Multiple Right-hand-side Implementation for ${\sf DD}\alpha{\sf AMG}$

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QFT

In path-integral formulation, all physical predictions in QFT is the generating functional:

$$Z=\int {\cal D}\phi {
m e}^{i{
m S}}.$$

In particular, for QCD,

$$S_{
m QCD} = \int d^4 x {\cal L}_{
m QCD} = S_\psi + S_G$$

where

$$S_\psi = \int d^4x \sum_f ar{\psi}_f \left(i \gamma_\mu D_\mu - m_f
ight) \psi_f$$
 and $S_G = S_G(A^a_\mu).$

The expectation value of an observable can be computed using

$$\langle O \rangle = rac{1}{Z} \int \mathcal{D} \phi O(\phi) e^{iS}.$$

Lattice QCD

Now the task is to evaluate $\langle {\cal O} \rangle.$ We do this non-perturbatively. To be more specific,

We perform Wick rotation from Minkovski to Eulcidean space-time to obtain

$$\langle O
angle = rac{1}{Z} \int \mathcal{D} \phi O(\phi) e^{-S_{\mathrm{E}}}$$

where $S_{\rm E}$ is a real-valued Euclidean action.

▶ Then, we use $e^{-S_{\rm E}}$ as a Monte-Carlo weight and have

$$\langle O \rangle = \lim_{N \to \infty} \frac{1}{N} \sum_{i}^{N} O_i \approx \frac{1}{N} \sum_{i}^{N} O_i.$$

To make the computation doable on a computer,

- We discretize the space-time into the lattice with its spacing a and dimension L³ × T.
- Fermion fields, ψ, live on a lattice site, and gauge fields are replaced by a link, U_μ, connecting the adjacent points.



The Problem

As the fermion fields in the action are Grassmann numbers, using its properties, we can take integration over fermion fields explicitly to obtain

$$\langle O \rangle = \frac{1}{Z} \int \mathcal{D} U O(D^{-1}, U) \mathrm{Det} D(U) e^{-S_G}$$

where D is a Dirac matrix in $S_{\psi} = \bar{\psi} D \psi = \sum_{f} \bar{\psi}_{f} \left(\not{\!\!\!D}_{E} + m_{f} \right) \psi_{f}$.

Evaluation of the expectation value requires

- computation of the inverse of the Dirac matrix repeatedly many times
- generation of gauge configurations according to the weight $e^{-S_c} \text{Det}D(U)$ via importance sampling, e.g., HMC methods

In lattice QCD calculations, the main computational task is to solve the system

$$Dx = b.$$

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The Matrix

- Dirac matrix: $D = \not{D}_E + m$
- Wilson Dirac operator: $D_{\rm W} = \frac{1}{2} \sum_{\mu=0}^{3} \left(\gamma_{\mu} (\Delta_{\mu} + \Delta^{\mu}) - a \Delta_{\mu} \Delta^{\mu} \right) + \frac{m}{a} \text{ where}$

$$\Delta^{\mu}\psi = \frac{1}{a} \left(U_{\mu}(x)\psi(x+\hat{\mu}) - \psi(x) \right)$$
$$\Delta_{\mu}\psi = \frac{1}{a} \left(\psi(x) - U^{\dagger}_{\mu}(x-\hat{\mu})\psi(x-\hat{\mu}) - \psi(x) \right).$$



4) Q (↓ 6 / 51 The Matrix for Twisted Mass Fermions The clover-improved Wilson Dirac operator:

$$D_{\mathrm{cW}} = D_{\mathrm{W}} - rac{c_{sw}}{32} \sum_{\mu,\nu=0}^{3} (\gamma_{\mu}\gamma_{\nu}) \otimes (Q_{\mu\nu}(x) - Q_{\nu\mu}(x))$$

here
$$Q_{\mu
u}(x) = \sum_{n=0}^{3} U_{\mu
u}^{\odot^{n}}(x)$$
 and
 $U_{\mu
u}(x) = U_{\mu}(x)U_{\nu}(x+\hat{\mu})U_{\mu}^{\dagger}(x+\hat{\nu})U_{\nu}^{\dagger}(x)$
 $U_{\mu
u}^{\odot}(x) = U_{\nu}(x+\hat{\mu})U_{\mu}^{\dagger}(x+\hat{\nu})U_{\nu}^{\dagger}(x)U_{\mu}(x)$



