# Data & Analysis Preservation: status update

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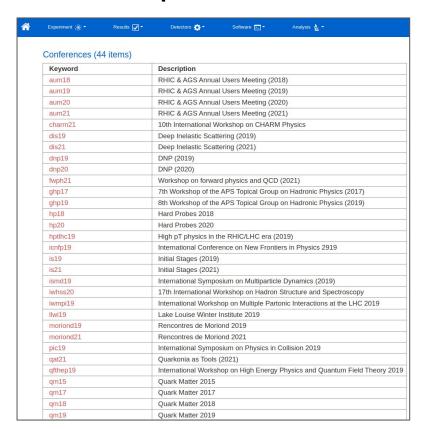


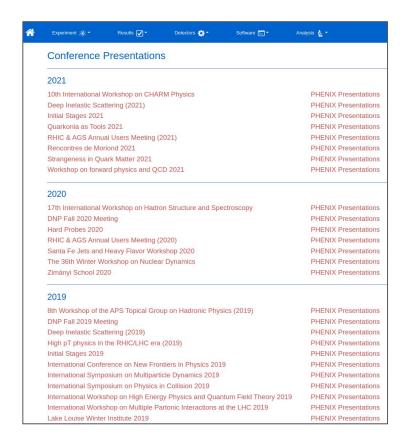
#### Overview

- Zenodo+Website
- HEPData
- Docker+REANA
  - CVMFS deployment
  - Progress with realistic payloads
  - Some residual technical work might be needed
- Summary and plans



## Zenodo uploads







#### Zenodo+Website

- Uploads ongoing (thanks Gabor)
  - Total of 44 conferences included
  - 2019, 2020 and 2021 are all covered
  - Six more conferences since last report + relevant keywords
- (Some) School'21 materials were added as well
  - DAP: 3 sets of slides also added to the PHENIX GitHub repo
- REANA page corrections





#### **HEPData**

- Master spreadsheet updated
- PPG071 (revision), PPG115 finalized
- Other active items in the pipeline: 026, 201, 238



## Docker: a recap (one last time)



- The core problem for containerizing the PHENIX software:
  - PHENIX software components evolved over ~20 years accumulating dependencies
  - Large number of dependencies on packages installed on the SDCC cluster resulted in two subsidiary issues
    - No "clean build" procedure available too many components to be covered
    - Large size of the sum of binaries too large for Docker
  - An attempt to build a smaller "custom image" by chasing down and isolating dependencies proved very time-consuming and not completely reliable
- The SDCC-provided images were developed in parallel now the preferred option
  - The idea is to have a complete snapshot of SDCC + PHENIX stack deployed to CVMFS
  - ...resulting in a compact image + 30GB of binaries available on demand
  - ...see next slide about CVMFS



#### **CVMFS**

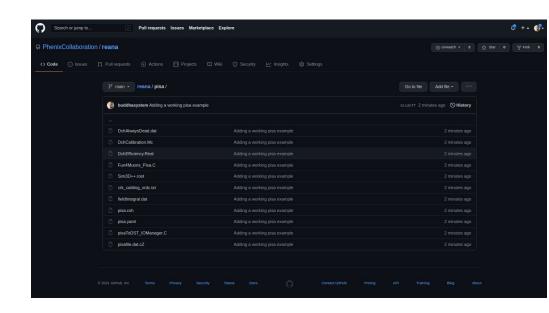


- The CernVM File System (originally developed to support the CERN VM project)
- A scalable, reliable and low-maintenance software distribution service
- The client uses outgoing HTTP connections only (no firewall issues)
- From the user perspective it is somewhat like read-only AFS
  - Key: globally visible (if mounted), can be used on clusters like REANA anywhere
  - Only admins can deploy to CVMFS
  - Can have a POSIX interface via FUSE (i.e. can browse it like most filesystems)
- Widely used in HEP everywhere
- Example of setting an environment variable in a user script:
   setenv OFFLINE\_MAIN /cvmfs/phenix.sdcc.bnl.gov/x8664\_sl7/release/release\_new/new

#### REANA: recent progress

reana

- REANA tutorials during the school went well
- SDCC made a few iterations of the PHENIX image with CVMFS-based provisioning of binaries - can be used both in REANA and Singularity
- The goal now is to expand validation with realistic macros and understand potential problems (e.g recently resolved the database driver issue)
- Added working examples to the repo
  - PISA to DST
  - Run 16 QA "prod\_output"





#### REANA "workspaces"



- The usual facility-wide filesystems are not visible from the worker nodes in the REANA cluster (i.e. no access to gpfs, nfs, afs etc)
- All the files needed by a workflow are meant to be staged in and staged out by the REANA client, before and after the workflow execution - and they reside in the scratch space known as "workspace"
- Each workspace is private for each instance of workflow
- The user can browse the workspaces using the Web browser and also the CLI client
- Example if you create multiple workflows under the same tag "my\_workflow", reana will instantiate instances indexed by an integer i.e. my\_workflow.1, my\_workflow.2, my\_workflow.3 etc. Still, each instance will only have access to its own workspace.



