

Computer Age Neuroscience and The Future of Science Process Management

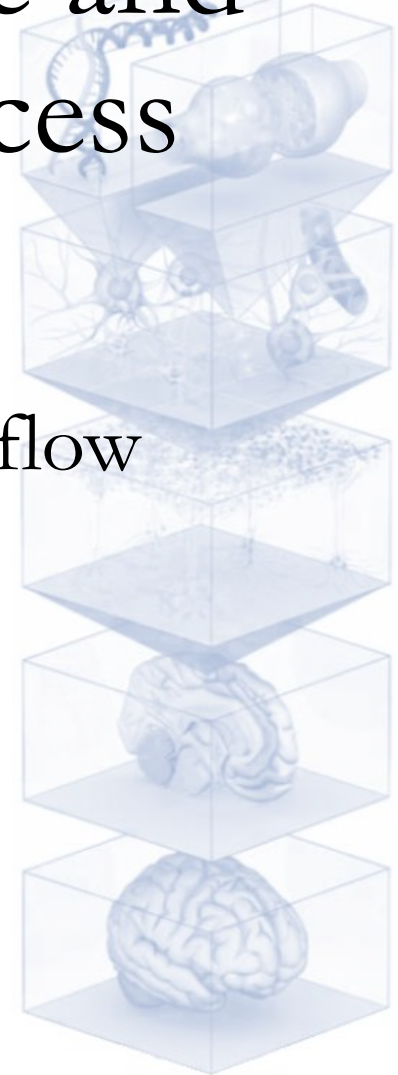
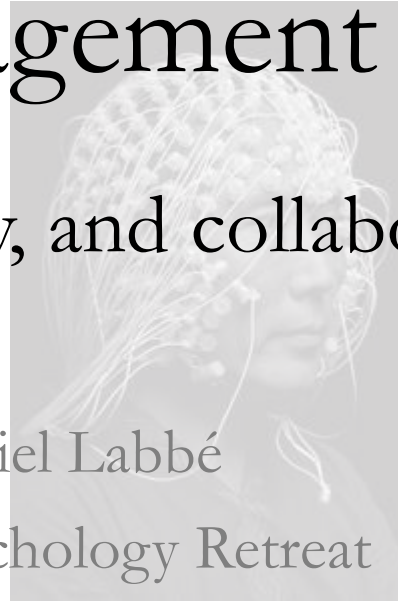
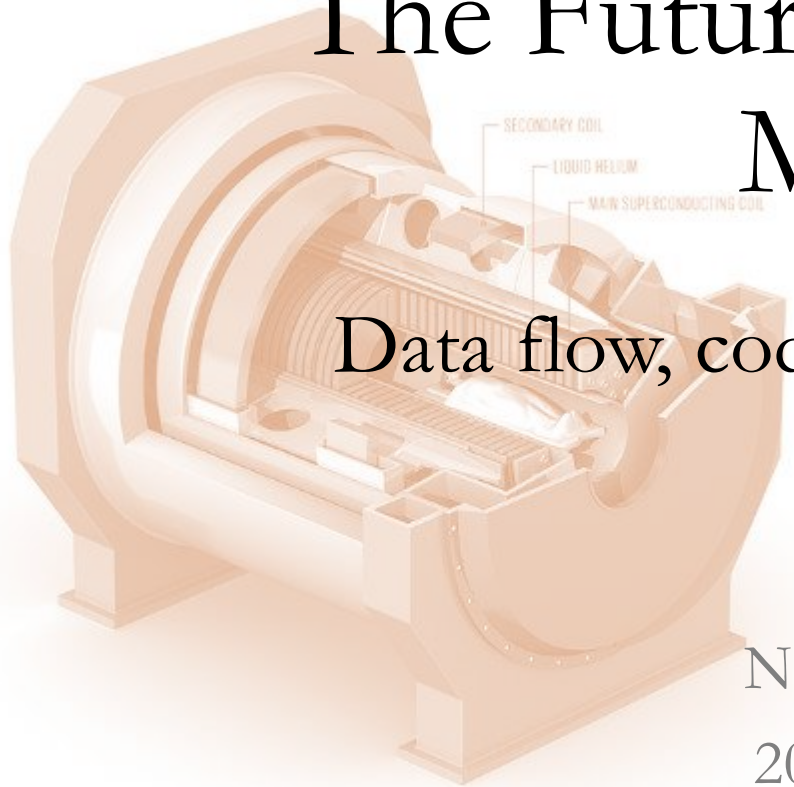
Data flow, code flow, and collaborative flow

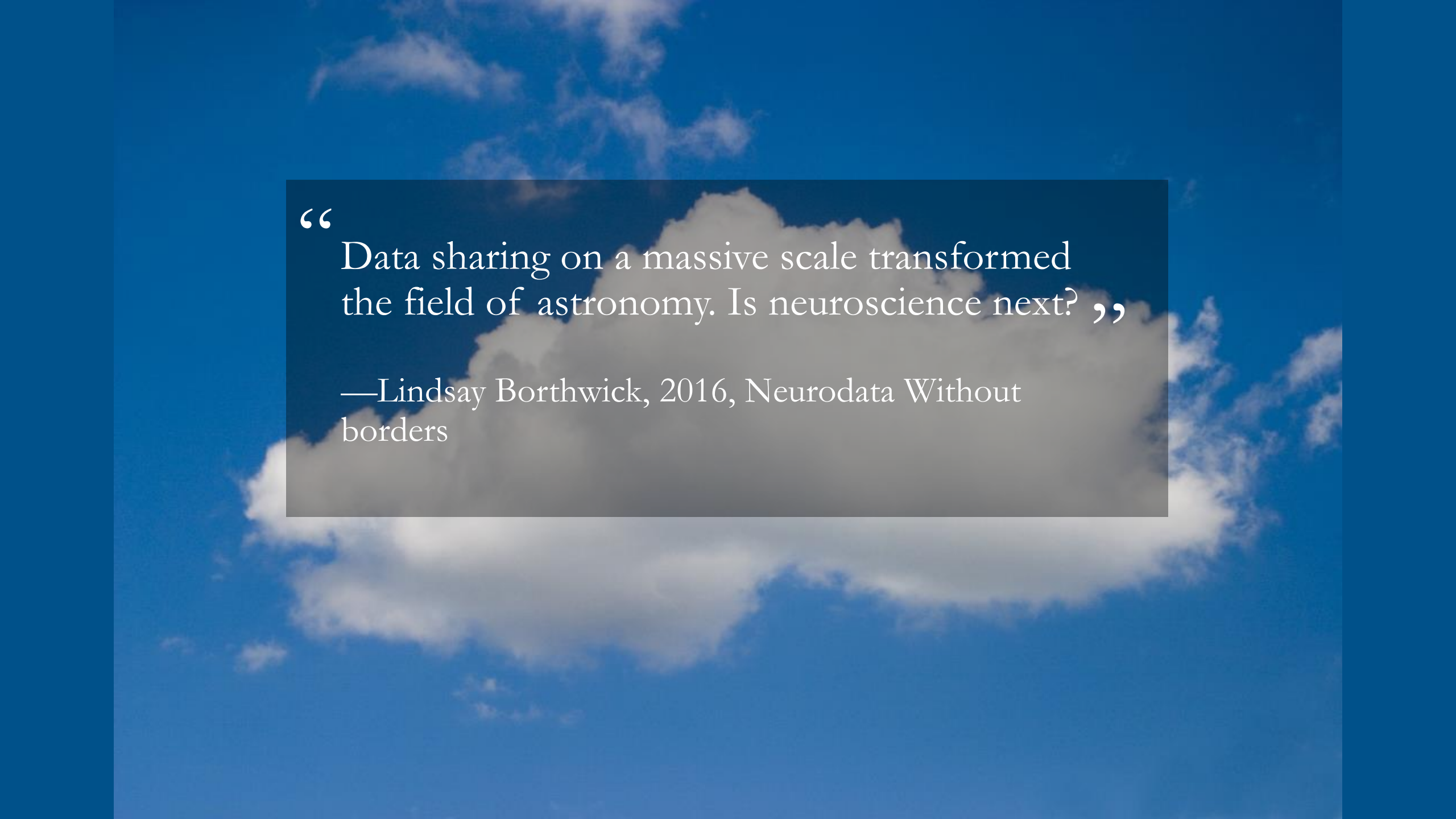
Daniel Labbé

Neuropsychology Retreat

2017-07-03 - 2017-07-05

daniel.labbe@rub.de





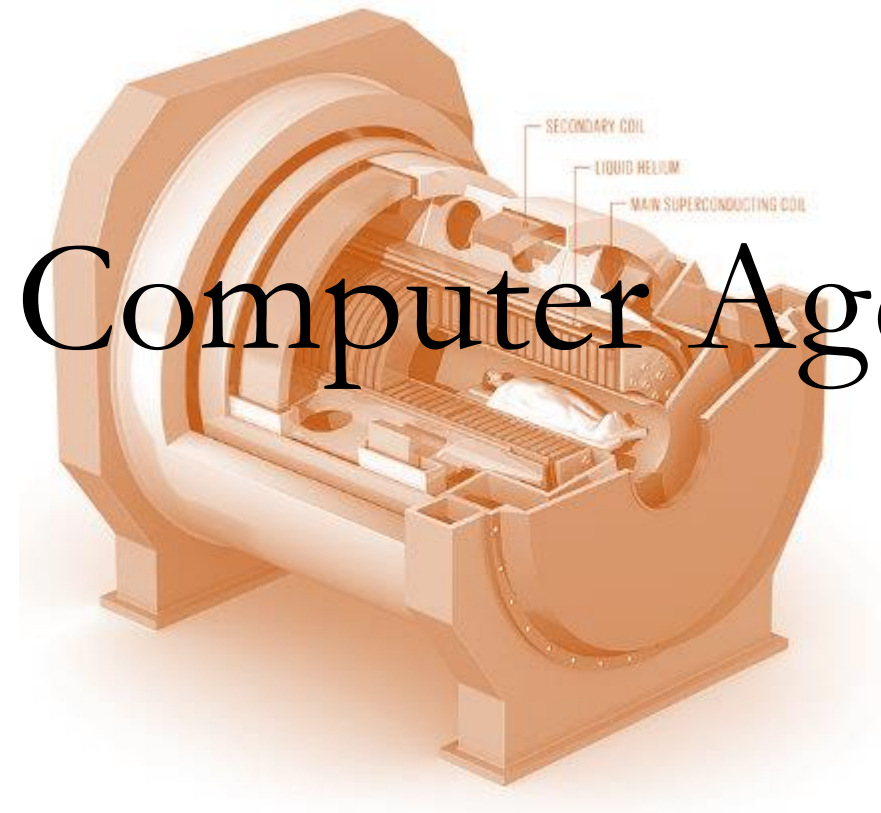
“Data sharing on a massive scale transformed the field of astronomy. Is neuroscience next? ”

