





Parallel Performance and Numerical Stability of Communication Hiding Pipelined Krylov Subspace Methods

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Motivation Exascale systems projection

	Today's Systems	Predicted Exascale Systems*	Factor Improvement
System Peak	10 ¹⁶ flops/s	10^{18} flops/s	100
Node Memory Bandwidth	10^2 GB/s	10 ³ GB/s	10
Interconnect Bandwidth	10 ¹ GB/s	10 ² GB/s	10
Memory Latency	$10^{-7} { m s}$	$5\cdot 10^{-8}\text{s}$	2
Interconnect Latency	$10^{-6} \mathrm{s}$	$5\cdot 10^{-7}~\text{s}$	2

*Sources: from P. Beckman (ANL), J. Shalf (LBL), and D. Unat (LBL)

- Data movement (communication) is much more expensive than flops (computation) in terms of both time and energy
- Reducing time spent moving data/waiting for data will be essential for exascale applications
 - ⇒ Communication avoiding / Communication hiding



Motivation Communication hiding vs. communication avoiding

Communication cost has motivated several approaches to reducing global synchronization cost in Krylov subspace methods:

Avoiding communication: s-step Krylov subspace methods * [A. Chronopoulous, J. Demmel, M. Hoemmen, E. Carson, L. Grigori, J. Erhel, . . .]

- Compute iterations in blocks of s (change of Krylov subspace basis)
- Reduces number of synchronizations per iteration by a factor of $\mathcal{O}(s)$

Hiding communication: Pipelined Krylov subspace methods * [P. Ghysels, W. Vanroose, S. C., P. Sanan, B. Gropp, I. Yamazaki, ...]

- Introduce auxiliary vectors to decouple SpMV and inner products
- Enables overlapping of communication and computation

^{*} Both equivalent to corresponding Krylov subspace methods in exact arithmetic





Iteratively improve an approximate solution of linear system Ax = b,

$$x_i \in x_0 + \mathcal{K}_i(A, r_0) = x_0 + \text{span}\{r_0, Ar_0, A^2r_0, \dots, A^{i-1}r_0\}$$





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- ▶ minimize certain error measure over Krylov subspace $K_i(A, r_0)$
- Krylov subspace methods:

Conjugate Gradients (CG), Lanczos, GMRES, MinRES, BiCG, CGS, BiCGStab, CGLS, ...

▶ Preconditioners:

AMG & GMG, Domain Decomposition Methods, FETI, BDDC, Incomplete factorization, Physics based preconditioners, ...





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- usually in combination with sparse linear algebra/stencil application
- three algorithmic building blocks:
 - i. dot-product
 - $\mathcal{O}(N)$ flops
 - global synchronization (MPI_Allreduce)
 - ii. SpMV
 - $\mathcal{O}(nnz)$ flops
 - neighbor communication only
 - iii. axpy
 - O(N) flops
 - no communication



Krylov subspace methods Classical CG

Algorithm CG	
1: procedure $CG(A, b, x_0)$ 2: $r_0 := b - Ax_0$; $p_0 = r_0$ 3: for $i = 0, \dots$ do	
$ \begin{aligned} 4 &: & s_i \coloneqq Ap_i \\ 5 &: & \alpha_i \coloneqq (r_i, r_i) / (s_i, p_i) \\ 6 &: & x_{i+1} \coloneqq x_i + \alpha_i p_i \\ 7 &: & r_{i+1} \coloneqq r_{i-1} - \alpha_i s_i \\ 8 &: & \beta_{i+1} \coloneqq (r_{i+1}, r_{i+1}) / (r_i, r_i) \\ 9 &: & p_{i+1} \coloneqq r_{i+1} + \beta_{i+1} p_i \\ 10 &: & \text{end for} \end{aligned} $ $ 11 :: \text{end procedure} $	dot-pr SpMV axpy

Hestenes & Stiefel (1952)

i. dot-products

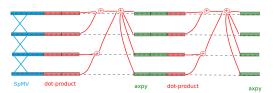
- ▶ 2 global reductions: latency dominated
- ► scales as log₂(#partitions)

ii. SpMV

- ► computationally expensive
- ▶ good scaling (minor comm.)

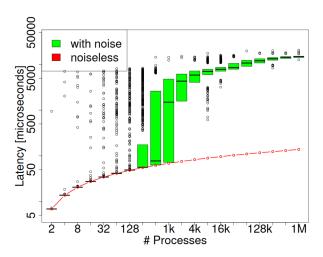
iii. axpy's

- vector operations (recurrence relations)
- perfect scaling (no comm.)





Krylov subspace methods Global reduction latency



T. Hoeffler, T. Schneider and A. Lumsdaine, SC10, 2010





```
Algorithm
                   Pipelined CG
 1: procedure PIPE-CG(A, b, x_0)
         r_0 := b - Ax_0; w_0 := Ar_0
         for i = 0, \dots do
 4: \gamma_i := (r_i, r_i)
 5: \delta := (w_i, r_i)
 6: q_i := Aw_i
             if i > 0 then
                  \beta_i := \gamma_i/\gamma_{i-1}; \alpha_i := (\delta/\gamma_i - \beta_i/\alpha_{i-1})^{-1}
 9.
              else
                  \beta_i := 0; \alpha := \gamma_i / \delta
10.
                                                             dot-pr
11-
              end if
                                                             SpMV
12: z_i := q_i + \beta_i z_{i-1}
13: s_i := w_i + \beta_i s_{i-1}
                                                             axpy
14: p_i := r_i + \beta_i p_{i-1}
15: x_{i+1} := x_i + \alpha_i p_i
16: r_{i+1} := r_i - \alpha_i s_i
17: w_{i+1} := w_i - \alpha_i z_i
18:
         end for
19: end procedure
```

Ghysels & Vanroose (2014)

