

Analysing Parallel I/O using Allinea Tools

Introduction to I/O profiling at scale

Keeran Brabazon, ARM Parallel I/O Tutorial, ISC,18th June 2017



Allinea – What is it?



 Help the HPC community develop and design the best applications and make the most use of HPC clusters

• HPC Tools company 2002 - 2016

- Part of ARM since December 2016
 - Continue to improve tools for new uses in HPC
 - Support for all HPC applications and architectures



Allinea Products



- Allinea Forge
 - Combined debugging (DDT) and profiling (MAP) in the same interface
 - Designed for application developers



- Summary of application performance
- Designed for system administrators



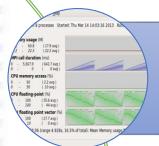


 For more information see www.allinea.com



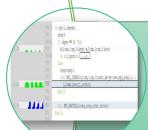
Performance Analysis with Allinea MAP





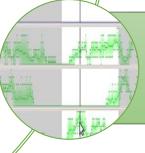
Low overhead measurement

- Accurate, non-intrusive application performance profiling
- Seamless no recompilation or relinking required



Easy to use

- Source code viewer pinpoints bottleneck locations
- Zoom in to explore iterations, functions and loops



Deep

- Measures CPU, communication, I/O and memory to identify problem causes
- Identifies vectorization and cache performance



Scalability



- Allinea tools are to be used at scale
 - Use on 100k+ cores
 - Profiles remain on the order of 10s of MB whether running on 100 processes for 30 seconds or 100k processes for 10 hours
 - When evaluating performance of an HPC application you should run at HPC scales





- Program should be compiled with compiler optimisations left on
 - Optimisation should be done on production run
- Debug symbols need to be included
 - -g flag in GNU and Intel compilers
 - Gfast compilation flag for Cray compiler





- Simply prefix map --profile to execution command
 - Also in batch submission script
- A .map is generated with filename including time-stamp in working directory
 - Use -o <output_fname>.map to write to named file









```
mpiexec -n 1024 ./my_mpi_exe command line args ...
```







Viewing Performance Data in Allinea MAP



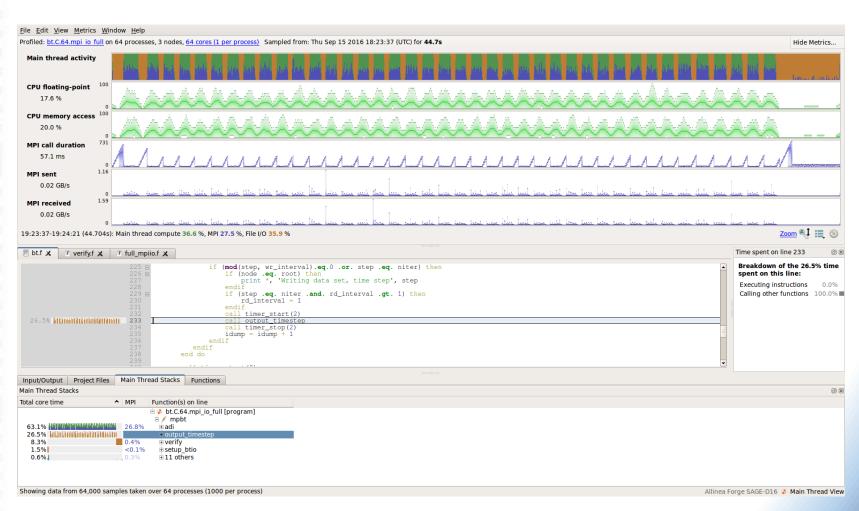
- Using the Allinea MAP GUI to view a single file
 map my_map_file.map
- Using scripts to view data from multiple profiles
 - 1. Export to JSON

2. Extract and view data. Example scripts at https://github.com/arm-hpc/allinea_json_analysis



Example MAP Profile









Visualising I/O Performance with Allinea MAP





Consider the application benchio from the earlier tutorial session

- Writes to file with:
 - 1 stripe
 - Lustre selected striping
 - Default striping (4 on Archer)



