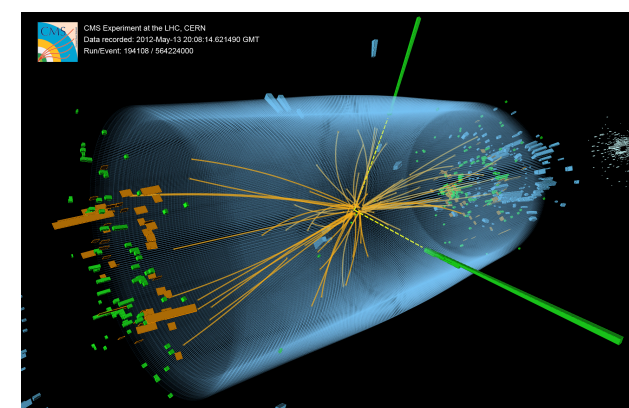




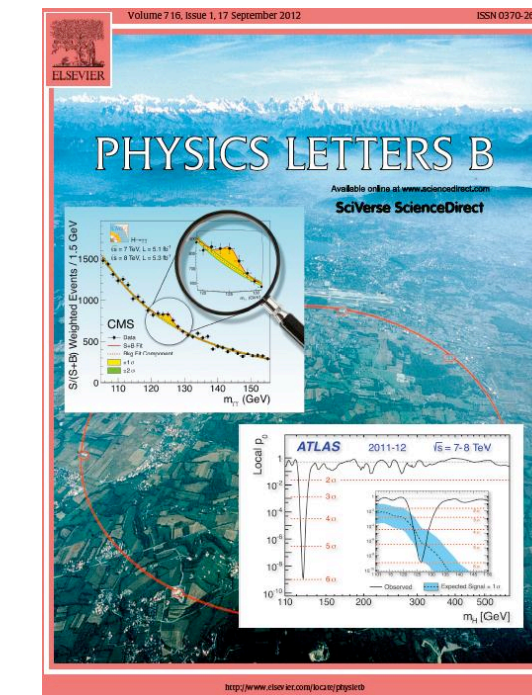
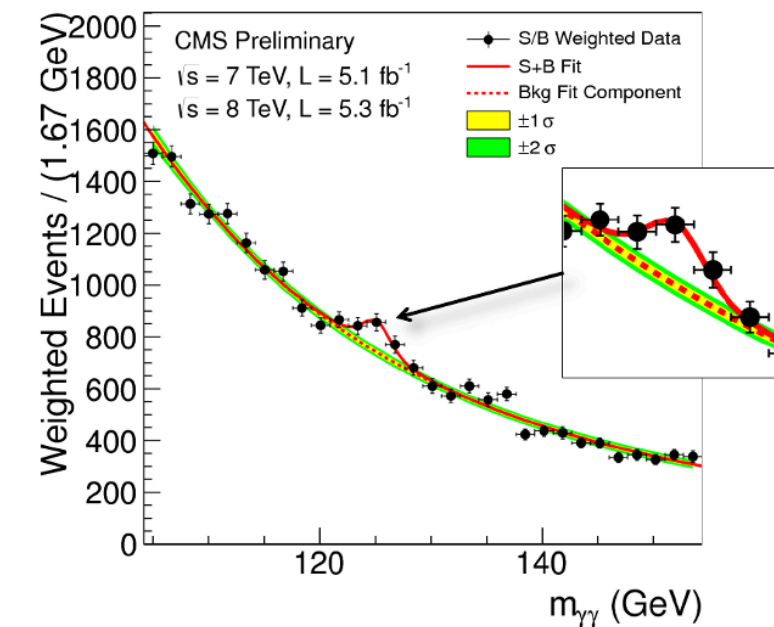
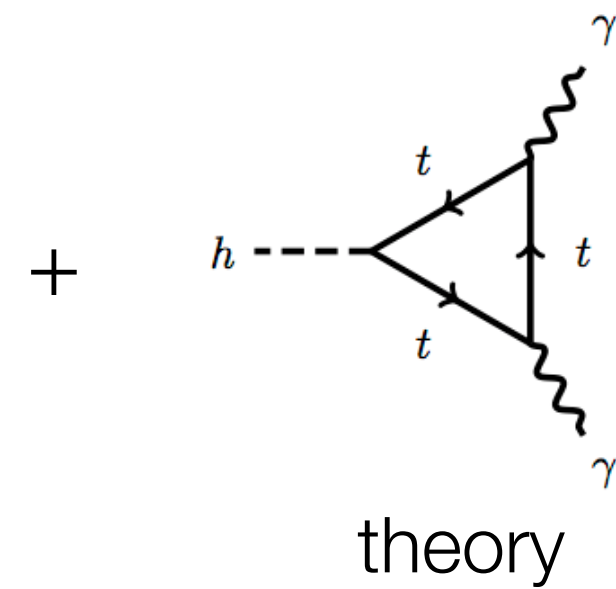
# Design and Execution of make-like distributed Analyses based on Spotify's Pipelining Package Luigi



> make



experiment

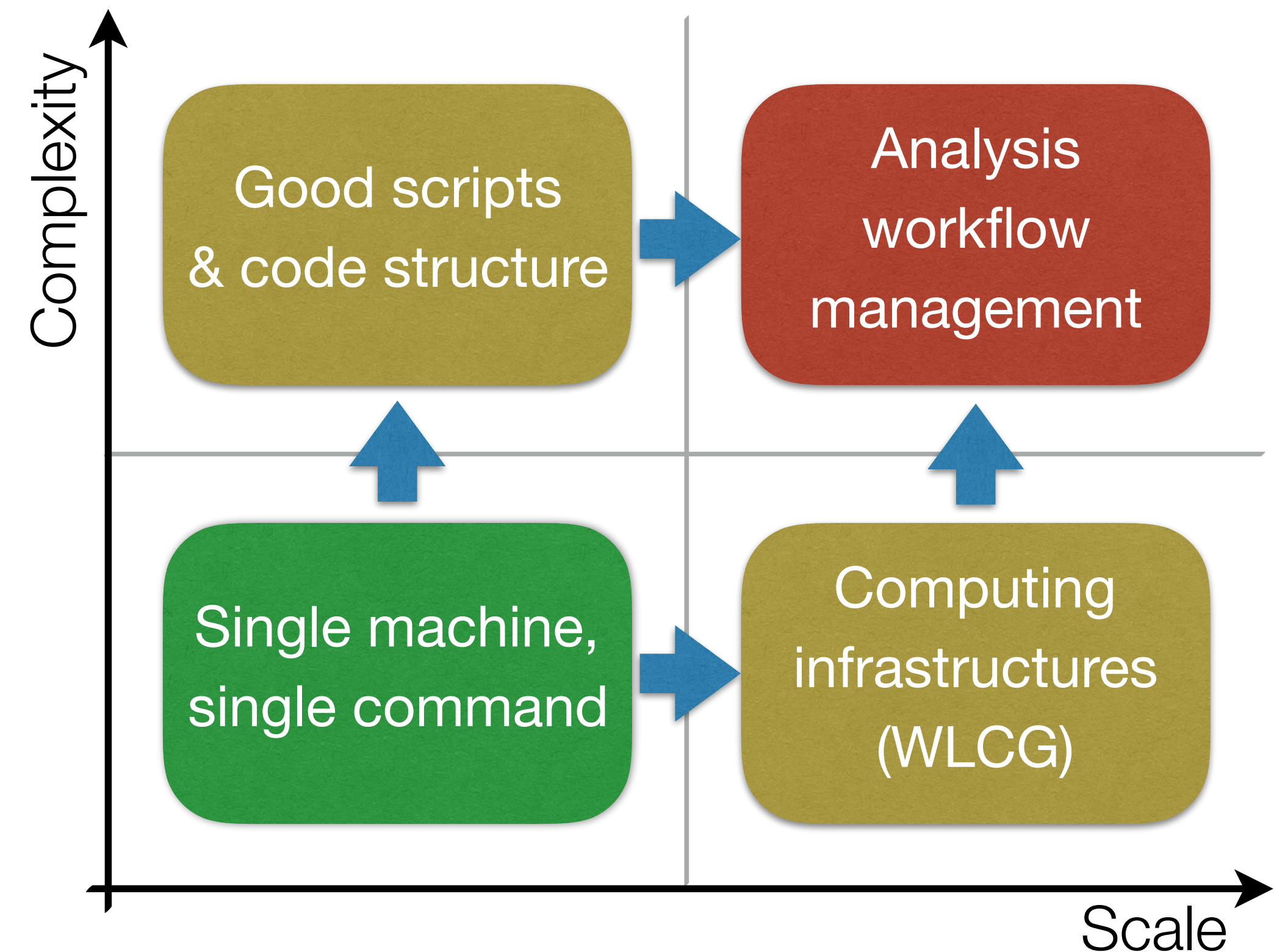


Marcel Rieger\*,  
Martin Erdmann, Benjamin Fischer, Robert Fischer

12/10/2016

# Landscape of Analyses in HEP

- Scale: measure of resource consumption and amount of data
- Complexity: measure of granularity and inhomogeneity of workloads
- Future analyses likely to be large *and* complex, bottlenecks:
  - Entangled and undocumented structure & requirements between workloads, only exists in the “physicist’s head”
  - Bookkeeping of code, data, versions, ...
  - Manual execution and steering of jobs
  - Error-prone & time-consuming



