

BIDS Statistical Models - An implementation-independent representation of General Linear Models

Christopher Markiewicz, Katherine Bottenhorn, Gang Chen, Alejandro de La Vega, Oscar Esteban, Camille Maumet, Thomas Nichols, Russell Poldrack, Jean-Baptiste Poline, Tal Yarkoni

▶ To cite this version:

Christopher Markiewicz, Katherine Bottenhorn, Gang Chen, Alejandro de La Vega, Oscar Esteban, et al.. BIDS Statistical Models - An implementation-independent representation of General Linear Models. OHBM 2021 - 27th Annual Meeting of the Organization for Human Brain Mapping, Jun 2021, Online, South Korea. pp.1-4. inserm-03478963

HAL Id: inserm-03478963 https://inserm.hal.science/inserm-03478963

Submitted on 14 Dec 2021

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers. L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

BIDS Statistical Models - An implementation-independent representation of General Linear Models

Authors:

<u>Christopher Markiewicz</u>¹, Katherine Bottenhorn², Gang Chen³, Alejandro De La Vega⁴, Oscar Esteban⁵, Camille Maumet⁶, Thomas Nichols⁷, Russell Poldrack¹, Jean-Baptiste Poline⁸, Tal Yarkoni⁹

Institutions:

¹Stanford University, Stanford, CA, ²Florida International University, Miami, FL, ³National Institutes of Health, Bethesda, MD, ⁴University of Texas Austin, Austin, TX, ⁵University Hospital of Lausannne and University of Lausanne, Lausanne, Switzerland, ⁶Inria, Univ Rennes, CNRS, Inserm, Rennes, France, ⁷University of Oxford, Oxford, United Kingdom, ⁸McGill University, Montreal, Quebec, ⁹University of Texas at Austin, Austin, TX

First Author:

Christopher Markiewicz Stanford University Stanford, CA

Co-Author(s):

Katherine Bottenhorn Florida International University Miami, FL Gang Chen National Institutes of Health Bethesda, MD Alejandro De La Vega University of Texas Austin Austin, TX Oscar Esteban University Hospital of Lausannne and University of Lausanne Lausanne, Switzerland Camille Maumet Inria, Univ Rennes, CNRS, Inserm Rennes. France Thomas Nichols University of Oxford

Oxford, United Kingdom <u>Russell Poldrack</u> Stanford University Stanford, CA <u>Jean-Baptiste Poline</u> McGill University Montreal, Quebec <u>Tal Yarkoni</u> University of Texas at Austin Austin, TX

Introduction:

The general linear model (GLM) is a mainstay of neuroimaging analysis, and particularly taskbased functional MRI (fMRI) analysis. As such, there is an independent implementation in all major tool suites. This proliferation provides access to the technique to researchers from varied backgrounds, but each implementation has its own input specification. Consequently, it is nontrivial to compare methods across studies that use different tool suites or to constructing the same model across suites in order to compare the tools themselves.

The Brain Imaging Data Structure (BIDS) is a standard for organizing data from a broad range of neuroscientific experiments [2,3,5,6], including neuroimaging data, experimental events and physiological recordings. Because experiments are often carefully designed to elicit neural responses corresponding to specific proposed cognitive mechanisms, it is useful to include the intended model in the dataset as a means of documenting the experiment and providing a software-independent guide to reproduce the analysis.

Here we present BIDS Stats-Models, a specification for describing how a GLM or similar model should be fit to a BIDS dataset.

Methods:

The BIDS Stats-Models specification was inspired by the OpenFMRI [7] structure, which included descriptions of events, run-level models and group-level models, formatted for easy integration with the FSL FEAT modeling tool [8,9].

Although experimental event descriptions were originally incorporated into BIDS, this did not abstract model descriptions away from the details of any one tool. A BIDS extension proposal (BEP) was initially drafted in September 2016, using a JSON format to declare the stages of the model, and assuming the input data is organized according to BIDS. In October 2018, a workshop of 20 attendees was hosted at Stanford University to assess and refine the specification.

Reference implementations have been used to drive development of the specification. The PyBIDS library [10] began to support components of Stats-Models in June 2017, and an application to run models (FitLins [4]) was begun in October 2017. The initial version of FitLins used the nilearn[1] estimator.

Results:

The BIDS statistical model is encoded as a JSON object describing the inputs to the model, and an array of steps corresponding to levels (run, session, subject or dataset) of analysis (Fig. 1a). Each step contains a model specification and contrast specification (Fig. 1b).

Image inputs to a Stats-Model may be selected via BIDS entities, such as "task", "run", "session" or "subject", if provided. These images, along with events and time series are provided as inputs to the initial (typically run-level) step. Later steps receive as inputs the statistical maps generated as outputs in the previous steps, along with variables defined at that level in the BIDS dataset. For example, if age or experimental group are defined in participants.tsv, these variables are available at the dataset level of analysis.

Within a step, a model can be specified to indicate the dependent variables, variance structure, error distribution and software-specific parameters.

Contrasts are named, weighted sums of parameter estimates, and may be of type "t", "F" or "FEMA" (fixed-effects meta-analysis). The resulting contrast maps are the inputs to the next level; hence, to pass a raw parameter estimate to the next level, a "dummy contrast" that shares a name with its lone input may be used.

Models can be included in a BIDS dataset, in a file according to the template models/model-_[desc-]_smdl.json.



a. A schematic model demonstrating the overall structure of a BIDS Stats-Model, including an optional name and description, inputselector, and steps. The steps are arranged in a "standard" order, fitting each run as a first level model, and combining at the session, subject and finally dataset levels. **b.** An example "Steps" section for a plausible multi-run study. A design matrix with face, place and aCompCor regressors is fit to each run in the study, and a contrast between face and place parameter estimates is calculated. At the subject level, runs are combined by a fixed-effects meta-analysis (FEMA) to estimate each subject's contrast and variance. Finally, a t-contrast is used at the dataset level to estimate the contrast with subject as a random effect.

(https://files.aievolution.com/prd/hbm2101/abstracts/abs_1174/fig1.png)

Conclusions:

Reproducibility and interoperability are core principles of open scientific practice, and common formats for representing data and algorithms are necessary to achieve these principles in practice.

The BIDS Stats-Model aims to provide an abstract, tooling-independent representation of the most common modeling approaches in neuroimaging.