

Presentation of the project 'Narps Open Pipeline'

Jeremy Lefort-Besnard, Camille Maumet, Boris Clenet, Elodie Germani

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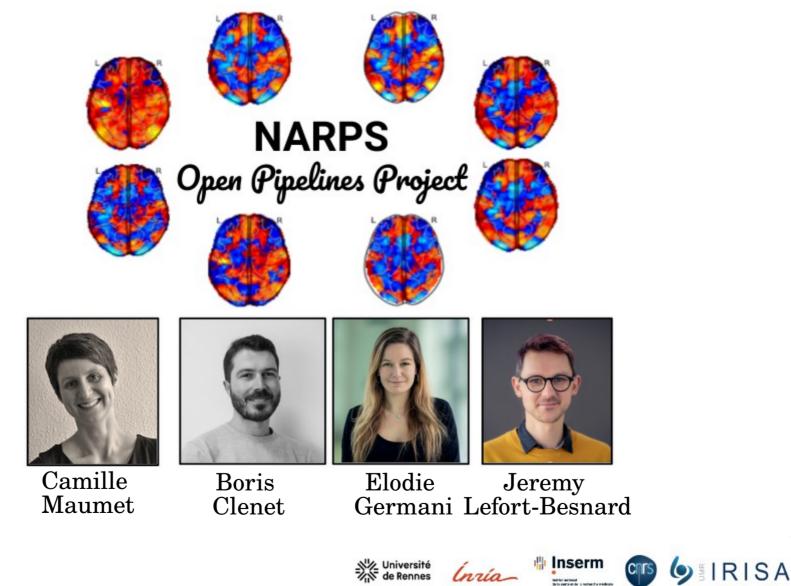
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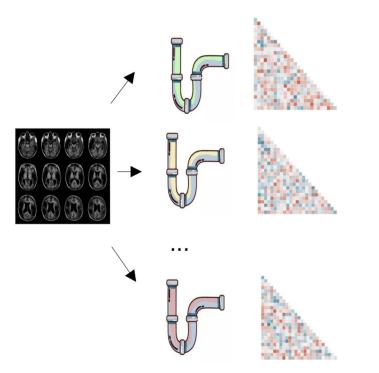


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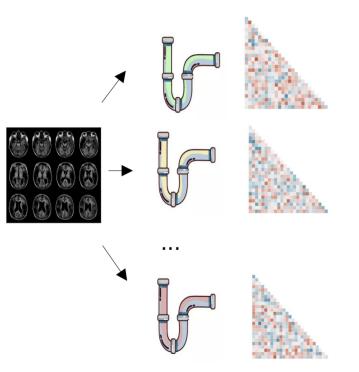


EMPENN

Analytical variability has often been overlooked assuming it was negligible compared to other sources of variability



Analytical variability has often been overlooked assuming it was negligible compared to other sources of variability



In 2020 a landmark paper in Nature challenged this status-quo:

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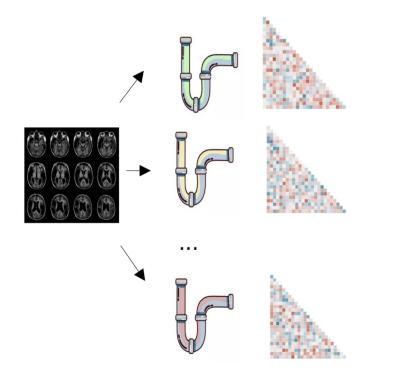
Variability in the analysis of a single neuroimaging dataset by many teams

Rotem Botvinik-Nezer, Felix Holzmeister, Colin F. Camerer, Anna Dreber, Juergen Huber, Magnus Johannesson, Michael Kirchler, Roni Iwanir, Jeanette A. Mumford, R. Alison Adcock, Paolo Avesani, Blazej M. Baczkowski, Aahana Bajracharya, Leah Bakst, Sheryl Ball, Marco Barilari, Nadège Bault, Derek Beaton Julia Beitner, Roland G. Benoit, Ruud M. W. J. Berkers, Jamil P. Bhanji, Bharat B. Biswal, Sebastian Bobadilla-Suarez, ... Tom Schonberg 🖾 + Show authors

Nature 582, 84-88 (2020) Cite this article

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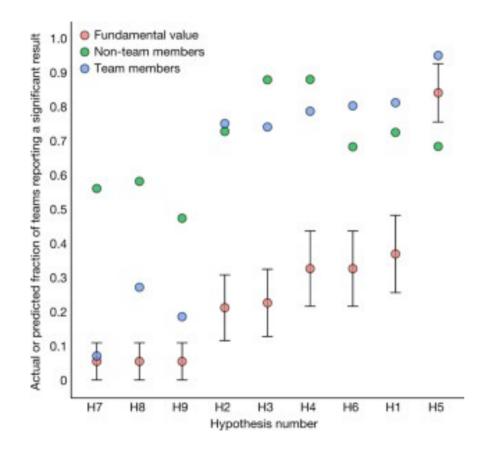
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70 expert teams (200 scientists) – Same yes/no research questions – Same dataset 1/5

