



PRUNE: A Preserving Run Environment for Reproducible Scientific Computing

-Peter Ivie



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- "[An article about computational science in a scientific publication is not the scholarship itself, it is merely advertising of the scholarship. **The actual scholarship is the complete software development environment and the complete set of instructions**]" —Jon Claerbout

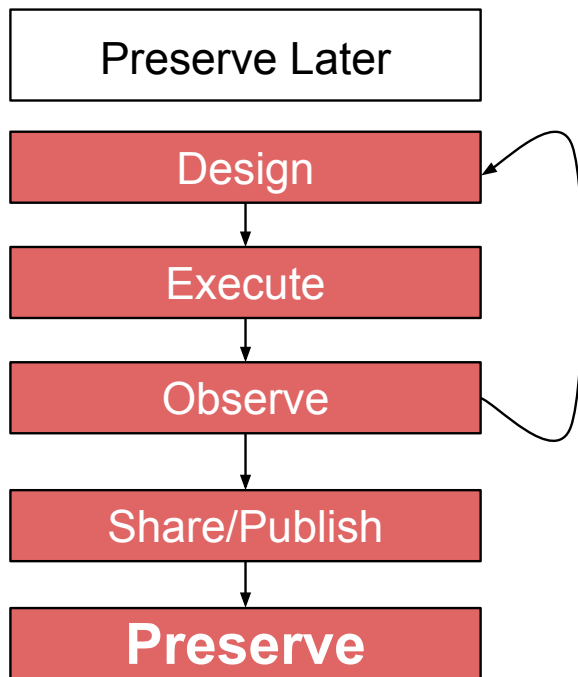
Verify and Extend

- Don't re-invent the wheel
- Stand on the shoulders of giants



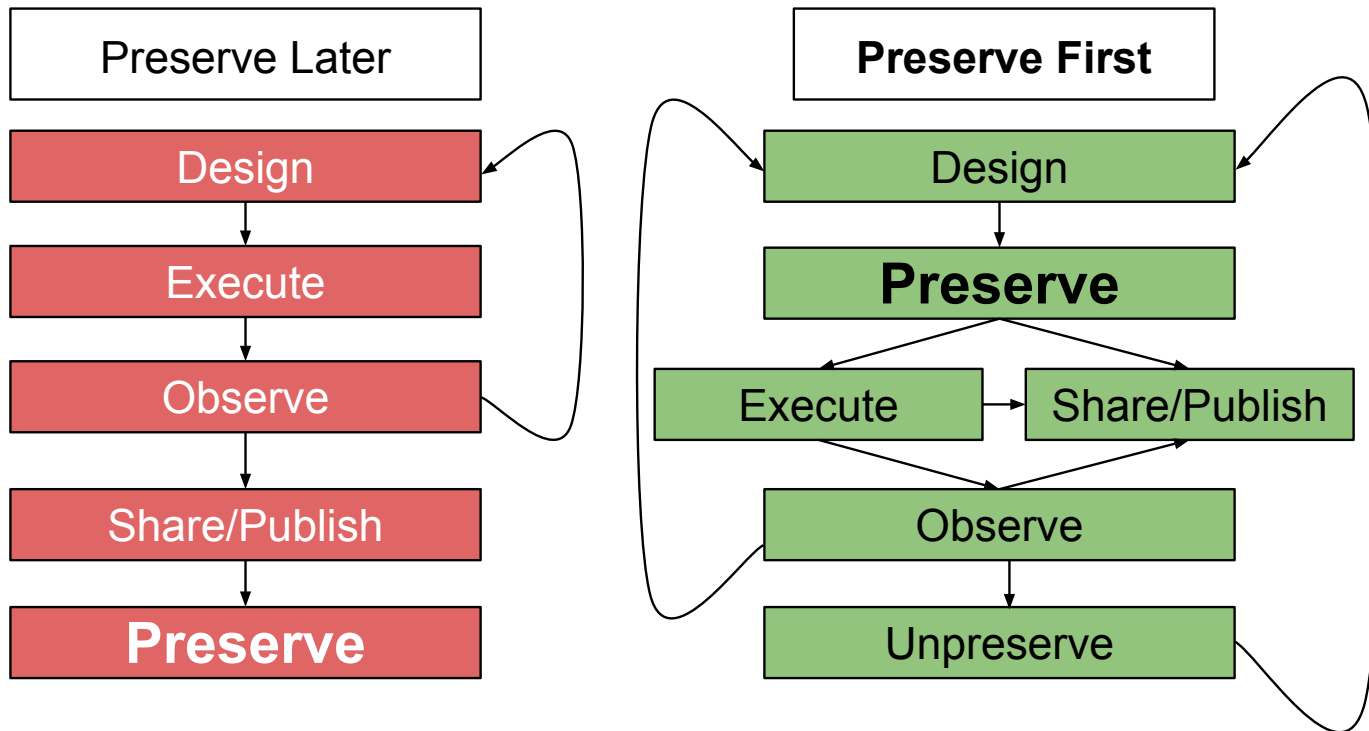
- Designed for Big Data
- Manage storage and compute resources
- Reproducible workflow specifications
- Share workflow with others
- Reshare changes back
- User defined granularity

Accepted philosophy



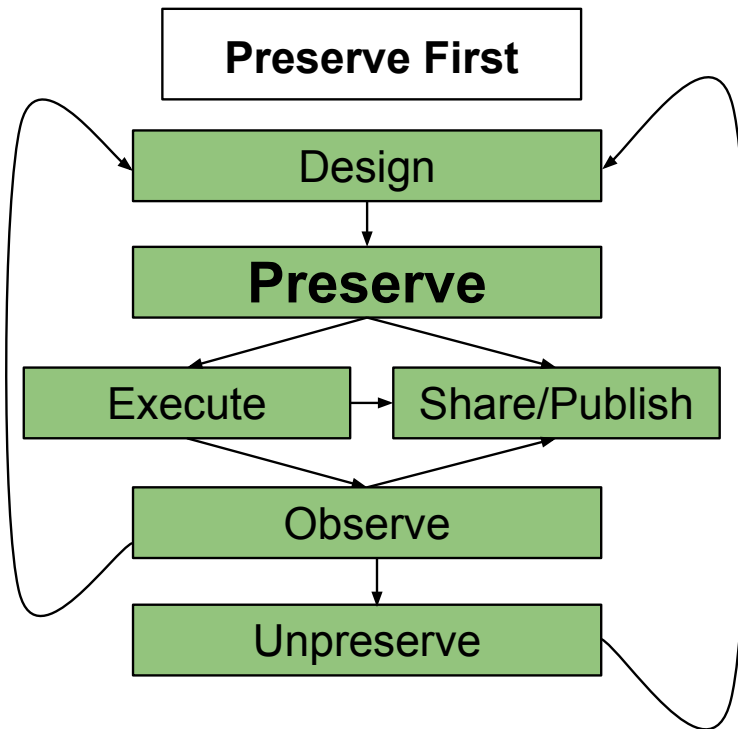
- Libraries
 - Hardware
 - Network
-
- System Administrators
 - Remote Collaborators
 - Graduated Students

