

Reviewing neuroimaging flexibility: Components and records of provenance

Camille Maumet

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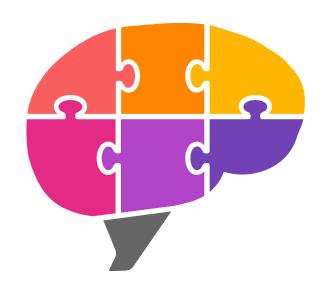
Submitted on 27 Jun 2022

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Symposium What should we do with neuroimaging analytical flexibility?





Reviewing neuroimaging flexibility

Components and records of provenance

Camille Maumet

Inria, Univ Rennes, Inserm, CNRS, IRISA



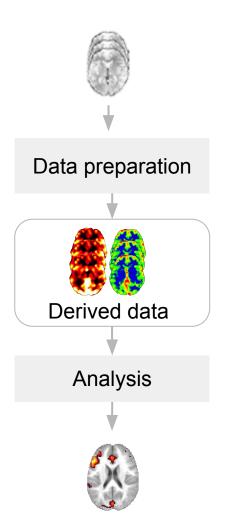


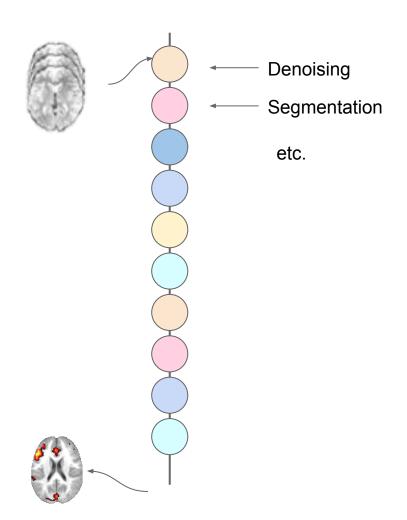


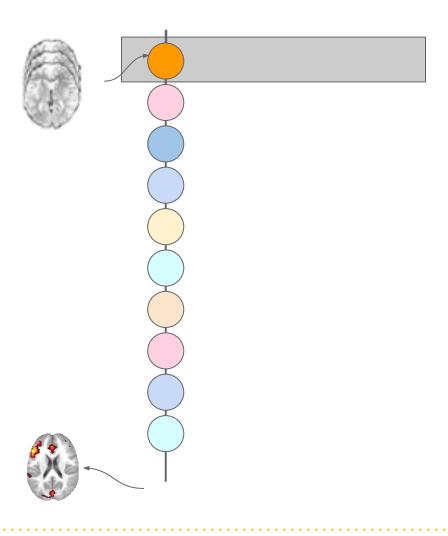


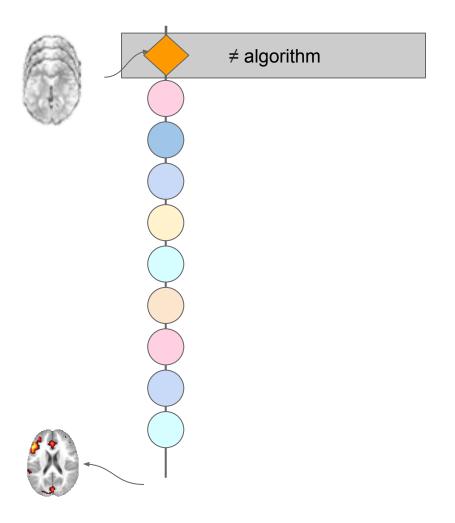


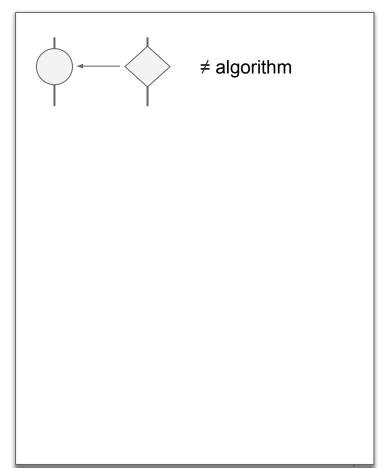
Different types of analytical variations

