Acceleration of Source Iteration using the Dynamic Mode Decomposition

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1 Introduction

- **2** The Dynamic Mode Decomposition
- **3** DMD for Accelerating Source Iteration
- 3.1 Slab geometry examples

4 Conclusions



- In scientific computing we are used to taking a known operator and making approximations to it.
- It is possible to use the action of an operator and use just the action of the operator to generate approximations to it.
 - This is the basis for many Krylov methods.
- In this talk I will detail how we can use the action of radiation transport operators to compute the slowly converging modes in source iteration to accelerate convergence without the need for diffusion-based preconditioning.
- The basis for this work is the dynamic mode decomposition (DMD). This method can
 - Estimate time eigenvalues present in a subcritical system (Ryan G. McClarren (2019) "Calculating Time Eigenvalues of the Neutron Transport Equation with Dynamic Mode Decomposition", *Nuclear Science and Engineering*, 193:8, 854-867), and
 - Be used to produce an inexpensive reduced-order model (Zachary K. Hardy, Jim E. Morel Cory Ahrens (2019) "Dynamic Mode Decomposition for Subcritical Metal Systems", Nuclear Science and Engineering)

We begin with a sequence of vectors related by an operator.



- Consider a sequence of vectors $\{y_0, y_1, \ldots, y_K\}$ where $y_k \in \mathbb{R}^N$.
- The vectors are related by a potentially unknown linear operator of size $N \times N$, A, as

$$y_{k+1} = A y_k$$

• If we construct the $N \times K$ data matrices Y_+ and Y_- ,

$$Y_{+} = \begin{pmatrix} | & | & & | \\ y_{1} & y_{2} & \dots & y_{K} \\ | & | & & | \end{pmatrix} \qquad Y_{-} = \begin{pmatrix} | & | & & | \\ y_{0} & y_{1} & \dots & y_{K-1} \\ | & | & & | \end{pmatrix}$$

we can write

$$Y_+ = AY_-.$$

- At this point we only need to know the data vectors y_k, they could come from a calculation, measurement, etc.
- As $K \to \infty$ we could hope to infer properties about A.



• We take the thin singular value decomposition (SVD) of Y_{-} to write

$$Y_{-}=U\Sigma V^{\mathrm{T}},$$

where U is a $N \times K$ orthogonal matrix, Σ is a diagonal $K \times K$ matrix with non-negative entries on the diagonal, and V is a $K \times K$ orthogonal matrix.

- The SVD requires $O(NK^2)$ operations to compute.
- Later, we will want $K \ll N$, if, for example, N is the number of unknowns in a transport calculation.
- Also, if the column rank of $Y_- < K$, then there is a further reduction in the SVD size.
- The matrix U has columns that forms an orthonormal basis for the row space of $Y_{-} \subset \mathbb{R}^{N}$.
- Using the SVD we get

$$Y_+ = AU\Sigma V^{\mathrm{T}}.$$

 If there are only r < K non-zero singular values in Σ, we use the compact SVD where U is N × r, Σ is r × r, and V is K × K.



 \bullet We can rearrange the relationship between Y_+ and Y_- to be

$$Y_+ = AU\Sigma V^{\mathrm{T}} \qquad \rightarrow \qquad U^{\mathrm{T}}AU = U^{\mathrm{T}}Y_+V\Sigma^{-1}.$$

- Define $\tilde{A} = U^{T}AU = U^{T}Y_{+}V\Sigma^{-1}$. This is a rank K approximation to A.
- Using the approximate operator \tilde{A} , we can now find out information about A.
- The eigenvalues/vectors of \tilde{A} ,

$$\tilde{A}w = \lambda w$$
,

are used to define the dynamic modes of A:

$$arphi = rac{1}{\lambda} U^{\mathrm{T}} Y_{+} V \Sigma^{-1} w.$$

• The dynamic mode decomposition (DMD) of the data matrix Y_+ is then the decomposition of into vectors φ . The mode with the largest norm of λ is said to be the dominant mode.

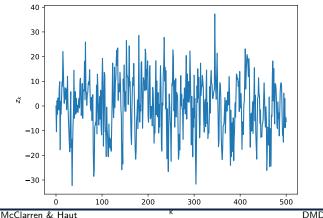


• Consider the sequence

$$\mathbf{z}_{k+1} = \mathbf{a}\mathbf{z}_k + \mathbf{n}_k,$$

where a = 0.5, and $n_k \sim \mathcal{N}(0, 10^2)$.

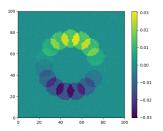
• Using K = 500, we estimate a = 0.506552 from the data below.



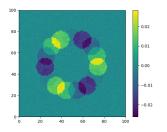


- Left: Data generated by moving a circle in a periodic motion with added noise. The data has two periods of motion.
- Right: Reconstruction generated by approximating \tilde{A} using one period of frames and starting from frame 1.

