

Effects of Climate Region, Reaction Temperature and Feed Composition on Microbial Community and Anaerobic Digestion Performance

Rui Chena, Mariana Murillo^b, Yuan Zhonga, Terry Marsha, Lorena Uribe Loriob, Lidieth Uribe Loriob, Dana Kirka, Wei Liaoa*

a. Anaerobic Digestion Research and Education Center (ADREC),
Michigan State University

b. Center of Microbiology, University of Costa Rica

*: Corresponding author

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- Objective
- Investigation of effects of climate and culture conditions on anaerobic microbial communities
- Pilot-scale digestion in Central America
- Conclusions



Investigate the effects of climate region and culture conditions on anaerobic microbial community to conclude the key factors that influence the digestion performance



1 L lab-scale anaerobic digesters



20 m3 pilot thermophilic digester at
Costa Rica

Lab experiment set-up

Experiment

- HRT: 20 days
- Temp.: 35 and 50°C
- Total solids: 5%
- pH: 7
- Feedstock: chicken litter, dairy manure, and food waste
- Location: Michigan and Costa Rica
- Experiment duration: 90 days



1 L anaerobic bioreactors

Parameters

- Biogas production
- TS reduction
- Carbohydrate reduction
- Microbial communities



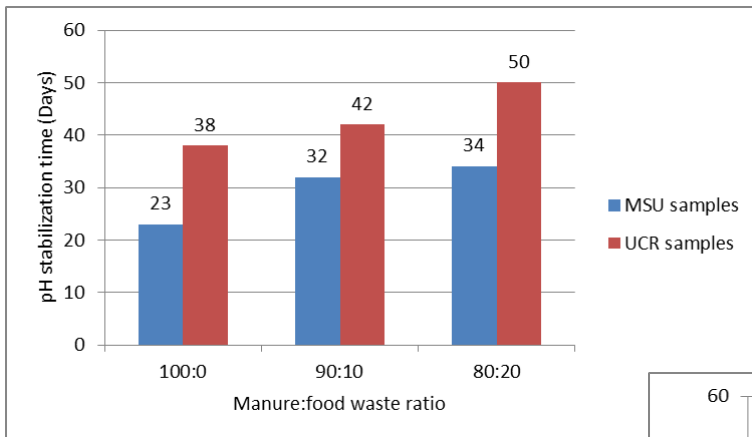
Anaerobic chamber for feeding and sampling

Feedstock characteristics*

		C(wt%)	N (wt%)	C/N	Glucan (wt%)	Xylan (wt%)	Lignin (wt%)
UCR	Dairy manure	40.6 ± 1.0	2.4 ± 0.1	16.7	14.7 ± 0.8	12.6 ± 1.3	27.3 ± 1.5
	Chicken litter	36.8 ± 0.9	3.3 ± 0.2	11.1	25.3 ± 1.0	9.3 ± 0.3	6.8 ± 0.3
	Food waste	44.5 ± 1.1	2.6 ± 0.1	17.1	37.3 ± 0.3	5.2 ± 0.3	16.6 ± 0.7
MSU	Dairy manure	43.7 ± 0.6	2.1 ± 0.2	20.6	22.7 ± 0.7	13.9 ± 0.5	28.4 ± 1.0
	Food waste	47.8 ± 0.1	5.3 ± 0.1	9.0	20.5 ± 2.3	4.7 ± 0.6	11.3 ± 1.1

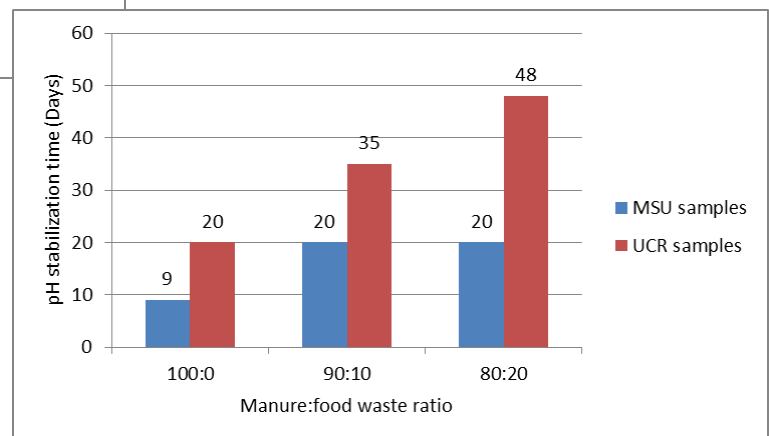
*: Data are average of three replicates with standard deviation