Pipelined CG = re-engineered version of CG for improved parallel performance





```
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    procedure PIPE-CG(A, b, x<sub>0</sub>)

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Re-ordering of operations requires new auxiliary variables:

$$s_i = Ap_i, w_i = Ar_i, z_i = As_i$$





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Ghysels & Vanroose (2014)

Pipelined CG = re-engineered version of CG for improved parallel performance

► Re-ordering of operations requires new auxiliary variables:

$$s_i = Ap_i, w_i = Ar_i, z_i = As_i$$

Derive recurrence relations to avoid computing additional SpMV's, e.g.:

$$s_i := Ap_i$$
 $p_i = r_i + \beta_i p_{i-1}$
 $\downarrow \downarrow$
 $s_i = w_i + \beta_i s_{i-1}$
with $w_i := Ar_i$



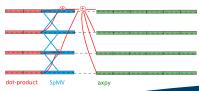


```
Algorithm
                  Pipelined CG

    procedure PIPE-CG(A, b, x<sub>0</sub>)

        r_0 := b - Ax_0; w_0 := Ar_0
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                 \beta_i := 0; \alpha := \gamma_i / \delta
10.
                                                           dot-pr
11-
             end if
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                                                           axpy
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15: x_{i+1} := x_i + \alpha_i p_i
16: r_{i+1} := r_i - \alpha_i s_i
17: w_{i+1} := w_i - \alpha_i z_i
18:
         end for
19: end procedure
```

- Communication avoiding: dot-products grouped in one global reduction phase per iteration
- ii. Communication hiding: overlap global synchronization with SpMV (+ Prec) computation
- iii. No free lunch: Additional recurrence relations, i.e. axpy's, for auxiliary var's $s_i = Ap_i$, $w_i = Ar_i$, $z_i = As_i$



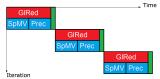


Krylov subspace methods Pipelined KSM with deep pipeline

Classic KSM:



Pipelined KSM:



Deep pipelined KSM:

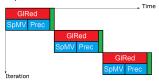




Classic KSM:



Pipelined KSM:



Deep pipelined KSM:



Krylov subspace methods Pipelined KSM with deep pipeline

Consider the Arnoldi relation

$$AV_i = V_{i+1}H_{i+1,i}$$

with V_i the Krylov subspace basis and $H_{i+1,i}$ upper Hessenberg. Introduce the auxiliary vectors $Z_{i+1} = [z_0, z_1, \dots, z_i]$ as

$$z_{j} := \begin{cases} v_{0} & j = 0, \\ P_{j}(A)v_{0} & 0 < j \leq I, \\ P_{I}(A)v_{j-1} & j > I, \end{cases}$$

with polynomials $P_l(t)$ of fixed order l, where l is the pipeline length

$$P_l(t) := \prod_{j=0}^{l-1} (t-\sigma_j).$$



Krylov subspace methods Pipelined KSM with deep pipeline

Writing the Arnoldi relation $AV_i = V_{i+1}H_{i+1,i}$ for basis vector v_{j-1} and applying $P_i(A)$ to both sides, we obtain recurrence relations for z_i :

$$v_{j-l} = \frac{Av_{j-l-1} - \sum_{k=0}^{j-l-1} h_{k,j-l-1}v_k}{h_{j-l,j-l-1}} \quad \Rightarrow \quad \mathbf{z}_j = \frac{A\mathbf{z}_{j-1} - \sum_{k=0}^{j-l-1} h_{k,j-l-1}\mathbf{z}_{k+l}}{h_{j-l,j-l-1}}.$$

This implies an Arnoldi-like relation for the auxiliary variables

$$AZ_i = Z_{i+1}B_{i+1,i}$$

with



Krylov subspace methods Pipelined KSM with deep pipeline

<u>Basis transformation:</u> Z_i and V_i both span the i-th Krylov subspace, thus

$$Z_i = V_i G_i$$

where G_i is an upper triangular matrix.

Recursive calculation of entries G_i :

$$g_{j,k-1} = (z_{k-1}, v_j) = \left(z_{k-1}, \frac{z_j - \sum_{m=0}^{j-1} g_{m,j} v_m}{g_{j,j}}\right)$$

$$= \frac{(z_{k-1}, z_j) - \sum_{m=0}^{j-1} g_{m,j} g_{m,k-1}}{g_{j,j}} \qquad (\forall j = 0, 1, \dots, k-1)$$

Diagonal element:

$$g_{j,j} = \frac{(z_j, z_j) - \sum_{m=0}^{j-1} g_{m,j} g_{m,j}}{g_{j,j}} \quad \Rightarrow \quad g_{j,j} = \sqrt{(z_j, z_j) - \sum_{m=0}^{j-1} g_{m,j}^2}.$$

Note that this can lead to breakdowns



Recursive calculation of the Hessenberg matrix $H_{i+1,i}$:

$$H_{k+1,k} = V_{k+1}^{T} A V_{k}$$

$$= V_{k+1}^{T} \underbrace{A Z_{k} G_{k}^{-1} \text{ using } Z_{k} = V_{k} G_{k}}$$

$$= \underbrace{V_{k+1}^{T} Z_{k+1} B_{k+1,k} G_{k}^{-1} \text{ using } A Z_{k} = Z_{k+1} B_{k+1,k}}_{A+1,k}$$

$$= G_{k+1} B_{k+1,k} G_{k}^{-1} \text{ using } G_{k+1} = V_{k+1}^{T} Z_{k+1}$$

$$= \begin{pmatrix} G_{k} & g_{:,k+1} \\ 0 & g_{k+1,k+1} \end{pmatrix} \begin{pmatrix} B_{k,k-1} & b_{:,k} \\ 0 & b_{k+1,k} \end{pmatrix} \begin{pmatrix} G_{k} & g_{:,k+1} \\ 0 & g_{k+1,k+1} \end{pmatrix}^{-1}$$

$$= \begin{pmatrix} G_{k} B_{k,k-1} & G_{k} b_{:,k} + g_{:,k+1} b_{k+1,k} \\ 0 & g_{k+1,k+1} b_{k+1,k} \end{pmatrix} \begin{pmatrix} G_{k-1}^{-1} & -G_{k-1}^{-1} g_{:,k} g_{k,k}^{-1} \\ 0 & g_{k,k}^{-1} \end{pmatrix}$$

$$= \begin{pmatrix} G_{k} B_{k,k-1} G_{k-1}^{-1} & \left(-G_{k} B_{k,k-1} G_{k-1}^{-1} g_{:,k} + G_{k} b_{:,k} + g_{:,k+1} b_{k+1,k} \right) g_{k,k}^{-1} \\ 0 & g_{k+1,k+1} b_{k+1,k} g_{k,k}^{-1} \end{pmatrix}$$

$$= \begin{pmatrix} H_{k,k-1} & \left(G_{k} b_{:,k} + g_{:,k+1} b_{k+1,k} - H_{k,k-1} g_{:,k} \right) g_{k,k}^{-1} \\ 0 & g_{k+1,k+1} b_{k+1,k} g_{k,k}^{-1} \end{pmatrix} H_{k,k-1} G_{k-1} = G_{k} B_{k,k-1}$$

Krylov subspace methods Pipelined GMRES with deep pipeline