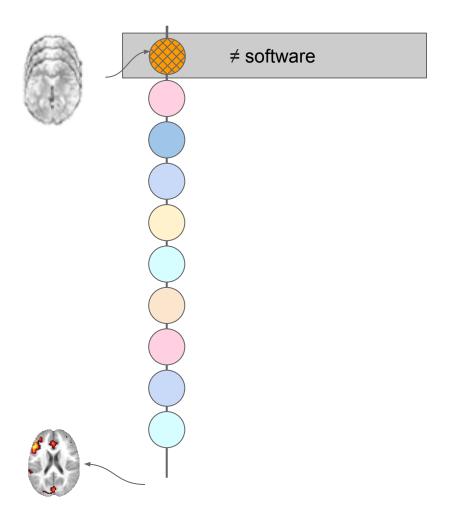


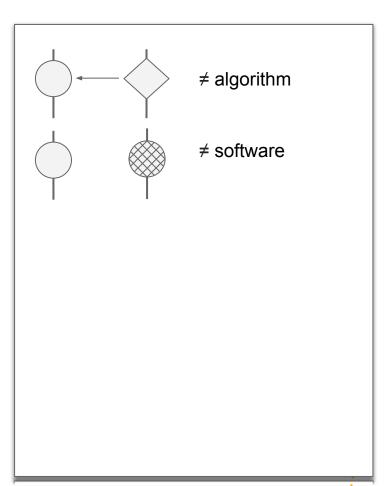


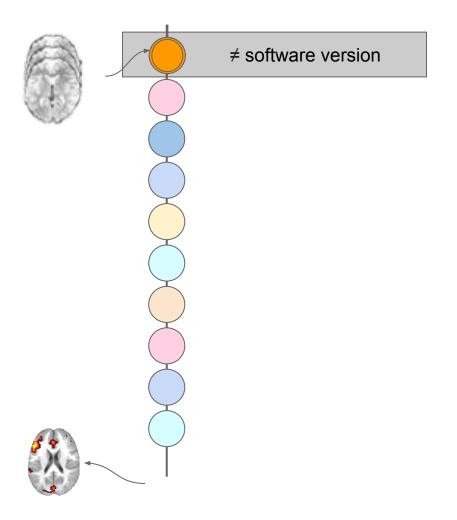


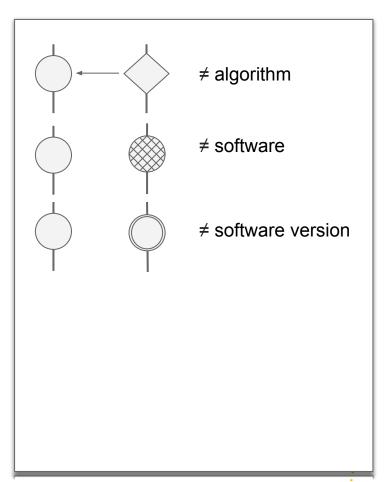


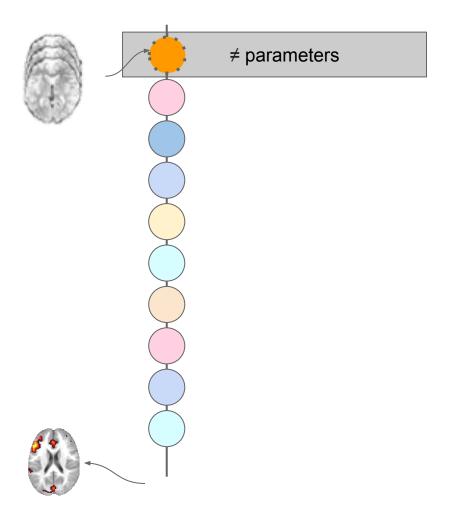


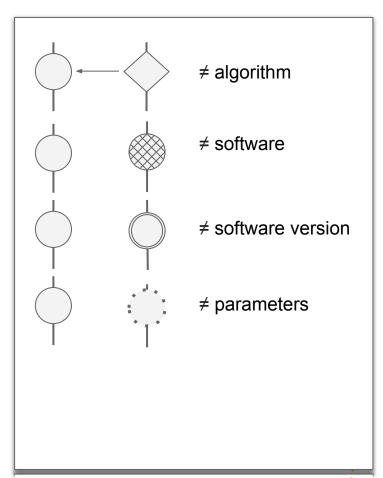


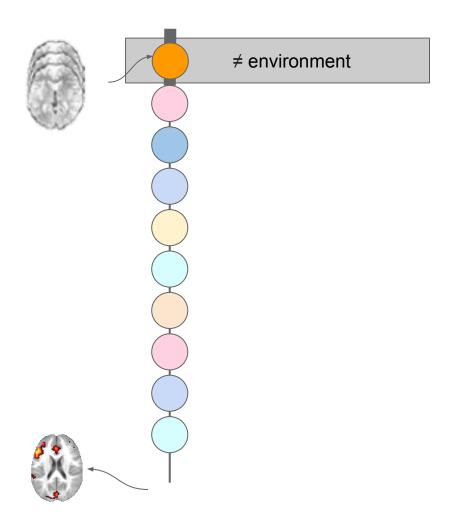