Proposed philosophy



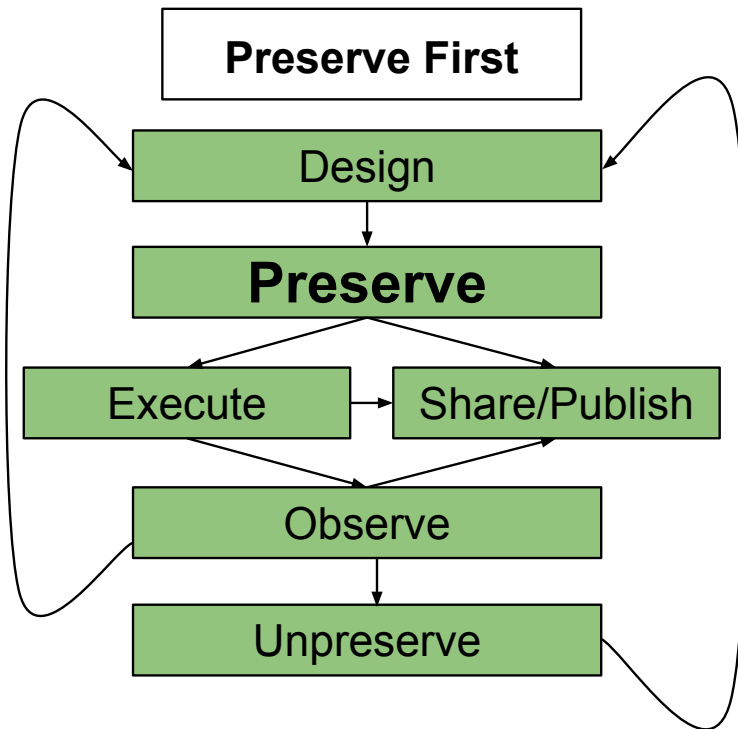
Differences

- Git: User decides when to preserve



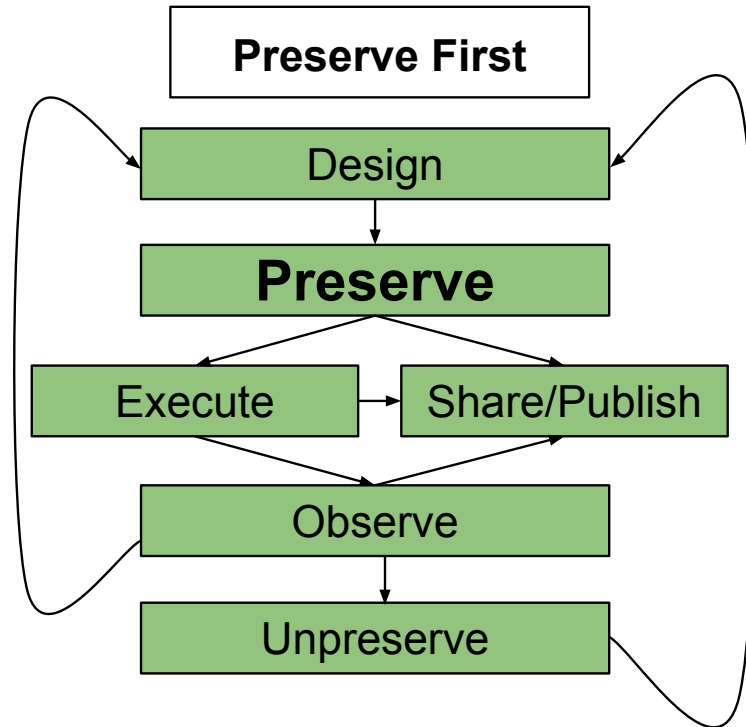
Differences

- ~~Git: User decides when to preserve~~
- Preserve ALL specification changes



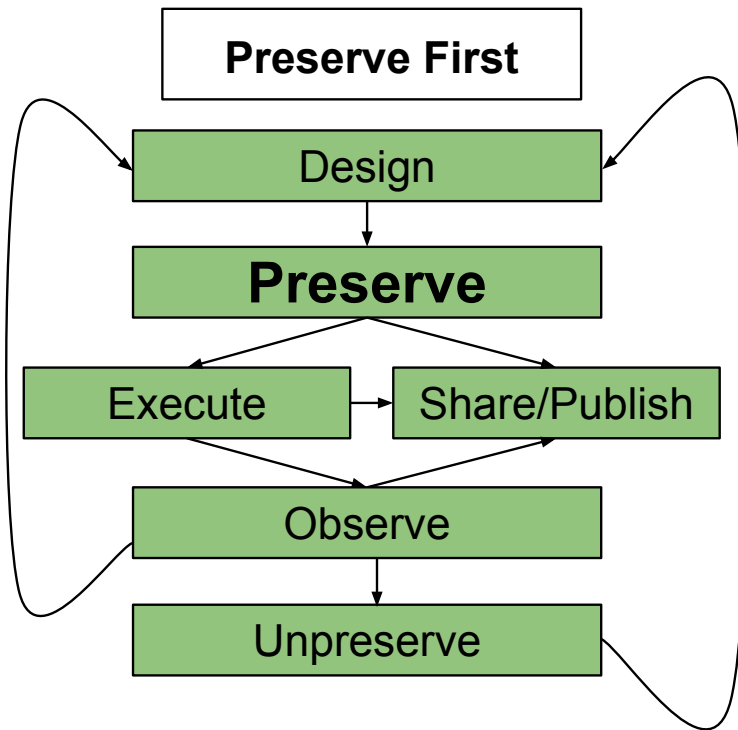
Differences

- ~~Git: User decides when to preserve~~
- Preserve ALL specification changes
- Git: Code Commits separate from Code Execution



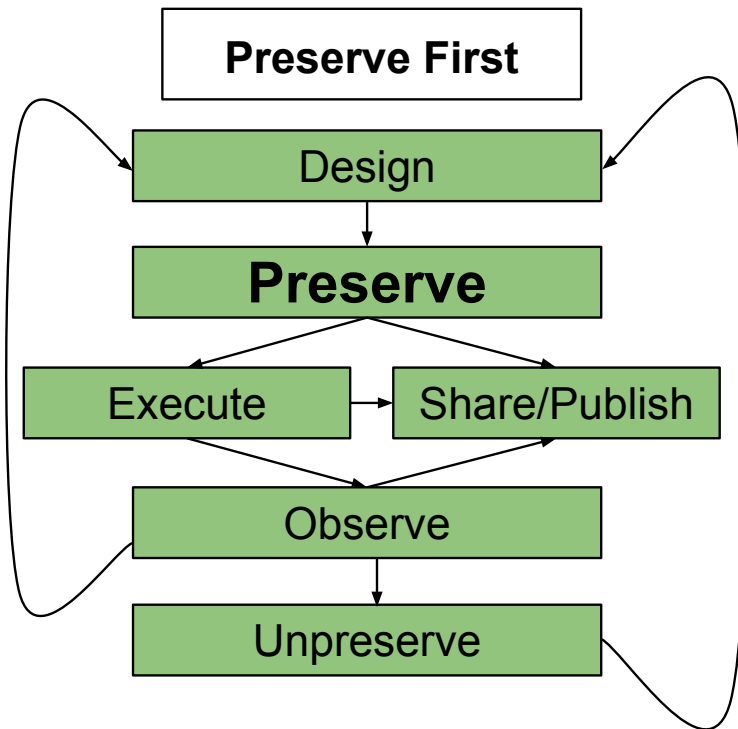
Differences

- ~~Git: User decides when to preserve~~
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- System Manages ALL computation

