Data Analysis and Visualization of "Big Data" from HPC

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Dave Pugmire (ORNL), George Ostrouchov (ORNL), Scott Klasky (ORNL) OLCF Workshop on Processing and Analysis of Very Large Data Sets 7 August 2013



What is "Big Data"?

• Big data is...

Extracting meaning from large sets of digital information

- "Large": Beyond the capabilities of "ordinary" techniques
- HPC has been doing this for decades:
 - Using datasets from large-scale simulation of physical phenomena



Data may be the "fourth pillar" of science





FOURTH PARADIGM

DATA-INTENSIVE SCIENTIFIC DISCOVERY

CHIED BY TONY HEY, STEWART TANSLEY, AND KRISTIN TOLLE



So what's all the hype about?

- Much of the hype of the last two years has been about extracting meaning...
 - ...from large unstructured and semi-structured datasets
 - ...from the Business Intelligence community
 - ...using scalable analysis and storage tools like: Hadoop, Cassandra, HBase, Hive, Pig, MongoDB, ...
- Some successes in scientific computing for unstructured data



Petascale machines are solving important scientific problems through the generation and analysis of large datasets



Physics of high-temperature superconducting cuprates



Chemical structure of HIV viral envelope



Global climate simulation of CO₂ dynamics and biochemistry



Biological conversion of cellulosic feedstock for biofuels



Three-dimensional simulation of blood flow



Combustion of turbulent lean fuel



Current success comes from having tools that scale to the HPC datasets

- Analysis methods are generally the same as in serial
- It's the tools and techniques that must scale
- SPMD data parallelism has taken us far -> Petascale+



Pipeline visualization



A lot of success has been through data flow networks (pipelines)

- Different input data formats:
 - NetCDF, HDF, text, CSV, PDB, ADIOS, ...
- Different types of data operations:
 - Slicing, resampling, mesh transforms, filtering, ...
- Different ways of drawing on-screen:
 - Pseudocolor, isosurfaces, volume rendering, ...

These are independent of each other



Make these independent modules

- Datareading
- Data operations
- Data plotting



This is a data flow network



Many software systems use data flow networks

• VTK



- Vislt, ParaView
- AVS/Express



- SCIRun
- COVISE





We've scaled to trillions of mesh cells

• Weak scaling study 2009 (isocontouring, volume rendering): ~63 million cells/core

Machine	Туре	Problem Size	# cores		
Jaguar	Cray XT5	2T	32k		
Franklin	Cray XT4	1T, 2T	16k, 32k		
Dawn	BG/P	4T	64k		
Juno	Linux	1T	16k, 32k		
Ranger Sun Linux		1T	16k		
Purple AIX 0.5T 8k					
Since this work, people have					

surpassed 8 trillion cells = $20,000^3$ cells.



2T cells, 32K procs



The "Three V's" of big data as applied to HPC



Validity,



The "Three V's" of big data as applied to HPC

- Volume:
 - Increasing mesh resolution
 - Increasing temporal complexity
- Variety:
 - Higher-dimensional data
 - Increasing multi-variance
 - Complex data models
 - Ensembles, parameter studies
- Velocity:
 - Future hardware constraints will limit I/O





Data scale limits scientific understanding

- Spatial resolution increasing in many domains
 - Too many zones to see on screen (with or without high-res Powerwall)
- Temporal complexity increasing
 - Climate simulations will have 100,000s of time steps
 - Manually finding temporal patterns is tedious and error-prone
- Multivariate overload
 - Climate simulations have 200-400 variables
 - Growing to thousands
- Issues of data models and domain-specific data
 - E.g., Multi-group radiation fields



Common Themes in Mathematics for Data

Computation Mode	Memory Architecture	Data Partition Availability	Data Passes
Serial or Multithreaded	contiguous	N/A (all)	many
Distributed	partitioned	all	many
Out-of-Core	partitioned	one-at-a-time	one or few
Streaming/ in situ	partitioned	one-at-a-time	one
Mathematics needed	partition coupling math	updating via partition coupling math	one-pass updating via partition coupling math

Minimizing data access (energy) pushes everyone toward one-pass (one-touch) updating.





