

# **ME493 - Methods of Data-Driven Control**

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# Table of contents

<b>Preface</b>	<b>3</b>
<b>1 The Four Fundamental Subspaces</b>	<b>4</b>
<b>2 Singular Value Decomposition</b>	<b>6</b>
2.1 Overview . . . . .	6
2.2 POD Analysis of Low Reynolds Number Pitching Airfoil DNS . . . . .	6
<b>3 Data-Driven Dynamical Systems</b>	<b>12</b>
3.1 Overview: Dynamic Mode Decomposition . . . . .	12
3.2 DMD Analysis of Low Reynolds Number Pitching Airfoil DNS . . . . .	13
3.3 Overview: Sparse Identification of Nonlinear Dynamics . . . . .	15
3.4 SINDy Implementation on Temporal Amplitudes . . . . .	16
<b>4 Model Reduction and Projection</b>	<b>18</b>
4.1 Overview . . . . .	18
<b>5 System Identification Techniques</b>	<b>19</b>
5.1 Overview: Eigensystem Realization Algorithm . . . . .	19
5.2 Overview: DMD with Control . . . . .	19
<b>References</b>	<b>20</b>

# Preface

Documentation of the independent study ME493: Methods of Data-Driven Control, taken during Spring 2024.

**Instructor:** Dirk M. Luchtenburg **Co-conspirators:** Sohaib Bhatti, Khushant Khurana, Eunky (Q) Kim, Sonam Okuda **Email:** dirk.luchtenburg@cooper.edu

**Description:** This independent study provides an introduction to state-of-the-art methods employed in the field of data-driven engineering. Data-driven methods such as the proper orthogonal decomposition and the dynamic mode decomposition are used in a variety of fields including fluid dynamics, climate analysis, mixing problems, the study of infectious diseases, etc. Data-driven methods rely heavily on concepts from linear algebra, calculus, probability, and statistics. We will review these essential elements and develop a hands-on understanding of a selection of data-driven methods. The study culminates with an application of some method(s) to a practical problem which is to be carefully detailed in a technical report.

Referenced texts include the following:

- Strang (2019)
- Brunton and Kutz (2022)
- Boyd and Vandenberghe (2018)
- Géron (2022)
- Sutton and Barto (2018)

# 1 The Four Fundamental Subspaces

In the Strang-ian view of linear algebra, an  $m$  by  $n$  matrix  $\mathbf{A}$  is associated with four fundamental subspaces - two of  $\mathbb{R}^m$  and two of  $\mathbb{R}^n$ . It's easiest to illustrate this using an example matrix:

$$\mathbf{A} = \begin{bmatrix} 1 & 2 & 3 \\ 4 & 5 & 6 \end{bmatrix} = \begin{bmatrix} \mathbf{r}_1 \\ \mathbf{r}_2 \end{bmatrix} = [\mathbf{c}_1 \quad \mathbf{c}_2 \quad \mathbf{c}_3]$$

where:

$$\mathbf{r}_1 = [1 \quad 2 \quad 3] \quad \mathbf{r}_2 = [4 \quad 5 \quad 6]$$

$$\mathbf{c}_1 = [1 \quad 4]^T \quad \mathbf{c}_2 = [2 \quad 5]^T \quad \mathbf{c}_3 = [3 \quad 6]^T$$

## 💡 Column Space

The column space (or range)  $R(\mathbf{A})$  contains all linear combinations of the column vectors of  $\mathbf{A}$ .

It only takes two linearly independent vectors to span  $\mathbb{R}^2$ , and we have three! Our column space is  $\mathbb{R}^2$ .

## 💡 Row Space

The row space  $R(\mathbf{A}^T)$  contains all linear combinations of the column vectors of  $\mathbf{A}^T$  (or equivalently, the row vectors of  $\mathbf{A}$ ).

Transposed row vectors  $\mathbf{r}_1^T$  and  $\mathbf{r}_2^T$  span the following plane:

$$\left\{ \begin{bmatrix} x_1 + 4x_2 \\ 2x_1 + 5x_2 \\ 3x_1 + 6x_2 \end{bmatrix} : \mathbf{x} \in \mathbb{R}^2 \right\}$$

### 💡 Null Space

The null space  $N(\mathbf{A})$  contains all solutions  $\mathbf{u}$  to  $\mathbf{A}\mathbf{u} = \mathbf{0}$ .

Solving this equation yields the nontrivial solution:

$$\mathbf{u} = \begin{bmatrix} 1 \\ -2 \\ 1 \end{bmatrix}$$

Our null space is the span of  $\mathbf{u}$ , or the following line:

$$\left\{ \begin{bmatrix} x \\ -2x \\ x \end{bmatrix} : x \in \mathbb{R} \right\}$$

### 💡 Left Null Space

The left null space  $N(\mathbf{A}^T)$  contains all solutions  $\mathbf{v}$  to  $\mathbf{A}^T\mathbf{v} = \mathbf{0}$ .

This equation has no nontrivial solutions, so the left null space is the zero subspace.

