



UPPSALA
UNIVERSITET

Molecular Biotechnology Programme

Uppsala University School of Engineering

UPTEC X 04 035	Date of issue 2004-09	
Author	Åsa Innings	
Title (English)	Bacterial species determination using bioinformatic tools and the Pyrosequencing technology	
Title (Swedish)		
Abstract	<p>Pyrosequencing™ is a technology for real-time DNA sequencing. It can be used in a variety of application including microbial identification. In this study methods for bacterial classification using the Pyrosequencing technology have been evaluated. The rRNA gene <i>rnpB</i> was used as target and members within the <i>Streptococcus</i> genus as model species. Bioinformatic tools were used to evaluate and find discriminatory regions of the target gene as well as developing result evaluation tools.</p> <p>The relatively short sequences that can be obtained by Pyrosequencing were found to be discriminatory enough to separate 27 of 48 streptococci species and subspecies. When a second target region was selected, 42 species could be distinguished. The Pyrosequencing assay designed for streptococci identification was highly accurate and reproducible.</p>	
Keywords	Pyrosequencing, DNA sequencing, bacterial identification, <i>Streptococcus</i> , <i>rnpB</i> , RNase P.	
Supervisor	Margareta Krabbe Biosystems, Biotage AB	
Scientific reviewer	Björn Herrmann Department of Clinical Microbiology, Uppsala University Hospital	
Project name	Sponsors	
Language	Security	
ISSN 1401-2138	Classification	
Supplementary bibliographical information	Pages 37	
Biology Education Centre Box 592 S-75124 Uppsala	Biomedical Center Tel +46 (0)18 4710000	Husargatan 3 Uppsala Fax +46 (0)18 555217