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Calibration Workflows with Cylc

10/8/2024

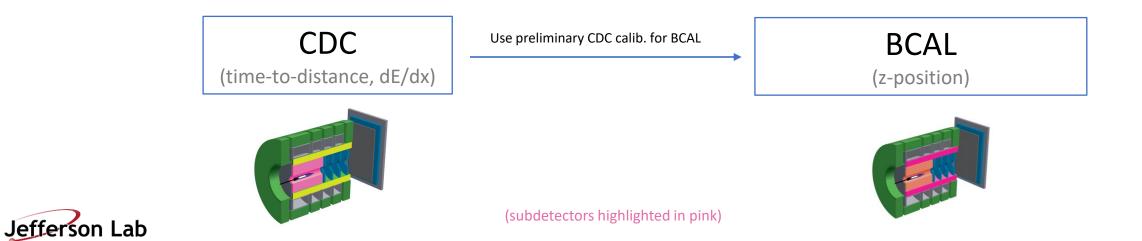
EPSC





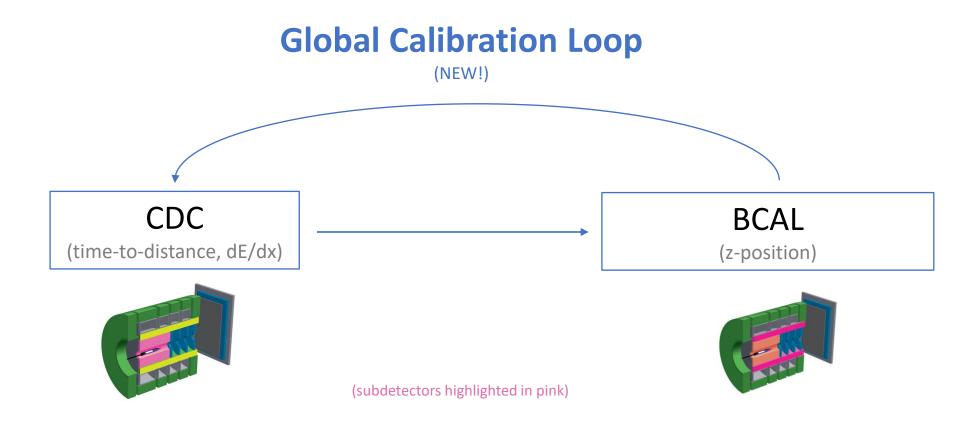
Calibration Motivation

- Often, calibrations need to be done sequentially
- Simple example: tracking \rightarrow EM calorimeters @ Hall D



Calibration Motivation

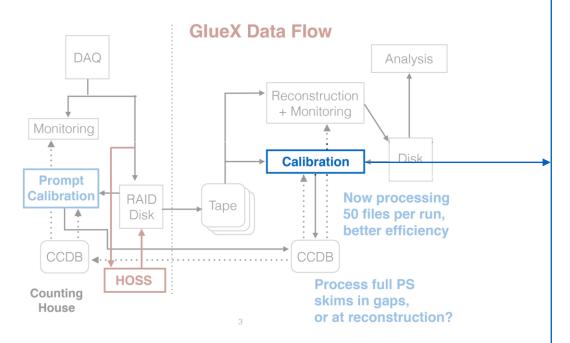
Will iterating over multiple subsystems improve calibrations?



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Calibration Motivation, cont.

Usually situation is more complex. Motivation to use cylc (pronounced "silk")



Run-dependent Calibrations

Calibration	Person
Overall Timing (first pass)	Sean Dobbs
Overall Timing (post-ST updates)	Sean Dobbs
CDC wire gains	Naomi Jarvis
CDC overall gains	Naomi Jarvis
CDC dE/dx	Naomi Jarvis
CDC time-to-distance	Naomi Jarvis
PS/PSC Timing	Olga Cortes
TAGH timewalks	Olga Cortes
TAGM timewalks	Sean Dobbs/Richard Jones
TOF	Beni Zihlmann
DIRC	Justin Stevens

