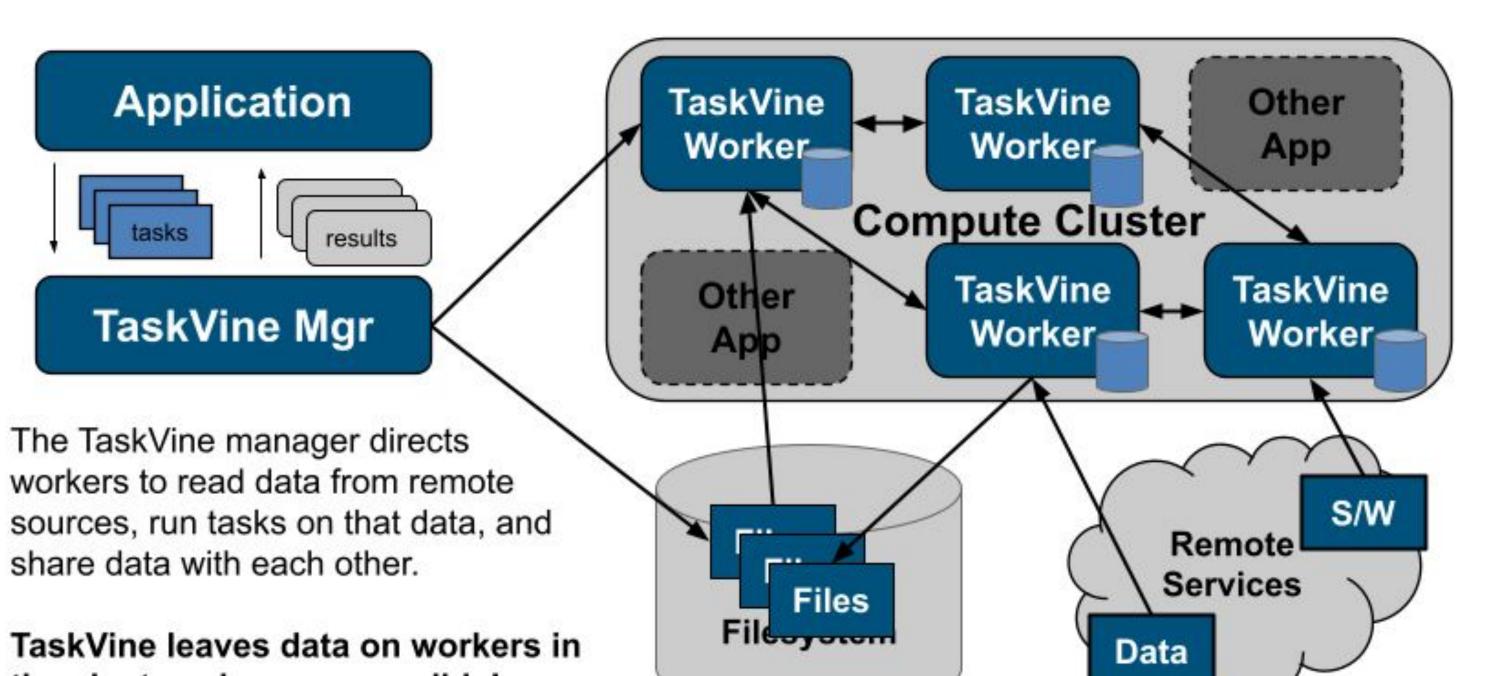


# CSSI Element: <del>DataSwarm</del>: TaskVine: A User-Level Framework for Data Intensive Scientific Applications PI: Douglas Thain, University of Notre Dame, Award #: 1931348



Many scientific applications are expressed as high-throughput workflows that consist of large graphs of data assets and tasks to be executed on large parallel and distributed systems. A challenge in executing these workflows is **managing data**: both datasets and software must be efficiently distributed to cluster nodes; intermediate data must be conveyed between tasks; output data must be delivered to its destination. Scaling problems result when these actions are performed in an uncoordinated manner on a shared filesystem. TaskVine is a system for exploiting the aggregate local storage and network capacity of a large cluster. TaskVine tracks the lifetime of data in a workflow – from archival sources to final outputs-- making use of local storage to



# Example of key TaskVine operaitons import taskvine **as** vine m = vine.Manager(9123)

# File objects are first class citizens. = m.declareFile("mydata.txt") file buffer = m.declareBuffer("Some literal data") = m.declareURL("https://nd.edu/data.tar.gz") lurl = m.declareTemp(); temp

# Peform a standard transformation on a file = m.declareUntar( url ) data software = m.declarePoncho( package )

#### distribute, and re-use data wherever possible.

the cluster wherever possible!



