

# Project 1

## FYS-STK4155

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This study revolves around the famous double-slit experiment and the wave-nature of particles. By employing the Crank-Nicholson scheme we managed to discretize the Schrödinger equation describing the evolution of a single particle in a two-dimensional box. This was used to perform simulations with single-, double-, and triple-slit configurations, storing the probability values associated with the particle's position within the box at each time step. Visualizing these results proved striking resemblance to the behavior of coherent waves passing through corresponding setups, as expected. Hence, we observed the probability distribution describing the particle's position to evolve and interact with obstacles like a wave packet, which is a clear indicator that an actual stream of such particles would too. By assuming that the particle eventually hits a detection screen on the other side of the slits, we plotted the probability values versus  $y$ -position at  $x = 0.8$  and found that the patterns all agreed with those famously observed during real life experiments. Furthermore, we determined that the real part of the wave packet itself was constantly phase-shifted  $\pi/2$  relative to its imaginary part throughout the double-slit simulation. This was expected based on the initial state, hence further validating our methods and results. We have explored various regression techniques and resampling methods within the context of machine learning, motivated by the need to develop robust models that can accurately predict and generalize from complex datasets. The main objective was to analyze and compare the performance of Ordinary Least Squares (OLS), Ridge, and Lasso regression in fitting synthetic and real-world data, focusing on the bias-variance tradeoff and model generalizability. We applied these regression techniques to the Franke function, a synthetic benchmark used in numerical analysis, and extended our analysis to cosmological N-body simulation data generated using the public GASOLINE2 code SPH code. To assess model performance and generalization, we employed resampling methods such as bootstrap and k-fold cross-validation, examining how they help to evaluate model accuracy under different training and test data conditions. Our study provided insights into how model complexity and regularization parameters affect the bias-variance tradeoff and prediction error, enabling us to critically evaluate each regression technique's performance using statistical and resampling approaches.

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<https://github.com/Oskar-Idland/FYS-STK4155-Projects>

### I. INTRODUCTION

In many scientific and practical applications, developing models that can accurately predict outcomes and generalize to new data is a critical challenge. Complex datasets often contain noise and exhibit nonlinear relationships, making it essential to employ robust regression techniques that balance model flexibility with the risk of overfitting. This work is motivated by the need to understand how different regression methods perform when applied to synthetic and real-world data, and how resampling techniques can aid in evaluating model accuracy and generalizability. The primary goal is to develop a solid understanding of the Ordinary Least Squares (OLS), Ridge and Lasso regression techniques, and their application to both synthetic and real-world data. We also intend to investigate how resampling methods such as bootstrap and cross-validation can help assess model performance.

We begin by applying the abovementioned methods to the Franke function, a widely used test function in numerical analysis, and extend the analysis to cosmo-

logical N-body simulation data made with the public version of the GASOLINE2 Smoothed Particle Hydrodynamics (SPH) code [1]. Our approach involves fitting polynomial regression models of varying complexity to the Franke function as well as the simulation data, and studying the bias-variance tradeoff, an essential concept in machine learning, in both cases. This allows us to evaluate the impact of model complexity, regularization parameters and the size of training data on the accuracy and generalizability of the models. The overarching aim is to gain insights into how different regression methods handle overfitting, model complexity and data variability, and to develop a framework for critically evaluating model performance using statistical and resampling techniques.

In section II we present relevant background theory, including central concepts such as OLS, Ridge and Lasso regression, model bias, variance and the bias-variance tradeoff, as well as two crucial resampling methods; bootstrapping and cross-validation, both of which will be implemented in this work. A few of the most important expressions introduced in this section are derived in ap-

pendix B. Our methodology is explained in section III, specifically how we define the essential design matrix, scale our data and implement the regression and resampling techniques. We also give an overview of our code structure and present the cosmological simulation data. In section IV we present, interpret and discuss the results of our analyses in light of expectations based on our previous knowledge of the implemented regression and resampling methods. Lastly, we summarize and conclude the main findings of our work in section V, and reflect.

## II. THEORY

### A. Regression Analysis

Regression analysis is a fundamental statistical technique used to model the relationship between one or more independent variables (also known as predictors or features) and a dependent variable (or target). The goal of regression analysis is to find the mathematical relationship that best explains the variation in the dependent variable based on the values of the independent variables.

Given a dataset consisting of  $n$  data points  $\{(\mathbf{x}_i, y_i)\}_{i=0}^{n-1}$ , where  $\mathbf{x}_i$  represents the input features (which is a vector in the case of multiple features), and  $y_i$  is the corresponding target value, the goal is to find a model that predicts  $\tilde{y}_i$  based on  $x_i$  such that it is as close to  $y_i$  as possible. For a linear model, the relationship between  $x_i$  and  $\tilde{y}_i$  can be expressed as

$$\tilde{y}_i = \mathbf{x}_i^\top \boldsymbol{\beta} = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \cdots + \beta_p x_{ip}, \quad (1)$$

where  $\mathbf{x}_i^\top$  is the vector of input features for the  $i$ -th data point and  $\boldsymbol{\beta} = [\beta_0, \beta_1, \dots, \beta_p]^\top$  is the vector of regression coefficients that we want to estimate. This can be written as a full-fledged matrix equation by using the so-called design or feature matrix  $\mathbf{X} \in \mathbb{R}^{n \times (p+1)}$ , which contains all input features and the bias term (so that the intercept  $\beta_0$  is included):

$$\tilde{\mathbf{y}} = \mathbf{X}\boldsymbol{\beta}. \quad (2)$$

write differently?

At its core, regression analysis seeks to find a mathematical function that relates the independent variables to the dependent variable. In its most basic form, the relationship between a dependent variable  $\mathbf{y}$  and an independent variable  $\mathbf{x}$  is modeled as

$$\mathbf{y} = f(\mathbf{x}) + \boldsymbol{\epsilon}, \quad (3)$$

where  $f(\mathbf{x})$  is the function we are trying to estimate, which represents the relationship between  $\mathbf{x}$  and  $\mathbf{y}$ . This is then what we approximate with our model  $\tilde{\mathbf{y}}$ . Furthermore,  $\boldsymbol{\epsilon}$  is the error term, representing the part of the variation in  $\mathbf{y}$  that is not explained by the model (due to noise or other unobserved factors).

### 1. Ordinary Least Squares

The Ordinary Least Squares (OLS) method is one of the most fundamental and widely used techniques in regression analysis. Its objective is to find the best-fitting line or curve for a given set of data by minimizing the sum of the squared differences between the observed values and the predicted values. These squared differences are called errors, hence the sum of squared errors (SSE):

$$\text{SSE} = \sum_{i=0}^{n-1} (y_i - \tilde{y}_i)^2. \quad (4)$$

Minimizing this sum is equivalent to minimizing the cost function

$$C(\mathbf{X}, \boldsymbol{\beta}) = (\mathbf{y} - \mathbf{X}\boldsymbol{\beta})^2 = (\mathbf{y} - \mathbf{X}\boldsymbol{\beta})^\top (\mathbf{y} - \mathbf{X}\boldsymbol{\beta}), \quad (5)$$

since the solution to this minimization problem gives the optimal values of  $\boldsymbol{\beta}$ .

To minimize  $C(\boldsymbol{\beta})$ , we take the derivative of the cost function with respect to  $\boldsymbol{\beta}$  and set it to zero:

$$\frac{\partial C(\boldsymbol{\beta})}{\partial \boldsymbol{\beta}} = -2\mathbf{X}^\top (\mathbf{y} - \mathbf{X}\boldsymbol{\beta}) = 0. \quad (6)$$

This gives the normal equation

$$\mathbf{X}^\top \mathbf{X}\boldsymbol{\beta} = \mathbf{X}^\top \mathbf{y}, \quad (7)$$

which, by solving for  $\boldsymbol{\beta}$  yields the OLS estimator:

$$\hat{\boldsymbol{\beta}} = (\mathbf{X}^\top \mathbf{X})^{-1} \mathbf{X}^\top \mathbf{y}. \quad (8)$$

This formula provides the best-fitting coefficients  $\boldsymbol{\beta}$  that minimize the sum of squared errors. In appendix B 1 we derive the following important results:  
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$$\mathbb{E}[\hat{\boldsymbol{\beta}}] = \boldsymbol{\beta}, \quad (9)$$

$$\text{Var}[\hat{\boldsymbol{\beta}}] = \sigma^2 (\mathbf{X}^\top \mathbf{X})^{-1}. \quad (10)$$

Here  $\mathbb{E}[\hat{\boldsymbol{\beta}}]$  and  $\text{Var}[\hat{\boldsymbol{\beta}}]$  are the expectation value and variance of the OLS estimator, respectively, while  $\sigma^2$  is the variance of the error term  $\boldsymbol{\epsilon}$ .

### 2. Ridge

While OLS provides a foundational method for regression, it can lead to problems when the data has high multicollinearity or when there are more features than data points, leading to overfitting. OLS attempts to minimize the sum of squared errors, but it does not impose any restrictions on the model complexity. This often results in high variance when the model learns to fit noise in the training data, especially when the design matrix  $\mathbf{X}$  is poorly conditioned (i.e., when columns

of  $\mathbf{X}$  are nearly linearly dependent). This is because the matrix  $\mathbf{X}^\top \mathbf{X}$  becomes close to singular, making its inverse highly sensitive to small changes in the data.

**remove last sentences?**

To mitigate the issue with OLS, Ridge regression introduces a regularization term to the cost function (5), penalizing large coefficients and preventing the model from becoming overly complex. The modified cost function is

$$C_{\text{Ridge}}(\mathbf{X}, \boldsymbol{\beta}) = \frac{1}{n} \|\mathbf{y} - \mathbf{X}\boldsymbol{\beta}\|_2^2 + \lambda \|\boldsymbol{\beta}\|_2^2, \quad (11)$$

where  $\lambda \geq 0$  is a hyperparameter that controls the strength of the regularization, and the subscripts 2 simply mean that we are taking the  $L^2$  norm. This is defined as

$$\|\mathbf{x}\|_2 = \sqrt{\sum_i x_i^2}.$$

We see that when  $\lambda = 0$ , Ridge regression reduces to OLS, as the regularization term disappears. On the other hand, when  $\lambda$  is large, the coefficients  $\boldsymbol{\beta}$  must shrink to minimize the cost function, potentially reducing overfitting. It is important to note, however, that if  $\lambda$  becomes too large the coefficients tend to zero, and we may end up with underfitting instead. We therefore need to determine the hyperparameter carefully in order to find the perfect balance between the two extremes.