Prediction of Ice and Climate Evolution at Extreme Scales (PISCEES)

- SciDAC Project that is:
 - Developing robust, accurate, and scalable dynamical cores for ice sheet modeling on a variety of mesh types
 - Evaluate ice sheet models using new tools for V&V and UQ
 - Integrate these models and tools into the Community Ice Sheet Model (CISM) and Community Earth System Model (CESM)
 - Simulate decade-to-century-scale evolution of the Greenland and Antarctic ice sheets





PI: William Lipscomb, Philip Jones (Acting)

Data set structure

- Structured NetCDF files
 - Ice defined by two variables: topg, thk
 - Ice flow sheets defined by N levels



land height



Flow Analysis in PISCEES





We have a scalable integral curve algorithm Parallelize over blocks Use a combination of seed and block parallelization Parallelize Dynamically steer computation over seeds Allocate resources where needed Load blocks from disk on demand Parallelize over seeds and blocks Goals: 1. Maximize processor utilization 2. Minimizes IO 3. Minimizes communication Works adequately in all use cases D. Pugmire, H. Childs, C. Garth, S. Ahern, G. Weber, "Scalable Computation of Streamlines on Very Large Datasets," Proceedings of Supercomputing 2009, Portland, OR, November, 2009

Parallel Integral Curve System (PICS)



PICS Filter Design



"Distance Traveled" operator





Results



1 year of travel

10 years of travel

Steady state flow distance traveled





Big Data Exploration Requires Flexibility in Analytics

- Analytics for not-so-big data:
 - Science produces great many algorithms
 - Often redundant and related
 - Statistics and mathematics prune, generalize, and connect
 - Still we have hundreds of statistics and mathematics books on data analytics



No reason to expect big data less needy than not-so-big data <u>Flexibility and a big toolbox</u> are critical for exploration <u>Need for analytics diversity</u> persists for big data



Programming with Big Data in R: pbdR

- R has unmatched diversity and flexibility for data
- Goals:
 - Shorten time from big data to science insight
 - Productivity, Portability, Performance
- Approach:
 - Implicit management of distributed data details
 - Scalable, big data analytics with high-level syntax
 - Identical syntax to serial R
 - Powered by state of the art scalable libraries
 - Free* R packages

*MPL, GPL and BSD licensed

pbdR Core TeamGeorge Ostrouchov^{1,2}, Team Lead Wei-Chen Chen¹Pragnesh Patel²Drew Schmidt²

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High-Level Syntax

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	Cov.X <- cov(X)	Cov.X <- cov(X)
Developer	N <- nrow(X)mu <- colSums(X) / NX <- sweep(X, STATS=mu, MARGIN=2)Cov.X <- crossprod(X.spmd) / (N-1)	N <- allreduce(nrow(X))mu <- allreduce(colSums(X) / N)X <- sweep(X, STATS=mu, MARGIN=2)Cov.X <- allreduce(crossprod(X)) / (N-1)

	Lm.X <- lm.fit(X, Y)	Lm.X <- lm.fit(X, Y)
Developer	tX <- t(X)A <- tX %*% XB <- tX %*% Yols <- solve(A) %*% B	tX <- t(X)A <- <mark>allreduce(</mark> tX %*% X <mark>)</mark> B <- <mark>allreduce(</mark> tX %*% Y) ols <- solve(A) %*% B



A High-Level Language for Big Data Analytics



Promising Scalability



Promising Scalability



Nautilus



Performance is highly dependent upon I/O rates

• Weak scaling study 2009 (isocontouring, volume rendering): ~63 million cells/core







Appx I/O time: 2-5 minutesAppx processing time: 10 seconds

Hardware technology curves at the exascale: ~2018

- Billion-way concurrency
- Constrained memory environment
- Very constrained I/O (in relation to compute capability)

NSF CIF21: "Computing power, data volumes, software, and network capacities are all on exponential growth paths."

