

Co-metabolism of mucins and dietary fibres by gut microbiota = Mutsiinide ja kiudainete kometabolism soolestiku mikrobioota poolt

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Comparison and applications of label-free absolute proteome quantification methods on Escherichia coli

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Dynamic single fermenter model for study of survival and growth of probiotics in human upper gastrointestinal tract

Sumeri, Ingrid; Arike, Liisa; Adamberg, Signe; Adamberg, Kaarel; Liidemann, G.; Stekolštšikova, Jelena; Laht, Tiit-Maie; Paalme, Toomas Book of abstracts : Gut Microbiology : Research to Improve Health, Immune Response and Nutrition : Aberdeen, Scotland, 21-23 June 2006 2006 / ? p

A dynamic single fermenter model of human upper gastrointestinal tract for study of probiotics

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Effect of stress pretreatment on survival of probiotic bacteria in gastrointestinal tract simulator

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Identification and relative quantification of proteins in Escherichia coli proteome by “up-front” collision-induced dissociation

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Metabolic changes underlying the higher accumulation of glutathione in saccharomyces cerevisiae mutants

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Metabolic changes underlying the higher accumulation of glutathione in Saccharomyces cerevisiae mutants

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