

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

Raba, Grete 2023 <https://doi.org/10.23658/taltech.11/2023> <https://digikogu.taltech.ee/et/Item/37ec835d-31e6-4779-97c5-f5c65d0dbe48>
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Comparison and applications of label-free absolute proteome quantification methods on Escherichia coli

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Corrigendum to: "Comparison and applications of label-free absolute proteome quantification methods on Escherichia coli" [J Proteomics 75 (17) (2012) 5437-5448]

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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

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Genome-wide investigation of mRNA lifetime determinants in Escherichia coli cells cultured at different growth rates

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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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