



# Deliverables: 30 October 2017



AVATR



# Sprint 1 DoDs

- DoD: Explore the current landscape
  - We have many Tech Evals and notes on the landscape, and decided on what to use
- DoD: Capability to upload and store annotations
  - FIJI + gen\_commands + ingest\_large\_vol, stores annotation images in the boss
- DoD: Capability to pull/upload data from the boss
  - Modified NDR pulls data from boss, organizes into folders, and preps for upload
- DoD: Basic unsupervised methods
  - No implementation that runs an algorithm on data (we just got one?). Next week make a plugin?
- DoD: View labels
  - We can load the labels in NeuroGlancer/NDVIS from NDWT link, but can't see
- DoD: DOCUMENTATION
  - We are writing a lot of documentation

Sprint	Due Date	Requirements
Sprint 1	11/10	<ul style="list-style-type: none"><li>• Exploration into current landscape:<ul style="list-style-type: none"><li>◦ OpenNeuro, Clowdr, Boutique, AWS, NDWebtools</li></ul></li><li>• Data Ingest Plugin<ul style="list-style-type: none"><li>◦ Capability to upload and store annotations</li><li>◦ Capability to pull/upload data from boss</li></ul></li><li>• Analysis Plugin<ul style="list-style-type: none"><li>◦ Basic unsupervised methods</li></ul></li><li>• NDVis Visualization Plugin<ul style="list-style-type: none"><li>◦ Can view labels</li></ul></li><li>• DOCUMENTATION!</li></ul>

# This Week:

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- Make Documentation for ingest\_large\_vol ✓
  - Rewrote a lot of the pipeline based on feedback ✓
- Get feedback on pipeline ✓
- Demo of ingest\_large\_vol in BOSS ✓
- Demo for NDM (rip)
- Tech Eval for COINS/LORIS ✓

# Annotation Workflow Feedback

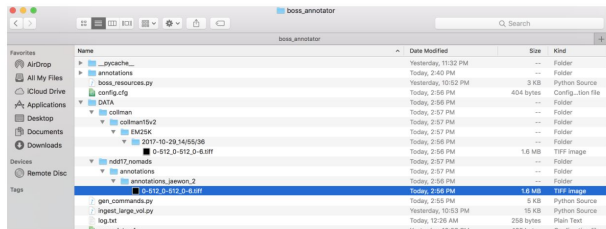
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- Hackathon feedback:
  - Doing stuff with their own collections?
    - Added stuff in documentation
      - LINE 80 IN INGEST\_LARGE\_VOL.PY
  - Part still not documented is with setting up annotation channels and the bugs along with it
    - Talk to Ben maybe
- Annotation squad:
  - INGEST\_LARGE\_VOL on next slide
  - Big point - saving the FIJI annotations as XML lets them pick up where they left off

# Updated docs: ingest\_large\_vol + DEMO

NOTE: At any point, you can export your annotations as an xml by the same method listed above. Opening the xml file will start you where you left off.

- k. A black screen will appear - these are your annotations, don't worry if you can't see them.
- l. Save your annotations in the correct directory *with the same name*, an example given below.



7. To push annotations to the BOSS, run `gen_commands.py`.
8. Paste command line output into terminal. If this doesn't work, you will probably have to change some parameters in `gen_commands.py`. Below is a list of all parameters:

## Parameters

Parameters	Description	Required	Tips and Examples
script	Path to ingest_large_vol.py script	Yes	Should not have to change.
source_type	Where the data is being ingested from	Yes	Either s3 or local.
s3_bucket_name	AWS S3 Bucket	No	Only specify if

## Feedback:

<http://neurodata-annotator.readthedocs.io/en/latest/>

- Annotators didn't like editing python code
- There wasn't documentation for people using different data who ran into other problems
- Including NDWebTools parts in documentation will help

## Demo:

- It looks like uint64 stuff doesn't work at all. So we just push annotations as uint8 since FIJI saves them like that.
- NDVis downsampling being weird

# Continued Issues with NeuroDataManager

- I think the documentation isn't exactly clear on what needs to be done in order to run DataManager. I think that:
  - I need to investigate the info manifest
  - Also structure the segmented directory? (does DM do this?)

```
(NDD) -----
l ~/Envs/NDD/DataManager/build @ Bijans-MacBook-Pro (bijanvarjavand)
l => bin/ndm -datadir ../../avatr-f17s18/source/group/base_annotator/collman_collman15v2_EM25K/ -input ../../avatr-f17s18/source/group/
base_annotator/DATA/collman_collman15v2_0_520_0_520_0_16_EM25K.tif -x 520 -y 520 -z 16 -scale 0 -gzip
[libc++abi.dylib: terminating with uncaught exception of type boost::exception_detail::clone_impl<boost::exception_detail::error_info_in
jector<boost::iostreams::gzip_error> >: gzip error: unspecified iostream_category error
*** Aborted at 1509214885 (unix time) try "date -d @1509214885" if you are using GNU date ***
PC: @ 0x7fffba197d42 __pthread_kill
*** SIGABRT (@0x7fffba197d42) received by PID 26757 (TID 0x7fffc2f863c0) stack trace: ***
@ 0x7fffba278b3a _sigtramp
@ 0x4c2f6a240 (unknown)
@ 0x7fffba0fd420 abort
@ 0x7fffb8c5094a abort_message
@ 0x7fffb8c75c17 default_terminate_handler()
@ 0x7fffb9785713 _objc_terminate()
@ 0x7fffb8c72d49 std::_terminate()
@ 0x7fffb8c729d2 __cxx_rethrow
@ 0x10fe8845d BlockManager_namespace::FilesystemBlock::load()
@ 0x10fe78ae6 BlockManager_namespace::Block::add<>()
@ 0x10fe775a2 BlockManager_namespace::BlockManager::Put<>()
@ 0x10fe765d5 main
@ 0x7fffb069235 start
@ 0xe (unknown)
Abort trap: 6
```

