

Portable and Productive Performance on Hybrid Systems with libsci_acc

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What is Cray Libsci_acc?

- CRAY
- Provide basic scientific libraries optimized for hybrid systems
 - Incorporate the existing GPU libraries into Cray libsci
- Independent to, but fully compatible with OpenACC
- Multiple use case support
 - Get the base use of accelerators with no code change
 - Get extreme performance of GPU with or without code change
- Provide additional performance and usability
- Two interfaces
 - Simple interface
 - Auto-adaptation
 - Base performance of GPU with minimal (or no) code change
 - Target for anybody: non-GPU users and non-GPU expert
 - Expert interface
 - Advanced performance of the GPU with controls for data movement
 - Target for CUDA, OpenACC, and GPU experts
 - Does not imply that the expert interfaces are always needed to get great performance

Why libsci_acc?



- Code modification is required to use existing GPU libraries!
- Several scientific library packages already exist
 - CUBLAS, CUFFT, CUSPARSE (NVIDIA), MAGMA (U Tennessee), CULA (EM Photonics)
- No Compatibility to Legacy APIs
 - cublasDgemm(....)
 - magma_dgetrf(...)
 - culaDgetrf(...)
 - Why not dgemm(), dgetrf()?
- Not focused on Fortran API (C/C++)
 - Require CUDA data types, primitives and functions in order to call them
- Performance

Auto-tuning



- Cray Autotuning framework has been built to tune BLAS for accelerators
 - GPU kernel codes are built using code generator
 - Enormous offline auto-tuning is used to build a map of performance to input
 - An adaptive library is built from the results of the auto-tuning
 - At run-time, your code is mapped to training set of input
 - Best kernel for your problem is used

Simple Interface

- CRAY
- Supports the standard API in the original form
- Will perform all GPU dirty-work for you
 - Initialize data structures on GPU
 - Split your problem into a CPU portion and GPU portion
 - Copy data to the GPU memory from CPU memory
 - Perform GPU and CPU operations
 - Copy data back to CPU memory
- Library-heavy codes can use GPUs with no code change
- Is not only a tool for simple usage
 - If you don't need the data on the GPU afterwards, use the simple interface
- Simple API has automatic adaptation

Adaptation in the Simple Interface



 You can pass either host pointers or device pointers with the simple interface

A is in host memory

```
dgetrf(M, N,(A,)lda, ipiv, &info)
```

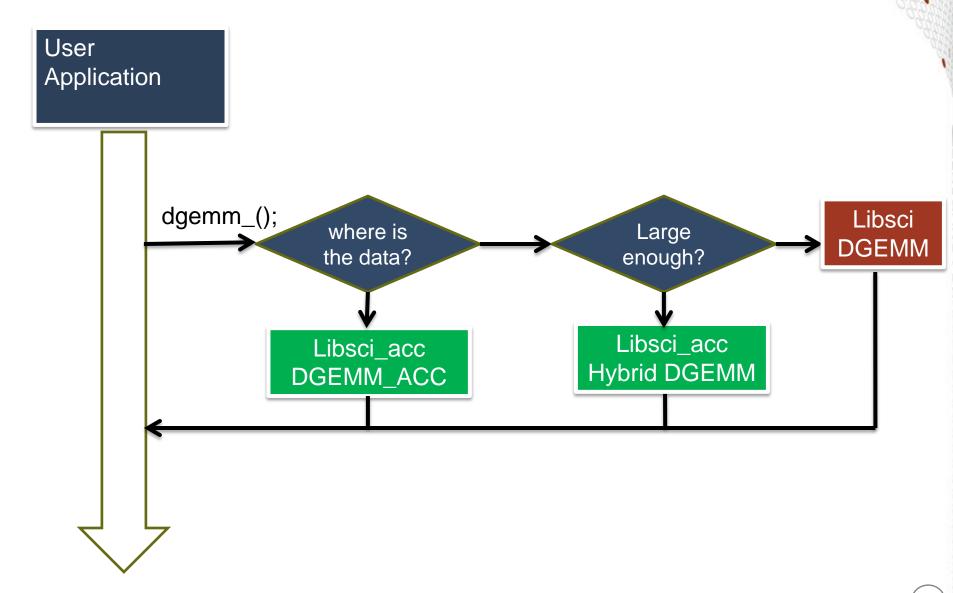
- Performs hybrid operation on GPU
- if problem is too small, performs host operation
- Pass Device memory

```
dgetrf(M, N, (d_A, lda, ipiv, &info)
```

