



HSF SEMINAR – SOFTWARE PACKAGING • 24 JUNE 2026

Packaging and distributing the HEP ecosystem on conda-forge

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CERN • University of Wisconsin–Madison

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Data Science Institute
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- Will break this down into three parts:



Tooling

install & manage
environments



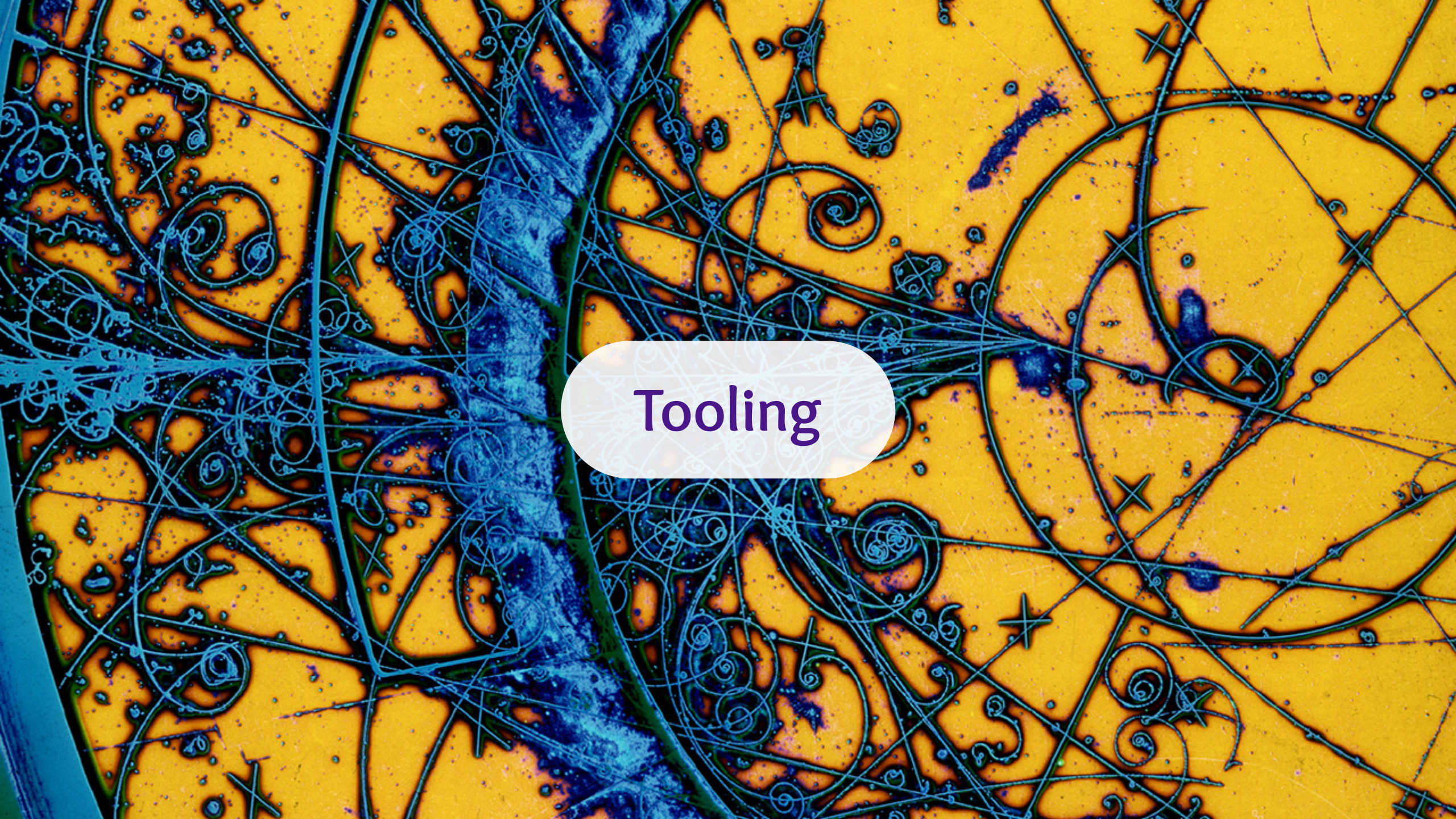
Packages

recipes & binaries



Development

build & iterate



Tooling

What is the conda ecosystem?

