



# HIGH PERFORMANCE SCIENCE

**Steffen Bollmann**

open and reproducible  
neuroscience image  
processing that scales?

# FROM DATA TO PUBLICATION



# WHAT CAN POSSIBLY GO WRONG?

- Scientists write software, but are not trained in software development
- There will be bugs –tricky to find if plausible results



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@sbollmann\_MRI

Steffen.Bollmann@cai.uq.edu.au



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# SELECTIVE DEBUGGING



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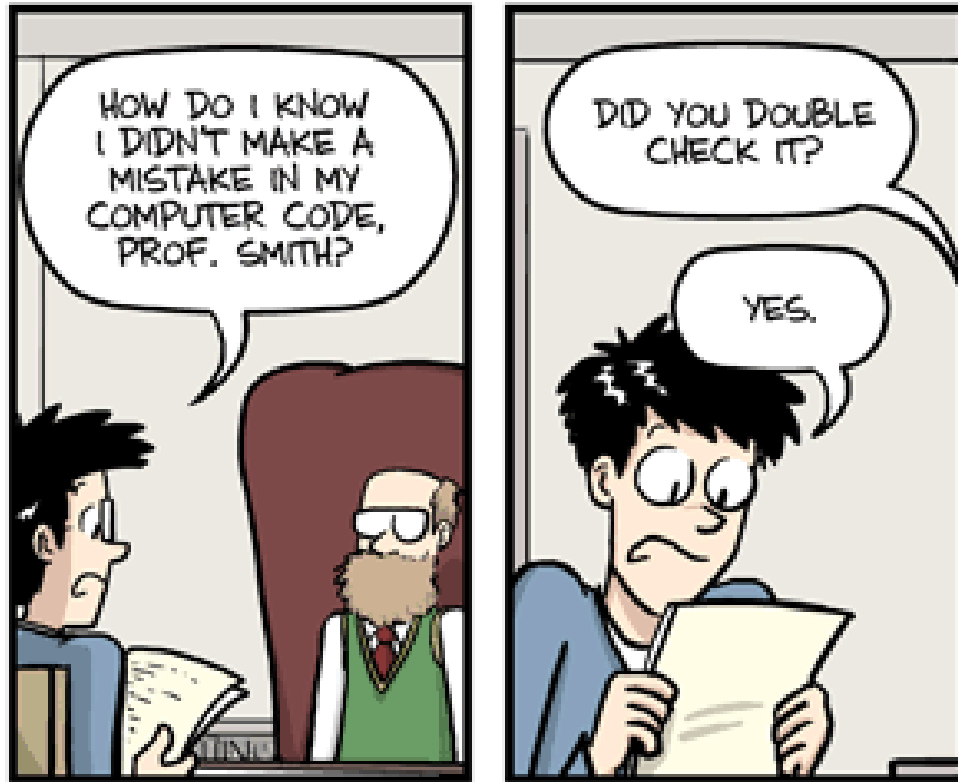
Steffen.Bollmann@cai.uq.edu.au



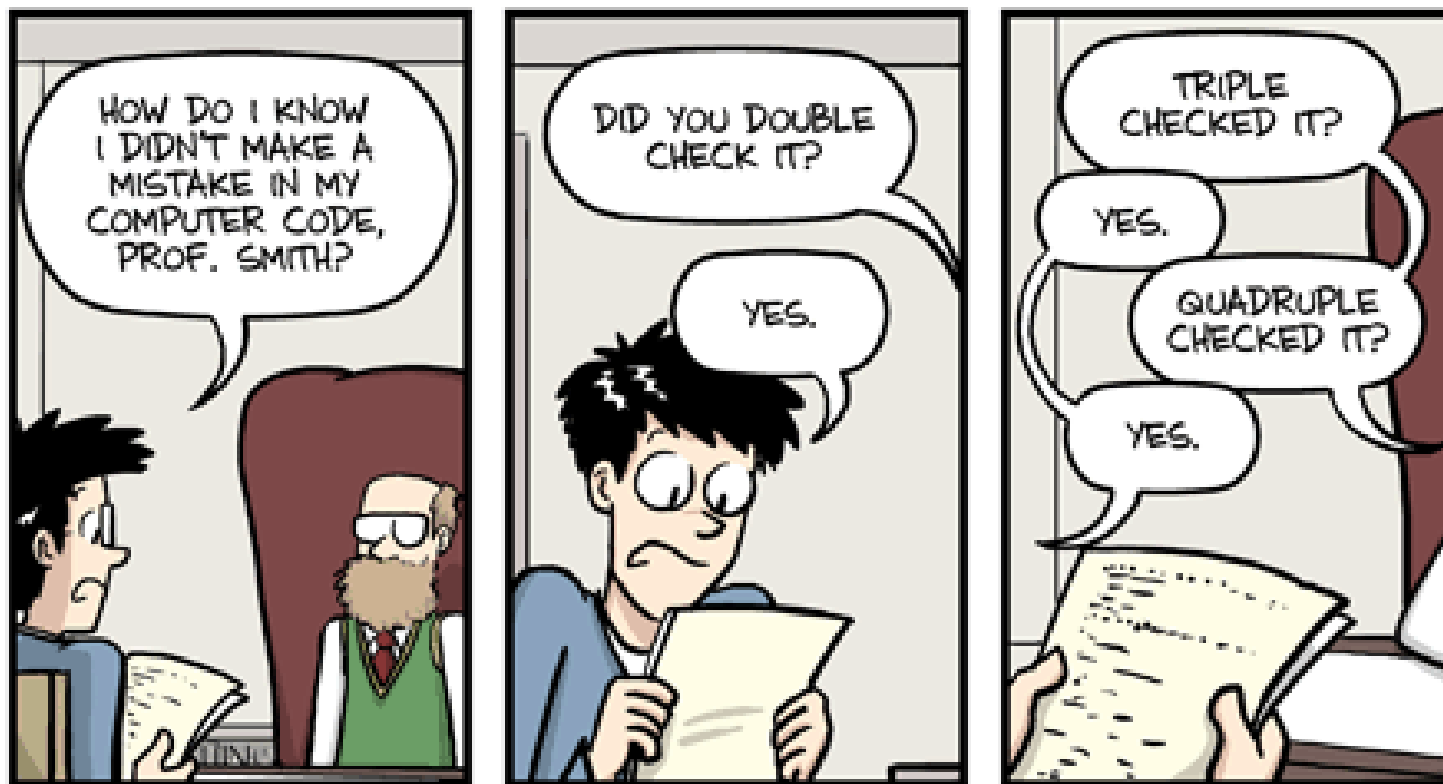
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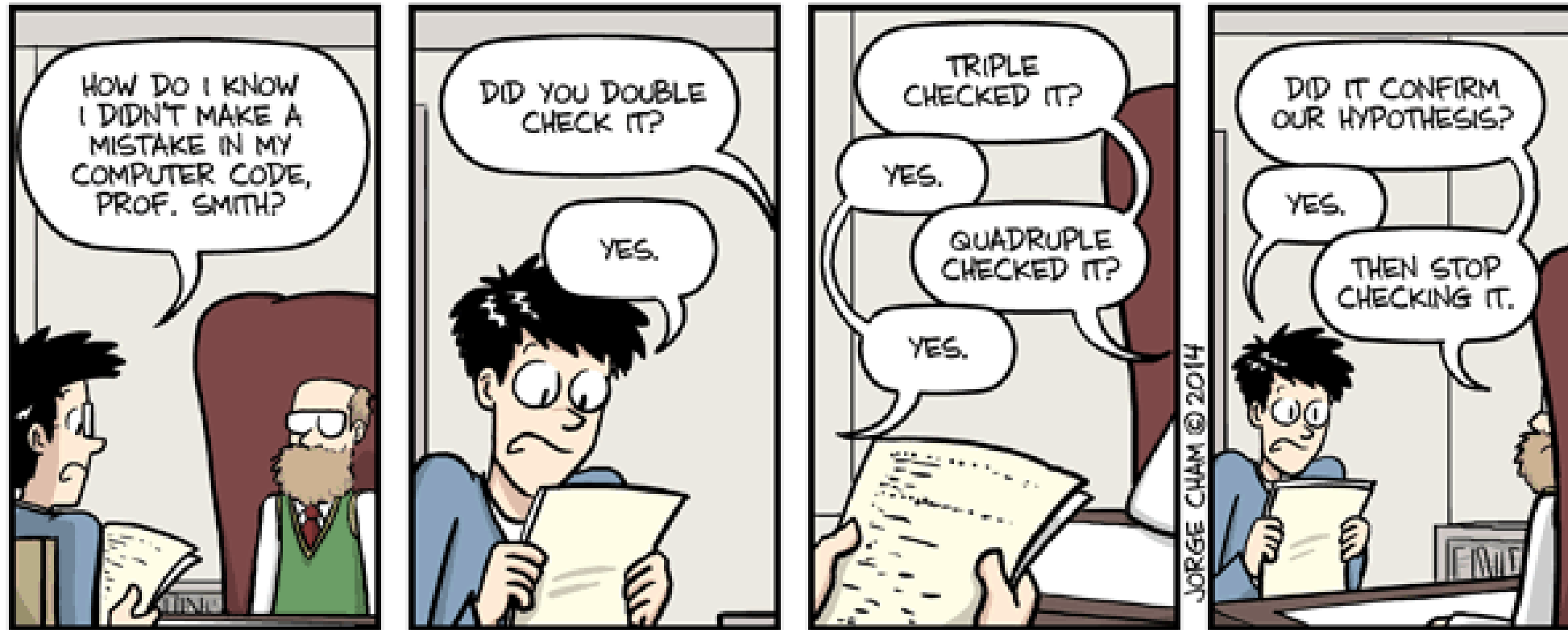
# SELECTIVE DEBUGGING