—Lindsay Borthwick, 2016, Neurodata Without borders

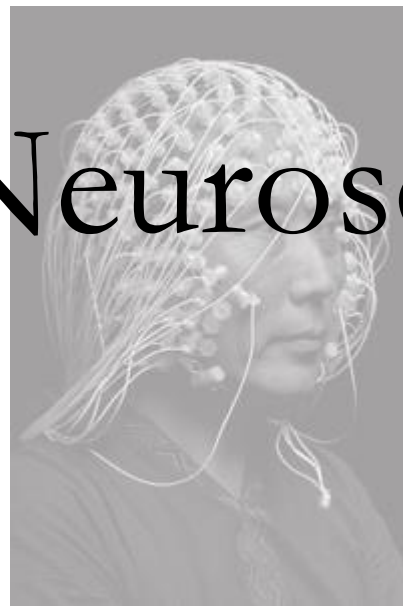
Outline

- Part I: Computer Age Neuroscience
- Part II: Science Process Management
- Discussion





Computer Age Neuroscience

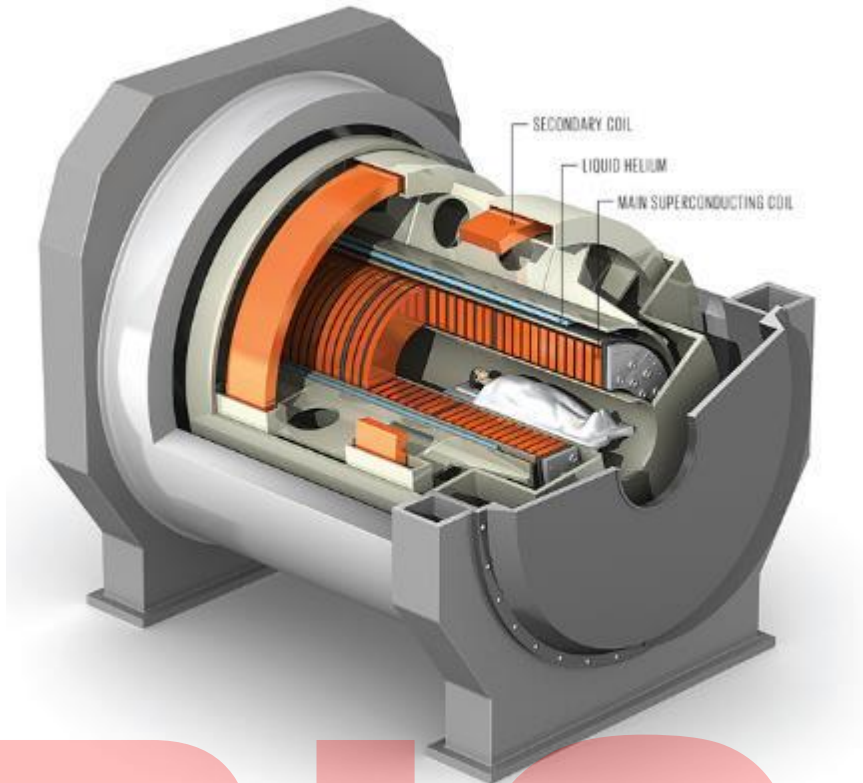


Computer Age Neuroscience

- How The End of Moore's Law Prompts Parallel Computing
- The Inescapable Imminence of the Cloud
- Cloud Neuroscience
- The International Brain Station (TIBS)



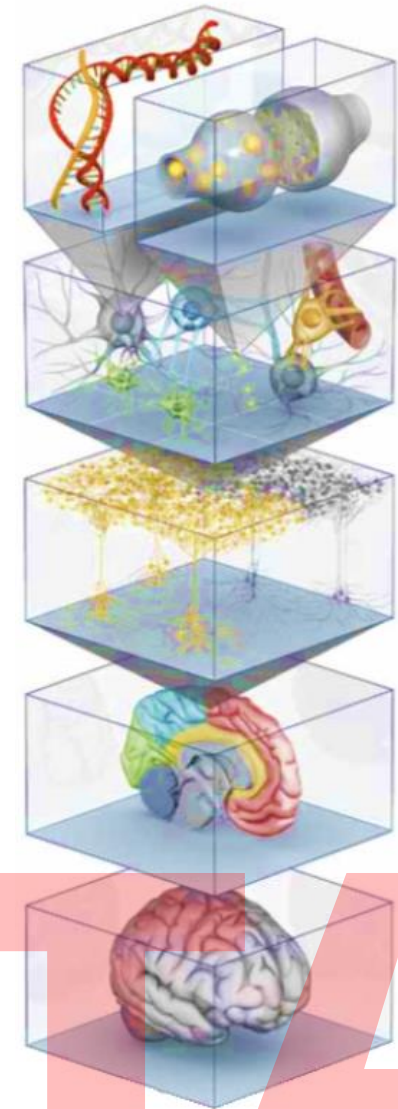
More Voxels, More Channels, and Multiple Levels of Simulation



11.7 Tesla



256 Electrodes



BIG DATA

Computer Age Neuroscience

- "...computer-based technology allows scientists to collect enormous data sets, orders of magnitude larger than those that classical statistical theory was designed to deal with; **huge data demands new methodology**, and the demand is being met by a burst of innovative computer-based statistical algorithms." —Bradley Efron and Trevor Hastie, 2016, p.4 *Computer Age Statistical Inference*. NY: Cambridge University Press. (My emphasis.)
- "large-scale prediction algorithms" such as "neural nets, deep learning, boosting, random forests, and support-vector machines" **became prominent due to the commercial value of prediction** (Efron & Hastie, 2016, p.446, my emphasis)
- **Data science** became more about the **performance** of algorithms for big data (Efron & Hastie, 2016)