**formulate differently?**

To minimize this altered cost function, we again take the derivative with respect to  $\boldsymbol{\beta}$  and set it to zero:

$$\frac{\partial C_{\text{Ridge}}(\boldsymbol{\beta})}{\partial \boldsymbol{\beta}} = -2\mathbf{X}^\top(\mathbf{y} - \mathbf{X}\boldsymbol{\beta}) + 2\lambda\boldsymbol{\beta} = 0. \quad (12)$$

Rearranging this equation gives us the Ridge regression normal equation:

$$(\mathbf{X}^\top \mathbf{X} + \lambda \mathbf{I})\boldsymbol{\beta} = \mathbf{X}^\top \mathbf{y}, \quad (13)$$

where  $\mathbf{I}$  is the identity matrix. Notice the term  $\lambda \mathbf{I}$  added to  $\mathbf{X}^\top \mathbf{X}$ , which makes the matrix invertible even when  $\mathbf{X}^\top \mathbf{X}$  is poorly conditioned. This leads to the Ridge regression estimator:

$$\hat{\boldsymbol{\beta}}_{\text{Ridge}} = (\mathbf{X}^\top \mathbf{X} + \lambda \mathbf{I})^{-1} \mathbf{X}^\top \mathbf{y}. \quad (14)$$

Here, the regularization term  $\lambda$  shrinks the coefficients and prevents them from becoming too large, thereby controlling the model's variance. **remove this paragraph?**

### 3. Lasso

Like Ridge regression, Lasso regression (Least Absolute Shrinkage and Selection Operator) is a regularization technique designed to improve the generalizability of the model by introducing a penalty term to the cost

function. While Ridge regression uses the  $L^2$  norm for regularization, Lasso regression employs the  $L^1$  norm, defined as

$$\|\mathbf{x}\|_1 = \sum_i |x_i|.$$

This leads to a different form of regularization, since the cost function now takes the form

$$C_{\text{Lasso}}(\mathbf{X}, \boldsymbol{\beta}) = \frac{1}{n} \|\mathbf{y} - \mathbf{X}\boldsymbol{\beta}\|_2^2 + \lambda \|\boldsymbol{\beta}\|_1. \quad (15)$$

The difference from Ridge regression has important implications, mainly because the  $L^2$  norm makes it impossible for any of the coefficients to vanish completely. **correct?** This can be problematic in high-dimensional datasets where we might expect many features to be irrelevant. By using the  $L^1$  norm, Lasso regression has the unique ability to drive some coefficients to exactly zero, effectively performing both regularization and feature selection. Thus, Lasso regression not only addresses overfitting but also simplifies the model by automatically excluding irrelevant features, making it particularly useful for sparse models where only a subset of features are truly important. **double check, introduce expressions, cite?**

## B. Properties of Predictive Models

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### 1. Predicted Values

In the context of predictive modeling,  $\tilde{\mathbf{y}}$  represents the model's predicted values of the target variable  $\mathbf{y}$ . These predictions are made based on the features and parameters learned from the training data. For new or unseen data, the true target values  $\mathbf{y}$  are often unknown, but the model generates an estimate  $\tilde{\mathbf{y}}$  that approximates these values.

The performance of the model's predictions can be analyzed by decomposing the error into several components. The error of  $\tilde{\mathbf{y}}$  stems from three main sources: noise variance  $\sigma^2$ , model bias and model variance. The latter two are expressed as

$$\text{Bias}[\tilde{\mathbf{y}}] = \mathbb{E}[(\mathbf{y} - \mathbb{E}[\tilde{\mathbf{y}}])^2], \quad (16)$$

$$\text{Var}[\tilde{\mathbf{y}}] = \mathbb{E}[(\tilde{\mathbf{y}} - \mathbb{E}[\tilde{\mathbf{y}}])^2]. \quad (17)$$

Each of these represents a different aspect of the total error that affects the model's performance.

### 2. Model Bias

From the expression (16) we see that the model bias measures how far the average model prediction  $\mathbb{E}[\tilde{\mathbf{y}}]$  is

from the true target value  $\mathbf{y}$ . More specifically, it refers to the systematic error introduced by the model's inability to capture the true underlying relationship between the independent and dependent variables. This error arises when the model makes incorrect assumptions about the data or oversimplifies the relationship. For example, a linear model trying to fit highly non-linear data will result in high bias because the model cannot flexibly represent the complexity of the data.

### 3. Model Variance

We see from the expression (17) for the model variance that it represents the variability in the model's predictions across different training sets. A high model variance indicates that the model is overly sensitive to the specific training data, leading to overfitting, where the model performs well on the training set but poorly on new, unseen data such as the test set. This happens because the model has "memorized" the noise in the training data rather than capturing the true underlying pattern. High variance typically arises in overly complex models that are capable of capturing minute details, which may not generalize well to new data.

### 4. Bias-Variance Tradeoff

Measures of a predictive model's total error and how well it is likely to predict future samples are commonly expressed in terms of the mean-squared error (MSE) and the score function, respectively. These are given by

$$\text{MSE}(\mathbf{y}, \tilde{\mathbf{y}}) = \frac{1}{n} \sum_{i=0}^{n-1} (y_i - \tilde{y}_i)^2, \quad (18)$$

$$R^2(\mathbf{y}, \tilde{\mathbf{y}}) = 1 - \frac{\sum_{i=0}^{n-1} (y_i - \tilde{y}_i)^2}{\sum_{i=0}^{n-1} (y_i - \bar{y})^2}, \quad (19)$$

where  $\bar{y}$  is the mean value of  $\mathbf{y}$  and  $n$  is the number of data points. In appendix B 2 [maybe move up here?] we show that the MSE alternatively can be expressed in terms of the model bias, model variance and noise variance as

$$\mathbb{E}[(\mathbf{y} - \tilde{\mathbf{y}})^2] = \text{Bias}[\tilde{y}] + \text{Var}[\tilde{y}] + \sigma^2. \quad (20)$$

Since the noise variance adds an irreducible contribution to the total model error, an ideal model strikes a balance between bias and variance, where it is flexible enough to capture the underlying patterns in the data but not so flexible that it fits the noise. This balance is referred to as the bias-variance tradeoff. It is a fundamental concept in machine learning, specifically when building predictive models, and will be studied in great detail throughout this work.

## C. Resampling Methods

Resampling methods are statistical techniques used to generate additional data samples from the available data. These methods are particularly useful in machine learning and data analysis when the dataset is limited, and we want to better assess the performance of a model. The primary goal of resampling is to estimate the accuracy of a model by splitting the data into different subsets or generating new samples of the data, repeatedly fitting the model, and evaluating its performance on different sets of data. In this work we implement two commonly used resampling techniques: bootstrap and cross-validation.

### 1. Bootstrap

The bootstrap method is a powerful resampling technique used to estimate the uncertainty and variability of a model by repeatedly drawing random samples, with replacement, from the original dataset. This resampling creates multiple "bootstrapped" datasets, each the same size as the original, but with some samples appearing multiple times and others potentially omitted. By computing the average performance of the multiple models created with the bootstrapped datasets it becomes easier to estimate the model's bias and variance.

The process of bootstrap resampling approximates the underlying distribution of a statistic, whether it's model performance, a parameter estimate or prediction error, without requiring strong parametric assumptions about the data. This makes it particularly useful when the theoretical distribution of a statistic is unknown or difficult to calculate. Bootstrap is especially convenient when the dataset is small, or when no explicit train-test split is available, as it makes the most out of the available data. It is also good for understanding the bias-variance trade-off since it can be used to determine a model's variability, and it allows for the estimation of confidence intervals for predictions and model parameters. The downside is that it involves training a model multiple times, which can be computationally expensive, especially with large datasets or complex models. [shorten or cite?]

### 2. Cross-Validation

Cross-validation is another resampling technique that involves partitioning the dataset into several distinct subsets (or "folds"), and then systematically training the model on one subset while testing it on another. A common form is  $k$ -fold cross-validation, where the dataset is divided into  $k$  equally-sized folds. The model is trained on  $k - 1$  folds and tested on the remaining fold. This process is repeated  $k$  times, with each fold serving as the test set exactly once.

Since all observations are used for both training and validation, cross-validation maximizes the use of avail-

able data, hence it is especially useful in the case of limited datasets. It also helps in managing the bias-variance tradeoff by evaluating the model’s performance on different subsets of data. By averaging the results across multiple folds, cross-validation reduces the variance of the model’s predictions, as it minimizes overfitting to any particular subset of the data. Additionally, by testing the model on unseen data, it ensures that the bias is not too high, making it a good indicator of model generalization. It does require multiple model trainings however (e.g., 10-fold cross-validation requires 10 models to be trained), making it computationally expensive for large datasets or complex models. [shorten or cite?](#)

#### D. The Franke Function

The Franke function is a widely used two-dimensional synthetic function in numerical analysis and computational mathematics, particularly in the fields of interpolation, regression analysis, and surface fitting. It is defined over the unit square  $[0, 1] \times [0, 1]$  and expressed as

$$\begin{aligned} f(x, y) = & \frac{3}{4} \exp\left\{\left(-\frac{(9x - 2)^2}{4} - \frac{(9y - 2)^2}{4}\right)\right\} \\ & + \frac{3}{4} \exp\left\{\left(-\frac{(9x + 1)^2}{49} - \frac{9y + 1}{10}\right)\right\} \\ & + \frac{1}{2} \exp\left\{\left(-\frac{(9x - 7)^2}{4} - \frac{(9y - 3)^2}{4}\right)\right\} \\ & - \frac{1}{5} \exp\{(-(9x - 4)^2 - (9y - 7)^2)\}, \quad (21) \end{aligned}$$

We will use the function multiple times throughout this work, as it produces a surface with both smooth and non-smooth regions, making it an excellent benchmark for testing regression and resampling algorithms. It simulates real-world data that may contain both complex variations and noise, mimicking scenarios that we will encounter as we move on to implementing the algorithms on actual simulation data.

