The PHENIX DAP update

- Workshops
- HEPData/GitHub
- Website updates
- Zenodo, Keywords and Conferences
- REANA plans
- Status of PHENIX access to the Open Data portal

Maxim Potekhin (BNL, NPPS) 12/10/2020

Workshops

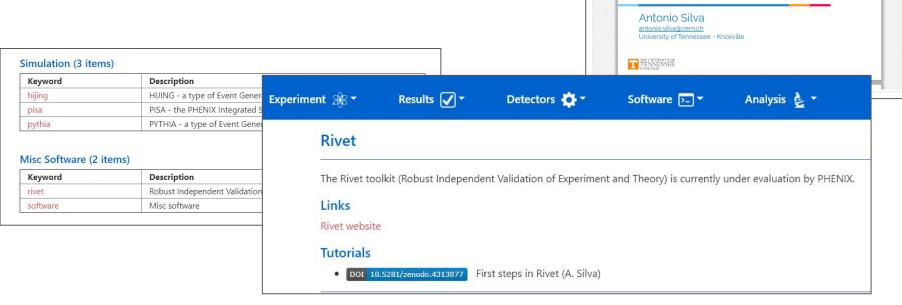
- Great many thanks from all of us to Christine for organizing these events for the benefit of the RHIC community
- Not lectures but hands-on workshops/tutorials on HEPData and Rivet
- HEPData: https://indico.bnl.gov/event/8843/timetable/ (11/10 & 11/17 2020)
 - o Org Committee: Christine Nattrass, Antonio Da Silva, Lauren Kasper, Andi Mankolli
- Rivet: https://indico.bnl.gov/event/8840/timetable/ (11/20-12/4 2020)
 - o Org Committee: Christine Nattrass, Antonio Da Silva, Christian Bierlich, Leif Lönnblad

Rivet, HEPData, YODA

- Integration with InspireHEP (ID)
- These topics are related since data exported from HEPData in the format called "YODA" are a part of input for Rivet. YODA is a set of analysis and histogramming classes, and the core of histogramming in Rivet
- There are *some* compatibility issues when exporting the data:
 - https://indico.cern.ch/event/808138/contributions/3364224/attachments/1851728/3040138/watt
 hepdata_may2019.pdf
 - https://conference.ippp.dur.ac.uk/event/875/contributions/4767/attachments/3898/4443/HepDat
 a Rivet 2020-01.pdf
- Discussions with Christine re: creating functional pages for our site (next slide)
- Interesting to see the scope of what's possible with Rivet in PHENIX
- Caveat: Rivet operates on fairly final data products which insulates it from the nitty-gritty of reconstruction and many analysis steps

Rivet content on the site

- Updated the Rivet page (in "Analysis")
- Uploaded slides to Zenodo
- Added "rivet" to the keyword list
- Work in progress, lots to do



