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Author Martin Dahlö		
Title (English) Bioinformatic analysis of RNA data from large-scale sequencing of adenovirus infected human cells		
Title (Swedish)		
Abstract <p>The gene usage in adenovirus infected human cells was compared to the gene usage in uninfected cells. This was done by studying the mRNA levels in the cells. The mRNA was extracted, sequenced and aligned to a human reference genome, and changes in coverage between the samples were found.</p>		
Keywords <p>Differential expression, mRNA, R, Bowtie, adenovirus, Solexa, time series, data analysis, human cells</p>		
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