GPFS at EBI

Facing performance degradation when using mmap based applications

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1. Who are EBI?

Europe's home for biological data, research and training

- A trusted data provider for life sciences.
- Part of the European Molecular Biology Laboratory (EMBL), an intergovernmental research organisation.
- International: 23 member states and 2 associate member states, 6 sites in Europe.
- EMBL-EBI manages a vast amount of public biological data, delivering it to the global life science community on demand, 24/7 without restriction or charge.



2. Systems at EBI

IT services at EBI are managed by the Technical Services Cluster (TSC)





3. Infrastructure

- Three datacentres with around 200 racks of equipment.
- More than 40.000 cores. Mostly ethernet based.
 - Multiple HPC-like clusters (mainly blades).
 - Hadoop cluster.
- Over 140 PB of raw storage capacity.
 - Multiple technologies: NFS, Object Store, GPFS, Lustre
 - Annual storage growth of 40-50%



4. Scenario for GPFS: EBI services





5. Scenario for GPFS: infrastructure





6. The problem: BLAST is slow

- BLAST (Basic Local Alignment Search Tools) is a software used to find regions of similarity between biological sequences.
- Developed by the NCBI and very popular in bioinformatics.
- Users reported running BLAST was extremely slow when source data was stored in GPFS.
- Some executions showed a performance 80 times slower compared to NFS.



6. The problem: BLAST is slow

BLAST RUNTIME AVG.





We tried multiples approaches without success.

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- **x** Update the GPFS client to the latest available version (5.0.0.1).
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8. The real evil: mmap

BLAST switches to using mmap for IO when it's executed with more than 1 thread.

- Performance degradation only occurs when running multiple threads in the same host.
- Analizing the execution of BLAST using strace we realized there were thousands of mmap calls.
- Many users reporting problems with parallel filesystems when using mmap, including GPFS.
- mmap causes IO to fallback to the default kernel page size (4K) instead of using the values defined for GPFS.



8. The real evil: mmap

BLAST switches to using mmap for IO when it's executed with more than 1 thread.

=== mmdiag: iohist ===

I/O history:

I/O start time	RW	Buf type o	disk:sectorNum	nSec	time ms	Туре	Device/NSD ID	NSD node
16:34:53.284905	R	data	2:338059951504	8	17.235	cli	AC110407:586CEE7B	10.7.74.15
16:34:53.302271	R	data	2:338059912744	8	4.659	cli	AC110407:586CEE7B	10.7.74.15
16:34:53.307139	R	data	1:338060022688	8	1.374	cli	AC110406:586CEE77	10.7.74.14

=== mmdiag: iohist ===

I/O history:

I/O start time	RW But	E type	disk:sectorNum	nSec	time ms	Туре	Device/NSD ID	NSD node
16:24:17.783564	R	data	2:214062309376	4096	22.108	cli	AC110407:586CEE7B	10.7.74.15
16:24:17.806245	R	data	1:214062120960	4096	23.263	cli	AC110406:586CEE77	10.7.74.14
16:24:17.830125	R	data	1:214062116864	4096	18.072	cli	AC110406:586CEE77	10.7.74.14



8. The real evil: mmap

BLAST switches to using mmap for IO when it's executed with more than 1 thread.

File objtools/blast/seqdb_reader/seqdbimpl.hpp

/// Set number of threads /// /// Set number of threads which will share this object. This /// permits use of an internal mmap for the threads. /// If the second parameter is 'false' (the default), /// the internal mmap is not used if num threads == 1. /// For certain applications where there are multiple CSegDB /// objects, each one accessed by only a single thread, /// setting num threads to 1 (thread per CSegDB) results in /// a performance hit by not using the mmap. /// In this case, force mt ("force multithreading") should /// be set to 'true' to allow use of the mmap when num threads /// == 1. For num threads > 1, force mt has no effect. /// /// @param num threads The number of threads which will share /// access to this CSeqDB object. [in] /// @param force mt Defaults to false, setting to true when /// num threads == 1 forces multithread /// internal mmap. [in]

void SetNumberOfThreads(int num_threads, bool force_mt = false);



9. Next step: help IBM!

No workaround could be provided:

- Running BLAST with 1 thread takes too much time.
- Switching to an MPI solution was too complex.

So we looked for help:

- Asked on the GPFS Users Group list
- Contacted with Sven Oehme
- Sven agreed on working on a fix for the problem





