justIN/GPUs mini tutorial + questions

Andrew McNab University of Manchester





GPUs mini tutorial + questions

- justIN recap
- GPU resources available
- Standard justIN hello world
- GPU hello world to the grid
- GPU hello world to NERSC/Perlmutter
- Questions from Computing to AI/ML user communities



justIN recap

- justIN was developed by DUNE as a workflow management system which was designed to work with MetaCat, Rucio, and GlideinWMS from day one
 - MetaCat = file catalogue: "what files are like this?"
 - Rucio = replica catalogue: "where are copies of this file?"
 - GlideInWMS = runs jobs at sites for justIN
- You ask justIN to run copies of your job to process all the files that match some query and it magically does it for you
- You don't need to know where things run or where files are



Rucio scopes and justIN

- justIN leverages Rucio's scopes to decide who can do what
 - Each file in Rucio is in a scope so we say SCOPE:FILENAME
- justIN says that each scope is owned by a group
 - Only people in that group can create workflows that save output files in the scope
- Currently two active groups, more to follow:
 - /dune = everyone in DUNE
 - /dune/production = just the production team
- Scopes will let us delegate control to working groups and users



GPU resources available

- GPUs are available via justIN at
 - NERSC/Perlmutter
 - and on the grid at Manchester and QMUL in the UK
 - RAL-PPD has them online and they are being added to OSG
 - More welcome please get in touch so we can add them
- At NERSC, DUNE has an annual allocation
 - Currently 20,000 GPU hours with A100s, but we can request more

- But only workflows writing to /dune/production scopes currently enabled
- At the Grid sites, it's first come first served,
 - They tend to be underused and all /dune can access them
 - eg Manchester has 36 Tesla T4s = 316,000 GPU hours spread over a year



Standard justIN hello world

- If you're going to try this out, please follow the justIN tutorial first
 - It includes a GPU section now
 - https://justin.dune.hep.ac.uk/docs/tutorials.dune.md
- In short, you submit a workflow by giving the jobscript you have written in Bash and specifying any options to justIN
- If you have a Hello World jobscript, you can run 10 copies like this:
 - justin simple-workflow --monte-carlo 10 --jobscript hello-world.jobscript
- --monte-carlo is a fake query that just gives you the numbers 01 to 10
- The --mql option can be used to query MetaCat give a list of files instead



Workflow outputs

- Each job in a workflow produces
 - A jobscript log file, the tail of which can be viewed in the dashboard
 - A .logs.tgz file put in Rucio managed storage for ~1 month
 - Contains all files matching *.log including jobscript.log
 - Any output files you nominate via patterns to be put in Rucio storage
 - (Or to dCache scratch disk at Fermilab)
- You can find all these files through the justIN dashboard or in Rucio datasets
 - See the tutorial and documentation for details



GPU hello world

- Fetch https://github.com/DUNE/dune-justin/blob/main/testing/hello-gpu.jobscript
- Run 10 copies like this:
 - justin simple-workflow --gpu --monte-carlo 10 --jobscript hello-gpu.jobscript
- This does the same as Hello World, but the jobs only run on worker nodes with GPUs
 - HTCondor/GlideInWMS is told to allocate 1 GPU to the job
 - You can request CPU memory and number of CPUs, but not yet:
 - GPU memory or GPU versions



GPU environment

- justIN asks HTCondor/GlideInWMS to allocate a GPU from the pool the pilot job has got from the local batch system
- The local batch system might hide other GPUs on the worker node from us, or it might not
- You might be running alongside other GPU jobs on the worker node from other experiments or inside the DUNE pilot job
- justIN runs your jobscript inside an Apptainer container with --nv
 - This maps the /dev/nvidia devices and correct CUDA files from the worker node into the container, and sets LD LIBRARY PATH
 - So we do not need to put NVIDIA licensed libraries in cvmfs



GPU job status page (top half)

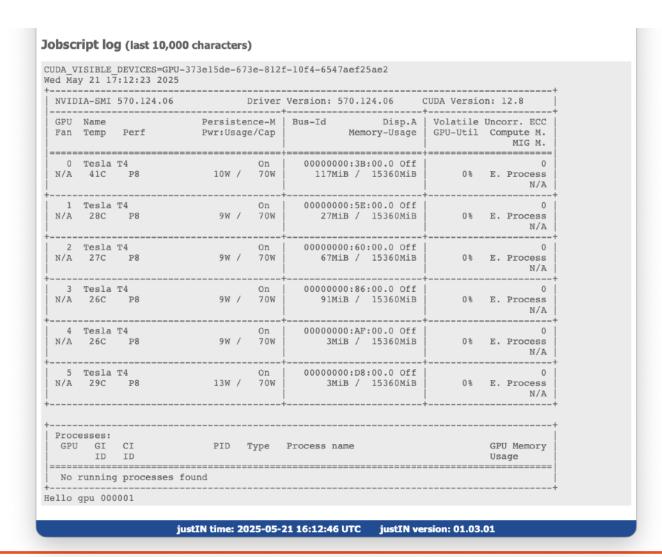


	I	
Site	UK_Manchester	
Entry	IceCube_UK_Manchester_ce01_gpu	
Last heartbeat	2025-05-21 16:12:40	
From worker node	Hostname	wn5915090.tier2.hep.manchester.ac.uk
	cpuinfo	Intel(R) Xeon(R) Silver 4210R CPU @ 2.40GHz
	OS release	Scientific Linux release 7.9 (Nitrogen)
	Processors	1
	RSS bytes	2097152000 (2000 MiB)
	Wall seconds limit	259200 (72 hours)
	GPU	Tesla T4 570.124.06 7.5 90.04.38.00.03 14913MiB
	Inner Apptainer?	True
Job state	finished	
Allocator name	justin-allocator-pro.dune.hep.ac.uk	



GPU job status page (bottom half)

- This is the output of the GPU Hello World jobscript
- It's mostly just these two commands:
- # Check the GPU environment
 printenv | grep -i cuda
 nvidia-smi
- CUDA_VISIBLE_DEVICES is set to tell you which of the GPUs you can use if more than one are visible
 - Depends on the local set up



GPU at NERSC hello world

- Run 10 copies like this:
 - justin simple-workflow --gpu --monte-carlo 10 --jobscript hello-gpu.jobscript --scope testpro --site US NERSC-GPU
- This forces the jobs to run with a GPU each at NERSC/Perlmutter
 - HTCondor / GlideInWMS / HEPCloud does some special magic!

- You can get the outputs as normal
- We currently have to give a Production scope (eg testpro)
 - Which means you need to be in the Production team
 - Since we have a finite amount of NERSC quota, we need ensure it does not get all used up accidentally in one go
 - With Grid you have more time to spot a mistake and kill a bad workflow



Questions

- What do you need in terms of limiting matches to
 - GPUs by GPU memory
 - GPUs by GPU hardware model
 - How to express that?
 - Other GPU/driver specific things?
 - This gets you less GPUs of course!
- Who needs to be able to use GPUs at NERSC?
 - Production team vs working groups' production teams vs individual users



Summary

- We have useful amounts of GPU capacity available
 - At multiple Grid sites
 - At NERSC/Perlmutter
- All of which can be accessed through the standard justIN interface with a trivial change to the submission command
- We will be able to use the existing justIN/Rucio mechanisms for controlling access via groups to limit usage of the DUNE NERSC quota
- But Computing needs help in gathering requirements and use cases to determine what other features are needed