# SELECTIVE DEBUGGING



# SELECTIVE DEBUGGING



# WHAT CAN POSSIBLY GO WRONG?

- Scientists write software, but are not trained in software development
- There will be bugs –tricky to find if plausible results
- Large amounts of data
- Analyses run on different hardware and operating systems



# GLIBC 2.5 VS 2.18

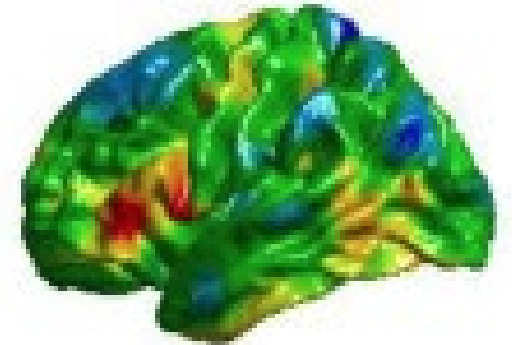
## Reproducibility of neuroimaging analyses across operating systems

*Tristan Glatard<sup>1,2</sup>, Lindsay B. Lewis<sup>1</sup>, Rafael Ferreira da Silva<sup>3</sup>, Reza Adalat<sup>1</sup>, Natacha Beck<sup>1</sup>, Claude Lepage<sup>1</sup>, Pierre Rioux<sup>1</sup>, Marc-Etienne Rousseau<sup>1</sup>, Tarek Sherif<sup>1</sup>, Ewa Deelman<sup>3</sup>, Najmeh Khalili-Mahani<sup>1</sup> and Alan C. Evans<sup>1\*</sup>*

```
expf(1.540518522262573242187500000000)  
=4.6670093536376953125000
```

```
expf(1.540518522262573242187500000000)  
=4.6670098304748535156250
```

- glibc 2.5 vs 2.18 deliver different floating -point results
- leads to significant differences in long pipelines



# WHAT CAN POSSIBLY GO WRONG?

- Scientists write software, but are not trained in software development
- There will be bugs –tricky to find if plausible results
- Large amounts of data
- Analyses run on different hardware and operating systems
- Difficult to share data and reproduce an analysis



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Essay

## Why Most Published Research Findings Are False

John P. A. Ioannidis

## Cluster failure: Why fMRI inferences for spatial extent have inflated false-positive rates

Anders Eklund<sup>a,b,c,1</sup>, Thomas E. Nichols<sup>d,e</sup>, and Hans Knutsson<sup>a,c</sup>

## RESEARCH ARTICLE SUMMARY

PSYCHOLOGY

## Estimating the reproducibility of psychological science

Open Science Collaboration\*



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# OUR JOURNEY BEGINS



<https://cai.centre.uq.edu.au/facilities/human-imaging/3t-magnetom-prisma>



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Steffen.Bollmann@cai.uq.edu.au