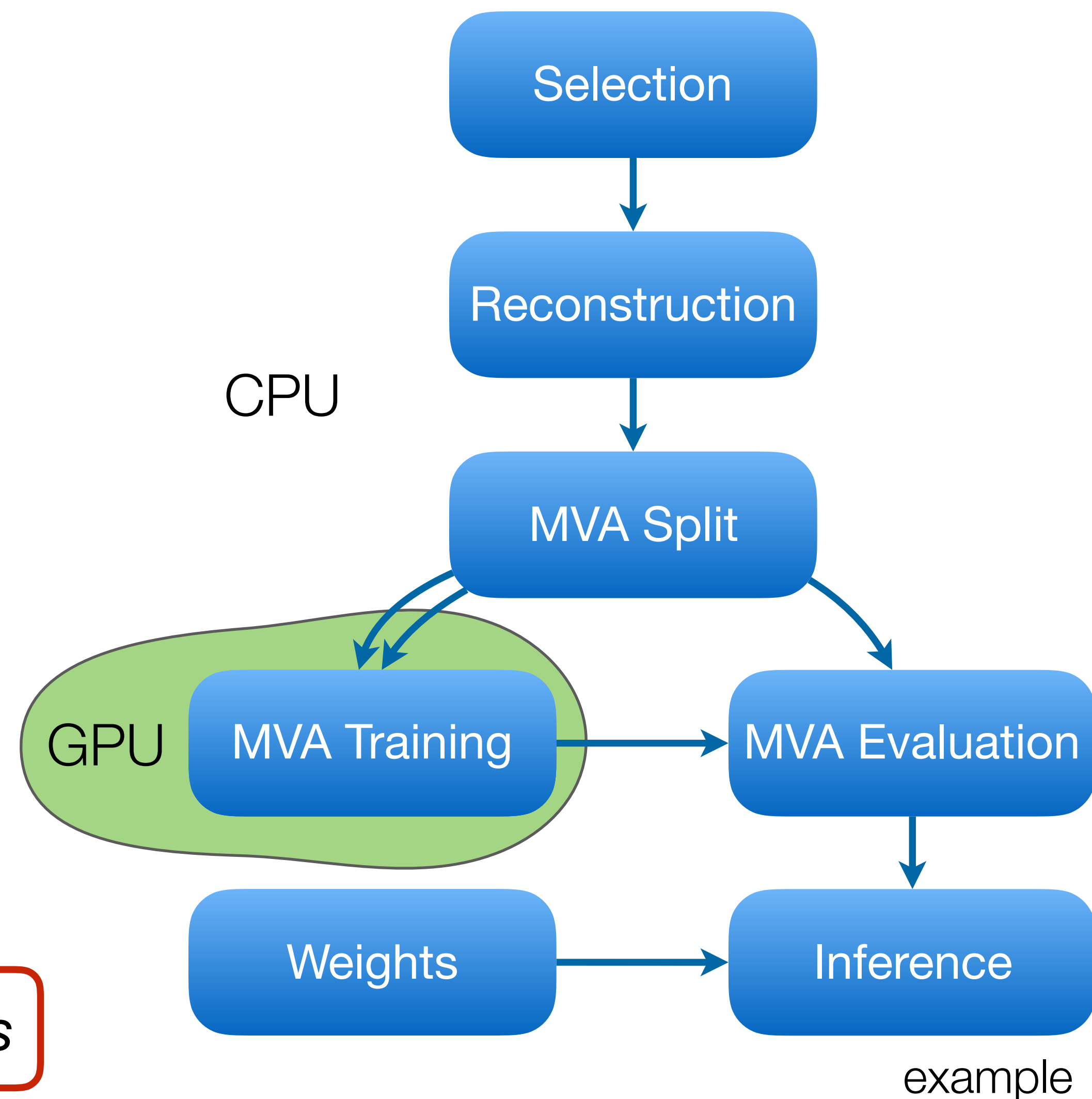
→ *Analysis workflow management essential for future measurements!*



# Abstraction: HEP Analysis

- Workflow, decomposable into particular workloads
- Workloads related to each other by common interface  
→ In/outputs define directed data flow
- Alter default behavior via parameters
- Computing resources
  - Run location (CPU, GPU, grid, ...)
  - Storage location (local, dCache, ...)
- Software environment
- Collaborative development and processing
- Reproducible intermediate and final results

→ *Large overlap with features of workflow systems*



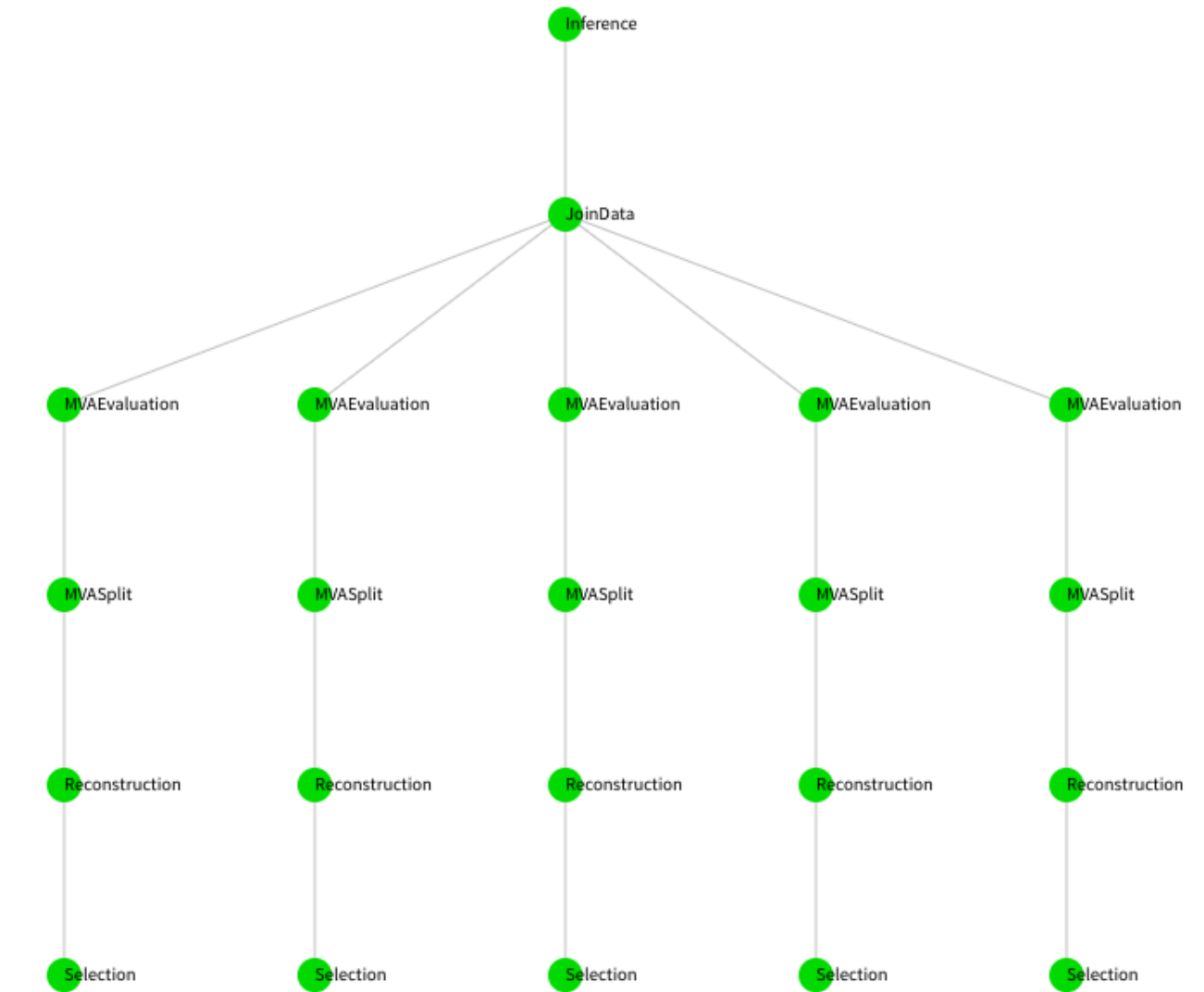
# Comparison of Workflow Management Systems (WMS)

	Existing WMS e.g. MC Management	Generic Analysis WMS
<b>Development Process</b>	final objective known in advance	iterative, final composition a priori unknown
<b>Workflow Structure</b>	chain structure, mostly one-dimensional	tree structure, arbitrarily branched
<b>Evolution</b>	static over time, recurrent execution	dynamic, fast R&D cycles
<b>Infrastructure</b>	specially tailored, e.g. storage systems, DBs	incorporate existing, quickly adapt to changes
<b>Applicability</b>	tuned to particular use case	flexible, able to model every possible workflow

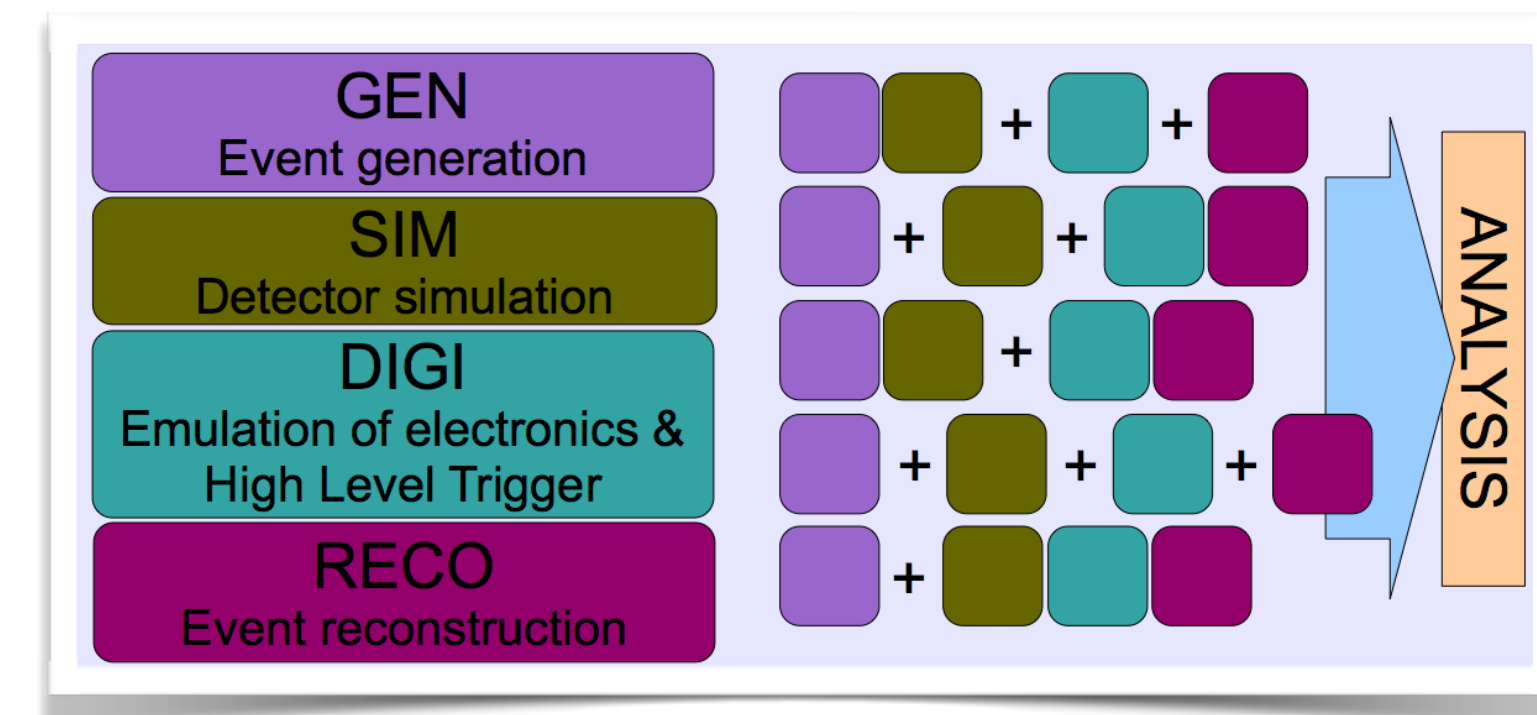
- Existing WMS highly specialized for designated use case
- Requirements for HEP analyses mostly orthogonal

→ *Toolbox for flexible analysis conception*

Typical analysis *tree*:



Typical MC *chain*:





