WEAVE data products and overall dataflow

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WEAVE Science Verification Community Workshop Tenerife 14th-15th November 2019

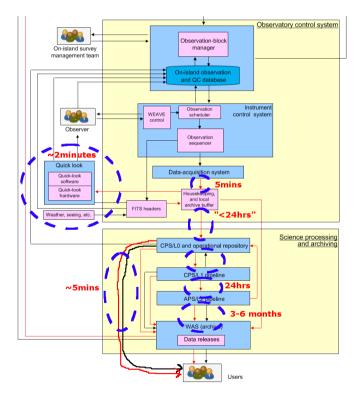


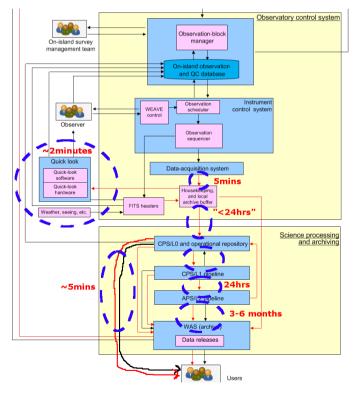
This session: *aim* and overview

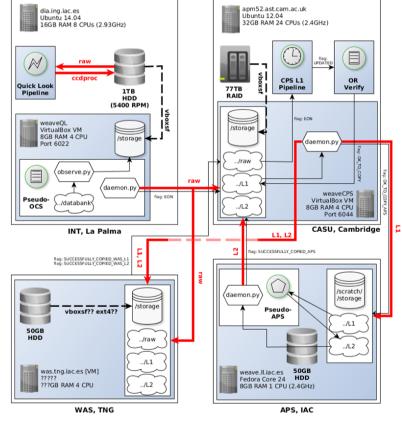
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- The main components within the WEAVE data model
- How they interact
- WEAVE data products: Input and Output
- How you get them

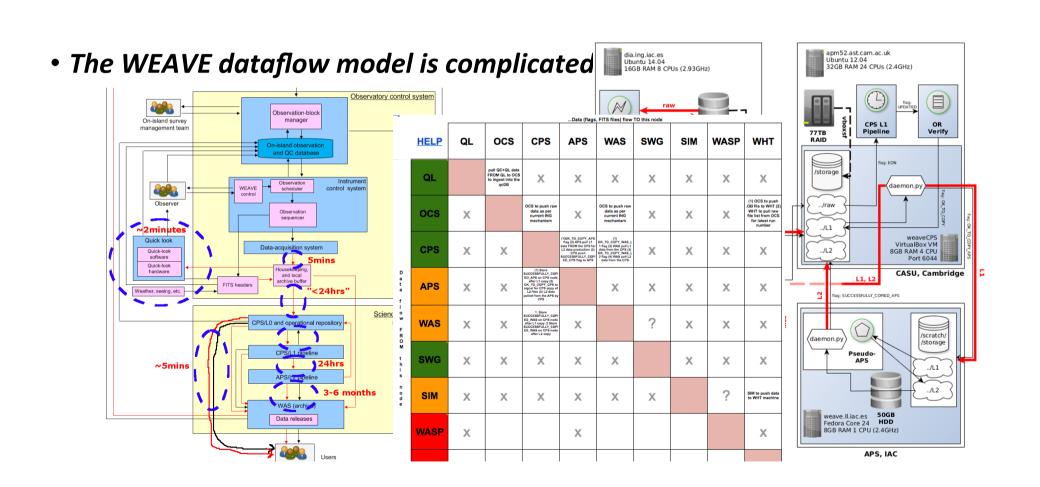
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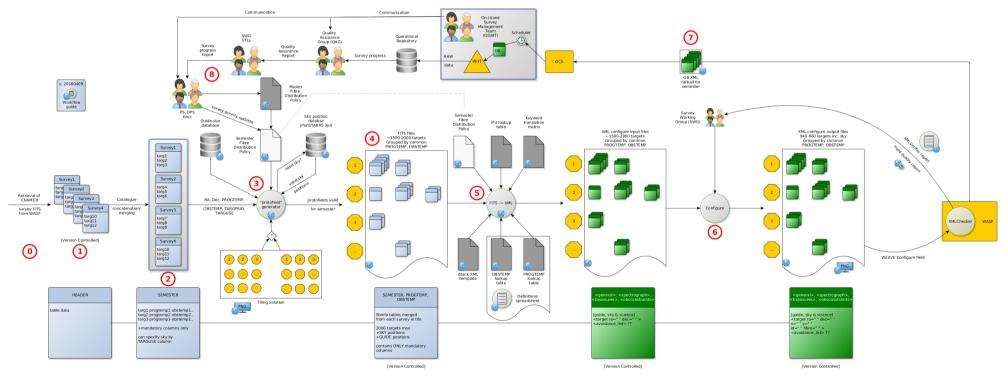
- Build on the broad outline of WEAVE dataflow given by Shoko
- The main components within the WEAVE data model
- How they interact
- WEAVE data products: Input and Output
- How you get them
- This isn't a lecture. Ask questions, ask for clarification etc etc

