

Allele distribution at nine STR loci - D3S1358, vWA, FGA, TH01, TPOX, CSF1PO, D5S818, D13S317 and D7S820 - in an Estonian population sample

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

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

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Microbiome of root vegetables - a source of gluten-degrading bacteria

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