Supporting Information for

Kenaf Addition Has Mixed Effects on Process Performance of Sequencing Batch Reactors Treating Municipal Wastewater

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Fig. S1 Relative abundance of control (C) and Kenaf-treated (K) microbiomes. Numbers after C or K indicate days of operation. Numbers indicate days of operation. Each microbiome was sampled and sequenced in replicate (indicated by -1 and -2). Bacterial and archaeal taxa are presented using a 3% relative abundance cutoff. Bars without labels represent less than 10% relative abundance. p_: phylum, c_: class, o_: order, f_: family, g_: genus.



Fig. S2 Relative amounts of *Nitrospira* DNA in kenaf and control reactors. The relative masses were calculated using the percentage of *Nitrospira* sequences to Bacterial sequences, and the total mass of DNA extracted.