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# Overview and status of the validation framework

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# Outline

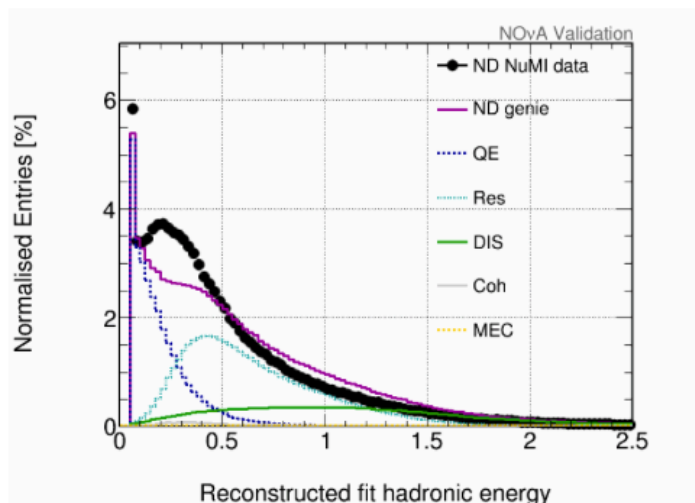
- Framework designed by Matthew Tamsett
- More details found here:  
<https://cdcv.s.fnal.gov/redmine/projects/novaart/repository/entry/trunk/Validation/analysis/README>
- Last update given by Bruno Zamorano in DocDB-[14824](#)
  - useful overview of the website
- Introduction to the validation framework
- Brief guide on how the framework works
- Histogram naming convention
- Current state
- To-do list

# The validation framework

- The validation framework was designed to lessen the load of validation plot making
- Instructions on installing and running the framework can be found here:  
<https://cdcv.s.fnal.gov/redmine/projects/novaart/repository/entry/trunk/Validation/analysis/README>
- Validation framework works by running a python script which points to a CAFAna macro via json configuration file
  - The json file contains configurations for the novasoft release, validation plot version, datasets and shell scripts to be run
  - Each run over a macro produces a root file containing histograms defined by the macro and dataset
- Example:
  - ***python \$NOVAANAVALID\_DIR/core/controller/analysis\_validation.py -c /nova/app/users/vinton/ana\_valid/nu\_mu/configurations/caf\_numu\_nd\_cutflow.json***
- The shell scripts each point to a CAFAna macro
  - For example, one of the shell scripts pointed to by **caf\_numu\_nd\_cutflow.json** is **nu\_mu/components/run\_caf\_numu\_nd\_cutflow.sh**, which runs: **cafe -bq caf\_numu\_nd\_cutflow.C**

# The validation framework

- A website displaying all validation components created by a user can be generated with: **`python $NOVAANAVALID_DIR/core/view/makeHTMLView.py`**
  - The website is created at:  
<http://nusoft.fnal.gov/nova/users/<USERNAME>/validation/>
- The website user can select the datasets they want to compare
- Cuts can be selected from a dropdown menu next to a plot (If the CAFAna macro produced hists for different cuts, see “naming convention” slide)



png

## Reconstructed fit hadronic energy

ND NuMI data: prod\_decaf\_S15-05-22a\_nd\_nuMI\_numu\_contain\_goodruns  
ND genie: rocco\_decaf\_S15-05-22\_nd\_genie\_fhc\_nonswap\_genierw\_numu\_contain

Dataset	Entries	Mean	Mean diff [%]
ND NuMI data	501,010	0.52	0.00
ND genie	1,510,188	0.61	17.07

## Controls

Log y:

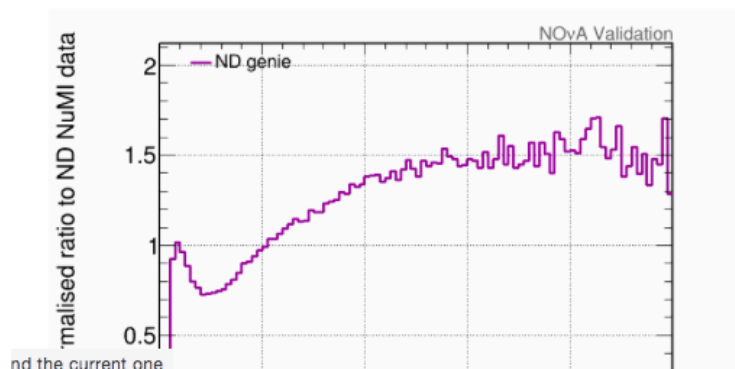
Normalisation:

Truth:

Ratio zoom:

Reco cuts:

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# Aside: naming convention

- All histograms must follow the naming convention:
  - **<category>-<variable>-<reco\_level>-<true\_level>**
    - **category**: logical category the plots fall into (reco, true, mixed, sam)
    - **variable**: observable
    - *reco\_level*: reconstruction cuts
    - *true\_level*: QE, Res, DIS, Coh
  - both *reco\_level* and *true\_level* are optional
  - Examples of acceptable names:
    - reco-reco\_numu\_energy
    - reco-reco\_numu\_energy-QE
    - true-numu\_energy
    - true-numu\_energy-n\_minus\_1\_preselection
- The website user can choose the truth and reco level for each plot, provided there are multiple reco\_levels and/or true\_levels per **<category>-<variable>** combination

# Changes needed to run art modules

- The validation [CAFAna](#) jobs currently dump output histograms in a flat format directly to a directory
  - However, [Art](#) modules dump the histograms into a subdirectory within a file. To enable use of the framework with art modules we should choose one of the two output file structures and apply to both art and [CAFAna](#) validation jobs
- Both the output file structure and the use of multiple module [Art](#) jobs need attention before art modules can be run with the framework

# Current state

- Website populated with image files ([eps](#))
- Can produce plots comparing variables in multiple datasets (data or MC)
  - validation with [CAFAna](#) working smoothly
  - **work needed to enable running art modules**
- Unfortunately, I don't have the time to develop this any further, framework needs a new champion
- further development will require some understanding of [json](#), [python](#) and [bash](#)

# Future plans

- framework needs adjusting to run art jobs
- I don't plan to have the time to develop this further
- **framework needs a new champion!**
  - Sussex group can provide support to get started
  - Framework is not complicated, get stuck in!
  - **Great** way to learn and hone **transferable** computing **skills**

# Backup



# The validation framework

- The validation framework was designed to lessen the load of validation plot making
- Instructions on installing and running the framework can be found here:  
<https://cdcv.s.fnal.gov/redmine/projects/novaart/repository/entry/trunk/Validation/analysis/README>
- Validation framework works by running a python script which points to a **CAFAna** macro via **json** configuration file
  - Each macro is run once for each dataset defined in the **json** file
  - Each run over a macro produces a **root** file containing histograms defined by the macro and dataset
- The website is created and populated with image files by running **“makeHTMLView.py”**
  - Multiple image files are created for each histogram type
    - Image files are made for all combinations of data/MC/Cuts for each detector
- The website user can select the datasets they want to compare
- Cuts can be selected from a dropdown menu next to a plot (If the **CAFAna** macro produced hists for different cuts, see “naming convention” slide)

# The validation framework

- Validation jobs are started by running the python validation script with a **json** file as argument, for example:
  - **`python $NOVAANAVALID_DIR/core/controller/analysis_validation.py -c /nova/app/users/vinton/ana_valid/nu_mu/configurations/caf_numu_nd_cutflow.json`**
  - The **json** file contains configurations for the novasoft release, validation plot version, datasets and shell scripts to be run
- The shell scripts each point to a **CAFAna** macro
  - For example, one of the shell scripts pointed to by **caf\_numu\_nd\_cutflow.json** is **nu\_mu/components/run\_caf\_numu\_nd\_cutflow.sh**, which runs: **`cafe -bq caf_numu_nd_cutflow.C`**
- A website displaying all validation components created by a user can be generated with: **`python $NOVAANAVALID_DIR/core/view/makeHTMLView.py`**
  - The website created at:  
**<http://nusoft.fnal.gov/nova/users/<USERNAME>/validation/>**