The “big picture of linear algebra,” as Gil Strang puts it, is that for an  $m$  by  $n$  matrix  $\mathbf{A}$ :

- The column space/range  $R(\mathbf{A})$  is perpendicular to the left null space  $N(\mathbf{A}^T)$  in  $\mathbb{R}^m$
- The row space  $R(\mathbf{A}^T)$  is perpendicular to the null space  $N(\mathbf{A})$  in  $\mathbb{R}^n$

### 💡 Rank

The rank  $r$  of a matrix  $\mathbf{A}$  is the number of independent rows/columns, i.e., the row space and column space/range have the same dimension  $r$ .

- The dimension of the null space  $N(\mathbf{A})$  is  $n - r$  and the dimension of the left null space  $N(\mathbf{A}^T)$  is  $m - r$ .

## 2 Singular Value Decomposition

### 2.1 Overview

The singular value decomposition (SVD) is a fundamental matrix factorization with numerous applications in data analysis and scientific computing. Mathematically, the SVD of an  $m \times n$  matrix  $\mathbf{X}$  is a factorization of the form:

$$\mathbf{X} = \mathbf{U}\mathbf{\Sigma}\mathbf{V}^*$$

where  $\mathbf{U}$  is an  $m \times m$  orthogonal matrix,  $\mathbf{\Sigma}$  is an  $m \times n$  diagonal matrix with non-negative real numbers on the diagonal, and  $\mathbf{V}$  is an  $n \times n$  orthogonal matrix.

The SVD is particularly useful for analyzing large, high-dimensional datasets that can be well-approximated by matrices of much lower rank. By extracting the dominant patterns in the data, the SVD enables efficient dimensionality reduction, noise removal, and data compression. It is the foundation of techniques like principal component analysis (PCA) and is widely applied in fields such as signal processing, machine learning, and image analysis.

Proper orthogonal decomposition (POD) modes are a set of orthogonal basis functions that optimally represent a given dataset in a least-squares sense. They are obtained by performing an SVD on a data matrix. POD modes form an orthonormal basis, meaning the modes are mutually orthogonal and have unit norm.

The modes are ranked by their energy content, with the first mode capturing the most energy and subsequent modes capturing progressively less energy.

### 2.2 POD Analysis of Low Reynolds Number Pitching Airfoil DNS

The [dataset](#) analyzed contains direct numerical simulations (DNS) of two-dimensional stationary and pitching flat-plate airfoils at a Reynolds number of 100. The dataset includes time-resolved snapshots of the velocity field, lift and drag coefficients, and airfoil kinematics spanning 40-100 convective time units. The cases consist of a stationary airfoil and eight different pitching frequencies. This dataset is part of a database intended to aid in the conception, training, demonstration, evaluation, and comparison of reduced-complexity models for fluid mechanics, created by Aaron Towne and Scott Dawson.

The dataset also includes a MATLAB function that provides a simple implementation of Dynamic Mode Decomposition (DMD), a data-driven method we'll get to later.

To analyze the DNS data using POD, I first extracted the velocity components from the provided snapshots. I then computed the mean-corrected snapshots by subtracting the mean of each snapshot and arranged them into a matrix  $\mathbf{X}$ .

Next, I performed an economy-sized SVD on  $\mathbf{X}$ . The squared singular values are plotted here, as a function of the mode index.