	1		107	
		"2018"		
System Parameter	2011	Swim Lane 1	Swim Lane 2	Factor Change
System Peak	2 Pf/s	1 E	f/s	500
Power	6 MW	≤ 20	MW	3
System Memory	0.3 PB	32-6	4 PB	100,200
Total Concurrency	225K	$1B \times 10$	1B×100	40,000-400,000
Node Performance	125 GF	1 TF	10 TF	0 00
Node Concurrency	12	1,000	10,000	83-830
Network BW	1.5 GB/s	100 GB/s	1000 GB/s	66-660
System Size (nodes)	18700	1,000,000	100,000	50 500
I/O Capacity	15 PB	300-10	00 PB	20-67
I/O BW	0.2 TB/s	20-60	TB/s	10-30



Embedded Visualization Computation 8 EB/s

Node Memory 400 PB/s

Co-Scheduled Visualization

Interconnect (10% Staging Nodes) 10 PB/s

Off-Line Visualization 60 TB/s

Implications for analysis & visualization

- Extremely limited ability to move data off HPC resource
 - The de-coupling of simulation codes and visualization codes can no longer continue
- Extremely limited memory space
- Extreme concurrency will break most communication models
 - Pure MPI already breaking at the petascale
- Fine degree of data parallelism will break most data processing models



Implications for visualization & analysis

Predicted Exascale Machines			
Node Concurrency	1,000 - 10,000		
Number of Nodes	1,000,000 - 100,000		
Total Concurrency	1 billion		

- Massive concurrency across nodes
 - New parallel methodologies avoid scaling limits
 - e.g., hybrid and multi-level parallelism
 - In situ analysis codes must be commensurate with simulation codes' parallelism
- Massive concurrency within nodes
 - Thread- and data-level parallelism
 - Accelerators/GPUs today also have discrete memory
 - Current vis GPU use generally limited to rendering



So what do we do?

- We've had a lot of success moving processing closer and closer to the data.
- Medium-term goal: Take the final step in moving processing closer to the data
 - Right to the source, in memory where it's generated
 - Reduce data every step of the way to the user
 - All data visualization (and analysis) becomes remote
- Long-term goal: Move data processing everywhere the data lives



Injection of analysis into I/O pipelines

- Leverages existing code methods for data movement
 - Lots of compute available
 - Exploits high-capacity HPC interconnect
- Possible to annotate, reduce, and reorder data in flight
- I/O is the primary bottleneck, so it's the logical place to begin adaptation
- Challenges:
 - Memory-constrained
 - Impossible to do serendipitous discovery with in situ alone
 - Temporal analysis can be difficult


Goal: Tightly-coupled analysis infrastructures



Simulation, analysis, visualization workflow

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ADIOS: ADaptable I/O System

- An I/O abstraction framework
- Provides portable, fast, scalable, easy-to-use, metadata rich output with a simple API
- Layered software architecture
- Change I/O method on-the-fly
- Abstracts the API from the method used for I/O
- Provides method for injection of analysis and visualization







ADIOS: I/O problems and their solutions

- I/O buffering and aggregation:
 - Buffering utilizes max I/O bandwidth at writing
 - Aggregation utilizes max I/O bandwidth at reading
- Data layout optimization:
 - Layout optimization improves read performance
- Reduce synchronization and contention:
 - Log-file format and aggregation decreases lock contention and decrease network movement
 - Scheduled staging decreases impact of asynchronous I/O







Success across many application domains

- Fusion:
 - Using staging towards predictive capabilities
 - Coupled with ParaView, Vislt, and dashboards
- Combustion:
 - Orders of magnitude speed improvement for checkpoint/restart
 - In situ feature detection
- More (astro, AMR, geophysics, ...)