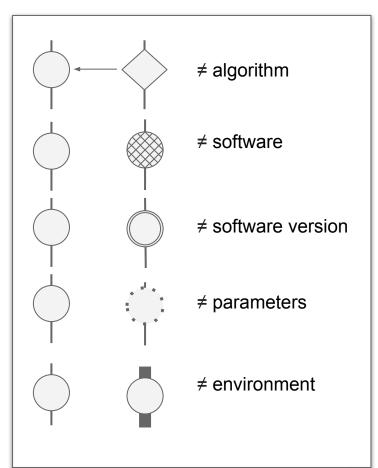


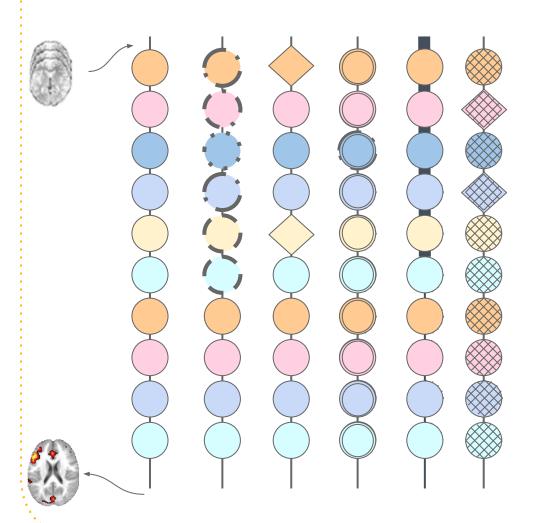


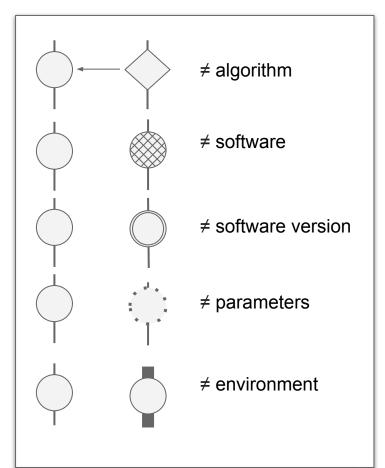










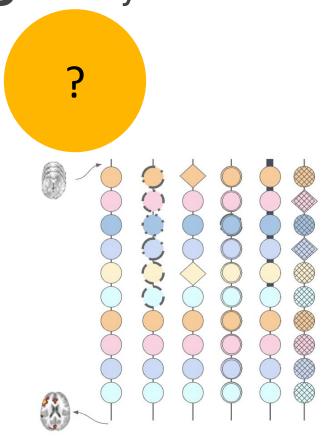


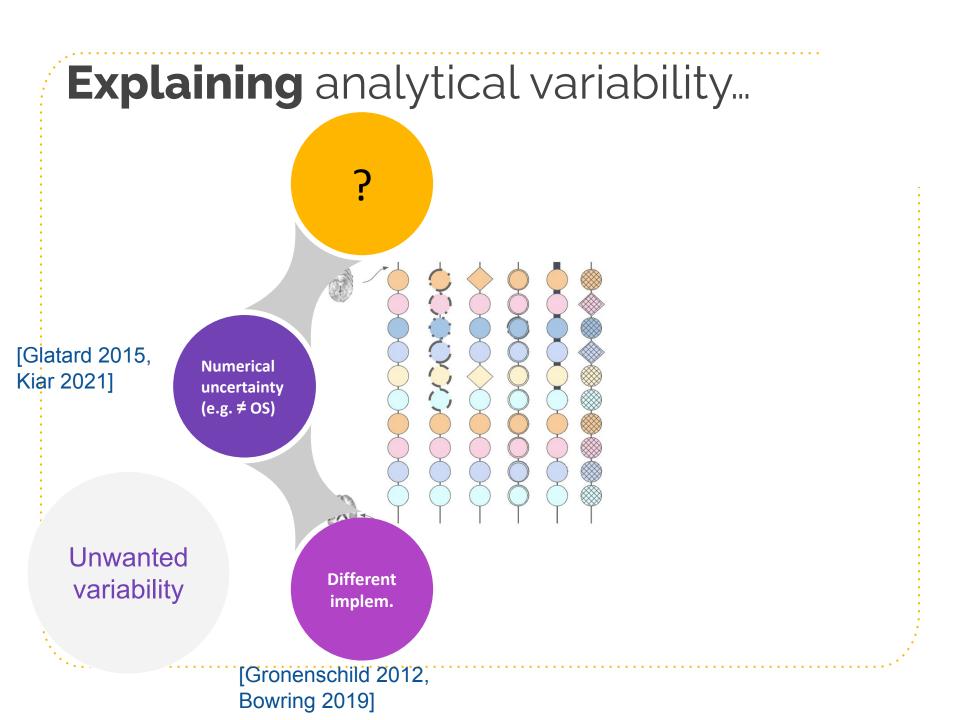
Can we explain this analytical variability?