The degenerated twisted mass operator:

w

$$egin{aligned} D_D(\mu) &= (D_{
m cW}\otimes I_2) + i\mu(\Gamma_5\otimes au_3) \ &= egin{pmatrix} D_{
m TM}(\mu) & 0 \ 0 & D_{
m TM}(-\mu) \end{pmatrix} \end{aligned}$$

where $D_{
m TM}(\mu)=D_{
m cW}+i\mu\Gamma_5$



Why do we need a linear solver?

- ► Typically, for modern simulations, the size of the lattice is $V = 128 \times 64^3 \sim 3 \times 10^7$.
- ► Then, a typical size of the matrix is $n \times n$ with $n = V \times \underbrace{4}_{\text{spins}} \times \underbrace{3}_{\text{colors}} \times \underbrace{2}_{\text{complex}} \sim 8 \times 10^8 \simeq 6.4 \text{GB}$
- For general matrix inversion, the computational cost is O(n^{2.376}) (Optimized CW-like algorithms)
- ▶ If we were to store all entries of the inverse, we need the memory space of $n \times n \simeq 6.5 \times 10^{17} \simeq 5 \times 10^9 \text{GB} = 5 \text{EB}$
- It is impractical in lattice computations to invert D explicitly and store all its entries.

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Why do we need a good linear solver?

• We switch to solving Dx = b.

- b is usually a random vector or a point source, i.e., b = δ_{xa,yb} where x, y are lattice sites and a, b are internal quantum numbers.
- In practice, computation of point-to-all propagators, i.e., the solutions of Dx = b with a point source, is often sufficient, e.g., for two-point correlation functions.
- The inversion needs to be done a lot of times.

For example, computation of nucleon structure requires $\underbrace{4}_{\rm spins} \times \underbrace{3}_{\rm colors} \times \underbrace{2}_{\rm flavors} \times \underbrace{5}_{\rm proj} \times \underbrace{200}_{\rm src\ pos} \times \underbrace{400}_{\rm configs} \sim 10^7 \text{ inversions.}$

We need a good solver!

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Krylov Solver

- Due to sparse nature of the matrix, the computational cost for matrix-vector multiplication is O(n)
- We use a Krylov method, which consists of constructing a Krylov subspace K_k = span{b, Db, D²b, D³b, ..., D^{k−1}b}, projecting the system onto K, and solving the projected system to obtain an approximate solution x_k.
- As it is a projection-based method, it does not require a large memory space.



Krylov Solver - Critical Slowing Down

- The condition number for D is roughly proportional to the inverse of the mass.
- The matrix D becomes singular as the mass approaches its critical value m_{critc}.
- Convergence to the solution slows down.
- Inversion on and generation of gauge configuration at physical light quark masses leads to a larger condition number and thus increasing solver time.



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Multigrid Solvers

Among various solvers, a class of solvers based on multigrid approaches in preconditioning Krylov subspace solvers has turned out to be very successful (Luscher 2007b; Luscher 2007a; Osborn et al. 2010; R. Babich et al. 2010; Frommer et al. 2013).

Several of the implementations for clover Wilson fermions that are openly available includes

- A two-level multigrid approach based on Lüschers inexact deflation (Luscher 2007b) available as OpenQCD at http://luscher.web.cern.ch/luscher/openQCD/
- Multigrid with generalized conjugate residual (MG-GCR) (J. Brannick et al. 2008; Clark et al. 2008; Ronald Babich et al. 2009; R. Babich et al. 2010) available as part of the USQCD package QOPQDP at https://github.com/usqcd-software/qopqdp and QUDA at http://lattice.github.io/quda/
- An aggregation-based domain decomposition multigrid (DDαAMG) approach (Frommer et al. 2013) available at https://github.com/DDalphaAMG/DDalphaAMG.