### III. METHODS & IMPLEMENTATION

write in past tense from here on!

#### A. Generating the Data

##### 1. Synthetic Data

The Franke function generated data used in this study was created by sampling the function on a grid of  $N \times N$  points, where  $N = 50$ . A visualization of the function as a surface is shown in fig. 1. Furthermore, to emulate inherent variability and uncertainty in real-world data we added a gaussian noise component with a zero mean

and standard deviation of  $\sigma = 0.1$ . The noise was added to simulate the inherent variability and uncertainty in real-world data.

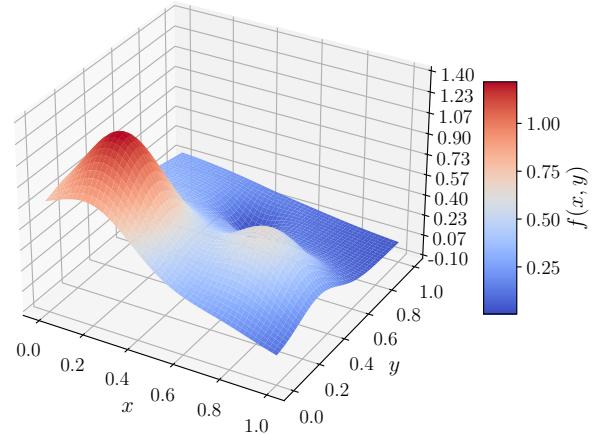


FIG. 1: Visualization of the two-dimensional Franke function expressed in (21).

##### 2. Cosmological Simulation Data

After implementing regression analysis and resampling methods using the Franke function we repeated the process using data from an N-body simulation of dark matter structure formation made with the public GASOLINE2 SPH code [1]. The initial conditions for the simulation were generated with MUSIC (MULTI-Scale Initial Conditions) [2], and standard Planck cosmology parameters  $\Omega_m = 0.3077$ ,  $\Omega_\Lambda = 0.6923$  and  $H_0 = 67.81 \text{ km/s/Mpc}$  were used. The simulation box has length, width and height  $L = 20 \text{ Mpc}$ , corresponding to a  $500 \times 500 \times 500$  grid, and contains  $64^3$  dark matter particles.

The output data includes 128 “snapshots” of the box through time, where the time parameter in the simulation is the scale factor, defined as

$$a = \frac{1}{1+z}.$$

Here  $z$  is the cosmological redshift, which is 0 today and thus in the 128th snapshot. We focused on the dark matter density  $\rho$ , and studied a snapshot at redshift  $z \approx 12.88$  because the distribution of  $\rho$  is smoother at larger  $z$ . To get a function of two variables we averaged over one of the three axes so that we get a 2D grid, and because the density is so large and vastly different at different points in the box we took  $\ln \rho$  to be our dependent variable instead of  $\rho$ . The resulting data is visualized as a contour plot and a surface plot in figure 2. We did not add a synthetic noise component to this data, i.e.  $\sigma = 0$  in this

case, since the simulation output can be interpreted as the true data if we were to perform real observations of dark matter density.

## B. Regression Analysis

explain implementation on both Franke and data

### 1. Creating the Design Matrix

Our first step when performing regression analyses on our data was to extend the design matrix  $\mathbf{X}$  to the situation when we have two independent variables  $\mathbf{x}$  and  $\mathbf{y}$  instead of one  $\mathbf{x}$ . We then use  $\mathbf{z}$  to denote our dependent variable. In the case of polynomial regression of degree  $p$  with one independent variable vector  $\mathbf{x} = (x_0 \dots x_n)^\top$  the design matrix has the form

$$\mathbf{X} = \begin{bmatrix} 1 & x_0 & x_0^2 & \dots & x_0^p \\ 1 & x_1 & x_1^2 & \dots & x_1^p \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ 1 & x_n & x_n^2 & \dots & x_n^p \end{bmatrix}.$$

The feature vector in this case is  $\beta = (\beta_0 \dots \beta_p)^\top$ , hence it is a column vector of length  $p + 1$ , while our design matrix has dimensions  $(n + 1) \times (p + 1)$

When we have two independent variables representing physical coordinates in a grid each combination of  $x_i$  and  $y_j$  corresponds to a unique component  $z_{ij}$  in  $\mathbf{z}$ . If  $\mathbf{x}$  and  $\mathbf{y}$  contain  $n_x + 1$  and  $n_y + 1$  components, respectively, the design matrix for polynomial degree  $p$  then becomes

$$\mathbf{X} = \begin{bmatrix} 1 & x_0 & y_0 & \dots & x_0^p & x_0^{p-1}y_0 & \dots & x_0y_0^{p-1} & y_0^p \\ 1 & x_1 & y_0 & \dots & x_1^p & x_1^{p-1}y_0 & \dots & x_1y_0^{p-1} & y_0^p \\ \vdots & \vdots & \vdots & \ddots & \vdots & \vdots & \ddots & \vdots & \vdots \\ 1 & x_{n_x} & y_0 & \dots & x_{n_x}^p & x_{n_x}^{p-1}y_0 & \dots & x_{n_x}y_0^{p-1} & y_0^p \\ 1 & x_0 & y_1 & \dots & x_0^p & x_0^{p-1}y_1 & \dots & x_0y_1^{p-1} & y_1^p \\ \vdots & \vdots & \vdots & \ddots & \vdots & \vdots & \ddots & \vdots & \vdots \\ 1 & x_{n_x} & y_{n_y} & \dots & x_{n_x}^p & x_{n_x}^{p-1}y_{n_y} & \dots & x_{n_x}y_{n_y}^{p-1} & y_{n_y}^p \end{bmatrix}.$$

We thus see that for polynomial degree 0 we have one column, three columns for first degree, six for second, ten for third, etc. This is a well known series of numbers, and we can express the number of features  $l$  in terms of the polynomial degree  $p$  as

$$l = \frac{1}{2}(p + 1)(p + 2).$$

Our design matrix thus has dimensions  $N \times l$  where  $N = (n_x + 1) \times (n_y + 1)$ . move to theory?

## 2. Preprocessing

For this study, data scaling was essential to ensure that the features were on a similar scale, which is critical for many machine learning algorithms. By utilizing the `StandardScaler` function from the scikit-learn library [cite](#), we standardized our datasets by subtracting the mean and dividing the standard deviation over the data set, for each feature. Although the data generated by the Franke function and the cosmological simulation data used in subsequent analyses are inherently two-dimensional and may not exhibit significant variation in scale between features, scaling still provides consistent benefits.

The primary motivation for applying this scaling function is to enhance the performance and stability of our algorithms. Scaling prevents features with larger magnitudes from disproportionately influencing the objective function and helps balance minor discrepancies between features that might affect the learning process. The performance of our regression models is significantly improved when the data is scaled due to ensuring faster convergence and reducing the risk of numerical instability.

To illustrate the impact of scaling, we plotted the beta coefficients for both a non-scaled and a scaled design matrix as a function of polynomial degree in our OLS model fits on the Franke function-generated data. This visualization allowed us to observe how scaling influenced the coefficients and affected the model's performance.

Another important preprocessing step was splitting the data into training and test sets. This step enables the evaluation of the model's performance on unseen data and helps in assessing its generalization capabilities. We used the `train_test_split` function from the scikit-learn library to randomly split the data into training and test sets, with a test size of 0.2. This ensured that 20% of the data was reserved for testing, while the remaining 80% was used to train the model. By splitting the data in this manner, we could evaluate the model's performance on new data and determine how well it generalized to unseen samples.

## 3. Parameter Dependencies

Investigating the dependency on polynomial degree was a critical aspect of understanding how the regression models performed under varying conditions. By increasing the polynomial order, we enhanced the models' capacity to fit more complex patterns in the data. However, this also raised the risk of overfitting, where the model might capture noise along with the underlying signal. To explore this, we evaluated the performance of the models across different polynomial degrees using metrics such as the MSE and  $R^2$  scores mentioned in the previous section. These evaluations allowed us to quantify how well each model balanced bias and variance.

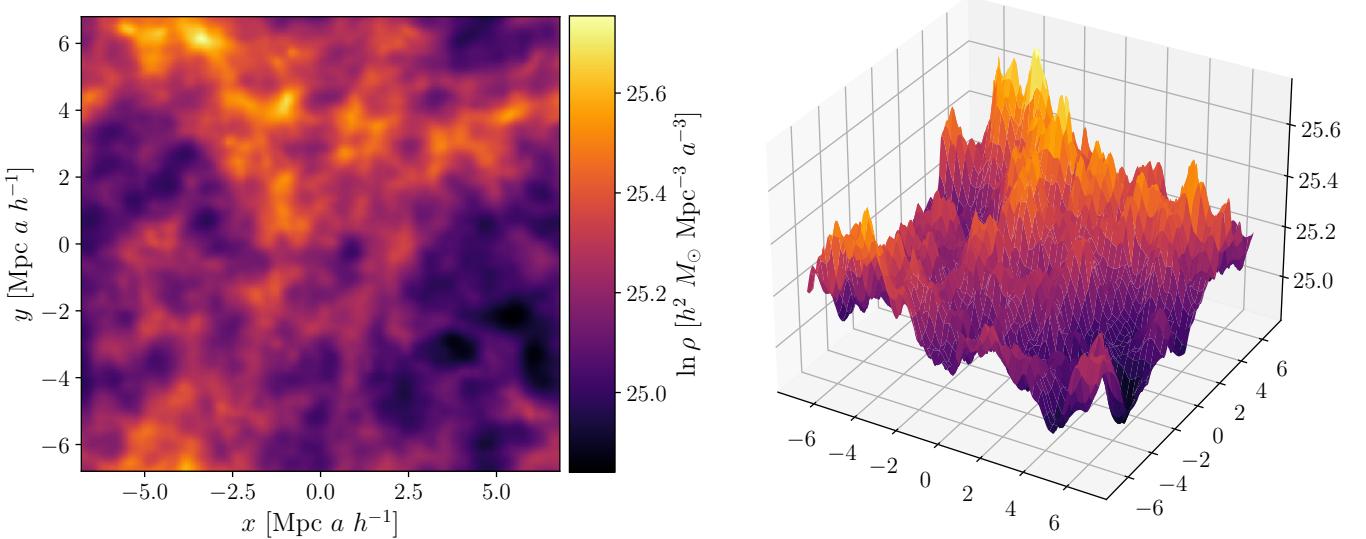


FIG. 2: Logarithm of dark matter density in the simulation box averaged over one of the three axes at redshift  $z \approx 12.88$ . Plotted as contour plot on the left and as surface plot on the right. Because the Universe expands as time goes on the box does as well, and we therefore scale the coordinates of the box by multiplying with  $a$ . Similarly, space between the dark matter particles also increases with time, which is why the density is scaled by multiplying with  $a^{-3}$ . GASOLINE2 uses the rest of the scaling factors by default.