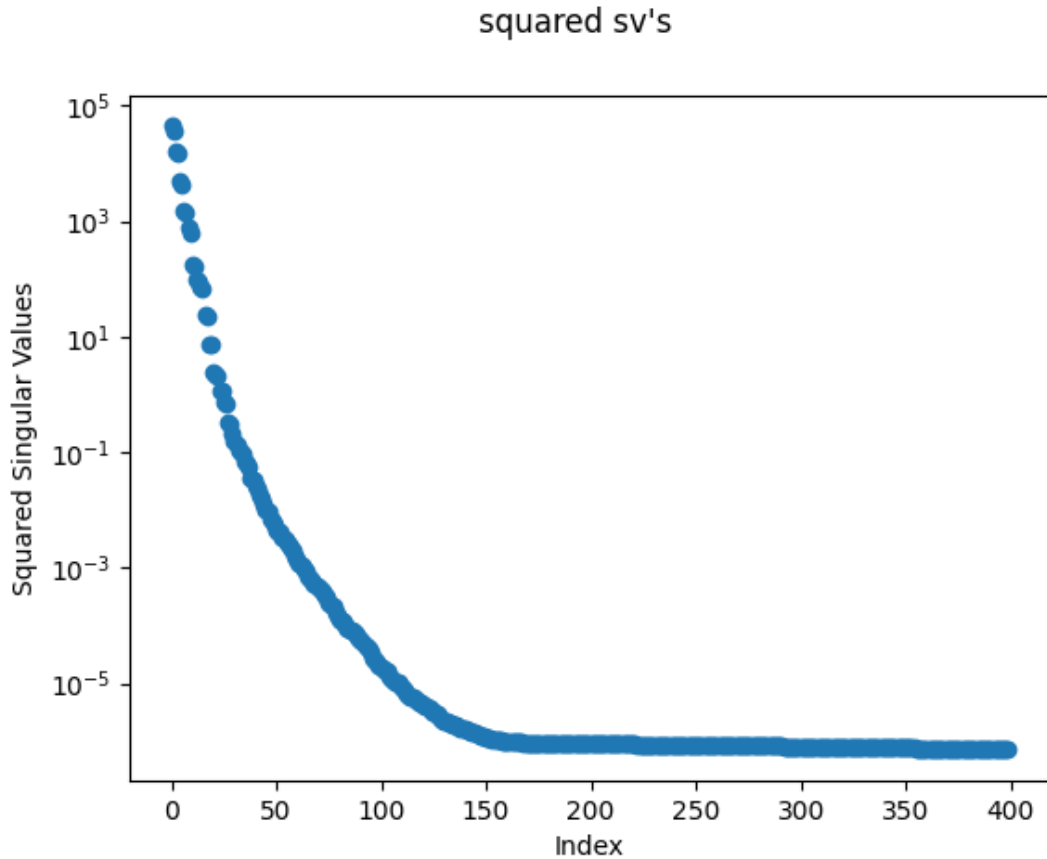


Figure 2.1: squared\_sv

This plot reveals how many individual POD basis vectors there are. The rapid decay of the singular values indicates that the flow is well-approximated by a low-rank subspace, with the first 16 modes capturing the majority of the energy.

Visualizing the first six POD modes for both  $u_x$  and  $u_y$  revealed the spatial structure of the dominant flow patterns. The oscillatory modes have a characteristic wavelength that can be estimated from the spatial distribution of the mode amplitudes.