Each team built a different pipeline

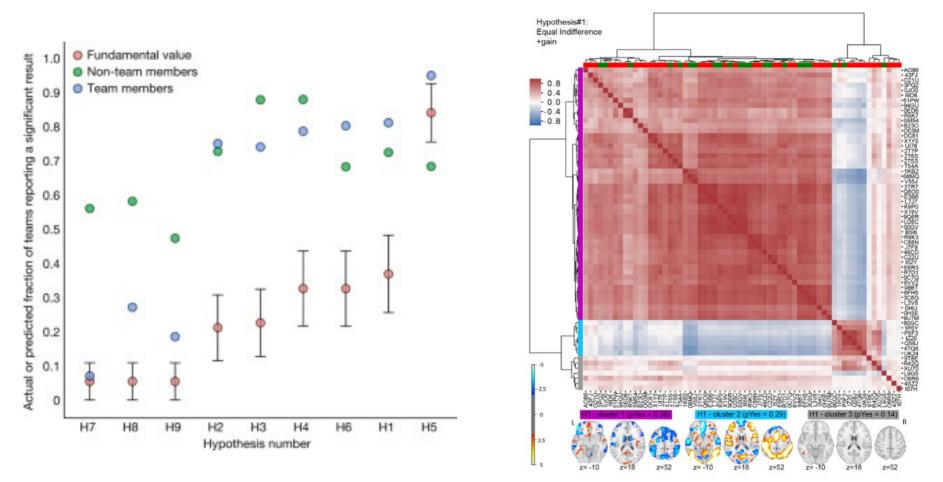
Each team built a different pipeline Differences in pipelines led to contradictory findings

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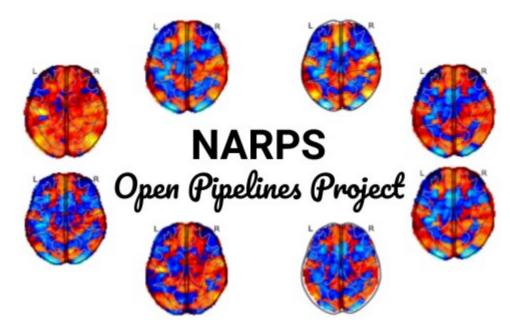


Figures from Botvinik-Nezer, R., Holzmeister, F., Camerer, C.F. et al. Variability in the analysis of a single neuroimaging dataset by many teams. Nature 582, 84–88 (2020). https://doi.org/10.1038/s41586-020-2314-9

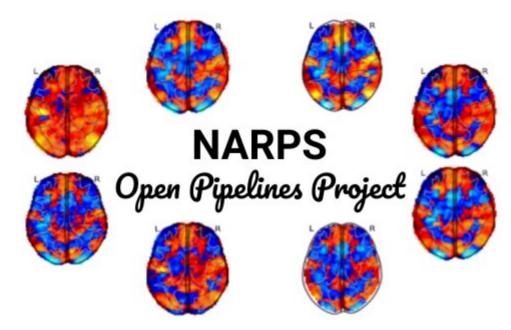
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Figures from Botvinik-Nezer, R., Holzmeister, F., Camerer, C.F. et al. *Variability in the analysis of a single neuroimaging dataset by many teams*. Nature 582, 84–88 (2020). https://doi.org/10.1038/s41586-020-2314-9



Create a codebase reproducing the 70 pipelines of the NARPS study and share this as an open resource for the community



Create a codebase reproducing the 70 pipelines of the NARPS study and share this as an open resource for the community



Github.com/Inria-Empenn/narps_open_pipelines



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Use a NARPS pipeline for your project:

narps_open_pipelines / narps_open / pipelines / team_2T6S.py		
Code	Blame Executable File · 664 lines (562 loc) · 27.8 KB	
	#!/usr/bin/python	
	# coding: utf-8	
	""" Write the work of NARPS team 2T6S using Nipype """	
	from os.path import join	
	from itertools import product	
	from nipype import Workflow, Node, MapNode	
	from nipype.interfaces.utility import IdentityInterface, Function	
11	from nipype.interfaces.io import SelectFiles, DataSink	
12	from nipype.interfaces.spm import (
	Smooth,	
	Level1Design, OneSampleTTestDesign, TwoSampleTTestDesign,	
	EstimateModel, EstimateContrast, Threshold	
	from nipype.algorithms.modelgen import SpecifySPMModel	
	from nipype.algorithms.misc import Gunzip	
	from narps open.pipelines import Pipeline	

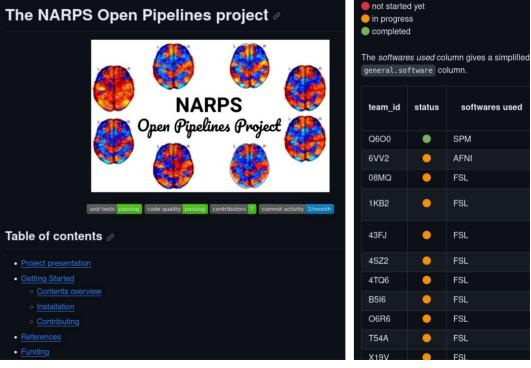


$Github.com/Inria-Empenn/narps_open_pipelines$

Use a NARPS pipeline for your project:

narps_open_pipelines / narps_open / pipelines / team_2T6S.py		
bclenet 2T6S pipeline reproduction (#75) -		
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Contribute to NARPS Open Pipelines:





This project is supported by Région Bretagne (Boost MIND) and by Inria (Exploratory action GRASP)





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This project receives contributions by Rémi Gau and during events such as OHBM Brainhack 2022 and 2023, e-ReproNim FENS NENS Cluster Brainhack 2023, ORIGAMI lab hackathon 2023 by : Arshitha Basavaraj, Trang Cao, Anna Menacher, Liz Bushby, Michael Dayan, Aimee Westbrook, Yaroslav Halchenko, Paul Taylor





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Come meet us and/or contribute during our next brainhack sessions:

- Monthly Narps Open hours
- Marseille brainhack 2023
- OHBM brainhack 2024