For the Franke function-generated data, we systematically varied the polynomial degree from 1 to 6 for OLS. We then assessed the model's performance using the MSE and  $R^2$  scores, plotting these metrics as a function of model complexity to determine the optimal polynomial degree for the regression method.

For the cosmological simulation data, we followed a similar approach. We varied the polynomial degree from 1 to 40, using only every other step, for OLS. The model's performance was evaluated using the same metrics as for the Franke function-generated data. We also plotted the predicted surfaces to visually assess the models' fit to the data and identify any potential overfitting or underfitting issues. Specifically, we analyzed the fits for polynomial degrees 5, 10, 20, 30, and 50. Additionally, a 3D surface plot of a 30-degree polynomial fit was created alongside the raw data to visualize the model's performance in capturing the underlying patterns.

For both Ridge and Lasso regression on the Franke function-generated data, we explored polynomial degrees ranging from 1 to 41, using every second polynomial degree for Ridge and every fourth polynomial degree for Lasso. These choices were motivated by the runtime constraints of the algorithms. For the cosmological simulation data, the polynomial degree range was from 1 to 42 for Ridge and from 1 to 21 for Lasso, with a step of every fourth polynomial degree for both methods.

In addition to polynomial degree, we examined the impact of the hyperparameter  $\lambda$  on the performance of our Ridge and Lasso regression models. Visualizing the MSE and  $R^2$  scores for Ridge and Lasso regression involved not only evaluating polynomial complexity but also varying  $\lambda$  values.

To thoroughly evaluate the models, we systematically tested a range of  $\lambda$  values, from 1e-05 to 1. For the Franke function-generated data, this range included 30 values, evenly spaced on a logarithmic scale, whereas 11 values were tested for the cosmological simulation data. Each  $\lambda$  value was tested for every polynomial degree within the ranges mentioned earlier.

The resulting MSE and  $R^2$  scores were plotted as functions of both  $\lambda$  and polynomial degree using contour plots. This dual-parameter visualization allowed us to identify the optimal combination of polynomial degree and  $\lambda$  for each model, thereby providing a comprehensive view of how model performance changes with varying complexity and regularization strength.

### C. Bias-Variance Tradeoff

we expect  $\text{MSE} = \text{bias} + \text{var}$  for data since noise var is 0

To showcase the bias-variance trade-off and validate our implementation, we imitated the analysis illustrated in ??Fig. 2.11 from Hastie, Tibshirani, and Friedman[fig:Hastie]. Specifically, we plotted the MSE for both the training and test sets of the Franke data using

OLS regression. We varied the polynomial degree from 1 to 81 in steps of 4. [Link to our result?](#)

The generation and splitting of the Franke data were conducted as described previously.

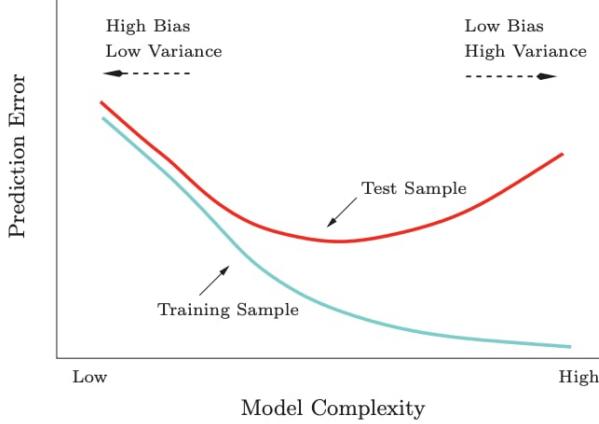


FIG. 3: Fig. 2.11 of Hastie, Tibshirani, and Friedman illustrating bias-variance trade-off as a function of model complexity [cite](#)

### 1. Bootstrap With OLS

Our implementation of the bootstrap technique involved resampling the training data multiple times and fitting an OLS model to each resampled dataset. This was accomplished using the resample function from the scikit-learn library. For each resampled dataset, we computed the MSE, bias, and variance to evaluate the model's performance.

For the Franke function-generated data, we designed four different scenarios, each varying a specific parameter to investigate the influence of the number of bootstraps, data points, model complexity, and test size separately.

The default values for each scenario were set as follows: 100 bootstraps, 50 data points, polynomial degree of 5, and a test size of 20%. The varying parameters for each respective scenario were:

- Number of bootstraps: [1, 2, 5, 10, 20, 50, 100, 200, 500, 1000, 2000, 5000, 10000]
- Number of data points: [10, 20, 30, 40, 50, 100, 150, 200, 300, 400, 500, 1000]
- Polynomial degree: from 1 to 41 with a step of 2
- Test size: ranging from 5% to 95% with a step of 5%

For each scenario we plotted the model evaluation metrics as a function of the varying parameter to analyze how the model's performance changed with different settings.

For the cosmological simulation data, we only did a single scenario with 100 bootstraps, 50 data points, test size of 20%, and a varying polynomial degree from 1 to 31 with a step of 3. We only plotted the MSE as a function of polynomial degree for this scenario. [Explain why?](#)

### 2. Cross-Validation

go into detail how we implement

compare with bootstrap include Ridge and Lasso

For our cross-validation analysis, we used the `KFold` function from the scikit-learn library to split the data into  $k$  folds, with the integer  $k \in [5, 15]$ . We iterated over a

## D. The Program

### 1. Code Structure

All source code for this project is available on our GitHub repository, linked in appendix A. The `README.md` contains the entire project structure. The code is divided into the following files and notebooks:

`data_analysis.ipynb`: Jupyter notebook containing the analysis of the dark matter density data. This is where we apply our regression models.

`franke.ipynb`: Jupyter notebook containing the analysis of the Franke function-generated data, to validate our implementation.

`functions.py`: Python script containing all the functions used in the notebooks. All properly documented.

### 2. Tools

Python [3] is used for creating our models, together with scikit-learn [4]. To vectorize our code we use `numpy` [5], and for visualization we use `matplotlib.pyplot` [6]. To make our plots prettier we use the colormaps module [7]. Code completion and debugging is done in Visual Studio Code [8] with additional assistance of GitHub Copilot [9]. We use `git` [10] for version control, and `GitHub` [11] for remote storage of our code.

raw and smoothed surfaces, explain reason/method

## IV. RESULTS & DISCUSSION

### A. Effects of Data Scaling

In fig. 4 we have plotted the estimated MSE and  $R^2$  scores evaluated on the Franke function test for polynomial degrees  $\in [1, 6]$ . The left and right columns show

the scores computed for the raw and scaled data, respectively. As expected there looks to be a constant factor between the raw and scaled MSE's, while the  $R^2$  scores are equal in both cases. This is because  $y_i$ ,  $\tilde{y}_i$  and  $\bar{y}$  (or in our case,  $z_i$ ,  $\tilde{z}_i$  and  $\bar{z}$ ) are scaled equally, and from eq. (19) we thus see that the scaling cancels out from the numerator and denominator in the second term.

One might expect to see the same effects of scaling for the cosmological simulation data, but in fig. 5 we see that this is not the case. Here we have plotted the MSE and  $R^2$  scores for 21 evenly spaced polynomial degrees in the interval [1, 41], and we see that for degrees  $\gtrsim 9$  the raw scores diverge from the expected trend. Instead of a decreasing MSE and increasing  $R^2$  as in the scaled case we get MSE's over 400 and very large, negative  $R^2$  scores, a metric that should never be negative in the first place. This clearly shows why it is beneficial to scale the data, since it is safe to assume that the raw data catastrophe occurs due to numerical instability caused by loss of floating point precision. correct? Moreover, we see that we never reach a turning point in the scores where they start to worsen with polynomial degree, which makes sense considering the complexity of our data set.

In fact, we see from fig. 6 that the scaled coefficients for the fits to the cosmological data (plotted in orange) grow in orders of magnitude with complexity (cirka up until degree 21, where stay around  $10^4$ ), while the raw coefficients (plotted in purple) fall to zero around polynomial degree 9. The reason for the strong increase in the  $\beta_{\text{scaled}}$ 's is that  $\mathbf{x}$  and  $\mathbf{y}$  (both raw and scaled) are centered at 0, and in the scaled case these are divided by their standard deviations, hence most of the highest order polynomial terms decrease with the degree. We thus need very large coefficients for their contributions to the fit to be noticeable. The  $\beta_{\text{raw}}$ 's must be much smaller since the highest order terms grow rapidly with degree at the edges of the box, and eventually we encounter floating point errors. This is contrary to the coefficients estimated for the Franke function, where generally  $\beta_{\text{raw}} \geq \beta_{\text{scaled}}$ , as seen in fig. 16 in appendix C. This is because the Franke function is defined over the unit square, hence the raw coefficients grow with polynomial degree as well. rewrite?

## B. Hyperparameter Dependency

### 1. Ridge

### 2. Lasso

## C. Bias-Variance Tradeoff

## D. Resampling Methods

### 1. OLS: Cross-Validation VS Bootstrap

### 2. Cross-Validation With Ridge and Lasso

## E. Fits to the Cosmological Data

### 1. Complexity Dependency for OLS

In fig. 14 we have plotted OLS fits to the cosmological data using polynomial degrees 5, 10, 20, 30 and 50

to further examine the effects of increasing model complexity. From degree 30 to 50 the fit actually seems to worsen, looking more smudged in the center, while ripple-like structures near the edges become more prominent. These are already visible for degree 30, but compared to the data it arguably doesn't make the fit look less accurate than for degree 20.