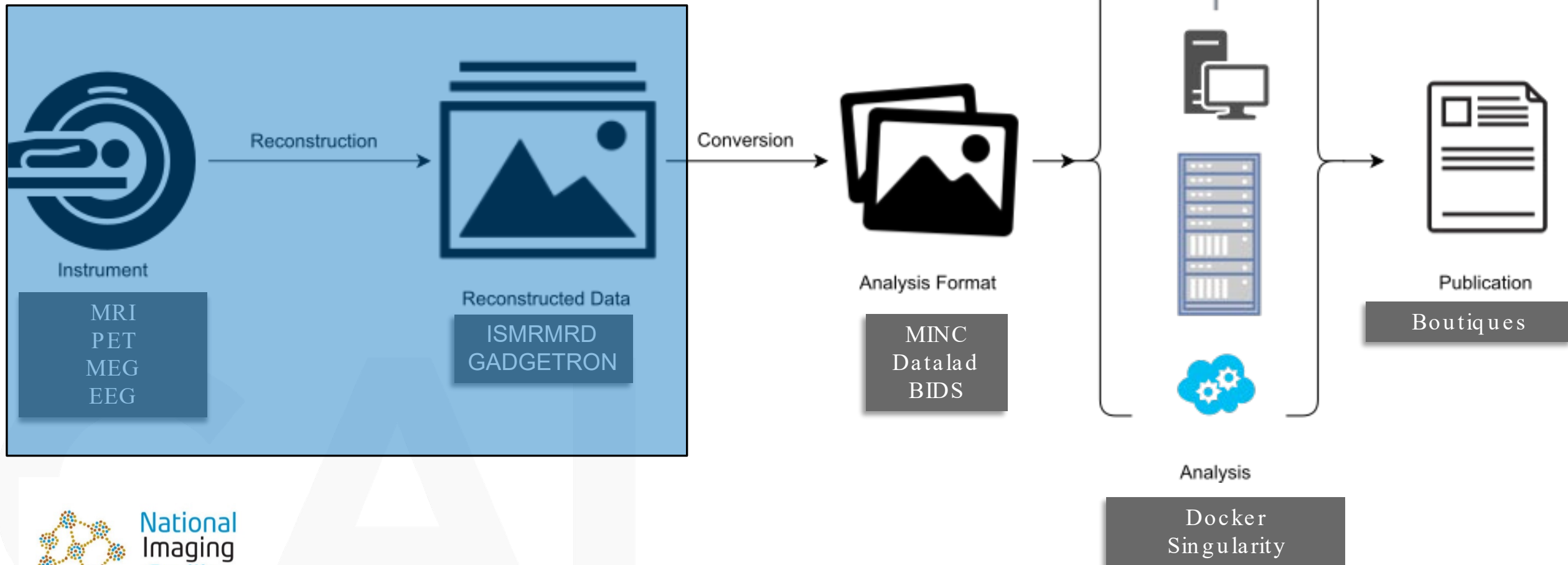


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# FROM DATA TO PUBLICATION



# ISMRRMRD & GADGETRON

## ISMRRMRD Raw Data Format: A Proposed Standard for MRI Raw Datasets

Souheil J. Inati,<sup>1</sup> Joseph D. Naegele,<sup>1</sup> Nicholas R. Zwart,<sup>2</sup> Vinai Roopchansingh,<sup>1</sup> Martin J. Lizak,<sup>3</sup> David C. Hansen,<sup>4</sup> Chia-Ying Liu,<sup>5</sup> David Atkinson,<sup>6</sup> Peter Kellman,<sup>7</sup> Sebastian Kozerke,<sup>8</sup> Hui Xue,<sup>7</sup> Adrienne E. Campbell-Washburn,<sup>7</sup> Thomas S. Sørensen,<sup>9</sup> and Michael S. Hansen<sup>7\*</sup>

### ISMRRMRD

- ISMRRMRD: Open format based on HDF5 + routines for converting from any closed source vendor format

<https://github.com/ismrmd>



## Gadgetron: An Open Source Framework for Medical Image Reconstruction

Michael Schacht Hansen<sup>1\*</sup> and Thomas Sangild Sørensen<sup>2,3</sup>

### Gadgetron

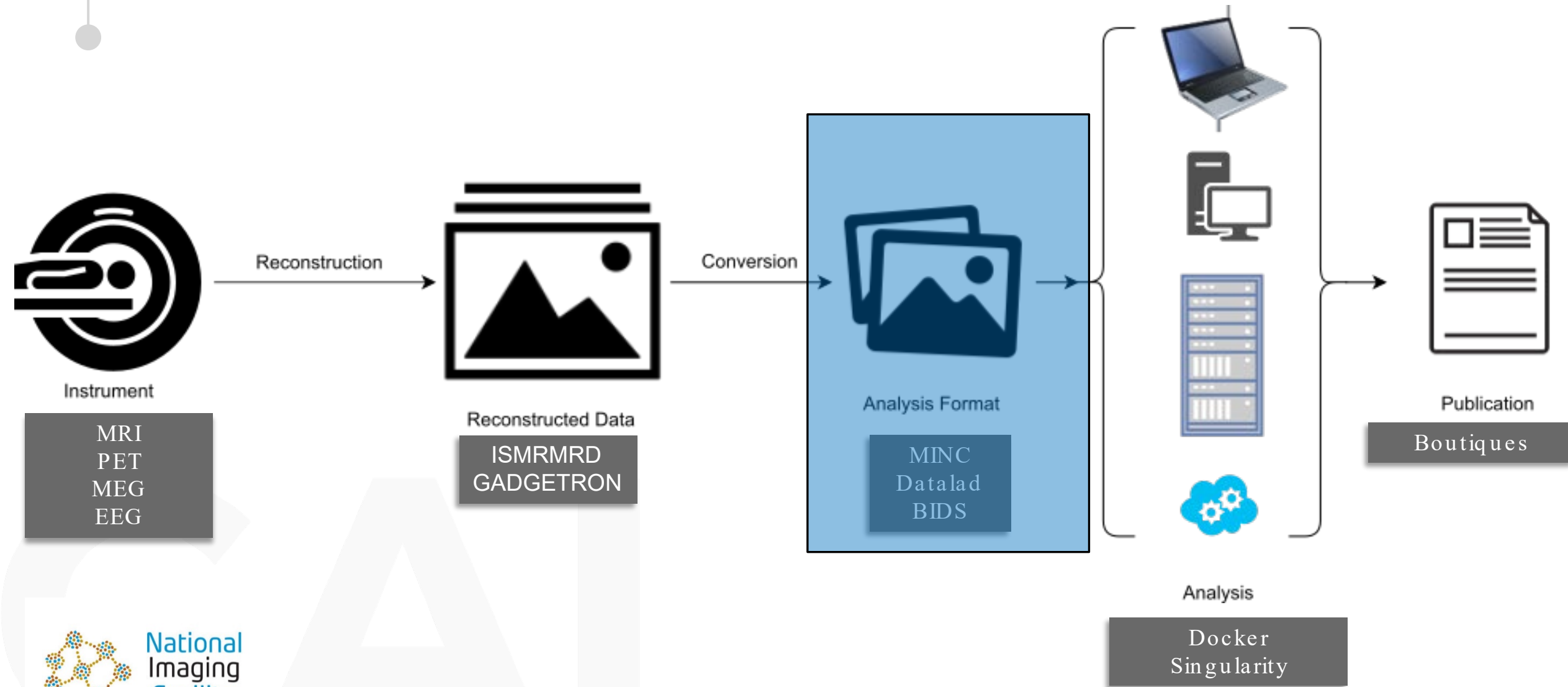
- Gadgetron : Open source reconstruction system compatible with ISMRRMRD

<https://github.com/gadgetron>



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# METROPOLITAN DATA CACHING INFRASTRUCTURE (MEDICI)



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<https://rcc.uq.edu.au/data> -storage

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# BRAIN IMAGING DATA STRUCTURE