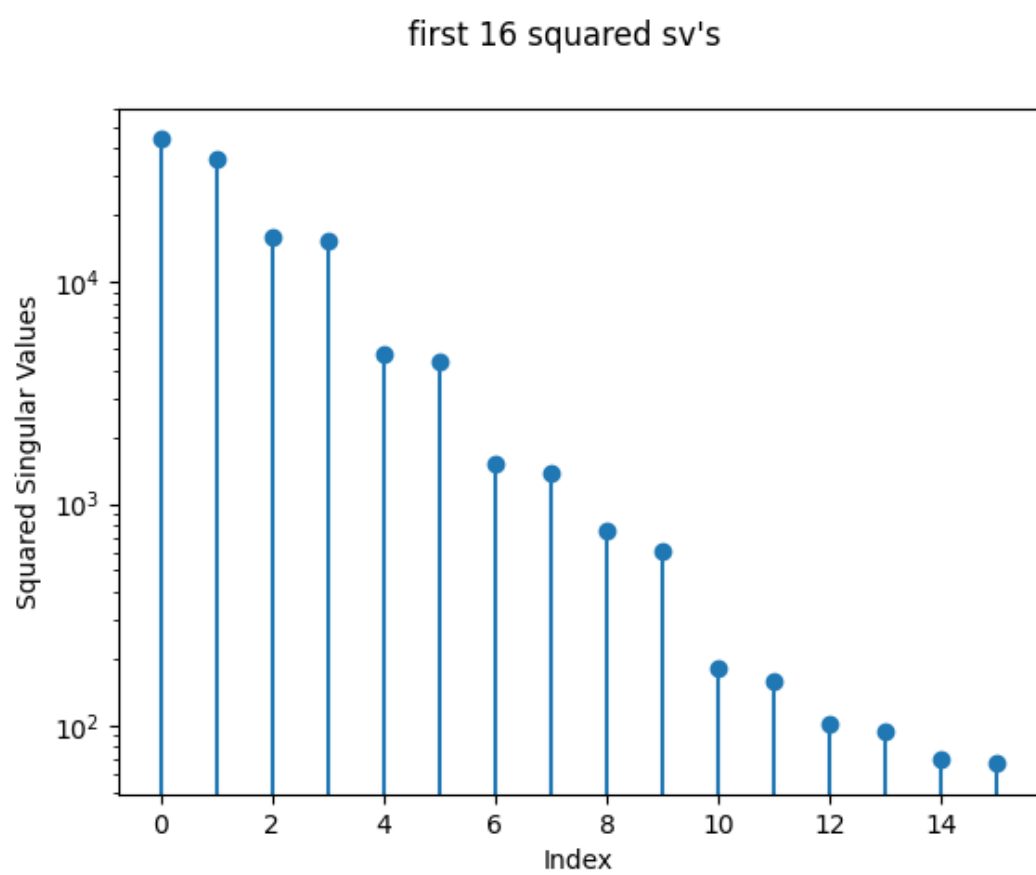


Figure 2.2: squared\_sv\_truncated



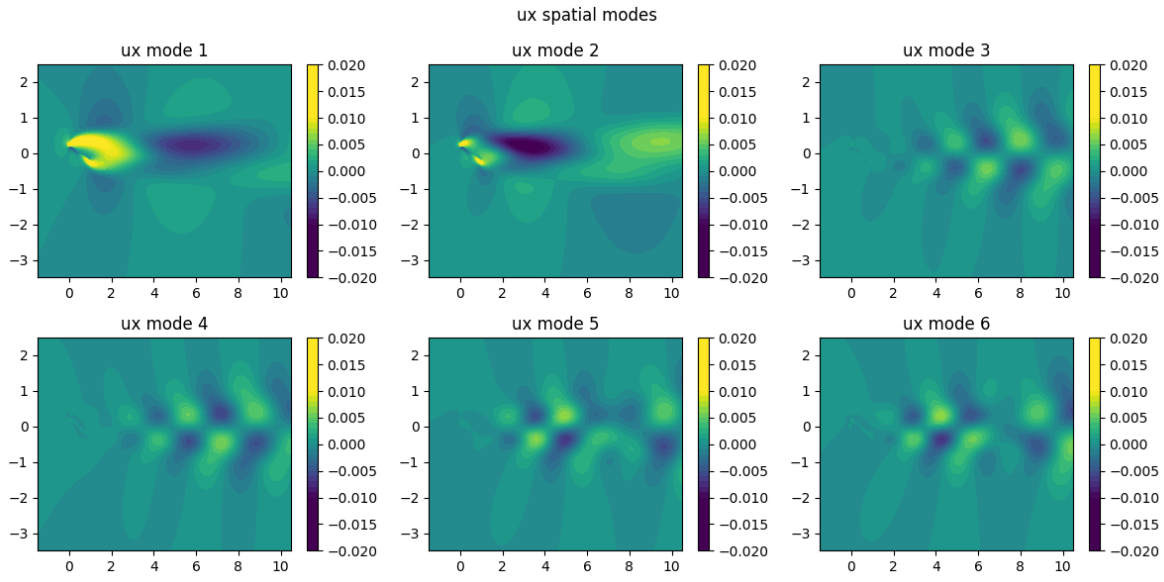


Figure 2.3:  $u_x$  spatial modes

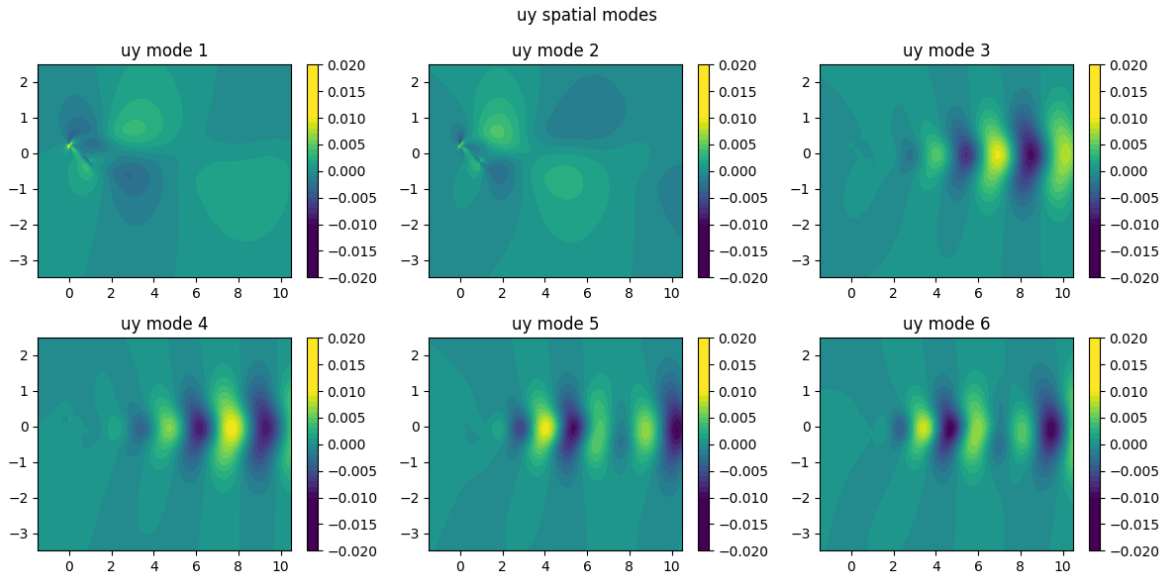


Figure 2.4:  $u_y$  spatial modes

Here are the temporal amplitudes associated with those spatial modes:

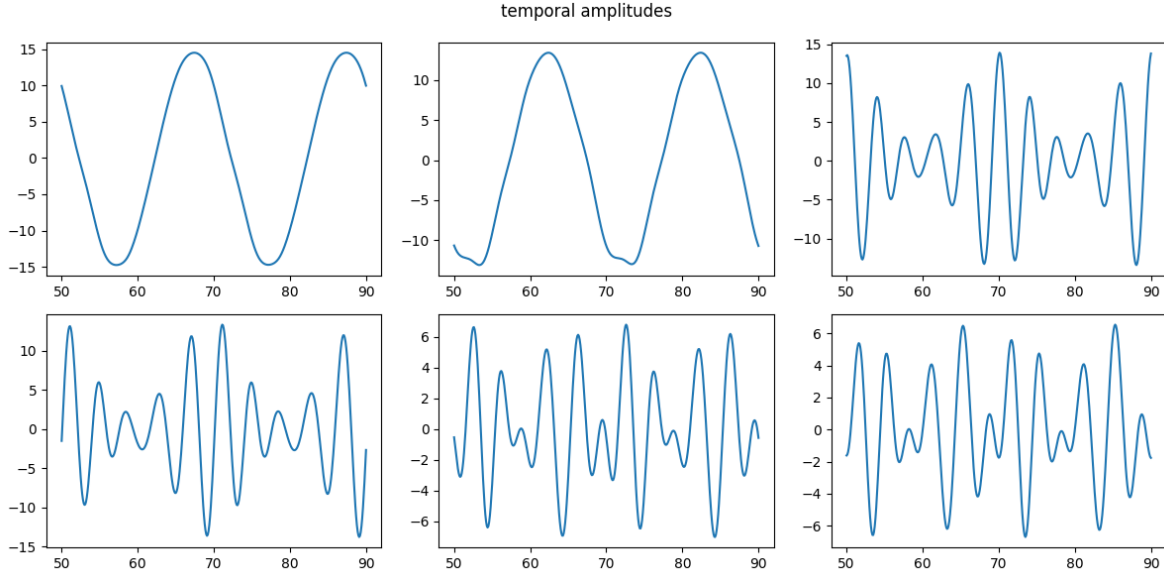


Figure 2.5: temporal\_amplitudes

Finally, I reconstructed the snapshots using a rank-4 approximation, which captures the most energetic flow structures. Comparing the reconstructed snapshots with the original data showed that the low-rank approximation successfully recovers the essential flow physics, such as the presence of coherent structures and the overall flow patterns.