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4. Tools for building packages:

- `conda-build`
- `rattler-build`



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- `conda-build`
- `rattler-build`

5. Distributed cyberinfrastructure for building conda packages coherently:

<https://github.com/conda-forge/>





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3. Anaconda, Inc. (the company)
 - conda is a community project with a varied elected steering council
 - conda-forge packages are **free and open source**
 - Anaconda supports conda-forge, but is only a small fraction of the community
 - Anaconda's paid offerings are not interesting to HEP users



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4. Something new

It's been a while...

“

Looking back at your PyHEP 2019 conda-forge talk (interesting to see how much has stayed the same over 6+ years)

”



Packaging for Python and Beyond — PyHEP 2019



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- Pixi provides a lot of "user experience" improvements

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- Workspace model of working
 - Add a `pixi.toml` Pixi manifest to describe the software you need
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 - Ensures that everyone has the same software installed^{*}
 - This should typically be committed to the repository[†]



^{*} Assuming the same OS (Linux/macOS) and CPU architecture.

[†] Tools like *renovate* can be used to automatically update the lock file periodically.

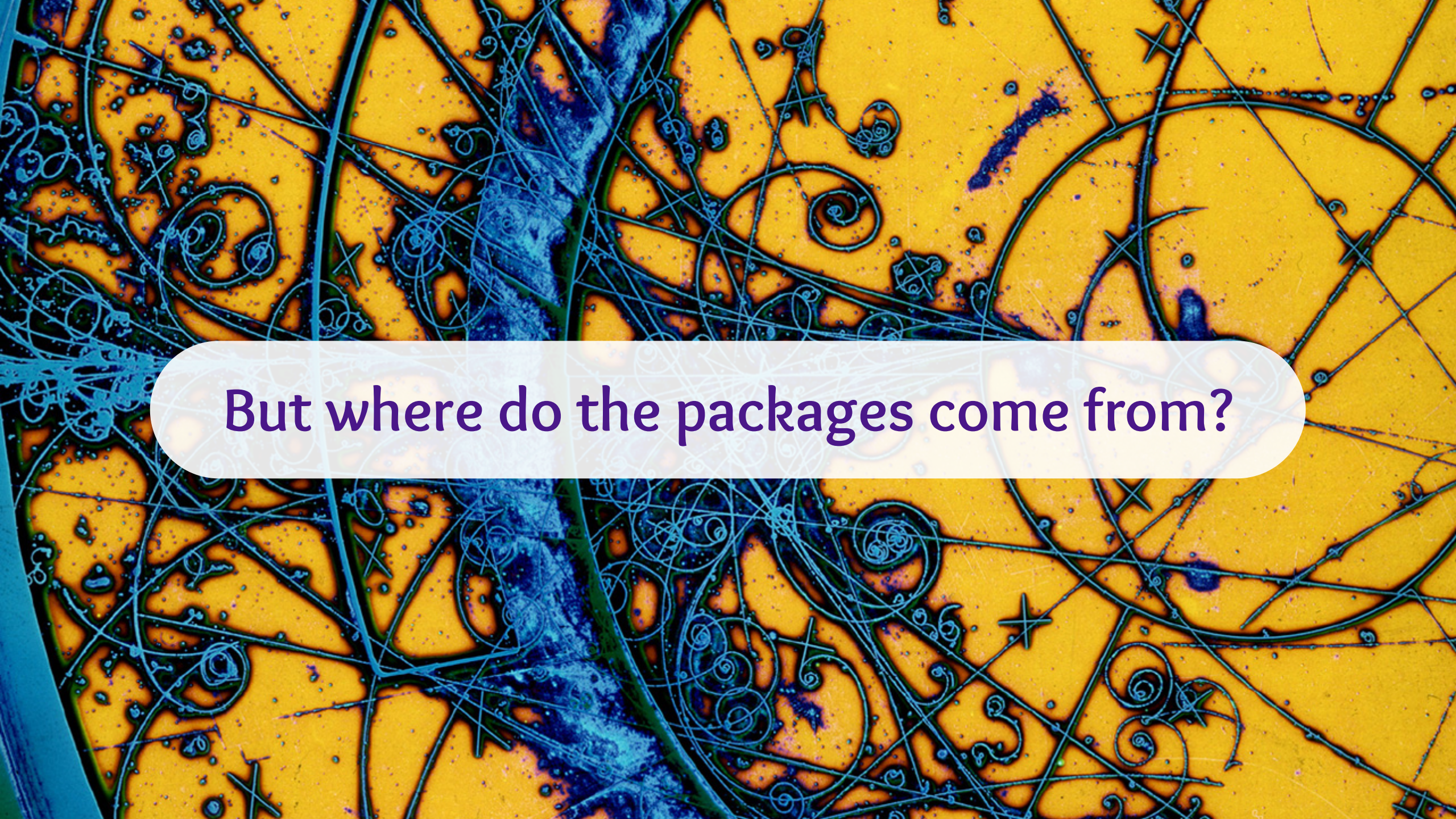
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- ✂️ Pixi takes care of managing all of the software environments for you ✂️