**OPEN**

SUBJECT CATEGORIES

- » Data publication and archiving
- » Research data

## The brain imaging data structure, a format for organizing and describing outputs of neuroimaging experiments

Received: 18 December 2015

Accepted: 19 May 2016

Published: 21 June 2016

Krzysztof J. Gorgolewski<sup>1</sup>, Tibor Auer<sup>2</sup>, Vince D. Calhoun<sup>3,4</sup>, R. Cameron Craddock<sup>5,6</sup>, Samir Das<sup>7</sup>, Eugene P. Duff<sup>8</sup>, Guillaume Flandin<sup>9</sup>, Satrajit S. Ghosh<sup>10,11</sup>, Tristan Glatard<sup>7,12</sup>, Yaroslav O. Halchenko<sup>13</sup>, Daniel A. Handwerker<sup>14</sup>, Michael Hanke<sup>15,16</sup>, David Keator<sup>17</sup>, Xiangrui Li<sup>18</sup>, Zachary Michael<sup>19</sup>, Camille Maumet<sup>20</sup>, B. Nolan Nichols<sup>21,22</sup>, Thomas E. Nichols<sup>20,23</sup>, John Pellman<sup>6</sup>, Jean-Baptiste Poline<sup>24</sup>, Ariel Rokem<sup>25</sup>, Gunnar Schaefer<sup>1,26</sup>, Vanessa Sochat<sup>27</sup>, William Triplett<sup>1</sup>, Jessica A. Turner<sup>3,28</sup>, Gaël Varoquaux<sup>29</sup> & Russell A. Poldrack<sup>1</sup>



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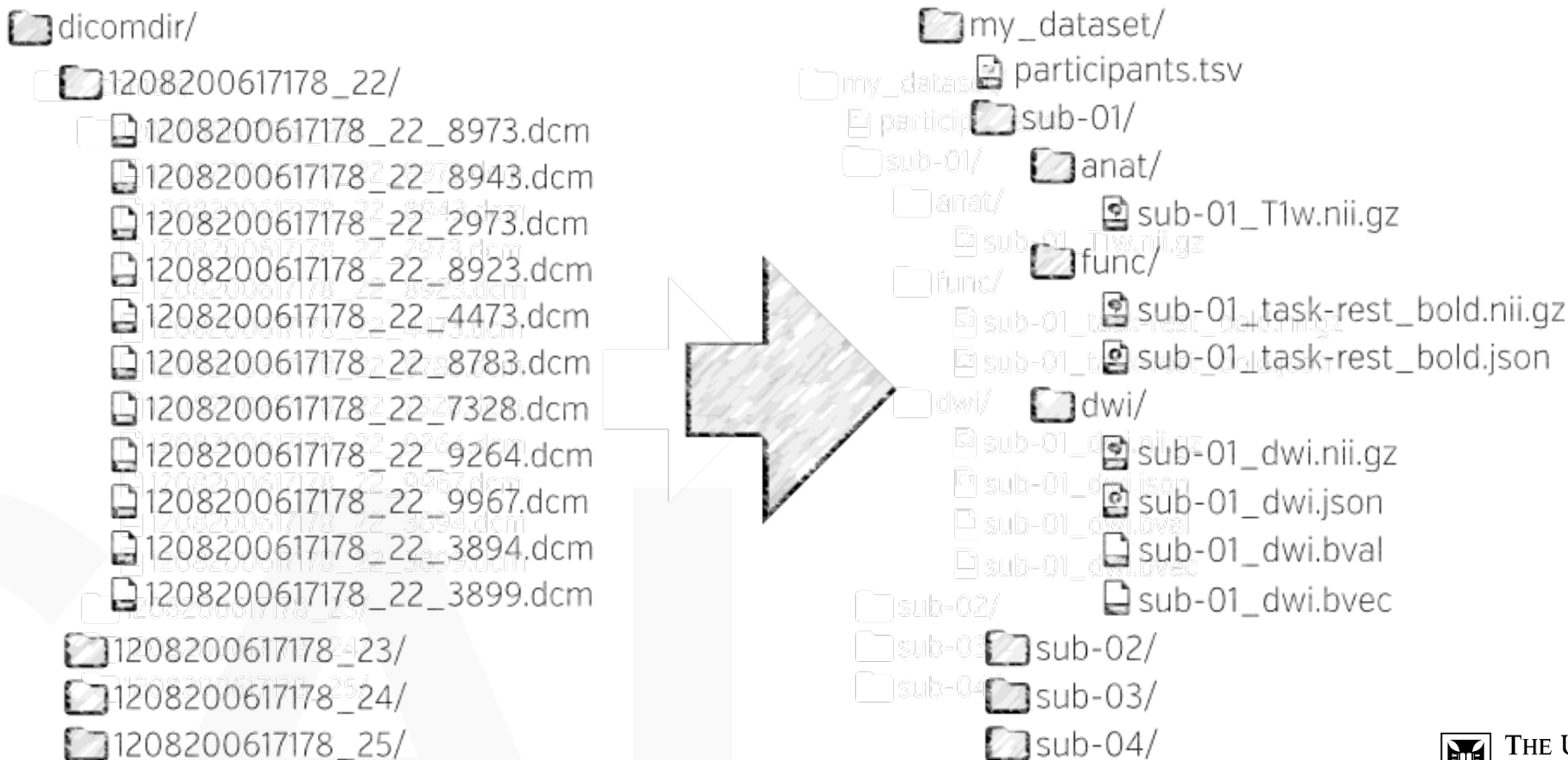


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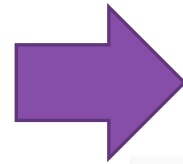
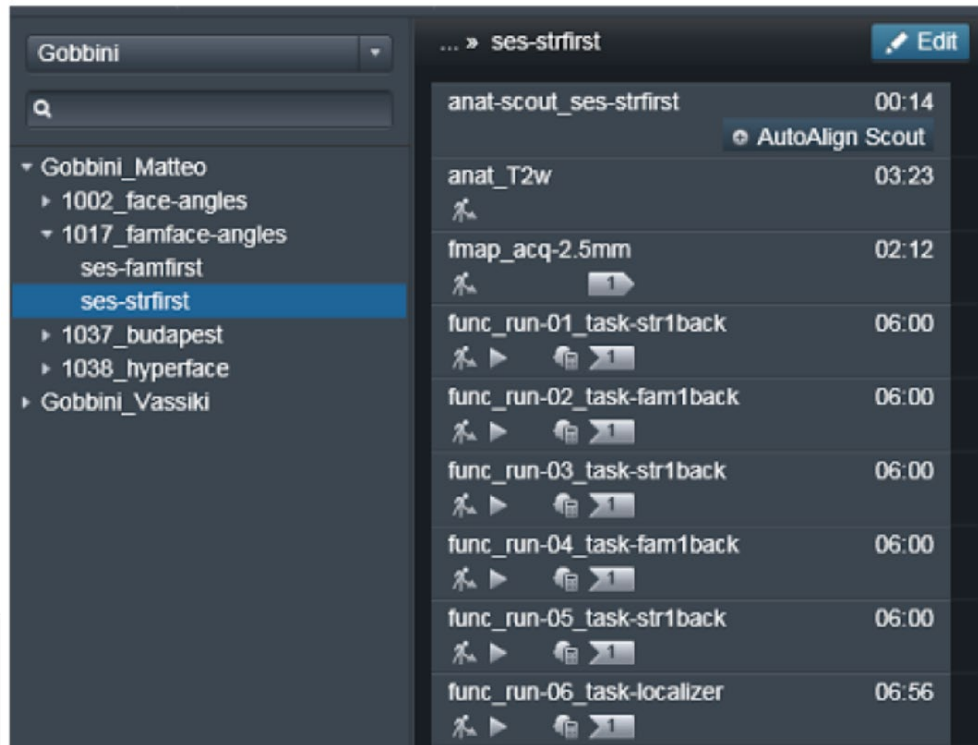