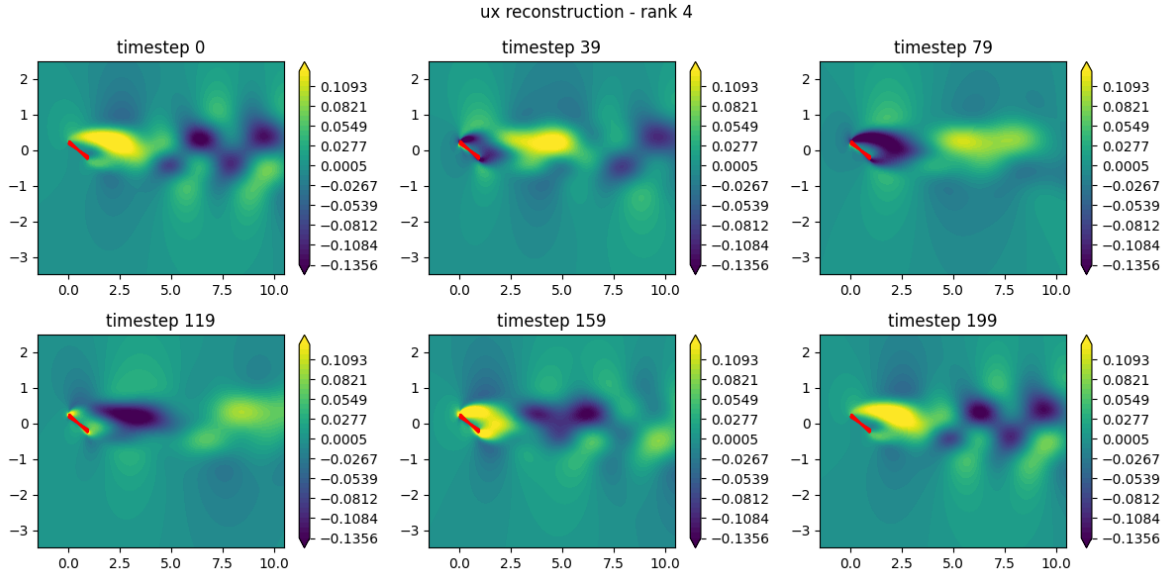


Figure 2.6:  $u_x$ \_reconstruction

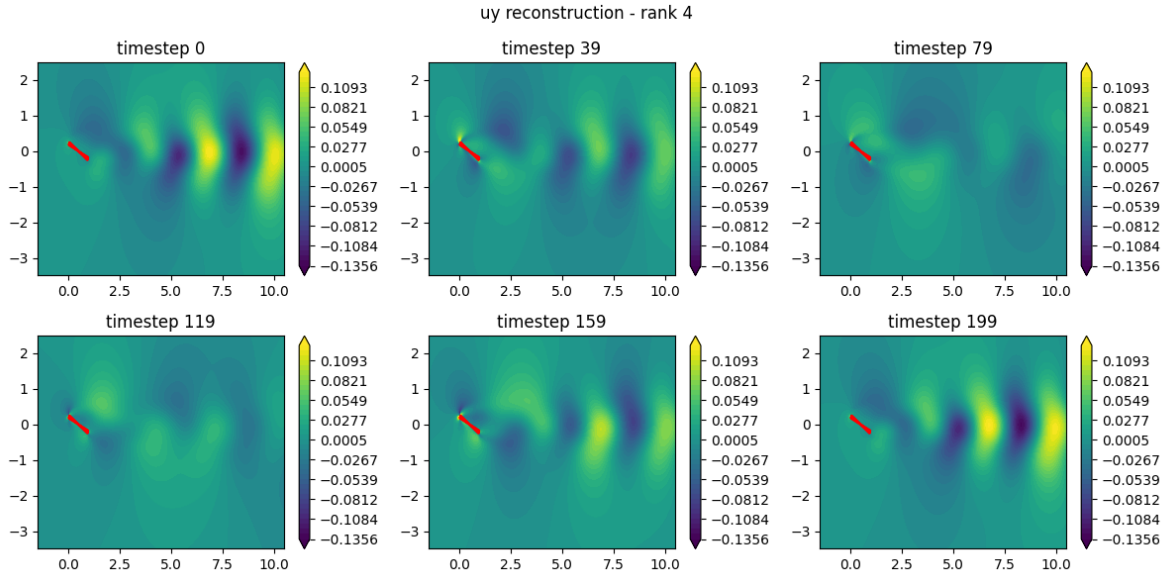


Figure 2.7:  $u_y$ \_reconstruction

## 3 Data-Driven Dynamical Systems

### 3.1 Overview: Dynamic Mode Decomposition

Dynamic mode decomposition (DMD) is a dimensionality reduction technique developed in the fluid dynamics community to extract spatiotemporal coherent structures from high-dimensional data. Unlike principal component analysis (PCA) or proper orthogonal decomposition (POD), which focus solely on spatial correlations or energy content, DMD provides a modal decomposition where each mode is associated with a specific oscillation frequency and growth/decay rate. This allows DMD to capture the temporal evolution of the system, in addition to reducing the dimensionality. DMD achieves this by approximating the Koopman operator, which describes the linear dynamics governing the system, rather than just identifying the dominant spatial patterns like PCA/POD. As a result, DMD modes can be more physically meaningful than the orthogonal modes generated by PCA, as they directly correspond to the intrinsic temporal behaviors of the system.