Extending Multigrid Solvers

Extension of DD α AMG to twisted mass fermions:

- C. Alexandrou, S. Bacchio, J. Finkenrath, A. Frommer, F. Kahl, and M. Rottmann, Adaptive Aggregation-based Domain Decomposition Multigrid for Twisted Mass Fermions, Phys. Rev. D 94, 11 (2016)
- The generalization to non-degenerate twisted mass fermions is discussed in (Alexandrou, Bacchio, and Finkenrath 2019)
- Publicly available at https://github.com/sbacchio/DDalphaAMG
- A version with new features at https://github.com/sy3394/DDalphaAMG

Extension to other fermions:

- ▶ MG-GCR to domain-wall fermions in (Cohen et al. 2011)
- **DD** α AMG to overlap fermions in (James Brannick et al. 2016)

$DD\alpha AMG$ - Basics

- Adaptive Aggregation-based Domain Decomposition Multigrid method
- ▶ The algorithm is designed to invert a large sparse matrix effectively
- In particular, the algorithm is applied to Dirac matrices for Wilson or twisted mass fermions
- It takes advantage of the sparse structure of the Dirac matrix.



$DD\alpha AMG$ - Preconditioners

- To circumvent the issue of critical slowing down and effectively invert the large sparse matrix, DDαAMG uses two preconditioners: a smoother and coarse grid correction
- For a smoother, we use red-black Schwarz Alternating Procedure (SAP) (Luscher 2007a).
- For coarse grid correction, we use Algebraic MultiGrid (AMG) (Wesseling 1995).



 $\label{eq:linear} \mbox{Picture Courtesy: Luke Olson, http://lukeo.cs.illinois.edu/cs556 + < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a > < a$

- The lattice is divided into red blocks and black blocks in a chessboard manner.
- Due to nearest neighbor interactions, block of a color couples to a block of the other color



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- ▶ In SAP, we neglect off-diagonal parts, *D_{rb}*, *D_{br}*, i.e., inter-block interactions.
- Then, within each group of blocks, we visit blocks sequentially and invert the block matrix locally.



- Depending on when we perform the global residual update, error propagates differently.
- In additive SAP, residual update, r ← b − Dx, is done once before local inversion over blocks.

$$\varepsilon \leftarrow \left(I - \sum_{i} D_{i}^{-1} D\right) \varepsilon$$

In multiplicative SAP, residual update is done every time before local inversion on each block.

$$\varepsilon \leftarrow \left[\prod_i (I - D_i^{-1}D)\right]\varepsilon$$

Information spread faster with multiplicative SAP.

In red-black SAP, within a group of blocks of each color, additive SAP is adopted, and over groups of blocks, multiplicative SAP is used.

Define

$$B_{rr}=egin{pmatrix} D_{rr}^{-1}&0\0&0\end{pmatrix}$$
 and $B_{bb}=egin{pmatrix} 0&0\0&D_{bb}^{-1}\end{pmatrix}.$

Then, error propagation is given by

$$\varepsilon \leftarrow (I - MD)\varepsilon = (I - B_{rr}D)(I - B_{bb}D)\varepsilon.$$

Since local block matrices live on a single MPI rank, their inversion does not call for global reduction so that SAP is suitable for parallelization. As it involves inversion of local block matrices, it removes UV-modes from the residual



Coarse Grid Correction

Basic Ideas:

- Start with Dx = b
- Restrict the problem onto a coarse grid
- Solve $D_c x_c = r_c$
- Prolong the solution back to the original grid



 $\label{eq:linear} \mbox{Picture Courtesy: Luke Olson, http://lukeo.cs.illinois.edu/cs550 \mbox{ } < \mbox{ } > \mbox{ } < \mbox{ } > \mbox{ } < \mbox{ } > \mbox{ } < \mbox{ }$

MG

To perform coarse-grid correction, we use a multigrid method. For that, we need

- the restriction operator, R, which maps the problem from a finer lattice to a coarser lattice
- the prolongation operator, P, which maps the problem from a coarser lattice to a finer lattice

$$\blacktriangleright RIP = I_c$$

The coarsened problem is significantly cheaper to solve.