Why is it useful to study analytical variability?

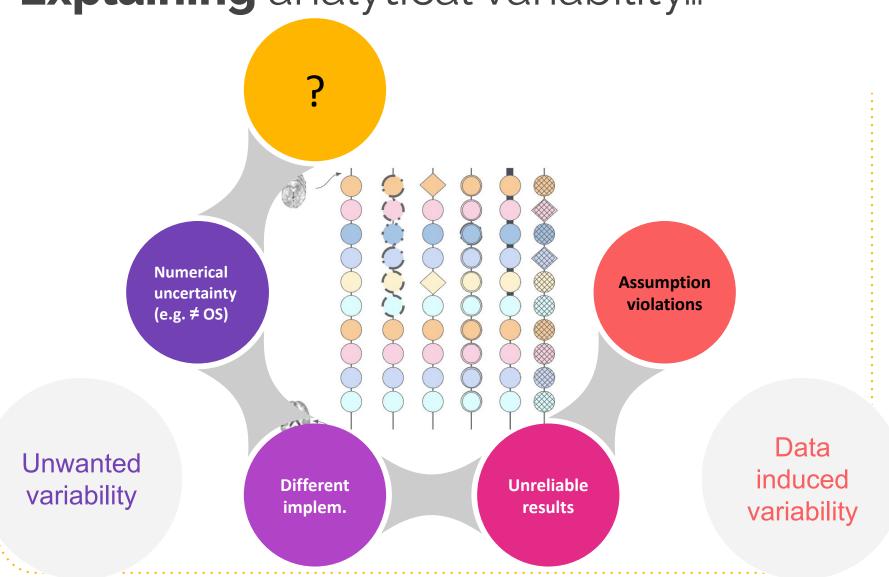
- Impacts the **reliability** of existing findings
- Impacts the reusability of derived datasets

Explaining analytical variability...

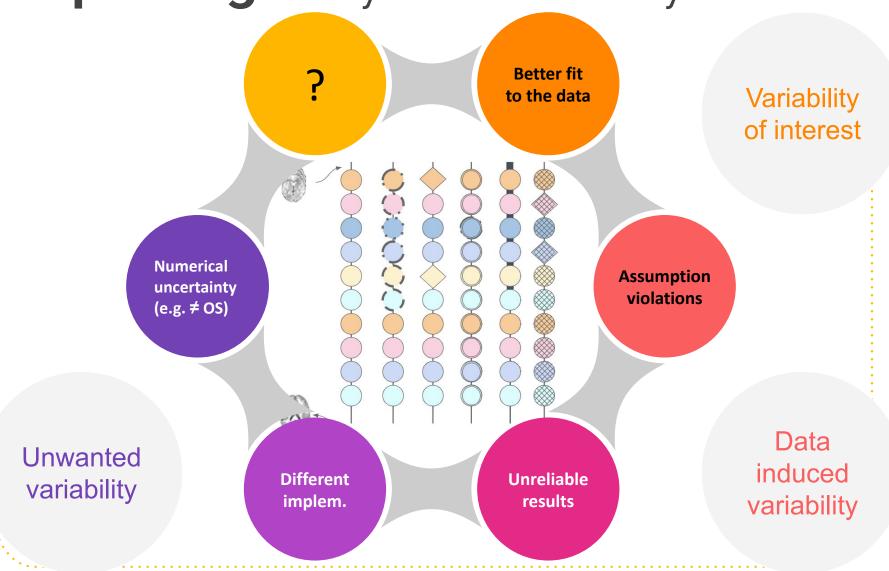




Explaining analytical variability...