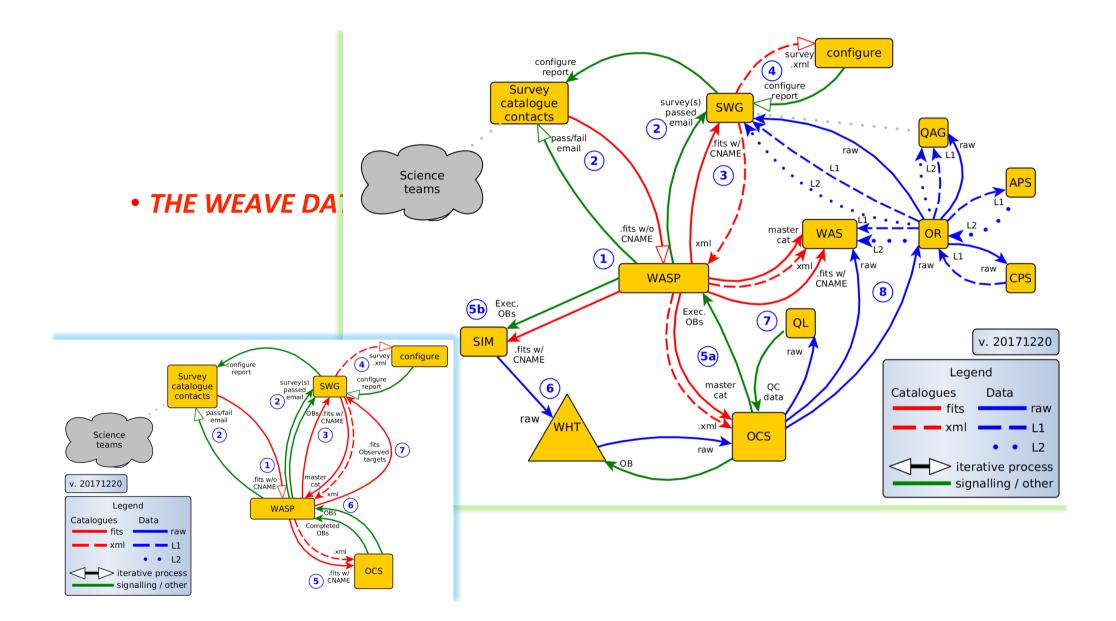


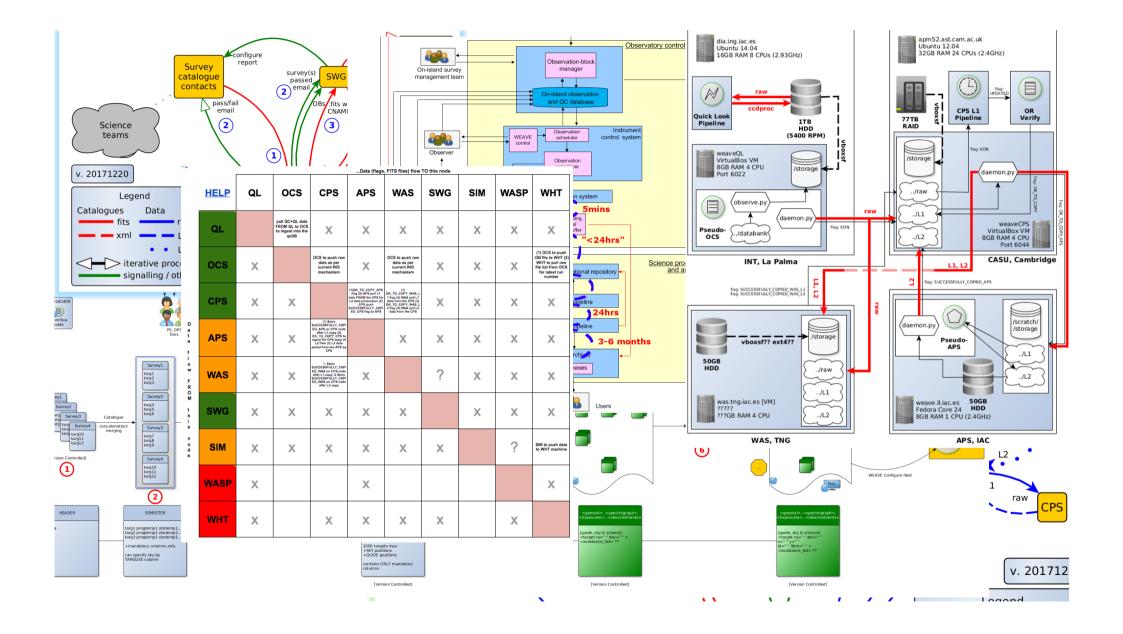


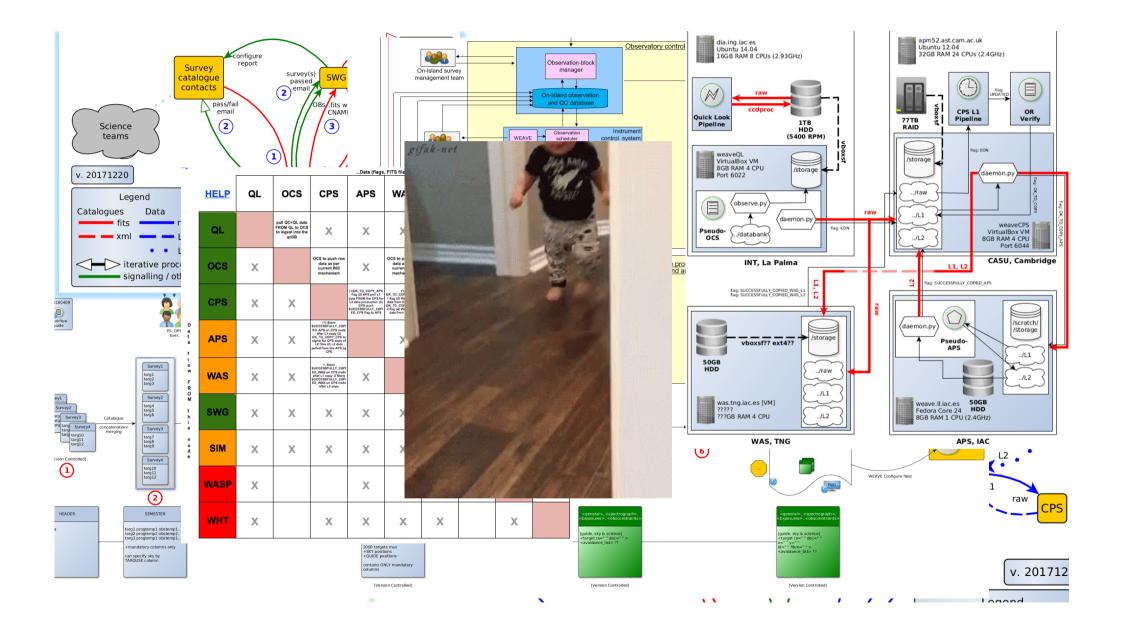












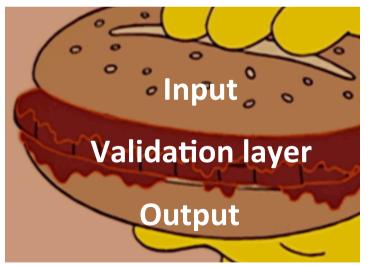
Our aims for this workshop

- The WEAVE dataflow model is complicated
- Our mission should be to make using WEAVE as easy to use as possible
- However... WEAVE is a complicated instrument multiple modes, spectrograph arms, resolutions, varied 1D, 2D, 3D data products, complex interplays between different systems
- The overall dataflow system must incorporate all of these, but under a "survey-mode" philosophy that demands science can be done even in adverse conditions, with fast data turnaround and often conflicting survey requirements.

WEAVE dataflow context

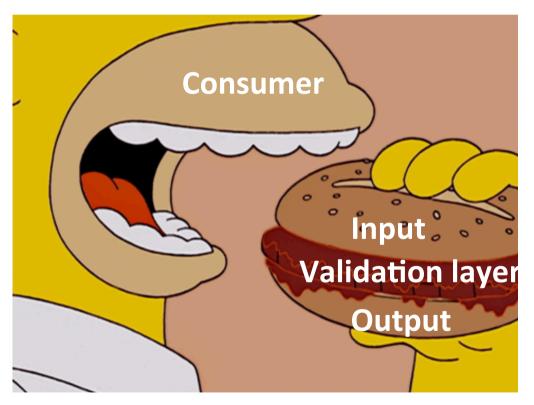
- The WEAVE dataflow model is complicated
- We have already implemented the dataflow design, and simulated realistic WEAVE operations under this model
- Implementation encompasses required inputs and the outputs generated by pipelines and analysis packages
- In devising the overall system, necessary balance between flexibility/ features for would-be users vs. the need for streamlined "surveymode observing" and data treatment
- It's always the small niggles that catch you (us).... No questions are stupid, so please ask: you might poke a hole in our model (good!)