DMD operates on a series of data snapshots, typically obtained from simulations or experiments. These snapshots are organized into a matrix  $\mathbf{V}$ , where:

$$\mathbf{V} = [\mathbf{v}_1 \quad \mathbf{v}_2 \quad \cdots \quad \mathbf{v}_N] \in \mathbb{R}^{M \times N}$$

Each  $\mathbf{v}_i \in \mathbb{R}^M$  represents a snapshot at a specific time. DMD attempts to find a linear operator  $\mathbf{A}$  that approximates the evolution of the system from one snapshot to the next.

First, the snapshot data is arranged into two matrices  $\mathbf{X}$  and  $\mathbf{X}'$ , where  $\mathbf{X}$  contains the first  $N-1$  snapshots and  $\mathbf{X}'$  contains the last  $N-1$  snapshots.

The SVD of  $\mathbf{X}$  is taken. To deal with high-dimensional data, a reduced order representation may be used, by truncating the SVD to the first  $r$  modes.

$$\mathbf{X} = \mathbf{U}\Sigma\mathbf{V}^* \approx \mathbf{U}_r\Sigma_r\mathbf{V}_r^*$$

The full matrix  $\mathbf{A}$  is found by computing the pseudo-inverse of  $\mathbf{X}$ :

$$\mathbf{A} = \mathbf{X}'\mathbf{V}_r\Sigma_r^{-1}\mathbf{U}_r^*$$

The next step involves computing the matrix  $\tilde{\mathbf{A}}$ , which is the projection of  $\mathbf{A}$  onto the subspace spanned by the columns of  $\mathbf{U}_r$ .

$$\tilde{\mathbf{A}} = \mathbf{U}_r^* \mathbf{A} \mathbf{U}_r = \mathbf{U}_r^* \mathbf{X}' \mathbf{V}_r \Sigma_r^{-1}$$

Then, we “eigendecompose” (is that a word?)  $\tilde{\mathbf{A}}$  to find its eigenvalues  $\Lambda$  and eigenvectors  $\mathbf{W}$ . These eigenvalues and eigenvectors are used to compute the DMD modes  $\Phi$ , which are given by:

$$\Phi = \mathbf{X}' \mathbf{V}_r \Sigma_r^{-1} \mathbf{W}$$

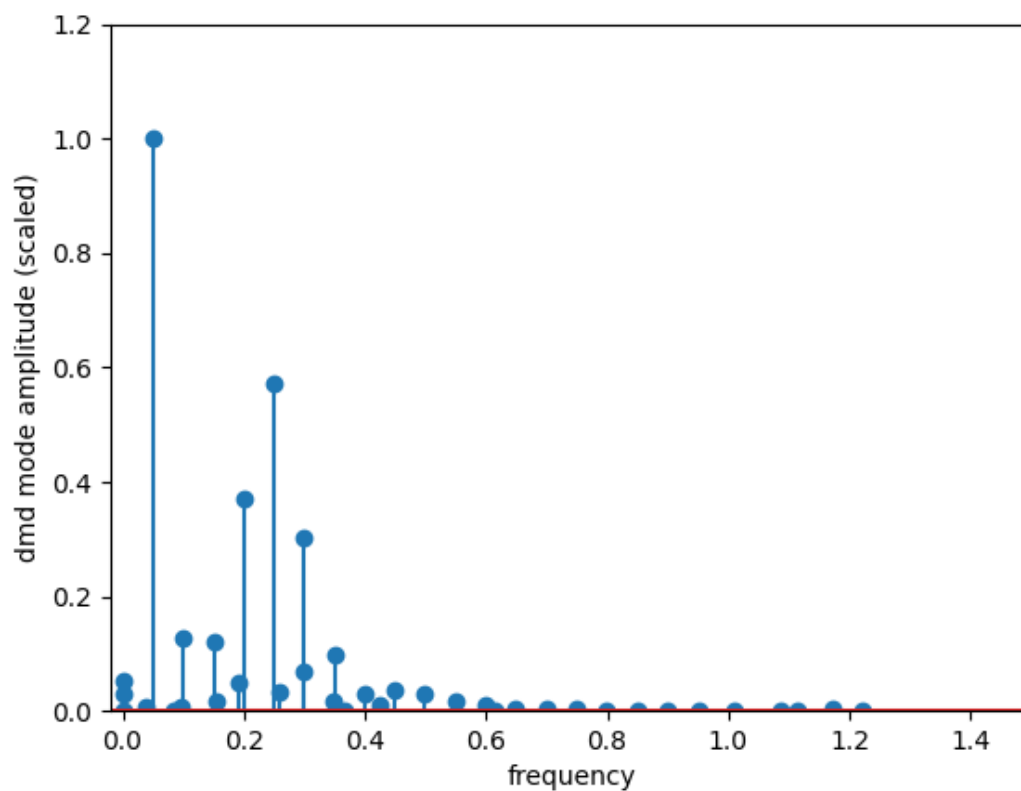
Notably, these high-dimensional DMD modes are also the eigenvectors of the full  $\mathbf{A}$  matrix corresponding to the eigenvalues in  $\Lambda$ .