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But where do the packages come from?



Getting packages

- How to build software ("recipes")
- How to avoid building software ("binaries")
- How to know it will work together
 - Build the entire universe for every cosmetic change? (à la Nix[†])
 - Rely on compatibility between binaries (à la conda-forge)

[†] Nix is an excellent tool and it's a perfectly valid approach, but I think it's the wrong one for HEP.



Binary distribution

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- Potentially a different model than what you're used to, but is a very pragmatic approach
- For two binaries to be compatible it doesn't matter:[†]
 - Which C++ compiler was used to build them
 - Which C++ standard is used
 - Which Linux distribution you use

[†] This is massively over simplified but we don't have time to go into the details here.



- Shared CI and distribution infrastructure.
- ~7800 contributors, 33,000+ packages, 43 billion+ downloads
 - contributors maintain one or more packages
 - core team maintains the infrastructure and keeps the ecosystem healthy
- Heavy use of automation to manage version updates, rebuilds, and ABI change "migrations"
- Isn't frozen: you can contribute to add / update / fix packages
 - Uploaded binaries are immutable, associated metadata isn't
- HEP leadership:
 - Chris Burr is member of core leadership team
 - Chris Burr, Matthew Feickert are members of conda-forge/staged-recipes review team

The HEP Packaging Coordination project

- A community project to get **as much HEP software as possible** onto conda-forge.
- Directed by: Chris Burr, Matthew Feickert, Lindsey Gray, Giordon Stark, ...you!
- Contributors across HEP: ATLAS, Belle II, CMS, DIRAC, IRIS-HEP, LEGEND, LHCb, ROOT, Scikit-HEP, SHiP, theory/pheno...
- **120+ HEP packages** already: ROOT, Pythia8, FastJet, Awkward Array, Rivet, CMS Combine, ...
- Installing should be trivial

The screenshot displays the 'Analysis' section of the hep-packaging-coordination website. It features a table with columns: Name, Feedstock, Output, Downloads, Version, Platforms, and Open PRs. The table lists various HEP packages such as collier, contur, cuttools, eko, emela, evermore, ff, iregi, lhpdf, looptools, mpfun90, ninja-hep-ph, nnpdf, oneloop, pineappl, pytha, pysiha, and qcdloop. Each row provides details about the package's feedstock, output, download count, version, supported platforms, and the number of open pull requests.