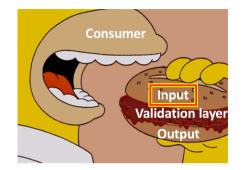
- We will discuss key parts of the dataflow today; in-depth look at SV-relevant components tomorrow.
- The WEAVE data sandwich:



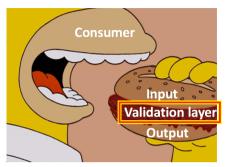
- Basic concept is that usergenerated input data are subject to validation against the WEAVE data model
- Data passed to WHT, observations made, output data are also validated (just in case) against the model...

- ... and then made available for consumption.
- There is more than one way to eat the WEAVE data sandwich – we will cover these today, and highlight the differences.
- Let's now look at these 4 components

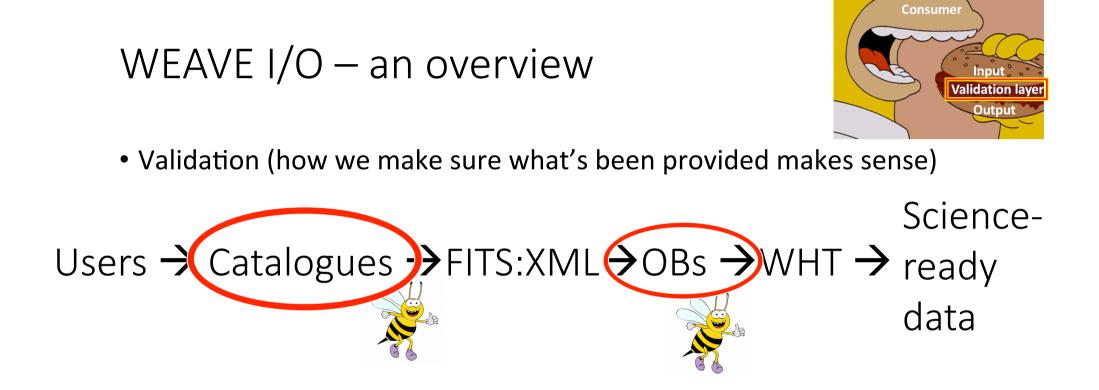




- Input (what users must provide)
 - Input FITS-format catalogues describing the targets you wish to observe
 - XML-format Observing Blocks (OBs) describing the fiber positions, instrument configuration, minimum observing conditions, etc
- For Science Verification:
 - PIs will generate their FITS catalogues
 - WEAVE members will help generate OBs on your behalf
 - PIs will review generated OBs to agree they meet aims of the proposal
- Under normal operations
 - PIs are responsible for FITS and OB submissions, provided with tools to help



- Validation (how we make sure what's been provided makes sense)
 - Upload facility to a validation platform (think ESO p2pp)
 - Pass / Fail with a generated report: both FITS and XML
 - Tells you the products are legal, but can't check for "common sense"
 - "I only want to allocate 5 of these 940 fibers to my targets" (you might want to !!)
 - "I meant to use NGC in my target names, but used NCG by mistake"
 - "I stuck that mIFU bundle on the wrong galaxy"
 - "This 20th mag galaxy can totally be observed with a sky brightness of 17.5 mags"
 - Warnings *are* issued in some cases, but no substitute for common sense!
 - This facility acts as a gatekeeper between users and the instrument



The WEAVE Automated Submission Platform (WASP)

Teaser slides for tomorrow



JobID 1573684536

#	ID	File	User	Field	RA	Dec	PROGTEMP	OBSTEMP	Plate	Sci fibres %	Process State	Result	Report	Download
1	2	3365.xml	weave	PS1_3557	27.0658	27.9602	41331	XECEC	LIFU	<lifu></lifu>	COMPLETED	FAILED	X	[<u>XML]</u>
2	3	3367.xml	weave	PS1_3557	27.0658	27.9602	41331	XECEC	LIFU	<lifu></lifu>	PROCESSING			
3	4	3366.xml	weave	PS1_3557	27.0658	27.9602	41331	XECEC	LIFU	<lifu></lifu>	PENDING			
4	5	3368.xml	weave	PS1_3557	27.0658	27.9602	41331	XECEC	LIFU	<lifu></lifu>	PENDING			