Explaining analytical variability...

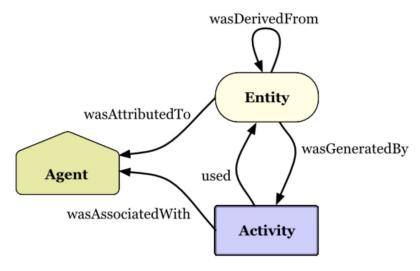


How can recording provenance help manage analytical variability?

Provenance

"Provenance describe how an artifact or set of artifacts was produced" Wikipedia

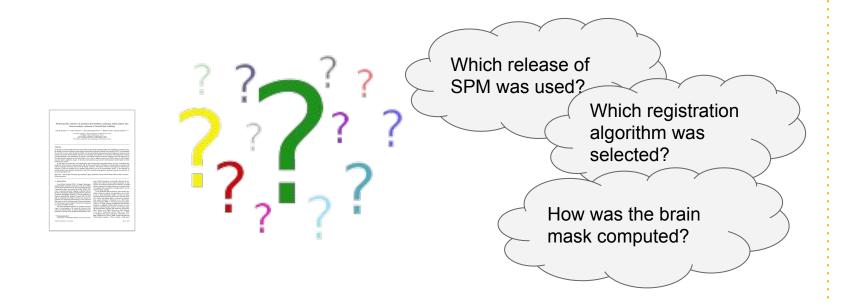
"Provenance is information about entities, activities, and people involved in producing a piece of data [...]" W3C PROV.



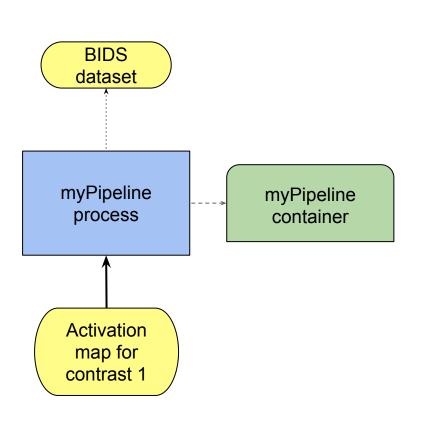
W3C PROV model

https://www.w3.org/TR/2013/NOTE-prov-primer-20130430/





reproduce
exactly the same
results
(same data, same
methods)?



Can we
reproduce
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results
(same data, same

(same data, same methods)?

Provenance Use Case 2 Dealing with unwanted variability

A previously
widely-used
parameter is no
longer considered
as best practice

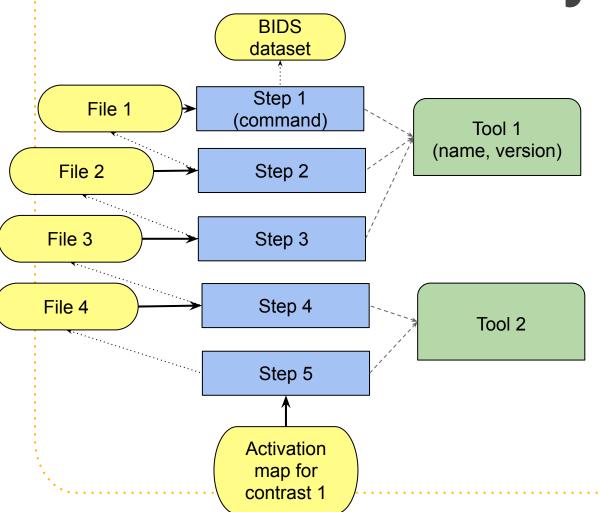
Provenance Use Case 2 Dealing with unwanted variability

A previously
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as best practice

A bug was found in a given tool version, can we retrieve all corresponding results?