#### REANA I/O



- When staging in files from a "/" path rather than the current directory, a complex path will be created, relative to the workspace
  - /afs/rhic.bnl.gov/phenix/etc/odbc.ini ⇒ ./afs/rhic.bnl.gov/phenix/etc/odbc.ini
  - This pattern will also apply when staging data directly from GPFS
- For multi-GB upload packages the transfer will take a few minutes in our current REANA environment
- A complete folder can be staged in using the "directory" attribute in YAML
  - Care must be taken since directories can be quite large beware AFS may take forever,
     and the overall quota may be exceeded (TBD)



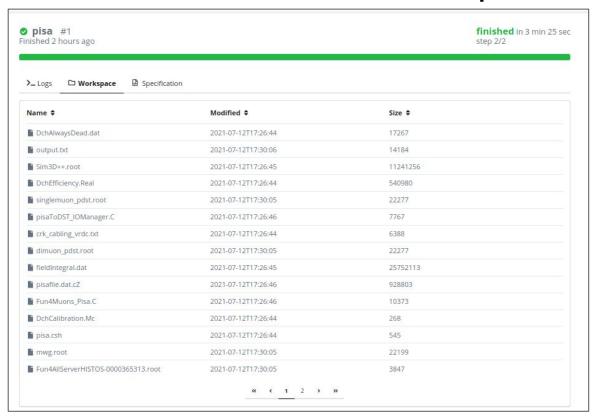
## REANA I/O (cont'd)



- The contents of a workspace can be updated as needed w/o having to create a completely new one - workflows can be restarted with new/updated data if needed
  - This is useful when there is a large amount of data already staged and the user needs to make a correction
- If the outputs are correctly defined, it takes a simple "download" command to get the results back to your workstation
- Alternatively, every individual file in the workspace can be downloaded if needed using the Web UI (through the browser)



## REANA: "PISA to DST workspace" example

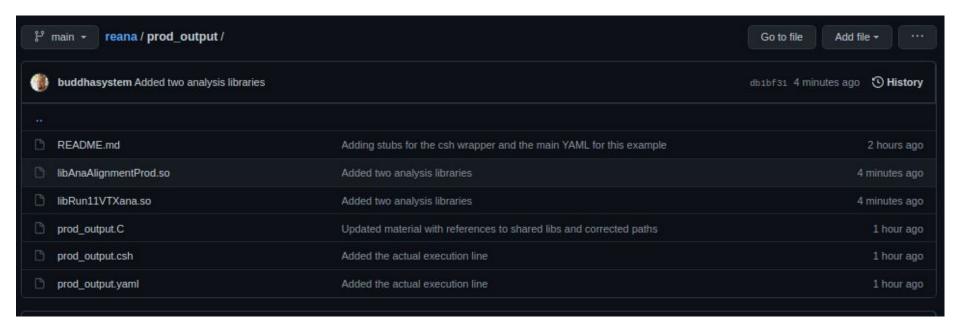


```
/afs/rhic.bnl.gov/phenix/etc/odbc.ini
- /afs/rhic.bnl.gov/phenix/etc/odbc.ini.phnxdbrcf2.orig
     /pisa.csh
/DchCalibration.Mc
     /pisafile.dat.cZ
specification:
     environment: 'registry.sdcc.bnl.gov/sdcc-fabric/rhic sl7 ext:1.3'
       - chmod +x ./pisa.csh
       - ./pisa.csh > output.txt
 - output.txt
  singlemuon_pdst.root
dimuon pdst.root
 - mwg.root
- Fun4AllServerHISTOS-0000365313.root
```

Files declared in the "outputs" section can be easily downloaded using the reana client: reana-client download -w pisa