Effects of location, feedstock composition and temperature on digestion stabilization *



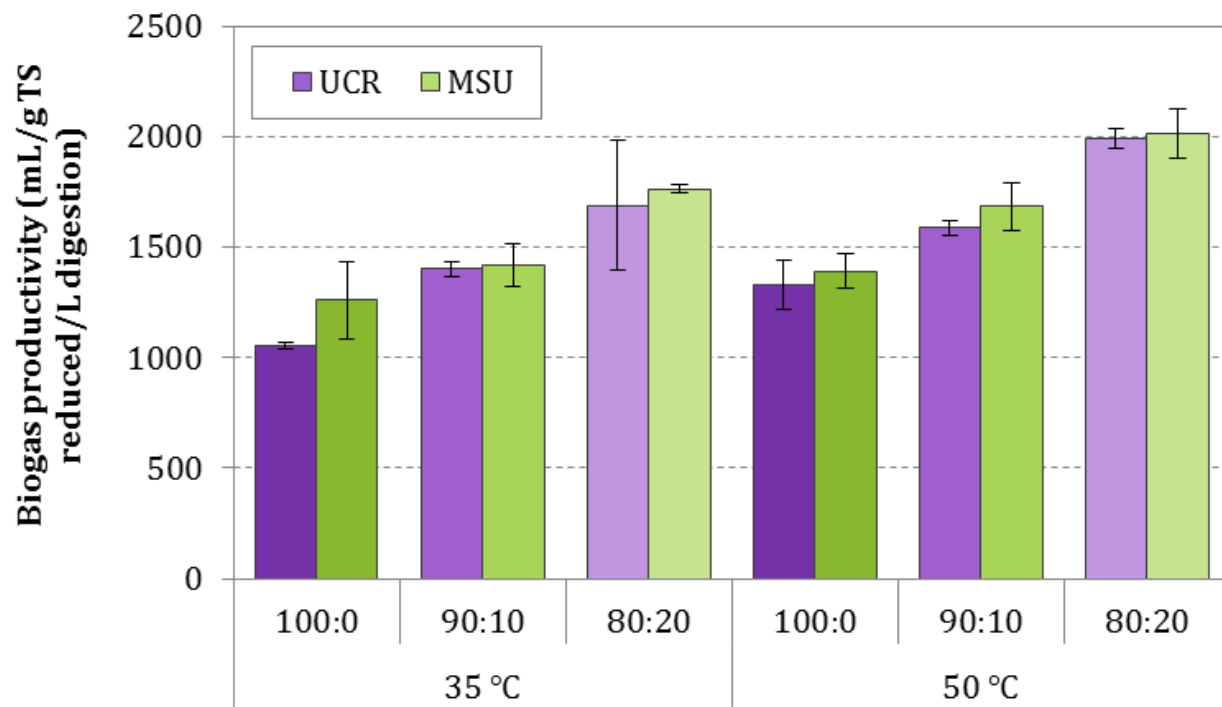
Culture temp.: 35°C

Culture temp.: 50°C

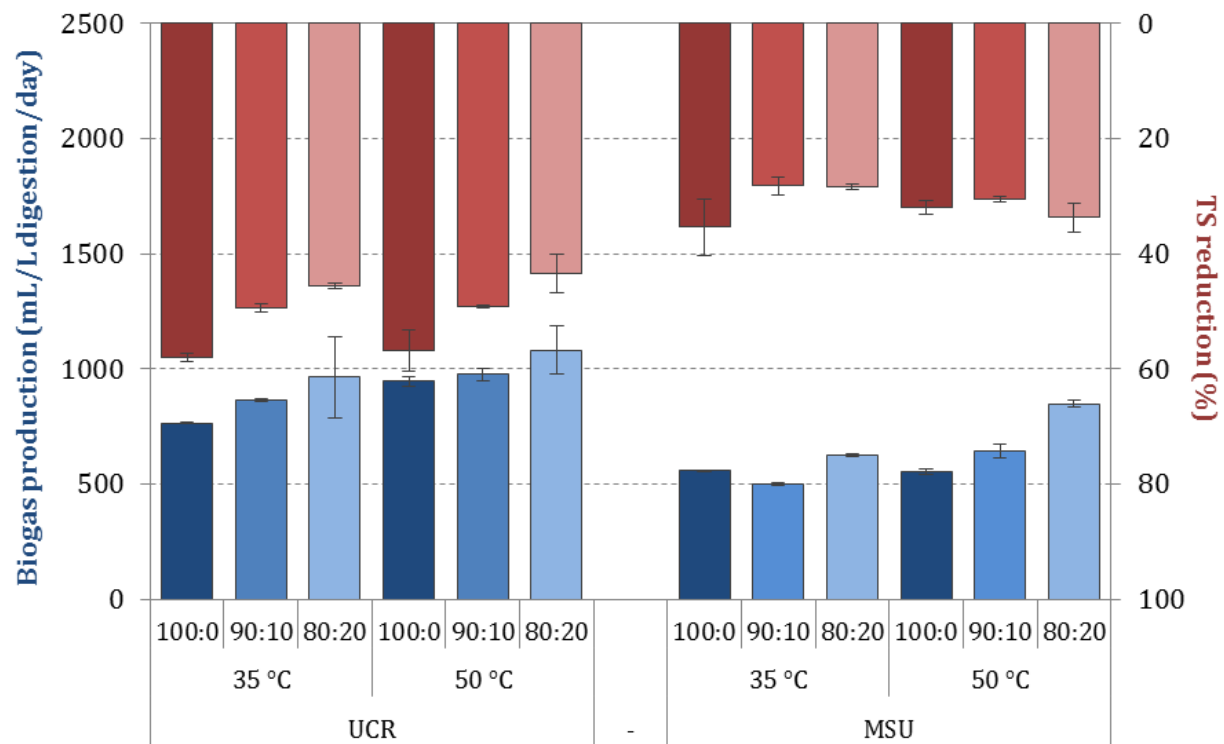


*: Digestion stability is based on pH

Effects of location, feedstock composition and temperature on digestion performance



Effects of location, feedstock composition and temperature on digestion performance



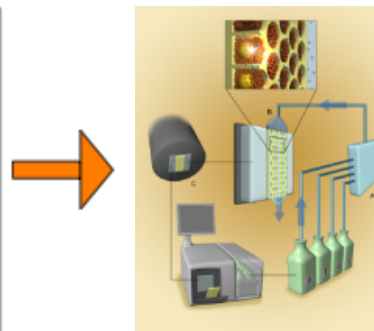
Digestion performance of gas production and fiber reduction*

			Daily biogas production (mL/L digestion/day)	CH ₄ content (%)	TS reduction (%)	Glucan reduction (%)	Xylan reduction (%)
UCR	35 °C	100/0	764.8 ± 1.5	62.0 ± 0.1	58.0 ± 0.8	77.5 ± 1.7	64.7 ± 0.6