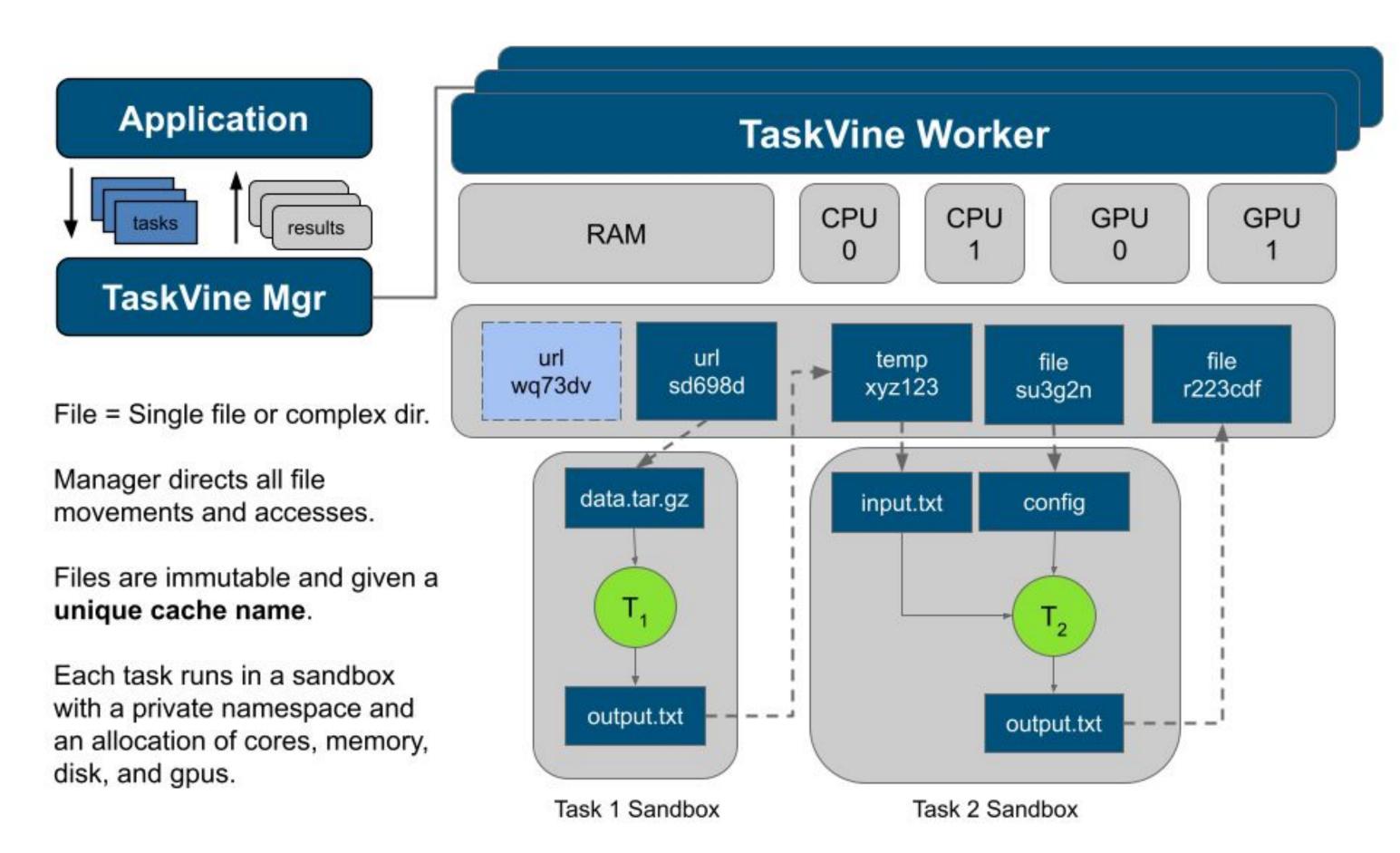
# Submit a standard executable task. task = vine.Task("mysim.exe -p 50 input.data -o output.data")

t.add\_input(url,"input.data") t.add\_output(temp, "output.data") t.set\_cores(4) t.set\_memory(2048) t.set\_disk(100) taskid = m.submit(t)

