Applying MPI to Manage HPC-scale Datasets

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Dataset file and byte counts increase as users run larger MPI jobs

Scatter plot of dataset bytes vs files top 20 Lustre users





Motivating example: Consider the task of copying a large dataset

Single directory

200,000 files

128 MB per file

24.4 TB total

Goal: cp -r dir dir2





Approach 1: Use cp on a login node

>>: date; cp -r dir dir2 Tue Jun 11 10:20:35 PDT 2019

... check in after 8 hours ...

>>: date; ls -ltr dir2/ | wc
Tue Jun 11 18:20:36 PDT 2019
 114352 1029161 7712420

Completed ~114,352 of 200,000 files in 8 hours.

Some math ... (200,000 / 114,352) * 8 hours ~ **14 hours to finish**



Approach 2: Use dcp on 8 compute nodes

>>: bsub -nnodes 8 -Is /bin/bash

>>: mpirun -np 320 dcp dir dir2

[2019-06-10T18:14:16] Started: Jun-10-2019,18:08:01 [2019-06-10T18:14:16] Completed: Jun-10-2019,18:14:16 [2019-06-10T18:14:16] Seconds: 374.766 [2019-06-10T18:14:16] Items: 200001 [2019-06-10T18:14:16] Directories: 1 [2019-06-10T18:14:16] Files: 200000 [2019-06-10T18:14:16] Links: 0 [2019-06-10T18:14:16] Data: 24.414 TB (26843545600000 bytes) [2019-06-10T18:14:16] Rate: 66.708 GB/s (26843545600000 bytes in 374.766 seconds)



Result: dcp is slightly faster than cp



dcp is ~140x faster





mpiFileUtils enables users to manage datasets with same resources they used to create them

- Large HPC jobs generate datasets while using thousands of compute nodes running tens of thousands of processes
- The problem:
 - Users must often manage those datasets using singlenode, single-process tools like cp, chgrp, and rm
 - Users create a dataset with 10,000 cores but then copy it with one.
- mpiFileUtils is designed to:
 - Let users perform common data management tasks with the same HPC resources they used to run their jobs
 - Scale to saturate available compute, network, and file system bandwidth
 - Operate across common HPC file systems like Lustre, GPFS, and NFS





Project origin: mpiFileUtils was created by extending existing tools - libcircle and dcp



- libcircle
 - Each process pulls work items from its local queue.
 - A process can add new work items to its local queue.
 - If a process runs out of work, it requests work from a random process.
 - "On distributed file tree walk of parallel file systems", Jharrod LaFon, Satyajayant Misra, and Jon Bringhurst, SC'12.
 - <u>http://dl.acm.org/citation.cfm?id=2389114</u>
 - <u>https://github.com/hpc/libcircle</u>
- dcp (original)
 - Built all copy functions on libcircle
 - walk, file create, data copy, metadata update
 - <u>https://github.com/hpc/dcp</u>



mpiFileUtils = Library + Tools + File Format



- Library common data structures and routines
 - Quickly create new tools
 - Apps can even invoke API directly



- Tools MPI-based versions of cp, rm, etc.
 - Apps create datasets with thousands of processes.
 - Why manage them using a single process?



- File format common format for interoperability
 - Allows one to compose tools into pipelines
 - Allows third-party software to generate input or process output (e.g., Hopper)



The file list (mfu_flist) is the primary data structure within the libmfu common library

The mfu_flist is a distributed list of stat-like info for each item





Collective mfu_flist operations operate across all processes, for example, to do a global sort

Sort items alphabetically by their full path





mpiFileUtils has a growing set of production and experimental tools (v0.11); 'd' is for distributed

- dbcast broadcast file to compute nodes
- **dbz2** compress/decompress file with bz2
- dchmod change perms/owner/group
- dcmp compare directories/files
- dcp copy data
- ddup find duplicate files
- **dfilemaker** generate test files
- **dfind** filter files
- **dreln** update symlinks

- drm delete files
- **dstripe** restripe files (Lustre)
- dsync synchronize directory trees
- dtar create / unpack tar files
- dwalk list, sort, summarize files
- dgrep parallel grep
- dparallel MPI-based parallel
- dsh interactively list, summarize, and remove files





The common file format lets one compose tools, e.g., purge all files last accessed over 180 days ago

```
# walk directory to stat all files, record list
in file
```

```
dwalk --output list.mfu /path/to/walk
```

filter list to identify all regular files
that were last accessed over 180 days ago

```
dfind \
  --input list.mfu \
  --type f --atime +180 \
  --output purgelist.mfu
```

```
# delete all files in the purge list
```

```
drm --input purgelist.mfu
```



Performance at scale





Sierra Scaling Study: dcp, dcmp, dchmod, drm

- Given our example dataset:
 - Single directory
 - 200,000 files
 - 128 MB per file
 - 24.4 TB total
- Do the following:
 - 1. dcp: Make a copy of the dataset
 - 2. dchmod: Change group on all files in the copy
 - 3. drm: Remove the copy
- Scale from 1 node (40 procs) up to 1k nodes (40k procs)





dcp <u>walks</u> source directory, <u>creates</u> destination files, <u>copies</u> data, then <u>updates</u> file metadata







The dcp write bandwidth increases with node count and holds steady at 320 GB/s on Sierra



Once the bandwidth is saturated, there is no value to scale to higher node counts.



dchmod changes group on files after the copy, a test that stresses file system metadata ops



Sierra GPFS, IBM Spectrum MPI, jsrun

Quartz Lustre, MVAPICH2 MPI, SLURM

For quick jobs, the MPI job launch time can be significant. Run those with smaller node counts.



drm removes the copy after the test



Sierra GPFS, IBM Spectrum MPI, jsrun

Quartz Lustre, MVAPICH2 MPI, SLURM

An example of bad scaling. It is unknown whether the cause in this case is mpiFileUtils or thrashing within the parallel file system.