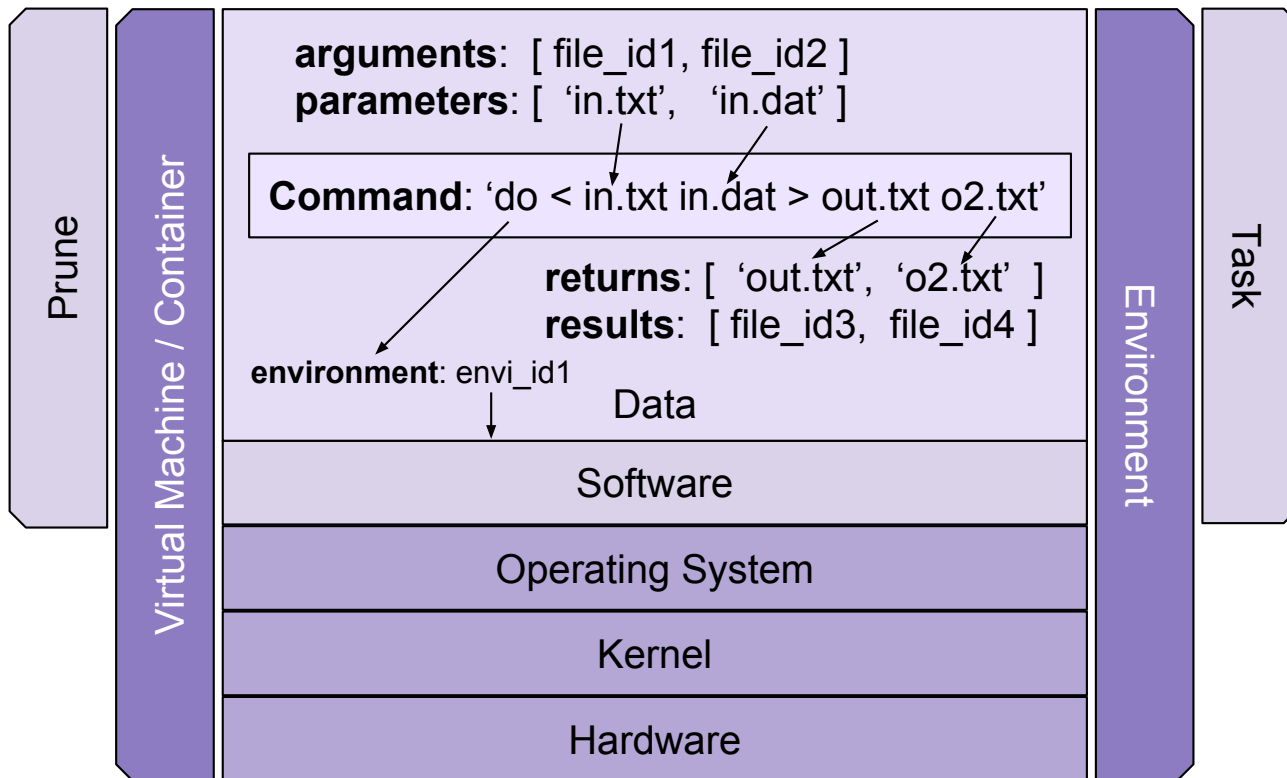


Differences

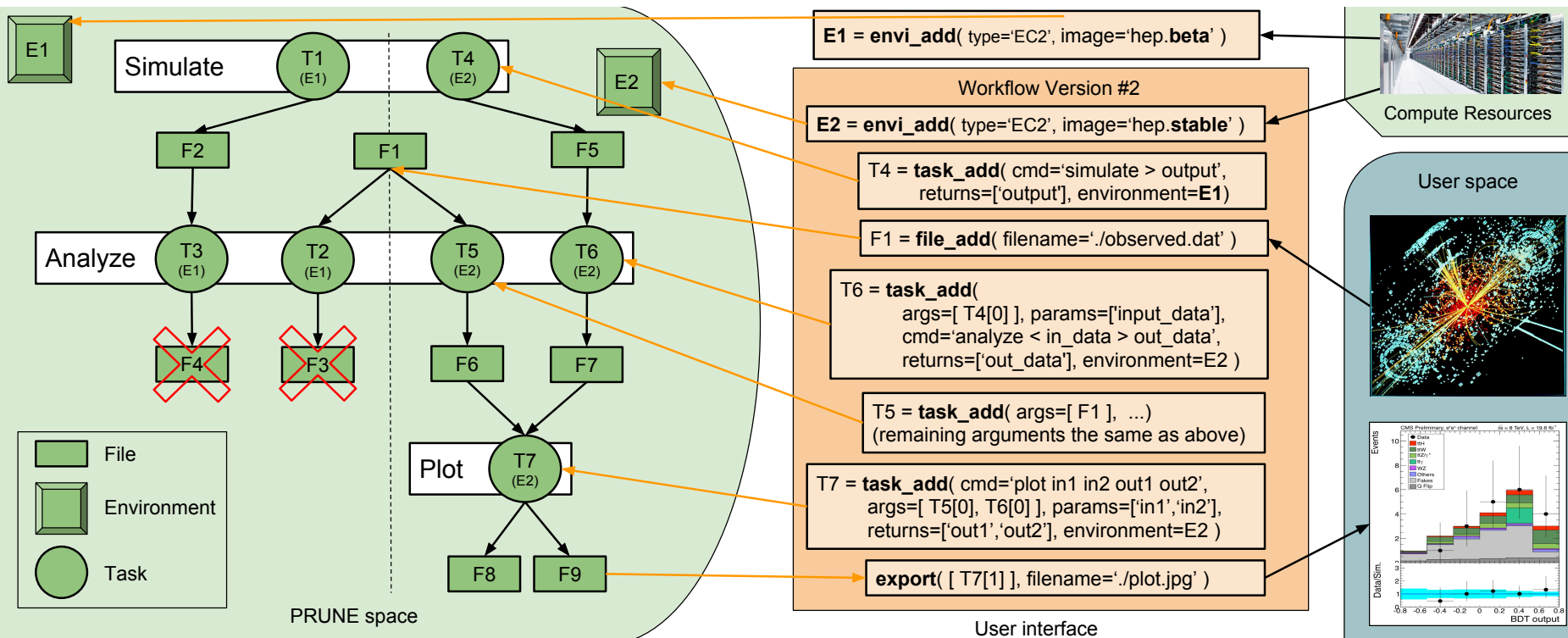
- ~~Git: User decides when to preserve~~
- Preserve ALL specification changes
- ~~Git: Code Commits separate from Code Execution~~
- System Manages ALL computation
- Remove unneeded items later on