```
Algorithm p(l)-GMRES
 1: r_0 \leftarrow b - Ax_0; v_0 \leftarrow r_0/||r_0||; z_0 \leftarrow v_0
 2: for i = 0, ..., m + l do
      w \leftarrow \begin{cases} (A - \sigma_i I) z_i, & i < l \\ A z_i, & i \ge l \end{cases}
                                                                                                                                           \leftarrow SpMV Az_i
         if a \ge 0 then
             g_{j,a+1} \leftarrow (g_{j,a+1} - \sum_{k=0}^{j-1} g_{k,i}g_{k,a+1})/g_{j,j}, j = a - l + 2, \dots, a
                                                                                                                                           \leftarrow scalar update G_i
             g_{a+1,a+1} \leftarrow \sqrt{g_{a+1,a+1} - \sum_{k=0}^{a} g_{k,a+1}^2}
             # Check for breakdown and restart or re-orthogonalize if necessary
             if a < l then
 9:
                 h_{j,a} \leftarrow (g_{j,a+1} + \sigma_a g_{j,a} - \sum_{k=0}^{a-1} h_{j,k} g_{k,a})/g_{a,a}, j = 0, \dots, a
10:
                                                                                                                                           \leftarrow scalar update H_{i+1,i}
                h_{a+1,a} \leftarrow q_{a+1,a+1}/q_{a,a}
             else
                h_{j,a} \leftarrow (\sum_{k=0}^{a+1-l} g_{j,k+l} h_{k,a-l} - \sum_{k=j-1}^{a-1} h_{j,k} g_{k,a})/g_{a,a}, \ j=0,\ldots,a
13:
                h_{a+1} = \leftarrow q_{a+1} + h_{a+1} + h_{a+1} = 1
14.
             end if
                                                                                                                                           \leftarrow axpv's V_{i-1}, Z_{i+1}
15:
           v_a \leftarrow (z_a - \sum_{i=0}^{a-1} g_{i,a} v_i)/g_{a,a}
16:
         z_{i+1} \leftarrow (w - \sum_{j=0}^{a-1} h_{j,a-1} z_{j+l})/h_{a,a-1} end if
17:
        g_{j,i+1} \leftarrow \begin{cases} \langle z_{i+1}, v_j \rangle, & j = 0, \dots, a \\ \langle z_{i+1}, z_j \rangle, & j = a+1, \dots, i+1 \end{cases}
                                                                                                                                           \leftarrow \mathsf{dot}\text{-}\mathsf{prs}\;(z_{i+1},v_i),
20: end for
21: y_m \leftarrow \operatorname{argmin} ||(H_{m+1,m}y_m - ||r_0||e_1)||_2
22: x \leftarrow x_0 + V_m y_m
```



Krylov subspace methods Pipelined GMRES with deep pipeline

```
Algorithm p(l)-GMRES
 1: r_0 \leftarrow b - Ax_0; v_0 \leftarrow r_0/||r_0||; z_0 \leftarrow v_0
 2: for i = 0, ..., m + l do
                                                                                                            \leftarrow SpMV Az_i
       if a \ge 0 then
          g_{j,a+1} (g_{j,a+1} - \sum_{k=0}^{j-1} g_{k,j} g_{k,a+1})/g_{j,j}, j = a - l + 2, \dots, a
 6:
                                                                                                            \leftarrow scalar update G_i
          g_{a+1,a+1} \leftarrow \sqrt{g_{a+1,a+1}} - \sum_{k=0}^{a} g_{k,a+1}^2
 7:
          # Check for breakdown and restart or re-orthogonalize if necessary
          if a < l then
 9:
10:
                                                                                                            \leftarrow scalar update H_{i+1,i}
             h_{a+1,a} \leftarrow A_{a+1,a+1} Results of dot-
12:
          else
             h_{j,a} \leftarrow \sum_{k=0}^{a+1-l} g products needed
13:
             h_{a+1,a} \leftarrow g_{a+1,a+1} / iterations later
14.
                                                                                                            \leftarrow axpv's V_{i-1}, Z_{i+1}
15:
          v_a \leftarrow z_a - \sum_{j=0}^{a-1} g_j to update G_{:,i-l+1}
16:
17:
18-
                                                                                                            \leftarrow dot-prs (z_{i+1}, v_i),
19:
21: y_m \leftarrow \operatorname{argmin} ||(H_{m+1,m}y_m - ||r_0||e_1)||_2
```

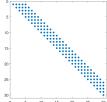
22: $x \leftarrow x_0 + V_m y_m$



Krylov subspace methods Pipelined CG with deep pipeline

In case of an SPD matrix A, the following observations simplify the algorithm:

- $ightharpoonup H_{i+1,i}$ is tridiagonal
- ► *G_i* has a (2*l* + 1)-nonzero diagonal band structure. E.g. *l* = 2:



- Cornelis et al. (2018)
- ▶ The solution x_i can be constructed using a recursively computed search direction p_i , instead of using the entire Krylov basis V_i :

$$p_i = (v_i - \delta_{i-1}p_{i-1})/\eta_i, \qquad x_i = x_{i-1} + \xi_{i-1}p_{i-1}.$$

Liesen & Strakos, Krylov Subspace Methods (2012)



Krylov subspace methods Pipelined CG with deep pipeline