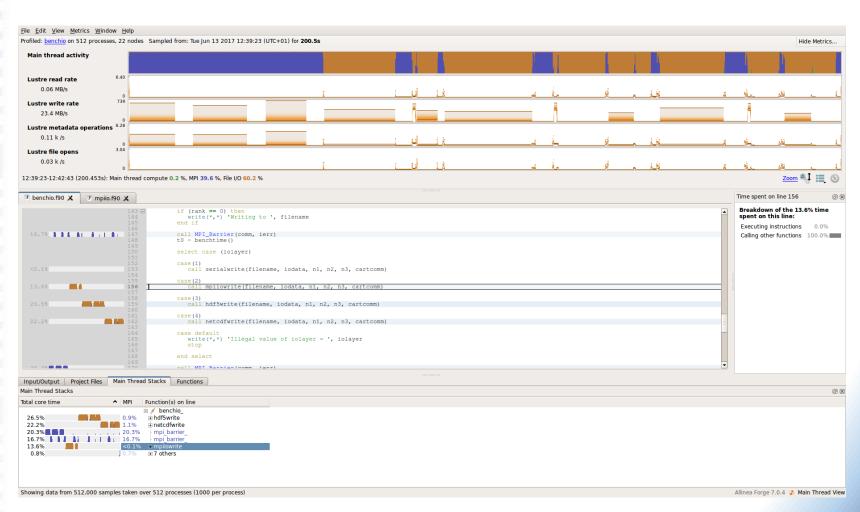


 Consider the application benchio from the earlier tutorial session

- Writes to file using:
 - Serial I/O
 - MPI I/O
 - HDF5
 - NetCDF







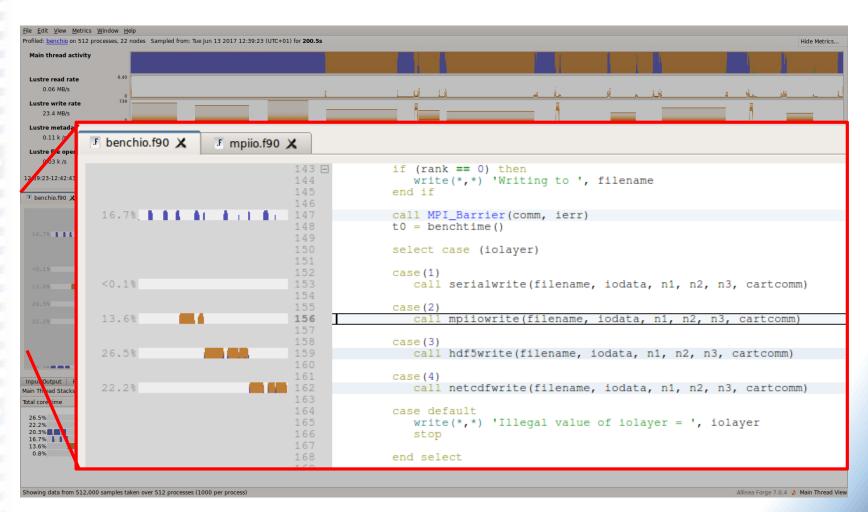






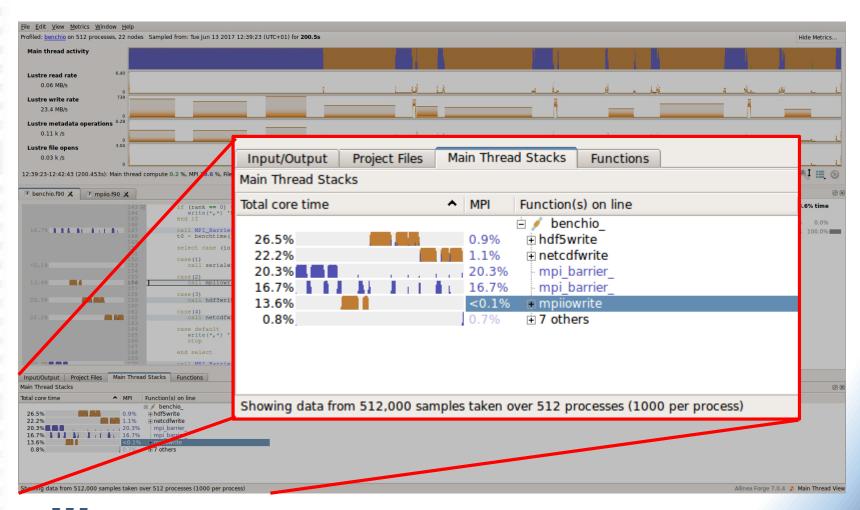






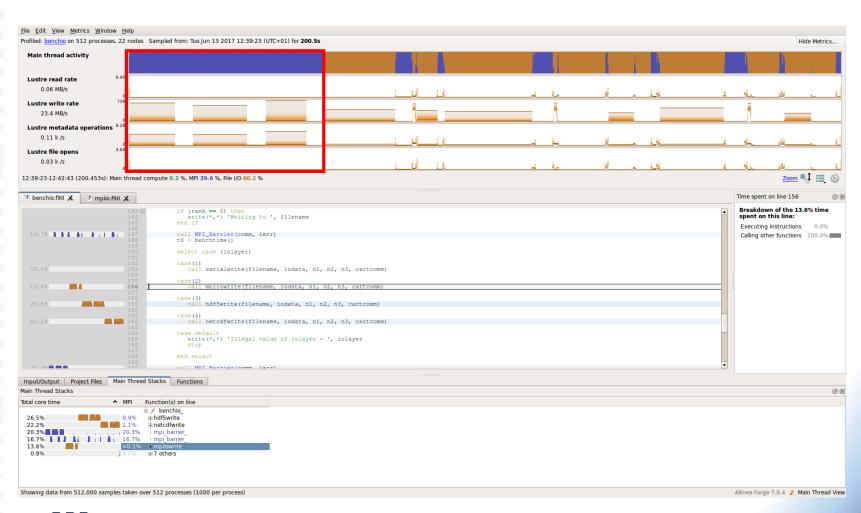