What to Preserve



Overview



Sample code: Merge sort

```
#!/usr/bin/env python
from prune import client
prune = client.Connect() #Use SQLite3

##### Import sources stage #####
E1 = prune.env_add(type=`EC2`,
                    image=`ami-b06a98d8`)
D1, D2 = prune.file_add( `nouns.txt`, `verbs.txt` )
```

Sample code: Merge sort

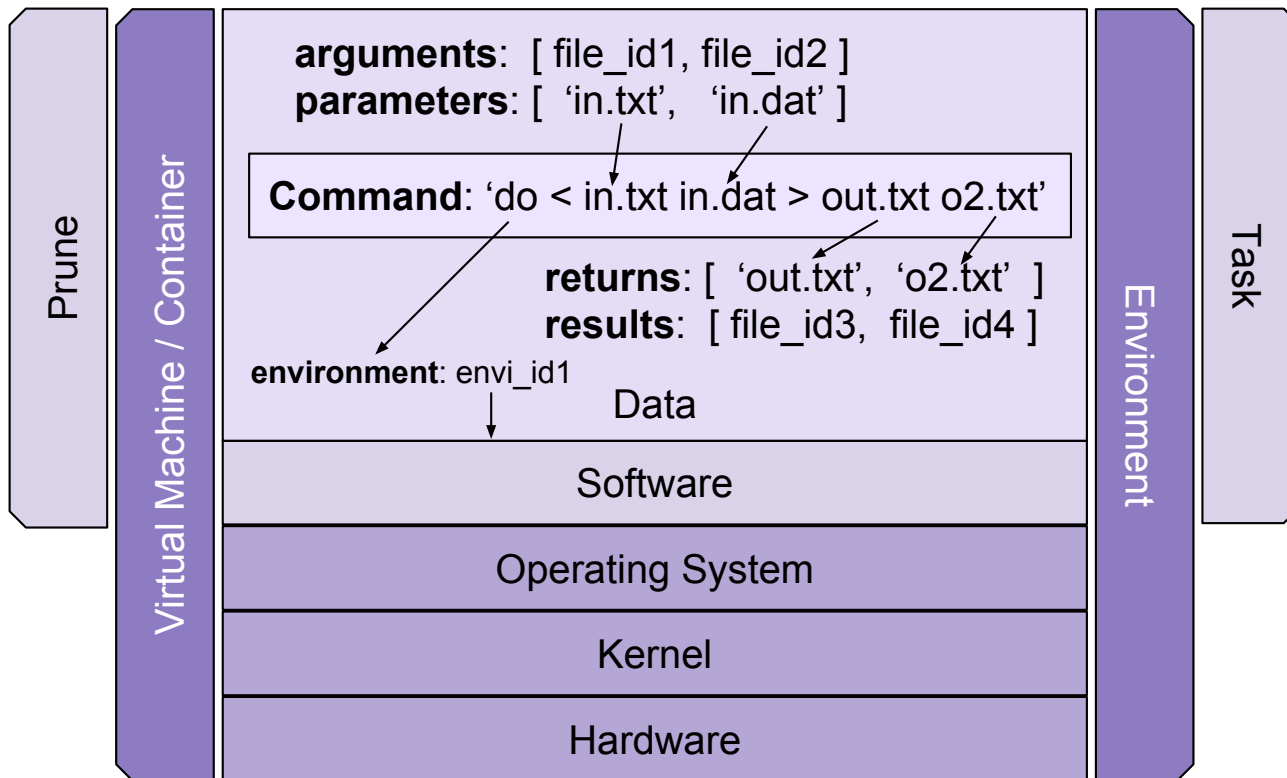
Sort stage

```
D3, = prune.task_add( returns=[`output.txt'],  
    env=E1, cmd=`sort input.txt > output.txt',  
    args=[D1], params=[`input.txt'] )  
D4, = prune.task_add( returns=[`output.txt'],  
    env=E1, cmd=`sort input.txt > output.txt',  
    args=[D2], params=[`input.txt'] )
```

Merge stage

```
D5, = prune.task_add(  
    returns=[`merged_out.txt'], env=E1,  
    cmd=`sort -m input*.txt > merged_out.txt',  
    args=[D3,D4], params=[`input1.txt',`input2.txt'] )
```

Prune Task



Sample code: Merge sort

Sort stage

```
D3, = prune.task_add( returns=[`output.txt'],  
    env=E1, cmd=`sort input.txt > output.txt',  
    args=[D1], params=[`input.txt'] )
```

```
D4, = prune.task_add( returns=[`output.txt'],  
    env=E1, cmd=`sort input.txt > output.txt',  
    args=[D2], params=[`input.txt'] )
```

Merge stage

```
D5, = prune.task_add(  
    returns=[`merged_out.txt'], env=E1,  
    cmd=`sort -m input*.txt > merged_out.txt',  
    args=[D3,D4], params=[`input1.txt',`input2.txt'] )
```

Sample code: Merge sort

Execute the workflow

```
prune.execute( worker_type='local', cores=8 )  
#prune.execute( worker_type='wq', name='myapp' )
```

Export

```
prune.export( D5, `merged.txt' ) # Final data  
prune.export( D5, `wf.prune', lineage=2 )
```

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Execute the workflow

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prune.execute( worker_type='local', cores=8 )  
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Export

```
prune.export( D5, `merged.txt' ) # Final data  
prune.export( D5, `wf.prune', lineage=2 )
```

Sharable workflow description file

```
{"body": {"args": ["f908ff689b9e57f0055875d927d191ccd2d6deef:0",  
"319418e43783a78e3cb7e219f9a1211cba4b3b31:0"], "cmd": "sort -m input*.txt > merged_output.txt", "env":  
"da39a3ee5e6b4b0d3255bfef95601890afd80709", "env_vars": {}, "params": ["input1.txt", "input2.txt"], "precise":  
true, "returns": ["merged_output.txt"], "types": [], "cbid": "e82855394e9dcdee03ed8a25c96c79245fd0481a", "size":  
322, "type": "call", "wfid": "a0230143-9b3a-4766-809d-5b7172e9b967", "when": 1476886144.7171359}  
{"body": {"args": ["29ae0a576ab660cb17bf9b14729c7b464fa98cca"], "cmd": "sort input.txt > output.txt", "env":  
"da39a3ee5e6b4b0d3255bfef95601890afd80709", "env_vars": {}, "params": ["input.txt"], "precise": true, "returns":  
["output.txt"], "types": [], "cbid": "f908ff689b9e57f0055875d927d191ccd2d6deef", "size": 241, "type": "call", "wfid":  
"a0230143-9b3a-4766-809d-5b7172e9b967", "when": 1476886144.484422}  
{"body": {"args": ["48044131b31906e6c917d857ddd1539278c455cf"], "cmd": "sort input.txt > output.txt", "env":  
"da39a3ee5e6b4b0d3255bfef95601890afd80709", "env_vars": {}, "params": ["input.txt"], "precise": true, "returns":  
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"wfid": "a0230143-9b3a-4766-809d-5b7172e9b967", "when": 1476886144.6183109}  
{"cbid": "29ae0a576ab660cb17bf9b14729c7b464fa98cca", "size": 144, "type": "file", "wfid":  
"a0230143-9b3a-4766-809d-5b7172e9b967", "when": 1476886144.2482941}
```