To explain the reason for the ripple effect, and why the smaller scale structures in the data don't show up more clearly as we continue increasing the model complexity we need to consider the columns of the scaled design matrix. As mentioned in section IV A most of the scaled box coordinates are less than 1. They are in fact close to zero in the center region and slightly larger than 1 in magnitude near the edges. Thus, as we move to higher order polynomials the elements of the rightmost columns in  $\mathbf{X}_{\text{scaled}}$  (corresponding to the highest order terms) diverge, some tend to zero while others grow in magnitude. To minimize the model error the coefficients cannot continue increasing because then the fit worsens in the edges, but they cannot decrease too much either as this leads to a less accurate fit in the center. EXPLAIN THE RIPPLE EFFECT

## 2. Comparing Regression Variants

## V. CONCLUSION

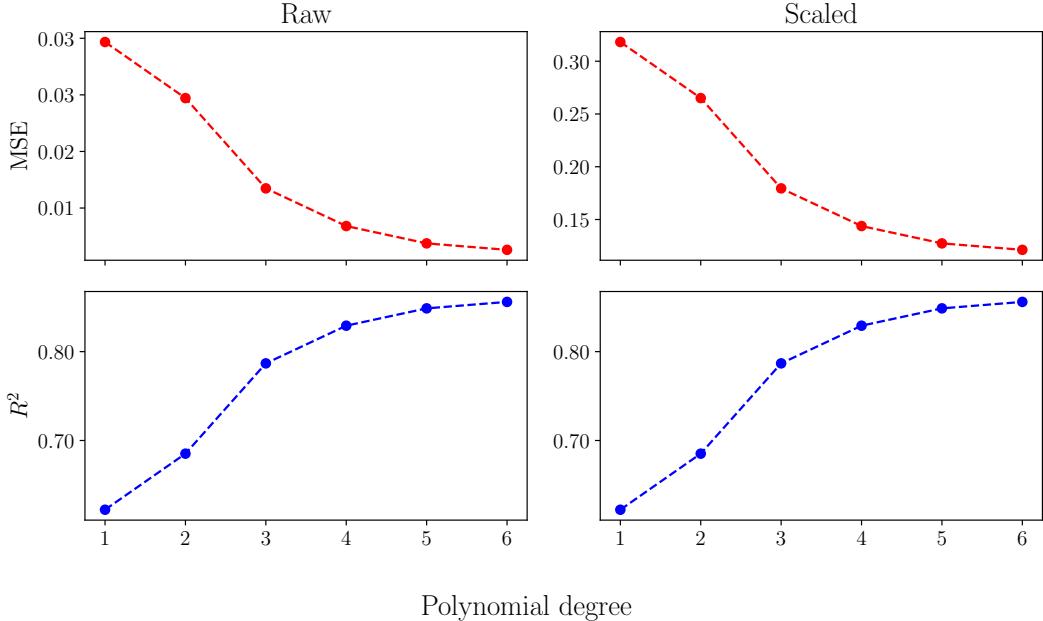


FIG. 4: MSE (upper row) and  $R^2$  (lower row) evaluated on the Franke function test set using unscaled (left column) and scaled (right column) values, for polynomial degrees in the interval  $[1, 6]$ .

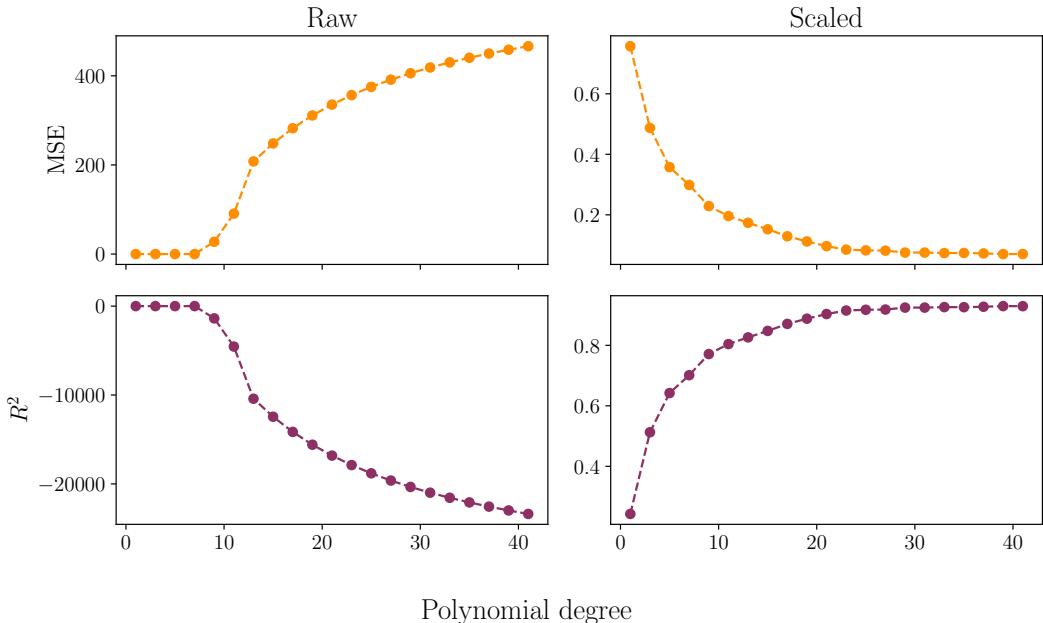


FIG. 5: MSE (upper row) and  $R^2$  (lower row) evaluated on the cosmological test set using unscaled (left column) and scaled (right column) values, for polynomial degrees in the interval  $[1, 41]$ .

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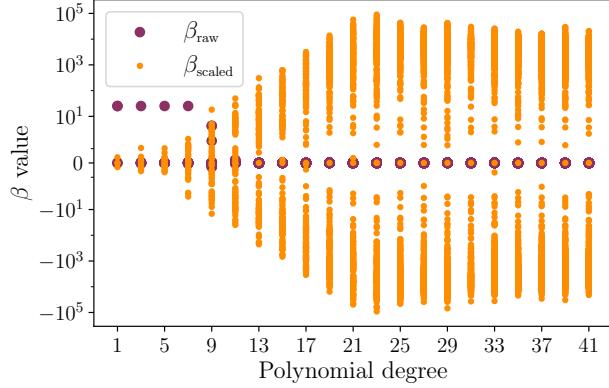


FIG. 6: The coefficients  $\beta$  for the cosmological fit estimated using unscaled (purple) and scaled (orange) data, plotted for polynomial degrees in the interval  $[1, 41]$ . We have used a logarithmic scale for the  $\beta$ 's.

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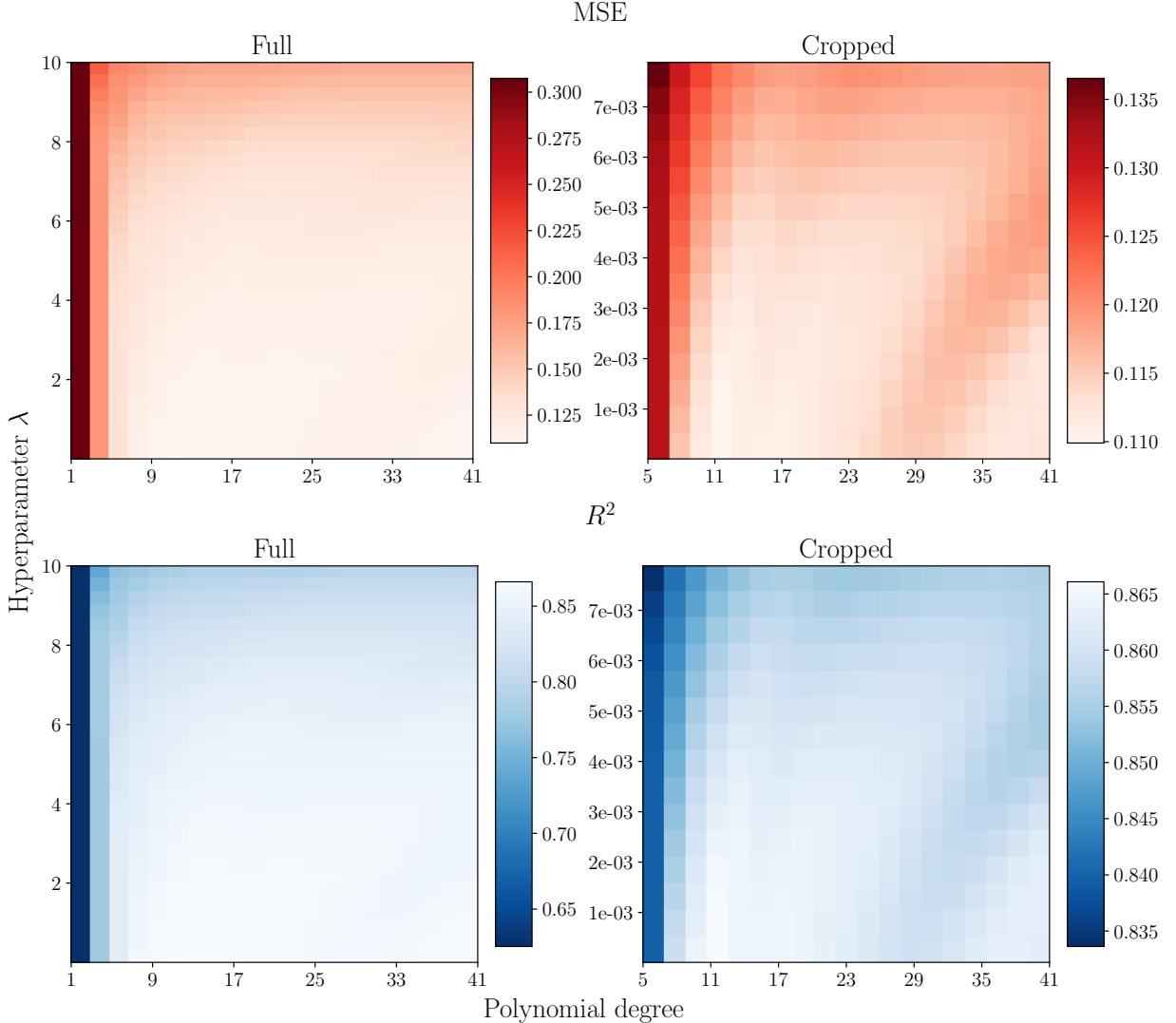


FIG. 7: MSE (upper row) and  $R^2$  (lower row) for Ridge regression evaluated on the Franke test data and contour plotted as functions of polynomial degree (horizontal axes) and hyperparameter  $\lambda$  (vertical axes). We have used 21 evenly spaced degrees in the interval  $[1, 41]$  and 30 logarithmically spaced values for  $\lambda$  in the interval  $[10^{-5}, 10^{-1}]$ .