# Submit a Python function execution t = vine.PythonTask( simulate\_func,molecule,parameters) taskid = m.submit(t)

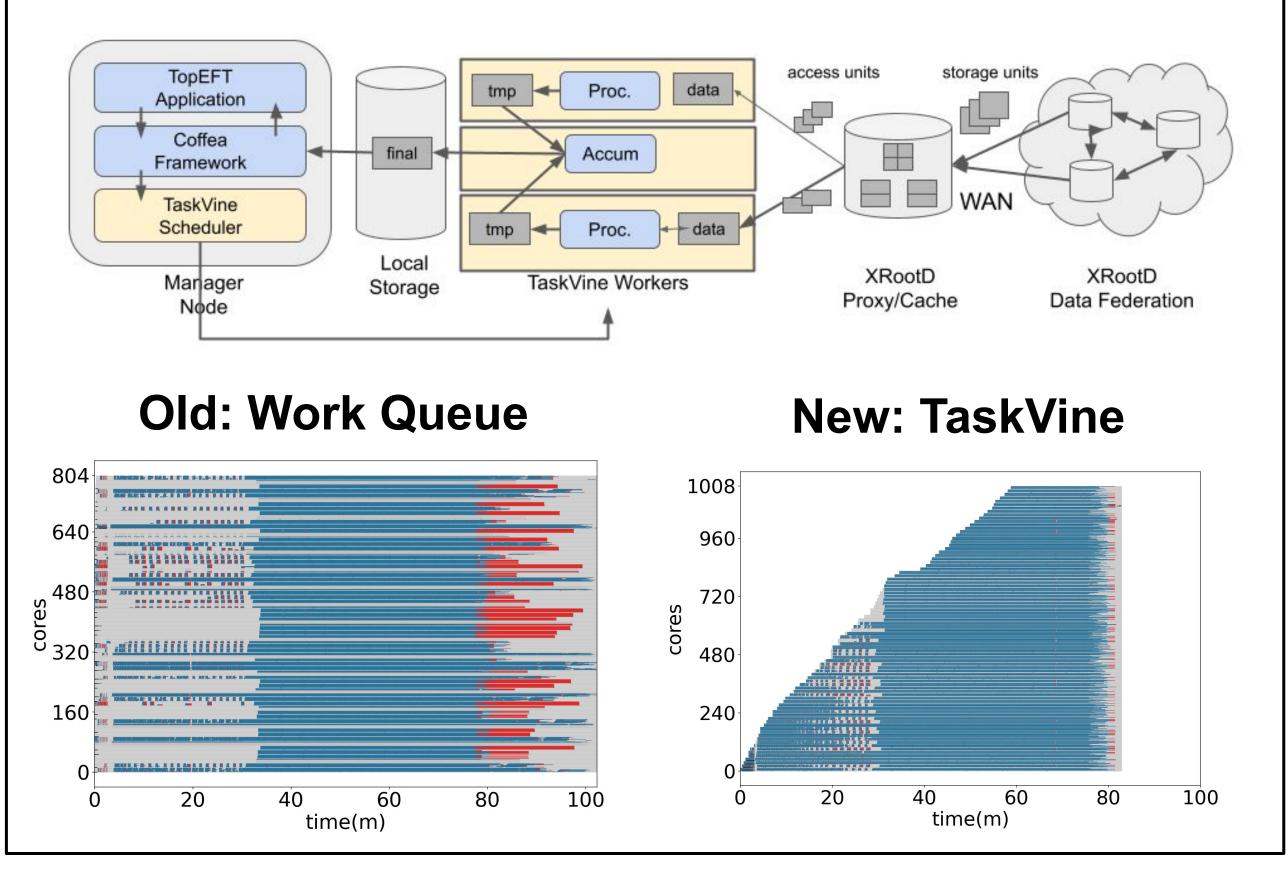
# Wait for any task to complete. = m.wait() print( t.output )

2.26 fb<sup>-1</sup>, 2015 (13 TeV sigle\_atop single\_top single\_tW wjets ttbar Data Makeflow Dask Parsl TaskVine Manager Worker Worker Worker Compute Cluster