- Performs hybrid operation on GPU
- BLAS 1 and 2 performs computation local to the data location
 - CPU-GPU data transfer is too expensive to exploit hybrid execution

Libsci_acc: Simple Interface for BLAS3 and LAPACK





Expert Device & CPU Interface



- Device interface gives higher degrees of control
- Allow users to explicitly specify the execution
 - Every routine in libsci has a version with _acc and _cpu suffix
 - e.g. dgetrf_acc, dgetrf_cpu
 - Simple API for device memory and _acc API are the same

Usage - Basics

CRAY

- Supports Cray and GNU compilers.
- Fortran and C interfaces (column-major assumed)
 - Load the module craype-accel-nvidia35.
 - Compile as normal (dynamic libraries used)
- To enable threading in the CPU library, set OMP_NUM_THREADS
 - E.g. export OMP_NUM_THREADS=16
- Assign 1 single MPI process per node
 - Multiple processes cannot share the single GPU
- Execute your code as normal

Libsci_acc with OpenACC



- If the code uses
 OpenACC, it's
 possible to use the
 library with directives
- All data management performed by OpenACC
- Calls the device version of dgemm
- All data is in CPU memory before and after data region

```
!$acc data copy(a,b,c)
!$acc parallel
!Do Something
!$acc end parallel
!$acc host data use device(a,b,c)
call dgemm acc('n','n',m,n,k,&
               alpha, a, lda, &
               b,ldb,beta,c,ldc)
!$acc end host data
!$acc end data
```

Libsci_acc with OpenACC



- Libsci_acc is a bit smarter that this
- Since 'a,' 'b', and 'c' are device arrays, the library knows it should run on the device
- So just dgemm is sufficient

```
!$acc data copy(a,b,c)
!$acc parallel
!Do Something
!$acc end parallel
!$acc host data use device(a,b,c)
call dgemm
               ('n','n',m,n,k,&
                alpha, a, lda, &
               b,ldb,beta,c,ldc)
!$acc end host data
!$acc end data
```

libsci_acc BLAS Routines Available



- BLAS 3 Full HYBRID Implementations
 - [s,d,c,z]GEMM
 - [s,d,c,z]GEMM
 - [s,d,c,z]TRSM
 - [z,c]HEMM
 - [s,d,c,z]SYMM
 - [s,d,c,z]SYRK
 - [z,d]HERK
 - [s,d,c,z]SYR2K
 - [s,d,c,z]TRMM
- The following are supported without HYBRID implementations because there is no performance advantage
 - All BLAS 2 Routines
 - All BLAS 1 Routines

libsci_acc LAPACK Routines Available



Full HYBRID Implementations:

- [d,z]GETRF (LU Factorization)
- [d,z]POTRF (Cholesky Factorization)
- [d,z]GETRS (System Solver)
- [d,z]POTRS (System Solver)
- [d,z]GESDD* (Generalized Singular Values)
- [d,z]GEBRD (Generalized Bidiagonalization)
- [d,z]GEQRF* (QR Factorization)
- [d,z]GELQF (LQ Factorization
- [d,z]GEEV (Non-symmetric Eigenvalues)
- DSYEVR* / ZHEEVR* (Hermitian/Symmetric Eigenvalues)
- DSYEV / DSYEVD (Hermitian/Symmetric Eigenvalues)
- ZHEEV / ZHEEVD (Hermitian/Symmetric Eigenvalues)
- DSYGVD / ZHEGVD (Hermitian/Symmetric Eigenvalue System Solver)

^{*} Include Cray Proprietary Optimizations

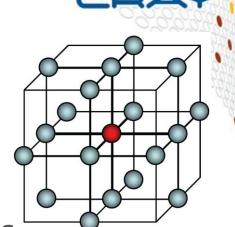
Summary



- Access to libsci_acc routines is simple
 - No need to explicitly link Programming Environment drivers (cc, ftn, CC) do this for you
 - Just target the GPU by loading module
- Can automatically take advantage of threading on CPU
 - Just set OMP_NUM_THREADS and run
- Simple interface available to enable hybrid, CPU or GPU execution of a routine depending on where memory pointers reside and problem size
- Interface for advanced control is also available