time

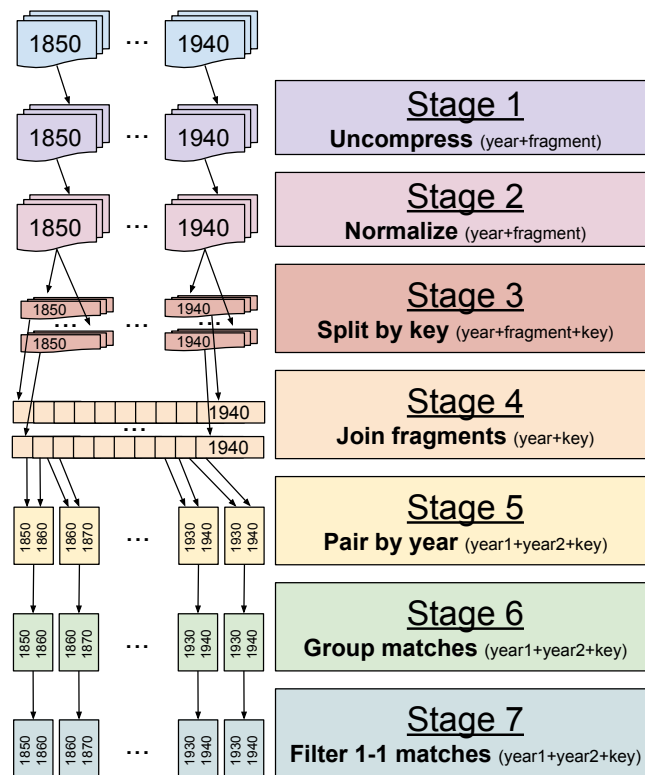
person

year

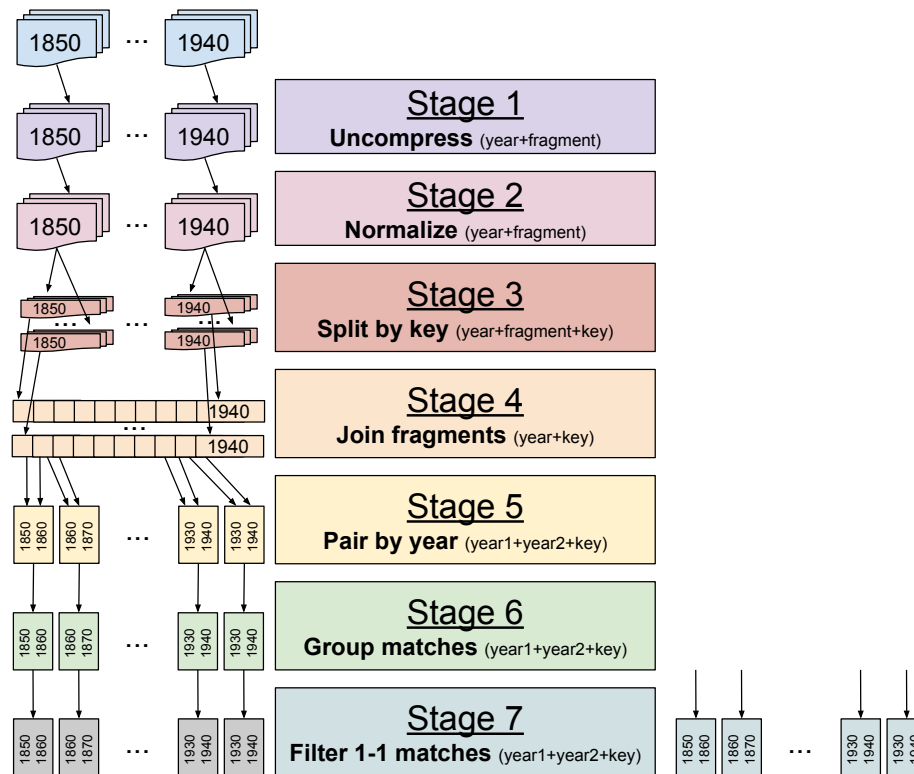
Way

...

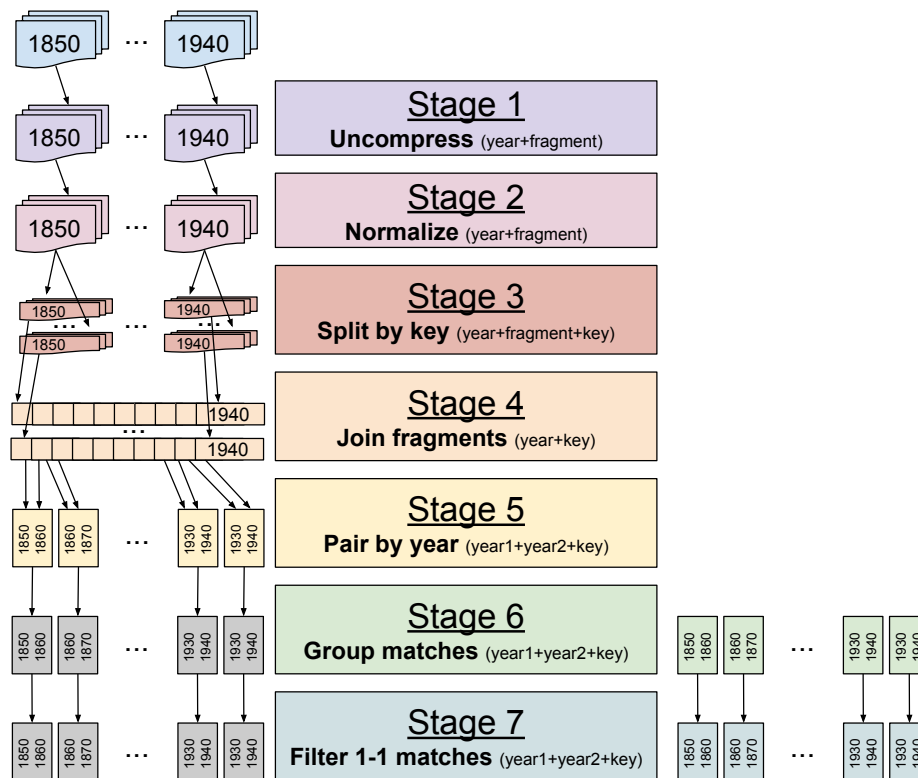
Workflow evolution (US Censuses)