		90/10	865.3 ± 5.4	59.4 ± 2.0	49.4 ± 0.8	74.9 ± 1.4	56.0 ± 0.7
		80/20	963.3 ± 177.8	59.1 ± 0.3	45.6 ± 0.5	74.7 ± 2.5	51.9 ± 1.6
	50 °C	100/0	946.1 ± 20.6	59.7 ± 1.2	56.9 ± 3.6	70.0 ± 1.6	59.5 ± 2.4
		90/10	976.3 ± 25.1	67.0 ± 0.4	49.2 ± 0.2	67.4 ± 0.7	51.5 ± 0.5
		80/20	1082.7 ± 107.0	68.2 ± 0.2	43.5 ± 3.3	65.5 ± 0.8	41.1 ± 0.6
MSU	35 °C	100/0	558.9 ± 2.8	58.3 ± 1.0	35.4 ± 5.0	35.6 ± 4.1	33.9 ± 4.8
		90/10	499.1 ± 8.3	65.2 ± 0.5	28.2 ± 1.5	36.8 ± 0.7	25.1 ± 0.5
		80/20	626.5 ± 5.7	60.1 ± 0.3	28.4 ± 0.5	34.3 ± 4.0	17.9 ± 0.3
	50 °C	100/0	554.4 ± 12.4	59.0 ± 1.7	31.9 ± 1.1	44.2 ± 1.1	27.7 ± 0.5
		90/10	642.1 ± 28.7	58.6 ± 1.7	30.5 ± 0.6	44.1 ± 6.0	29.1 ± 6.6
		80/20	848.8 ± 16.2	67.6 ± 0.1	33.7 ± 2.5	40.3 ± 4.9	23.4 ± 3.9