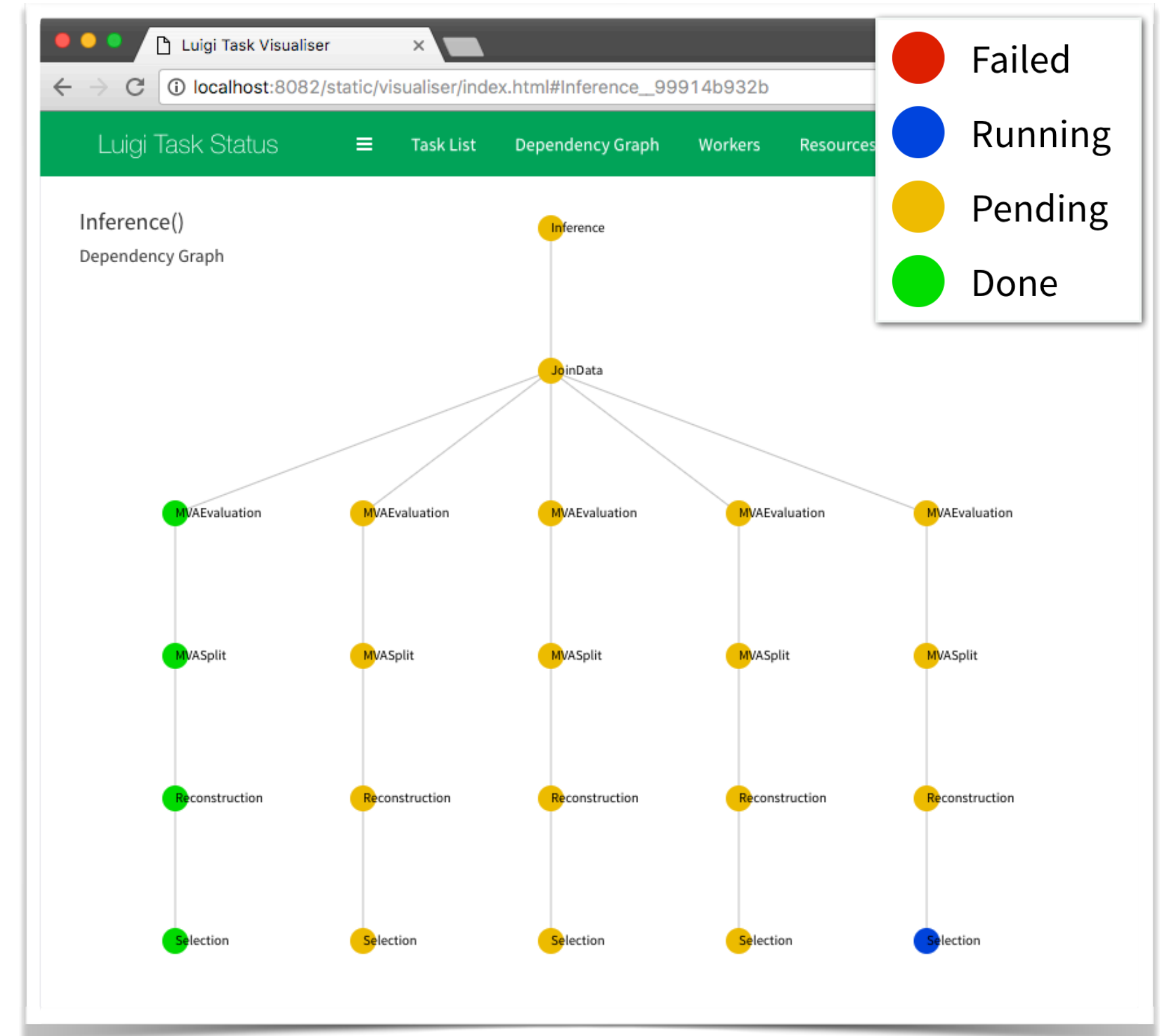
- Python package for building complex pipelines
- Development started at Spotify, now open-source and community driven

## Building blocks

1. Workloads defined as *Task* classes
2. Tasks *require* other tasks & output *Targets*
3. *Parameters* customize and control task behavior

- Task execution → builds up dependency tree, only computes what is necessary
- Web interface, error handling, command line tools, collaborative features, ...

