

Introduction to the Cray Accelerated Scientific Libraries

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What makes Cray libraries special?



1. Node performance

Highly tuned routines at the low-level (ex. BLAS)

2. Network performance

- Optimized for network performance
- Overlap between communication and computation
- Use the best available low-level mechanism
- Use adaptive parallel algorithms

3. Highly adaptive software

 Use auto-tuning and adaptation to give the user the known best (or very good) codes at runtime

4. Productivity features

Simple interfaces into complex software

LibSci Usage



LlbSci

- The drivers should do it all for you. No need to explicitly link.
- CCE will automatically pattern match to select scientific libraries
- For threads, set OMP_NUM_THREADS
 - Threading is used within libsci.
 - If you call within a parallel region, single thread used

FFTW

module load fftw (there are also wisdom files available)

PETSc

- module load petsc (or module load petsc-complex)
- Use as you would your normal PETSc build

Trilinos

- module load trilinos
- CASK no need to do anything you get optimizations free

Check you got the right library!



- Add options to the linker to make sure you have the correct library loaded.
- -WI adds a command to the linker from the driver
- You can ask for the linker to tell you where an object was resolved from using the –y option.
 - E.g. –WI, -ydgemm_

```
.//main.o: reference to dgemm_
/opt/xt-libsci/11.0.05.2/cray/73/mc12/lib/libsci_cray_mp.a(dgemm.o):
definition of dgemm_
```

Note: explicitly linking "-lsci" is bad! This won't be found from libsci 11+ (and means single core library for 10.x!)

LibSci for Accelerators: libsci_acc



- Provide basic libraries for accelerators, tuned for Cray
- Must be independent to OpenACC, but fully compatible
- Multiple use case support
 - Get the base use of accelerators with no code change
 - Get extreme performance of GPU with or without code change
 - Extra tools for support of complex code
- Incorporate the existing GPU libraries into libsci
 - CUBLAS
 - Magma
 - Cray Implementation BLAS/LAPACK
- Provide additional performance and usability
 - OpenACC support
 - CUDA support
- Maintain the Standard APIs where possible!

Why libsci_acc?



- Code modification is required to use existing GPU libraries!
- Several scientific library packages are already there
 - 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

Autotuning



Cray Autotuning framework has been built to tune all BLAS for accelerators

- GPU kernel codes are built using code generator
- Enormous offline autotuning is used to build a map of performance to input
- An adaptive library is built from the results of the autotuning
- At run-time, your code is mapped to training set of input
- Best kernel for your problem is used

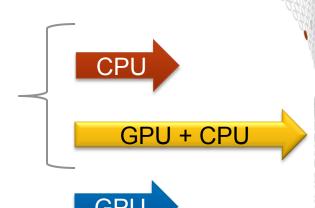
Three interfaces for three use cases



Simple interface



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



Device interface

dgetrf_acc(M, N, d_A, lda, ipiv, &info)



CPU interface

dgetrf_cpu(M, N, A, lda, ipiv, &info)



Simple interface



- You can pass either host pointers or device pointers to simple interface
- Host memory pointer
 - Performs hybrid operation on GPU
 - If problem is too small, performs host operation
- Device memory pointer
 - Performs operation on GPU
- BLAS 1 and 2 perform computation local to the data location
 - CPU-GPU data transfer is too expensive to exploit hybrid execution

Device interface



- Device interface gives higher degrees of control
- Requires that you have already copied your data to the device memory
- API
 - Every routine in libsci has a version with _acc suffix
 - E.g. dgetrf_acc
 - This resembles standard API except for the suffix and the device pointers

CPU interface



- Sometimes apps may want to force ops on the CPU
 - Need to preserve GPU memory
 - Want to perform something in parallel
 - Don't want to incur transfer cost for a small op
- Can force any operation to occur on CPU with _cpu version
- Every routine has a _cpu entry-point
- API is exactly standard otherwise

Usage - basics



- 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 DGEMM example

- Starting with a code that relies on dgemm.
- The library will check the parameters at runtime.
- If the size of the matrix multiply is large enough, the library will run it on the GPU, handling all data movement behind the scenes.
- NOTE: Input and Output data are in CPU memory.

call dgemm('n','n',m,n,k,alpha,&
a,lda,b,ldb,beta,c,ldc)

