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Free Download. Aerobic granular sludge cultivation of
the algal-bacterial mixed cultures derived from
macroalgae-enriched activated sludge. Bacterial species
in mixed cultures with the potential ability of anaerobic
digestion of mixed macroalgae-enriched activated
sludge and the algal species observed in food
manufacturing industries were investigated by 16S rRNA
gene sequencing, Fluorescence in situ hybridization
(FISH) techniques, and functional gene assay. Partial
sequencing of 16S rRNA genes demonstrated that
different bacterial groups were present in the mixed
cultures. Cluster analysis revealed that the mixed
cultures were affiliated to the phyla Proteobacteria,

Bacteroidetes, Actinobacteria, Firmicutes, Chloroflexi, Acidobacteria, Gemmatimonadetes, and Planctomycetes. Microbial composition of bacterial groups based on the FISH technique and functional gene assay showed that Gram-positive, Gram-negative, methanogenic, and sulfate reducing bacterial populations were present. Micro-aerobic and anaerobic methylotrophic bacteria were also present. The microbial composition of the activated sludge mixture was influenced by the different type of macroalgae species added to the activated sludge. The bacterial community structure was modified after the addition of the algal species. The characterization of the activated sludge mixture after the amendment of a macroalgae-enriched activated sludge showed that the biological stability and nutrient removal capacity were increased after the addition of the macroalgae. These findings highlight the potential application of the microbial communities derived from the activated sludge as model or surrogate active bioreactor in drinking water production and biogas production.

JACKSON, MI -- A Jackson man left his dead girlfriend's body at the school yard of his son's elementary school in the weekend's cold snap, police say

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