

## Sample of English Editing Level 2

Field: Water management and microbiology Topic: Community dynamics of Methanomicrobials, Methanosarcinales and Methanobacteriales associated with different loading rates

## 1. Introduction

In-During anaerobic digestion, crude organic matters are is converted to their monomers, short chained volatile fatty acids (VFAs) and along with hydrogen/ and carbon dioxide. These substrates can then be converted to methane by through the activity of a complex microbial consortium consisting of hydrolytic/fermentative acidogenic bacteria, acid-oxidizing bacteria and methanogenic archaea (Schink, 1988). Due to their different growth requirement and kinetics, a two-stage anaerobic digestion (AD) process has been proposed for an enhanced performance to improve performance by providing different operational conditions for eachin two reactors; one for acidogens and the other for methanogens (Ghosh et al., 1987; Demirer and Chen, 2005). No matterRegardless of what the types of AD process, however, methanogens play a key role\_in an overallprocess since the final methanogenic step affects the finalstabilization of organic matters (Mladenovska et al., 2003). Methanogens are also of great interest to researchers due to their unique physiological characteristics (e.g., low growth rate, high

susceptibility to external conditions and limited substrate utilization range), which <u>make render</u> the whole process sensitive to environmental changes (Hori et al., 2006; Lee et al., 2009b). Variations of operational conditions such as hydraulic retention time (HRT), substrate composition and concentration <u>would can</u> result in variance variability in composition of trophic levels <u>and</u>; thus influence the entire methanogenic community structure and <u>population\_dynamics</u>. Therefore, the an understanding of the-<u>methanogens</u> in anaerobic processes is an important requirement for effective operation and reliable control of anaerobic reactors.

In a continuous mode of bioreactor operation, microbial growth is affected by HRT or given (the inverse of the flow rate), Tand he different microbial growth stages (i.e., lag, exponential, stationary, and decay-stages) can be correlated with different HRTs (Shuler and Kargi, 2002). The growth stage of methanogens, for example, is likely to be in the decay phase at long HRTs, where the residual substrate concentration is low. When HRT consecutively decreases, growth rates of microorganisms increases and shows <u>aits</u> maximum nearatclose to the washout point (Shuler and Kargi, 2002; Madigan et al., 2003). Consequently, This shows that the diversity of methanogens in the <u>a</u> bioreactor would greatly bewould be greatly affected by HRT change because anaerobic digestion usually involves many different **Comment [JN1]:** CHECK: By 'population', do you mean 'population levels'?

**Comment [JN2]:** CHECK: Please ensure this edit retains your intended meaning: 'the inverse of the flow rate' as an alternate explanation for HRT. methanogenic species with differen<u>t</u>ee growth kinetic properties (Yu et al., 2002; Lee et al., 2009a).

Flow rate also affects the loading rates of substrate and as well as other components in the wastewater to the microbial growthmicroorganisms, and tThis would affect microbial diversity based depending on their positive and/or negative effects of the chemicals on the target microorganisms. Loading increaseIncreasing the concentrations of bio-energetically favoredable substrates, such as acetic acid, for example, would be beneficial for methanogenic growth up to a certain level. However,On the other hand increased loading increase of ammonia may be inhibitory due to its toxicity (Hansen et al., 1998; Fujishima et al., 2000).

**Comment [BK3]:** CHECK: Please check that this phrase is correct. Did you mean '...other components in the wastewater which in turn affects microbial growth'?