Name	Feedstock	Output	Downloads	Version	Platforms	Open PRs
collier	feedstock/collier	recipe/collier	downloads: 164	conda-forge/collier		
collier		recipe/collier	downloads: 94	conda-forge/collier		
contur	feedstock/contur	recipe/contur	downloads: 284	conda-forge/contur		
cuttools	feedstock/cuttools-static	recipe/cuttools-static	downloads: 154	conda-forge/cuttools		
eko	feedstock/eko	recipe/eko	downloads: 534	conda-forge/eko		
emela	feedstock/emela	recipe/emela	downloads: 114	conda-forge/emela		
evermore	feedstock/evermore	recipe/evermore	downloads: 744	conda-forge/evermore		
ff	feedstock/ff-static	recipe/ff-static	downloads: 744	conda-forge/ff		
iregi	feedstock/iregi-static	recipe/iregi-static	downloads: 844	conda-forge/iregi		
lhpdf	feedstock/lhpdf	recipe/lhpdf				
looptools	feedstock/looptools-static	recipe/looptools-static				
mpfun90	feedstock/mpfun90	recipe/mpfun90				
ninja-hep-ph	feedstock/ninja-hep-ph	recipe/ninja-hep-ph				
nnpdf	feedstock/nnpdf	recipe/nnpdf				
oneloop	feedstock/oneloop	recipe/oneloop				
oneloop		recipe/oneloop				
pineappl	feedstock/pineappl	recipe/pineappl				
pytha	feedstock/pytha	recipe/pytha				
pysiha	feedstock/pysiha	recipe/pysiha				
qcdloop	feedstock/qcdloop	recipe/qcdloop				

The 'Simulation' section is also visible, showing packages like bxdecay0, delphes, fastjet, fastjet-contrib, and fastjet-contrib. The 'Grid' section lists packages like aptainer, ca-policy-lcg, davix, dirac-grid, gfal2, python-gfal2, rucio-mcp, voms-lsc, and xrootd.

[hep-packaging-coordination](https://hep-packaging-coordination.github.io/)



Development



What is development?

- Someone writing
 - libraries for general use (e.g. ROOT, FastJet, Rivet, Pythia, ...).
 - software for a specific experiment (e.g. LHCb, ATLAS, CMS, Belle II, ...).
 - a niche analysis tool for use inside an experiment



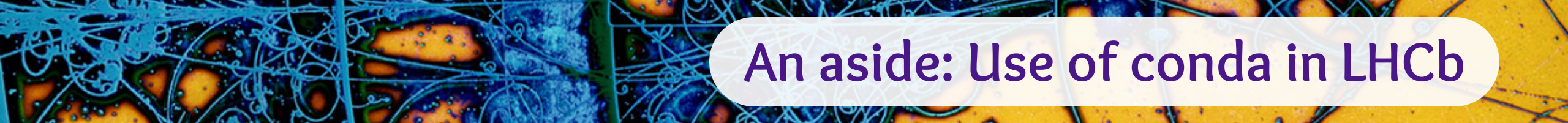
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 - a niche analysis tool for use inside an experiment
- Someone doing
 - analysis for physics
 - quick one-off detector study

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 - Point to local clones of your dependencies
- Kind of like Python's editable installs except for any language



An aside: Use of conda in LHCb

- Most LHCb software is on conda-forge (even extremely LHCb-specific software)
- Already used for:
 - User login environment on CVMFS/lxplus
 - All grid middleware
 - Hosting web services
 - User analysis environments on CVMFS (lb-conda)
 - Local user environments
- Notable exception is the "physics stack"
 - Have been experimenting with this more recently
 - Extremely promising, main questions are around the nicest way to integrate it