→ *Suitable tool to manage complexity*



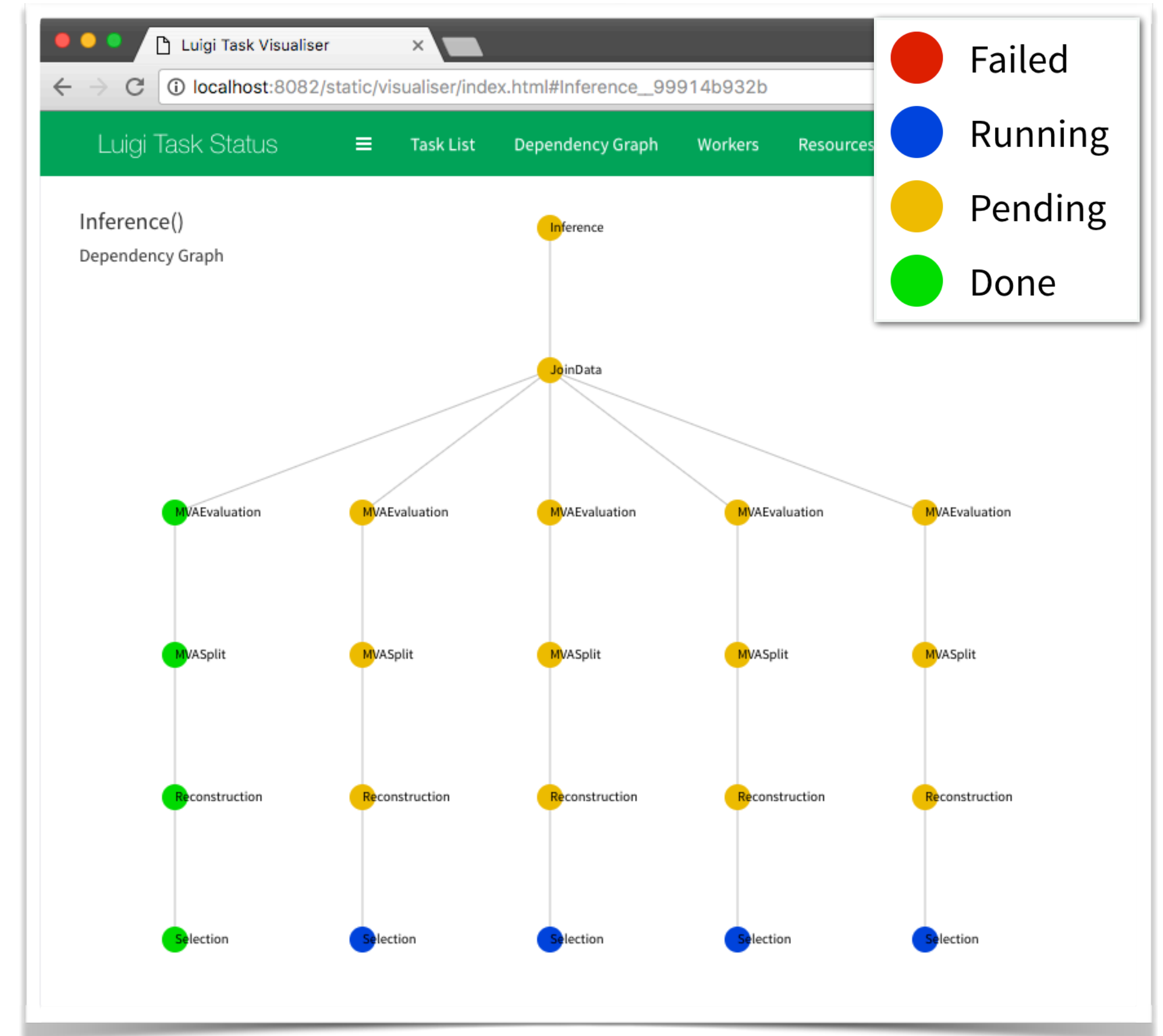
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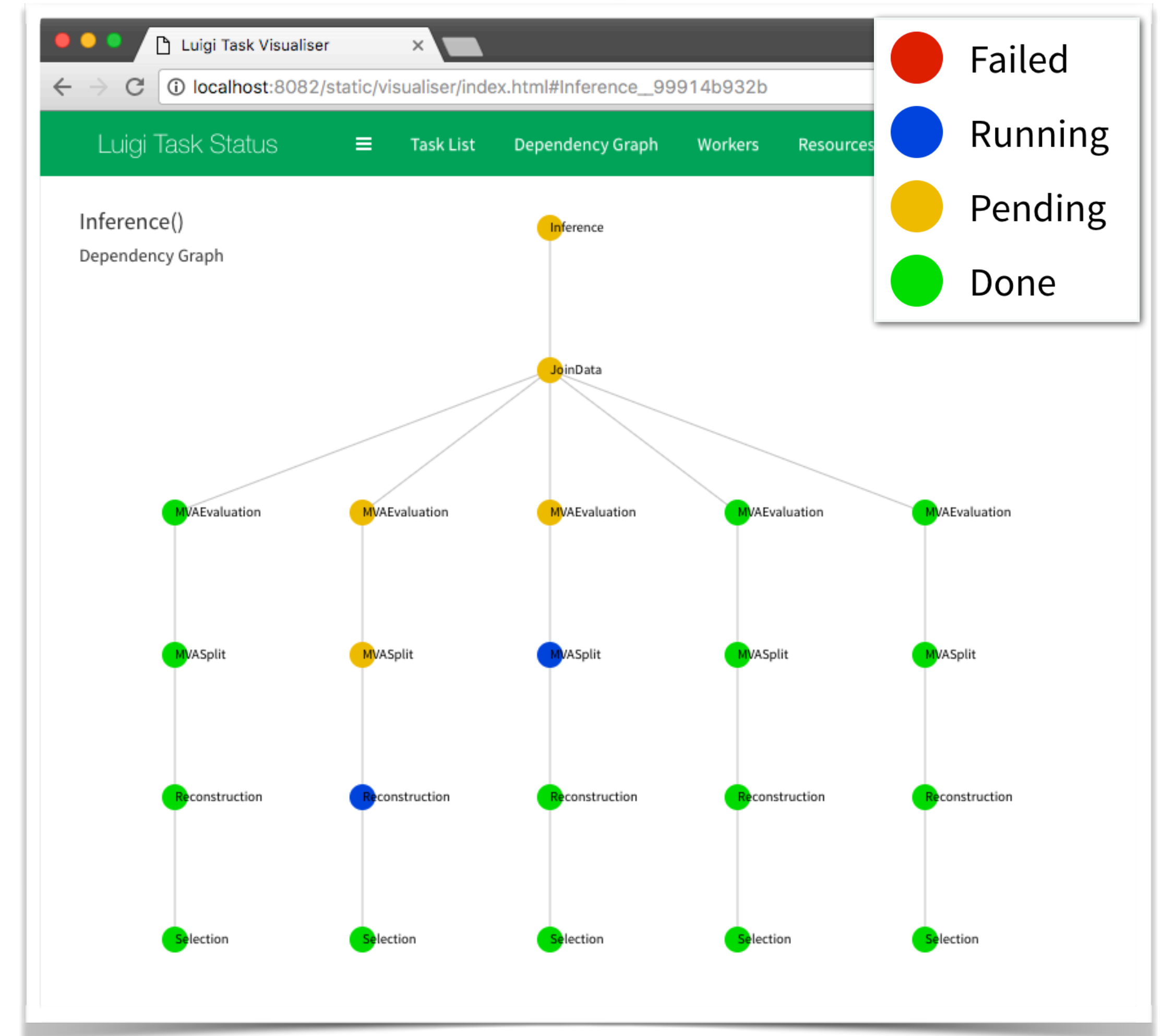
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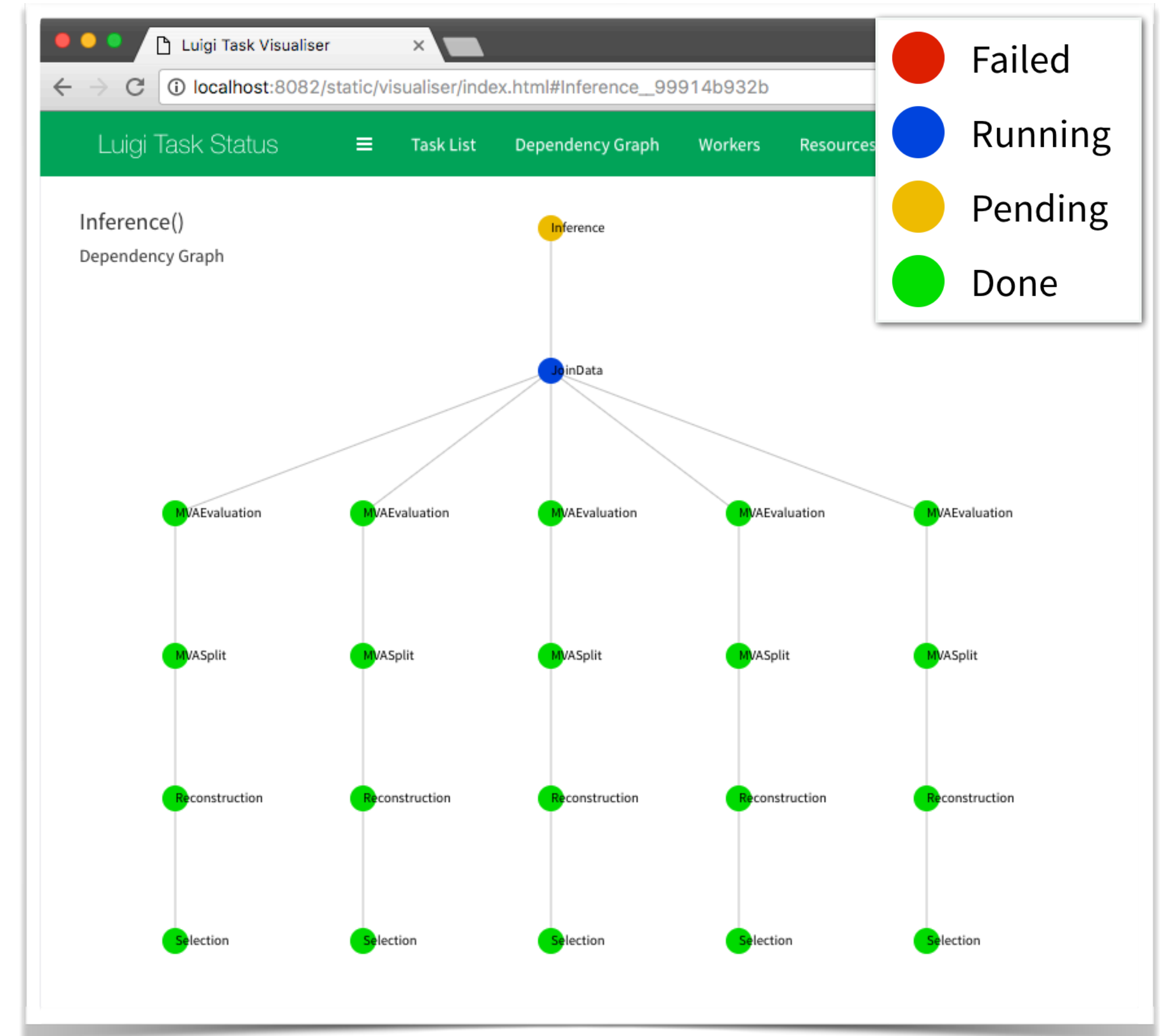
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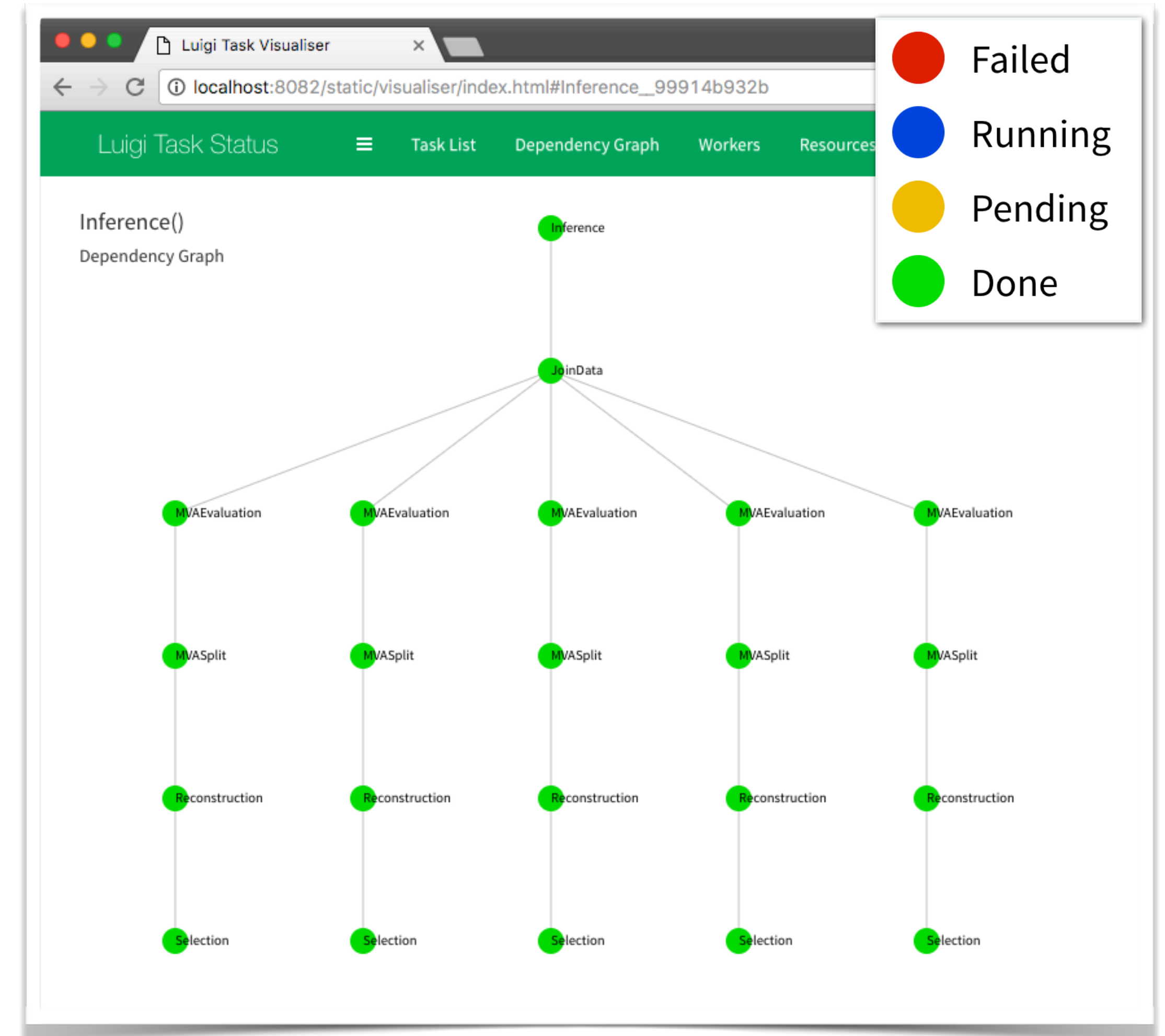
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# Adding Scalability: Luigi and the WLCG

- Example for implementation of abstract run & storage locations

## 1. Submit tasks as jobs to *computing elements*

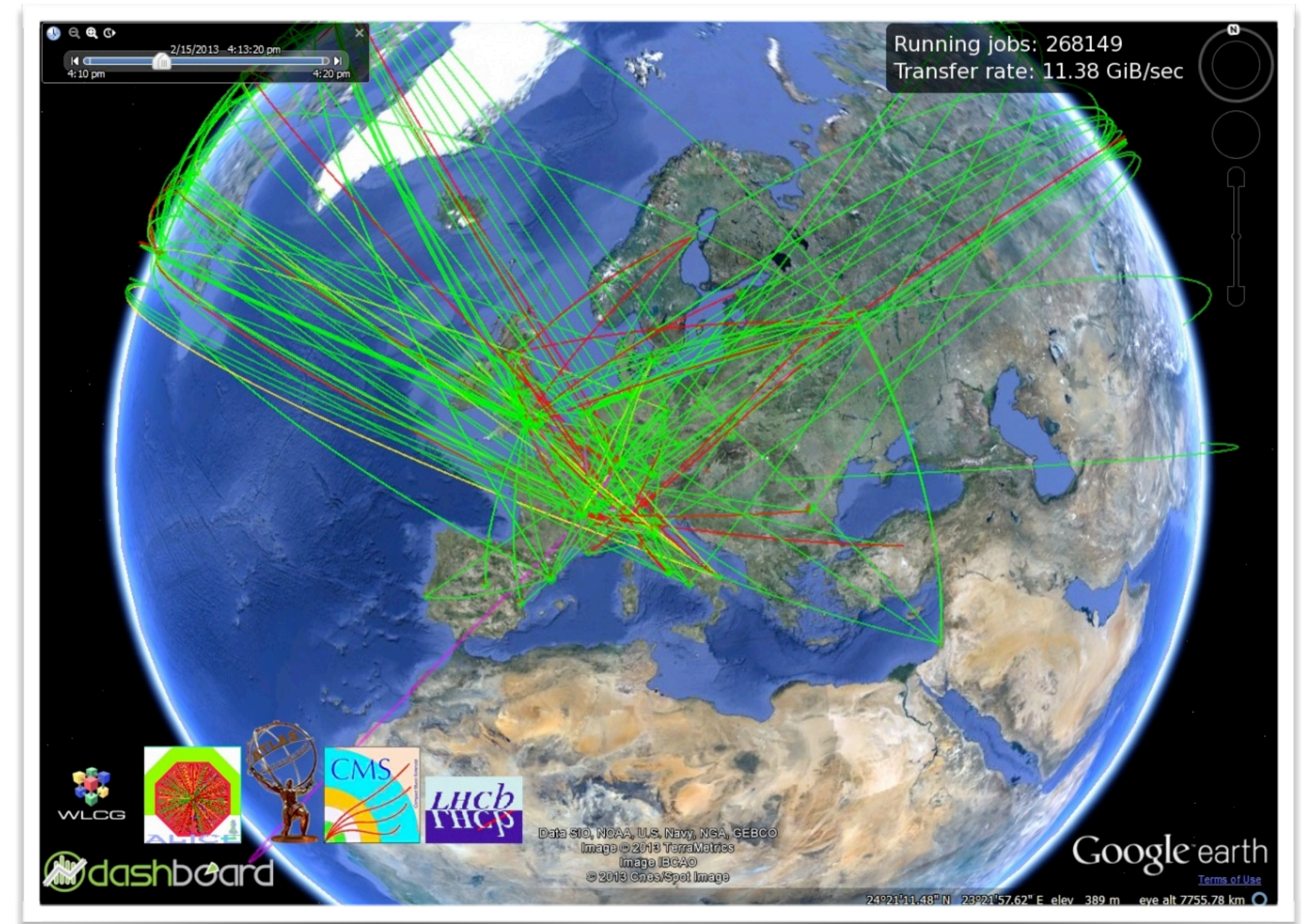
- Simple usage, transparent Luigi integration
- Actual run location (local, CE) not hard-coded, decision made at execution time
- Mandatory features like pilot jobs, automatic resubmission, or batch submission