Case Study: the Himeno Benchmark

- Parallel 3D Poisson equation solver
 - Iterative loop evaluating 19-point stencil
 - Memory intensive, memory bandwidth bound
- Fortran, C, MPI and OpenMP implementations available from http://accc.riken.jp/HPC_e/himenobmt_e.html
- Strong scaling benchmark
 - XL configuration: 1024 x 512 x 512 global volume
 - Expect halo exchanges to become significant
 - Use asynchronous GPU data transfers and kernel launches to help avoid this



Porting Himeno to the Cray XK6

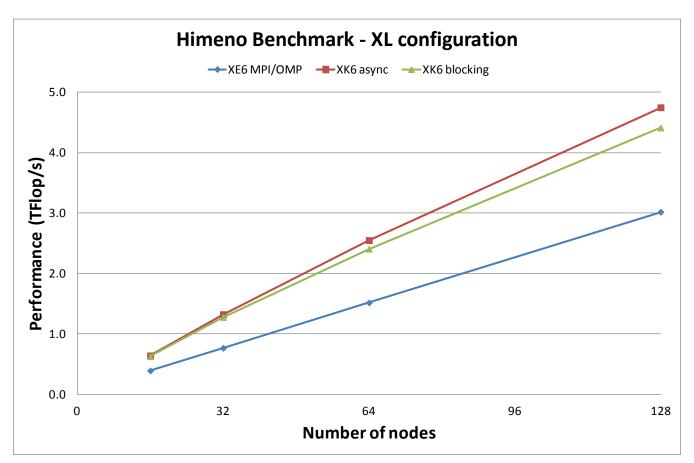


- Several versions tested, with communication implemented in MPI and Fortran coarrays
- GPU version using OpenACC accelerator directives
 - Total number of accelerator directives: 27
 - plus 18 "end" directives
- Arrays reside permanently on the GPU memory
- Data transfers between host and GPU are:
 - Communication buffers for the halo exchange
 - Control value
- Cray XK6 timings compared to best Cray XE6 results (hybrid MPI/OpenMP)

Himeno performance

CRAY

- XK6 GPU is about 1.6x faster than XE6
- OpenACC async implementation is ~ 8% faster than OpenACC blocking



CloverLeaf



2D hydro code, with several stencil-type operations

Developed by AWE

- Using to explore programming models
- to be released as Open Source to the Mantevo project hosted by Sandia (miniapps)

Current performance for 87 steps

Mesh	CUDA	OpenACC
960x960	2.44	2.03
3840x3840	37.42	31.77

GAMESS



- Computational chemistry package suite developed and maintained by the Gordon Group at Iowa State University
 - http://www.msg.ameslab.gov/gamess/
- ijk-tuples kernel Source changes
 - CUDA 1800 lines of hand-coded CUDA
 - OpenACC approximately 75 directives added to the original source
- Performance of ijk-tuples on 16 XK6 Nodes with Fermi
 - CPU Only (16 ranks per node) 311 Seconds
 - CUDA 134 seconds
 - OpenACC 138 seconds
 - CUDA was only ~3% faster than OpenACC
- Performance of ijk-tuples on 16 XK6 Nodes with Kepler
 - CPU Only (16 ranks per node) 311 Seconds
 - CUDA 76.6 seconds
 - OpenACC 68.1 seconds
 - OpenACC was ~12.5% faster than CUDA !!

Summary

- Cray provides a high level programming environment for acceletate Computing
 - Fortran, C, and C++ compilers
 - OpenACC directives to drive compiler optimization
 - Compiler optimizations to take advantage of accelerator and multi-core X86 hardware appropriately
 - Cray Reveal
 - Scoping analysis tool to assist user in understanding their code and taking full advantage of SW and HW system
 - Cray Performance Measurement and Analysis toolkit
 - Single tool for GPU and CPU performance analysis with statistics for the whole application
 - Parallel Debugger support with allinea DDT
 - Auto-tuned Scientific Libraries support
 - Getting performance from the system ... no assembly required