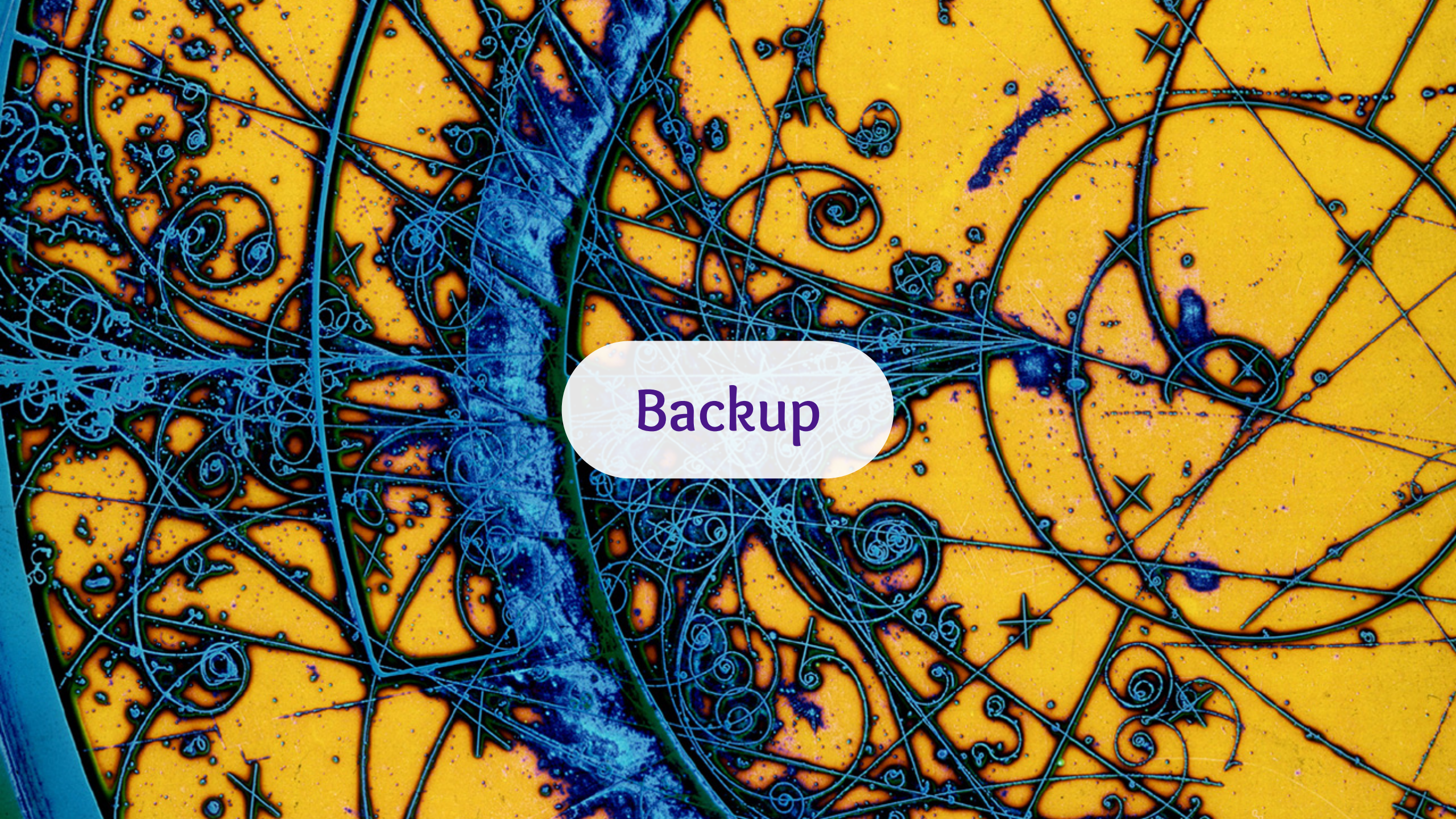
- conda-forge provides the recipes/binaries
- The conda community provides the tooling
- HEP Packaging Coordination coordinates efforts within HEP

Links:

- HEP Packaging Coordination — <https://hep-packaging-coordination.github.io/github/>
- Matthew's CHEP 2026 talk — <https://matthewfeickert-talks.github.io/talk-chep-2026/>
- "Conda, Pixi and RattlerFS" — <https://talks.chrisburr.me/2026-06-22-pixi-and-rattlerfs/>



Questions?



