

TRANSITION OF MICROBIAL COMMUNITIES DURING THE ADAPTION TO ANEROBIC DIGESTION OF CARROT WASTE Sarahi L. Garcia^{a,b}, Kamlesh Jangid^b, William B. Whitman^b and K.C. Das^a

^a Department of Biological and Agricultural Engineering, and ^b Department of Microbiology, University of Georgia, Athens, GA, 30602-4435 USA

Introduction

Although anaerobic digestion has been studied intensively for greater than 50 years, monitoring and controlling the anaerobic digestion from a biological perspective has not been fully possible. In this study, pyrosequencing was applied to study the acclimatization of microbial communities to a specific substrate such as carrot pomace.



Trends during adaption by microbial community



of the enrichments Schematics carried out in

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Generations

- Bacteria. Rumen derived enrichments
- -Bacteria. Mixed inoculum derived enrichments
- Archaea. Rumen derived enrichments

Archaea. Mixed inoculum derived enrichments

Phylogenetic distribution of archaeal community



triplicate. Inoculations for following enrichment were done after methane was detected in the previous.



Thermoplasmatales —Unclassified

Figures show the average distribution of the five major phylogenetic groups for the rumen fluid enrichments (D) and the mixed-inoculum enrichment with low methane (•M).

* Major bacterial groups remaining after enrichment were *Bacilli* (31% -45.3%), Porphyromonadaceae (12.1% - 24.8%) and Spirochaetes (12.5% -18.5%)

Figures show methane production of the triplicates for the enrichments (D for rumen, M for mixture).

enrichment with low methane. * Bars represent initial COD and grey lines represent time at which reinoculation occurred

Conclusions

- Communities adapted to changes in their environment while maintaining methane production as a function.
- Diversity was greatly reduced when acclimatizing to anaerobic digestion.
- Bacterial and archaeal populations exhibit different tendencies: Die out, fluctuate, become diluted and become enriched.
- A reactor failure involves community shifts.

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