# For now using Boss is okay (since NDM not ready yet).

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- For NDM, I'm trying to copy converts .tif files to a NeuroGlancer precomputed datastore (which should be able to work for S3 and Google Cloud).
  - A few issues I've run into which hopefully I can get ironed out by talking to Alex and Ben
  - Mostly just setting up my manifest and directory structure properly
- Cubes in boss have NeuroGlancer links, so we can view stuff with that for now. We can also pull stuff from that
  - Demo
- Why? (I spent too much time working on ingest\_large\_vol demo)

# COINS is a cool LIMS with some features we need...

LIMS (Laboratory Information Management System) is software designed for data storage, exchange, tracking and workflows (pipelines).

What it has that we care about:

1. Data sharing and collaboration (Portal)
2. Data exporting and downloading (QB)
3. Data uploading (DICOM Receiver)
4. Dashboard for viewing analyses results
5. Logs data history

The image displays three screenshots of the COINS LIMS interface, labeled A, B, and C.

**Screenshot A:** Shows an 'Analysis Report' for 'All URSI Reports'. It includes a table with columns for 'Incomplete', 'DTI', 'VBM', 'PS', 'GAMBLING', 'TASTE', and 'HRT'. The table contains data for various sessions and tasks, with cells colored green, yellow, and red.

**Screenshot B:** Shows a 'Filtered By' section with options for 'Show rows with image only', 'lowQA: 0', and 'highQA: 1'. Below this is a table with columns for 'URSI', 'SESSION', 'TASK', 'Segmentation', 'QA', and 'ANALYSIS FILES DOWNLOAD'. The table shows data for a specific session (20080117 143936) and task (mfrage).

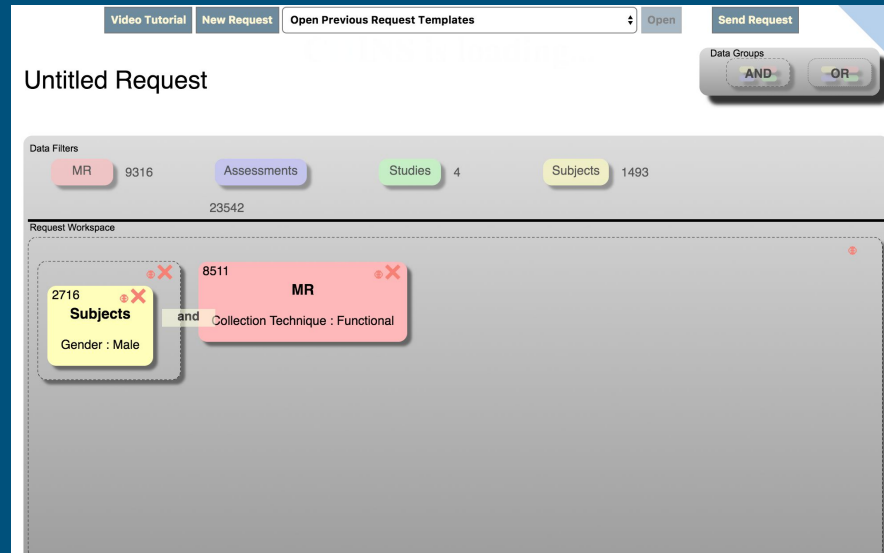
**Screenshot C:** Shows a 'Filtered By' section with options for 'Show rows with image only', 'lowQA: 0', and 'highQA: 1'. Below this is a table with columns for 'URSI', 'SESSION', 'TASK', 'Segmentation', 'QA', and 'ANALYSIS FILES DOWNLOAD'. The table shows data for various sessions and tasks, with cells colored green, yellow, and red. To the right of the table is a grid of brain scan images.



# ... but does not go well with what we need.

Cons for us:

- Only supports MRI, MEG, EEG scans and “clinical assessments”
  - So most of its features won't work for NOMADS, Clarity
- Used mainly to protect patient privacy and data sharing permissions
- Made for doctors or clinical researchers, not data scientists.



# For Next week:

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Wednesday: Hopefully figure out DataManager stuff. Plan out LIMS based off of what we saw in COINS and OpenNeuro

- Make Documentation for in Sphinx
  - Using NDWebTools
  - Using NDM
- Demo of storing things in NeuroDataManager
  - Images
  - Annotations
  - Objects?
- Demo of a preliminary LIMS?