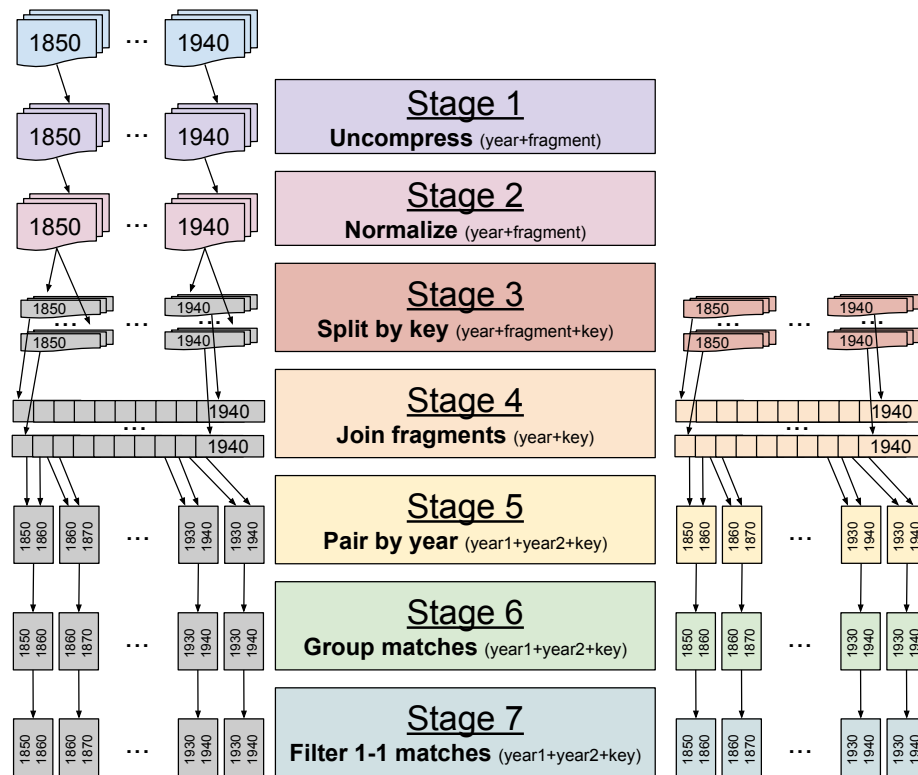
Redefine filter criteria



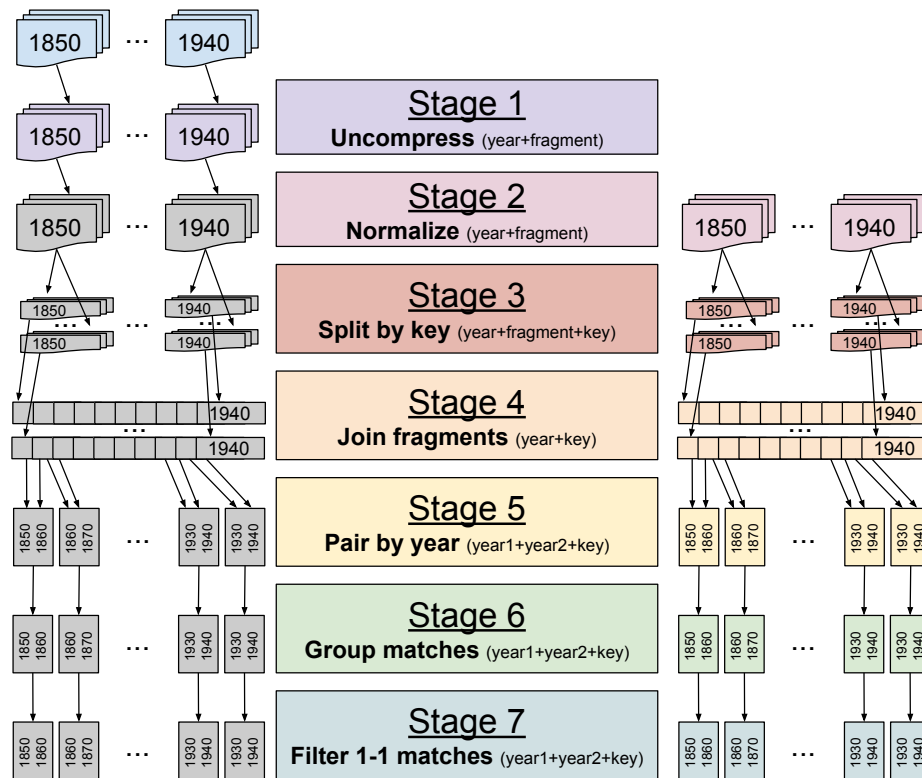
Redefine match criteria