Things to keep in mind since mpiFileUtils are MPI programs

- Typically need to run in a job allocation
 - The sweet spot for most tools is about 2-4 nodes.
 - Grab more nodes for large datasets.
- Launch your job with mpirun
 - Plan to max out the CPU cores.
 - Leave a few cores idle on each node for the file system client processes.
- Most tools do not checkpoint their progress
 - Be sure to request sufficient time in your allocation.
 - You may need to start over from the beginning if a tool is interrupted.
- Cannot pipe output of one tool to the input of another
 - The --input and --output file options are good approximations.
- Cannot easily check the return codes of tools
 - Check the output for errors.



mpiFileUtils are designed to be portable across all HPC sites and file systems

- Designed to work on any HPC system
 - Built with basic C, MPI, and POSIX/libc, so uses programming language and programming models that are ubiquitous in HPC

- Available in Spack

spack install mpifileutils

- Designed to work on all HPC file systems
 - Lustre, GPFS, NFS have been targets from start
 - Intel actively working to add DAOS
 - Default algorithms should work with any POSIX-compliant file system
 - One may include file-system specific optimizations and algorithms



Take away: For most of your work, use standard tools like cp. Too slow? Then try mpiFileUtils.

- Resources:
 - Documentation: <u>https://mpifileutils.readthedocs.io</u>
 - Code: <u>https://github.com/hpc/mpifileutils</u>
 - New users and new collaborators are always welcome!
- From its beginning, mpiFileUtils has been a multi-organizational open-source effort
 - Numerous people have contributed over time
 - <u>https://github.com/hpc/mpifileutils/blob/master/AUTHORS</u>
 - <u>https://github.com/hpc/mpifileutils/graphs/contributors</u>
 - Particularly: Jon Bringhurst, Jharrod LaFon, Danielle Sikich, Dalton Bohning, Elsa Gonsiorowski, Feiyi Wang, Xi Li, and Zheng Gu



Thank you MVAPICH team!

- The MPI programming model has enabled HPC users to continually scale up their problems for decades ...
- ... but only because of the hard work done by the people who implement MPI, and in particular due to the leading R&D from the MVAPICH team.

MPI = scalability.h MVAPICH = scalability.c









Frequently used mfu_flist functions. Some are collectives, some are local to the calling process.

- walk recursively walk one or more paths to create a list
- stat given a list, execute stat on each item
- filter apply find-like tests to find a matching subset of items
- depth split list into sublists dividing items by depth
- remap/spread reassign elements to different ranks
- sort globally order elements across ranks, based on item properties (path, name, size, user, access time)
- copy copy items in source list to a destination on the file system

- mkdir/mknod create directories or inodes on file system
- unlink remove all items named in a list from the file system
- read/write load list from a file, write list to a file
- pack/unpack serialize an element for network transfer
- query properties of global list, such as global list size and offset of local rank
- iterate over local elements
- query/set properties of a local element



An example tool: walk a path, write out list of all regular files having more than one hardlink

```
mfu flist flist = mfu flist new();
                                                            walk path(s) to get
mfu_flist_walk param paths(numpaths, paths,
                                                            initial list of items
    walk opts, flist);
                                                            create subset list
mfu flist flist links = mfu flist subset(flist);
uint64 t size = mfu flist size(flist);
                                                            iterate over all local
for (uint64 t idx = 0; idx < size; idx++) {
                                                            items in initial list
    mfu filetype type = mfu flist file get type(flist, idx);
    if (type != MFU TYPE FILE)
                                                            check whether item is a
        continue:
                                                            regular file
    const char* file = mfu flist file get name(flist, idx);
    mfu lstat(file, &statbuf);
                                                            copy item to subset if
    if (statbuf.st nlink > 1)
                                                            hardlink count is more
        mfu flist file copy(flist, idx, flist links);
                                                            than one
mfu flist summarize(flist links);
                                                            finalize subset list
mfu flist write cache (outputname, flist links);
                                                            write subset list out
                                                            to a file
mfu flist free(&flist links);
mfu flist free(&flist);
                                                            free lists
```



Same tool in six lines of python

```
import os
import stat
import mpifileutils as mfu
                                     import mpifileutils
flist = mfu.FList("/path/to/walk") walk target path
hardlinks = flist.subset(lambda f: sublist of
  f.type == mfu.TYPE FILE and
                                      reg files with
 os.stat(f.name)[stat.ST NLINK] > 1) hardlink > 1
```

hardlinks.write("hardlinks.mfu") write sublist to file





dcmp checks the new copy with the original, reads back and compares all bytes



It is recommended to run dcmp after dcp to verify contents.



dcmp saturates read bandwidth and holds steady at 1 TB/s on Sierra



dcmp is read heavy and can often reach peak file system bandwidth.





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