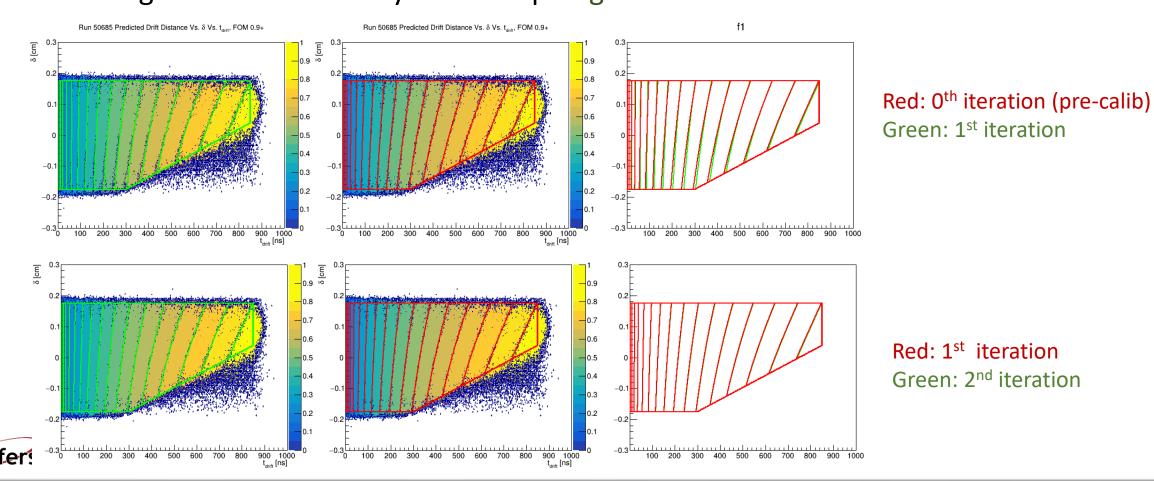
Overall Calibrations

Calibration	Person	Status
BCAL channel timing	Mark Dalton	
BCAL attenuation length/gain ratio	Mark Dalton	
BCAL z-position	Mark Dalton	
BCAL gains/non-linearities	Karthik Suresh	
FCAL gains/non-linearities	Colin Gleason	
FCAL timing	Colin Gleason	
SC Timewalks	Rupesh Dotel	Done
SC Propagation Time	Rupesh Dotel	



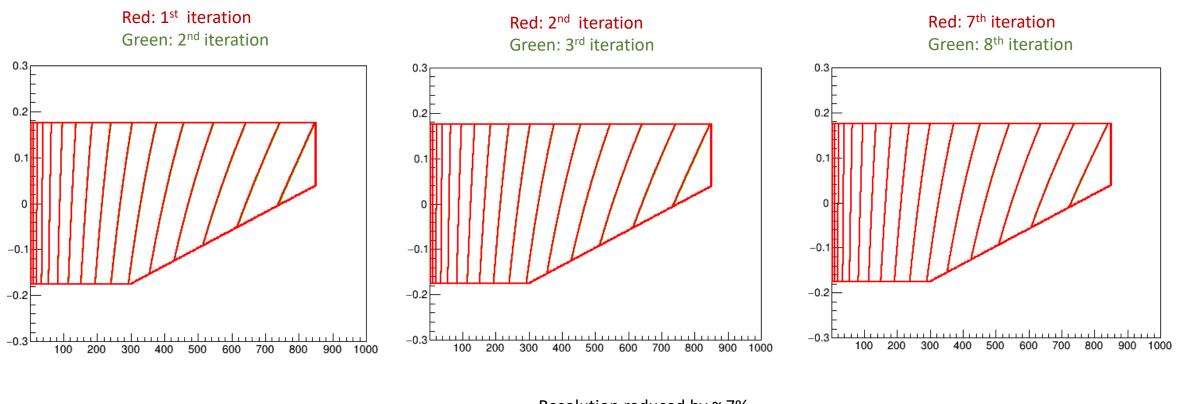
Case Study: CDC Time-to-Distance (TTOD)

Sometimes procedure need to iterate/converge, too Converged when red totally lies on top of green



Case Study: CDC Time-to-Distance (TTOD)

Here: converged when red totally lies on top of green



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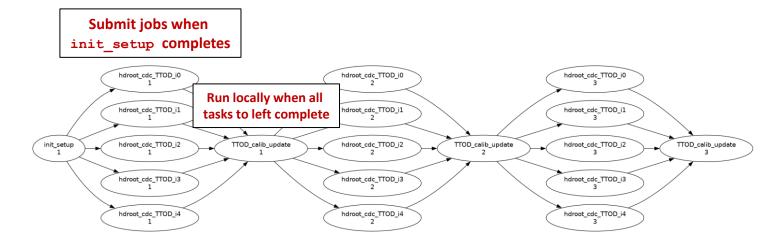
Resolution reduced by $\sim 7\%$ (1st to 8th iteration) Calibration Workflows with cylc

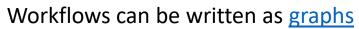
(pronounced "silk")

Punchline: I did all this from a single terminal command!