```
Algorithm
                     Deep pipelined Conjugate Gradients (p(l)-CG)
 1: r_0 := b - Ax_0;
 2: v_0 := r_0/||r_0||_2;
 3: z_0 := v_0; g_{0,0} := 1;
 4: for i = 0, ..., m + l do
                                                                                                                       \leftarrow \mathsf{SpMV} \; \mathsf{Az}_i
 7.
         if a \ge 0 then
              g_{j,a+1} := (g_{j,a+1} - \sum_{k=a+1-2l}^{j-1} g_{k,j}g_{k,a+1})/g_{j,j};
                                                                                   j = a - l + 2, ..., a
              g_{a+1,a+1} := \sqrt{g_{a+1,a+1} - \sum_{k=a+1-2l}^{a} g_{k,a+1}^2};
                                                                                                                       \leftarrow scalar update G_i
              # Check for breakdown and restart if required
10:
              if a < l then
11:
                   \gamma_a := (q_{a,a+1} + \sigma_a q_{a,a} - \delta_{a-1} q_{a-1,a})/q_{a,a};
12:
13:
                   \delta_a := q_{a+1,a+1}/q_{a,a};
                                                                                                                       \leftarrow scalar update H_{i+1,i}
              else
14:
                   \gamma_a := (g_{a,a}\gamma_{a-l} + g_{a,a+1}\delta_{a-l} - \delta_{a-1}g_{a-1,a})/g_{a,a};
15:
                   \delta_a := (q_{a+1,a+1}\delta_{a-1})/q_{a,a}
16:
             end if
17:
              v_{a+1} := (z_{a+1} - \sum_{j=a-2l+1}^{a} g_{j,a+1}v_j)/g_{a+1,a+1};
                                                                                                                       \leftarrow \text{axpy's } V_{i-l+1}, Z_{i+1}
18:
19:
              z_{i+1} := (z_{i+1} - \gamma_a z_i - \delta_{a-1} z_{i-1})/\delta_a;
          end if
20 \cdot
21.
          if a < 0 then
              g_{j,i+1} := (z_{i+1}, z_j); j = 0, ..., i + 1
22:
                                                                                                                       \leftarrow dot-prs (z_{i+1}, v_i),
23.
          else
              g_{j,i+1} := \begin{cases} (z_{i+1}, v_j); & j = \max(0, i-2l+1), \dots, a+1 \\ (z_{i+1}, z_j); & j = a+2, \dots, i+1 \end{cases}
24:
```

25:



Krylov subspace methods Pipelined CG with deep pipeline

