



Supplementary Materials: The Performance and Microbial Community Identification in Mesophilic and Atmospheric Anaerobic Membrane Bioreactor for Municipal Wastewater Treatment Associated with Different Hydraulic Retention Times

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Figure S1. Rarefaction curves base on Illumina-MiSeq sequencing of bulk sludge and cake sludge bacterial communities in the M-AnMBR and A-AnMBR. The OTUs were defined by 3% distances.



Figure S2. Principal component analysis (PCA) of bacterial communities from bulk sludge and cake sludge in the M-AnMBR and A-AnMBR based on Illumina-MiSeq sequencing of 16S rRNA gene (**a**) and unweighted pair group method with arithmetic mean (UPGMA) tree showing relationships between samples according to bacterial profile in the M-AnMBR and A-AnMBR (**b**). Operational taxonomic units (OTUs) were determined based on 97% similarity of reads.



Figure S3. Venn diagram showing the overlap of OTUs from bulk sludge (**a**) and cake sludge (**b**) in the M-AnMBR (1) and A-AnMBR (2).