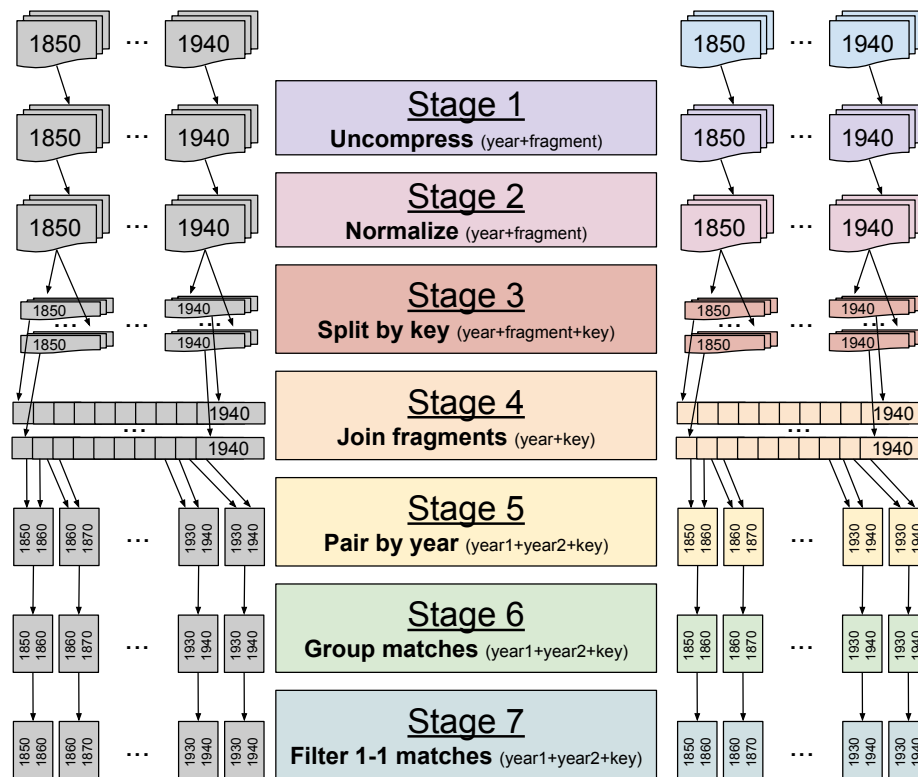
New key function chosen



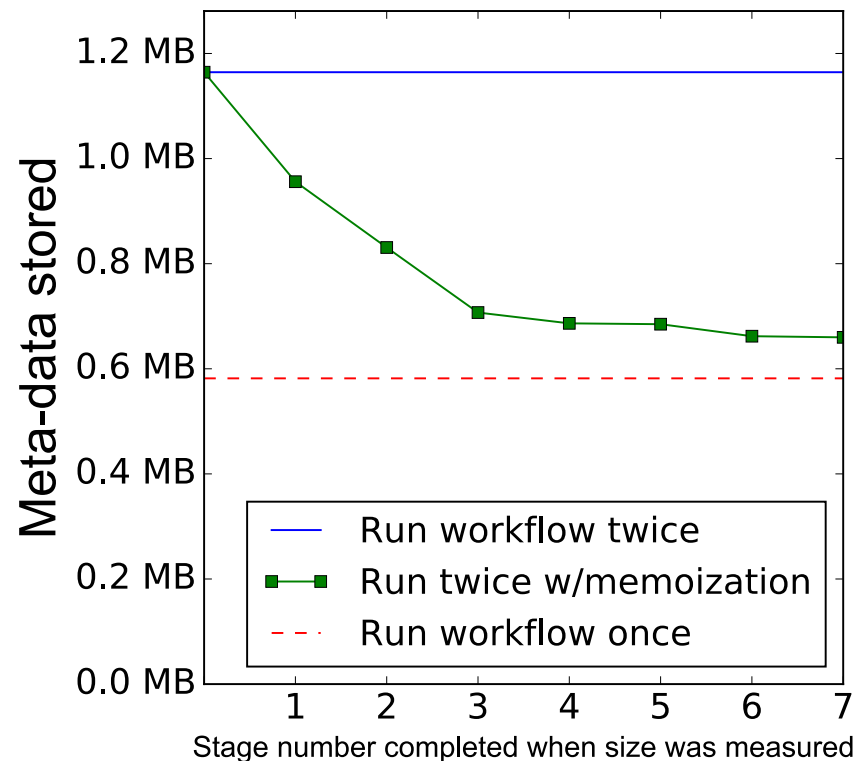
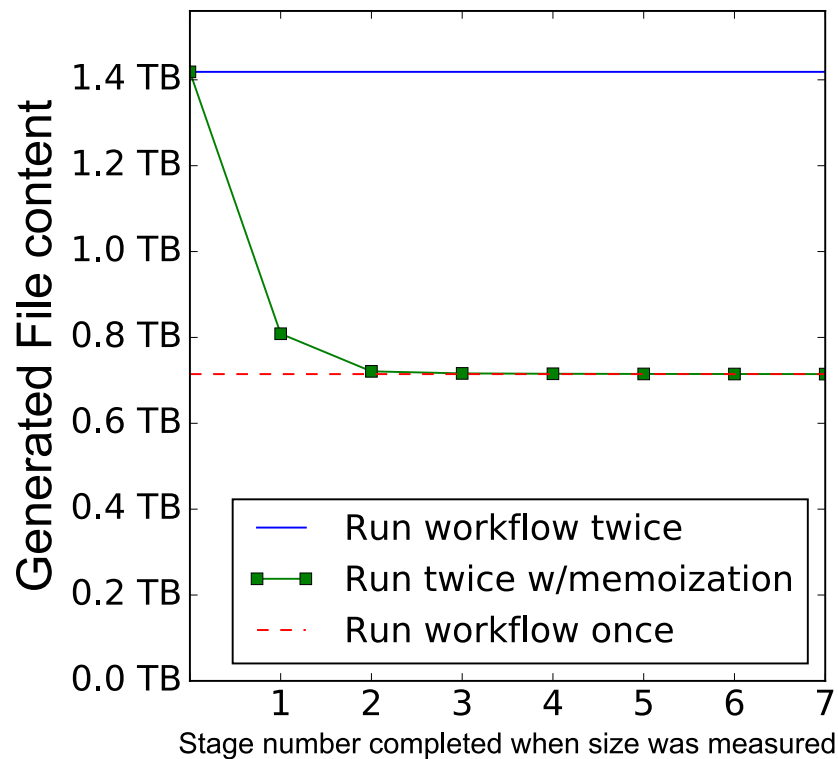
Re-normalize