BRADLEY EFRON
TREVOR HASTIE

COMPUTER AGE STATISTICAL INFERENCE

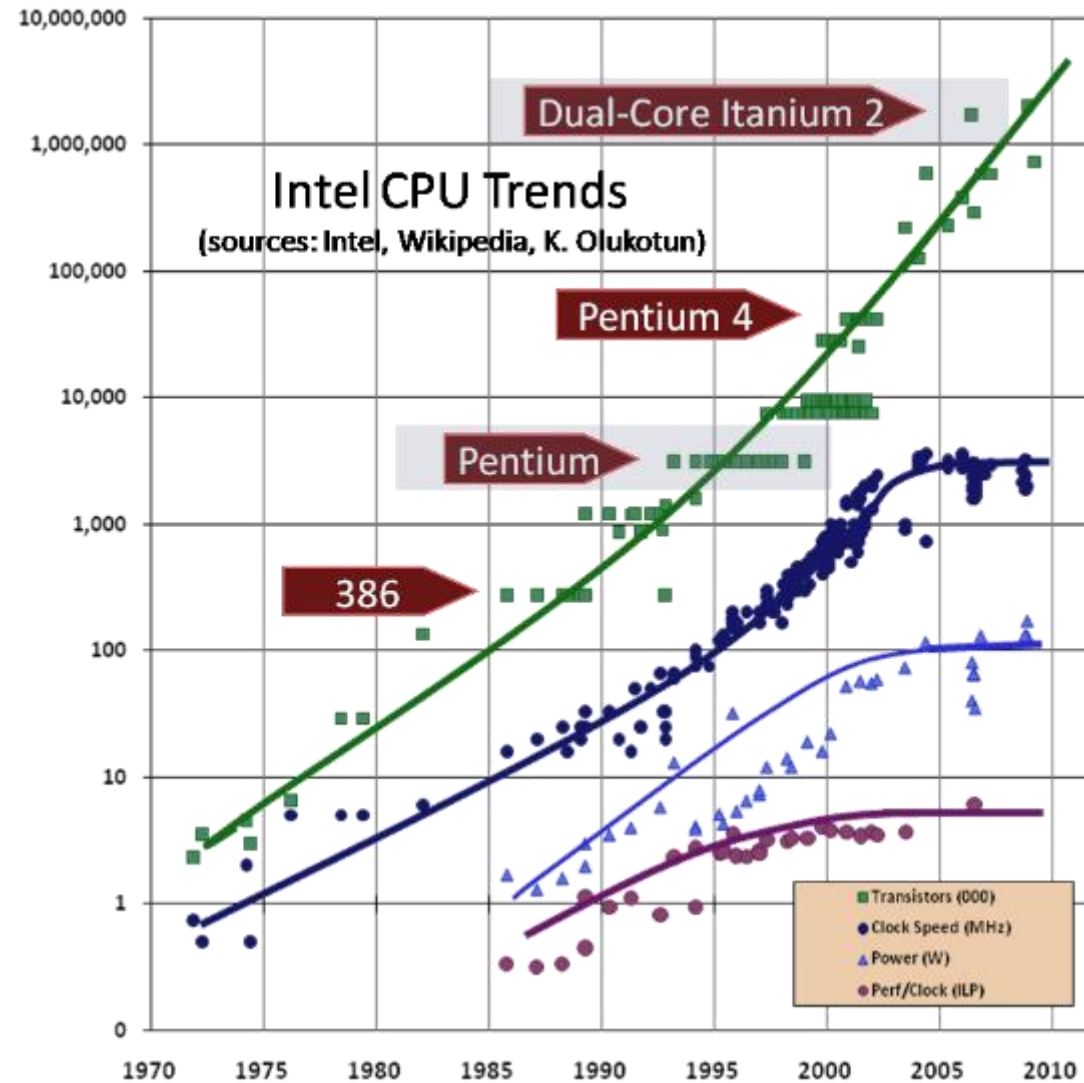
ALGORITHMS, EVIDENCE, AND DATA SCIENCE

How The End of Moore's Law Prompts Parallel Computing



<http://www.nature.com/news/the-chips-are-down-for-moore-s-law-1.19338>

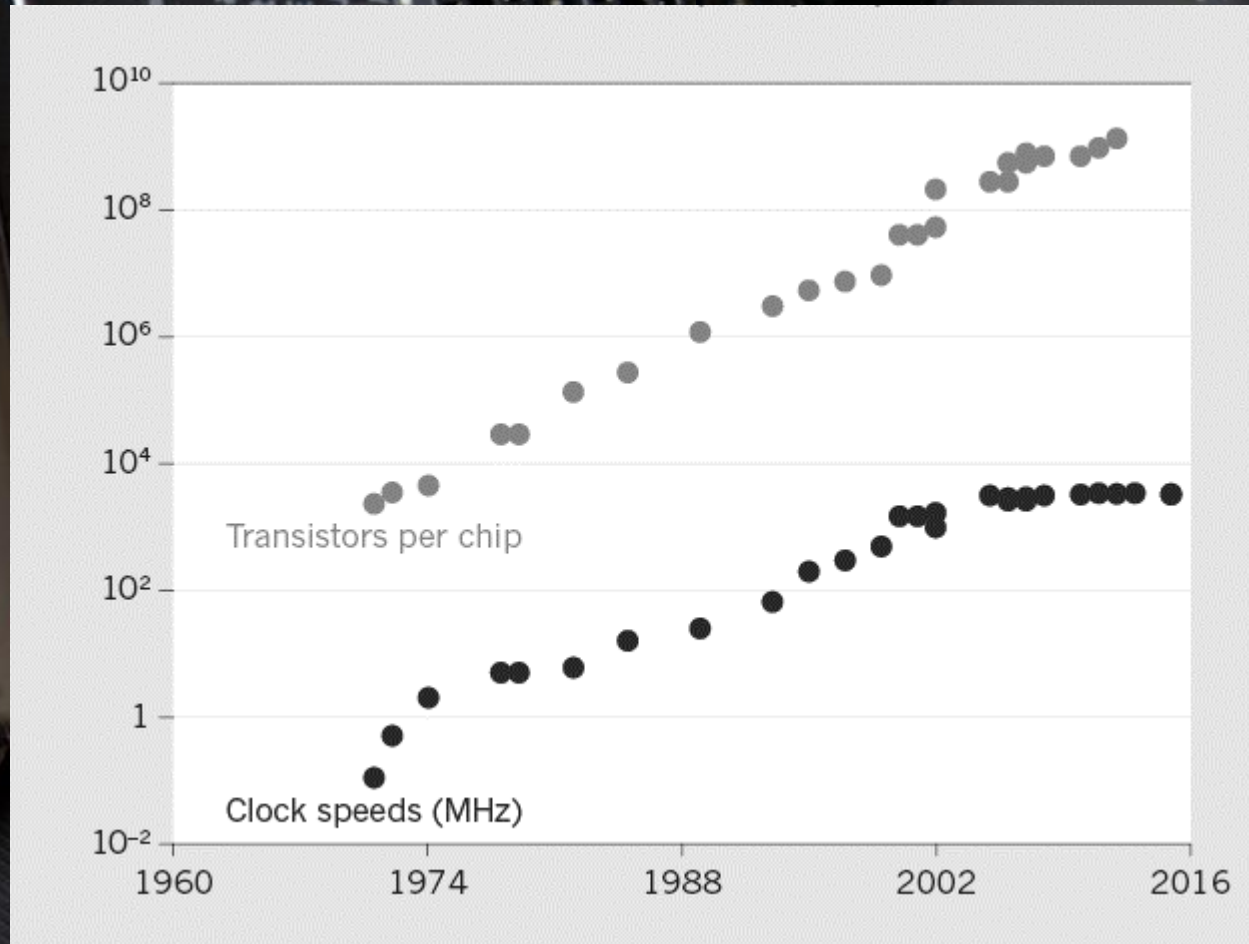
CPU Clock Speed Peaked in the Early 2000nds



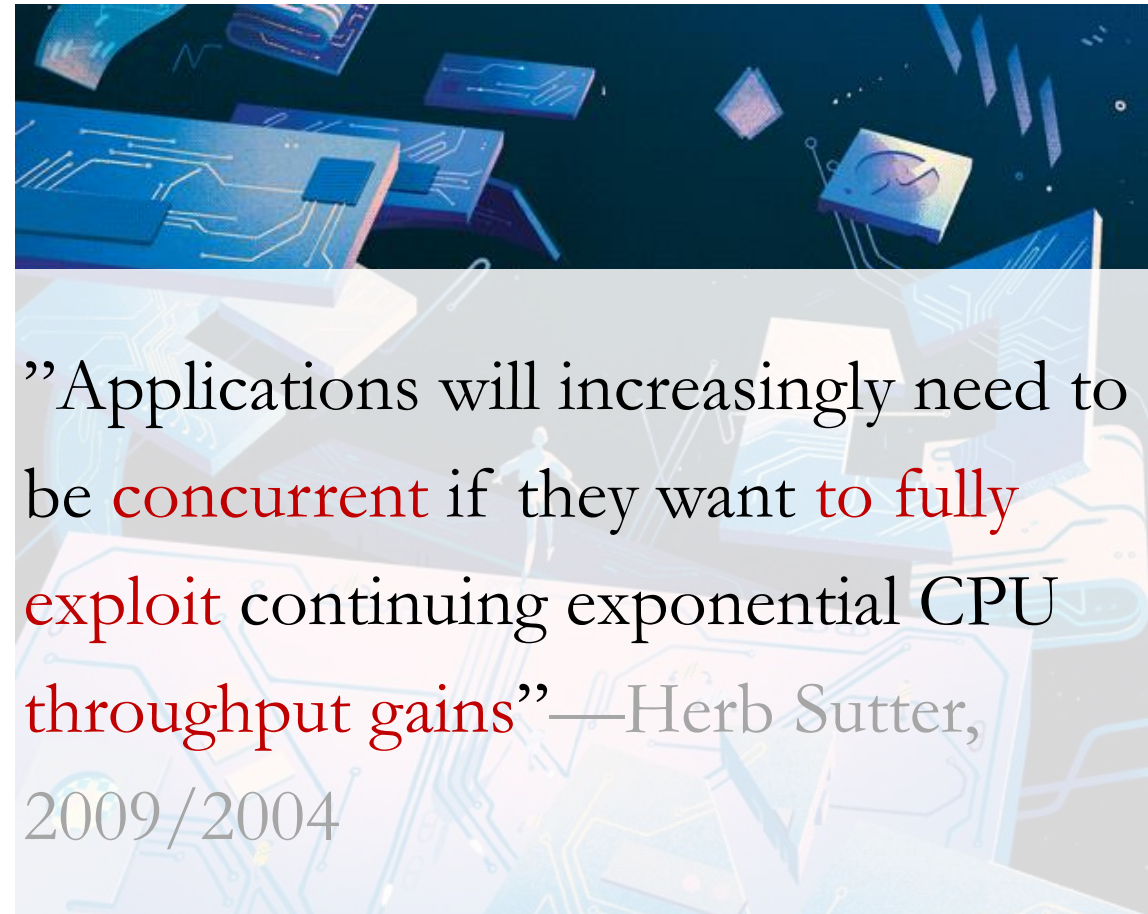
Sutter, H. (2009, 2005, 2004). Dr. Dobb's Journal. [Online Access]

Instruction-level
parallelism (ILP)

Heat does not Dissipate Fast Enough in High Density
Processing Arrays: Using multiple cores is a workaround



Waldrop, M. M. (2016).[News Feature] The chips are down for Moore's law, *Nature*.



”Applications will increasingly need to be **concurrent** if they want to **fully exploit** continuing exponential CPU **throughput gains**”—Herb Sutter, 2009/2004



The Inescapable Imminence of the Cloud

“[for] an application that runs well on today’s hardware and will just naturally run faster or better on tomorrow’s hardware, you need to write an app with lots of juicy latent parallelism...”

Costs

- Extra development effort
- Extra code complexity
- Extra testing effort

The Inescapable Imminence of the Cloud

The Good News

“(cont.) For many classes of applications **the extra effort will be worthwhile**, because concurrency will let them fully exploit the **exponential gains** in compute throughput that will continue to grow strong and fast **long after Moore’s Law** has gone into its sunny retirement, as we continue to mine the **cloud for the rest of our careers.**”—Sutter, H. (2012) (My emphasis.)

