

Multivariate analysis with the Brede **Toolbox**

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Summary

The Brede Toolbox is a Matlab toolbox that implements some of the standard multivariate analysis algorithms, such as K-means, ICA and NMF. The Matlab functions operate on annotated matrices and plotting and HTML generation functions can use this annotation when reporting analysis results, e.g., such that results of hierarchical NMF can be plotted with a single function. The Brede Toolbox has been used for text mining PubMed abstracts, analysis of Wikipedia citations and for analyzing the content of the neuroinformatics Brede Database. The latter analysis involves the formation of multiple kernel density estimates of brain coordinates and subsequent multivariate analyses of these densities with the results containing 3D visualization in brain space added to the Brede Database homepage.

Matlab structures

Most of the functions in the Brede Toolbox typi-cally operates with structures, and, e.g., the ma-trices are represented in a "mat" structure where rows and columns are annotated either with simple types or with other structures. Example:

type: matrix: :: [586x7752 double] :: {586x1 cell} :: [3x7752 double] :: 'Voxelized experime

Here the rows are structures of functional neuroimaging "experiments".1

Non-negative matr Neuroimaging article NMF component Data entry of brain coordinates 3D visualization Matrix of kernel densities One row Kernel density XML storage

Multivariate analysis

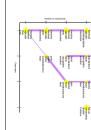
The Matlab structures take care of bookkeep the context information, so that corner cube valization² of results from non-negative matrix torization³ on a stack of kernel density estimation of the structure of the structure

load wobibs
E = brede_bib_bib2exp(B);
E = brede_wexp_exp2mat(E);
M = brede_met_mmf(N);
brede_mat_2mat2html(W,H);

Several multivariate analysis algorithms are immented in the Brede Toolbox: NMF, ICA, SY K-means clustering, canonical correlation. Intain contexts we apply "non-negative partial k squares" which is simply NMF of the inner procof two matrices:

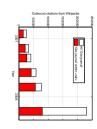
$WH = nmf(X^TY)$

One matrix may be from KDE, the other a bag words matrix of article abstract words. "Hie chical NMF" is simply independent varying si NMFs, which result may be visualized with a c ter bush visualization. Here with a bag-of-wo



Wikipedia citation clustering

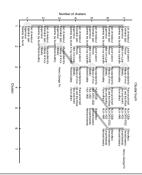
Wikipedia articles may cite scientific articles via the cite journal template. From XML dumps of Wikipedia it is possible to extract citation data from Wikipedia pages where the citations occur and match journal title to variation recorded in an ontology of journals.⁷



With the extracted citations a data matrix built be

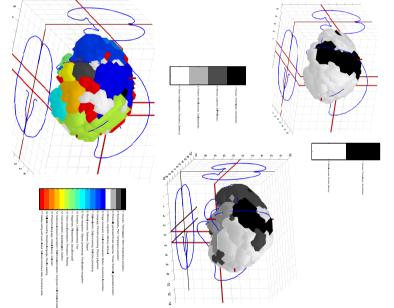
 $\mathbf{X}(\mathsf{Wikipedia\ pages} \times \mathsf{cited\ journals})$

and this matrix can subsequently be subjected to hierarchical NMF and visualized with a *cluster bush* plot as implemented in the Brede Toolbox:^{6,8}



These clusters may display groups of journals and groups of Wikipedia articles, e.g., in neurology.

Multivariate analysis with the Brede Database brain function ontology



Hierarchical non-negative partial least squares with two matrices: One "recursed" adjacency matrix constructed from the brain function ontology, and another matrix constructed from kernel density estimates of the brain coordinates.

Data is available

Data from Web: the Brede Database is available or

Also results from multivariate analysis is avai on that site as well as pages for the individual in the Brede Database, e.g., here part of the for a paper by Epstein et al.: http://neuro.imm.dtu.dk/services/brededatal



Also the extracted data from the tering is available for download the Wikipedia



Thanks to the Lundbeck Foundation for fur and Lars Kai Hansen and Daniela Balslev.



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