```
Deep pipelined Conjugate Gradients (p(l)-CG)
Algorithm
 1: r_0 := b - Ax_0;
 2: v_0 := r_0/||r_0||_2;
 3: z_0 := v_0; g_{0,0} := 1;
 4: for i = 0, ..., m + l do
                     (A - \sigma_i I)z_i,
                                                                                                                          C - 1/1/ / 1-
                           Az_i,
 6.
         if a \ge 0 then
 7.
             g_{j,a+1} := (g_{j,a+1} - \sum_{k=a+1-2l}^{j-1} g_{k,j}g_{k,a+1})/g_{j,j};
              g_{a+1,a+1} := \sqrt{g_{a+1,a+1} - \sum_{k=a+1-2l}^{a} g_{k,a+1}^2};
              # Check for breakdown and restart if required
10:
11:
              if a < l then
                  \gamma_a := (q_{a,a+1} + \sigma_a q_{a,a} - \delta_{a-1} q_{a-1,a})/q_{a,a};
12:
13:
                  \delta_a := q_{a+1,a+1}/q_{a,a};
             else
                                                                                                      Z0 Z1 ...
14:
                  \gamma_a := (g_{a,a}\gamma_{a-l} + g_{a,a+1}\delta_{a-l} - \delta_{a-1}g_{a-1,a})/g_{a,a};
15:
                  \delta_a := (q_{a+1,a+1}\delta_{a-1})/q_{a,a}
16:
             end if
17:
              v_{a+1} := (z_{a+1} - \sum_{j=a-2l+1}^{a} g_{j,a+1}v_j)/g_{a+1,a+1};
18:
19:
              z_{i+1} := (z_{i+1} - \gamma_a z_i - \delta_{a-1} z_{i-1})/\delta_a;
20 \cdot
         end if
21.
         if a < 0 then
             g_{j,i+1} := (z_{i+1}, z_j); j = 0, ..., i + 1
22:
23.
         else
                            (z_{i+1}, v_j); j = \max(0, i - 2l + 1), ..., a + 1
```

 $i = a + 2, \dots, i + 1$

Shorter recurrences compared to p(I)-GMRES

- computation: 2I + 2 axpy's
- storage: 3l+2 basis vectors

Fewer dot-products compared to p(I)-GMRES

• 2l + 1 band structure of G_i

(41+1,41)

 $(z_{i+1}, z_i);$

24:

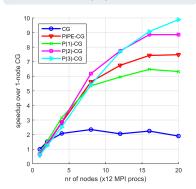
25:

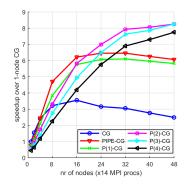


Krylov subspace methods Parallel performance of pipelined CG

Strong scaling experiments - PETSc 3.6.3/3.7.6 library - MPICH 3.1.3/3.3a2

Two 6-core Intel Xeon X5660 Nehalem 2.80 GHz per node - 2D Poisson (5pt) - 1 million unknowns Two 14-core Intel E5-2680v4 Broadwell 2.40 GHz per node - 2D Poisson (5pt) - 3 million unknowns







Krylov subspace methods Overview of pipelined Krylov methods

► Pipelined GMRES

Ghysels et al. (2013)

- depth-one pipeline: p-GMRES
- deep pipelines: $p(\ell)$ -GMRES: compute ℓ new Krylov subspace basis vectors (SpMV's) during global communication and orthogonalize after ℓ iterations.

$$V_{i-\ell+1} = [v_0, v_1, \dots, v_{i-\ell}]$$

$$Z_{i+1} = [z_0, z_1, \dots, z_{i-\ell}, \underbrace{z_{i-\ell+1}, \dots, z_{i}}_{\ell}]$$

- ► Pipelined CG
 - depth-one pipeline: p-CG
 - deep pipelines: p(ℓ)-CG
- ► Pipelined BiCGStab
- ► Pipelined Block-CG
- ► Pipelined BiCG/CGLS/LSQR
- ► Preconditioned pipelined variants
- ► Augmented/deflated/hybrid s-step pipelined methods

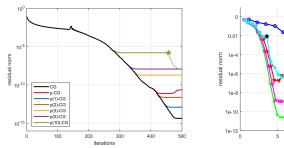
 Yamazaki et al. (2017)

- Ghysels et al. (2014)
- Cornelis et al. (2018)
- © C. & Vanroose (2017)
 - (work in progress)
 - (work in progress)



Pipelined Krylov subspace methods Numerical stability in finite precision

CPU time

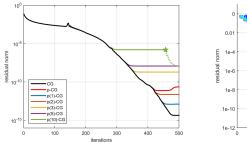


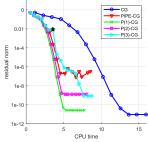
Pipelined KSM produce identical iterates to classic KSM in exact arithmetic, but finite precision computations introduce roundoff errors. This has two effects:

- 1. Delay of convergence, caused by loss of basis orthogonality
- 2. Loss of attainable accuracy, caused by local rounding errors in recurrences



Pipelined Krylov subspace methods Numerical stability in finite precision





Pipelined KSM produce identical iterates to classic KSM in exact arithmetic, but finite precision computations introduce roundoff errors. This has two effects:

- 1. Delay of convergence, caused by loss of basis orthogonality
- 2. Loss of attainable accuracy, caused by local rounding errors in recurrences

Severity of loss of attainable accuracy depends strongly on

- ▶ pipeline length I
- ▶ system size *N*
- \triangleright choice of polynomial $P_l(A)$



Rounding errors due to recurrence relations for residual and auxiliary variables:

$$\bar{x}_{i+1} = \bar{x}_i + \bar{\alpha}_i \bar{p}_i + \xi_{i+1}^x, \qquad \bar{r}_{i+1} = \bar{r}_i - \bar{\alpha}_i A \bar{p}_i + \xi_{i+1}^r.$$



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Residual deviates from the true residual $b - A\bar{x}_{i+1}$ in finite precision:

$$f_{i+1} = (b - A\bar{x}_{i+1}) - \bar{r}_{i+1}$$

$$= b - A(\bar{x}_i + \bar{\alpha}_i \bar{p}_i + \xi_{i+1}^{\times}) - (\bar{r}_i - \bar{\alpha}_i A \bar{p}_i + \xi_{i+1}^{r})$$

$$= f_i - A\xi_{i+1}^{\times} - \xi_{i+1}^{r}$$

$$= f_0 - \sum_{k=0}^{i} (A\xi_{k+1}^{\times} + \xi_{k+1}^{r}).$$
Sleijpen & van der Vorst (1995)