We have cropped away degrees 1 and 3 and the 15 largest  $\lambda$  values purely for visualization purposes.

#### Appendix A: Code

<https://github.com/Oskar-Idland/FYS-STK4155-Projects>

#### Appendix B: Derivations

##### 1. Expectation value and variance of the OLS estimator $\hat{\beta}$

rewrite to fit report?

Since the error in  $\mathbf{y}$  is normal distributed as  $\boldsymbol{\varepsilon} \sim N(0, \sigma^2)$  we know that the expectation value and variance of the  $i$ 'th element of  $\boldsymbol{\varepsilon}$  is  $\mathbb{E}(\varepsilon_i) = 0$  and  $\text{Var}(\varepsilon_i) = \sigma^2$ . Thus, since we approximate  $f(\mathbf{x})$  with  $\tilde{\mathbf{y}} = \mathbf{X}\boldsymbol{\beta}$  the expectation value of  $\mathbf{y}$  becomes

$$\mathbb{E}(\mathbf{y}) = \mathbb{E}(\mathbf{X}\boldsymbol{\beta}) + \mathbb{E}(\boldsymbol{\varepsilon}) = \mathbb{E}(\mathbf{X}\boldsymbol{\beta}), \quad (\text{B1})$$

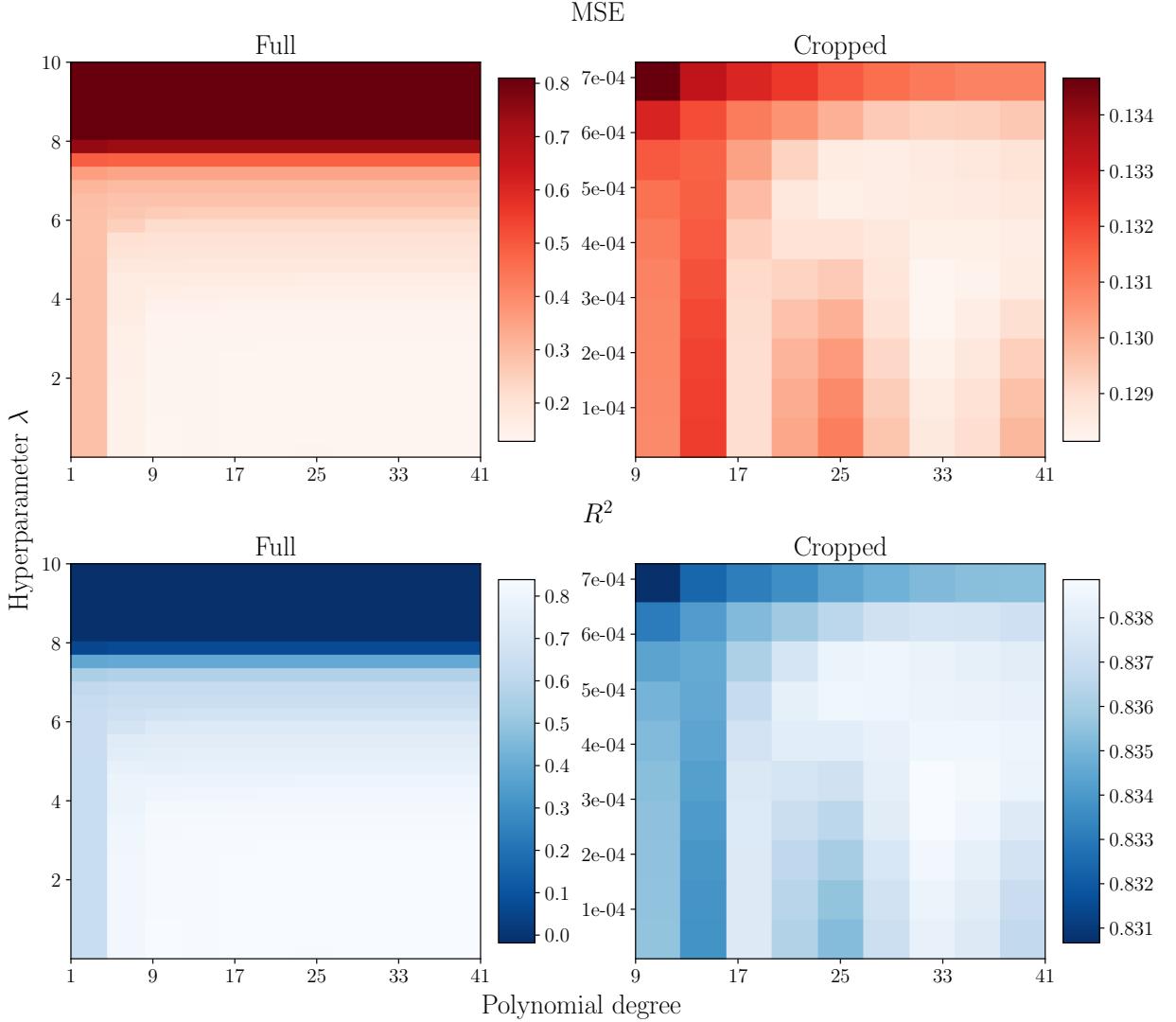


FIG. 8: MSE (upper row) and  $R^2$  (lower row) for Lasso regression evaluated on the Franke test data and contour plotted as functions of polynomial degree (horizontal axes) and hyperparameter  $\lambda$  (vertical axes). We have used 11 evenly spaced degrees in the interval  $[1, 41]$  and 30 logarithmically spaced values for  $\lambda$  in the interval  $[10^{-5}, 10^{-1}]$ , but in the right column we have cropped away degrees 1 and 5 and the 20 largest  $\lambda$  values.

which gives us the expectation value of  $\mathbf{y}$  for a given element  $i$ :

$$\mathbb{E}(y_i) = \mathbb{E} \left( \sum_j X_{ij} \beta_j \right) = \sum_j X_{ij} \beta_j = \mathbf{X}_{i,*} \boldsymbol{\beta}. \quad (\text{B2})$$

Here we have used that the sum  $\sum_j X_{ij} \beta_j$  is known to be the value of  $\tilde{y}_i$ , hence its expectation value is itself. Moreover, we can similarly find the variance of  $\mathbf{y}$  for a given element  $i$  by using that the terms in the aforementioned sum are known to be  $\tilde{y}_i$  for all  $i$ , i.e.  $\text{Var}(\mathbf{X}_{i,*} \boldsymbol{\beta}) = 0$ . Thus, we have

$$\text{Var}(y_i) = \text{Var}(\mathbf{X}_{i,*} \boldsymbol{\beta}) + \text{Var}(\varepsilon_i) = \sigma^2, \quad (\text{B3})$$

and consequently

$$y_i \sim N(\mathbf{X}_{i,*} \boldsymbol{\beta}, \sigma^2). \quad (\text{B4})$$

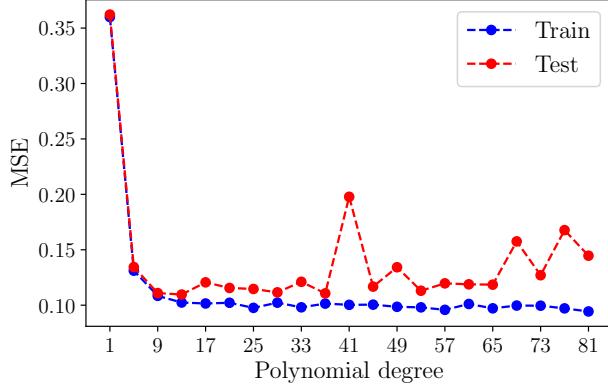


FIG. 9: MSE evaluated on the Franke function training set (blue) and test set (red) for 21 evenly spaced polynomial degrees in the interval [1, 81].

