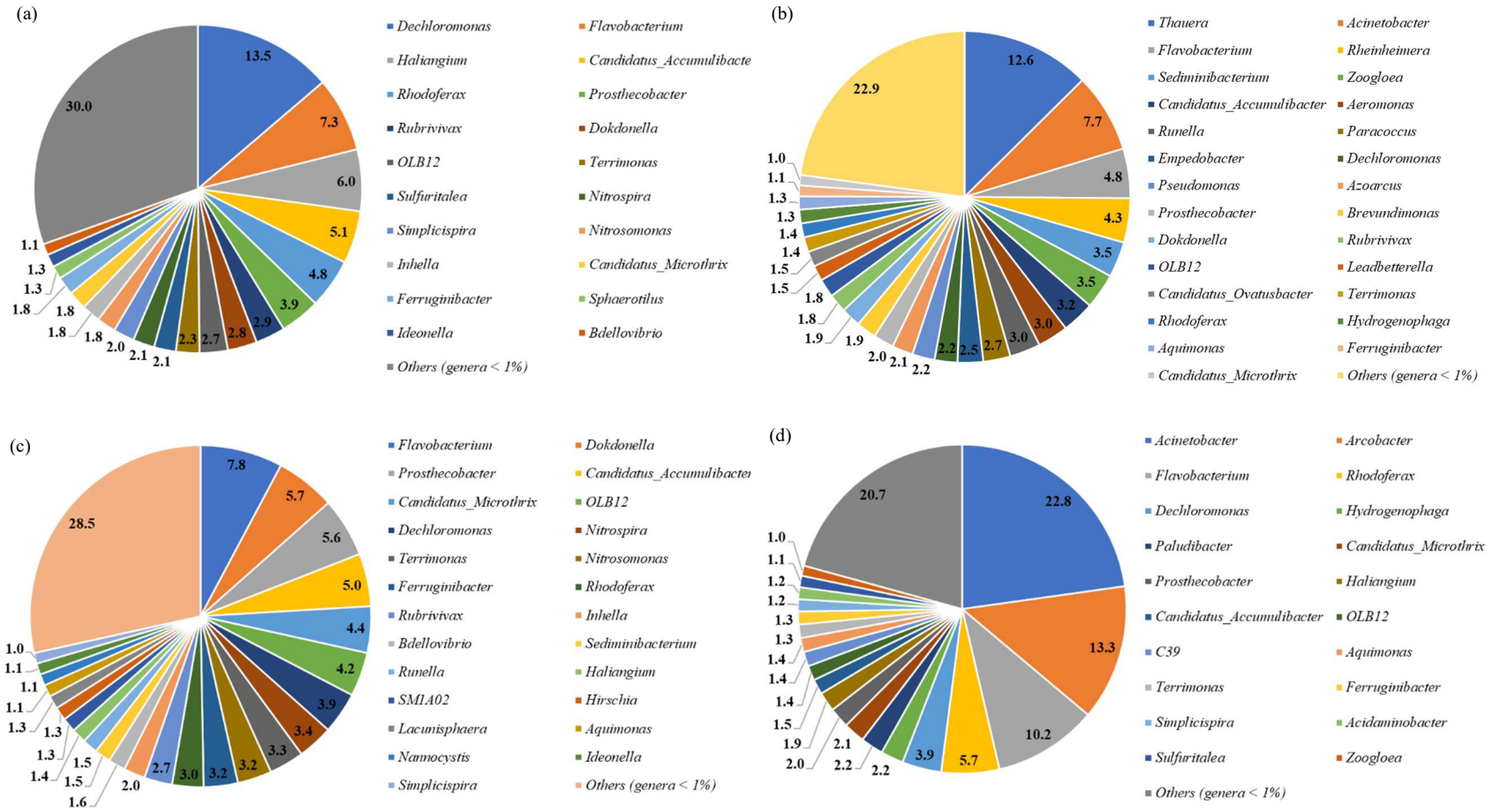

Fig. S3. Bacterial assembly of (a) inoculum; (b) heterotrophic nitrification (HN) AS biomass; (c) autotrophic nitrification (AN) AS biomass; (d) extended anaerobic (BioP_ExAN) AS biomass; (e) alternating (AN/O/AX/O_SBR) granular (AGS) biomass; and (f) purely aerobic (O_SBR) granular (AGS) biomass at genus level (relative abundances ≥ 1%)

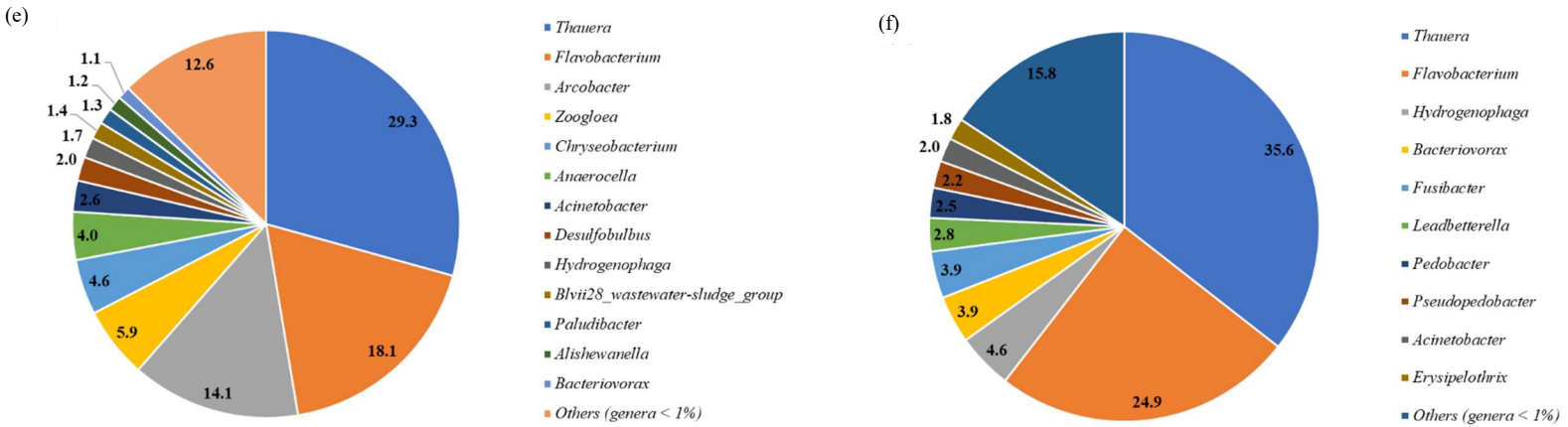
(e)
29.3
18.1
14.1
5.9
4.6
4.0
2.6
2.0
1.7
1.4
1.3
1.2
1.1
12.6
Thauera
Flavobacterium
Arcobacter
Zoogloea
Chryseobacterium
Anaerocella
Acinetobacter
Desulfobulbus
Hydrogenophaga
Blvii28_wastewater-sludge_group
Paludibacter
Alishewanella
Bacteriovorax
Others (genera < 1%)
(f)
35.6
24.9
4.6
3.9
3.9
2.8
2.5
2.2
2.0
1.8
15.8
Thauera
Flavobacterium
Hydrogenophaga
Bacteriovorax
Fusibacter
Leadbetterella
Pedobacter
Pseudopedobacter
Acinetobacter
Erysipelothrix
Others (genera < 1%)

Fig. S3. Cont'd