Absolute quantification of viable bacteria abundances in food by next-generation sequencing : quantitative NGS of viable microbes

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Co-culture process development using individual substrates

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Detailed analysis of metabolism reveals growth-rate-promoting interactions between Anaerostipes caccae and Bacteroides spp

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Development of consortia bioprocesses for efficient production of probiotics

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Evaluating the effect of different fibers on a simplified gut microbiota

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