libsci_acc interaction with OpenACC



- If the rest of 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 interaction 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
```

Advanced controls

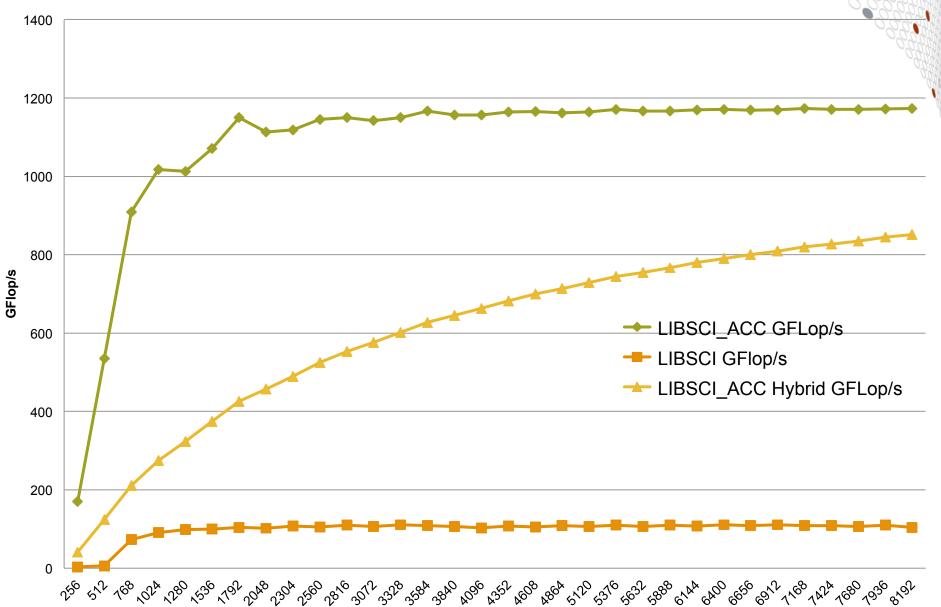


The communication avoidance (CA) version of DGETRF/ ZGETRF can be enabled by setting the environment variable LIBSCI ACC DLU = CALU / LIBSCI ACC ZLU = CALU

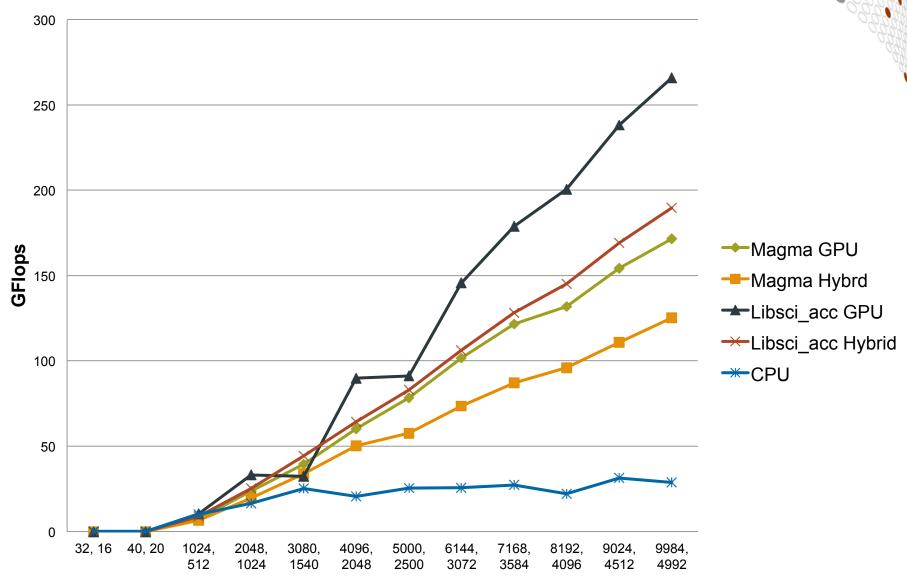
- Change Split Ratio of Hybrid GEMM routines
 - LIBSCI SGEMM SPLIT=0.9
 - LIBSCI DGEMM SPLIT=0.8
 - LIBSCI CGEMM SPLIT=0.9
 - LIBSCI ZGEMM SPLIT=0.8
- Force simple API to always call CPU routine
 - CRAY LIBSCI ACC MODE=2

Matrix Multiplication :: Double (DGEMM)

XK7 Kepler :: Nov 2012

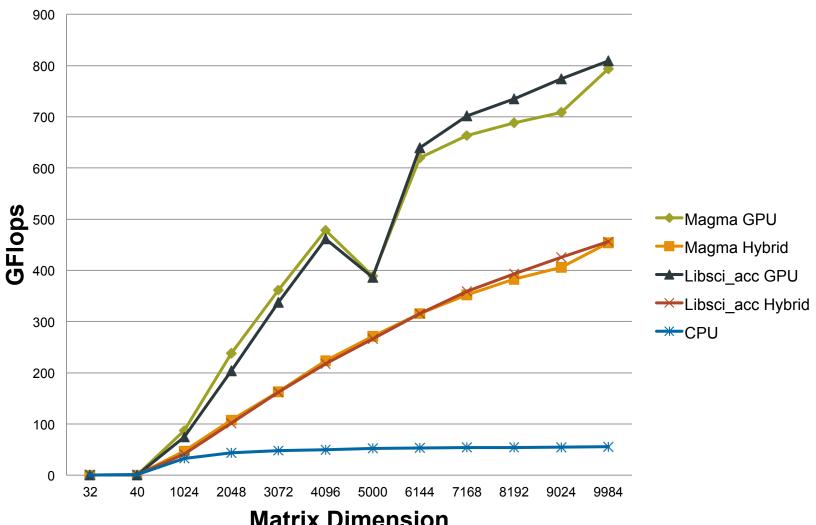






LAPACK LU factorization :: double complex (ZGETRF) XK7 Kepler :: Nov 2012





Summary



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

Tuning requests



- CrayBLAS is an auto-tuned library
 - Generally, excellent performance is possible for all shapes and sizes
- However, the adaptive CrayBLAS can be improved by tuning for exact sizes and shapes
- Send your specific tuning requirements to

crayblas@cray.com

Send the routine name and the list of calling sequences



Questions ?

Targets



- BLAS
 - [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
 - All level 2 BLAS
 - All level 1 BLAS

- LAPACK
 - [d,z]GETRF
 - [d,z]GETRS
 - [d,z]POTRF
 - [d,z]POTRS
 - [d,z]GESDD
 - [d,z]GEBRD
 - [d,z]GEQRF
 - [d,z]GELQF

Planned



- LAPACK Eigenvalue
 - DSYEVR
 - ZHEEVR
 - DSYGVR
 - ZHEGVR
 - DGEEV
 - ZGEEV

- LAPACK Least squares
 - DGELS/ZGELS
- LAPACK Linear solvers
 - DGESV/ZGESV
 - DPOSV/ZPOSV
 - DSYSV/ZSYSV