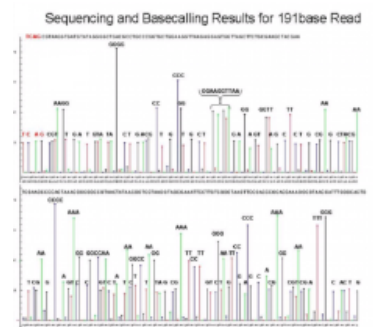
*: Data are average of three replicates with standard deviation

Microbial community in anaerobic digestion

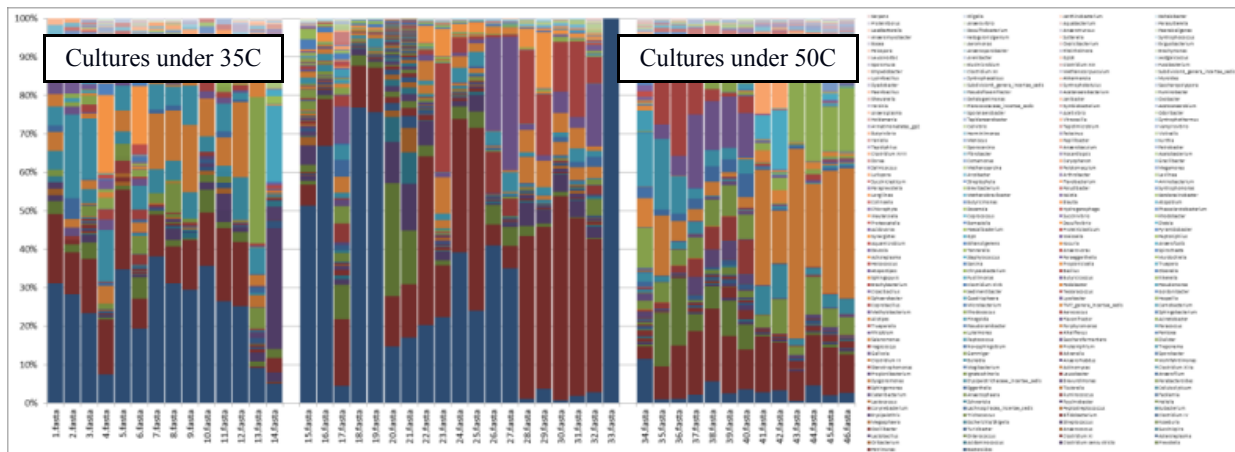
- **Primers for PCR amplification 16S rRNA gene:**
Universal bacterial primers 357f (5'-CCTACGGGAGGCAGCAG-3') and 926r (5'-CCGTCAATTCMTTTRAGT-3')
- **Primers for 454 sequencing:**
Human Microbiome Project (HMP) primers targeting the V3-V5 region of 16S rRNA gene