zenodo

First steps in Rivet

An introduction to Rivet, with use cases emphasizing RHIC experiments

First steps in Rivet

Nov. 30, 2020

1 of 50 - + Automatic Zooms

December 9, 2020

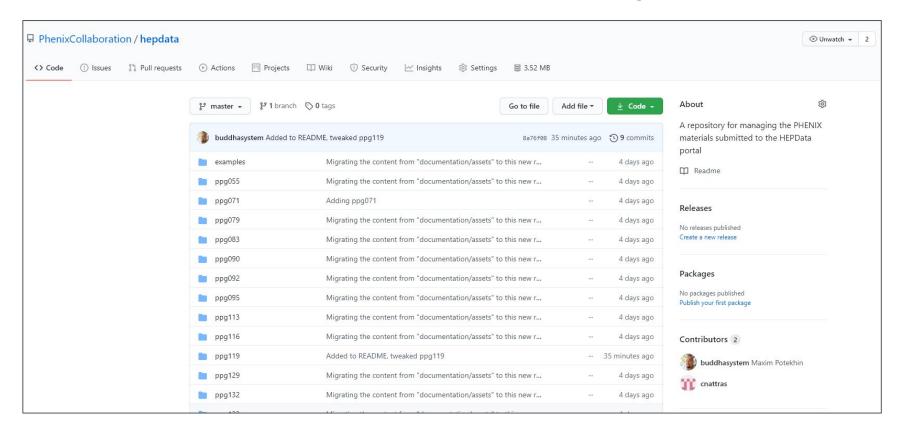
Presentation Open Access

22 8 B

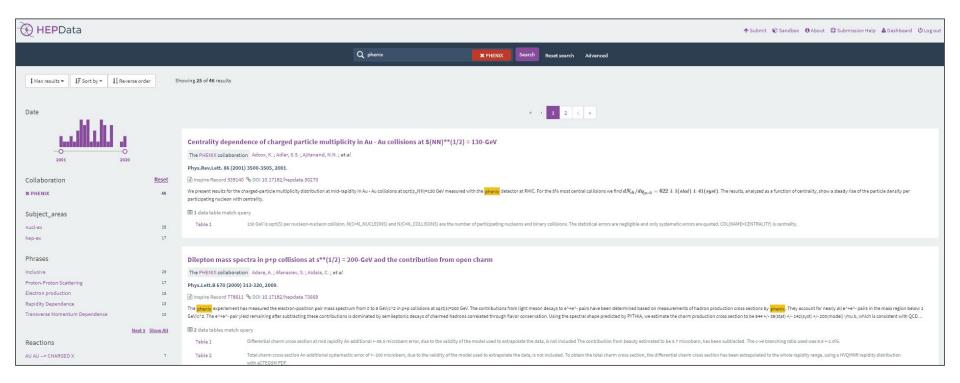
New Repository for HEPData

- HEPData work is progressing, active participants clone/fork the existing repo
 - Six active people in the past week
- The default "documentation" repo is an umbrella for all sorts of materials
- It has grown to a large size thanks in part to Stacyann's valuable effort to get our theses' materials organized in conjunction with Zenodo uploads
- But the repo doesn't scale and working with HEPData has become more cumbersome than it should be
- Solution: create a dedicated repo owned by PHENIX for clean and efficient access
 - It has been done: https://github.com/PhenixCollaboration/hepdata
 - Uniform naming scheme for the folders i.e. ppgXXX
- Links updated on the website

Repo contents: 25 submission packages+examples



Our HEPData content (46 items right now)



HEPData experience

- HEPData: the devil is in the detail. Progress has been made with users creating submission packages, but errors are still common and take time to fix
 - Quality of typeset (LaTeX) in various places on the page e.g. abstract, figure captions
 - More importantly, consistency of the numerical data components
 - Consistency (and availability) of thumbnails and main images needs to be checked
- Many careful "sandbox" checks are necessary to QA the materials
- In one case we had to update 3 times to get everything right (we want to avoid this, of course)
- Discovered that an older item (ppg071) needs some improvement, work in progress - thanks Krista!

HEPData: website updates etc

- Finally we started keeping track of the Inspire IDs as a part of a submission, as well as the designated IRC member
- Caveat: for smooth operation the reviewer needs to be registered on the portal under the same email as was used when the submission is defined
- Performed more updates of the HEPData policy page based on recent work and communication with participants of the effort
- Keeping track of progress in the README file in the repository

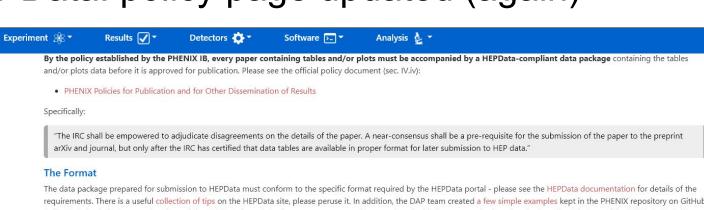
List of completed uploads

055, 079, 090, 092, 095, 113, 116, 129, 132, 133, 138, 146, 174, 182, 227, 228

In progress/recently finished

- 182: A.Khatiwada (prep, reviewer): khatiwada.ajeeta@gmail.com note: finalized
- 119: K.Smith (prep), Matt matthew.g.wysocki@gmail.com (reviewer) note: pending the reviewer's decision
- 071: K.Smith (prep), Tony afrawley@fsu.edu(reviewer) note: existing HEPData item being reviewed for completeness

HEPData: policy page updated (again)



requirements. There is a useful collection of tips on the HEPData site, please peruse it. In addition, the DAP team created a few simple examples kept in the PHENIX repository on GitHub to illustrate basic features and options of the HEPData format. Beginners are encouraged to experiment with these examples by using the "sandbox" feature of the HEPData Portal (see Appendix B below).

The basic idea of how a submission is structured is as follows. Data contents of each item included in the package (e.g. a plot) are described in a corresponding individual file formatted as YAML (e.g. if there are 5 plots in the paper you are expected to provide 5 YAML data files). In addition, a special YAML file submission.yaml describes the submission as a whole e.g. provides the names of the data and optional image files, list of keywords etc. It also contains an abstract (typically imported as LaTeX from the publication material); unfortunately, not every LaTeX feature will work correctly on HEPData and the output will need to be checked (see the "sandbox" reference below).

Since YAML allows comments - lines starting with a "#" sign - it is very easy to add any sort of extra information to submission.yaml that may be helpful for communication with members of the Collaboration, reviewers and for the workflow of the submission process in general. For example, it is necessary to provide the Inspire ID of the paper for the HEPData submission. It can be placed in a comment line. Also, including the PHENIX-internal PPG identifier is highly recommended as it reduces the chances of human error and facilitates communication. Both Inspire ID and the PPG identifier can be easily incorporated in the comment lines of the submission.yaml file mentioned above (i.e. in lines of text starting with "#"). There can be any number of comment lines. Including information about the designated reviewer (member of the IRC for the paper) as an additional comment line is encouraged. The following pattern of the top of the submission.vaml file may help illustrate this:

```
# PPG999
# InspireHEP: 99999
# Reviewing IRC member: M.Phenix mphenix@bnl.gov
```