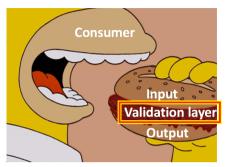
Cambridge Astronomy Survey Unit – Institute of Astronomy <u>The WEAVE Consortium</u> casuhelp@ast.cam.ac.uk



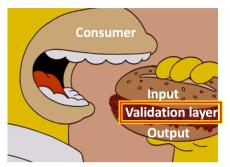
WEAVE Automated Submission Platform (WASP)

XML report for 3367.xml (jobID 1573684536, uid 3)

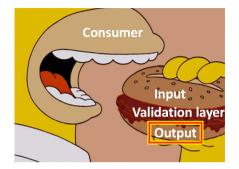
Result for file 3367.xml: Failed # of invalid entries: 2013 # of warnings: 1651	ement Logout
<pre>XML File Verification for WASP Compliance Current Version = OpR3d Current trimester = 2019A2 Started at: Wed Nov 13 22:37:01 2019 0/ FileIO and XML parsing</pre>	Download
VALID: File passed preliminary IO and parsing checks 1/ XML validation	(XML) (XML) (XML)
INPUTFILE: 3367.xml TEMPLATEFILE: 'BlankXMLTemplate.xml' (version 1.13) WARNING: Version of submitted XML (1.10) does not match version of current template (1.13) 2/ Baseline structure check against template XML	(XML) elp@ast.cam.ac.uk
VALID: Submitted XML matches structure and content of template XML 3/ Attribute check against template XML (based on LIFU observations)	
INVALID: For IFU submissions, XML should contain XMLIFU code version in a root:comment attribute. This attribute was not for INVALID: Attribute 'casuid' is missing from element 'observation' in submitted XML file INVALID: Attribute 'linkedgroup' is missing from element 'observation' in submitted XML file INVALID: Attribute 'obsgroup_validity' is missing from element 'observation' in submitted XML file INVALID: Attribute 'trimester' is missing from element 'observation' in submitted XML file INVALID: Attribute 'trimester' is missing from element 'observation' in submitted XML file INVALID: Illegal attribute 'semester' exists in element 'observation' in submitted XML file INVALID: Illegal attribute 'max_sky' (value:'100') in element 'observation:configure' does not match template XML value (



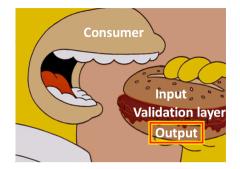
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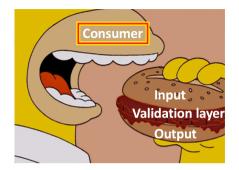
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 - Warnings *are* issued in some cases, but no substitute for common sense!
 - This facility acts as a gatekeeper between users and the instrument
 - Validation platform is called WASP developed and hosted by CASU
 - Comes with a variety of useful tools for preparation and management of observations
 - Test channel is available to check submissions well in advance of deadlines



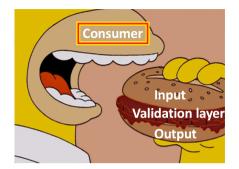
- Output (what data you will have access to)
 - Data from the instrument, including calibration data (arcs, flats)
 - Science frames processed with the WEAVE reduction pipeline:
 - Bias subtracted, flatfielded ("2D image processing")
 - Spectral extraction, wavelength resampling, flux calibration, sky subtraction
 - Creation of deep stacks from exposures within the 1hr OB and even-deeper stacks from observations over multiple OBs
 - Generation of data cubes for IFU observations (both mini-IFU and Large IFU)
 - Data products derived from analysis of these reduced products:
 - Target classification
 - Model template fitting (stellar and galaxy)
 - Redshift estimation, stellar parameter estimates (Teff, log g), abundances, indices
 - User selection of modules that allow analysis through additional software packages



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- Generation of data cubes for IFU observations (both mini-IFU and Large IFU)
- Data products derived from analysis of these reduced products:
- APS "L2"
- Target classification
- Model template fitting (stellar and galaxy)
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- User selection of modules that allow analysis through additional software packages



- Consumption / access (how you get the data)
 - There are two access points for WEAVE data, but they are different in design and purpose. SV users will have access to both.
 - Operational Repository is hosted at CASU
 - Tracking survey progress, OB status, data processing status
 - Access to most recent data available (RAW, L1, L2), driven by files, not targets
 - NOT part of the data release cycle: data is subject to change!
 - WEAVE Archive System hosted at Telescopio Nazionale Galileo (TNG)
 - Provides target-driven searches of **stable** data releases
 - Customisable and flexible searches, user interface NOT designed by David Murphy
 - Data visualisation, bulk downloading
 - SV papers must use data from the WAS, not the OR



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WEAVE I/O – what do the products look like?