Cloud Neuroscience

“The kernel of the idea is based on a view of the scientific process as an ‘upward spiral’: a collective effort where each new experiment yields data, upon which analysis is performed, leading to new or refined models, which suggest novel experiments. The goal of The International Brain Station (TIBS) is to build technology that would democratize brain science, so that all brain scientists—professional and otherwise—can build on the shoulders of one another. To make that a reality, TIBS would enable “cloud neuroscience,” meaning that the **data**, the **code** and the **analytic results** from neuroscience experiments all live in the cloud together, where it could be accessed by anyone.”

—Joshua T. Vogelstein [Interview], *Neurodata Without borders*. My emphasis.

<http://www.nwb.org/2016/11/11/unlocking-the-brain-with-open-data/>

The International Brain Station (TIBS)

“[A] global neuroscience collaboration that would link [neuroscientists] efforts and rival big science investments in astronomy and physics”—

Underwood, E., Science, [In Depth][Summary] International Brain Projects Proposed. doi: 10.1126/science.352.6283.277

“This resource will realize a new era of brain sciences, one in which the bottlenecks to discovery transition away from data collection and processing to data enriching exploring, and modeling.”—Global Brain Workshop

2016 Attendees, Grand Challenges for Global Brain Sciences. <http://brainx.io/>

Scaling Up Neuroscience

Scaling Up Neuroscience is a series of commentaries published in Nature Reviews Neuroscience.

“Several **large-scale international research initiatives** have recently been launched, fuelling substantial financial investments in neuroscience and raising expectations for the development of new knowledge and therapies. Meeting these expectations will require **global coordination** of stakeholders and the **adoption of team-based approaches that are not yet the norm for neuroscience.**” —Olds, J. L., [Abstract] Nature Reviews Neuroscience [Comment]. My emphasis.

BIG SCIENCE

Summary Part I: Computer Age Neuroscience

- Big Data emphasizes performance
- Vertical limit for processing units (CPUs, GPUs) reached
- Horizontal programming (parallelism, concurrency) required to future-proof software
- Cloud Neuroscience
 - Costs extra
 - Exponential gains
 - You cannot run from the cloud → Embrace the cloud
- Big Science requires a collaborative effort (team neuroscience)

How do we get there?

Science Process Management

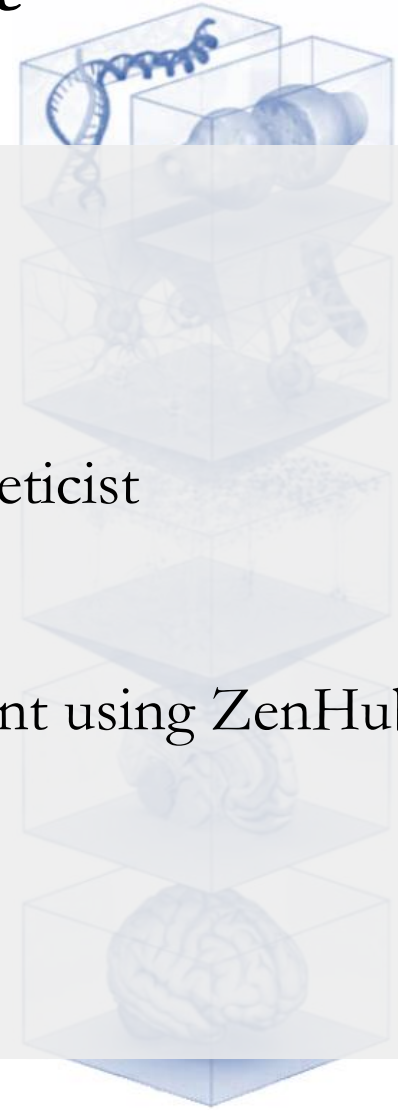
Optimizing & Maintaining Data Flow, Code Flow, and Collaborative Flow

Three Challenges

- Data Flow: Parallelization & Scalability
- Code Flow: Sharing Code and Analysis in the Clouds
- Collaborative Workflow: Team Neuroscience

Part II: Science Process Management

- Definitions
- Software Examples
 - Pipelines—How to Achieve **Data Flow** Like a Molecular Geneticist
 - Version Control—**Code Flow** with Git
 - Lean Process Management—NASA Style Process Management using ZenHub:
Integrating **code flow** with **workflow**
 - The Digital Notebook—Transparent **Knowledge Transfer**



Science Process Management

- **Scientific workflow**
”...a formal description of a process for accomplishing a scientific objective, usually expressed in terms of tasks and their dependencies” (Ludäscher et al. 2009, section 13.1)
- **Scientific workflow system** (Ludäscher et al., 2009)
 - Automatization
 - Documentation
 - Monitoring and recording (the process is requires regular self-regulation)
 - Optimization
 - Reusability
- Example: A paralell processing pipeline

The Human Process

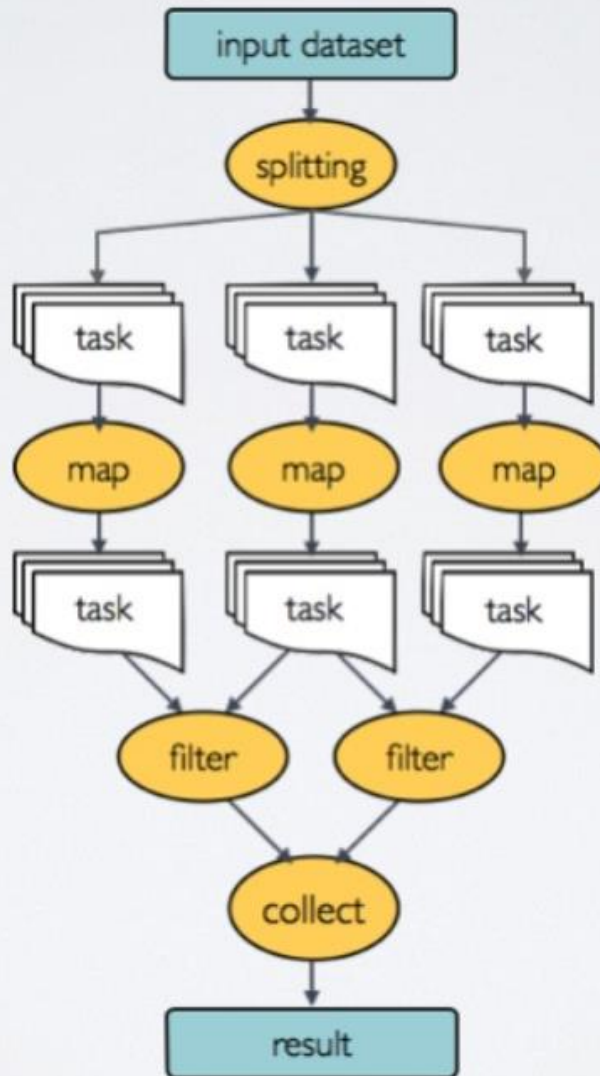
“ ‘human cycles’ are a sometimes neglected resource, which can and should be optimized as well. For example, the use of data and workflow provenance information can be used for traditional purposes (such as optimizing system performance or improving fault-tolerance [citation]), but also to enhance the scientist’s insights when trying to understand or debug scientific workflow results [citation].”

—Bertram Ludäscher et al., 2009, section 13.7, Scientific Process Automation and Workflow Management. In: Scientific Data Management. Chapman and Hall/CRC. doi: 10.1201/9781420069815