## 2. Store targets on *storage elements* (e.g. dCache)

- Built on top of GFAL2 Python bindings, transparent Luigi integration
- Mandatory features like automatic retries, local caching, or batch transfers

Luigi

GFAL2



→ *WLCG implementations provide scalability in the HEP context*



# Adding Scalability: Luigi and the WLCG

- Example for implementation of abstract run & storage locations



**GFAL2**

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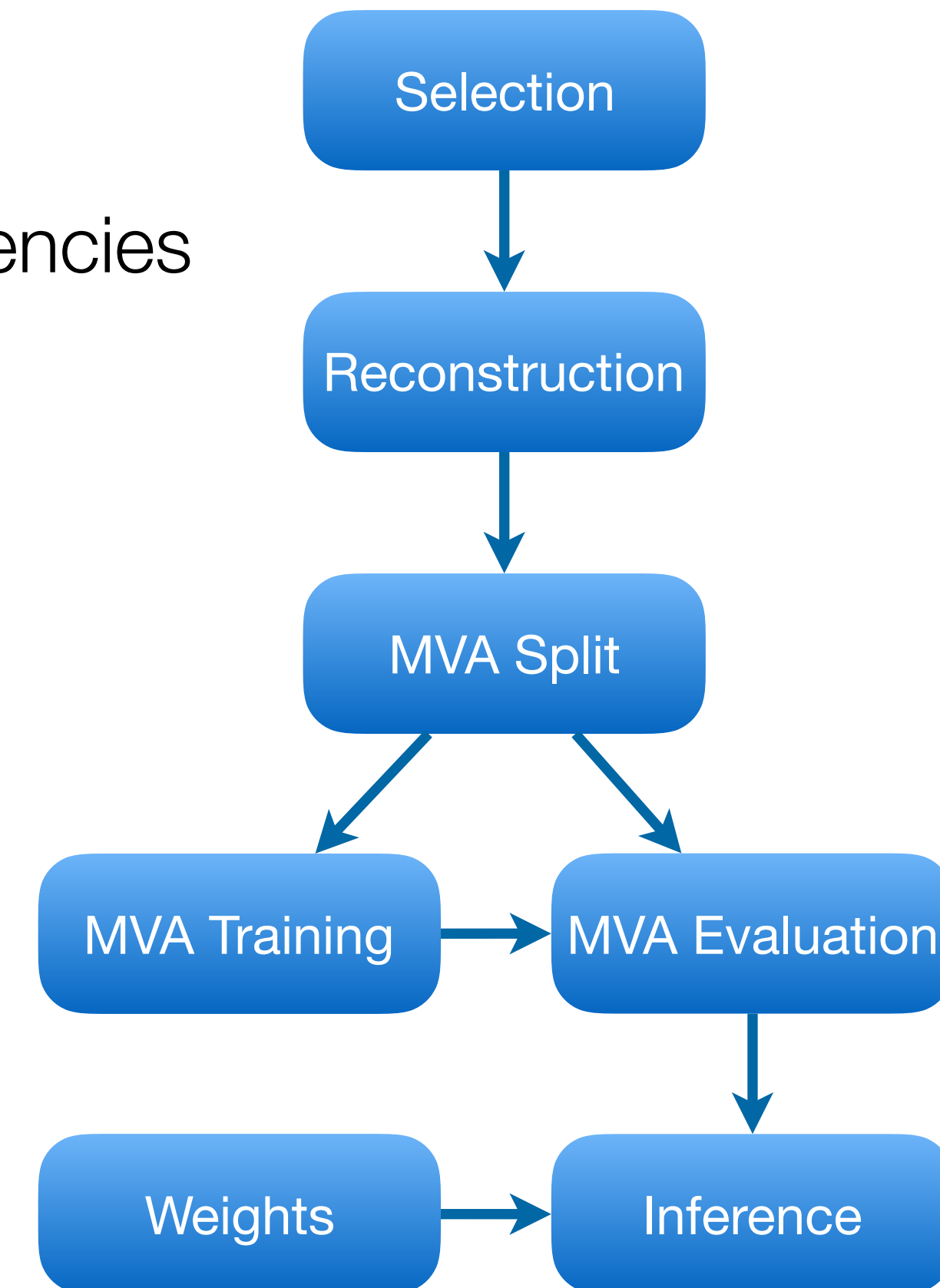
```
> pyl Reconstruction --v test1 --local  
> pyl Reconstruction --v prod1 --ce RWTH
```

```
target = DCacheTarget("/path/to/file.txt")  
  
with target.open("w") as f:  
    f.write("some result")
```

→ *WLCG implementations provide scalability in the HEP context*

# Direct Consequences and Benefits

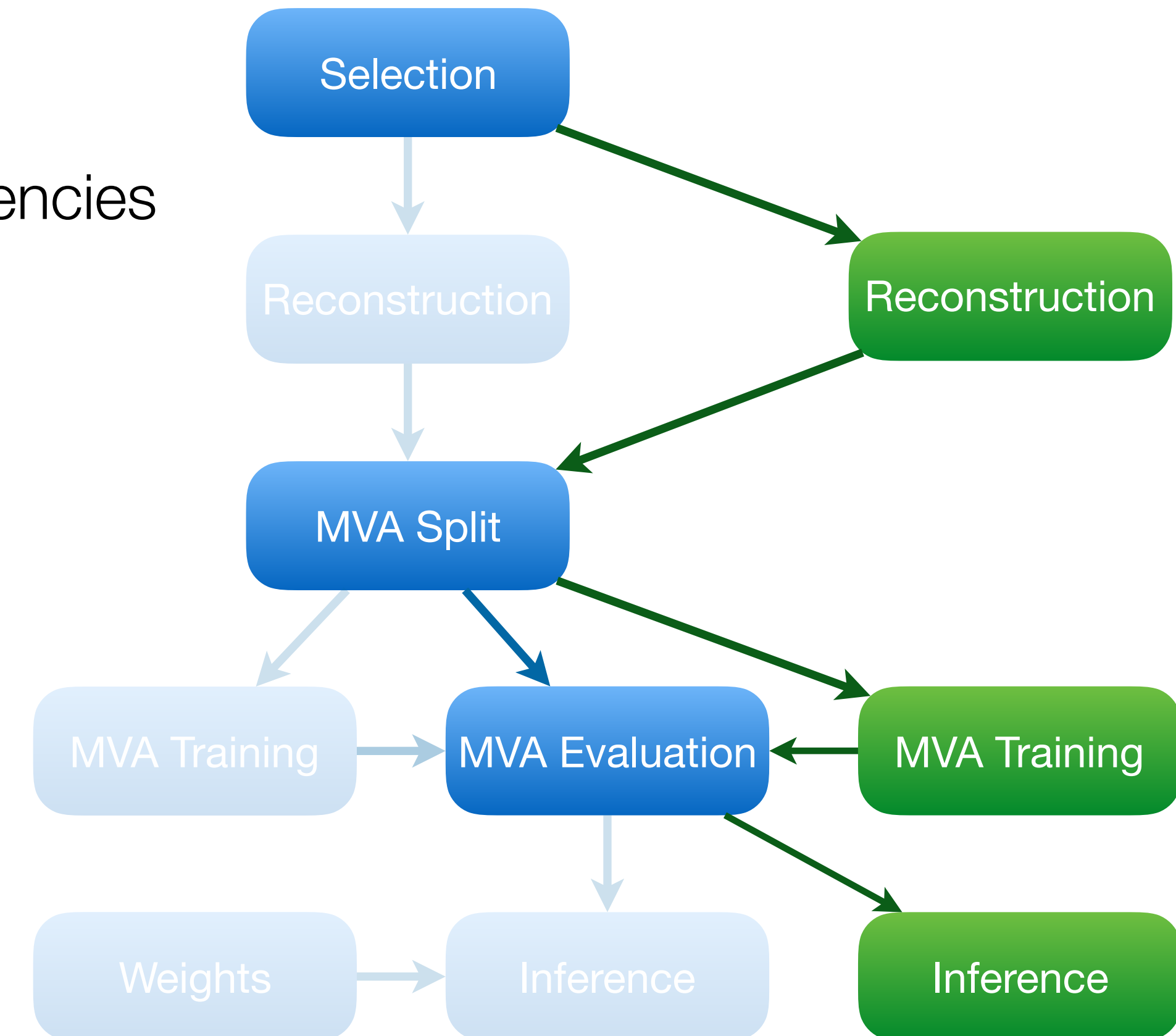
- Toolbox providing building blocks for analyses, *not a framework*
  - Permissive, non-restrictive *design pattern*  
(e.g. no constraint on language or data structure)
- *All* information transparently encoded in tasks, targets & dependencies
  - Results *reproducible* by developer, groups, reviewers, ...
  - Documentary benefits, enables *analysis preservation*
- make-like execution across distributed resources
  - Reduces overhead of manual management
  - Improves cycle times & error-proneness
- Expansion of the concept of *collaboration*
  - Clear structure lowers entry barrier
  - Modularization allows re-use of tasks & intermediate results