	RAW (one file / arm)	L1 (CPS; 1 file / arm)	L2 (APS; 1 file / exposure)*	
	FITS file, image + tables	FITS file, image + tables	FITS file, tables	
n s	PHU	PHU (inherited)	PHU (inherited)	
0	Image data Det1	Sky-subtracted 1D spectra	Fibre information table	
n s i	Image data Det2	inv. variance	Classification spectral fit	Classification extracted parameters
x t e		No sky-subtracted 1D spectra	FERRE stellar spectral fit	FERRE extracted parameters
Ш		No sky-subtracted inv. Variance	RVSPECFIT stellar spectral fit	RVSPECFIT extracted parameters
ΙTS	Fibre information table	Fibre information table (extras added)	Galaxy model spectral fit	Galaxy extracted parameters
ш		IFU: white light image for cube	* In some instances, per- arm too	

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WEAVE I/O – the fibre information table

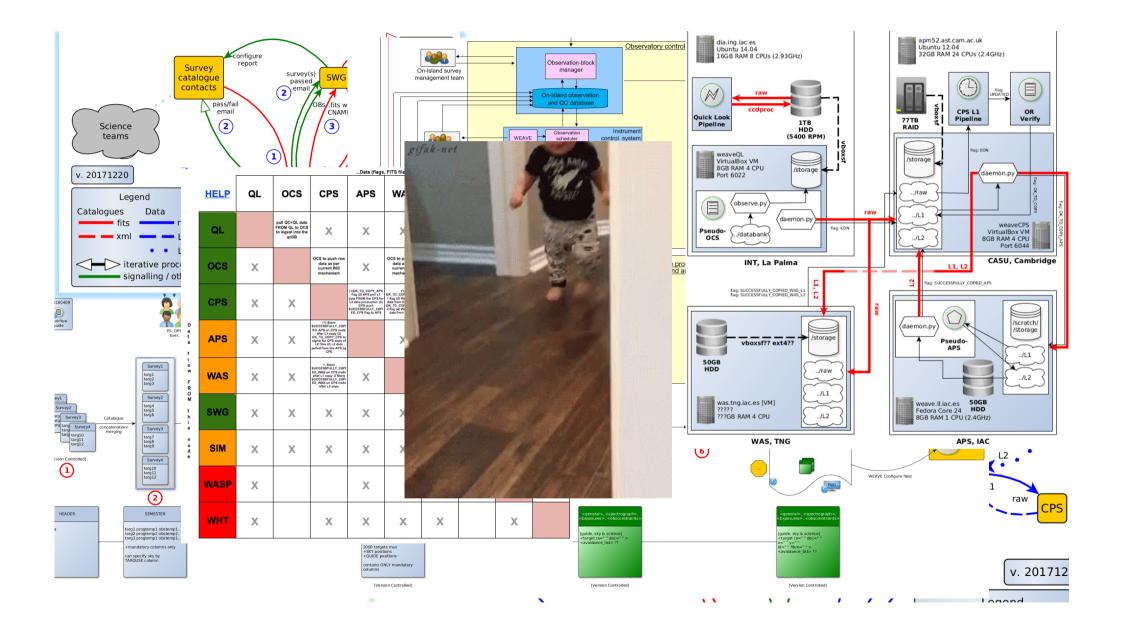
- Describes what the fibers are pointing at in each data product
- Inherits important data from the FITS target catalogue the user supplies
- This information will be ingested into the (target driven) WEAVE Archive System, and to a lesser extent the Operational Repository
- The **fib**re **info**rmation (fibinfo) table must include important columns that allow us to identify:
 - What target?
 - How was it observed? What observing conditions were requested?
 - How was it analysed?

WEAVE Targets: useful things to care about

- CNAME maps position on sky \rightarrow ID
 - All science targets, calibration targets and sky fibers have a CNAME
 - 01:33:50.9 +30:39:35.8 → WVE_01335090+3039358
- Instrument configuration: "PROGTEMP" how was this target observed?
- Observing constraints: "OBSTEMP" what were the minimum conditions requested?
- Analysis flags: "APS_FLAG" are extra L2 analysis packages required?
- Applies to exposures, OB-stacks, "super-stacks"*
 - *when a target is observed over multiple Obs, this is a stack of all observations
- A search in the WEAVE Archive for a target can yield >1 result. Why?
 - The same target could be observed in MOS in low-res AND high-res
 - Target "uniqueness" is generally:
 - CNAME + configuration (+survey in some cases)