> cylc vip gx_ttod

- Runs 20 jobs per iteration
- Then repeat ×8 iterations
- Read/write ccdb (local copy)
- Automatically resubmits failed jobs (but only if I asked it to)



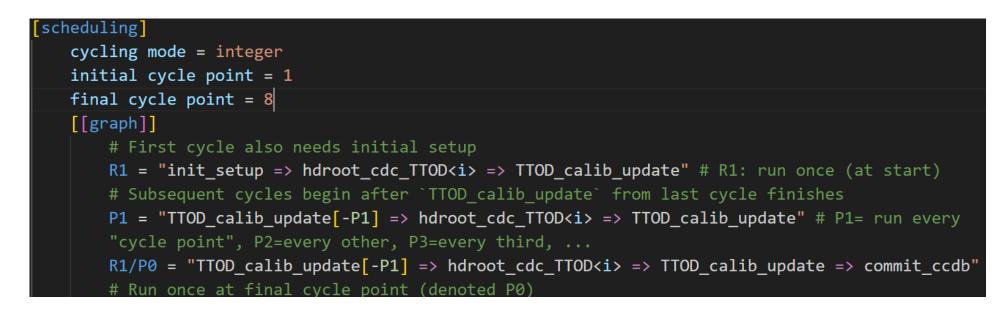




Defining Workflows

Workflows described in file called flow.cylc

Gives graph of tasks to run





Defining Workflows, cont.

Workflows described in file called flow.cylc

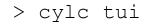
Add inherit = gx_recontask to run as batch job (otherwise runs locally)

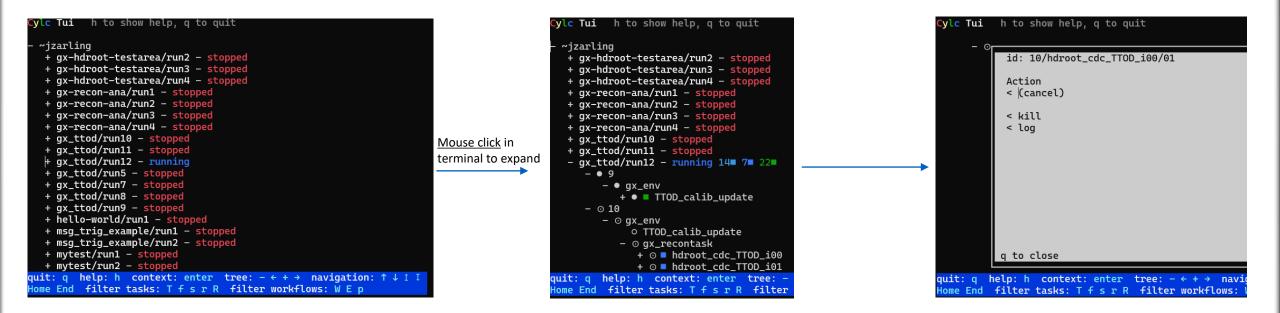
```
[[hdroot_cdc_TTOD<i>]]
inherit = gx_recontask
script = """
source gx_env.sh
hd_root --config=${jana_config} ${evio_folder}/hd_rawdata_0${run}_${fnum}.evio
mv hd_root.root ${ttod_topdir}/root/hd_root_TTOD_${run}_${fnum}_CP$
{CYLC_TASK_CYCLE_POINT}.root
"""
```





Terminal interface (interactive!):





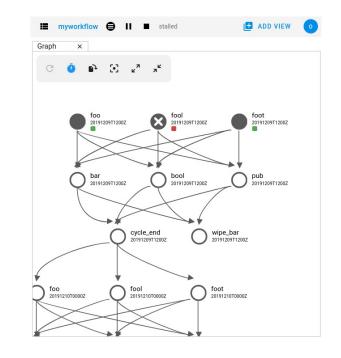


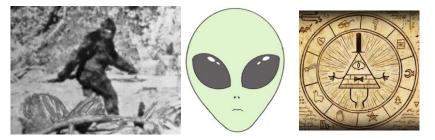
Maybe One Day?

An online dashboard for calibrations in process

- Multiple users can access, monitor, modify, etc.
- Different users can have different privilege levels
 - Person A: global control
 - Person B: can start/stop/modify jobs related to their subdetector
 - Person C: can look at monitoring plots

Web GUI via Jupyter Hub setup:







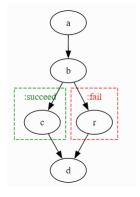
Further inspiration

Other Nifty Features

- Supports job submission to remote machines
- Workflows can support:

Trigger startup: on regular clock cycle
Trigger startup: on file appearance
Branching logic
Workflows nested in other workflows