At the same time: Exploit fine-grained parallelism

- Memory space per core may be down in the megabytes to 100s of kilobyte range
 - (Though HMC's may alleviate the memory wall and capacity.)
- Data parallelism becomes much more complex at this fine level of detail:
 - How to create and exploit search data structures
 - How to harness communication hierarchies
 - How to handle "ghost" region between data blocks
 - Iterative algorithms (e.g., streamlines) become extremely inefficient
- No analysis or visualization algorithms are written this way. **Everything must be rewritten**.



Extreme-scale Analysis and Visualization Library: EAVL

- Update traditional data model to handle modern simulation codes and a wider range of data. Extend current analysis and vis tools to data that cannot be approached today.
- Investigate how an updated data and execution model can achieve the necessary computational, I/O, and memory efficiency.
- Explore methods for visualization algorithm developers to achieve these efficiency gains and better support exascale architectures. Allow easy gradual adoption into existing analysis/vis tools.





Promising results to date

- Data thresholding is more memory- and computation-efficient.
- Coordinate space transformation moves from O(n) to O(1).
- Structured grids are an order of magnitude more space efficient.
- Efficient execution on heterogenous architectures
- Temporal parallelism provides scalability benefits
- In-place processing provides significant memory benefits





HPC data is becoming more complex

- Workflow tools are becoming more common:
 - Pegasus, Swift, HTCondor, Kepler, Eden, ...
- Ensembles are growing larger
- Parameter studies and Design of Experiments are becoming much more common
- Tracking *provenance* is important
- There are no general tools for processing large runs of simulations

The concept of a "data set" is evolving ...becoming more unstructured.



Turbo Compressor Innovation Breakthrough aerodynamic design optimization

INCITE Project Allan Grosvenor, Ramgen Power Systems Allocated hours: 40M Used hours: 36M

Science Objectives and Impact

- Ramgen Power Systems is developing shock wave compression turbo machinery to meet DOE goals for reducing Carbon Capture and Sequestration (CCS) costs
- Complementary goal: design a gas turbine with dramatically lower costs and higher efficiency
- Compressing CO₂ to the required 100 atmospheres represents approximately 33 percent of the total cost of Sequestration

OLCF Contribution

- 50x improvement in code scalability with more efficient memory utilization
- Accelerated I/O by 10x with optimizations and ADIOS

 Intelligent use of ensembles to explore parameter space using 240,000 cores

Results

- Transformed Ramgen's aerodynamic design process
- Observed designs with valuable new characteristics, from ensembles not possible without Jaguar
- Created a new workflow paradium that accelerate
- Created a new workflow paradigm that accelerates design of compressors
- Accelerated computational design cycle for turbo machinery from months to 8 hours!





Leadership-scale ensemble runs on Jaguar support intelligently-driven design optimization





Nuclear Physics The Limits of the Nuclear Landscape

Science Objectives and Impact

- Predict how many protons and neutrons can be bound within a nucleus.
- Identify the nuclear drip lines that denote the limits of nuclear binding.
- Project advances toward the vision of "designer nuclei," with uses ranging from potential cancer treatments to a better understanding of superconductivity.

OLCF Contribution

 Custom code (co-authored by OLCF staff member Hai Ah Nam) used density functional theory to solve the nuclear structure, which involved a large algebraic non-linear eigenvalue problem.

• Each ensemble of nuclei took about two hours to calculate on 224,256-processors in Jaguar system.

Each run evaluated about 250,000 possible nuclear configurations.

Compute Usage 70% 35% 0% -20% 20-60% >60% INCITE Project James Vary, Iowa State University Allocated hours: 37M Used hours: 60M



Map of bound even-even nuclei as a function of \angle and *N*. Shown are stable nuclei (black), radioactive nuclei (green), mean drip lines with uncertainties (red), and two-neutron separation line (blue). The inset shows the irregular behavior of the two-neutron drip line around Z = 100.