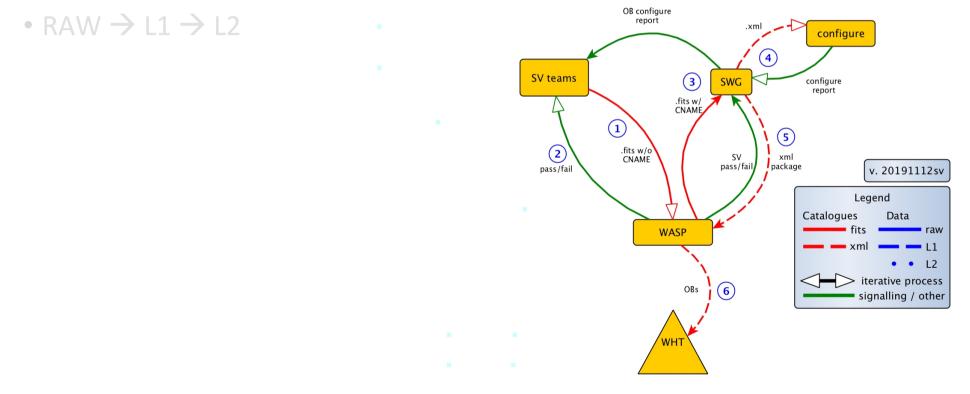
Let's tie this all together....

• ...with some simple plots

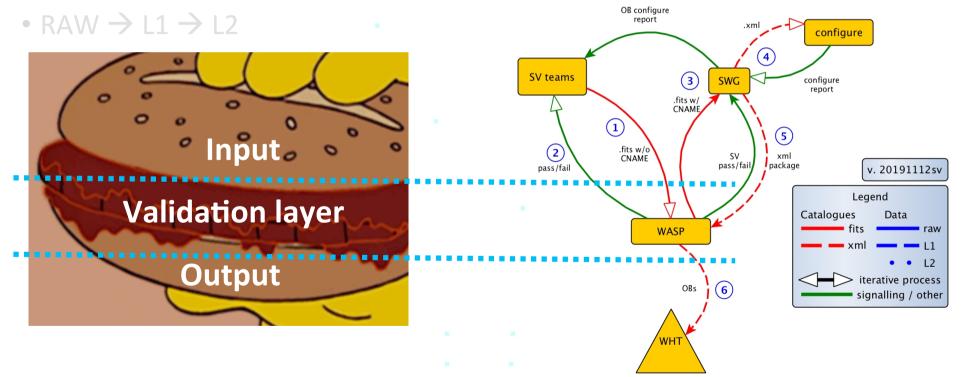


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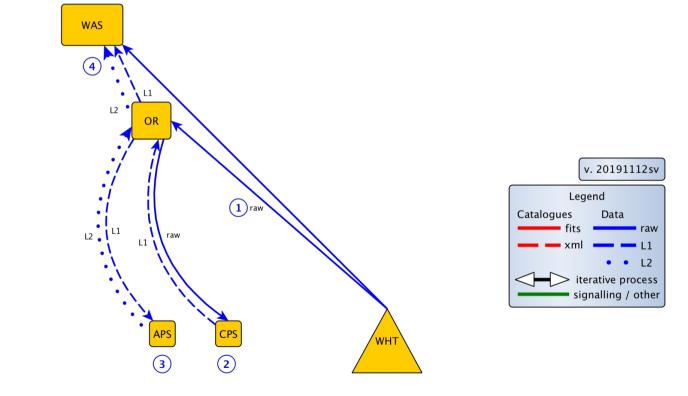
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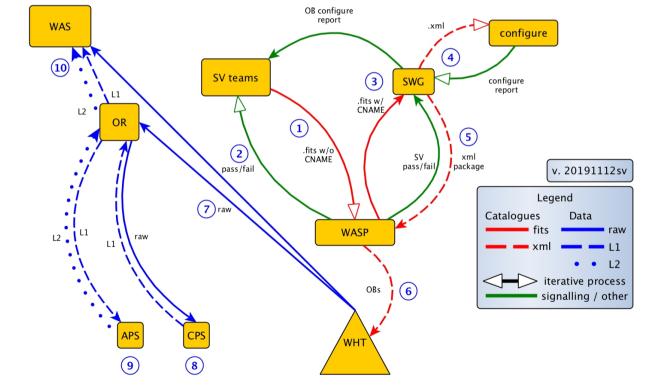
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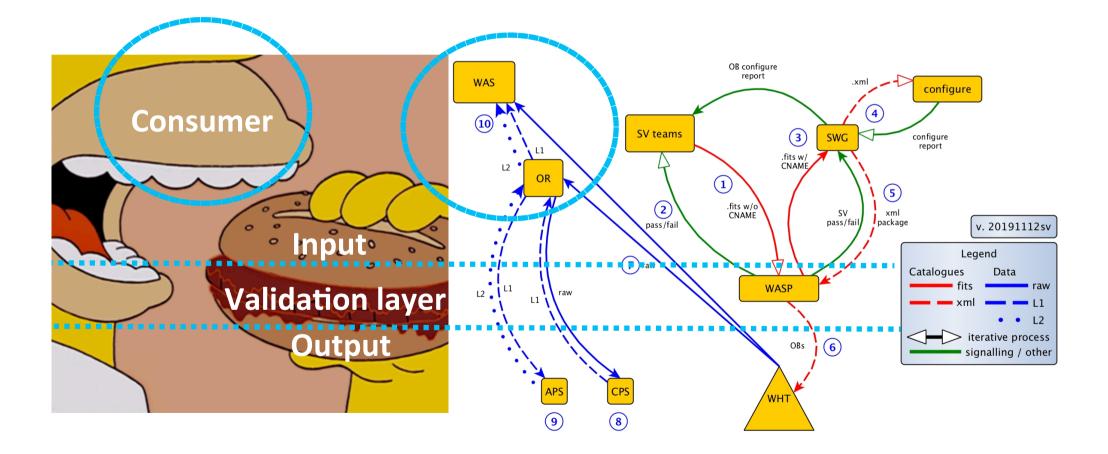