Pipelines

Paralellism and Concurrency using NextFlow

REACTIVE NETWORK



nextflow



Zero config



Polyglot

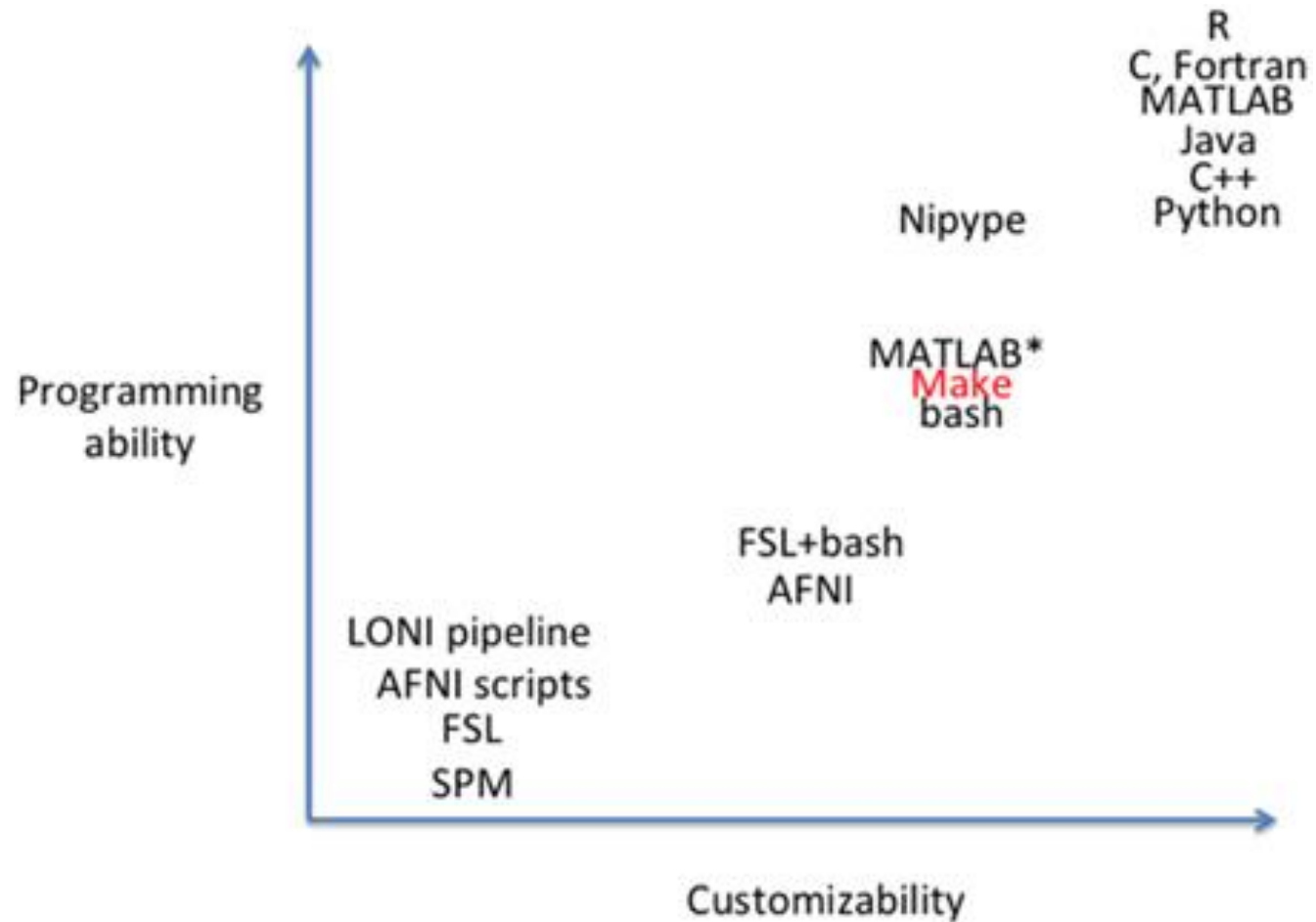


Concurrency



Scale easily

The next few slides are mainly from Paolo Di Tommaso, Center for Genomic Regulation, Barcelona (Switzerland HPC Conference 2016.)



* to call neuroimaging programs

Askren et al. (2016). Using Make for Reproducible and Parallel Neuroimaging Workflow and Quality-Assurance. *Frontiers in Neuroinformatics*.
<https://doi.org/10.3389/fninf.2016.00002>

Campagne Laboratory

institute for **computational biomedicine**

Weill Cornell Medical College



Search here...



Laboratory focus

Blog

Lab Members

Publications

Software

Opportunities

PI Contact

RSS

Nextflow Workbench



Helping biologists and clinicians learn how to write or reuse reproducible analysis pipelines

- [Nextflow Workbench Roadmap](#)
- [Installation instructions](#)
- [Workflow Execution](#)
 - [Executions on Cloud Cluster](#)
 - [Local Workflow Executions](#)
 - [Executions on Remote Cluster](#)
 - [Provision a Cloud Cluster](#)
- [Change Log](#)
- [Instructions for Workflow Tutorial](#)
- [Training](#)

>MCHU - Calmodulin

ADQLTEEQIAEFKEAFSLFDKDGDTITTKELGTVMRSLGQNPTEAELQDMINEVDADGNGTID
FPEFLTMMARKMKDTDSEEEIREAFRVFDKDGNGYISAAELRHVMTNLGEKLTDEEVDEMIREA
DIDGDGQVNYEEFVQMMTAK*



```
<no description>
Workflow workflowBash

with input:
  fastafile = [ ${org.campagnelab.workflow.home}/data/sample1.fasta ]

do:
  'input' = fastafile << ... >> splitSequence -> splitfile : 'seq_#' will run n times <<options>>;
  record = splitfile << ... >> reverse -> result : 'finaleres.txt' will run n times <<options>>;

and report:
  for each value of result << ... >> do: (c)->void {
    System.out.println("This workflow has produced this file: " + c.getAbsolutePath());
  }
```

WHO IS USING NEXTFLOW?



Weill Cornell Medical College

UT Southwestern
Medical Center

UCSF



SciLifeLab



UiO : University of Oslo



Version Control

Using Git

ISSUE:

RECENT UPDATE BROKE
SUPPORT FOR HARDWARE
I NEED FOR MY JOB.

WORKAROUND:

IF WE WAIT LONG ENOUGH,
THE EARTH WILL EVENTUALLY
BE CONSUMED BY THE SUN.



ChangeDetectionHub

ColorJudgementTask

Undo

Redo

Pull

Push

Branch

Stash

Pop

Viewing 7/7 Show All

Filter (Ctrl + Alt + f)

LOCAL1/1

ColorJudgementTask1

REMOTE6/6

origin

ColorJudgementTask

develop

Lab

master

py2to3

SubliminalCDT

PULL REQUESTS0

TAGS0/0

SUBMODULES0

develop

ColorJudgementTask

develop

Lab

master

py2to3

SubliminalCDT

Kotating color wheel

misc

removed pyc

new instructions for color wheel

fix

ignore pyc in subfolders

renamed subliminal to masking

Renamed subliminal to masking

added color wheel

Merge pull request #3 from cogmind/Lab

changed items

Main

Merge remote-tracking branch 'origin/Lab' into Lab

Fixed config

Fixed triggers for subliminal

Merge pull request #2 from cogmind/Lab

Merge branch 'Lab' of https://github.com/cogmind/Chan...

Merge remote-tracking branch 'origin/Lab' into Lab

fixed stimulus trigger

get_trigger now accepts frame

Added contrast to header

added contrast data to row_data

fixed trials

fixed items from 2 4 to 2 4 5

40 p trials

pycs

cmd line

Added cmd line option

Compiled

Fixed Data

Merge pull request #1 from cogmind/develop

Merge pull request #2 from cogmind/Lab

Lab

Daniel Labbé

commit: 5a0c78

authored 19.5.2016 @ 13:46

parent: 1892fa,206f52

15 modified

Name

Full Path

src/configuration.py

src/configuration.pyc

src/controller.py

src/controller.pyc

src/experiment_logic/factory_supplier.py

src/experiment_logic/factory_supplier.pyc

src/experiment_logic/trial.py

src/experiment_logic/trial.pyc

src/experiment_logic/triallogic.py

src/experiment_logic/triallogic.pyc

src/experiment_logic/trigger_manager.py

src/main.py

src/prepare_scenario.py

src/prepare_scenario.pyc

src/view.pyc

Version Control



Companies & Projects Using Git

Google

facebook

Microsoft

twitter

LinkedIn

NETFLIX

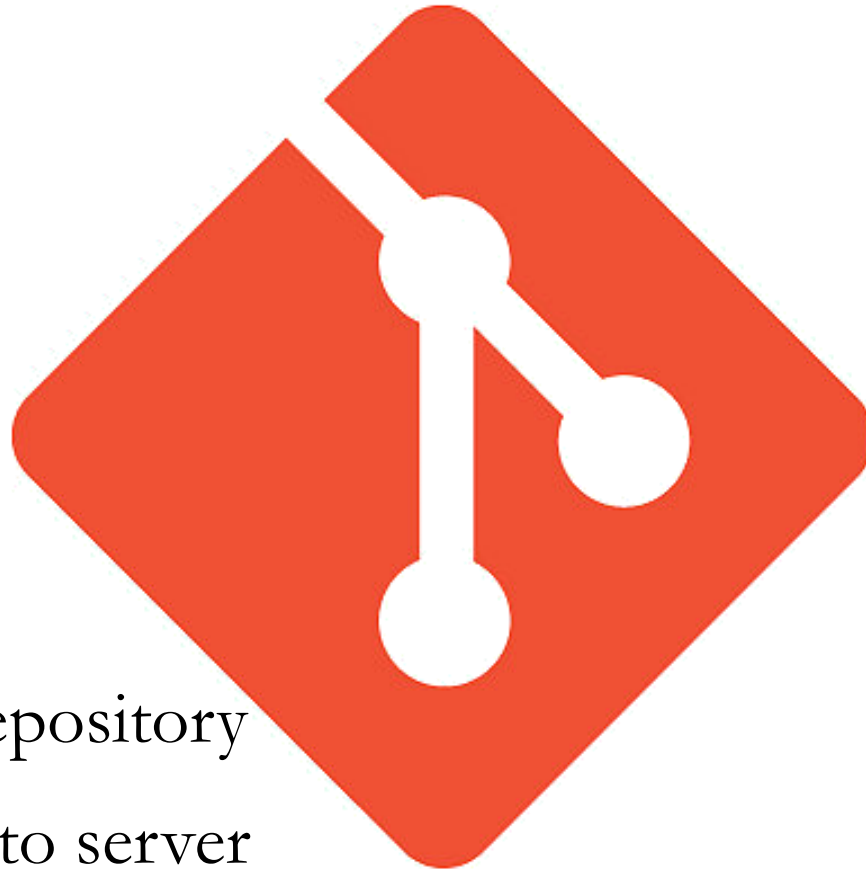


PostgreSQL



Git

- Enables
 - Versioning
 - Branching
 - Distributed code repository
 - Secure connection to server
 - Intermediate documentation for changes (commit messages)
- Accessible through Git client



GitHub



- GitHub is an online code repository service that uses Git
- The biggest of several providers (other examples: GitLab, Bitbucket)
- Open source projects are free
- Private repositories for a monthly fee
- GitHub can also be connected to the Open Science Framework (OSF), an open data platform for publishing data, code, and for managing collaborations

Open Science Framework

A scholarly commons to connect the entire research cycle



Lean Process Management

Using ZenHub

ZenHub

- A browser/API add-on for GitHub
- Suitable for science process management
- Scalable from project level to organization



Jet Propulsion Laboratory
California Institute of Technology



SONY

vmware



Panasonic



ZenHub

- Lean process management
 - Register tasks, assign people
 - Implements an advanced, and customizable, Kanban system
 - Set **milestones**, estimate relative effort/**story points**, and review progress in **burn-down charts**
- Shared planning
 - Collaborations
 - Individual projects (you can merge planning from two different repositories)
 - Team Neuroscience
- Horizontal knowledge transfer

Kanban System

The screenshot displays a GitHub Kanban board for the repository `cogmind / ChangeDetectionHub`. The interface includes a top navigation bar with links for Pull requests, Issues, Marketplace, Gist, and ToDo. Below this, the repository name and a search bar are visible. The main section shows a horizontal navigation bar with tabs for Code, Issues (6), Pull requests (0), Boards (selected), Reports, Projects (0), Wiki, and Insights. Below the navigation bar, there are several filter buttons: View, Repos (1/1), Show one, Labels, Milestones, Assignees, Epics, and Releases. A search bar and a 'New Issue' button are also present. The Kanban board itself consists of five columns: Icebox, Backlog, In Progress, Review/QA, and Done. Each column has a header indicating '0 Issues - 0 Story Points' and a vertical list of issues. The 'Icebox' column has a vertical label 'New Issues' on the left. The 'Done' column has a vertical label 'Closed' on the right. The 'Add' button is located at the bottom right of the board.

