Microbial abundance in response to varied waste introduction methods in onsite sanitation systems

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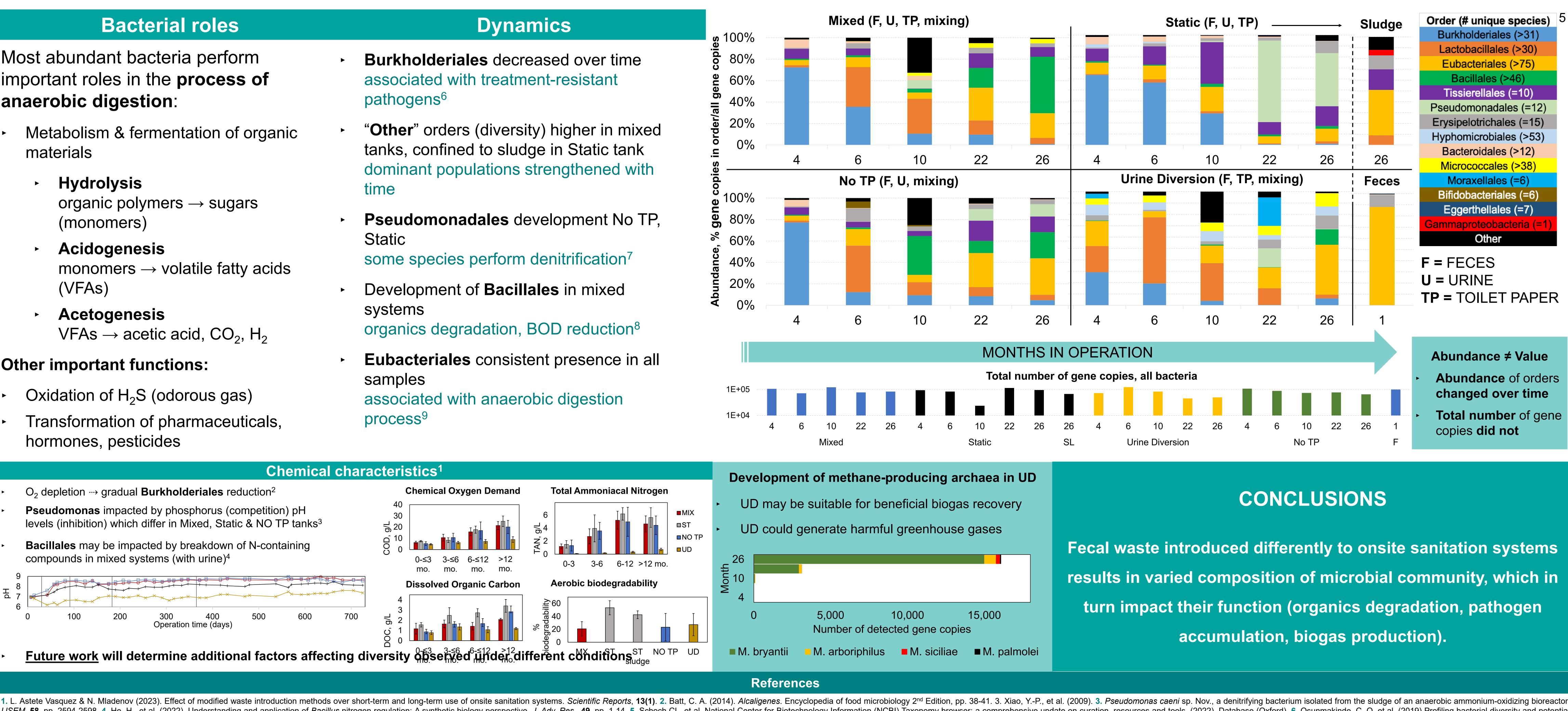
Hypothesis: Bacterial communities responsible for disease, the stabilization of greenhouse gases in onsite sanitation systems change in response to time of use and modifications to how toilets are used.

Background¹

Onsite sanitation systems used by 43% of global population:

- Unhoused encampments (use of 0-2 mo.)
- Refugee camps & disaster relief zones (use of 2-12 mo.)
- Individual households (use of ≥ 12 mo.)

Most abundant bacteria perform important roles in the process of anaerobic digestion:	►	Burkhe associa pathog	
 Metabolism & fermentation of organic materials Hydrolysis 		"Other" tanks, o domina time	
organic polymers → sugars (monomers) ► Acidogenesis	►	Pseud Static	
monomers \rightarrow volatile fatty acids (VFAs)	►	some s Develo	
• Acetogenesis VFAs \rightarrow acetic acid, CO ₂ , H ₂		system organic	
Other important functions:	►		
 Oxidation of H₂S (odorous gas) 		sample associa proces	
 Transformation of pharmaceuticals, hormones, pesticides 			
Chemical characterist			



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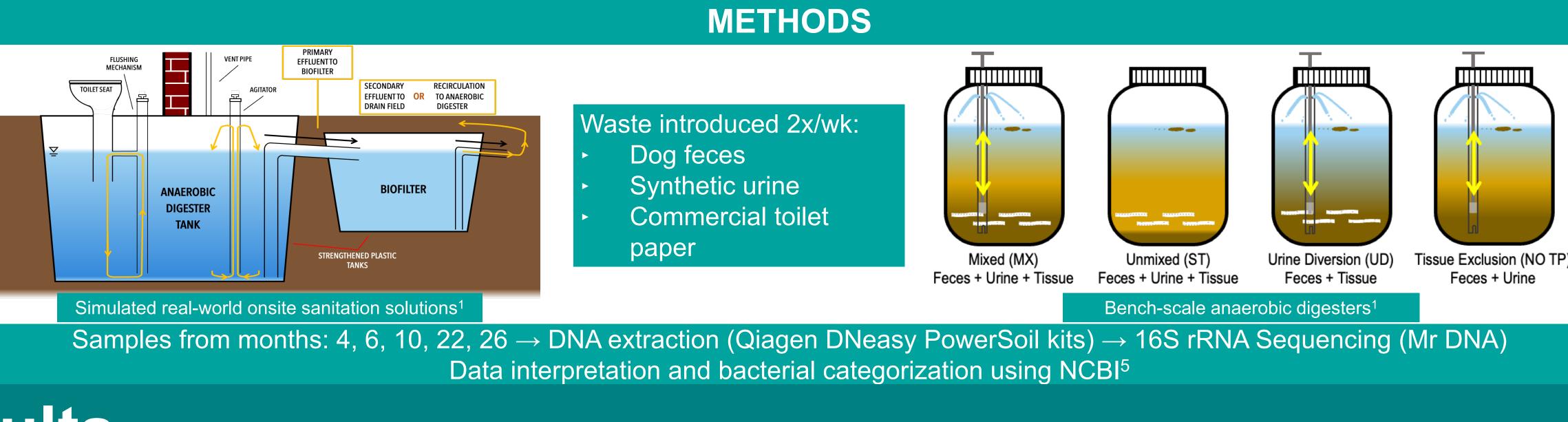




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- **Considers scientifically and culturally relevant** modifications:
- Mixing of fecal waste (increased access to substrate)
- Urine diversion (reduced nutrient concentrations)
- Exclusion of toilet paper (lower C:N ratios)



Results