Provenance Use Case 2 Dealing with

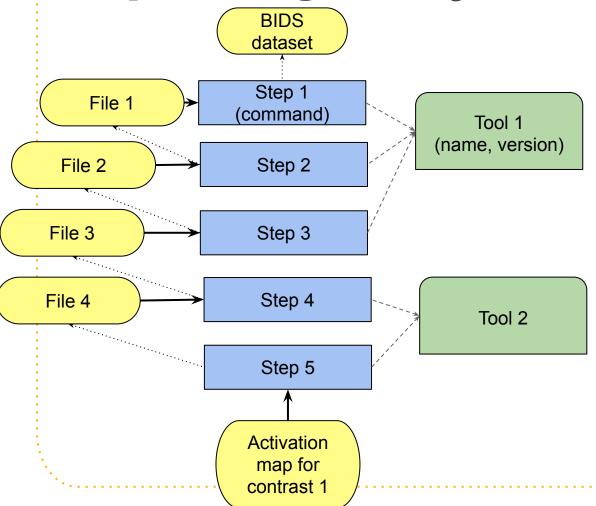
unwanted variability



A previously
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A bug was found in a given tool version, can we retrieve all corresponding results?

Provenance Use Case 3 Exploring factors impacting analytical variability



Low level: Are my conclusions dependant on the size of the smoothing kernel used at Step 5?

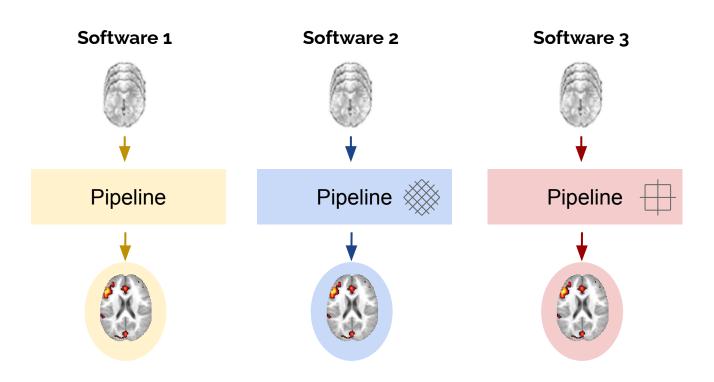
Provenance Use Case 3 Exploring factors impacting analytical variability

Variability across software

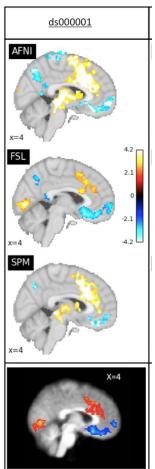
- Reproduced 3 published functional MRI studies
- Using 3 different neuroimaging software packages







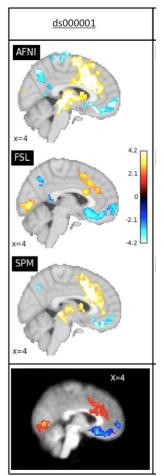
Variability across software

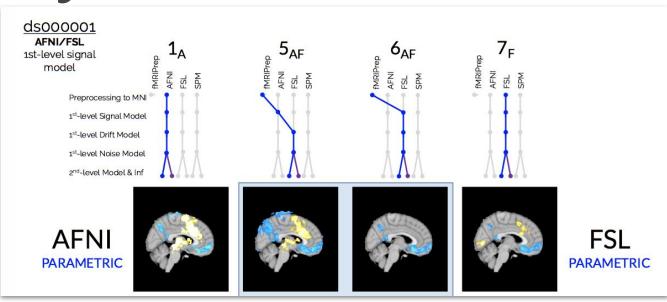


Thresholded maps in AFNI, FSL, SPM + original

[Bowring et. al, HBM 2019]

Variability across software





[Bowring et. al, HBM 2021]

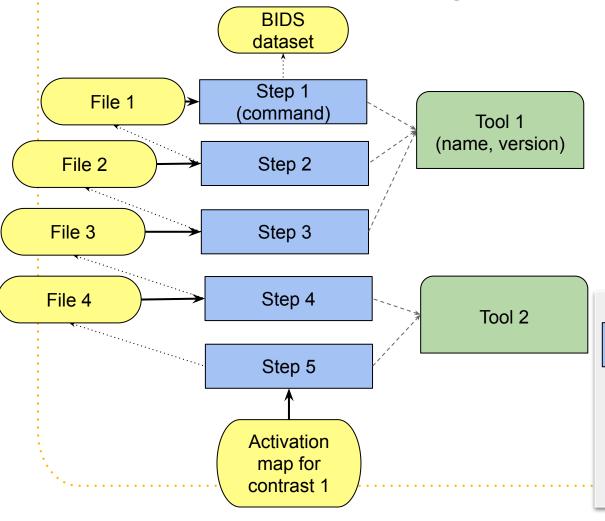
Thresholded maps in AFNI, FSL, SPM + original

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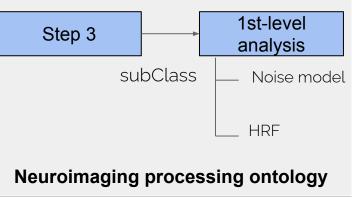
Provenance Use Case 3 Exploring factors impacting analytical variability

High level: Are my conclusions dependant on the type on the noise model in the 1st-level analysis?