Science Results

- Accurate calculation of the number of bound nuclei in nature.
- Calculations predicted about 7,000 possible combinations of protons and neutrons allowed in bound nuclei with up to 120 protons.
- Several leading models of nuclear interaction shown to be largely in agreement.

J. Erier, et al., "The Limits of the Nuclear Landscape" Nature, June 2012.



Sea-Level Rise Inevitable

Aggressive Greenhouse Gas Mitigation Can Help Slow Rate

Science Objectives and Impact

- Determine how much sea level will rise based on three climate-change mitigation scenarios
- Examine mechanisms involved in sealevel rise, including ice melt and thermal expansion of water
- Quantify sea-level rise in response to mitigation strategies



Sea level will continue to rise due to thermal expansion into 2300 under the mos aggressive mitigation scenario, (cooling), but the rise will be slowed enough to implement adaptation measures. With less aggressive mitigation (stable) and (warming), there would be less time for adaptation.

OLCF Contribution

• Used CCSM4 to conduct simulation ensembles for each mitigation scenario

 Simulation of 15 separate solutions, 4,500 total simulation years

Science Results

- Sea level will continue to rise in the future, even with aggressive CO₂ mitigation scenarios.
- Aggressive mitigation measures strongly affect the rate of increase
- Mitigation buys important time to implement adaptation measures for inevitable sea-level rise

G.A. Meehl, et al., "Relative outcomes of climate change mitigation...," Nature Climate Change, August 2012.



Computational and Molecular Biophysics High-Performance Computing for Accelerated Drug Discovery

Discretionary Project Jerome Baudry, UTK/ORNL Allocated hours: 7M Used hours: 8M

Science Objectives and Impact

- Test millions of compounds against a specific protein receptor using simulations.
- Find the best match between a compound and its targeted protein receptor, in hopes of creating drugs with a higher degree of specificity and less cross-reactivity, as well as exploring alternative uses for existing drugs.
- Impact: Drastically decrease the time and money spent bringing new, improved drugs to market.



Computational approaches are used to describe how molecular compounds of a drug candidate (displayed in colored spheres) bind to its specific protein receptor target. Image courtesy of Jerome Baudry, UTK & ORNL

OLCF Contribution

Director's Discretion Project

Usage between 1/1/2012–9/30/2012 Used 7.6 M hours of allocation, or 110%

Used variations of AutoDock, a General Public License, open-source software, commonly used for molecular 3D modeling in protein-docking simulations.

Science Results

- Successfully screened 2 million compounds against a targeted receptor in a matter of days, as opposed to months using computing clusters or longer with test-tube methods.
- Allowed scientists to account for specific binding in protein receptors as well as structural variations within the receptor.
- Enabled search of a vast library of molecular compounds to find alternative uses for existing drugs (i.e. "repurposing").

S. Ellingson. "Accelerating virtual high-throughput ligand docking," Presented at HPDC12, 2012.



Southern California Earthquake Center (SCEC)

CyberShake workflow

A single large HPC simulation leads to *millions* of serial analysis jobs



What does the future hold for big data from HPC?

- Evolving nature of data sets means new research and development directions. We have no general tools to handle these!
- Workflow tools, ensemble processing, and parameter studies ->
 - Need to store "meta" information about simulation results.
 - Promising: Database storage and query methods. SciDB?
- Scalability limitations of traditional files ->
 - Need to store scientific information with new methods.
 - Promising: Object stores and access methods
- Fusion of HPC data and observational data ->
 - Need for algorithms and methods to expand into new data types.
- New data types and descriptions ->
 - Requirement for processing beyond simply distributed-parallel methods
 - Promising: "business analytics" methods for unstructured data



We've got a lot of work to do!

Thanks for your time!

