

# **Enabling Graph-Based Profiling Analysis using Hatchet** Student: lan Lumsden<sup>1</sup> Mentors: Stephanie Brink<sup>2</sup>, Michael R. Wyatt II<sup>1</sup>, Todd Gamblin<sup>2</sup>, Michela Taufer<sup>1</sup>

# Introduction

- Profiling is a way to measure the performance of code and how code runs on systems in high-performance computing (HPC)
- Numerous tools for HPC profiling (e.g. TAU, Caliper, HPCToolkit) have custom data format and analysis tools
- Users are locked into types of analysis dictated by the provided tools • Hatchet [1] is a new, general data analysis tool that can read HPC profiling data from **different profilers** 
  - Store the raw performance data into a pandas DataFrame
- Represent the relational *caller-callee* data with a directed acyclic graph
- Hatchet restricts users to table-based analysis of the raw performance data
- Hatchet does NOT support analysis of the relational data collected by HPC profilers

1. Collect Data • Run each of the following benchmarks with MVAPICH2 (M) and Spectrum-MPI (S) using 64, 128, 256, and 512 MPI ranks, and profile the runs with HPCToolkit

- AMG2013
- Kripke
- Lammps
- Load the generated profiles into Hatchet
- Perform the benchmarking on LLNL's Lassen supercomputer

### 2. Extract MPI Layer

- Filter the *call graphs* to get subgraphs rooted at standard MPI function calls
- Query Path used to extract MPI Layer: [{"name": "P?MPI\_.\*"}, "\*"]





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# 3a. Calculate Percent MPI Time for each MPI Function<sup>+</sup> Kripke 40 <u>کی</u> 20 512 512 256 Number of MPI Ranks

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### Our graph-based filtering query language consists of:

- User Input: A Query Path represented as a list of abstract graph nodes
- Algorithm:
- Read and parse the user's query path
- Match real nodes in the graph being filtered to the abstract nodes in the query path
- Collect all graph paths that match the full query path
- Create a new graph containing only the nodes in the matched paths



<u>References</u> [1] A. Bhatele et al. 2019. Hatchet: Pruning the Profiles. In *Proc. of SC19* runtime.

**†** "Remaining MPI Time" consists of all MPI functions that contribute less than 5% of the overall MPI runtime **‡** "Remaining MPI Time" consists of all child functions Overgrowth in Parallel that contribute less than 10% of the overall MPI





<b>MPI Function Calls</b>				
	MPI_Finalize		MPI_Send	
	MPI_Allreduce		MPI_Wait	
	MPI_Allgather		MPI_Waitany	
	MPI_Waitall		MPI_Alltoallv	
	MPI_Testany		Remaining MPI Time	





# Methods



Figure 2: Example of filtering a graph using the new query language

# Lessons Learned

## Using the query language and Hatchet, we were able to: • Extract all call paths specific to a given library Determine the performance contributions of function calls used by these libraries • Correlate children function calls to specific important library API calls in an application Use this correlation to determine children function calls that contribute the most to the performance of the targeted library API call • Compare the correlation of children and API calls across libraries to determine possible causes for performance differences In our tests with MVAPICH2 and Spectrum-MPI, we learn that: • The **pthread\_spin\_lock** function is consistently one of the most impactful in the performance on MPI functions for both libraries In AMG2013, the worse performance of MPI Allgather in Spectrum-MPI can possibly be attributed to pthread\_spin\_lock **Child Function Calls**

unknown file> [libmlx5.so.1.0.0]:1133		<unknown file=""> [libmlx5.so.1.0.0]:0</unknown>				
pthread_spin_lock.c:26		syscall-template.S:81				
pml_pami_send.c:0	pml_pami_init.c:0					
memset.S:1133		Geometry.h:0				
malloc.c:0		Remaining MPI Time				
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