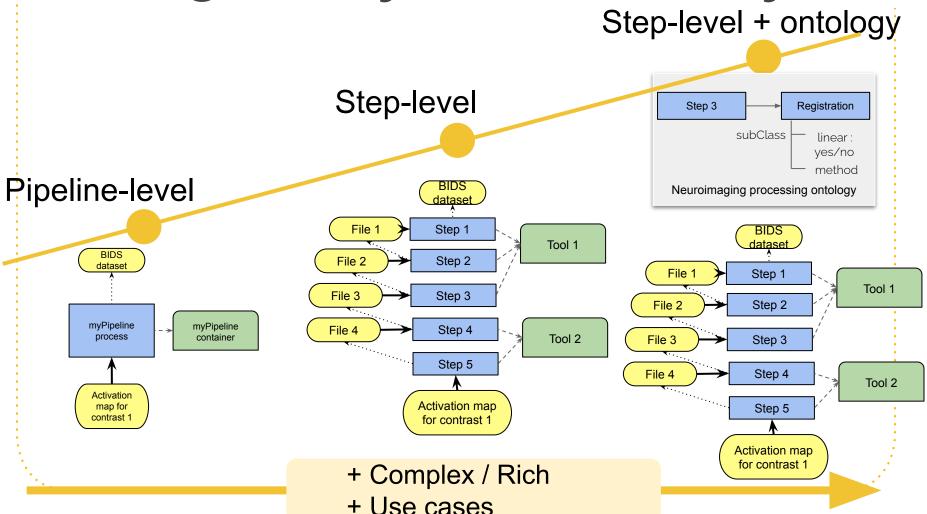
Provenance Use Case 3 Exploring factors impacting analytical variability



High level: Are my conclusions dependant on the type on the noise model in the 1st-level analysis?



Recording provenance to help manage analytical variability



Recording provenance to help manage analytical variability

Step-level + ontology **BIDS-Prov** Step-level Step 3 Registration subClass linear: yes/no method Pipeline-level Neuroimaging processing ontology dataset Step 1 File 1 Tool 1 dataset **BIDS** Step 2 File 2 dataset File 1 Step 1 Tool 1 File 3 Step 3 File 2 Step 2 myPipeline myPipeline File 4 Step 4 process container Tool 2 File 3 Step 3 Step 5 File 4 Step 4 Tool 2 Activation Activation map map for Step 5 contrast 1 for contrast 1 Activation map for contrast 1 + Complex / Rich + Use cases



BIDS Extension Proposal 28

Moderators: Satra Ghosh et Camille Maumet

Contributors: Rémi Adon, Tibor Auer, Stefan Appelhoff, Michael Dayan, Eric Earl, Yaroslav Halchenko, Matthieu Joulot, Chris Markiewicz, Cyril R. Pernet, Jean-Baptiste Poline, Omar El Rifai, Sarah Saneei, Ghislain Vaillant.

Thank you!

BIDS-Prov is open for community feedback https://github.com/bids-standard/BEP028 BIDSprov

steering committee: Guiomar Niso, Melanie Ganz, BIDS Robert Oostenveld, Russell Poldrack, Ariel Rokem

BIDS maintainers: Stefan Appelhoff, Chris Markiewicz, Franklin Feingold, Taylor Salo, Rémi Gau, Ross Blair, Anthony Galassi, Eric Earl

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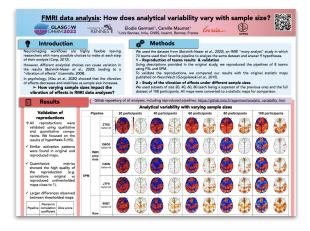
Camille Maumet



Poster WTh872 How does analytical variability vary with sample sizes? **Elodie Germani**

Thank you!







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