Repository: `cogmind / ChangeDetectionHub` (Private)

Navigation: Pull requests, Issues, Marketplace, Gist, ToDo

Repository Actions: Unwatch (2), Star (0), Fork (0)

Navigation Bar: <> Code, Issues (6), Pull requests (0), Boards (selected), Reports, Projects (0), Wiki, Insights

Filter Buttons: View, Repos (1/1), Show one, Labels, Milestones, Assignees, Epics, Releases

Search: Search

New Issue

Kanban Board Columns:

- Icebox: 0 Issues - 0 Story Points
- Backlog: 0 Issues - 0 Story Points
- In Progress: 0 Issues - 0 Story Points
- Review/QA: 0 Issues - 0 Story Points
- Done: 0 Issues - 0 Story Points

Vertical Labels:

- Icebox: New Issues
- Done: Closed

Add

ZenHub Example

- Case Study: Supervision of a BSc student
- Two repositories
 - Writing up
 - Analysis

<> Code

Issues 16

Pull requests 0

Boards

Reports

Projects 0

Wiki

Insights



View

Repos (1/2)

Show all

Labels

Milestones

Assignees

Epics

Releases

Search

New Issue



8 Issues - 52 Story Points

Icebox



3 Issues - 0 Story Points

Backlog



3 Issues - 27 Story Points

In Progress



1 Issue - 2 Story Points

Review/QA




1 Issue - 0 Story Points

Done




Closed


7+ Issues - 3 Story Points

 bsc_Britta_Hardwigen #21
Anmeldung BA bis zum ??

1 deliverable

 bsc_Britta_Hardwigen #18
"Das Verfassen wissenschaftlicher Texte"

Gold Plating

 bsc_Britta_Hardwigen #4
Ergebnisse - First draft
Ergebnisse

21 deliverable

 bsc_Britta_Hardwigen #5
Diskussion - First draft
Diskussion


21 deliverable

 bsc_Britta_Hardwigen #6
Abstract - First draft
Abstract


2 deliverable

 bsc_Britta_Hardwigen #22
Literatur
Anhang

3 deliverable

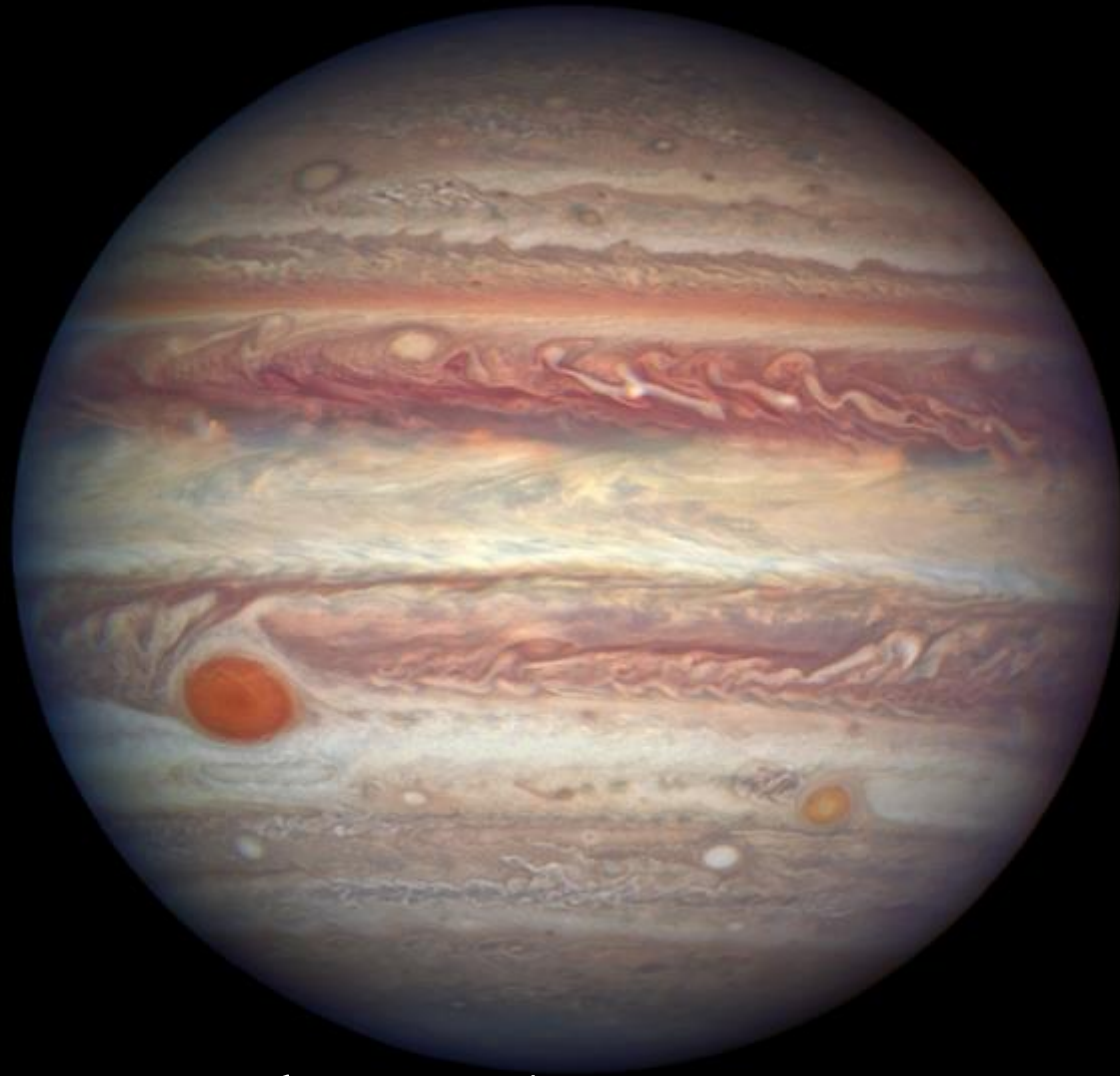
 bsc_Britta_Hardwigen #7
Anhang bsc_Britta_Hardwigen #9
Config - Location
Methodenteil bsc_Britta_Hardwigen #15
[Colors] Alpha, Gamma bsc_Britta_Hardwigen #20
Screenshot(s) for Methods
Methodenteilapa ftw deliverable help wanted
Priority 2 simplify wontfix bsc_Britta_Hardwigen #2
Einleitung - First draft
Einleitung

21 deliverable discuss Priority 1

 bsc_Britta_Hardwigen #3
Methoden - First Draft
Methodenteil5
American Psychological Association
Manual of Style?
apa ftw APA missing apa?
help wanted more APA
needs more APA Priority 1 bsc_Britta_Hardwigen #23
Distraktor? Interferenz? - Wort
gesucht!1 discuss help wanted Priority 1
question bsc_Britta_Hardwigen #16
Offtopic: Bescheinigung
Praktikum
Einleitung

2 deliverable Priority 1

 bsc_Britta_Hardwigen #17
Einleitung - Literatur
Einleitungdiscuss help wanted Priority 1
question



Digital Notebooks

Enabling Knowledge Transfer



Currently in use at

Google

Microsoft

IBM

Bloomberg

O'REILLY

CONTINUUM
ANALYTICS

Rackspace
the managed cloud company

SOUNDCloud



NetApp

software
carpentry

hhmi janelia
Research Campus

<CODE NEURO>

N-Site LLC

COCA.LC

BRYN
MAWR
COLLEGE

CAL POLY
SAN LUIS OBISPO

Berkeley
UNIVERSITY OF CALIFORNIA

The
University
Of
Sheffield

THE GEORGE
WASHINGTON
UNIVERSITY
WASHINGTON, DC

CLEMSON
UNIVERSITY

MICHIGAN STATE
UNIVERSITY

Northwestern
University

NYU

NASA

IPython
Interactive Computing


```
D:\analysis\alpha\eeg>cd mne
```

```
D:\analysis\alpha\eeg\mne>dir
```

```
Volume in Laufwerk D: hat keine Bezeichnung.
```

```
Volumeseriennummer: 860C-4E82
```

```
Verzeichnis von D:\analysis\alpha\eeg\mne
```

```
2017-06-26 16:27 <DIR> .
2017-06-26 16:27 <DIR> ..
2017-06-26 16:59          922 mne_start.py
                1 Datei(en),          922 Bytes
                2 Verzeichnis(se), 471,515,373,568 Bytes frei
```

```
D:\analysis\alpha\eeg\mne>jupyter notebook
```

```
[I 14:17:10.352 NotebookApp] [nb_conda_kernels] enabled, 7 kernels found
```

```
[W 14:17:14.900 NotebookApp] Error loading server extension nbpresent
```

```
Traceback (most recent call last):
```

```
File "c:\Anaconda2\lib\site-packages\notebook\notebookapp.py", line 1046, in init_server_extensions
```

```
    mod = importlib.import_module(modulename)
```

```
File "c:\Anaconda2\lib\importlib\__init__.py", line 37, in import_module
```

```
    __import__(name)
```

```
File "c:\Anaconda2\lib\site-packages\nbpresent\__init__.py", line 5, in <module>
```

```
    from nbconvert.exporters.export import exporter_map
```

```
File "c:\Anaconda2\lib\site-packages\nbconvert\__init__.py", line 4, in <module>
```

```
    from .exporters import *
```

```
File "c:\Anaconda2\lib\site-packages\nbconvert\exporters\__init__.py", line 1, in <module>
```

```
    from .export import *
```

```
File "c:\Anaconda2\lib\site-packages\nbconvert\exporters\export.py", line 9, in <module>
```

```
    import entrypoints
```

```
File "c:\Anaconda2\lib\site-packages\entrypoints.py", line 16, in <module>
```

```
    from backports import configparser
```

```
ImportError: cannot import name configparser
```