GT 454 FLX sequencer

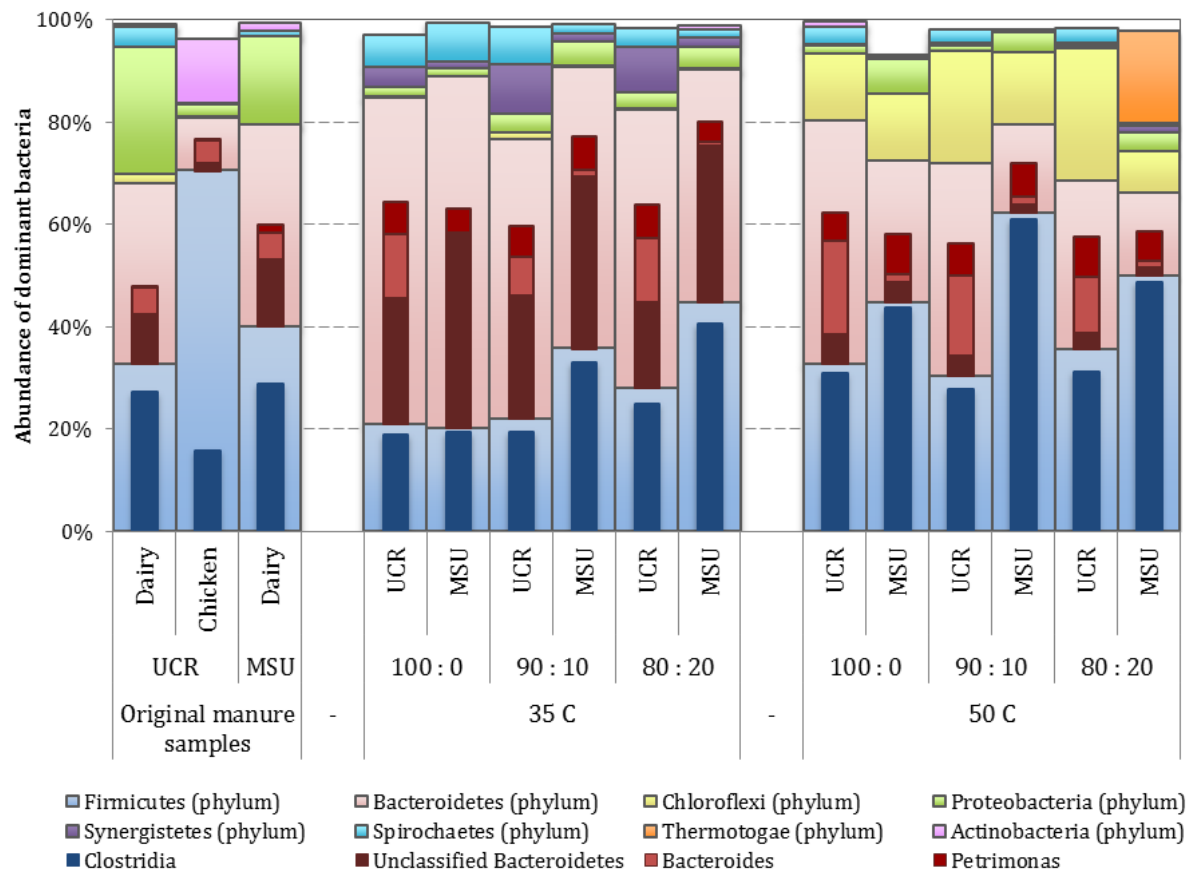


Readings from the sequencer

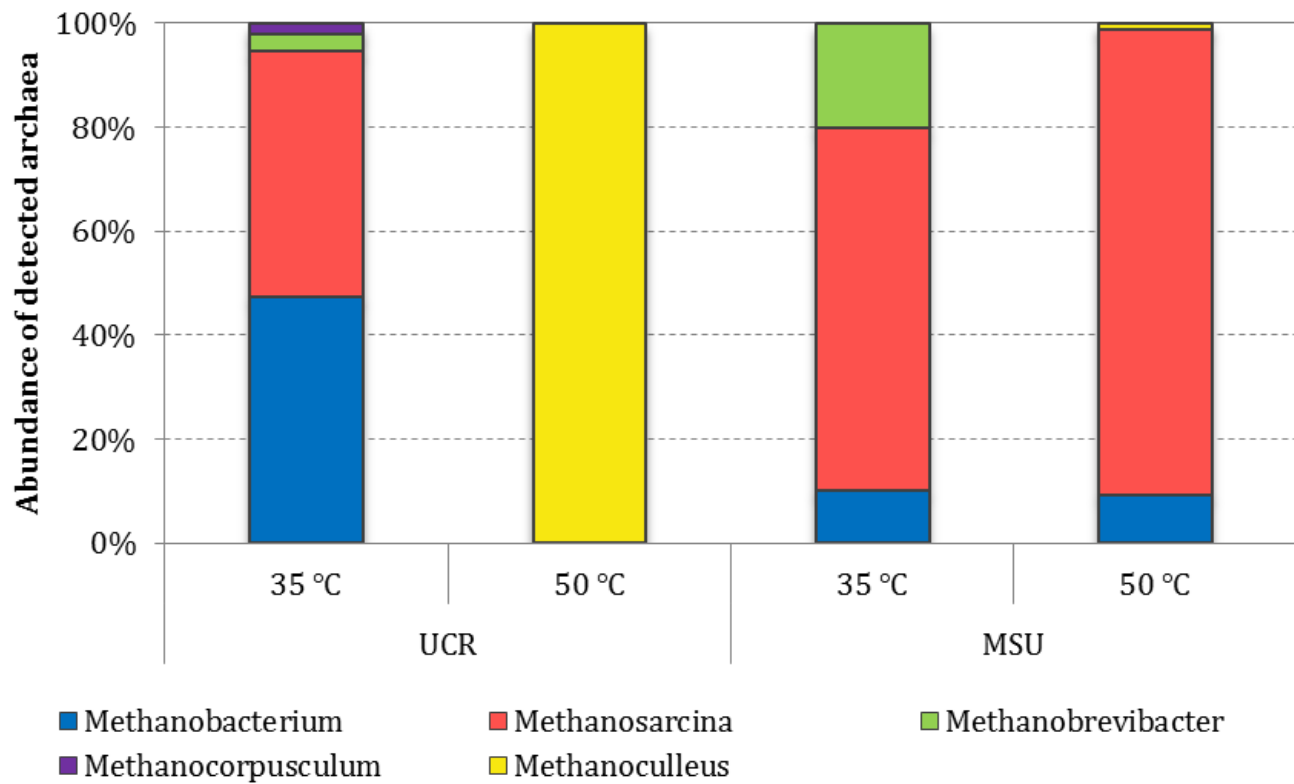


Abundances
of bacterial
community

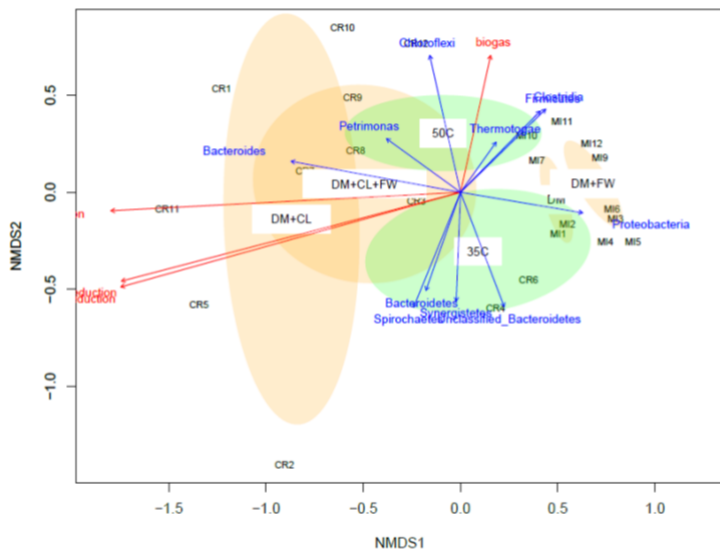
Abundance of dominant bacteria