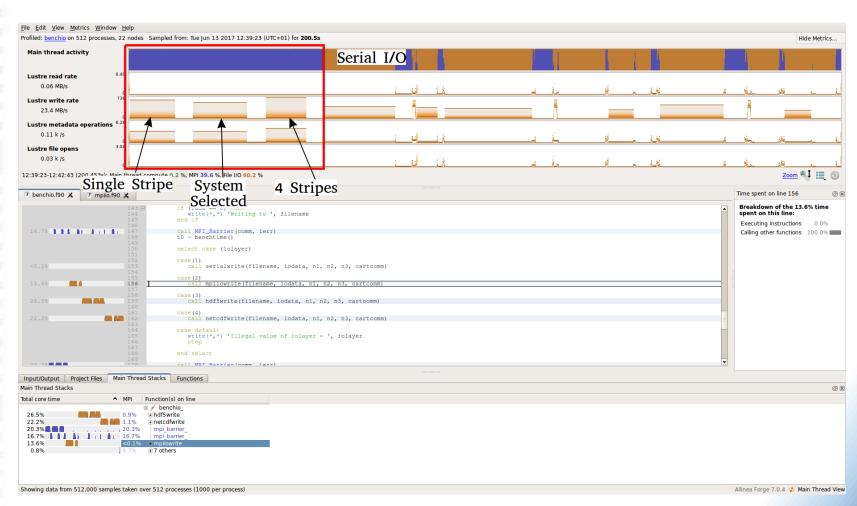






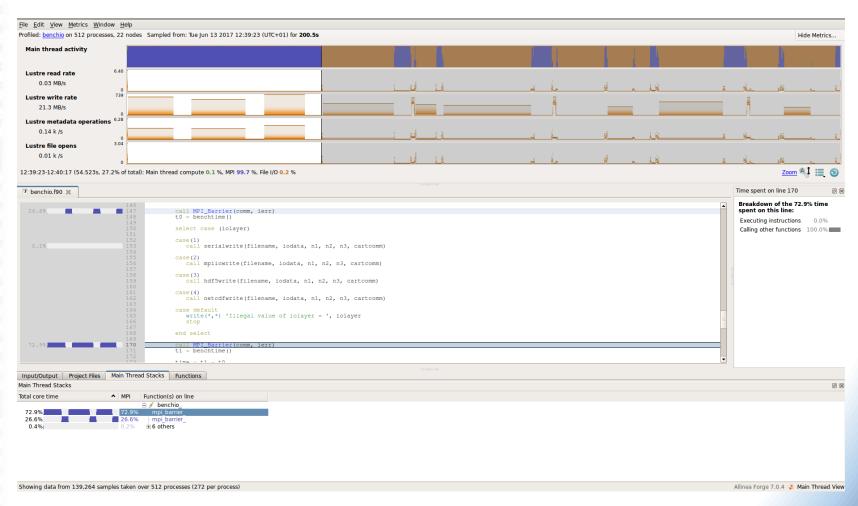






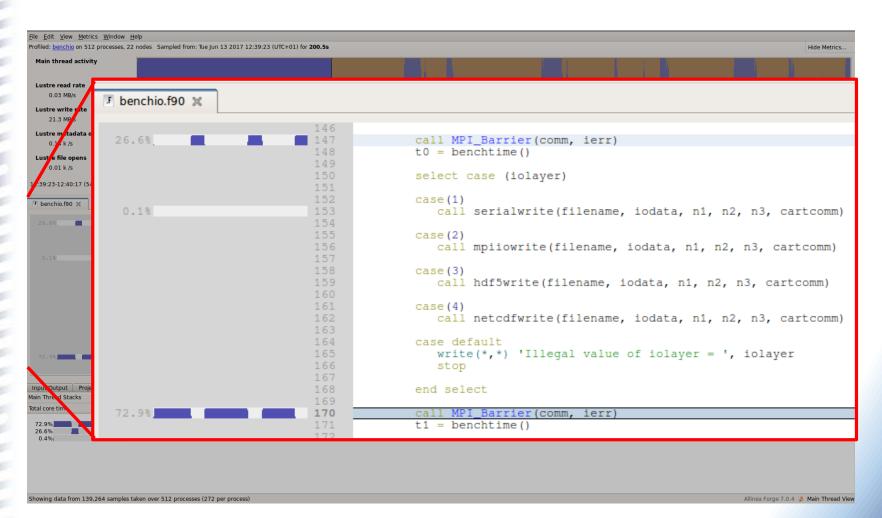






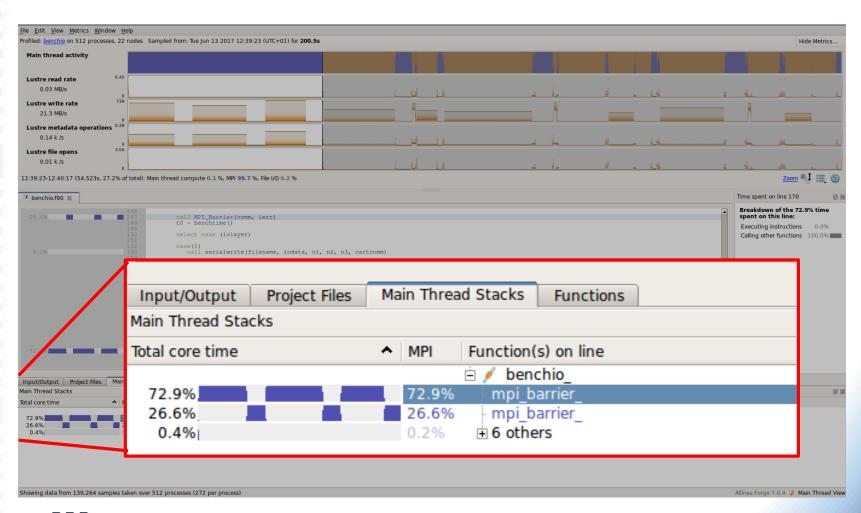






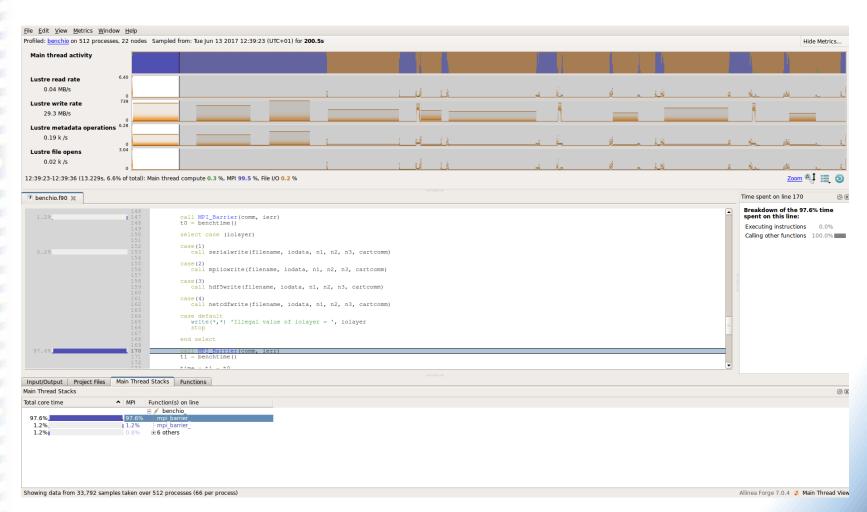






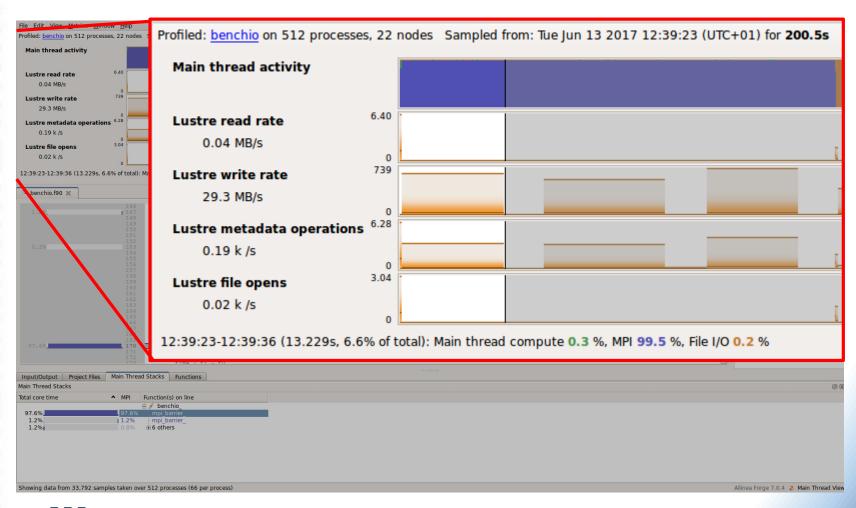














Summary



Interesting behaviour can be viewed in a single profile

 Viewing data in a timeline can show extra information with respect to the benchmark output



Summary



- Further investigation of the profiles to be done in the hands-on sessions
- Consider not only the bandwidth, but the overhead, which is shown as 'gaps' in the write rate in the profiles collected
- To draw conclusions on the performance of different paradigms need a set of runs, at different scales with different volumes of data written (in total and per-process)





Questions?

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