#### Dynamic Mode Decomposition Discussion

Zihao Wang, Sam Minkowicz, April Zhou, Zhongsheng Sang, Avinash Karamchandani February 2020

# 1. Explain equations (and the relationship between equations) 20.1.5, 20.1.6, and 20.1.10. What do these equations assume about the data and what do they capture?

$$\mathbf{x}_{i+1} = \mathbf{A}\mathbf{x}_i,\tag{20.1.5}$$

where **A** is linear, time-independent Koopman operator,  $x_j$  and  $x_{j+1}$  are vectors of data collected at times  $t_j$  and  $t_{j+1}$ . The matrix

$$\mathbf{X}_1^{M-1} = \begin{bmatrix} \mathbf{x}_1 & \mathbf{x}_2 & \mathbf{x}_3 & \cdots & \mathbf{x}_{M-1} \end{bmatrix}$$
 (20.1.6)

is constructed the vector data collected at different times ( $\mathbf{x}_j$  for j=1:M-1). With the assumption of equation (20.1.5), we can deduce the equation (20.1.6) to the equation (20.1.10). Actually equation (20.1.10) gives us a recursion formula for the next time step.

$$\mathbf{A}\mathbf{X}_1^{M-1} = \mathbf{X}_2^M \tag{20.1.10}$$

#### 2. What is the relationship between 20.1.7, 20.1.8, and 20.1.11?

Making use of equation (20.1.5), we are able to rewrite  $\mathbf{X}_1^{M-1}$  as follows:

$$\mathbf{X}_{1}^{M-1} = [\mathbf{x}_{1} \ \mathbf{A}\mathbf{x}_{1} \ \mathbf{A}^{2}\mathbf{x}_{1} \ \dots \ \mathbf{A}^{M-2}\mathbf{x}_{1}]. \tag{20.1.7}$$

We try to represent the final data point  $\mathbf{x}_M$  as a linear combination of the previous M-1 data points, as best as possible (in an L2 sense):

$$\mathbf{x_M} = \sum_{m=1}^{M-1} b_m x_m + r, \tag{20.1.8}$$

where  $b_m$  are the coefficients of the Krylov space vectors and  $\mathbf{r}$  is the residual term (which is orthogonal to the Krylov space). Making use of equation (20.1.8), equation (20.1.10) can be rewritten as

$$\mathbf{X}_{2}^{M} = \mathbf{X}_{1}^{M-1}\mathbf{S} + \mathbf{r}e_{M-1}^{*} \tag{20.1.11}$$

where  $e_{M-1}^*$  is the (M-1)th unit vector and **S** is the coefficient matrix.

### 3. If $x_k$ is a snapshot of state-space at time k and N spatial points, what are the sizes of $\mathbf{X}_1^{M-1}$ , $\mathbf{X}_2^M$ , r, and S?

 $\mathbf{X}_1^{M-1}$  and  $\mathbf{X}_2^M$  are  $N \times (M-1)$  matrices,  $\mathbf{r}$  is a  $N \times 1$  vector and  $\mathbf{S}$  is a  $(M-1) \times (M-1)$  matrix.

#### 4. What is the relationship between S and A?

Based on the linearity assumption and Krylov space method, we rewrite the relation between  $\mathbf{X}_2^M$  and  $\mathbf{X}_1^{M-1}$  as equation (20.1.11). In this way, the eigenvalues of  $\mathbf{S}$  approximate some of the eigenvalues of the unknown Koopman operator  $\mathbf{A}$ .

Comparing 20.1.10 and 20.1.11 we see that **A** is being right multiplied while **S** is being left multiplied therefore **A** is  $N \times N$  and **S** is a  $(M-1) \times (M-1)$ . In fluids applications the data is of much higher spatial resolution than temporal resolution, N >> M. Thus, **S** will be smaller than **A**.