Abundance of dominant archaea

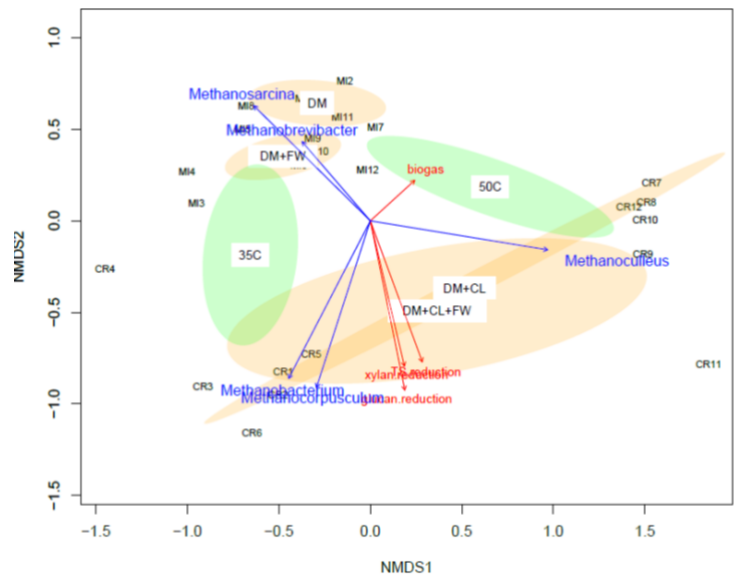


Non-metric Multi-dimensional Scaling (NMDS) analysis of microbial community



Interaction between **Bacteria** and digestion performance

Interaction between **Archaea** and digestion performance