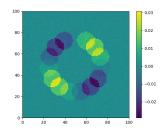
The DMD modes of the inferred operator are what we would expect.



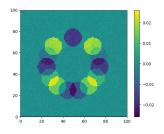
Dominant DMD mode: $U\varphi_1$



Third DMD mode: $U\varphi_3$



Second DMD mode: $U\varphi_2$



Fourth DMD mode: $U\varphi_4$



- The discrete ordinates method for transport is typically solved using source iteration (Richardson iteration) and diffusion-based preconditioning/acceleration.
- Source iterations converge quickly for problems with a small amount of particle scattering.
- For strongly scattering media, the transport operator has a near nullspace that can be handled using a diffusion preconditioner.
- However, the question of efficiently preconditioning/accelerating transport calculation on high-order meshes with discontinuous fine elements is an open area of research.
- The approximate operator found from DMD can be used to remove this same near nullspace and improve iterative convergence without the need for a separate preconditioner or diffusion discretization/solve.

We consider source iteration for a simple transport problem.

- ¢<mark>**</mark>3
- The steady, single group transport equation with isotropic scattering can be written as

$$L\psi=\frac{c}{4\pi}\phi+\frac{Q}{4\pi},$$

where \boldsymbol{c} is the scattering ratio, \boldsymbol{Q} is a prescribed source, and the streaming and removal operator is

$$L = (\Omega \cdot \nabla + 1)$$
.

• $\psi(\mathbf{x}, \Omega)$, $\Omega \in \mathbb{S}_2$,

$$\phi(\mathbf{x}) = \int_{4\pi} \psi \, d\Omega = \langle \psi \rangle.$$

• Source iteration solves this problem using the iteration strategy

$$\phi^{\ell} = \left\langle L^{-1} \left(\frac{c}{4\pi} \phi^{\ell-1} + \frac{Q}{4\pi} \right) \right\rangle,$$

where ℓ is an iteration index.

- One iteration is often called a "transport sweep".
- A benefit of source iteration is that the angular flux, ψ does not have to be stored.
- As $c \rightarrow 1$, the convergence of source iteration can be arbitrarily slow.

We can write source iteration as a sequence of vectors related by a linear operator.

• Rearranging the transport equation we see that source iteration is an iterative procedure for solving

$$\phi - \left\langle L^{-1} \frac{c}{4\pi} \phi \right\rangle = L^{-1} Q,$$

or

$$(I-A)\phi=b.$$

• Therefore, the source iteration vectors are

$$\phi^{\ell+1} = A\phi^\ell + b,$$

or

$$\phi^{\ell+1} - \phi^{\ell} = A(\phi^{\ell} - \phi^{\ell-1})$$

• Therefore, we can cast the difference between iterates in a form that is amenable to the approximation of A using DMD, $Y_+ = AY_-$,

$$Y_{+} = \left[\phi^{2} - \phi^{1}, \phi^{3} - \phi^{2}, \dots, \phi^{K} - \phi^{K-1}\right],$$
$$Y_{-} = \left[\phi^{1} - \phi^{0}, \phi^{2} - \phi^{3}, \dots, \phi^{K-1} - \phi^{K-2}\right]$$



Source iteration can be accelerated by taking several iterates and approximating the solution as $\ell\to\infty$



• As before we define an approximate A as the $K \times K$ matrix:

$$\tilde{A} = U^{\mathrm{T}} A U = U^{\mathrm{T}} Y_{+} V \Sigma^{-1},$$

• We can use \tilde{A} to construct the operator $(I - \tilde{A})^{-1}$ and use this to approximate the solution:

$$(I - A)(\phi - \phi^{K-1}) = b - (I - A)\phi^{K-1}$$

= $b - \phi^{K-1} + (\phi^{K} - b)$
= $\phi^{K} - \phi^{K-1}$.

• The difference $\phi - \phi^{K-1}$ is the difference between step K-1 and the converged answer. We define a new vector Δy as the length K vector that satisfies

$$\phi - \phi^{K-1} = U \Delta y. \tag{1}$$

• We then substitute and multiply by U^{T} to get

$$(I - \tilde{A})\Delta y = U^{\mathrm{T}}(\phi^{K} - \phi^{K-1}).$$
⁽²⁾

This is a linear system of size K that we can solve to get Δy and then compute the update to ϕ^{K-1} as