## 3.2 DMD Analysis of Low Reynolds Number Pitching Airfoil DNS

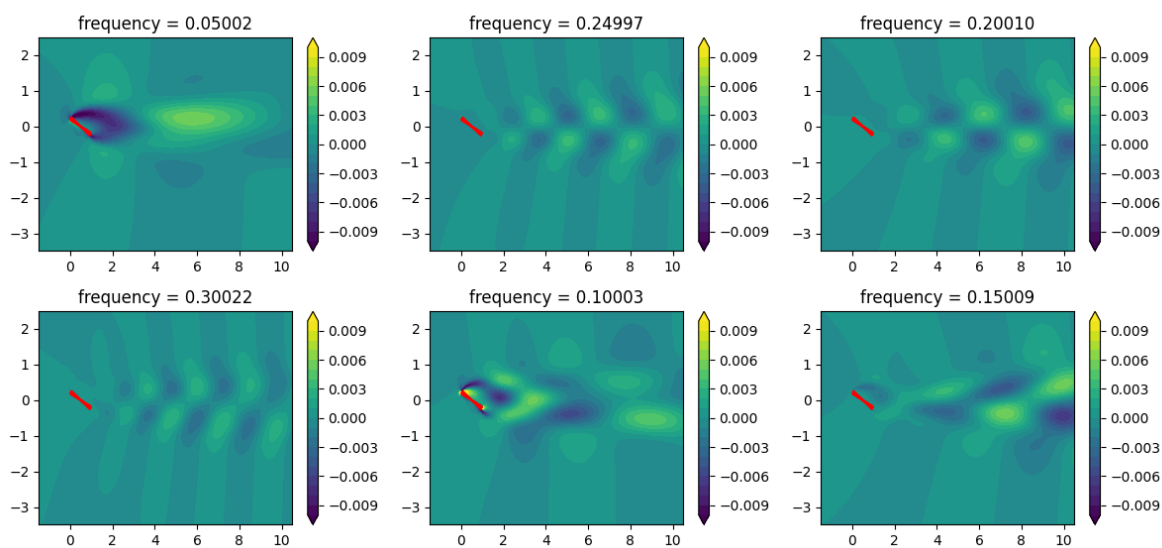
We now return to the dataset analyzed using POD, but now we use the dynamic mode decomposition to analyze this data.

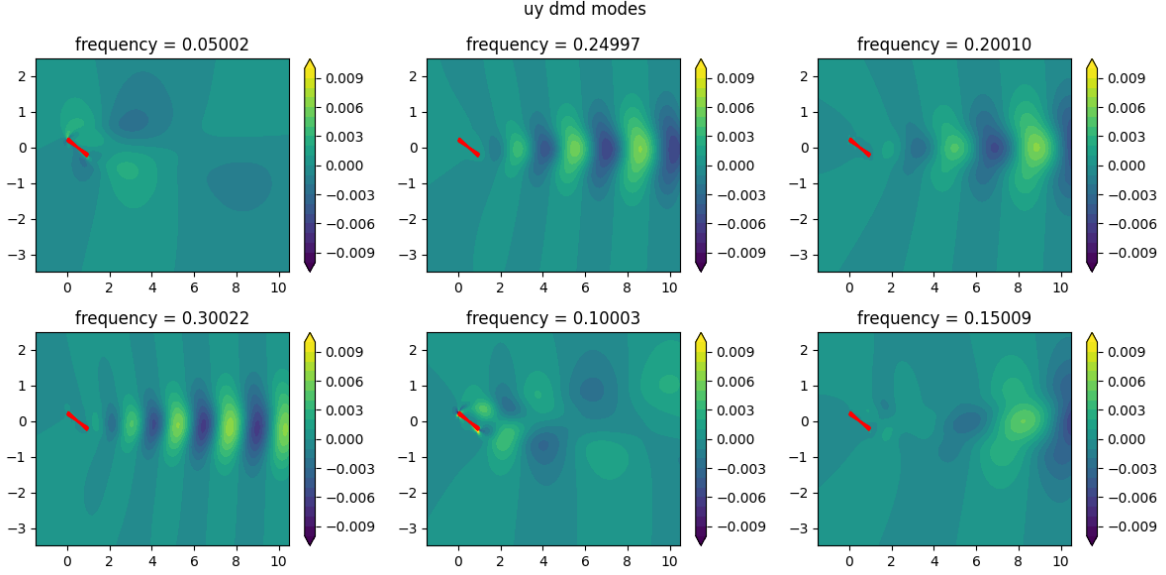
The modal amplitudes, and first six  $u_x$  and  $u_y$  modes are displayed below. It is clearly apparent that the modes are **not** identical to the POD modes. Due to the non-orthogonal nature of its modes, DMD-based representations can be less concise compared to those generated by PCA. However, DMD modes often provide greater physical insight, as each mode corresponds to a damped or driven sinusoidal time behavior.

dmd mode amplitudes w/ truncation value 75



ux dmd modes





Here, the spatial modes reveal the dominant patterns or structures within the flow. Each spatial mode corresponds to a specific frequency and growth/decay rate, providing insights into how different flow features contribute to the overall dynamics and how they change spatially across the domain.

