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Finding related functional neuroimaging volumes

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Abstract

We describe an information retrieval system for finding related experiments in functional neuroimaging based on the activation volumes. Both sets of locations (e.g., local maxima) as well as summary images can be used.

Introduction

Identification of related research in functional neuroimaging can be done, e.g., by searching in bibliographic databases such as PubMed, browsing "table of contents" of scientific journals or searching BrainMapTM [5] with, e.g., behavioral or location criteria. Here we describe an image retrieval method based on activation information in 3D Talairach space. The information might either come in the form of a list of points representing activation hot spots or it might come as an activation volume, e.g., a volume of t-values from a statistical analysis of a functional neuroimaging data set. Our first goal is to establish a service comparable to "Related Articles" of PubMed.

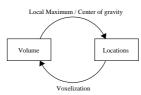
Some image retrieval systems are based on text description of images, but others have combined it with extracted features from the image, e.g., color, texture, shape and keywords. We work entirely with the gray level of the image.

Earlier descriptions of our work are available in [6].

Which space?

The distances between the volumes can be computed in several different spaces

- Voxel space where the distance is aggregated from the distance between the two corresponding voxel values from the two
- · Projected space where the distance is computed in a subspace, e.g., if number of volumes N is smaller than the number of voxels ${\cal P}$ then the volumes can be represented in a small N-dimensional space rather than the large P-dimensional space.
- In location space the distance between volumes is computed as the distance between the hot spots (e.g., local maxima) in the functional volume



Translation from volume/voxel to location representation is usually done by thresholding the volume followed by identification of local maxima (or minima) above the threshold. An other method finds the center of gravities of the connected components in a thresholded volume. Translation from the (set of) locations to the volume/voxel representation is done by voxelization.

Voxelization

For voxelization we convolve every location x with a Gaussian distribution with isotropic covariance – a method associated with kernel density estimation also called Parzen window or Specht kernel estimation [7]. Each location (represented by a 3-dimensional vector \mathbf{x}) is convoluted with an isotropic 3-dimensional Gaussian probability density function (PDF) with a width of σ^2 , presently fixed to $\sigma=10$ mm. The combined PDF is the sum of all the Gaussian PDFs

There are several issues which should be ad

- Each volume may have one or several locations associated with it. We can choose to normalize with the number of locations (so the volumes become a probability volumes).
- A magnitude is often associated with the location. In BrainMap TM this can have different units, e.g., "percent change of rCBF" or "ml/100g/min".
- A sign is sometimes associated with the value of the location. This could indicated a deactivation or a reversed contrast in a t-test. Other method, e.g., an F-test, will not direct produce a sign.

For the present application we normalize with the number of locations, we ignore the magnitude and handle the sign simply by negating the Gaussian PDF for "negative" locations

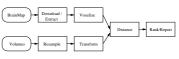
Distance/similarity

We want to rank the volumes according how similar they are with each other. There are multiple ways in which the distance/similarity between the volumes can be measured, e.g.,

- If the volumes is are density volumes we can use information theoretic measures such as the Kullback-Leibler distance
- By a correlation measure such as the inner product or the correlation coefficient [11].



As our primary data we will use the BrainMap™ database [5, http://ric.uthscsa.edu] which we download from the web. Bibliographic and location data is extracted and the location data are feed to the voxelization stage before computation of distances and ranking of related volume for each



We supplements $BrainMap^{TM} \quad with$ the location data from with a few summary images from a functional neuroimaging study with cluster The volumes are resampled and converted from MNI to the Talairach space [2]

Example

HTML web-pages with the resulting lists of related volumes are generated. The unrelated (or rather "antirelated") volumes are also reported. Also added are short bibliographic information, the list of locations and a Corner Cube visualization. An example from one of the 797 experiments presently used is shown below with an experiment from [3].



(95) Shift L in R vis. field A PET shock of visuospan

It shows two clusters of activations: one in the parietal lobe and an other in the frontal lobe. This pattern is repeated for the five most related experiments: Two others from the same study, two from [8] and one from [9].

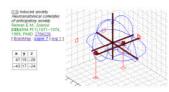
The volume representation requires large amount of data. To reduce the amount of memory used a coarse sampling of 8mm is employed resulting in only 7752 voxels, the volumes × voxels matrix being $\mathbf{X}(797{\times}7752)$. With large data sets it might be an advantage to measure distance in the smaller location space rather than in the large voxel space.

Other processing

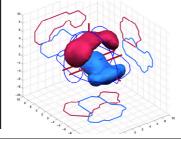
Once the data is in a common space other processing can be made. Indices based on image statistics can give a entry to or overview of the data. An example is the measure of the departure of a volume from the mean volume (across all volumes), — a novelty measure. The upper part of a list with ranked novelties for all volumes is shown below



One of the top entries is $\left[10\right]$ that associates the temporopolar cortex with anxiety. A later experiment found the signal could be extracranial [4].



An other method that can give an overview of the data is singular value decomposition. Isosurfaces in the 2nd eigenimage are shown below with thresholds at 5% and 95% of the number of voxels



Concluding remarks

- · We have devised a method for finding related volumes in voxel and in "location" representation.
- We have applied it on data from primarily BrainMapTM.
- A number of parameters can be optimized e.g., kernel type, width estimation, handling of sign and magnitude information and distance measure.
- Some entries in the BrainMapTM database are meta-analyses and unpublished studies These could be given small weight in the distance measure if they are not wanted.

More information

- Some of the tools this for a na lysis is implemented the Brede in toolbox Matlah available http://hendrix.imm.dtu.dk/software/brede/.
- The generated lists of related volavailable from are presently umes http://hendrix.imm.dtu.dk/services/jerne/.
- Lists of related volumes are also generated for the (anatomical) "volumes of interest" (see poster 490/10497).

Acknowledgment

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