Pilot bioreactor system at the UCR Fabio Agricultural Experiment Station



Feeding unit



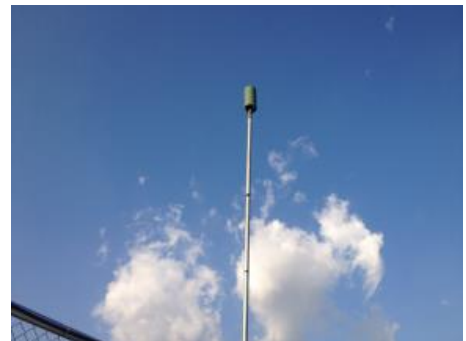
Bioreactor (20 m3)



Engines (16 kw x 2)



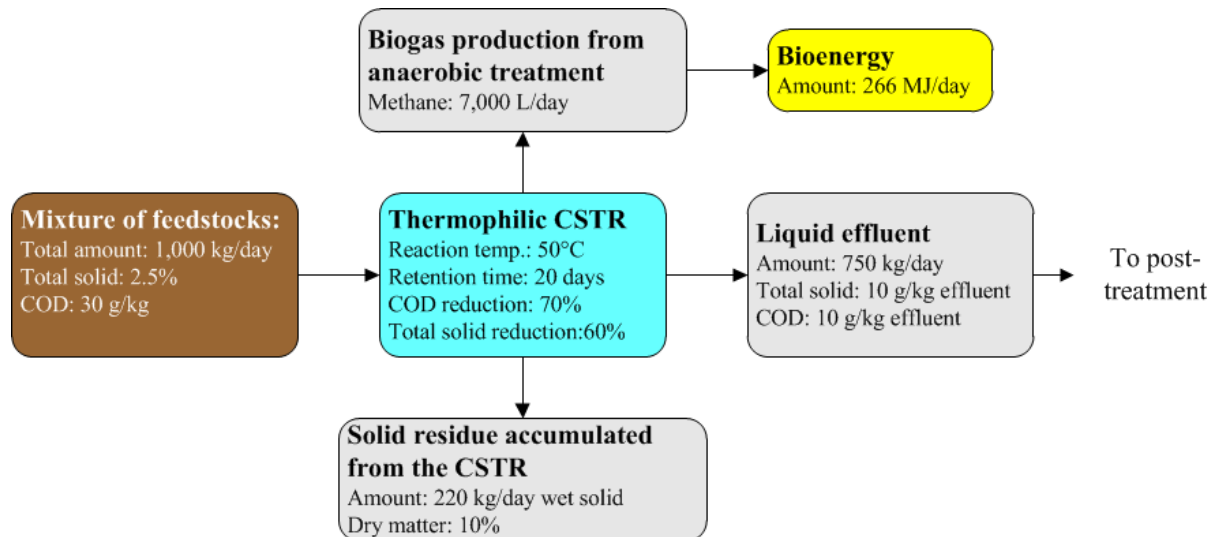
Biogas storage (60 m3)