# 5. What are the characteristics of the SVD decomposition in 20.1.9? (i.e. what are the structure, dimensions, and orthogonal properties of the matrices U, $\Sigma$ , and V?

$$\mathbf{X}_{1}^{M-1} = \mathbf{U}\boldsymbol{\Sigma}\mathbf{V}^{*} \tag{20.1.9}$$

U is an  $N \times K$  matrix, and V is an  $M-1 \times K$  matrix,  $\Sigma$  is an  $K \times K$  rectangular diagonal matrix with non-negative real numbers on the diagonal. The diagonal entries of  $\Sigma$  are known as the singular values of  $X_1^{M-1}$ . The columns of U and V are called the left-singular vector and right-singular vector of  $X_1^{M-1}$ , respectively.

6. The textbook skips some steps in deriving 20.1.13. Show that you can get to this approximation of  $\tilde{S} = U^*SU$  and the SVD to solve 20.1.10 for the least-squares solution.

From equations (20.1.10) and (20.1.9), we have that

$$\begin{aligned} \mathbf{A} &= & \mathbf{X}_2^M (\mathbf{X}_1^{M-1})^{-1} \\ &= & \mathbf{X}_2^M (\mathbf{U} \boldsymbol{\Sigma} \mathbf{V}^*)^{-1} \\ &= & \mathbf{X}_2^M \mathbf{V} \boldsymbol{\Sigma}^{-1} \mathbf{U}^*. \end{aligned}$$

Let  $\tilde{\mathbf{S}} = \mathbf{U}^* \mathbf{A} \mathbf{U}$  be the similarity transformation of  $\mathbf{A}$ . Then, A can be approximated by

$$\tilde{\mathbf{S}} = \mathbf{U}^* \mathbf{A} \mathbf{U}$$

$$= \mathbf{U}^* (\mathbf{X}_2^M \mathbf{V} \mathbf{\Sigma}^{-1} \mathbf{U}^*) \mathbf{U}$$

$$= \mathbf{U}^* \mathbf{X}_2^M \mathbf{V} \mathbf{\Sigma}^{-1}.$$
(20.1.13)

Eigenvlaues and eigenvectors are preserved under a similarity transform.

### 7. How are eigenvectors and eigenvalues used to estimate the DMD solution?

We solve the eigenvalue problem:

$$\tilde{\mathbf{S}}\mathbf{y}_k = \mu_k \mathbf{y}_k \quad \text{for} \quad k = 1, 2, ..., K$$
 (20.1.14)

Where K is the rank of the approximation we make  $(K < rank(\mathbf{X}^M))$ . We then use the eigenvalues  $\mu_k$  to construct the DMD modes

$$\psi_k = \mathbf{U}\mathbf{y}_k. \tag{20.1.15}$$

Finally the approximate solution at all future times is given by

$$\mathbf{x}_{DMD}(t) = \sum_{k=1}^{K} b_k(0)\psi_k(\mathbf{x}) \exp(\omega t) = \mathbf{\Psi} \operatorname{diag}(\exp(\omega t))\mathbf{b}$$
 (20.1.16)

where  $\omega_k = \ln(\mu_k)/\Delta t$ ,  $b_k(0)$  is the initial amplitude of each mode,  $\mathbf{\Phi}$  is the matrix whose columns are the eigenvectors  $\phi$ , diag( $\omega t$ ) is a diagonal matrix whose entries are the eigenvalues  $\exp(\omega_k t)$ 

#### 8. How do you calculate the initial condition for the solution?

Evaluating the approximate solution at the first time point gives  $x_1 = \Psi b$ . Since  $\Psi$  contains a truncated set of the eigenvectors of A, it will not be of full rank. To determine b, we use a psuedo-inverse:  $b = \Psi^{\dagger} x_1$ .

## 9. What makes the solution unstable and when does it become unstable?

Since approximate solution takes the form

$$\mathbf{x}_{DMD}(t) = \sum_{k=1}^{K} b_k(0) \psi_k(\mathbf{x}) \exp(\omega_k t),$$

for the solution to be stable, we need  $\omega_k < 0$  for all k. And since  $\omega_k = \ln |\mu_k|/\Delta t$ , this means  $|\mu_k| < 1$ .