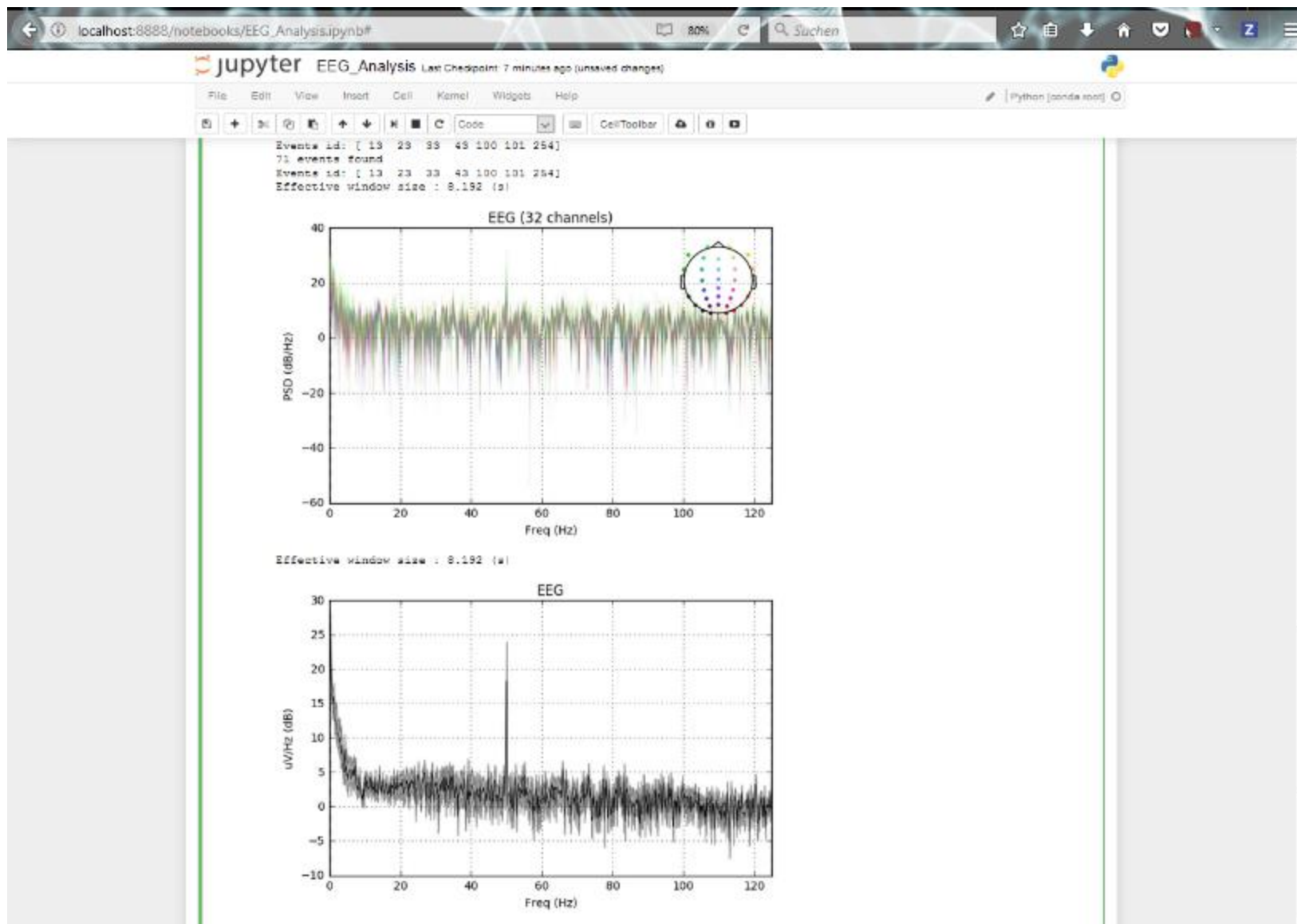
```
[I 14:17:14.961 NotebookApp] [nb_conda] enabled
```

```
[I 14:17:16.605 NotebookApp] [nb_anacondacloud] enabled
```

```
[I 14:17:17.344 NotebookApp] Serving notebooks from local directory: D:\analysis\alpha\eeg\mne
```

```
[I 14:17:17.345 NotebookApp] 0 active kernels
```

```
[I 14:17:17.345 NotebookApp] The Jupyter Notebook is running at: http://localhost:8888/
```



Branch: master

vector / eeg_analysis.ipynb

Find file

Copy path

cogmind Add files via upload

1e7aeed 21 days ago

1 contributor

389 lines (388 sloc) 11 KB

Raw

Blame

History



```
In [1]: # Binomatic
#
# Description= Generates ERPLAB binlister files
#
# Author= Daniel Labbé
# Date= 2017-06-09

# Based on https://github.com/lucklab/erplab/wiki/Assigning-Events-to-Bins-with-BINLISTER
#
```

```
In [2]: from itertools import count, product
```

```
In [22]: # GRAMMAR
_TAGS = '<' + '{}' + '>'

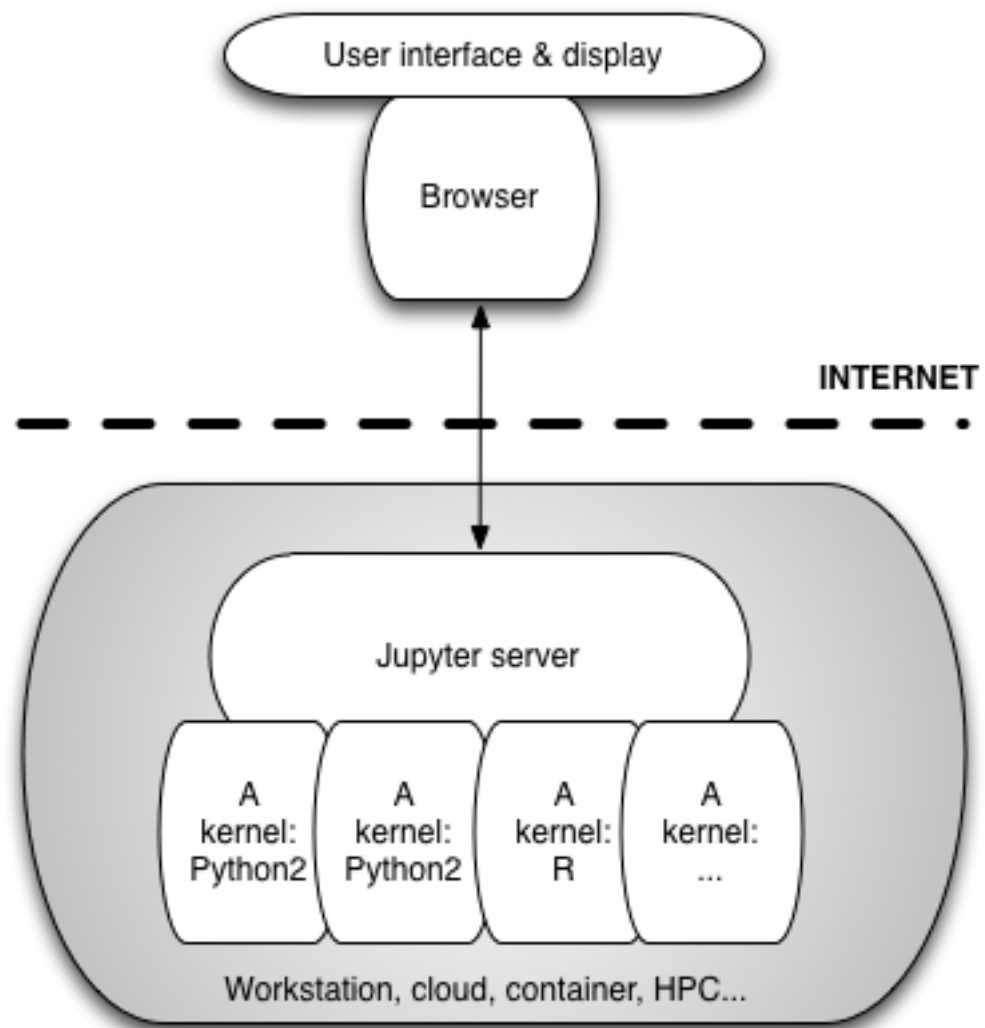
# Please note: "101;102" should be entered as a string and not a tuple

g = dict(
    EVENT = '{{{0}}}', # {} acts as escape for {
    PRE = '.' + '{}',
    POST = '{}' + '.',
    NOT = '~' + '{}',
    OR = '{}' + ';' + '{}',
    FROM_TO = '{}' + ':' + '{}',
    RANGE = '{}-{}',
    TAGS = _TAGS,
    TIME = 't' + _TAGS + '{}', # 't<' + '{}' + '>' + '{}'
    RT = '{}' + ':rt' + _TAGS,
    WA = '{}' + ':wa' + _TAGS,
    FA = '{}' + ':fa' + _TAGS,
    WB = '{}' + ':wb' + _TAGS,
    FB = '{}' + ':fb' + _TAGS
)
```

```
In [28]: def parse(stimulus, syntax):
    #print(type(stimulus))
    assert isinstance(stimulus, str) or isinstance(stimulus, int) or isinstance(stimulus, tuple)
    e)
    assert isinstance(syntax, str)

    if isinstance(stimulus, str):
        p = g[syntax].format(stimulus)
    elif isinstance(stimulus, tuple):
```





Discussion

Technical Debt

DEFINITION

The extra work required to bring a rushed piece of code up to standard with current practises or your own understanding of the code.