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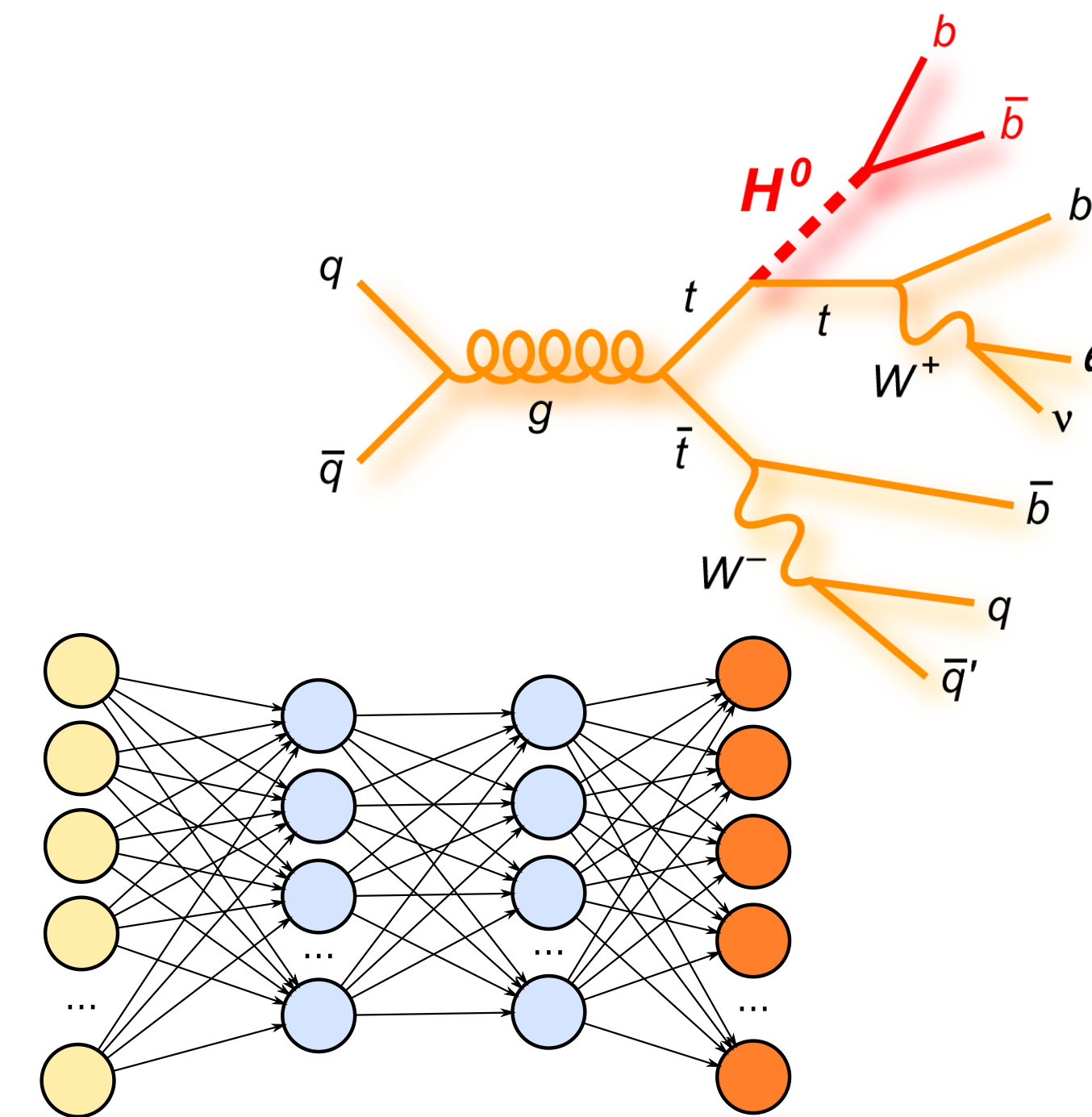
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# Example Application: $t\bar{t}H$ Analysis @ CMS

- $t\bar{t}H$ : measurement of Higgs  $\leftrightarrow$  quark Yukawa coupling
  - large-scale: ~30k input files, ~50 TB of storage, ~1000 unique Tasks
  - complex: irreducible backgrounds, ~40 systematic variations, DeepLearning/BDTs, multiple categorization schemes
- Run locations: 7 CEs, local machines, GPU machines
- Storage locations: 2 SEs (dCache), local disk, Dropbox, CERNBox
- Aware of entire workflow structure at all times, fast evaluation & revision
- Group of 5 people, clear allocation of duties and their interface
- Yet, entire analysis *operable by everyone* at all times
- Setup allows for *execution with a single command*

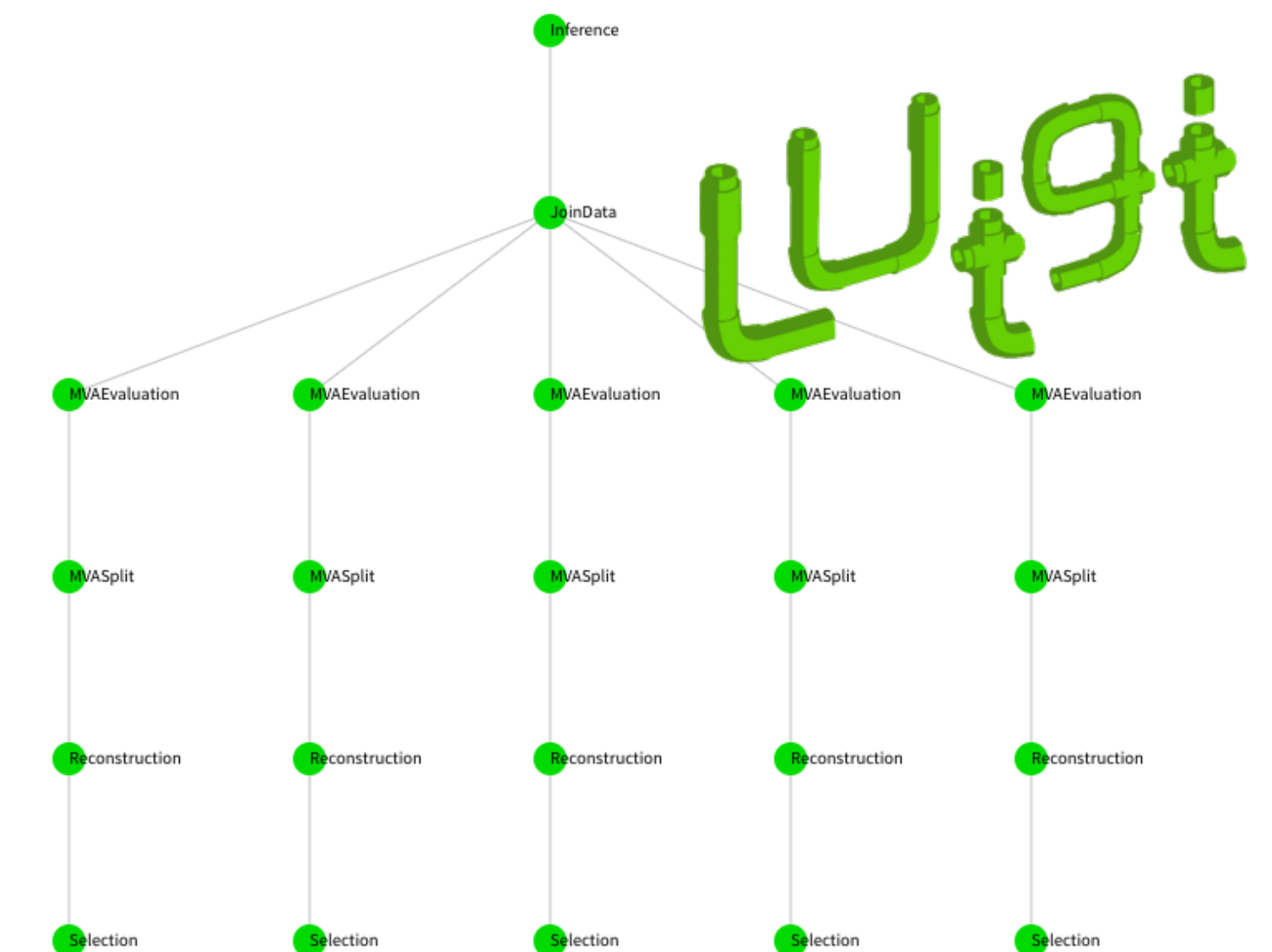
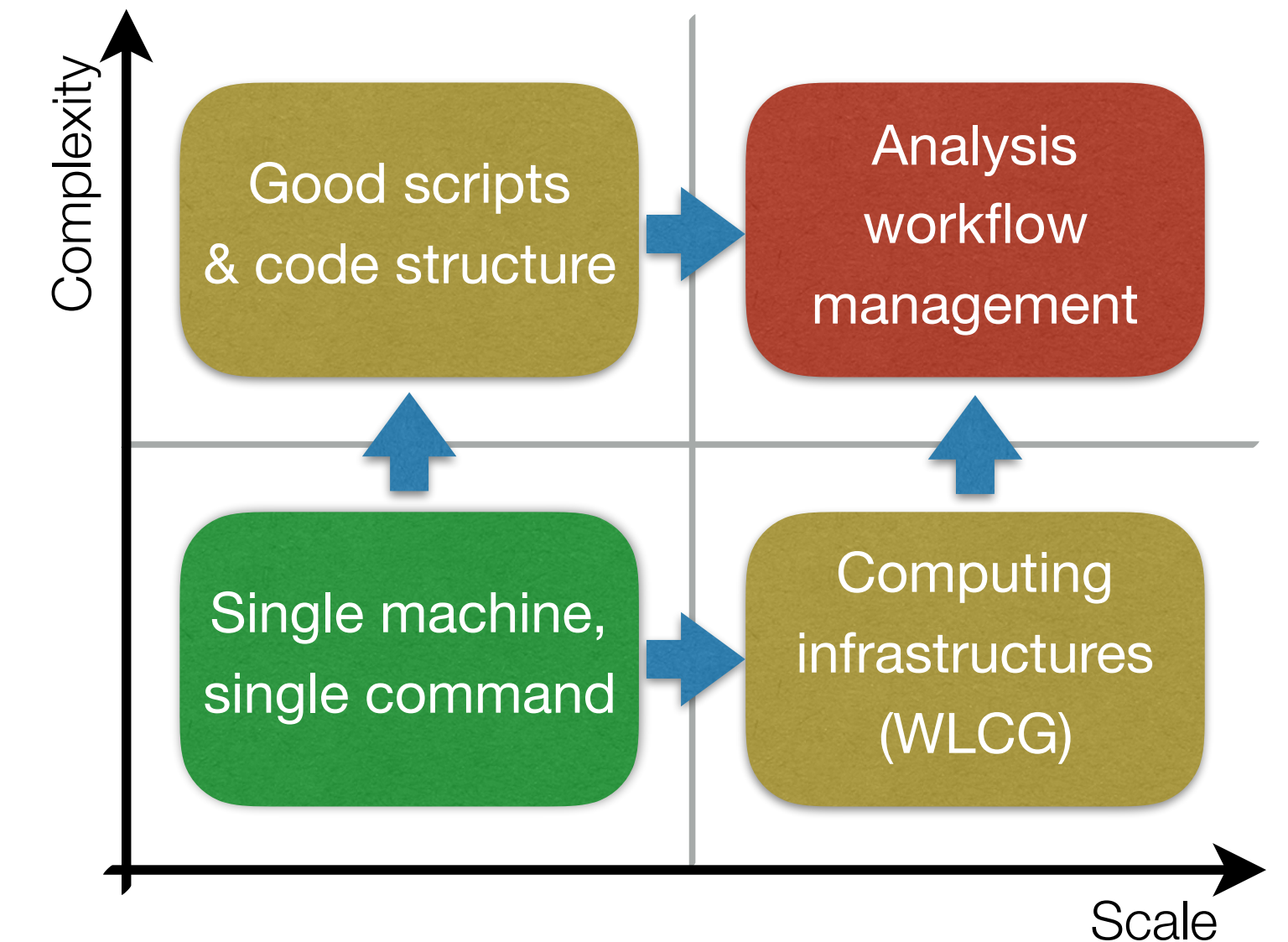
→ *Successful proof of usability & suitability*



workflow image: Kevin Boucher, lansa.com

# Summary

- HEP analyses likely to increase in scale and complexity  
→ Analysis workflow management *essential for success* of future measurements
- Divergent requirements of existing, specialized management systems and those for “end-user” analyses  
→ Need for a *toolbox* providing a *design pattern, not a framework*
- Luigi provides a promising way to model even *complex* workflows
- WLCG extension introduces *scalability* in the HEP context
- Increased *transparency & reproducibility* → *analysis preservation*
- Encourages collaboration beyond code sharing
- Successfully applied in actual *ttH* analysis with CMS