## REANA: run16 QA example on GitHub

https://github.com/PhenixCollaboration/reana/tree/main/prod\_output



#### REANA submission for run16 QA

output.txttest 459208.root



```
version: 0.0.1
inputs:
 files:
 - /afs/rhic.bnl.gov/phenix/etc/odbc.ini
 - ./prod_output.csh
 - ./prod_output.C
  - ./libAnaAlignmentProd.so # any analysis-specific shared libraries can be organized optimally as folders etc
  - ./libRun11VTXana.so
  - /phenix/crs/agg/run16/run16AuAu_200GeV_CA_pro111_agg/CNT_MB/CNT_MB_run16AuAu_200GeV_CA_pro111-0000459208-9000.root
 - /phenix/crs/agg/run16/run16AuAu_200GeV_CA_pro111_agg/DST_SVX_MB/DST_SVX_MB_run16AuAu_200GeV_CA_pro111-0000459208-9000.root
  - /phenix/crs/agg/run16/run16AuAu_200GeV_CA_pro111_agg/DST_EVE_MB/DST_EVE_MB_run16AuAu_200GeV_CA_pro111-0000459208-9000.root
workflow:
type: serial
 specification:
  steps:
   - environment: 'registry.sdcc.bnl.gov/sdcc-fabric/rhic sl7 ext:1.3'
    commands:
    - chmod +x ./prod output.csh
    - mv ./phenix/crs/agg/run16/run16AuAu 200GeV CA pro111 agg/*/*.root .
    - rm -fr ./phenix
    - Is > output.txt
    - ./prod output.csh >> output.txt
outputs:
 files:
```



## **Automating REANA submission**

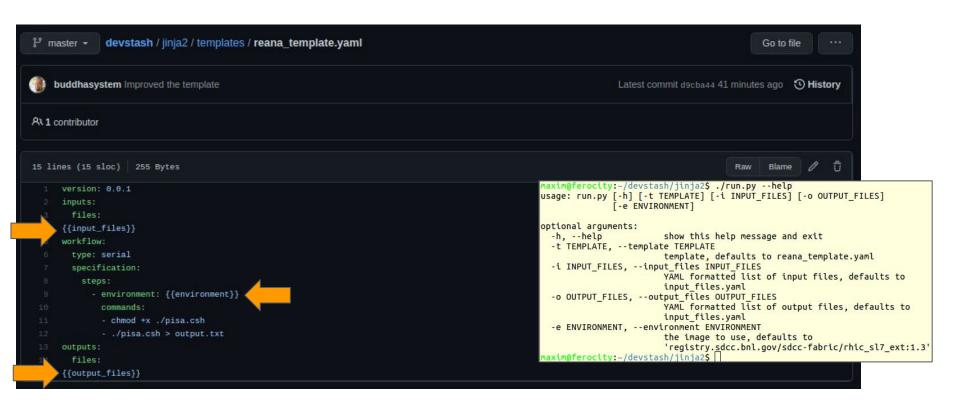


- If the number of input/output files is large, manual tweaking of the YAML file for REANA submission becomes tedious and error prone
- An initial simple prototype script has been created (and can be vastly improved)
  which allows to automatically fill out a template with blocks in input and output file
  definitions; based on the Jinja2 template engine which allows for rich logic to be
  included in the templates (inheritance, conditions, loops etc)
- Can be extended to handle wildcards in a folder, based on some criteria
  - o e.g. file0001-9999 can be included as inputs, and output names can be inferred
  - Need some use cases to move forward with implementation; what patterns in folder names and filenames need to be parsed?
- Total storage on the REANA cluster is limited so scripting will be needed for staging the data in and out in any case, and will need to be mindful of quotas



#### REANA templates and the helper script





## REANA submission: a possible design



- Create a base template specific to a particular analysis
  - e.g. references to specific macros and conditions, all local to the current folder
  - This makes the analysis case self-contained and suitable for preservation
  - o Input files (and references) are not included in the template (can be restored later if deleted)
- Have a companion script generate lists of inputs (and if needed outputs) based on wildcards or other similar criteria
- Have the template extrapolation script generate REANA submission files for large groups of inputs; record groupings in a unified log
- Submit workflows to REANA, perhaps with an additional wrapper
- Monitor execution and download outputs before staging in more data

#### REANA roadmap



- There has been a substantial investment of effort invested in the images and REANA
- Conservatively 80% of work on tuning Docker images has been done
- Simple but realistic PHENIX examples work
- To start seeing dividends of this work we need wider community engagement really the key to completing this work area and making it meaningful
- Should we organize a REANA seminar for the PHENIX community, as a follow up to the School tutorials?
- Can we identify ~2 PPGs who can be asked to "reanify" some parts of their analysis?
  - There are some direct benefits for the groups (self-documented reproducible procedures)
  - REANA take-up can be facilitated by providing templates for submission files



## Summary

- Complete Docker images good progress has been made
  - Initial REANA testing promising
  - Focus: setting up REANA environments for various use cases + automation
  - Participation of people doing analyses is crucial
  - Suggestions for use cases are welcome and needed
- HEPData, steady state effort (take some "pinging" to keep things moving)
- Presentations at the School, AUM and DPHEP were well received
- The PHENIX section of the NPP PAC is very positive, including the DAP mention
- Due to Gabor's effort there is progress in migration of conference materials to Zenodo, with curated keywords dependency on the legacy catalog is reduced
- Any new ideas re: new OpenData entries?