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$$\begin{split} f_{i+1} &= (b - A\bar{x}_{i+1}) - \bar{r}_{i+1} \\ &= b - A(\bar{x}_i + \bar{\alpha}_i \bar{p}_i + \xi_{i+1}^{\times}) - (\bar{r}_i - \bar{\alpha}_i A \bar{p}_i + \xi_{i+1}^{r}) \\ &= f_i - A\xi_{i+1}^{\times} - \xi_{i+1}^{r} \\ &= f_0 - \sum_{k=0}^{i} (A\xi_{k+1}^{\times} + \xi_{k+1}^{r}) \,. \end{split}$$
 Sleijpen & van der Vorst (1995)

Matrix notation:
$$\mathcal{F}_{i+1} = [f_0, \dots, f_i]$$
, $\Theta_i^{\mathsf{x}} = [0, \xi_1^{\mathsf{x}}, \dots, \xi_{i-1}^{\mathsf{x}}]$, $\Theta_i^{\mathsf{r}} = [f_0, \xi_1^{\mathsf{r}}, \dots, \xi_{i-1}^{\mathsf{r}}]$
 $\mathcal{F}_{i+1} = -(A\Theta_{i+1}^{\mathsf{x}} + \Theta_{i+1}^{\mathsf{r}}) U_{i+1}$,

with U_{i+1} an upper triangular matrix with all entries one.



Rounding errors due to recurrence relations for residual and auxiliary variables:

$$\bar{x}_{i+1} = \bar{x}_i + \bar{\alpha}_i \bar{p}_i + \xi_{i+1}^x, \qquad \bar{r}_{i+1} = \bar{r}_i - \bar{\alpha}_i A \bar{p}_i + \xi_{i+1}^r.$$

Residual deviates from the true residual $b - A\bar{x}_{i+1}$ in finite precision:

$$\begin{split} f_{i+1} &= (b - A\bar{x}_{i+1}) - \bar{r}_{i+1} \\ &= b - A(\bar{x}_i + \bar{\alpha}_i \bar{p}_i + \xi_{i+1}^{\times}) - (\bar{r}_i - \bar{\alpha}_i A \bar{p}_i + \xi_{i+1}^{r}) \\ &= f_i - A\xi_{i+1}^{\times} - \xi_{i+1}^{r} \\ &= f_0 - \sum_{k=0}^{i} (A\xi_{k+1}^{\times} + \xi_{k+1}^{r}) \,. \end{split}$$
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 $\mathcal{F}_{i+1} = -(A\Theta_{i+1}^x + \Theta_{i+1}^r) U_{i+1}$,

with U_{i+1} an upper triangular matrix with all entries one.

Only accumulation of local rounding errors in classical CG, no amplification.

- Gutknecht & Strakos (2000)
- avan der Vorst & Ye (2000)



Additional recurrence relations in pipelined CG all introduce local rounding errors:

$$\begin{split} \bar{x}_{i+1} &= \bar{x}_i + \bar{\alpha}_i \bar{p}_i + \xi_{i+1}^{x}, & \bar{s}_i &= \bar{w}_i + \bar{\beta}_i \bar{s}_{i-1} + \xi_i^{s}, \\ \bar{r}_{i+1} &= \bar{r}_i - \bar{\alpha}_i \bar{s}_i + \xi_{i+1}^{r}, & \bar{w}_{i+1} &= \bar{w}_i - \bar{\alpha}_i \bar{z}_i + \xi_{i+1}^{w}, \\ \bar{p}_i &= \bar{r}_i + \bar{\beta}_i \bar{p}_{i-1} + \xi_i^{p}, & \bar{z}_i &= A\bar{w}_i + \bar{\beta}_i \bar{z}_{i-1} + \xi_i^{z}, \end{split}$$



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Residual gap is coupled to the gaps on the other auxiliary variables:

$$\begin{split} &f_{j} = (b - A\bar{x}_{j}) - \bar{r}_{j} = f_{0} - \sum_{k=0}^{j-1} \bar{\alpha}_{k} g_{k} - \sum_{k=0}^{j-1} \left(A \xi_{k+1}^{x} + \xi_{k+1}^{r} \right), \\ &g_{j} = A\bar{p}_{j} - \bar{s}_{j} = \left(\prod_{k=1}^{j} \bar{\beta}_{k} \right) g_{0} + \sum_{k=1}^{j} \left(\prod_{l=k+1}^{j} \bar{\beta}_{l} \right) \left(A \xi_{k}^{p} - \xi_{k}^{s} \right) + \sum_{k=1}^{j} \left(\prod_{l=k+1}^{j} \bar{\beta}_{l} \right) h_{k}, \\ &h_{j} = A\bar{r}_{j} - \bar{w}_{j} = h_{0} - \sum_{k=0}^{j-1} \bar{\alpha}_{k} e_{k} + \sum_{k=0}^{j-1} \left(A \xi_{k+1}^{r} - \xi_{k+1}^{w} \right), \\ &e_{j} = A\bar{s}_{j} - \bar{z}_{j} = \left(\prod_{k=1}^{j} \bar{\beta}_{k} \right) e_{0} + \sum_{k=1}^{j} \left(\prod_{l=k+1}^{j} \bar{\beta}_{l} \right) \left(A \xi_{k}^{s} - \xi_{k}^{z} \right). \end{split}$$



Additional recurrence relations in pipelined CG all introduce local rounding errors:

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\bar{p}_i = \bar{r}_i + \bar{\beta}_i \bar{p}_{i-1} + \xi_i^{\mathsf{p}},
\bar{z}_i = \bar{x}_i + \bar{\beta}_i \bar{z}_{i-1} + \xi_i^{\mathsf{p}},
\bar{z}_i = \bar{x}_i + \bar{\beta}_i \bar{z}_{i-1} + \xi_i^{\mathsf{p}},
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Amplification of local rounding errors possible, depending on $\bar{\alpha}_i$'s and $\bar{\beta}_i$'s.

- Carson et al. (2017)
- © C. & Vanroose (2017)



Additional recurrence relations in pipelined CG all introduce local rounding errors:

$$\begin{split} \bar{x}_{i+1} &= \bar{x}_i + \bar{\alpha}_i \bar{p}_i + \xi_{i+1}^x, \\ \bar{r}_{i+1} &= \bar{r}_i - \bar{\alpha}_i \bar{s}_i + \xi_{i+1}^r, \\ \bar{p}_i &= \bar{r}_i + \bar{\beta}_i \bar{p}_{i-1} + \xi_i^p, \end{split}$$

$$\begin{split} \bar{s}_i &= \bar{w}_i + \bar{\beta}_i \bar{s}_{i-1} + \xi_i^s, \\ \bar{w}_{i+1} &= \bar{w}_i - \bar{\alpha}_i \bar{z}_i + \xi_{i+1}^w, \\ \bar{z}_i &= A \bar{w}_i + \bar{\beta}_i \bar{z}_{i-1} + \xi_i^z, \end{split}$$

Matrix notation:

$$\begin{split} \mathcal{F}_{j+1} &= (A\Theta_{j+1}^{x} + \Theta_{j+1}^{r}) \ U_{j+1} - \underbrace{\mathcal{G}_{j+1}}_{\mathcal{G}_{j+1}} \mathcal{A}_{j+1} \\ \underline{\mathcal{G}_{j+1}} &= (A\Theta_{j+1}^{p} + \Theta_{j+1}^{s}) \ \mathcal{B}_{j+1}^{-1} + \mathcal{H}_{j+1} \widehat{\mathcal{B}_{j+1}^{-1}} \\ \mathcal{H}_{j+1} &= (A\Theta_{j+1}^{u} + \Theta_{j+1}^{w}) \ U_{j+1} - \mathcal{E}_{j+1} \mathcal{A}_{j+1} \\ \mathcal{E}_{j+1} &= (A\Theta_{j+1}^{q} + \Theta_{j+1}^{z}) \ \mathcal{B}_{j+1}^{-1} \end{split}$$

Amplification of local rounding errors possible, depending on $\bar{\alpha}_i$'s and $\bar{\beta}_i$'s.

☐ Carson et al. (2017)