*Backup*

# Luigi - An Introduction

- Package for building complex pipelines
- Development started at Spotify, now open-source and community driven
- Simple core API:

```
# reco.py
```

```
import luigi
```

```
from analyses.ttH.tasks import Selection
```

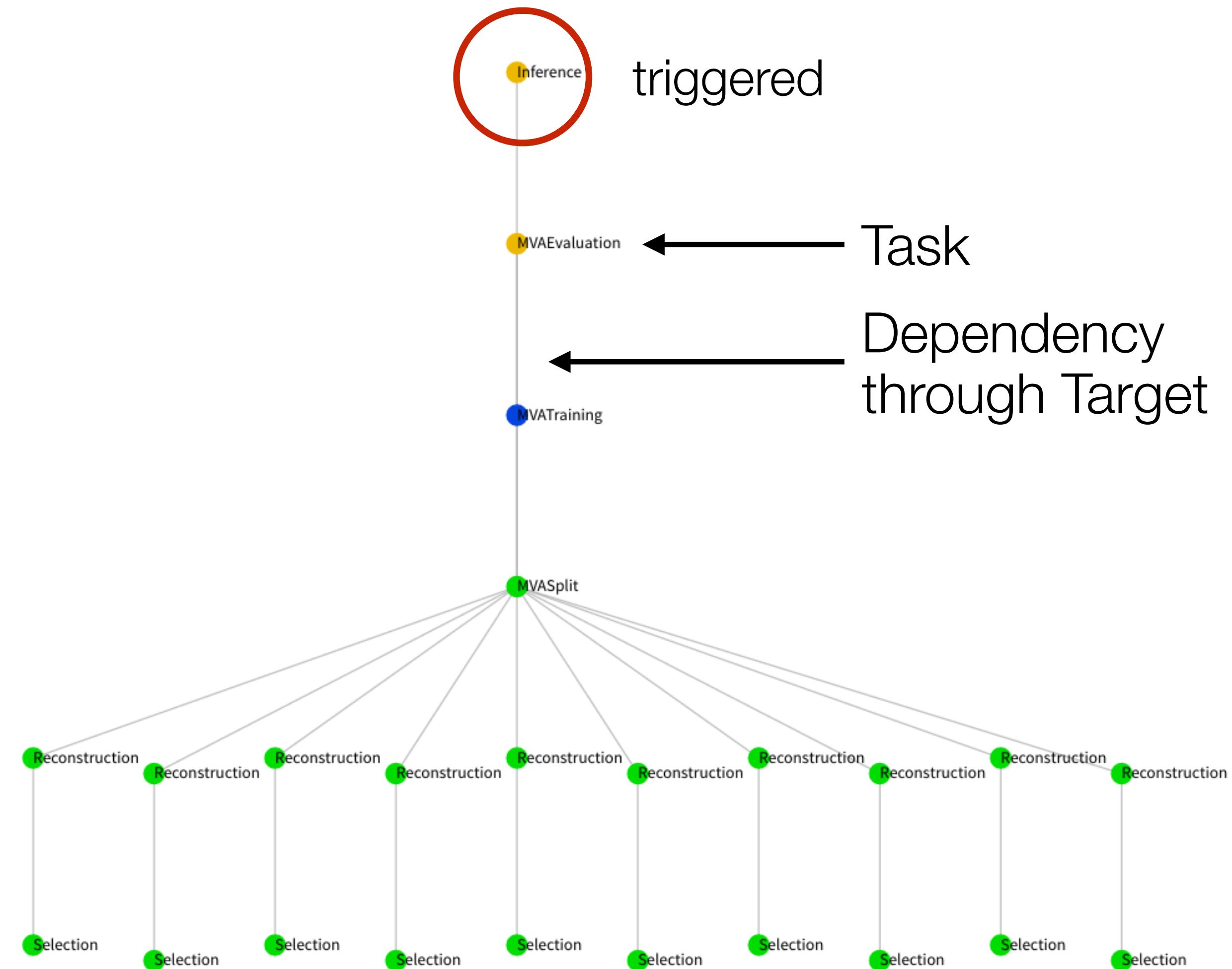


- Workloads are written in Task's  $\longrightarrow$  `class Reconstruction(luigi.Task):`
- Tasks are configured with Parameter's  $\longrightarrow$  `dataset = luigi.Parameter(default="ttH125")`
- Tasks can **require** other tasks,  $\longrightarrow$  `def requires(self):`  
defines (multiple) dependencies `Selection(dataset=self.dataset)`
- Tasks produce Target's, output  $\longrightarrow$  `def output(self):`  
representations with an `exist()` method `return luigi.LocalTarget("reco_%s.root" \`  
`% self.dataset)`
- Actual workload defined in `run()` method,  $\longrightarrow$  `def run(self):`  
completely flexible via python code `# do whatever a reconstruction does`  
`...`

```
> python reco.py Reconstruction --dataset ttH125
```

# Luigi - make-like Execution

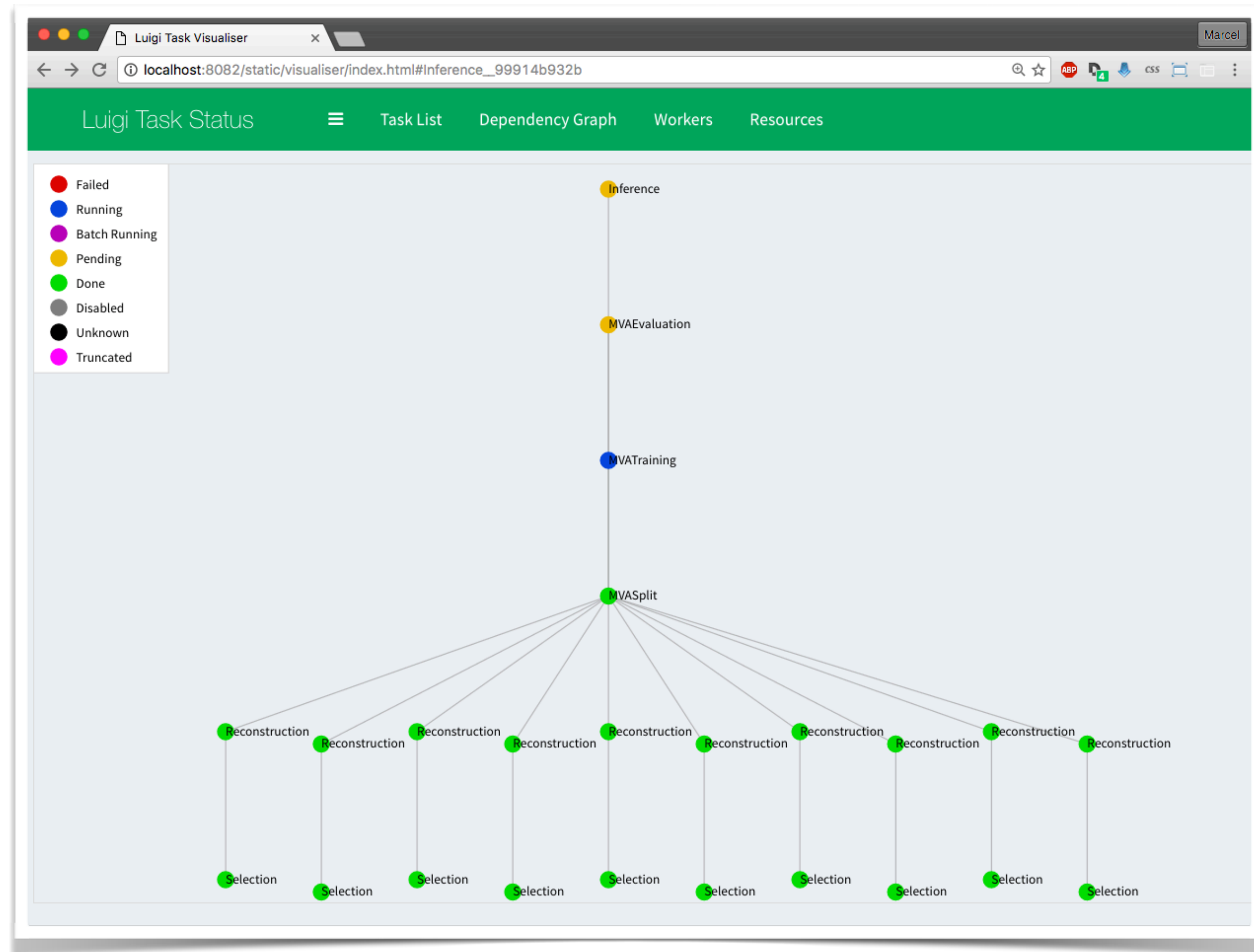
- Luigi's execution system is **make**-like, it only processes what is really necessary
  1. For the triggered task, create the dependency tree
  2. Determine tasks to actually run:
    - 2.1. Walk down the tree
    - 2.2. For each path, stop when all output targets of a task *exist*
  3. Run tasks in  $n$  workers
- Very clear & scalable through simple structure
- Error handling & automatic re-scheduling
- Command line integration & tools
- Central scheduling & visualization





# Luigi - Central Scheduler

- Not a “scheduler” in HEP language, scheduling takes place on worker
- Think of it as a “global task lock”
- Optional, but powerful when working in teams / collaborations
  - Same task should not run twice
  - Saves resources but also ensures target/ data integrity
- Dependency, status & resource visualization
- Control of running workers (add, abort, ...)
- Custom status messages & task history