MG

Using those two operators, we construct a coarse-grid Dirac matrix, $D_c = RDP$, and coarse-grid vectors, $x_c = Rx$, $r_c = R(b - Dx)$. Then,

• Coarse-grid inversion: $D_c x_c = r_c$

- Coarse-grid correction: $x \leftarrow x + PD_c^{-1}R(b Dx)$
- Error propagation: $\varepsilon \leftarrow (I PD_c^{-1}RD)\varepsilon$

In multi-level approaches, we generalize the coarse-grid correction to

$$\triangleright R_I I_I P_I = I_{I+1}$$

$$\blacktriangleright D_{I+1} = R_I D_I P_I \text{ with } D_1 = D$$

$$D_{l+1}x_{l+1} = r_{l+1} = R_l(b_l - D_lx_l)$$

$$> x_l \leftarrow x_l + P_l D_{l+1}^{-1} R_l (b - D_l x_l)$$

 $\triangleright \varepsilon_{l} \leftarrow (I_{l} - P_{l}D_{l+1}^{-1}R_{l}D_{l})\varepsilon_{l}$

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AMG

- For a standard MG methods, restriction and prolongation are defined geometrically, for example using block aggregation.
- In algebraic multigrid approaches, we do not base the definition of P and R on geometry of the lattice.
- Instead, we use eigenvectors of the Dirac operator with small eigenvalues to define P and R based on partition of the lattice into aggregates.
- This constructed P effectively captures the low-modes of D due to local coherence (Luscher 2007b).



AMG

So in AMG, we define a coarse-grid Dirac matrix directly from the Dirac matrix on the fine lattice.



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Coarse-grid correction removes low-modes from the residual.

Coarse grid correction



SAP+AMG

So when we combine the two preconditioners as follows,

$$\varepsilon_{I} \leftarrow (I - M_{I}D_{I})^{k}(I_{I} - P_{I}D_{I+1}^{-1}R_{I}D_{I})(I_{I} - M_{I}D_{I})^{j}\varepsilon_{I},$$

the low-modes and high-modes of the error are both suppressed.



$\mathsf{DD}\alpha\mathsf{AMG}$ with SAP and AMG

Iterative adaptive setup phase:

- We construct *P* and so the coarse Dirac operator at each level.
- Instead of applying eigensolver to D to get a few approximate eigenvectors with small eigenvalues, we construct them in an adaptive manner using two preconditioners.
- First, start with a set of random vectors, v_i 's, solve $Dv_i = 0$ using only smoother to reduce the high modes from v_i , and construct initial multigrid hierarchy.
- Then, compute $D_l^{-1}v_i$ with post-smoothing at each level and update v_i as well as P and D_c .

$\mathsf{DD}\alpha\mathsf{AMG}$ with SAP and AMG

- Solver phase:
 - We combine coarse-grid correction and Krylov method with a cycling strategy called "K-cycle" to obtain a recursive Krylov solver.
 - Residuals, max Krylov space size, and a number of restarts can be set to different values at different levels.
 - In particular, the residual of 10^{-1} is sufficient at the bottom even when the goal residual at the top is 10^{-10} .



$DD\alpha AMG$ - Performance

- MG solvers outperforms traditional Krylov subspace solvers like the conjugate gradient solver at small quark masses
- DDalphaAMG for twisted mass fermions is two orders of magnitude faster than CG



$DD\alpha AMG$ - Performance



► HMC simulation:

$DD\alpha AMG$ - Scaling

- Bottlenneck of Multigrid methods is the scalability
- Ideal scaling breaks down, and performance stagnates for parallelization above 125 Skylake nodes in case of a 3-level MG approach
- With the current hardware trends higher core counts per node the scalability window will even shrink further



Figure 2: A scaling plot on the ensemble of $N_f = 2 + 1 + 1$ twisted mass clover with $a \sim 0.07$ fm and $V = 80^3 \times 160$ at physical point simulated on SuperMUC-NG (Intel Xeon ("Skylake")) at LRZ

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Multiple R.H.S. - Objectives

Originally,

- the code inverted each rhs one by one
- also, loops are vectorized manually using instruction sets for a specific SIMD extension

However,

- We can perform multiple inversions more efficiently.
- We also want to improve portability of our code letting compilers perform optimization analysis and vectorization.