# BRAIN IMAGING DATA STRUCTURE



# REPROIN

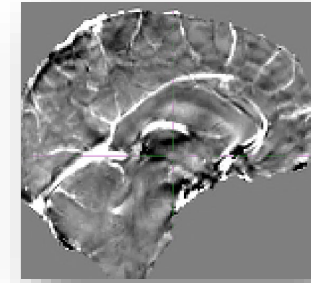
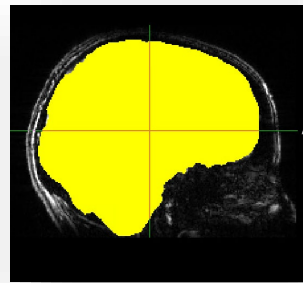
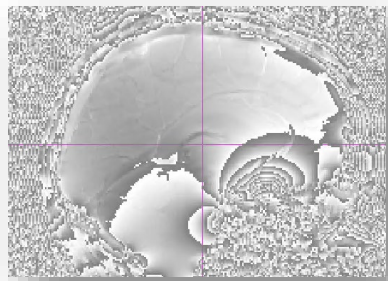
- setting up protocols on scanner to enable automatic conversion to bids



```
anat
  sub-sid000005_ses-strfirst_T2w.json
  sub-sid000005_ses-strfirst_T2w.nii.gz
fmap
  sub-sid000005_ses-strfirst_acq-25mm_magnitude1.json
  sub-sid000005_ses-strfirst_acq-25mm_magnitude1.nii.gz
...
func
  sub-sid000005_ses-strfirst_task-fam1back_run-02_bold.json
  sub-sid000005_ses-strfirst_task-fam1back_run-02_bold.nii.gz
  sub-sid000005_ses-strfirst_task-fam1back_run-02_events.tsv
...
sub-sid000005_ses-strfirst_scans.tsv
```

# DATA PROVENANCE

- Data Provenance / Lineage = keeping track of what happens to data during an analysis



```
[fsluser@localhost data]$ bet mag.nii.gz mask -n -m -R -f 0.1 -g 0.0
```

```
[fsluser@localhost TGVQSM-master-011045626121baa8bfdd6633929974c732ae35e3]$  
tgv_qsm -p ../phase.nii.gz -m ../mask_mask.nii.gz -f 2.89 -t 0.02 -s -o qsm
```

# MINC: BUILD -IN PROVENANCE

## MINC 2.0: A Flexible Format for Multi-Modal Images

*Robert D. Vincent<sup>1</sup>, Peter Neelin<sup>2</sup>, Najmeh Khalili-Mahani<sup>1</sup>, Andrew L. Janke<sup>3</sup>, Vladimir S. Fonov<sup>1</sup>, Steven M. Robbins<sup>1</sup>, Leila Baghdadi<sup>4</sup>, Jason Lerch<sup>4,5</sup>, John G. Sled<sup>4,5</sup>, Reza Adalat<sup>1</sup>, David MacDonald<sup>6</sup>, Alex P. Zijdenbos<sup>7</sup>, D. Louis Collins<sup>1,8</sup> and Alan C. Evans<sup>1\*</sup>*

```
$ minchistory FA.reg.mnc
--- History of B027915.01_DTI.B0.reg.clp.mnc ---
[01] Thu Nov 21 14:28:14 2002>>> mincaverage -clobber \
    /usr/people/steve/data/stroke/B027915/01_DTI/B027915.01_DTI.frame001.mnc \
    /usr/people/steve/data/stroke/B027915/01_DTI/B027915.01_DTI.frame002.mnc \
    /usr/people/steve/data/stroke/B027915/01_DTI/B027915.01_DTI.frame003.mnc \
    /usr/people/steve/data/stroke/B027915/01_DTI/B027915.01_DTI.frame008.mnc \
    /usr/people/steve/data/stroke/B027915/01_DTI/B027915.01_DTI.B0.mnc
[02] Mon Nov 3 16:54:24 2003>>> mincresample -clobber -like \
    /home/rotor/data/stroke/B027915/01_DTI/B027915.01_DTI.B0.mnc \
    -transformation \
    /home/rotor/data/stroke/B027915/xfms/B027915.01_DTI.midline-align.xfm \
    /home/rotor/data/stroke/B027915/01_DTI/B027915.01_DTI.B0.mnc \
    /home/rotor/data/stroke/B027915/reg/B027915.01_DTI.B0.reg.mnc
[03] Mon Nov 10 23:50:21 2003>>> minccalc -clobber -outfile value \
```