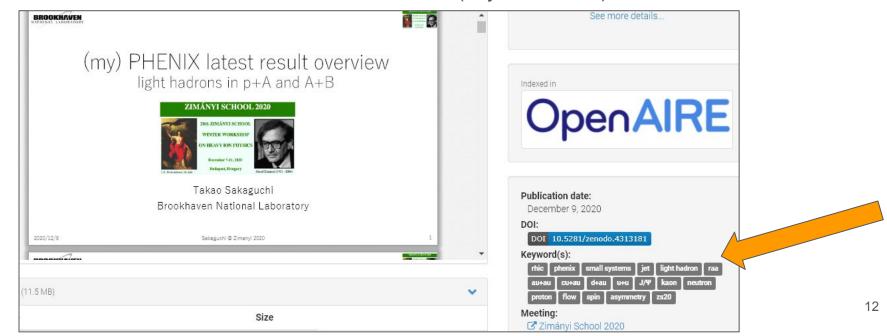
This is not to be confused with the comment attribute of the YAML file which almost always contains the abstract of the published paper, typically typeset in LaTeX:

HEPData content

- HEPData: naively, items simply reflect contents of papers
- What about supporting materials/data?
- There is no hard and fast policy suggested or enforced on the site
- In fact, different preparers can have different approaches
- Discussions took place lately
- Do we need to create basic guidelines and put them on the web page?

Keywords and Conferences

- Thanks to Gabor for reviewing Zenodo uploads and normalizing the list of keywords, the web page has been updated correspondingly
- Added Zimányi School 2020 to the conference page, seeded the Zenodo uploads for this conference with a PHENIX overview talk (keyword: zs20)



REANA revisited

- A couple of slides presented in the DAP meeting on 10/01/2020
 - See links to misc presentations in the slides from Oct.1 for more info
- http://reanahub.io/ "Reproducible Research Data Analysis Platform"
- Structured, unambiguous description of analysis workflows
 - There are two available workflow languages/schemas to choose from
 - Used in real-life complex analyses
 - Software environment (not just code) packaged as an image
 - Needs to run as a container on a cluster, currently at CERN, an option for local installation at BNL TBD
 - Helpful GUI

REANA - "Hello World" example: the description

version: 0.3.0 inputs: - code/helloworld.py - data/names.txt directories: - workflow/cwl parameters: input: workflow/cwl/helloworld-job.yml workflow: type: cwl file: workflow/cwl/helloworld.cwl outputs: files: - outputs/greetings.txt

sleeptime: 0
helloworld:
 class: File
 location: code/helloworld.py
inputfile:
 class: File
 location: data/names.txt

REANA - an example of a real-world analysis

```
version: 0.4.0
inputs:
files:
    - config/geantSim_TrackerPerformance.py
    - config/single_particle_trackFits.py
    - script/numHitsPerTrack.C
    - script/plot_single_particle_resolutions.py
parameters:
    events: 5000
    seed: 0123456
    particle: 13
    etamin: 0
    etamax: 6
    pt: 1000 2000 5000 10000 1000000 10000000
```

```
- environment: 'gitlab-registry.cern.ch/vavolkl/fcc-ubuntu:latest'
                                                                      Image
 - fccrun.py config/geantSim TrackerPerformance.py
   -N $events -s $seed --outName muons for seeding discrete pt.root
   --singlePart --particle $particle --etaMin $etamin --etaMax $etamax
   --discretePt --pt $pt
  | tee simulation.log 2> simulation.err
 - root -b 'script/numHitsPerTrack.C("muons for seeding discrete pt.root")'
  | tee plot-tracker-hits.log 2> plot-tracker-hits.err
 - fccrun.py config/single particle trackFits.py
   --outputfile single particle resolutions.root
  I tee fit.log 2> fit.err
  | tee plot-reconstructed-tracks.log 2> plot-reconstructed-tracks.err
```

REANA - another complex example



REANA: plans

- REANA is a complex system, need to evaluate cost-benefit for PHENIX
 - Learning curve is a little steep
 - State of the art, well supported
 - Part of a wider question of the DAP priorities
 - Potential for cross-experiment aspect ???(sPHENIX, STAR, EIC)??? recent Rivet/HEPData workshops are encouraging
- Ran a few simple tests on the CERN cluster, need to develop expertise
- Since PHENIX analysis are BNL-local due to various dependencies, the task is to build images that can run locally (AFS etc)
- Currently looking at Singularity as the main BNL framework to run containers
- If enough progress is made within weeks (perhaps months due to complexity) a request may be made to SDCC to create a local REANA cluster

Open Data (the CERN portal)

- Since the last DAP meeting we have received storage allocation: /eos/opendata/phenix/upload/
- I have yet to test uploads, on my to-do list, will report

Summary

- Steady progress with HEPData
 - Dedicated repository created
 - Website pages updated with latest instructions
 - A few packages are in the pipeline
- Two recent helpful workshops on HEPData and Rivet (C.Nattrass)
- Keyword updates, additions
- Good progress with conference material uploads to Zenodo (G.David)
- Thesis uploads and archiving on GitHub (Stacyann)
- REANA initial exploration done, evaluation for PHENIX to follow
- The CERN Open Data portal access: storage defined, testing to be done