$$\phi \approx \phi^{K-1} + U\Delta y. \tag{3}$$

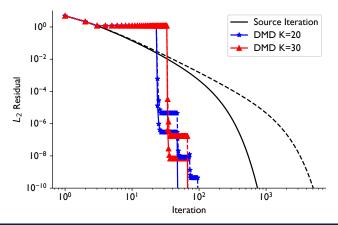


- The algorithm is as follows
 - **()** Perform *R* source iterations: $\phi^{\ell} = A\phi^{\ell-1} + b$.
 - **2** Compute K source iterations to form Y_+ and Y_- . The last column of Y_- we call ϕ^{K-1} .
 - **3** Compute $\phi = \phi^{K-1} + U\Delta y$ as above.
- Each pass of the algorithm requires R + K source iterations.
- The *R* source iterations are used to correct any errors caused by the approximation of *A* using the SVD.
- It is easiest to assess convergence between the source iterations.
- This works regardless of the spatial discretization used.
- Other algorithms are possible:
 - Rather than extrapolate to an infinite number of iterations, we can use \tilde{A} to approximate a finite number of source iterations.
 - We could use a coarsened vector $\bar{\phi}$ in the DMD procedure to reduce the memory/computational cost.

DMD works perfectly on a homogenous slab, the ur-demonstration problem for acceleration schemes.



- We consider a slab with vacuum boundaries and a scattering ratio of c = 0.99 and 1.0 and 400 spatial zones, S_8 angular discretization, and the diamond difference spatial discretization.
- Solid lines are c = 0.99 results and dashed lines are c = 1.0





• On the same problem set up, the number of iterations to converge is shown below.

K/c	0.1	0.5	0.9	0.99	0.999	0.9999	0.99999	0.999999
3	8	15	39	70	70	70	70	70
5	10	11	28	90	90	90	90	90
10	15	15	29	60	140	140	140	140
20	25	25	25	49	74	76	76	76
50	55	55	55	56	57	57	57	57
SI	6	17	89	637	2439	3681	3889	3911

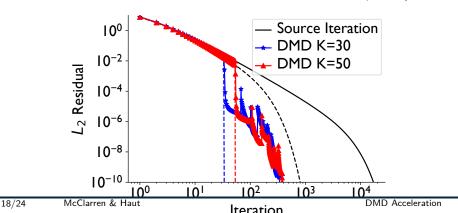
Performance does degrade on an insanely heterogeneous problem.



- We consider a problem with vacuum boundaries, 1000 cells, unit domain length, with c=0.9999 and

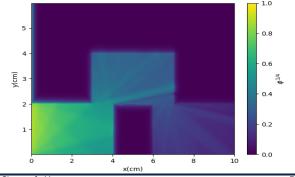
$$\sigma_{\rm t} = \begin{cases} 2^{p} & \text{cell number odd} \\ 2^{-p} & \text{cell number even} \end{cases}$$

• Below we see convergence for p = 5 (dashed) and p = 8 (solid), a factor of about 1000 and 6.5×10^4 between thick and thin cells, respectively.





- We solve a linear, xy-geometry version of the crooked pipe problem where all materials have a scattering ratio of 0.988 (to simulate a realistic sized time step).
- The density ratio between the tick and thin material is 1000.
- Problem solved using fully lumped, bilinear discontinous Galerkin in space and S_8 product quadrature.



The number of iterations required increases slowly with mesh refinement.



• The number of iterations for source iteration and DMD-accelerated calculations with K = 10 and R = 3.

$(N_x \times N_y)$	DMD	SI
25 imes15	53	811
50 imes25	52	873
100 imes 60	78	974
150 imes90	91	∞_{RML}
200 imes 120	104	∞_{RML}

 $\infty_{\text{RML}} =$ functionally infinite on my laptop.

• The increase seems to be the resolution to the 1/2 power (square root of the number of cells per dimension).

This can work for power iteration as well.



A recent paper, Roberts, Jeremy A., et al. "Acceleration of the Power Method with Dynamic Mode Decomposition." arXiv preprint arXiv:1904.09493 (2019), uses these ideas for power iterations. Here is Fig. 3 from that paper:

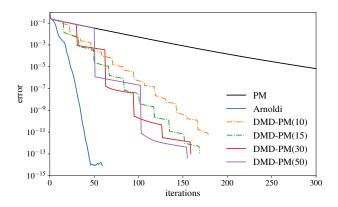


Fig. 3: The error in the predicted eigenmode for DMD-PM(n), where n is the number of power iterations performed. Errors are also included for the power method (PM) and Arnoldi's method.



- We could use DMD acceleration to compute a low-order transport acceleration (the so-called TSA method). In this case the we would use low-order in angle transport sweeps to estimate the slowly converging modes.
- Additionally, it is possible to estimate \tilde{A} using independently generated vectors. This would enable the Y_{\pm} matrices to be generated using sweeps computed in parallel.
- The big win could be from applying this to other iterative components:
 - Energy group iterations
 - Temperature iterations in radiative transfer.
- The performance of DMD on meshes with cycles is also a possible impact area.



- Using a DMD approach to compute approximate operators gives one the ability to
 - Estimate eigenvalues for the system, and
 - Accelerate calculations.
- There is much further research to be done, but progress is exciting.



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