First coined by Ward Cunningham, inventor of the wiki.

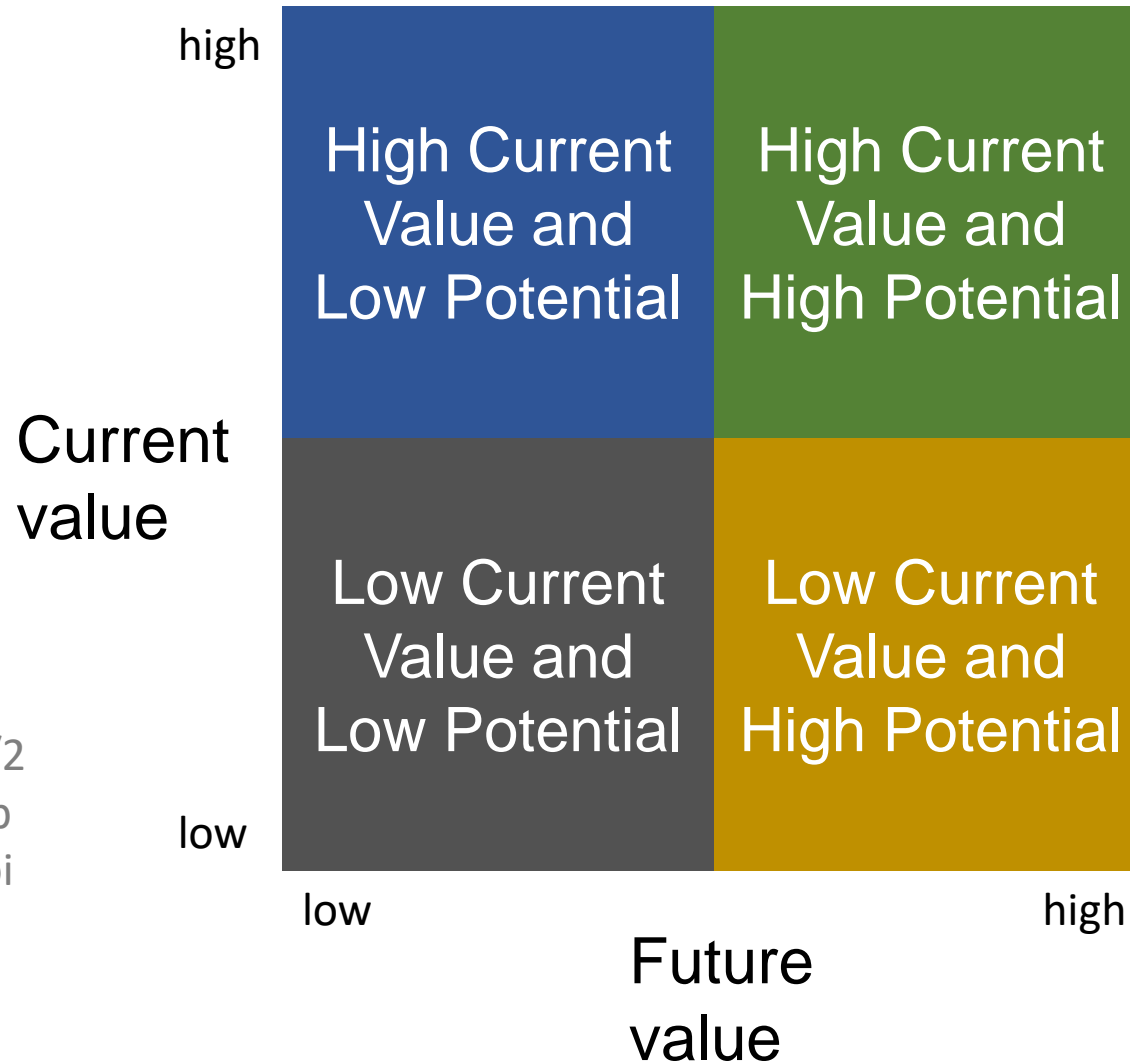
vs.

YAGNI

You ain't gonna need it...

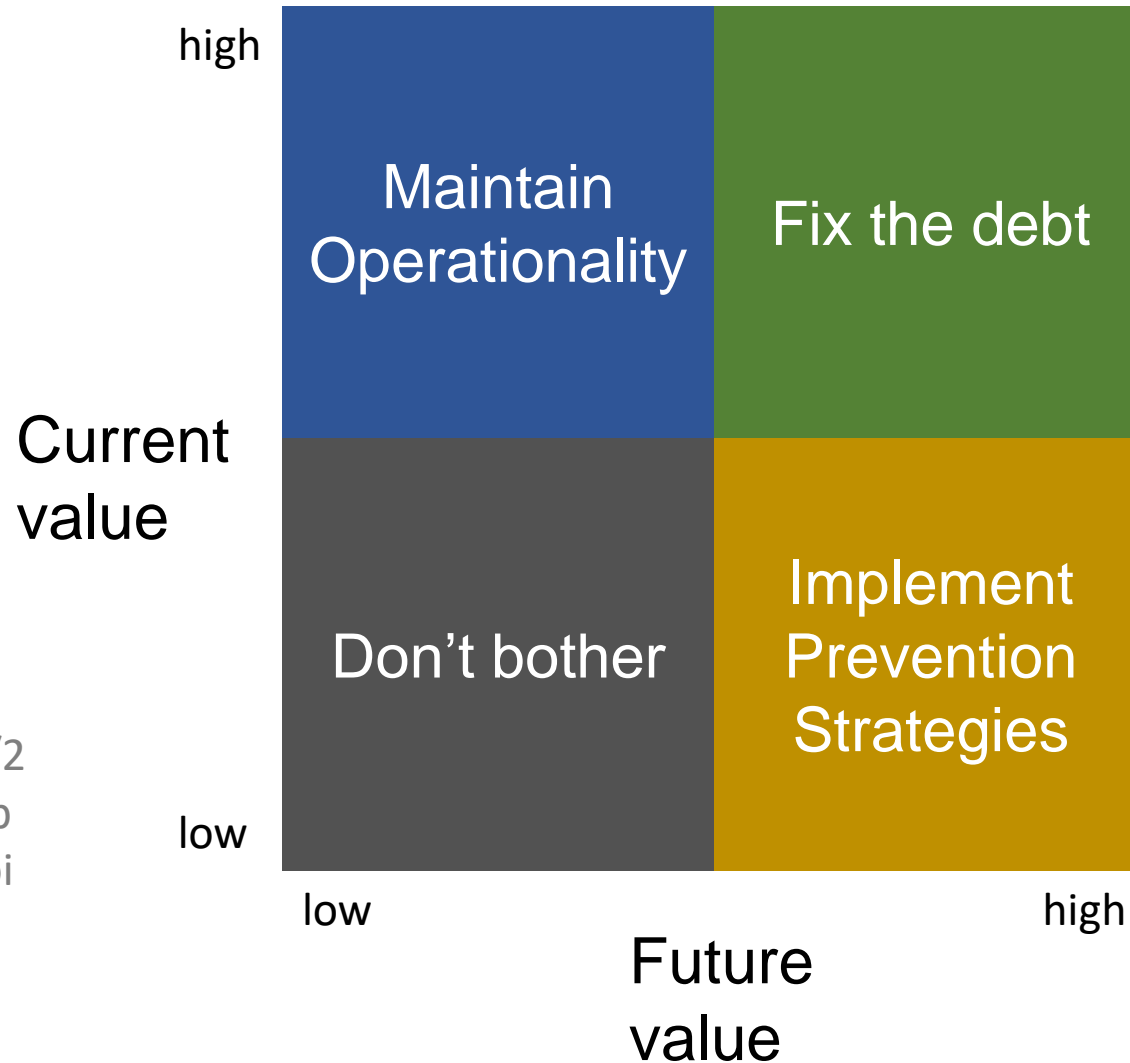


A Method to Determine when to Adapt Technology (Refactoring) by **Value**



<http://blogs.ripple-rock.com/SteveGarnett/2013/03/05/TechnicalDebtStrategiesTacticsForAvoidingRemovingIt.aspx>

A Method to Determine when to Adapt Technology (Refactoring) by **Value**



High Current Value and Low Potential	High Current Value and High Potential
Low Current Value and Low Potential	Low Current Value and High Potential

<http://blogs.ripple-rock.com/SteveGarnett/2013/03/05/TechnicalDebtStrategiesTacticsForAvoidingRemovingIt.aspx>

Six Open Questions

- Technical debt. Should we adapt?
- Are our efforts future proof?
- What are the specific challenges for cognitive neuroscience when it comes to sharing data and open data?
- How can cognitive neuroscientists improve collaborations with computational and basic/systems neuroscientists?
- What are the consequences for neuroscience education?
- What is our Mars expedition?

Main Literature

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