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Author Bengt Anell		
Title (English) Curation Master - a graphical analysis and annotation tool for biological data		
Title (Swedish)		
Abstract <p>An interactive and graphical software tool has been developed to facilitate the annotation and analysis process of biological data. Three applications, pairwise sequence alignment using BLAST, multiple sequence alignment using ClustalW, and literature search and annotation programs written in Java and Perl were implemented. All applications are accompanied with a graphical interface operable via a web browser. The software is currently under further development at The Arabidopsis Information Resource, Carnegie Institution of Washington, Stanford University.</p>		
Keywords Arabidopsis thaliana, sequence alignment, annotation, data mining, Bioinformatics, genome sequencing, Java, Perl, BLAST, ClustalW, literature search		
Supervisors Dr. Sue Rhee Department of Plant Biology, Carnegie Institution of Washington, Stanford University		
Examiner Docent Björn Andersson Institutionen för Genetik och Patologi, Uppsala Universitet		
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Biology Education Centre Box 592 S-75124 Uppsala	Biomedical Center Tel +46 (0)18 4710000	Husargatan 3 Uppsala Fax +46 (0)18 555217