Brief introduction to justIN



New concepts

- I highly encourage you to read about these new concepts
- Location management **Rucio** <u>https://rucio.github.io</u>/
- Data catalog management MetaCat <u>https://metacat.readthedocs.io/en/latest/</u>
- Workflow management justIN <u>https://justin.dune.hep.ac.uk/docs/</u>







A few changes

- As you may have noticed, we are in a transition period at the production level we are transitioning from POMS to justIN
- At the user level we are transitioning from SAM data catalog to **MetaCat** (**Rucio**)
- To retrieve file location users used to do samweb get-file-access-url
- SAM locations are not supported anymore
- Instead, user needs to get the file location from Rucio \bullet
- rucio -a \$USER list-file-replicas fardethd:nu dune10kt 1x2x6 1062 166 20230823T120323Z gen g4 detsim hitreco 2024 0222T231742Z reco2.root --pnfs

91/66/

231742Z reco2.root root://fndca1.fnal.gov:1094/pnfs/fnal.gov/usr/dune/tape backed/dunepro// fardet-hd/full-reconstructed/2024/mc/out1/fd mc 2023a reco2/00/00/10/62/ 231742Z reco2.root 1709158151

- root://dcdndoor.sdcc.bnl.gov:1094/pnfs/sdcc.bnl.gov/data/dune/RSE/fardet-hd/
- nu dune10kt 1x2x6 1062 166 20230823T120323Z gen g4 detsim hitreco 20240222T
- nu dune10kt 1x2x6 1062 166 20230823T120323Z gen g4 detsim hitreco 20240222T



justIN

- justIN is a workflow system that processes data by satisfying the requirements of data location/data catalog, software distribution and job submission to the grid
- There is documentation on how to get started with justIN at https://justin.dune.hep.ac.uk/docs/tutorials.dune.md
- If you are a member of DUNE and have a Fermilab computing account, you should be able to run this brief tutorial
- Before this, please READ the prerequisites carefully and check if you can login https://justin.dune.hep.ac.uk/docs/tutorials.dune.md and CONTINUE to read the DUNE tutorial to make sure you can run the examples
- This tutorial will repeat every single step discussed in DUNE justIN tutorial with specific DUNE examples



- We will show how to do three basic tasks
 - release and you don't actually modify any of it
 - release and you want to use a customized FCL file
 - release and you want to use customized code

• Process data (submit a job to the grid) if you are using code from the base

• Process data (submit a job to the grid) if you are using code from the base

• Process data (submit a job to the grid) if you are NOT using code from the base



- actually modify any of it
 - data sets
 - Let's say you want to run mergeana for electron neutrinos,
 - Where is the data?
 - In DUNE we provided datasets to easily identify a collection of files

fardet-hd:fardet-hd fd mc 2023a reco2 fullue dune10kt 1x2x6 out1 validation

Process data (submit a job to the grid) if you are using code from the base release and you don't

• Once you have identified what data you want to process, you can see the most recent data (official data) sets available at https://wiki.dunescience.org/wiki/Data Collections Manager/

reconstructed v09 81 00d02 standard reco2 dune10kt nu 1x2x6 prodgenie n

Dataset names tend to be self explanatory and includes the type of detector, which fcls files were used to produce, software version, data tier, and tag, in this case, the tag is validation



- and you don't actually modify any of it
 - Lets try to process mergeana in the first 100 files that in the data sets,
 - first 100 files of a given data set the query would be something like

"files from fardet-hd:fardet-hd fd mc 2023a reco2 fullreconstructed v09 81 00d02 standard reco2 dune10kt nu 1x2x6 pro dgenie nue dune10kt 1x2x6 out1 validation limit 100 ordered"

- The flag "ordered" is crucial to ensure reproducibility
- Now, how do you process this using justIN?
- You need to provide a jobscirpt and run a justIN command line

• Process data (submit a job to the grid) if you are just using code from the base release

MetaCat relies on MQL queries to select a collection of files in this case to select the



- release and you don't actually modify any of it
 - The basic tasks of your jobscript are:
 - Set software env
 - Find where the data is
 - Process the data
 - Save the output in a given location
 - Examples of jobscripts can be found at https://github.com/DUNE/dune-prod-utils/tree/main/justIN-examples

• Process data (submit a job to the grid) if you are just using code from the base





- release and you don't actually modify any of it
 - An example of a jobscript can be found at https://github.com/DUNE/dune-prod-utils/blob/main/justIN-examples/submit_ana.jobscript
 - Lets look at some of the code there