<https://bic-mni.github.io/>

Steffen.Bollmann@cai.uq.edu.au



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# GIT | GIT-ANNEX | DATALAD

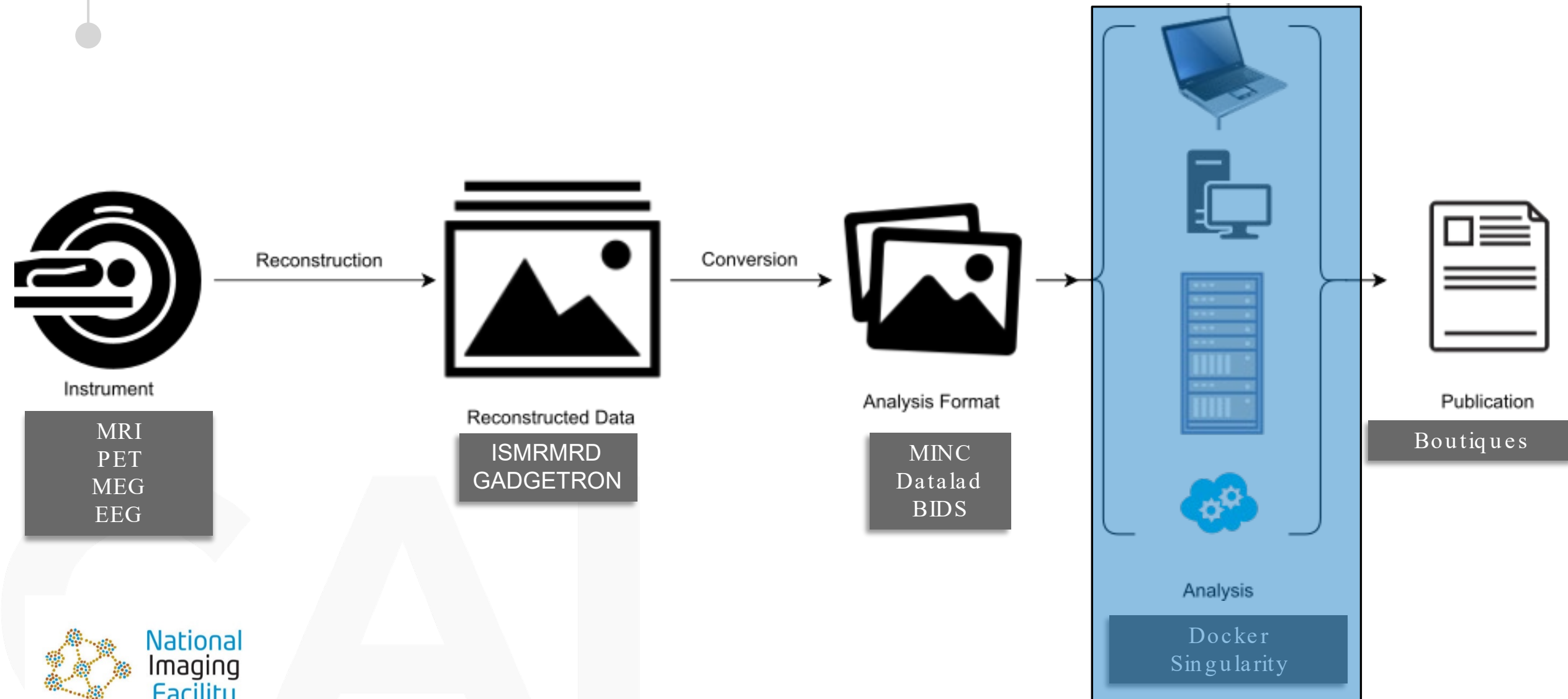
Feature	Git	git-annex	DataLad
Version control (text, code)	✓	✓	✓
Version control (binary data)	(not advised)	✓	✓
Auto-crawling available resources		✓ RSS feeds	✓ flexible
Unified dataset handling			✓
<ul style="list-style-type: none"> <li>• recursive operation on datasets</li> </ul>			✓
<ul style="list-style-type: none"> <li>• seamless operation across datasets boundaries</li> </ul>			✓
<ul style="list-style-type: none"> <li>• metadata support</li> </ul>		✓ per-file	✓
<ul style="list-style-type: none"> <li>• metadata aggregation</li> </ul>			✓ flexible
Unified authentication interface			✓

<http://datalad.org/for/git> -users



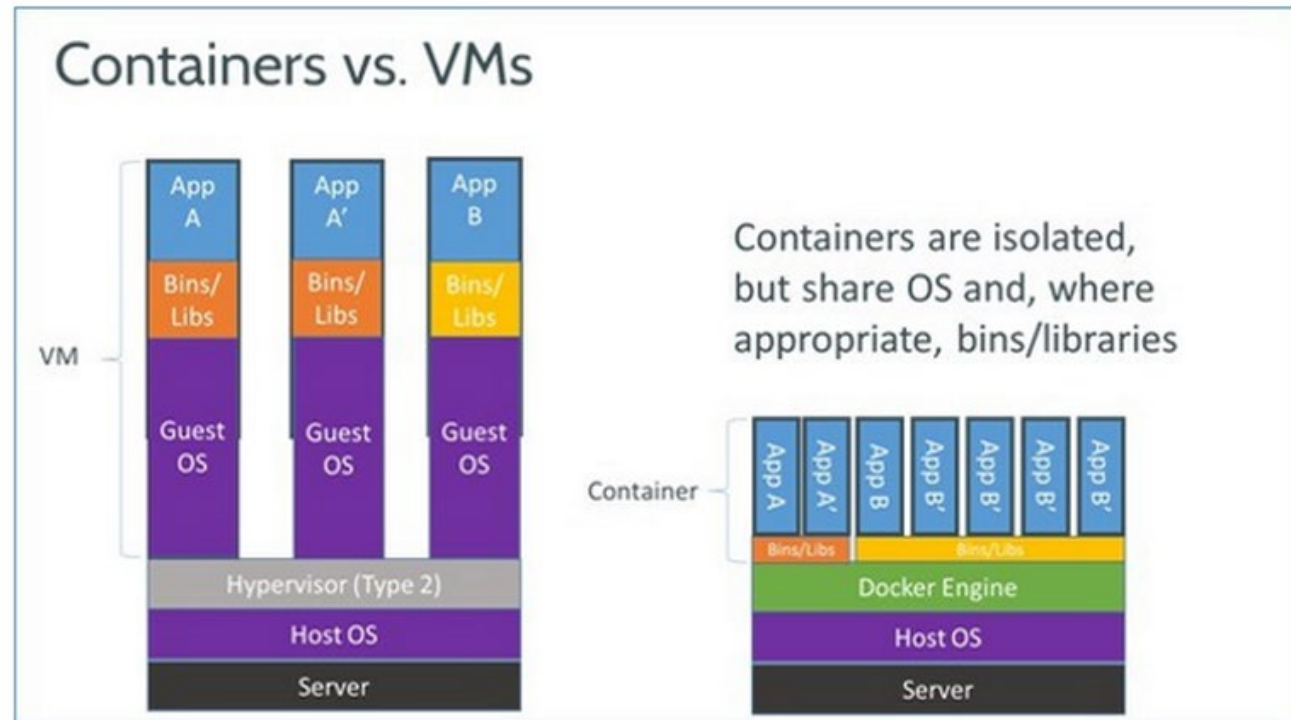
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# FROM DATA TO PUBLICATION



# DOCKER

- Containers are lighter and allow easier sharing of analysis pipelines
- Neurodocker and ReproZip help with building and deploying imaging pipelines
- But: Security on shared HPC systems is a problem



<http://www.zdnet.com/article/what-so-darn-popular/>

-is-docker -and -why -is-it-

<https://github.com/kaczmarj/neurodocker>

# SINGULARITY

RESEARCH ARTICLE

## Singularity: Scientific containers for mobility of compute

Gregory M. Kurtzer<sup>1</sup>, Vanessa Sochat<sup>2\*</sup>, Michael W. Bauer<sup>1,3,4</sup>

### Interactive Development

```
sudo singularity build --sandbox tmpdir/ Singularity
```

```
sudo singularity build --writable container.img Singularity
```

**BUILD ENVIRONMENT**

### Build from Recipe

```
sudo singularity build container.img Singularity
```

### Build from Singularity

```
sudo singularity build container.img shub://vsoch/hello-world
```

### Build from Docker

```
sudo singularity build container.img docker://ubuntu
```

### Container Execution

```
singularity run container.img  
singularity shell container.img  
singularity exec container.img ...
```

