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Introduction

- Databases of task-based fMRI results, such as **Brainmap.org** and **Neurosynth.org**, are commonly used for meta-analytic approaches, including activation likelihood estimation (ALE) [1] or meta-analytic connectivity modelling (MACM) [2].
- Results from such meta-analyses are often used for further research; for instance to define seed regions, volumetric masks, or as points of comparison for new results.
- The online **ANIMA** database (<http://anima.fz-juelich.de>) provides a means for researchers to share, query, and retrieve published meta-analytic results through an intuitive web interface.
- The accompanying stand-alone **Volume Viewer** application allows users to easily organize, visualize, and operate on downloaded ANIMA studies.

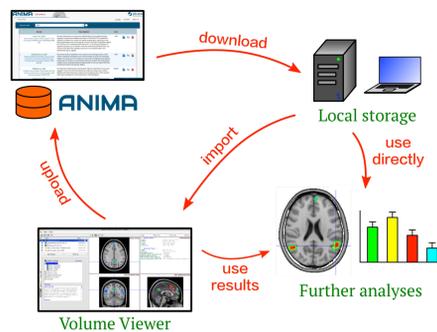
Click this QR code to visit the ANIMA site!



Concept

The ANIMA database was designed according to the following principles:

- Freely available** to the community (no registration required unless you want to submit your data)
- Intuitive interfaces** for retrieving, perusing, and submitting data
- Accessible tools** for organizing, visualizing, and processing retrieved data (Volume Viewer software)
- Rich data specification**, using standardized data formats
- Provenance tracking** (versioning), allowing for proper referencing in journal articles



Querying the Database

The **query page** (above) allows users to query, select, and download studies (24 are presently available). Studies can be filtered by author, journal, publication date, and keywords. Detailed metadata can be perused for each study, and neuroimage elements can be viewed via the Papaya plugin (below).

Volume Viewer

Volume Viewer is an **open source**, Java-based graphical user interface (GUI) for neuroimaging data, based upon the **ModelGUI** API [3]. Volume Viewer was designed to interact seamlessly with ANIMA data, through XML-format study files.

The Volume Viewer interface (above) offers numerous features:

- Organization of ANIMA studies into a **Library** panel
- Fast, intuitive **colour mapping** and **compositing** of multiple images
- Automatic **interpolation** into a common image space
- Availability of common **atlas** and **template** images with name mapping
- Tools** for extracting "blobs", computing peaks and centroids, and performing mathematical operations
- Ability to save entire **sessions** and share as part of an ANIMA study

References

- Eickhoff, S.B., Bzdok, D., Laird, A.R., Kurth, F., Fox, P.T., 2012. Activation likelihood estimation meta-analysis revisited. *Neuroimage* 59, 2349–2361.
- Eickhoff, S.B., Bzdok, D., Laird, A.R., Roski, C., Caspers, S., Zilles, K., Fox, P.T., 2011. Co-activation patterns distinguish cortical modules, their connectivity and functional differentiation. *Neuroimage* 57, 938–949.
- Reid A.T., Evans A.C. ModelGUI: an open-source, community-based approach to neuroscience software. *F1000Posters* 6: 113 (poster)

Submitting Your Data

On the **submit page**, users can register for an ANIMA account (above), and then submit data from their own published meta-analytic studies (below). Studies consist of metadata and individual data elements, which can be neuroimaging volume files, PDF or text documents, images, or Volume Viewer files.

Submitted studies are **reviewed** to ensure quality and suitability, and substantive changes are versioned to facilitate referencing.