Backup

What does typical end-user use look like?

```
$ pixi init example && cd example # create workspace
$ pixi add contur # declaratively add tools
✓ Added contur >=3.1.4,<4
$ pixi run contur ... # execute commands or tasks
$ pixi list rivet # inspect environments
```

Name	Version	Build	Size	Kind	Source
rivet	4.1.3	py314h9404863_2	53.69 MiB	conda	https://conda.anaconda

```
$ pixi shell # drop into interactive subshells

(debug) $ command -v contur
/tmp/example/.pixi/envs/default/bin/contur
```

pixi global install

- Install CLI tools that are always available, each in its **own isolated environment**
- Only the exposed executables land on your PATH — no dependency clashes between tools
 - Could have a pyroot640 and pyroot642 installed at the same time, for example

```
~> pixi global install root --with ipython
~> pixi global expose add --environment root pyroot=ipython
✓ Exposed executable pyroot from environment root.

~> which pyroot
/Users/cburr/.pixi/bin/pyroot           # just a wrapper on PATH

~> pyroot
IPython 9.14.1 -- An enhanced Interactive Python.
```

- Run a command in a **throwaway environment** — nothing is installed
- Pixi fetches what's needed, runs it, then it's gone — ideal for one-offs & CI

```
~> pixi exec root -l -b -q -e '1+1'  
(int) 2
```



Multiple channels

- There is precedent for using multiple conda channels together
- Most significant ones use conda-forge as a base:
 - [Bioconda](#) for bioinformatics software
 - [RoboStack](#) for robotics software
 - [Emscripten Forge](#) for the emscripten-wasm32 platform
- For HEP I think it's better to work within conda-forge
 - Get a lot of benefits from the shared infrastructure
 - Avoids overlap with multi-purpose software (ROOT, Geant4, ...)
- I think it can make sense to have smaller channels for niche software
 - LHCb will likely end up with a channel for their physics stack, for example



RattlerFS: What is the problem?

- The main trade off for conda is disk space and IO usage
 - No different from containers in that regard
 - Pixi tries to be smart with reflinking/hardlinks, laziness, etc.
- It's fine on a local NVMe drive (e.g. your laptop)
 - On hard drives it's not ideal but manageable
 - On AFS it's almost unusable
 - Having it in every grid job would be a disaster (there is a reason we have CVMFS!)

RattlerFS Primer 1: What does "install" mean?

- What does "installing a conda package" mean?
 1. Download and extract the package to the local cache
 2. "Copy" the files to the install location
 3. Apply any necessary fixes (e.g. shebangs, hard coded paths, python stuff, ...)
- RattlerFS is a virtual filesystem which implements step 2+3
 - Proxies data from the local cache to the install location on demand
 - No need to copy files, no disk usage or IO overhead!

RattlerFS Primer 2: One step further with CVMFS

“Download and extract the package to the local cache”

- We already have a tool for that: CVMFS!
- Cache all conda-forge packages on CVMFS
 - RattlerFS can proxy them to the install location on demand:

```
$ ls /cvmfs/conda-cache.cern.ch/prototype-v2/*  
linux-64/  noarch/   osx-arm64/
```



RattlerFS: Status

- This works on Linux/macOS/Windows using FUSE/NFS/ProjFS*
- The upstream package that provides conda tooling (`rattler`) is interested
- "Just" need to open the (large) pull request...

** Not all backends work on all operating systems*