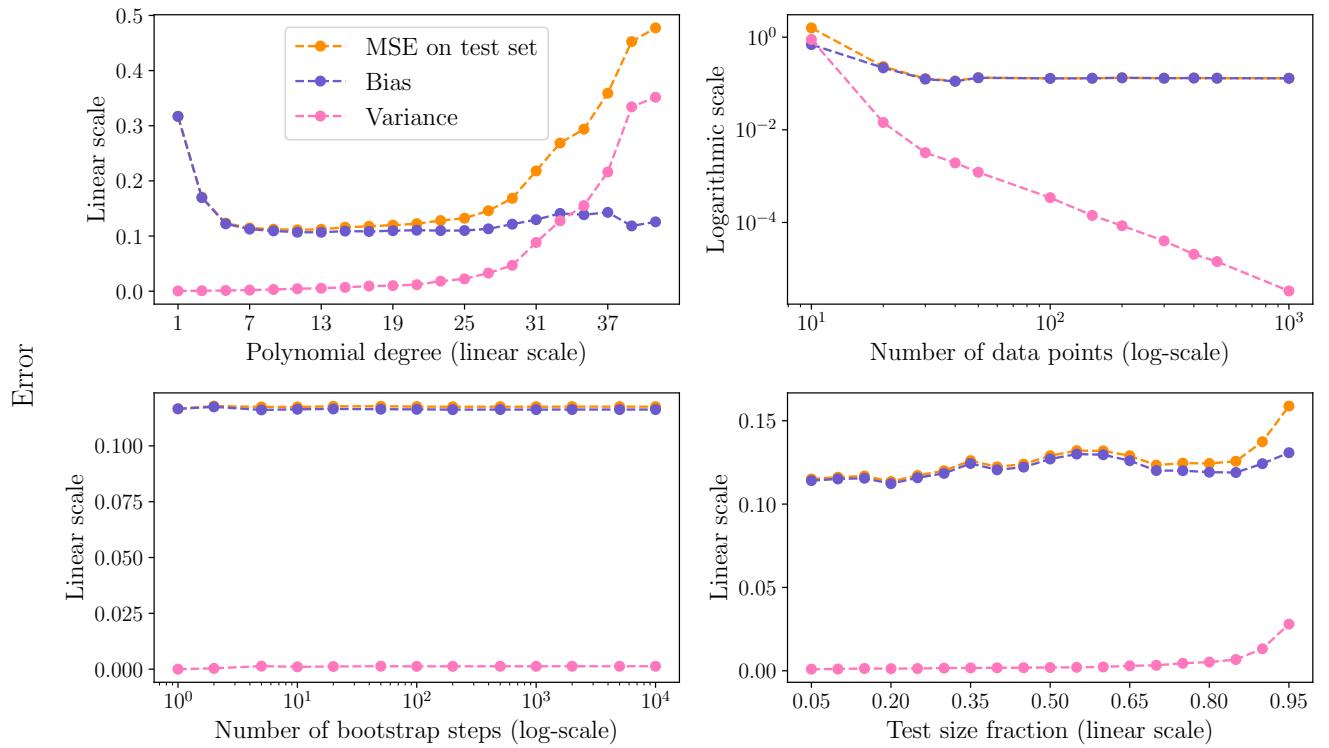


FIG. 10: MSE (orange), bias (blue) and variance (pink) for OLS regression evaluated on Franke test data and plotted as functions of polynomial degree (upper left), number of data points (upper right), number of bootstrap steps (lower left) and test size fraction (lower right). When these are not varied we have used degree 5, 50 data points in one direction (i.e. 2 500 in the grid), 100 bootstrap steps and a test size fraction of 0.2. Note that some axes are scaled logarithmically and some have a linear scale.

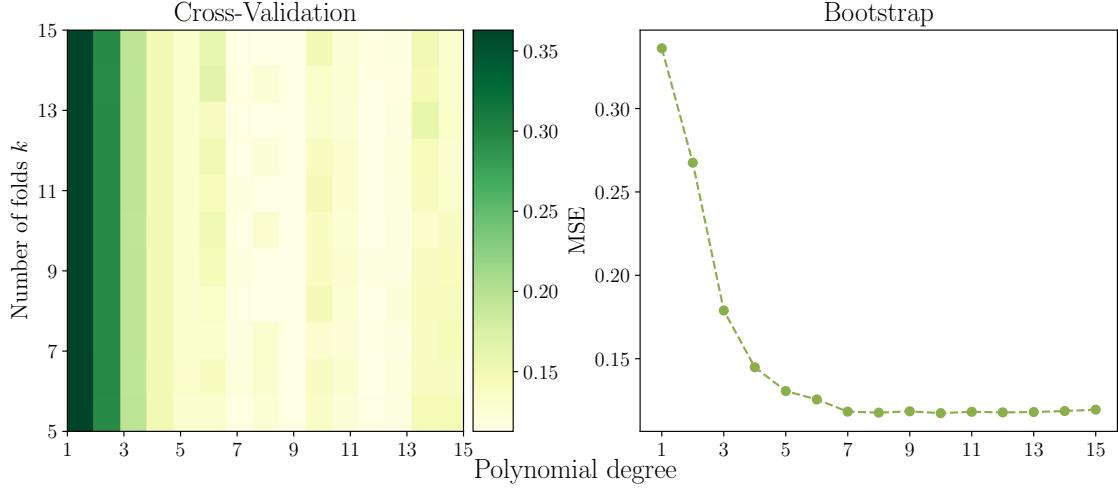


FIG. 11: MSE evaluated on Franke test data as function of polynomial degrees in the interval [1, 15] and number of folds  $k \in [5, 15]$  used in cross-validation (contour plot on the left). On the right we have plotted the MSE purely as function of complexity when using the bootstrap resampling method with 100 steps.

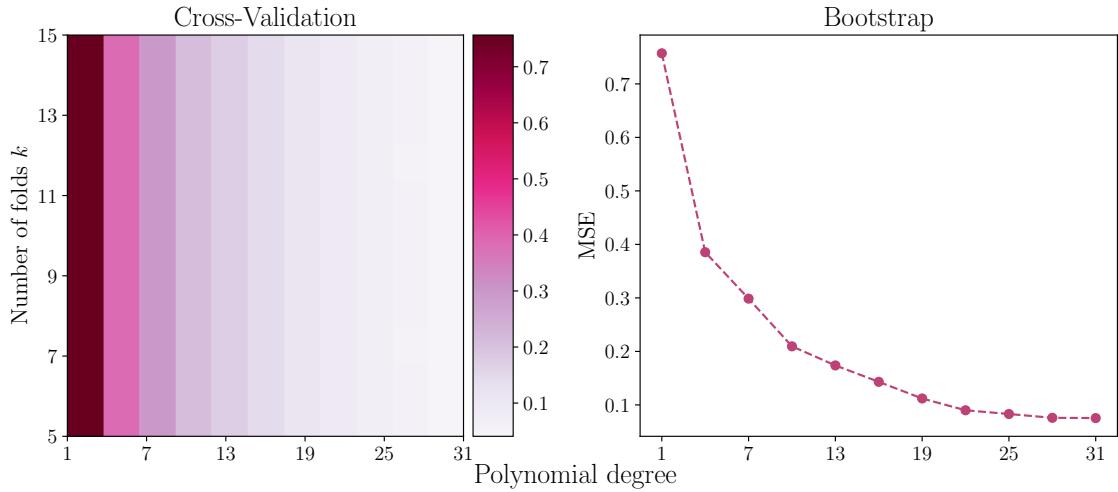


FIG. 12: MSE evaluated on cosmological test data as function of polynomial degrees in the interval [1, 31] and number of folds  $k \in [5, 15]$  used in cross-validation (contour plot on the left). On the right we have plotted the MSE purely as function of complexity when using the bootstrap resampling method with 100 steps.

Now, using that the OLS estimator is given by  $\hat{\beta} = (\mathbf{X}^\top \mathbf{X})^{-1} \mathbf{X}^\top \mathbf{y}$ , its expectation values become

$$\begin{aligned}
\mathbb{E}(\hat{\beta}) &= \mathbb{E} \left\{ (\mathbf{X}^\top \mathbf{X})^{-1} \mathbf{X}^\top \mathbf{y} \right\} \\
&= \mathbb{E} \left\{ (\mathbf{X}^\top \mathbf{X})^{-1} \mathbf{X}^\top \right\} \mathbb{E}(\mathbf{y}) \\
&= (\mathbf{X}^\top \mathbf{X})^{-1} \mathbf{X}^\top \mathbf{X} \beta \\
&= \beta.
\end{aligned} \tag{B5}$$

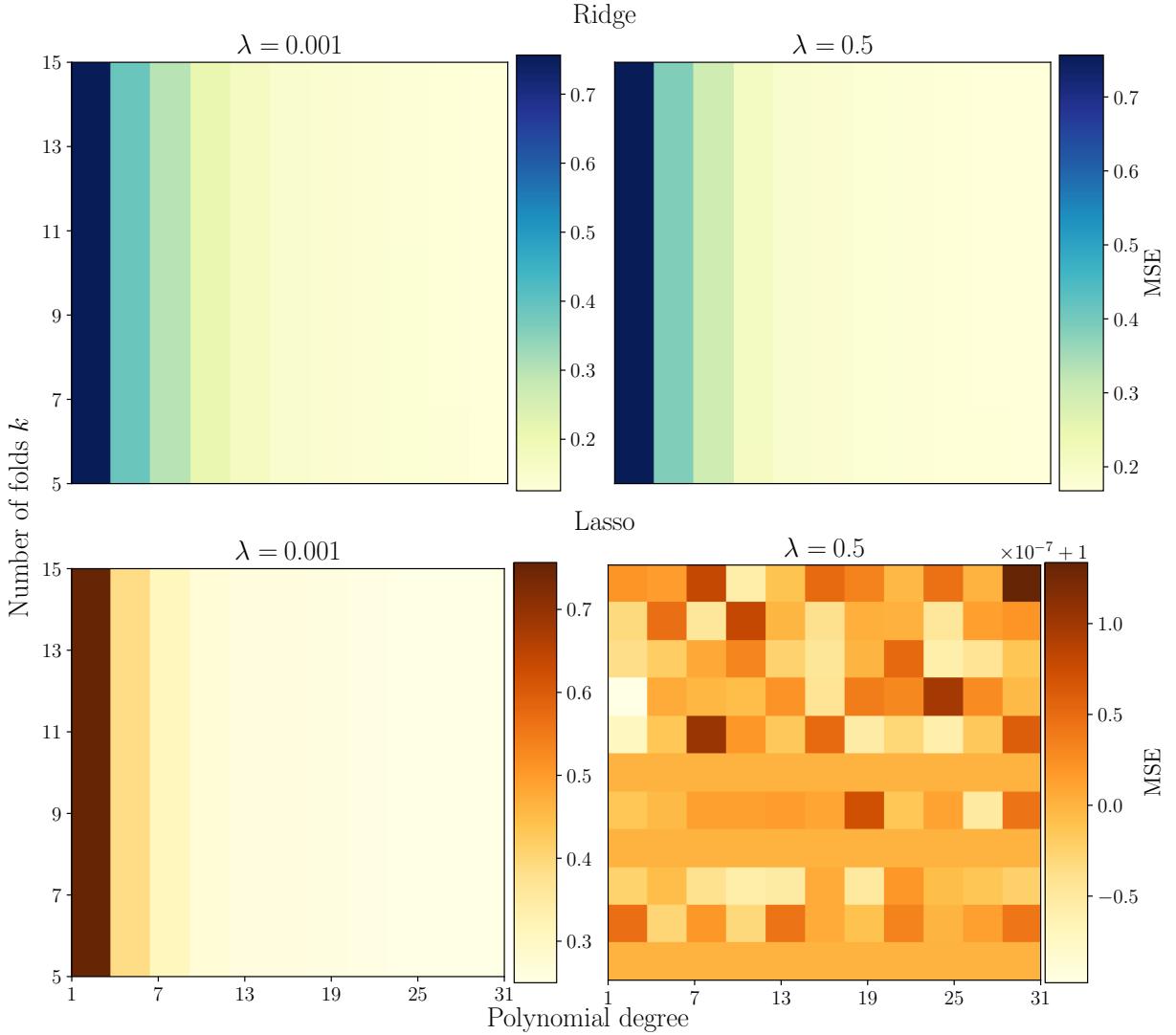


FIG. 13: MSE evaluated on cosmological test data as function of polynomial degrees in the interval  $[1, 31]$  and number of folds  $k \in [5, 15]$  used in cross-validation, for Ridge (upper row) and Lasso (lower row) regression. We have used  $\lambda = 0.001$  and  $\lambda = 0.5$  in the left and right columns, respectively.