19	<pre># fcl file and DUNE software version/qual</pre>
20	FCL_FILE=\${FCL_FILE:-standard_ana_dune10}
21	DUNE_VERSION=\${DUNE_VERSION:-v09_81_00d02
22	DUNE_QUALIFIER=\${DUNE_QUALIFIER:-e26:prot
23	
44	# Setup DUNE environment
45	<pre>source /cvmfs/dune.opensciencegrid.org/p</pre>
46	setup dunesw "\$DUNE_VERSION" -q "\$DUNE_G
47	
61	
62	lar -c \$FCL_FILE \$events_option -o \$outF:
63)

• Process data (submit a job to the grid) if you are just using code from the base

lifier to be used kt_1x2x6.fcl} 2} f}

products/dune/setup_dune.sh QUALIFIER"

ile "\$pfn" > \${fname}_ana_\${now}.log 2>&1



- modify any of it
 - As always, before sending 100s of jobs to the grid, it is highly recommended to test it interactively
 - the DUNE justIN tutorial
 - blob/main/justIN-examples/submit ana.jobscript

```
$ source /cvmfs/dune.opensciencegrid.org/products/dune/setup_dune.sh
$ setup justin
$ justin-test-jobscript --mql "files from fardet-hd:fardet-
hd fd mc 2023a reco2 full-
kt 1x2x6 out1 validation skip 10 limit 1 ordered " --jobscript
submit ana.jobscript --env NUM EVENTS=3
```

Process data (submit a job to the grid) if you are just using code from the base release and you don't actually

AT THIS POINT, WE EXPECT THAT YOU HAVE SUCCESSFULLY REPRODUCED THE EXAMPLES IN

To test your jobscript interactively you can use the example from <u>https://github.com/DUNE/dune-prod-utils/</u>

reconstructed v09 81 00d02 standard reco2 dune10kt nu 1x2x6 prodgenie nu dune10



- actually modify any of it
 - If your interactively test was successful you can proceed to send your jobs

• Crucial, make sure you have a valid proxy \$ export ROLE=Analysis \$ voms-proxy-init -rfc -noregen -voms=dune:/dune/Role=\$ROLE -valid 120:00

- scratch/users/\$USER
- \$ USERF=\$USER
- FNALURL='https://fndcadoor.fnal.gov:2880/dune/scratch/users' Ş

\$ justin simple-workflow --mql "files from fardet-hd:fardethd fd mc 2023a reco2 fullreconstructed v09 81 00d02 standard reco2 dune10kt nu 1x2x6 prodgenie nu dune10 kt 1x2x6 out1 validation skip 5 limit 5 ordered " --jobscript submit ana.jobscript --rss-mb 4000 --output-pattern "*_ana_*.root:\$FNALURL/\$USERF"

Process data (submit a job to the grid) if you are just using code from the base release and you don't

Remember you have to specify where the output should go, for now the output can only go to /pnfs/dune/



actually modify any of it

```
$ USERF=$USER
```

\$ FNALURL='https://fndcadoor.fnal.gov:2880/dune/scratch/users' \$ justin simple-workflow --mql "files from fardet-hd:fardethd fd mc 2023a reco2 fullreconstructed v09 81 00d02 standard reco2 dune10kt nu 1x2x6 prodgenie nu dune 10kt 1x2x6 out1 validation skip 5 limit 5 ordered " --jobscript submit ana.jobscript --rss-mb 4000 --output-pattern "* ana *.root:\$FNALURL/ \$USERF"

method=list-workflows

Process data (submit a job to the grid) if you are just using code from the base release and you don't

• You can look at your job status by using justIN dashboard <u>https://justin.dune.hep.ac.uk/dashboard/?</u>



- customized FCL file
- tutorial
- the DUNE justIN tutorial you need to tar the files needed and put them in cvmfs

```
$ tar cvz my fcls.tar my fcls
 source /cvmfs/dune.opensciencegrid.org/products/dune/setup dune.sh
 setup justin
Ş
$ rm -f /tmp/x509up u`id -u`
$ kx509
$INPUT TAR DIR LOCAL=`justin-cvmfs-upload my fcls.tar`
```

Wait a few minutes to check the files

\$ ls -l \$INPUT TAR DIR LOCAL

Process data (submit a job to the grid) if you are using code from the base release and you want to use a

• To do that, the best is to use the Rapid Code Distribution Service (RCDS) via cvmfs as explained in the

Let's say you have a customized FCL file that you need to run over some datasets. As per instruction from