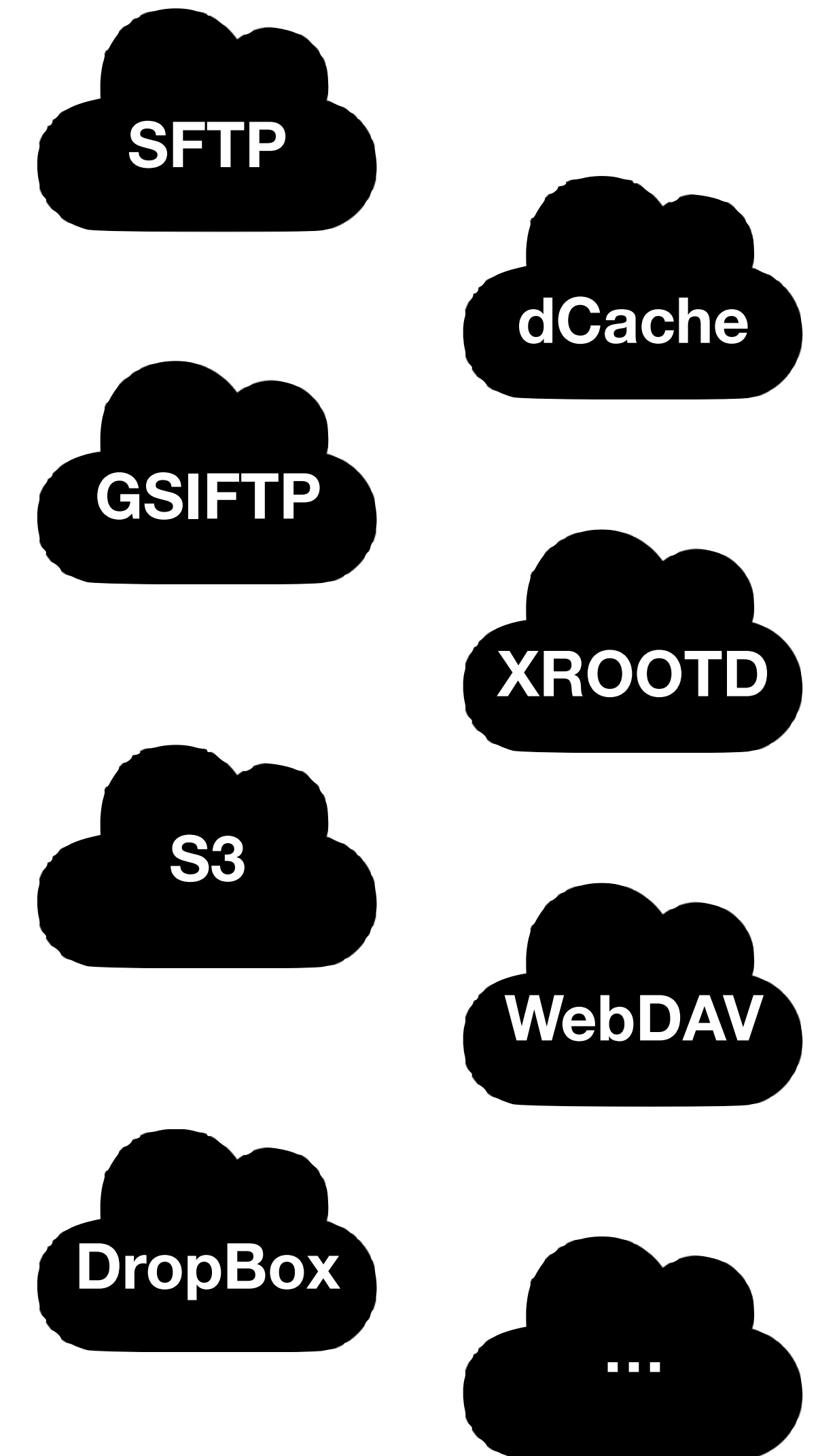
# HEP Layer - GFAL Targets

- When running on the WLCG, use of storage elements is a necessity
- Fortunately, there is GFAL (Grid File Access Library)
  - Developed by Data Management Clients group at CERN
  - Command line tools & python bindings
  - Handles all file transfer protocols of the HEP community

→ *Combine GFAL with Luigi targets*

- Simple API, batch transfers, validation, auto-retry, local caching, ...
- Usage equivalent to local targets

```
def output(self):  
    return DCacheTarget("/path/to/file.txt")  
  
def run(self):  
    self.output().parent.touch()  
  
    with self.output().open("w") as f:  
        f.write("measurement results: ...")
```





# Application: Implementation of Systematics

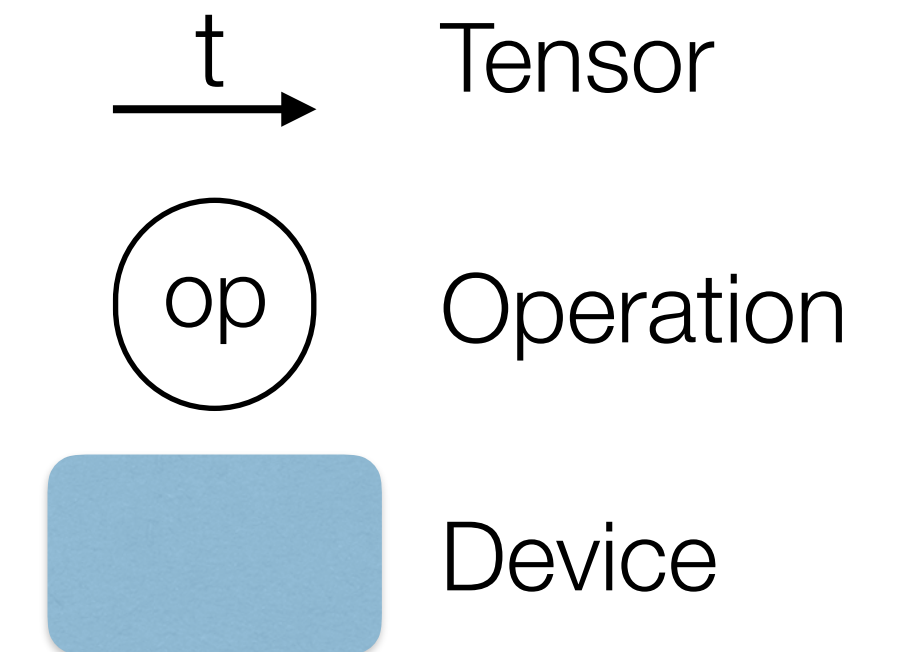
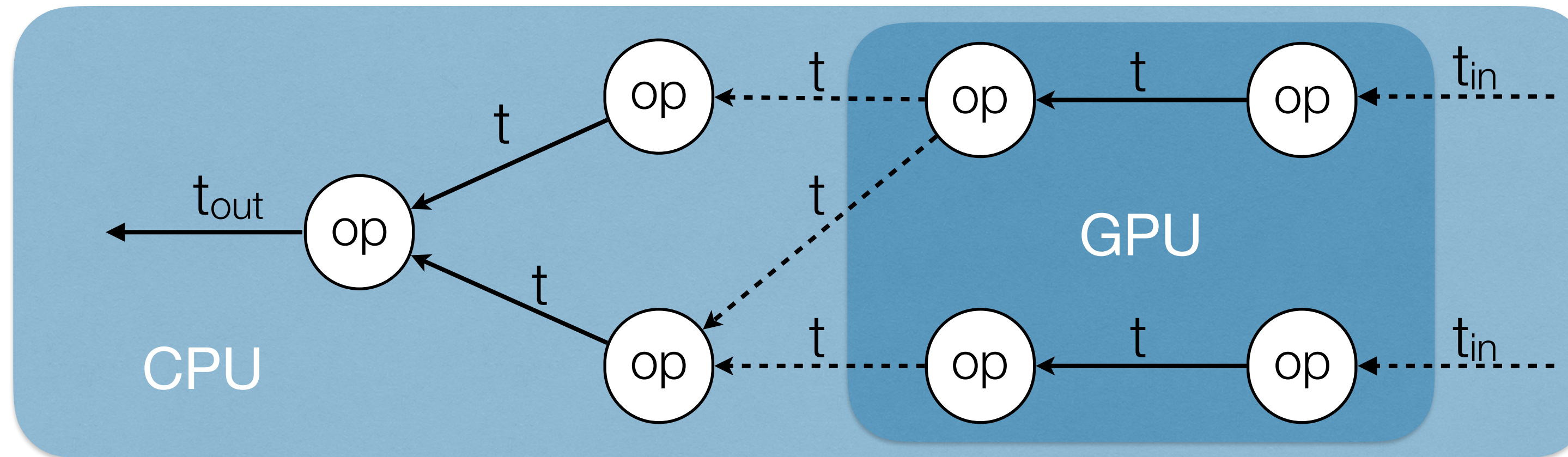
Systematics	“ShiftTask”			“AnalysisTask”
	Selection	Reconstruction	Evaluation	Inference
nominal	✓	✓	✓	✓
JES	☆	✓	✓	✓
JER	☆	✓	✓	✓
PDF	☆	☆	✓	✓
Q <sup>2</sup>	☆	☆	✓	✓

- ✓ implements
- ✓ bubbles up / effective: nominal
- ✓ requires
- ☆ saved
- “implement as late as possible”

————— direction of processing —————→

# tfdeploy (1)

- tensorflow *graphs* consist of *operations* and *tensors*



- Examples:  $t_3 = add(t_1, t_2)$ ,  $t_2 = softmax(t_1)$
- Ops are bound to devices (CPU/GPU), tensors are transferred if needed
- **tfdeploy**:
  - Implements tree structure in pure python
  - Tensors = numpy arrays
  - Ops = vectorized numpy calls, need to implement **all** tensorflow ops
  - Works in all environments, even in C++ with Python C-API, helpful for sharing



# tfdeploy (2)

riga / tfdeploy

Unwatch 10

Star 161

Fork 13

<> Code

Issues 1

Pull requests 0

Projects 0

Wiki


Pulse

Graphs

Settings

Deploy tensorflow graphs for fast evaluation and export to tensorflow-less environments running numpy. — Edit

README.md



build passing

docs latest

pypi package 0.2.4

Deploy [tensorflow](#) graphs for *fast* evaluation and export to *tensorflow-less* environments running [numpy](#).

Evaluation usage

```
import tfdeploy as td
import numpy as np

model = td.Model("/path/to/model.pkl")
inp, outp = model.get("input", "output")

batch = np.random.rand(10000, 784)
result = outp.eval({inp: batch})
```

# Modular Analysis with VISPA & PXL

The screenshot displays the VISPA web interface. The top navigation bar includes a 'Save' button and a 'Main' panel with a table of properties. The central area shows a workflow diagram with nodes: 'Read Data', 'Event Selection', 'Merge Muons', 'Plot Z Mass', and 'Write Rejected Events'. The 'Event Selection' node is highlighted in orange and has a 'passed' output leading to 'Merge Muons' and a 'veto' output leading to 'Write Rejected Events'. The bottom panel shows a console output with event selection criteria and a runtime of 3.26 s.

**Main**

Property	Value
Name	Event Selection
Type	PyModule
Version	0.0.0
Enabled	true
Compilation date	Jun 17 2015
Compilation time	09:33:06

**Options**

Property	Value
filename	/home/public/CERN/examp
script	
parameter	
class	
leptonName	Lepton

```
* particle name: Lepton
* minimal transverse momentum: 30.0 GeV
* maximum pseudo rapidity: 2.1
* minimal number of leptons: 2
* maximal number of leptons: 2
Info in <TCanvas::Print>: png file ./Z_mass.png has been created

runtime: 3.26 s
```