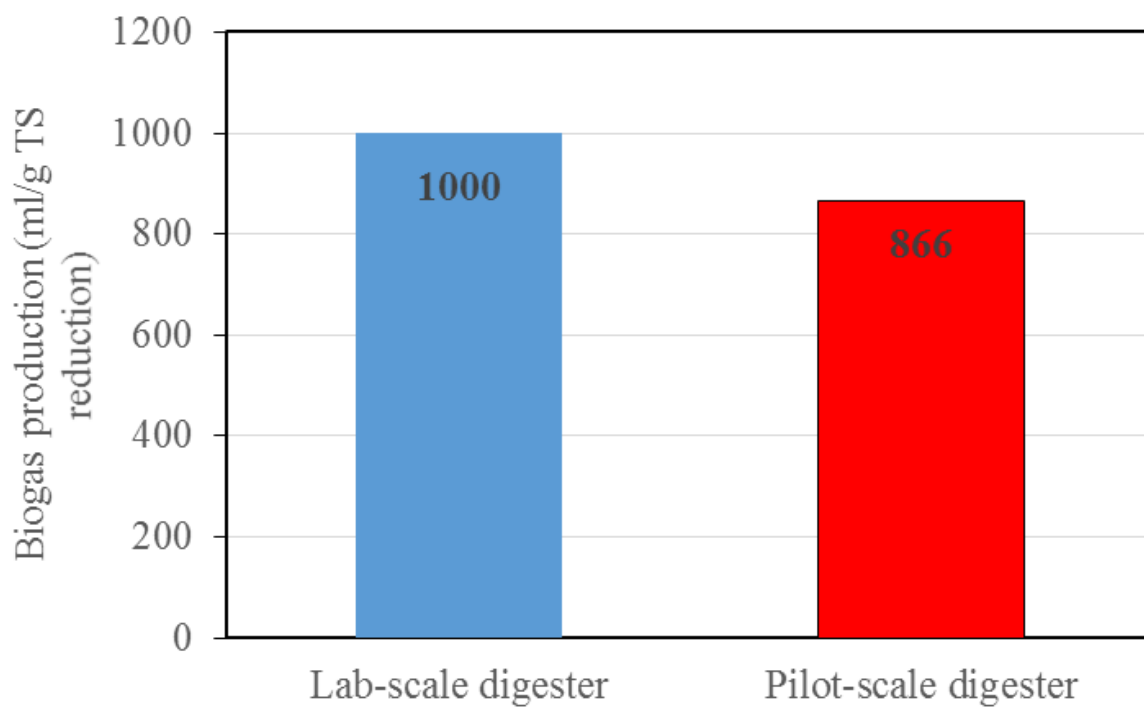
Flare

Mass balance for the pilot scale digester

Mass balance for the solar-bio system on 1,000 kg of mixed chickendairy manure and food wastes



- **Generating 25 kWh electricity per day**
- **Producing 2 gasoline gallon equivalent (GGE) renewable fuel per day**

Performance comparison between lab- and pilot-scale *

*: Under the same operational conditions

1. **Biogas productivity was more dependent on the reaction temperature than climate region.**
2. **The manure-to-food waste ratio in feedstock from the same climate region did not have any significant impact on microbial structure.**
3. Both bacterial and archaeal communities were distinctly different between locations and temperature settings, and they are significantly correlated with biomass reduction rate.
4. The assembly of dominant bacteria (*Bacteroidetes*, *Clostridia*, *Bifidobacterium*) and methanogenic archaea (*Methanobacterium* and *Methanosarcina*) proved that microbial structure shifted corresponding to the change in temperature and climate region.
5. Applying the optimal conditions concluded from the lab study to the pilot-scale digester demonstrated a consistent performance in the tropic region.

Technical Supports

MSU Research Technology Support Facility (MSU RTSF)
UCR Fabio Baudrit Agricultural Experiment Station

Financial Supports

The U.S. Department of State



The MSU Anaerobic Digestion Research and Education Center



Main building



High-bay area



Wet labs



Hot room



CSTR system
(2000 m³, 0.5 MW)



Plug flow system
(1000 m³)



Algal race-way system
(1,600 m² pond)



Solar panels

Homepage: <http://www.egr.msu.edu/bae/adrec/>