Thus,

• We solve the system of equations with multiple right-hand sides (rhs) simultaneously $(b \rightarrow \mathbf{b})$.

Multiple R.H.S. - Implementation

We define a new data structure for a bundle of vectors.

- Vectors in the bundle are ordered in such a way that the index on vectors runs the fastest.
- ► All low-level routines are rewrited to respect the new structure.
- We process a bundle of right-hand vectors simultaneously using SIMD vectorization of loops.



Multiple R.H.S. - Implementation

- Instead of manually vectorizing the loops using instruction sets, we auto-vectorize the loops using pragmas: _Pragma("unroll"), _Pragma("vector aligned"), and _Pragma("ivdep").
- These pragmas are applied to a for-loop of a pre-determined iteration length: for(jj=0; jj<num_loop; jj++).</p>
- The number of rhs are assumed to be multiple of num_loop.
- This shifts vectorization from 128 bit to 256 bit

Num. R.H.S.	1 rhs		4 rhs		8 rhs	
Instruction Mix	SP Flops	DP Flops	SP Flops	DP Flops	SP Flops	DP Flops
						0.000/
128-bit	95.26%	86.59%	23.41%	4.99%	24.92%	3.60%
256-bit	2.58%	1.26%	60.68%	78.13%	74.02%	94.76%
Total	97.26%		84.03%		98.81%	

Table 1: Vectorization Reports

Scaling

Conclusion:

Breakdown of strong scaling can be pushed to higher parallelization, mutiple rhs shows scalability up to 512 nodes



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Incorporating Block Krylov Solver

- Fast Accurate Block Linear krylOv Solver (Fabulous) is an external library implementing block Krylov solvers such as GMRES and GCR (Robbé and Sadkane 2006; Morgan 2005; Agullo, Giraud, and Jing 2014)
- It combines BGMRES with detection of inexact breakdown, deflation, and incremental QR factorization.
- It provides several different orthogonalization schemes.
- ► We linked the DDαAMG code to Fabulous and make it available non-block GMRES or one of the solvers provided via fabulous library at each level as a solver.
- Our implementation of multiple r.h.s. stultifies inexact breakdown.

Parameters

For Fabulous,

- Solvers: BGMRES, BGCR, BGMRES with deflated restarting (DR), BGMRES with incremental QR factorization (QR), BGMRES with DR and QR (DRQR)
- The number of deflating eigenvectors (for DR methods).
- Orhogonalization schemes: Classical Gram-Schmidt (CGS), Modified Gram-Schmidt (MGS), Iterative CGS (ICGS), Iterative MGS, each possibly with blocking

For $\mathsf{DD}\alpha\mathsf{AMG}$,

- \blacktriangleright 0th-level (bottom) and 1st-level (intermediate) residuals
- 0th-level (bottom) and 1st-level (intermediate) max Krylov space size and number of restarts

Tuning Constants

The systems used for tuning:

- Lattice: $48^3 \times 98$ at physical point
- System: Cyclone (Intel Xeon Gold 6248) at The Cyprus Institute
- Used 6 nodes with 32 cores each

Fixed solver parameters:

- (The number of levels) = 3.
- ► The solver at the top: FGMRES
- ► The solver at the level 1: FGMRES

Tuning Orthogonalization Schemes





Tuning Solvers

Best Solver: BGMRES with QR and DR



Tuning Residuals

Solver: BGMRES



Tuning Max Krylov Space Size

Solver: BGMRES with QR



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Tuning Mu Factor

• The mu factor is the factor multiplying μ in $D_{\rm TM}$ at the bottom $(\mu \to \delta_\mu \mu)$

Solver: BGMRES with QR



Preliminary Tuning Results



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- Scalability is extended by around a factor 5.
- We also want to test out code on different architectures like AMD Epyc's and ARM chips, e.g., Fujitsu A64FX.
- We will test reusable deflation space
- We consider a pipeline version of the solver to speed up the coarse grid correction for extending the scaling region
- ► DDαAMG using a fabulous solver at the bottom showed a promising result.
- We need to tune with more right-hand sides to confirm the trend.

Thank you!



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