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• Don't worry, stay hydrated!



• WEAVE(!) been doing this for quite a while, and it's still challenging to us

Inputs	Validation	Outputs	Access
FITS catalogues (targets)	WASP platform	RAW data	Operational Repository (survey/processing progress, bleeding-edge data)
XML observing blocks	WASP platform	L1 processed data ("spectra")	WEAVE Archive System (proper data releases, designed for science end-users)
		L2 analysis ("spectral analysis, redshifts, abundances)	

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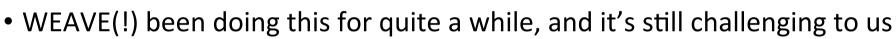
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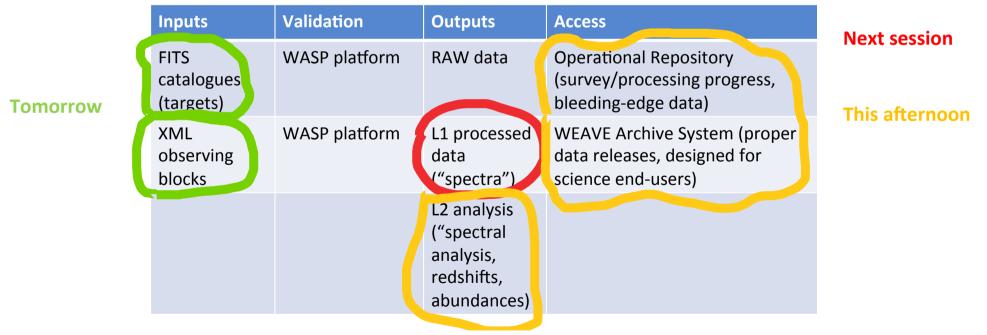


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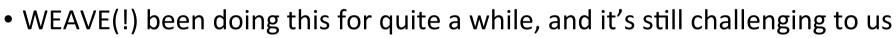
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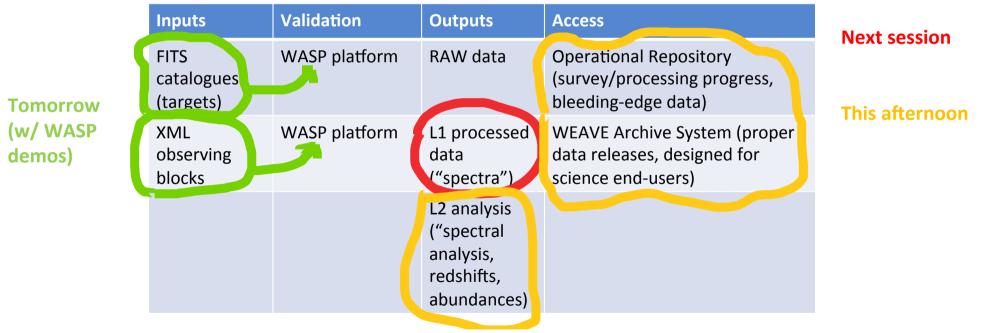
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