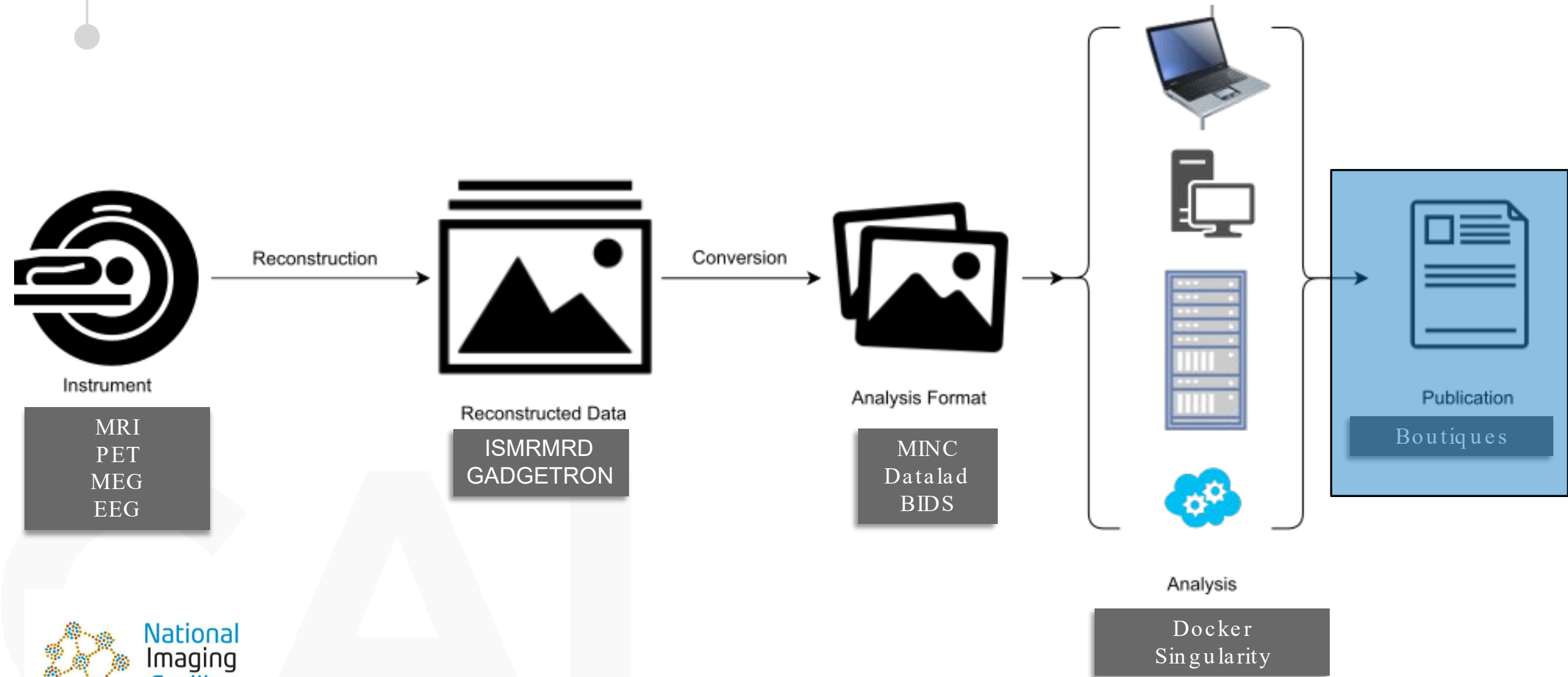
### Reproducible Sharing

```
singularity pull shub://...  
singularity pull docker://... *
```

