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Title (English) Analys of biomarker data for multiple sclerosis in a workflow environment		
Title (Swedish)		
Abstract In this project data from patients with multiple sclerosis was analysed for biomarkers using multivariate analysis methods. The analysis was conducted in a workflow environment, InforSense Knowledge Discovery Environment (KDE), to evaluate its performance in comparison to the computational programming language MATLAB.		
Keywords Multiple sclerosis, multivariate analysis, proteomics, biomarker, workflow environment		
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