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**Evaluation of combined anaerobic  
membrane bioreactor and downflow  
hanging sponge reactor for treatment of  
synthetic textile wastewater**

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- 1. Introduction
- 2. Materials and methods
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- 4. Conclusion

CRediT authorship contribution statement

Declaration of Competing Interest

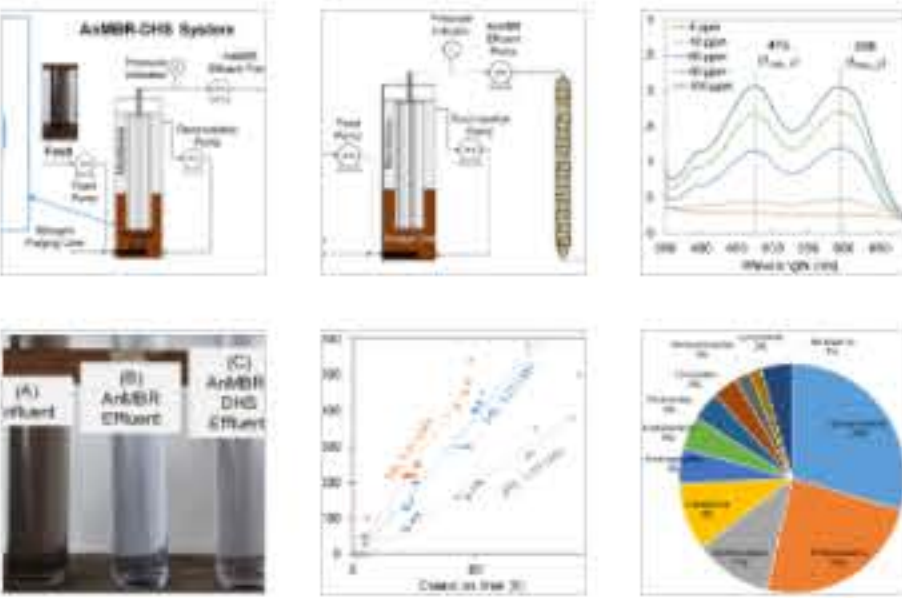
Acknowledgment

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# Evaluation of combined anaerobic membrane bioreactor and downflow hanging sponge reactor for treatment of synthetic textile wastewater

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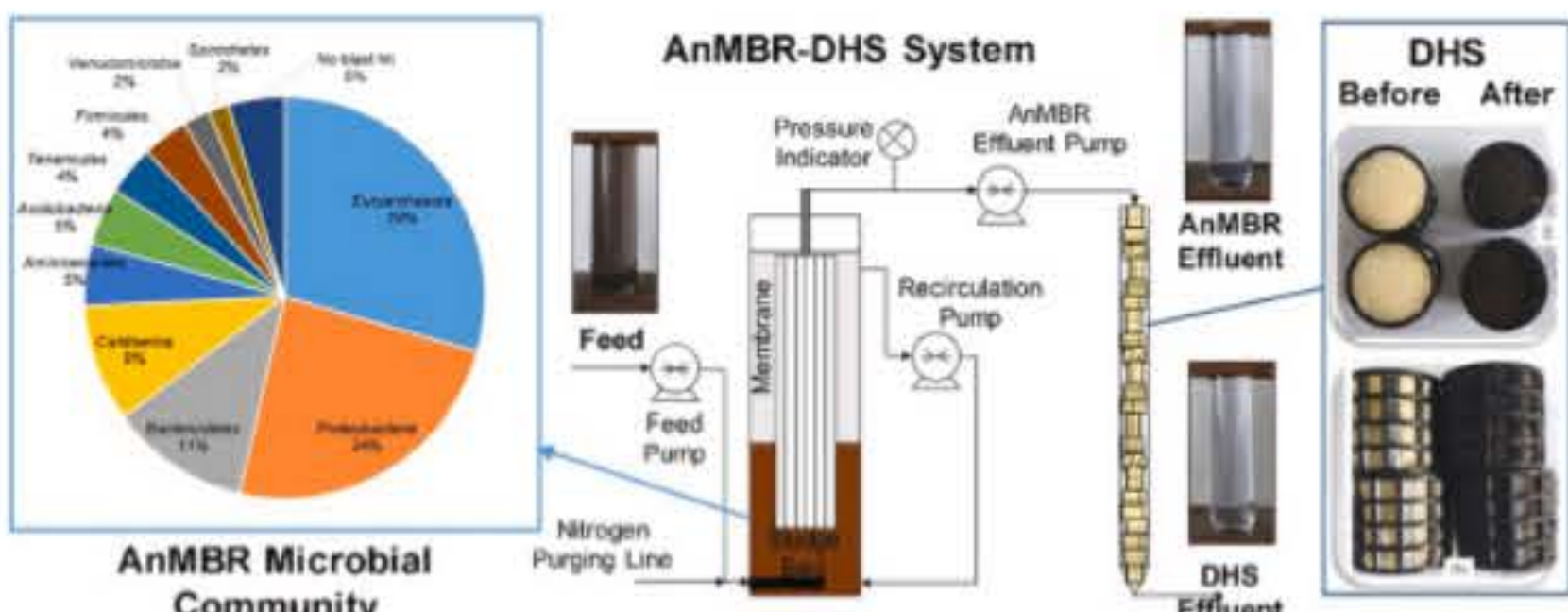
## Highlights

- The combination of AnMBR-DHS significantly reduced the COD, BOD, and color with the most reduction occurred in the AnMBR.
- The HRT and membrane flux did not significantly affect the AnMBR-DHS performances.
- Analysis of microbial in the AnMBR and DHS help to determine the potential microorganism groups for the degradation dye.

## Abstract

Various toxic chemicals in textile wastewater can cause serious problems for the ecosystem and human health if it is discharged without proper treatment. In this study, the performance of textile wastewater treatment using an anaerobic membrane bioreactor (AnMBR) combined with a downflow hanging sponge (DHS) reactor is evaluated. An AnMBR with a working volume of 5 L and a third-generation DHS with a working volume of 0.5 L is used to treat synthetic textile wastewater containing Reactive Black 5 azo dye. The experiment is performed under ambient conditions using AnMBR hydraulic retention times (HRTs) of 12 and 24 h, DHS HRTs of 1.4 and 2.8 h, and membrane fluxes of 2.65 and 5.21 LMH. The AnMBR-DHS combination significantly reduces the biochemical oxygen demand, chemical oxygen demand, and color of the synthetic textile wastewater by approximately  $97.3 \pm 1.8\%$ ,  $94.4 \pm 4.8\%$ , and  $95.0 \pm 1.6\%$ , respectively. Most of the reduction occurs in the AnMBR. The HRT and membrane flux do not significantly affect the performance of the AnMBR-DHS system. The microbial community in the AnMBR is dominated by the phyla *Euryarchaeota*, *Caldiserica*, and *Proteobacteria*, whereas that in the DHS is dominated by *Proteobacteria*. Some of the genera found in the AnMBR can reportedly reduce azo dyes, whereas some of those found in the DHS can reportedly degrade sulfonated aromatic amines.

## Graphical Abstract



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## Keywords

Anaerobic aerobic process; Biofilm; Microbial community; Reactive Black 5; Textile wastewater

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