### 3.3 Overview: Sparse Identification of Nonlinear Dynamics

The SINDy (Sparse Identification of Nonlinear Dynamics) algorithm is a technique designed to uncover the governing equations of a dynamical system from observed data. It works by identifying a sparse (or concise) representation of the system's dynamics, focusing on the most relevant terms that describe the system's behavior.

The process begins with collecting time-series data of the state variables of the system. From this data, a comprehensive library of candidate functions, which may include polynomials, trigonometric functions, and other nonlinear terms, is constructed. These functions represent potential components of the system's underlying equations.

SINDy then employs sparse regression techniques to sift through this library, selecting only the most pertinent functions. The goal is to find a minimal set of terms that can accurately describe the dynamics, resulting in a sparse representation of the system's governing equations. This approach ensures that the resulting model is both interpretable and parsimonious, capturing the essential dynamics without unnecessary complexity.

### 3.4 SINDy Implementation on Temporal Amplitudes

We attempt to fit a dynamic system  $\dot{\mathbf{x}} = f(\mathbf{x})$  to the POD temporal amplitudes using the SINDy algorithm. We select a number of temporal amplitudes to consider (6, in this case) and construct the matrix of temporal amplitudes  $\mathbf{tempamps}$  by scaling the right singular vectors  $\mathbf{V}$  with the singular values  $\Sigma$ .

Next, we compute the time derivatives of the temporal amplitudes using finite differences. We then pool the data to form a library of candidate functions  $\Theta$ , including polynomial terms up to the second order, using a function that constructs a matrix where each column is a candidate term for the dynamics, such as constant, linear, and quadratic terms of the state variables.

We perform sequential thresholding least squares to find a sparse matrix  $\Xi$  that best fits the time derivatives  $\dot{x}$  to the candidate functions in  $\Theta$ . The sparsification process involves iteratively zeroing out small coefficients (below the threshold value) and refitting the remaining coefficients.

The system is then integrated using cumulative trapezoidal numerical integration. This was initially attempted using Runge-Kutta methods, but these methods proved ineffective likely because the system is multi-scale (and rather complicated), leading to instability and inaccuracies in the solutions.

We then compare these SINDy-derived amplitudes with the original POD temporal amplitudes by plotting them. The plots show the effectiveness of the SINDy model in capturing the dynamics.

Usage of the [PySINDy](#) library was also experimented with - it yielded identical results.



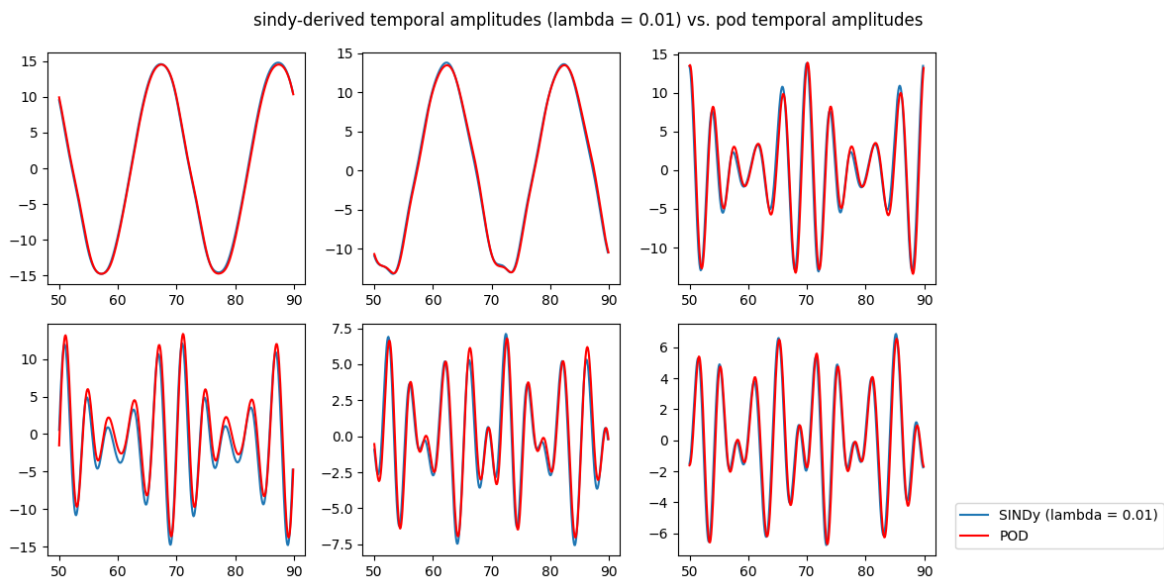


Figure 3.1: `sindy_fit`

## **4 Model Reduction and Projection**

### **4.1 Overview**

## **5 System Identification Techniques**

### **5.1 Overview: Eigensystem Realization Algorithm**

### **5.2 Overview: DMD with Control**

# References

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