**TopEFT Physics Data Analysis Application** 

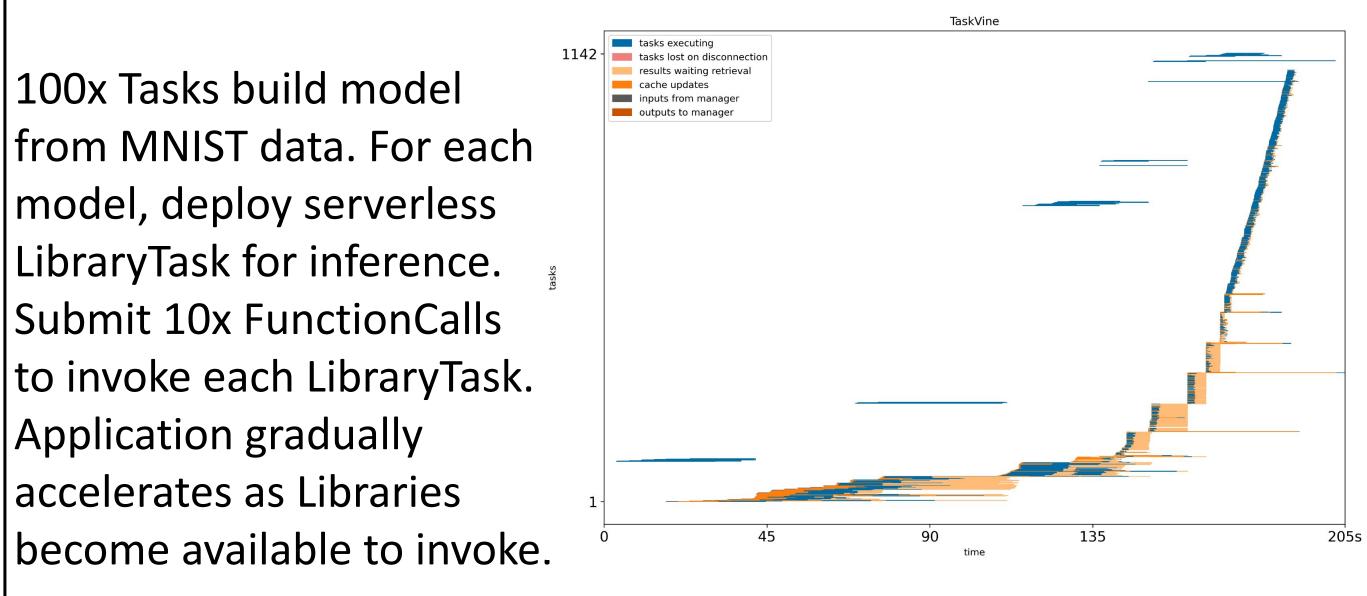
TaskVine Storage vs Shared Filesystem



#### Effect of TaskVine's data management and distribution method compared to relying on a shared filesystem for data staging. Workflow: Training 2048 NN configurations, each 2 cores/2GB, with 8 tasks Workflow execution time per worker on an vine-hot 7000 HTCondor pool 6000 using either 5000 Panasas or (s) o 4000 TaskVine for E 3000 data mgmt. 2000 Software deps are 4.4GB of 1000 -Python libraries Number of workers per node!

#### **Multi-Modal Serverless Workflows**

Simply converting "import tensorflow" into the preamble of a Library task saves **1.2GB** of Python libraries, **30K** metadata system calls, and **5-10s** latency per FunctionCall. We can mix standard Tasks, Libraries, and FunctionCalls into a multi-modal workflow:



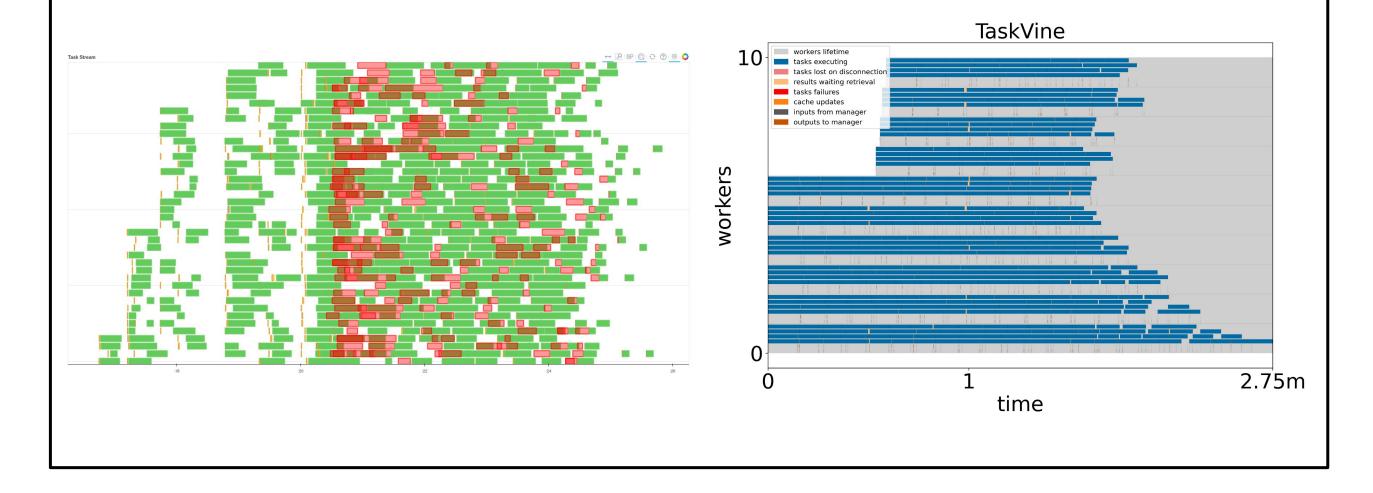
### Dask.Distributed vs. Dask.TaskVine

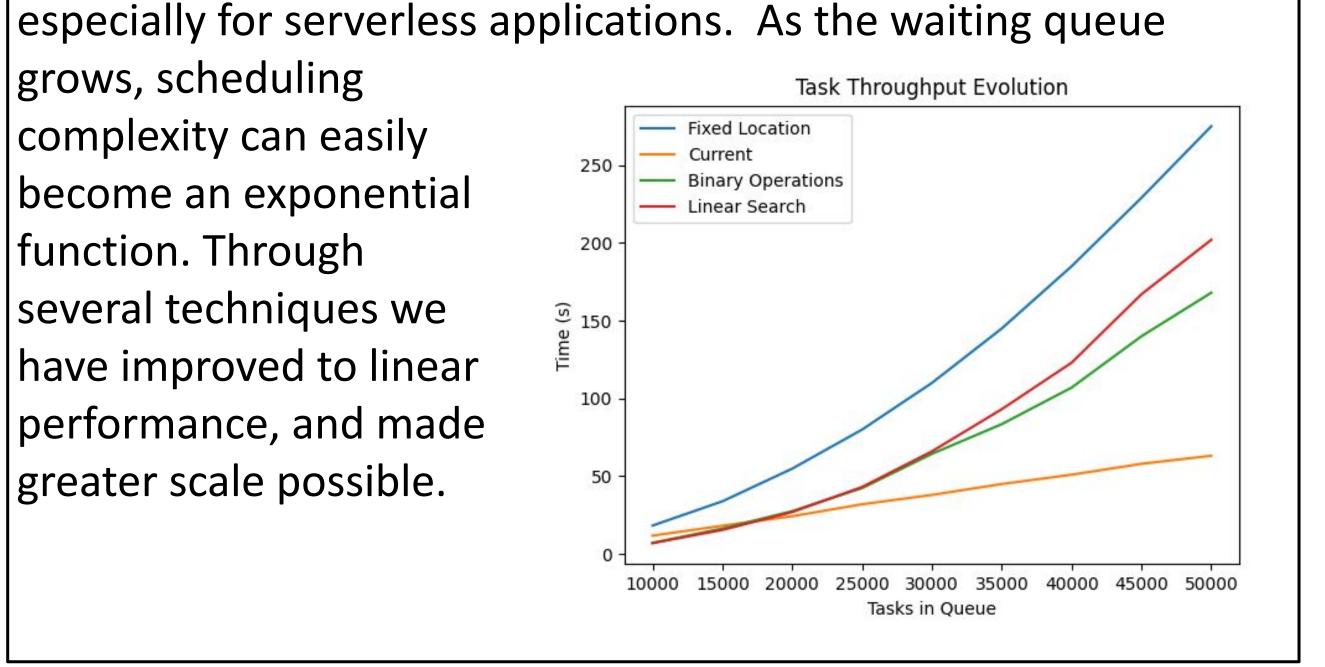
For some workflows, TaskVine significantly outperforms Dask native distributed workers. The following workflow performs a series of matrix multiplication tasks forming a tree of task executions. 256 initial multiplication tasks feed into new tasks and son on, generating a final matrix. Each matrix is 1.2MB which is transferred between workers when needed.

## TaskVine Scheduler Scaling

The TaskVine scheduler must consider a number of aspects while making task placement and data movement decisions without compromising high-throughput task dispatch capabilities,

Barry Sly-Delgado, Thanh Son Phung, Colin Thomas, David Simonetti, Andrew Hennesse, Ben Tovar, and Douglas Thain, "TaskVine: Managing In-Cluster Storage for High Throughput Data Intensive Workflows", WORKS Workshop at Scupercomputing, November 2023.





#### **Cooperative Computing Lab at the University of Notre Dame**