© C. & Vanroose (2017)



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\bar{z}_i = \bar{x}_i + \bar{\beta}_i \bar{z}_{i-1} + \xi_i^{\mathsf{p}},
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Amplification of local rounding errors possible, depending on $\bar{\alpha}_i$'s and $\bar{\beta}_i$'s.

- Carson et al. (2017)
- © C. & Vanroose (2017)



Additional recurrence relations in pipelined CG all introduce local rounding errors:

$$\bar{x}_{i+1} = \bar{x}_i + \bar{\alpha}_i \bar{p}_i + \xi_{i+1}^{\mathsf{x}}, \qquad \bar{s}_i = \bar{w}_i + \bar{\beta}_i \bar{s}_{i-1} + \xi_i^{\mathsf{s}},
\bar{r}_{i+1} = \bar{r}_i - \bar{\alpha}_i \bar{s}_i + \xi_{i+1}^{\mathsf{r}}, \qquad \bar{w}_{i+1} = \bar{w}_i - \bar{\alpha}_i \bar{z}_i + \xi_{i+1}^{\mathsf{w}},
\bar{p}_i = \bar{r}_i + \bar{\beta}_i \bar{p}_{i-1} + \xi_i^{\mathsf{p}}, \qquad \bar{z}_i = A \bar{w}_i + \bar{\beta}_i \bar{z}_{i-1} + \xi_i^{\mathsf{z}},$$

Matrix notation:

$$\mathcal{F}_{j+1} = (A\Theta_{j+1}^{\mathsf{x}} + \Theta_{j+1}^{\mathsf{r}}) \, U_{j+1} - \mathcal{G}_{j+1} \, (A_{j+1})$$

$$\mathcal{G}_{j+1} = (A\Theta_{j+1}^{\mathsf{x}} + \Theta_{j+1}^{\mathsf{s}}) \, (B_{j+1}^{-1}) \, \mathcal{H}_{j+1} \, (B_{j+1}^{-1}) \, (B_{j+$$

Amplification of local rounding errors possible, depending on $\bar{\alpha}_i$'s and $\bar{\beta}_i$'s.

- Carson et al. (2017)
- © C. & Vanroose (2017)



True basis vector $\bar{\mathbf{v}}_{j+1}$ satisfies:

$$Aar{V}_{j} = ar{V}_{j+1}ar{H}_{j+1,j} + (ar{V}_{j+1} - ar{V}_{j+1})ar{\Delta}_{j+1,j}$$



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$$A\bar{V}_{i} = \bar{V}_{i+1}\bar{H}_{i+1,i} + (\bar{V}_{i+1} - \bar{V}_{i+1})\bar{\Delta}_{i+1,i}$$

Computed basis vector \bar{v}_{i+1} satisfies:

$$ar{Z}_j = ar{V}_j ar{G}_j + \Theta_j^v$$



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Thus the basis vector gap $\mathcal{F}_{j+1} = oldsymbol{ar{V}}_{j+1} - ar{V}_{j+1}$ can be calculated as

$$\mathcal{F}_{j+1} = (\Theta_j^{\bar{z}} \bar{G}_j^{-1} - A \Theta_j^{\bar{v}} \bar{G}_j^{-1} + \Theta_{j+1}^{\bar{v}} \bar{B}_{j+1,j} \bar{G}_j^{-1}) \bar{\Delta}_{j+1,j}^{-1}$$



True basis vector $\bar{\mathbf{v}}_{j+1}$ satisfies:

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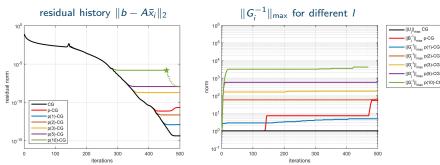
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Amplification of local rounding errors possible, depending on \bar{G}_i^{-1} .

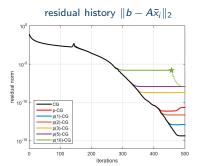
- Carson & Demmel (2014)
- C. (2018)

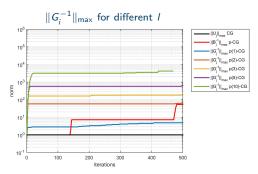




Maximum norm $\|G_i^{-1}\|_{\max}$ provides a measure for the impact of local rounding error propagation on maximal attainable accuracy in p(I)-CG







Maximum norm $\|G_i^{-1}\|_{\text{max}}$ provides a measure for the impact of local rounding error propagation on maximal attainable accuracy in p(I)-CG

Rounding error analysis explains loss of attainable accuracy with respect to

- ▶ pipeline length /
- ► system size N
- ▶ choice of polynomial $P_I(A)$



• Idea: improve accuracy by replacing \bar{r}_i , \bar{s}_i , \bar{w}_i and \bar{z}_i by their true values in selected iterations:

$$\bar{r}_{i+1} = \mathsf{fl}(b - A\bar{x}_{i+1}), \quad \bar{w}_{i+1} = \mathsf{fl}(A\bar{r}_{i+1}), \quad \bar{s}_i = \mathsf{fl}(A\bar{p}_i), \quad \bar{z}_i = \mathsf{fl}(A\bar{s}_i).$$

avan der Vorst & Ye (2000)



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$$\bar{r}_{i+1} = \mathsf{fl}(b - A\bar{x}_{i+1}), \quad \bar{w}_{i+1} = \mathsf{fl}(A\bar{r}_{i+1}), \quad \bar{s}_i = \mathsf{fl}(A\bar{p}_i), \quad \bar{z}_i = \mathsf{fl}(A\bar{s}_i).$$

- van der Vorst & Ye (2000)
- Choose when to replace based on estimate of $||f_i|| = ||(b A\bar{x}_i) \bar{r}_i||$; replacement criterion:

$$||f_{i-1}|| \le \tau ||\bar{r}_{i-1}||$$
 and $||f_i|| > \tau ||\bar{r}_i||$ with $\tau = \sqrt{\epsilon}$.

- ▶ Replace sufficiently often such that $||f_i||$ remains small
- ► Don't replace too often to limit additional computation cost of SpMV's
- ▶ Don't replace when $\|\bar{r}_i\|$ is too small, which may cause delay of convergence
- Sleijpen & van der Vorst (1996)
- Strakos & Tichy (2002)



• Idea: improve accuracy by replacing \bar{r}_i , \bar{s}_i , \bar{w}_i and \bar{z}_i by their true values in selected iterations:

$$\bar{r}_{i+1} = \mathsf{fl}(b - A\bar{x}_{i+1}), \quad \bar{w}_{i+1} = \mathsf{fl}(A\bar{r}_{i+1}), \quad \bar{s}_i = \mathsf{fl}(A\bar{p}_i), \quad \bar{z}_i = \mathsf{fl}(A\bar{s}_i).$$

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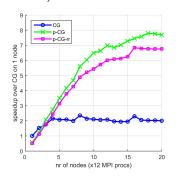
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- Sleijpen & van der Vorst (1996)
- Strakos & Tichy (2002)
- Estimate of ||f_i|| is computed at runtime (inexpensive). Accuracy is improved to comparable level as classical CG method in many cases.
 - C. & Vanroose (2017)

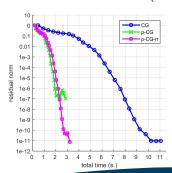


- ▶ PETSc implementation using MPICH-3.1.3 communication