- customized FCL file
- <u>submit_local_fcl.jobscript</u>
- The key part of the code is the following

19	<pre># fcl file and DUNE software ver</pre>
20	FCL_FILE=\${FCL_FILE:-\$INPUT_TAR_
21	<pre>DUNE_VERSION=\${DUNE_VERSION:-v09</pre>
22	DUNE_QUALIFIER=\${DUNE_QUALIFIER:
23	

\$ justin simple-workflow --mql "files from fardet-hd:fardethd fd mc 2023a reco2 fullkt 1x2x6 out1 validation skip 5 limit 5 ordered " --jobscript submit local fcl.jobscript --rss-mb 4000 --env INPUT TAR DIR LOCAL="\$INPUT TAR DIR LOCAL"

Process data (submit a job to the grid) if you are using code from the base release and you want to use a

• You can look at the example at <u>https://github.com/DUNE/dune-prod-utils/blob/main/justIN-examples/</u>

```
rsion/qualifier to be used
_DIR_LOCAL/my_code/my_ana.fcl}
9_81_00d02}
:-e26:prof}
```

```
reconstructed v09 81 00d02 standard reco2 dune10kt nu 1x2x6 prodgenie nu dune10
```



- Process data (submit a job to the grid) if you a to use customized code
- Probably you are developing some reconstruction alg and you want to check the results in a large sample, before committing your software to GitHub
- You can use your customized software (e.g. local installation of dunereco) and use justIN to process the data with your new LArSoft module
- Similar to the previous part, you will need to provide all pieces in a tar file and put them in cvmfs

\$ tar cvz my_code.tar my_code

• Here my_code.tar includes a directory with my_fcls files and one with my local products (e.g. localProducts_larsoft_v09_85_00_e26_prof) this is similar to what you used to do when using jobsub and using customized code

Process data (submit a job to the grid) if you are NOT using code from the base release and you want



to use customized code

```
• Next step is to send your tar file to cvmfs
$ tar cvz my code.tar my code
 source /cvmfs/dune.opensciencegrid.org/products/dune/setup dune.sh
Ş
 setup justin
Ş
$ rm -f /tmp/x509up u`id -u`
$ kx509
$INPUT TAR DIR LOCAL=`justin-cvmfs-upload my code.tar`
```

- Wait a few minutes to check the files
- \$ ls -l \$INPUT TAR DIR LOCAL

Process data (submit a job to the grid) if you are NOT using code from the base release and you want



- use customized software
- Next step is to send your tar file to cvmfs
- Once you files appear in cvmfs you can try use the following example

19	<pre># fcl file and DUNE software version/quali</pre>
20	<pre>FCL_FILE=\${FCL_FILE:-\$INPUT_TAR_DIR_LOCAL/</pre>
21	<pre>DUNE_VERSION=\${DUNE_VERSION:-v09_85_00d00}</pre>

- 22 DUNE_QUALIFIER=\${DUNE_QUALIFIER:-e26:prof}
- # Setup DUNE environment 44
- source /cvmfs/dune.opensciencegrid.org/products/dune/setup_dune.sh 45
- 46
- setup dunesw "\$DUNE_VERSION" -q "\$DUNE_QUALIFIER" 47 48 mrbslp

Process data (submit a job to the grid) if you are just using code from the base release and you want to

https://github.com/DUNE/dune-prod-utils/blob/main/justIN-examples/submit_local_code.jobscript

fier to be used my_code/fcls/my_reco.fcl}

source \$INPUT_TAR_DIR_LOCAL/my_code/localProducts_larsoft_v09_85_00_e26_prof/setup



- to use customized software
- Next step is to send your tar file to cvmfs
- Once your files appear in cvmfs you can try the following example

\$ justin simple-workflow --mql "files from fardet-hd:fardethd fd mc 2023a hithiguera/data collections/justIN/submit local code.jobscript --env INPUT TAR DIR LOCAL="\$INPUT TAR DIR LOCAL"

Process data (submit a job to the grid) if you are NOT using code from the base release and you want

```
<u>https://github.com/DUNE/dune-prod-utils/blob/main/justIN-examples/submit_local_code.jobscript</u>
```

```
reconstructed v09 78 01d01 standard recol dune10kt 1x2x6 prodgenie nu dune10k
t 1x2x6 out1 v1 official limit 10 ordered" --jobscript /exp/dune/data/users/
```



Questions

- Use the #computing-worfklow slack channel
- Justin commands
- \$ export RUCIO_ACCOUNT= \$USER
- \$ finish-workflow --workflow-id <ID>