• Local database to store/query workflow info maintained







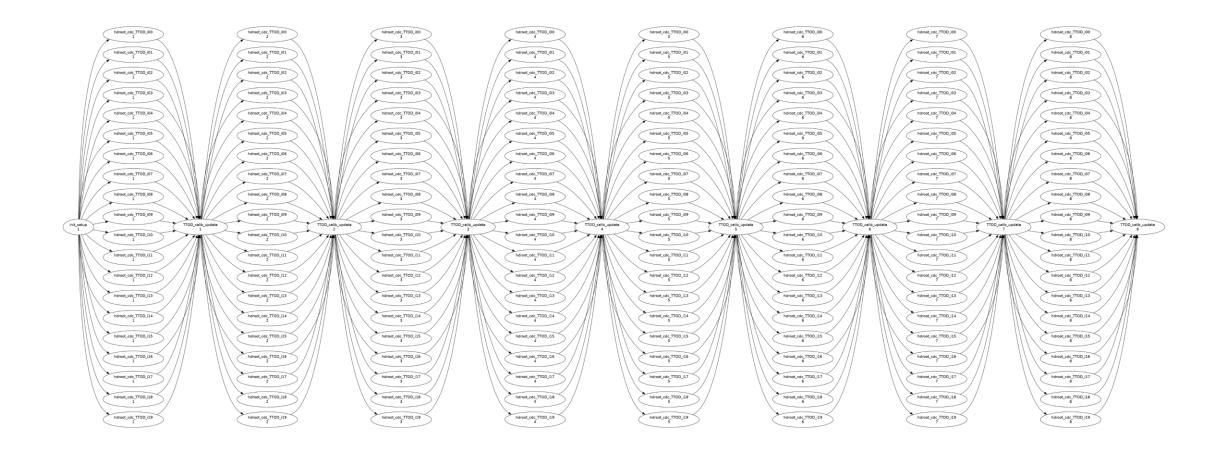
Summary

- Cylc seems like a great multipurpose tool https://cylc.github.io/
 - Functionality of swif2 (+more)
- Here: use for calibration workflows
 - A few more details here: <u>https://wiki.jlab.org/epsciwiki/index.php/File:Jz_gluex_calib_10.2.24.pdf</u>
- Potential for other use cases
 - MCWrapper offloading from OSG?
 - Hydra?
- Plan to put examples on wiki, Github
 - Let me know if there's interest in starting sooner



C Y I C

Backup: Full Workflow Graph

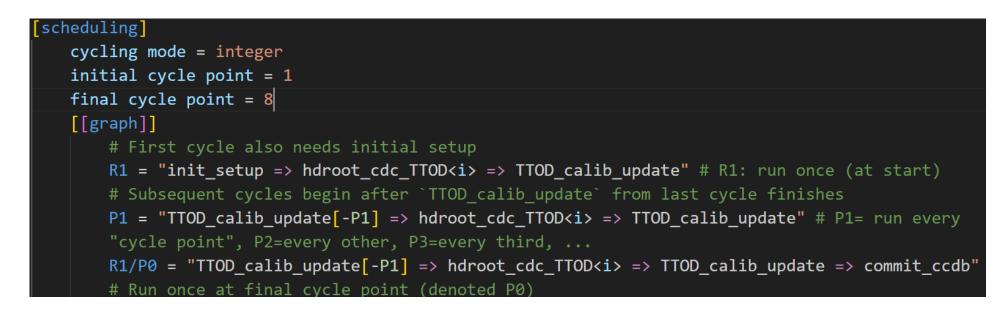




Backup: Defining Workflows

Workflows described in file called flow.cylc

Gives graph of tasks to run





Backup: Defining Workflows, cont.

Workflows described in file called flow.cylc

A single "task"

57	<pre>[[init_setup]]</pre>
58	inherit = gx_env
59	<pre>script = """</pre>
60	source gx_env.sh
61	ccdb_LocalUpdate.sh
62	cd \${CYLC_WORKFLOW_SHARE_DIR}
63	<pre>mkdir -p ccdb/ccdb_add ccdb/ccdb_dump \${ttod_topdir}/root</pre>
64	нин



Backup: Defining Workflows, cont.

Workflows described in file called flow.cylc

Defining how to run batch jobs

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```
# Reconstruction default job (MANY-THREADED!)
[[gx_recontask]]
    inherit = gx_env
    platform = jlab_slurm # Defined in my global.cylc file
    execution retry delays = 3*PT10S # I allow 3 timeout retries before declaring "failed"
    # Submit to the host system
    # job requires 5000MB of RAM and TMP disk, 16 CPUs
    [[directives]]]
        --mem = 5000 # MB
        --ntasks = 16 # Threads
        --time = 420 # walltime (minutes)
        --nodes = 1-1 # -N, --nodes=N, number of nodes on which to run (N = min[-max])
```

Backup: Defining Workflows, cont.

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Add inherit = gx_recontask to run as batch job (otherwise runs locally)

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