- ▶ Benchmark problem: 2D Laplacian model, 1,000,000 unknowns
- ▶ System specs: 20 nodes, two 6-core Intel Xeon X5660 Nehalem 2.8GHz CPUs/node

Speedup over single-node CG (12-240 cores)



Accuracy i.f.o. total time spent (240 cores)



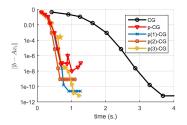


Pipelined Krylov subspace methods Numerically stable variant of p(I)-CG

Use idea similar to residual replacement to improve stability: replace recurrence for v_{i+1} by the Arnoldi relation:

Original p(I)-CG:

$$v_{j+1} = \left(z_{j+1} - \sum_{k=j-2l+1}^{j} g_{k,j+1} v_k\right) / g_{j+1,j+1}$$





Pipelined Krylov subspace methods Numerically stable variant of p(I)-CG

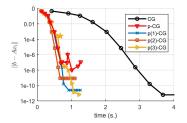
Use idea similar to residual replacement to improve stability: replace recurrence for v_{i+1} by the Arnoldi relation:

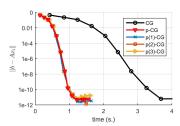
Original p(1)-CG:

$$v_{j+1} = \left(z_{j+1} - \sum_{k=j-2l+1}^{j} g_{k,j+1} v_k\right) / g_{j+1,j+1}$$

Stabilized p(I)-CG:

$$v_{j+1} = (Av_j - \gamma_j v_j - \delta_{j-1} v_{j-1})/\delta_j$$







Pipelined Krylov subspace methods Numerically stable variant of p(I)-CG

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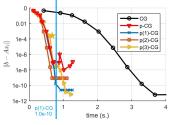
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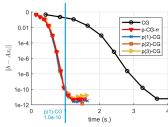
Stabilized p(1)-CG:

$$v_{j+1} = (Av_j - \gamma_j v_j - \delta_{j-1} v_{j-1})/\delta_j$$

Attainable accuracy is improved for all pipeline lengths I, but extra SpMV increases computation time

⇒ trade-off between numerical stability and performance







Conclusions and takeaways Wrap-up of this talk

- Pipelined Krylov subspace methods are a promising approach to reduce synchronization cost in linear solvers for large-scale problems
 - By adding auxiliary variables and recurrences it is possible to hide communication latency behind computational kernels
 - Deep pipelines allow to hide global reductions behind multiple SpMV's
 - Asynchronous implementation: dot-products can take multiple iterations to complete, in an overlapping manner
 - Improved scaling over classic KSMs in strong scaling limit, where global reduction latencies rise and volume of computations per core diminishes



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 - Improved scaling over classic KSMs in strong scaling limit, where global reduction latencies rise and volume of computations per core diminishes
- Finite precision behavior of communication reducing and hiding algorithms should be carefully monitored!
 - ► Rounding error analysis allows to explain observed loss of attainable accuracy
 - Residual replacement -type techniques can be applied to improve numerical stability, but at a (slight) increase in computational cost



- Work in progress (not covered in this talk):
 - ► Impact of *hard faults* & *soft-errors* on pipelined Krylov subspace methods (INRIA Bordeaux)
 - Resilience of pipelined Krylov subspace methods to system noise (UChicago/ETHZ/Rice/UIIIinois)
 - Pipelined Blocked Krylov subspace methods for systems with multiple rhs (UAntwerp/INRIA Bordeaux)
 - Pipelined Lanczos for eigenvalue calculation in graph partitioning algorithms (LBNL)
- Many interesting open problems and challenges remain as we push toward exascale-level computing!



Conclusions and takeaways Contributions to PETSc

Open source HPC linear algebra toolkit: https://www.mcs.anl.gov/petsc/



- ► KSPPGMRES: pipelined GMRES (thanks to J. Brown)
- ► KSPPIPECG: pipelined CG
- ► KSPPIPECGRR: pipelined CG with automated residual replacement
- ► KSPPIPELCG: pipelined CG with deep pipelines
- ▶ KPPPIPECR: pipelined conjugate residuals
- ► KSPGROPPCG: asynchronous CG variant by W. Gropp and collaborators
- ► KSPPIPEBCGS: pipelined BiCGStab

We are soliciting for feedback from your applications!



Conclusions and takeaways Our main publications



P. Ghysels, T. J. Ashby, K. Meerbergen, and W. Vanroose, Hiding Global Communication Latency in the GMRES Algorithm on Massively Parallel Machines SIAM J. Sci. Comput., 35(1), pp. C48C71, 2013.



P. Ghysels and W. Vanroose, Hiding global synchronization latency in the preconditioned Conjugate Gradient algorithm, Parallel Computing, 40(7), pp. 224-238, 2014,



S. Cools, E.F. Yetkin, E. Agullo, L. Giraud, W. Vanroose, Analyzing the effect of local rounding error propagation on the maximal attainable accuracy of the pipelined Conjugate Gradient method. SIAM J. on Matrix Anal. Appl., 39(1), pp. 426-450, 2018.



S. Cools, W. Vanroose, The communication-hiding pipelined BiCGStab method for the parallel solution of large unsymmetric linear systems. Parallel Computing, 65, pp. 1-20. Elsevier, 2017.



J. Cornelis, S. Cools, W. Vanroose, The communication-hiding Conjugate Gradient method with deep pipelines. Submitted to SIAM J. Sci. Comput., 2018, Preprint available at arXiv:1801.04728.



Thank you!

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