**PRODUCTION ENVIRONMENT**

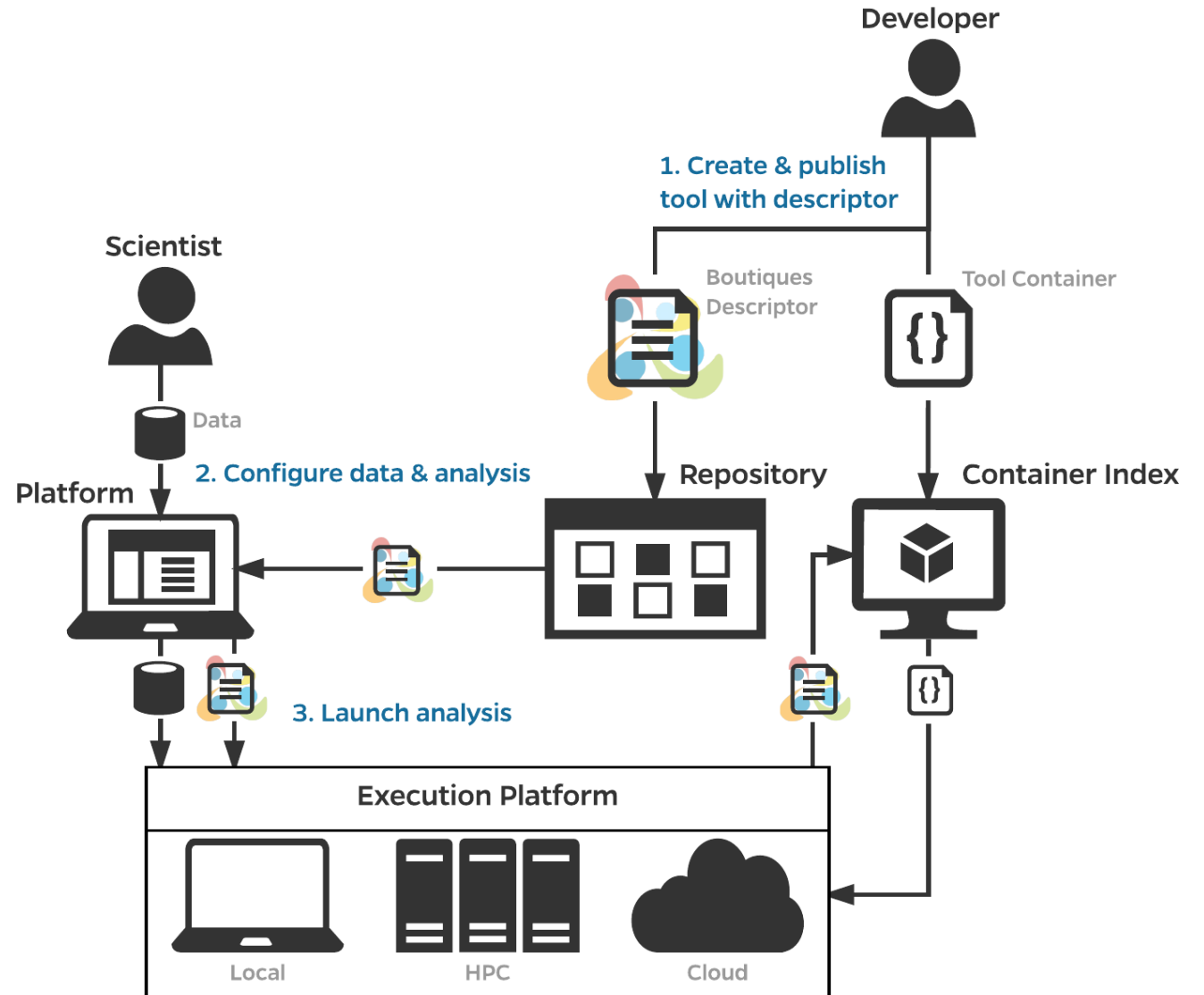


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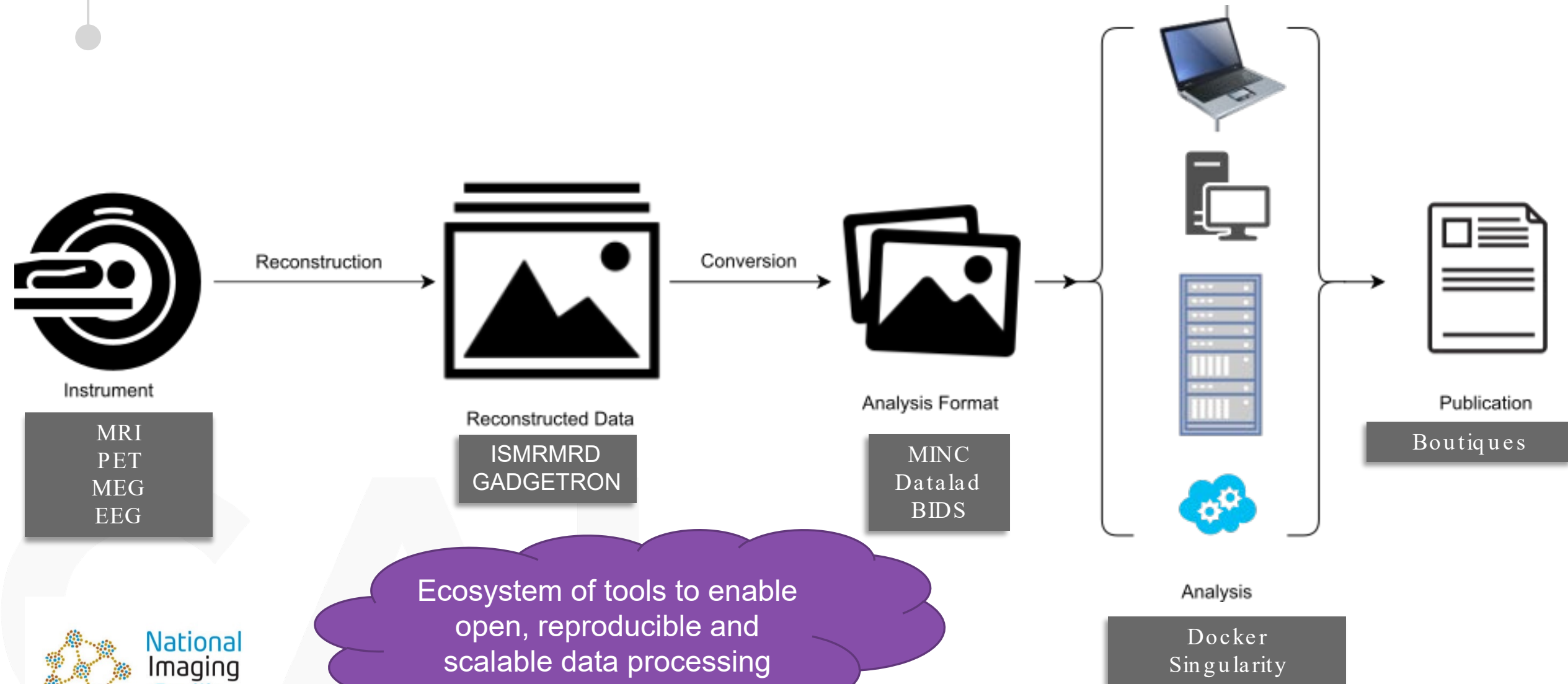


# BOUTIQUES

- automatically publish, integrate, and execute applications
- applications summarized in a JSON description



# FROM DATA TO PUBLICATION



# Thank you

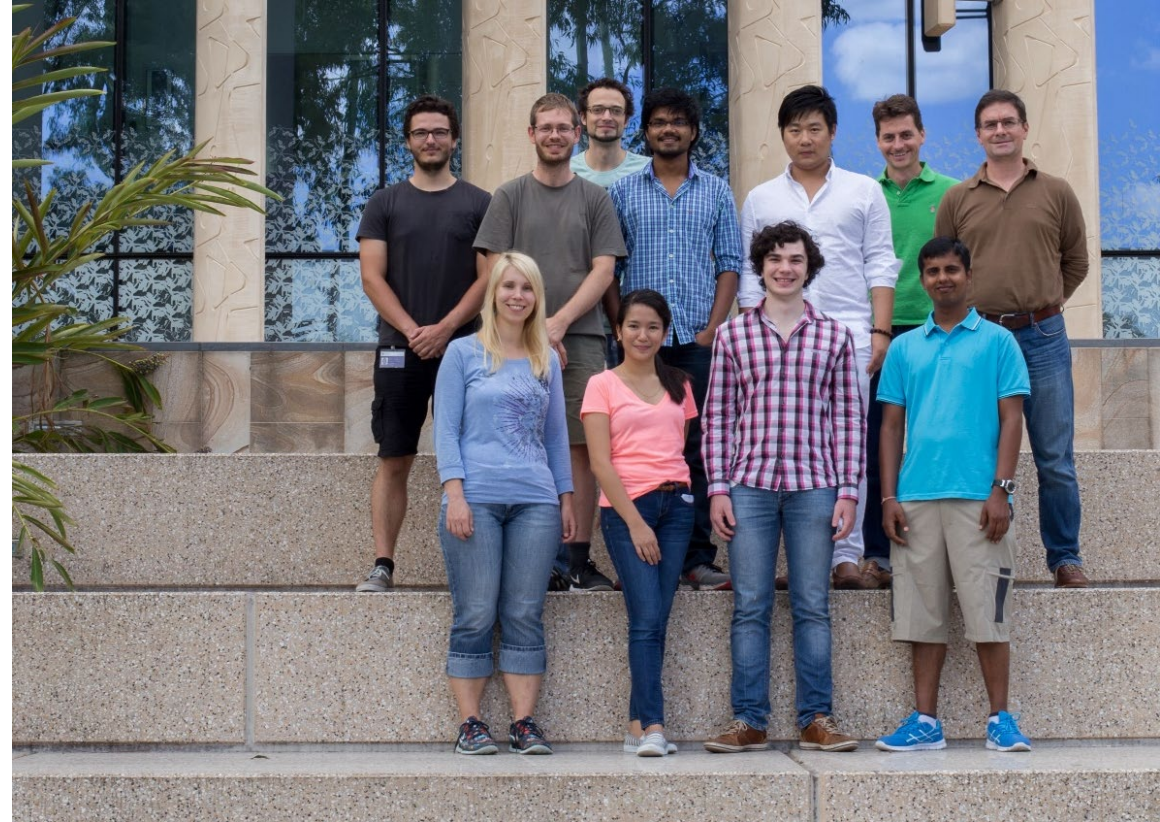


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[www.cai.uq.edu.au/bollmann](http://www.cai.uq.edu.au/bollmann)



[steffen.bollmann@cai.uq.edu.au](mailto:steffen.bollmann@cai.uq.edu.au)



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