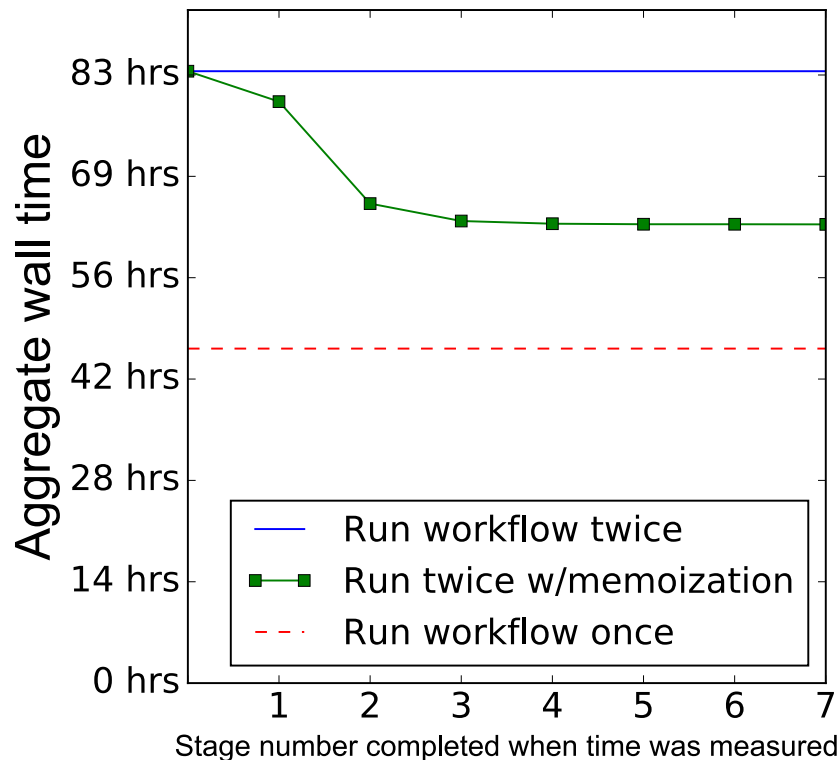
New input data



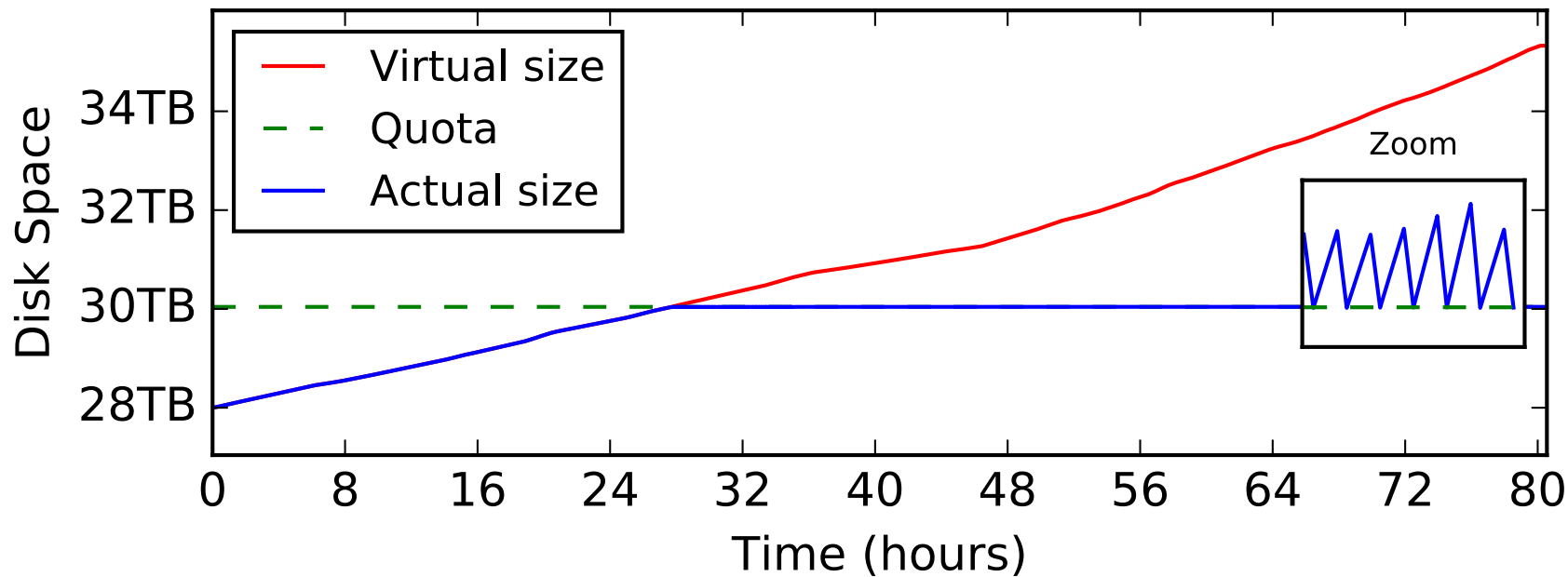
Derivation History = Cachable Results



Execution time cut in half for run #2

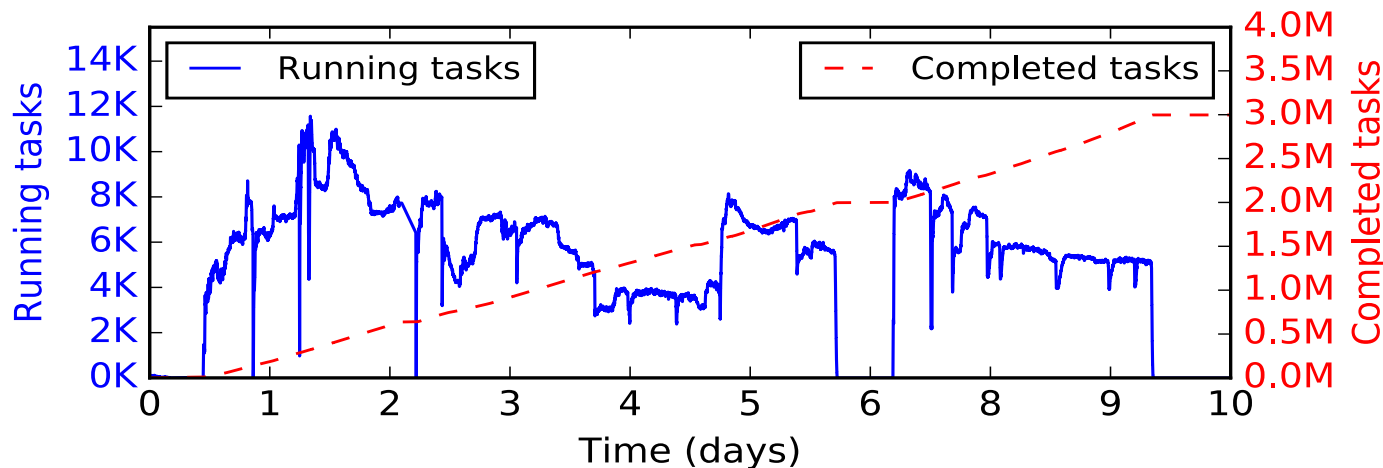


Quotas



Scalability

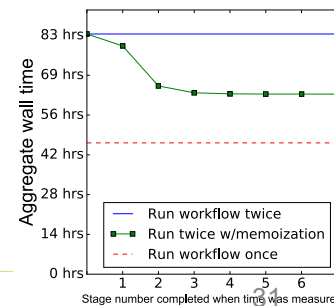
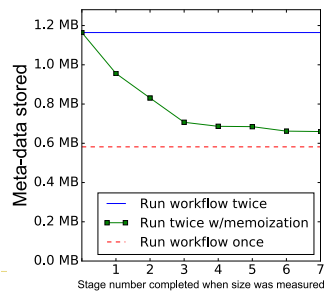
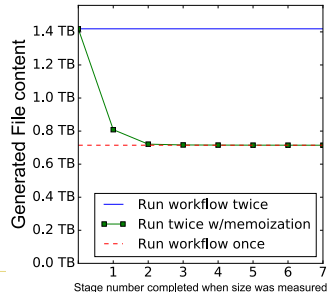
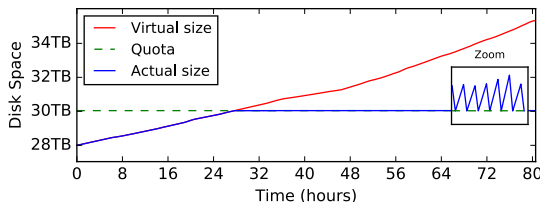
- ~12,000 parallel cores
- ~3 million tasks
- Overhead
~1% above
native wall clock



Sharing workflow between users

```
{
  "body": {
    "args": [
      "f908ff689b9e57f0055875d927d191ccd2d6deef:0",
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    "params": [
      "input1.txt",
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    ],
    "precise": true,
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      "merged_output.txt"
    ],
    "types": [],
    "cbid": "e82855394e9dcdee03ed8a25c96c79245fd0481a",
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  "when": 1476886144.2482941
}
```

time
person
year
Way
...





<http://ccl.cse.nd.edu/research/papers/>

- Sample workflows
- <http://ccl.cse.nd.edu/software/prune/prune.html>
 - Merge sort
 - Pairwise comparisons (US Censuses)
 - High-energy Physics

For more information: pivie@nd.edu



Thank You!



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