Furthermore, if  $x$  and  $y$  are two independent variables, the variance of their product is given by

$$\begin{aligned}
\text{Var}(xy) &= \mathbb{E}(x^2y^2) - (\mathbb{E}(xy))^2, \\
&= \mathbb{E}(x^2)\mathbb{E}(y^2) - (\mathbb{E}(x))^2(\mathbb{E}(y))^2, \\
&= [\mathbb{E}(x^2) - (\mathbb{E}(x))^2 + (\mathbb{E}(x))^2] [\mathbb{E}(y^2) - (\mathbb{E}(y))^2 + (\mathbb{E}(y))^2] - \mathbb{E}(x^2)\mathbb{E}(y^2), \\
&= [\text{Var}(x) + (\mathbb{E}(x))^2] [\text{Var}(y) + (\mathbb{E}(y))^2] - \mathbb{E}(x^2)\mathbb{E}(y^2), \\
&= \text{Var}(x)\text{Var}(y) + \text{Var}(x)(\mathbb{E}(y))^2 + \text{Var}(y)(\mathbb{E}(x))^2 + \mathbb{E}(x^2)\mathbb{E}(y^2) - \mathbb{E}(x^2)\mathbb{E}(y^2), \\
&= \text{Var}(x)\text{Var}(y) + \text{Var}(x)(\mathbb{E}(y))^2 + \text{Var}(y)(\mathbb{E}(x))^2,
\end{aligned}$$

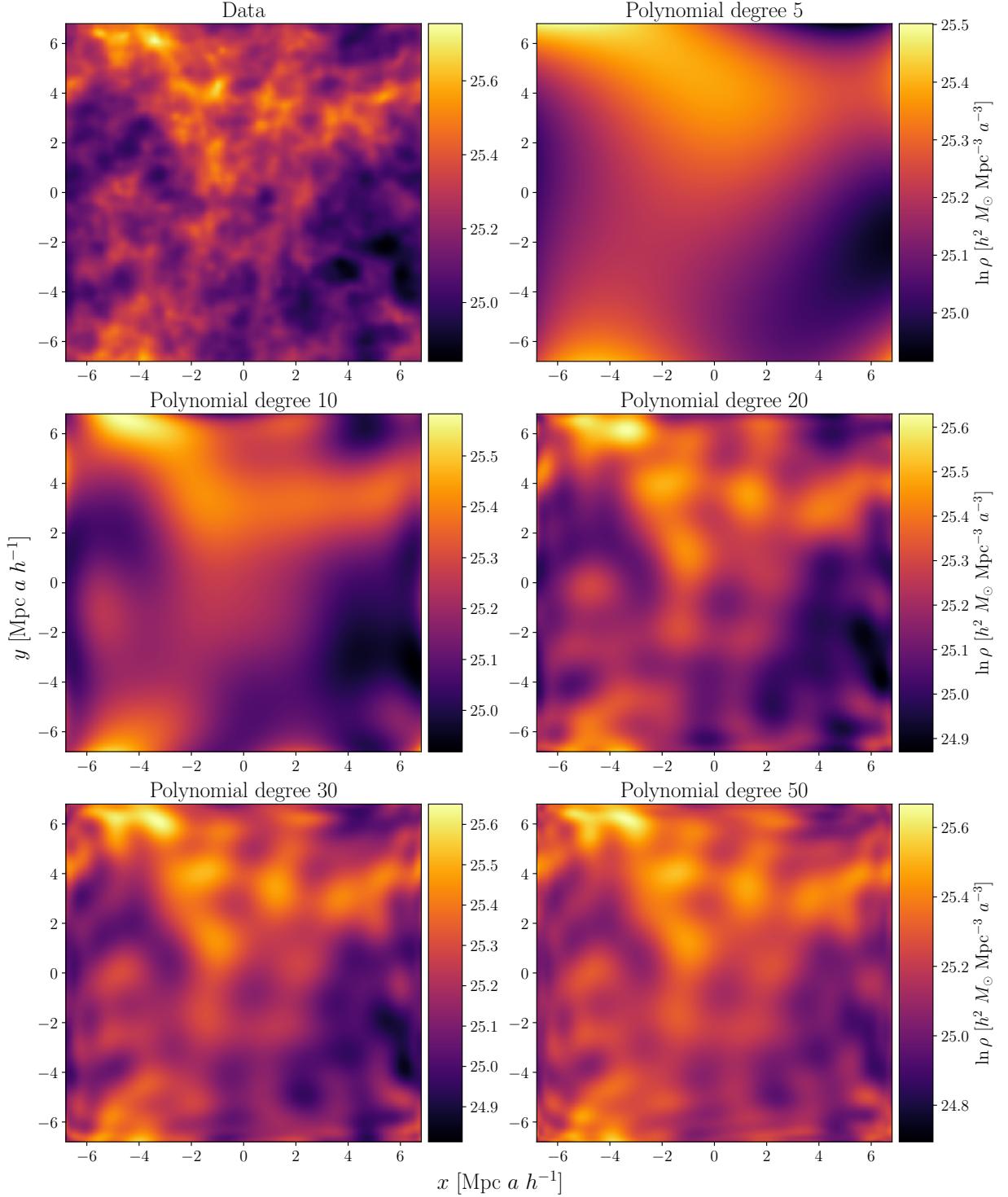


FIG. 14: Contour plots of the cosmological simulation data (upper left) and the OLS fits for polynomial degrees 5 (upper right), 10 (middle left), 20 (middle right), 30 (lower left) and 50 (lower right).

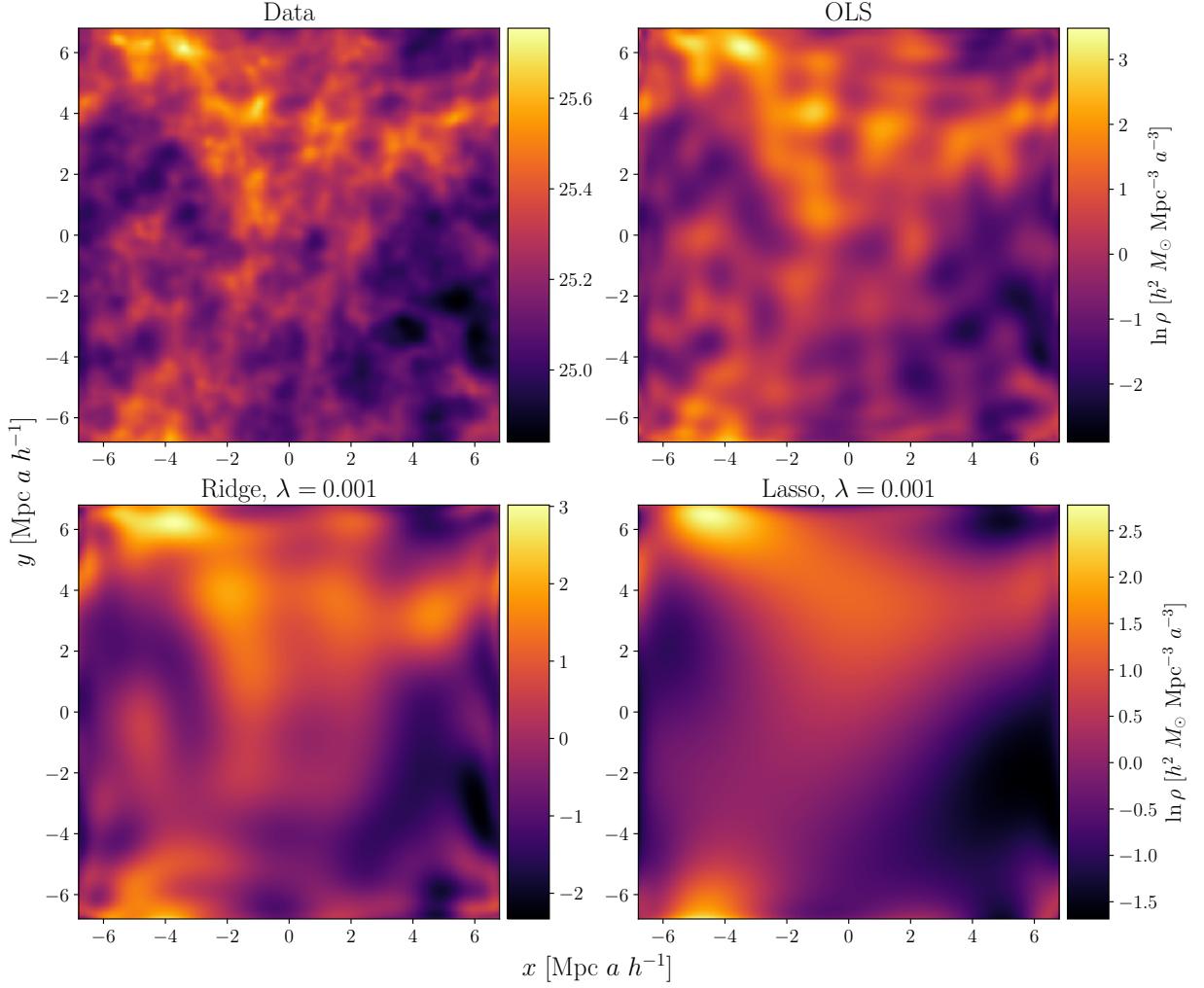


FIG. 15: Contour plots of the cosmological simulation data (upper left) and OLS (upper right), Ridge (lower left) and Lasso (lower right) regression fits for polynomial degree 31 and  $k = 15$  number of folds in cross-validation. For Ridge regression we used `lambda` and for Lasso we used `lambda`.

so if we now set  $x = (\mathbf{X}^\top \mathbf{X})^{-1} \mathbf{X}^\top$  and  $y = \mathbf{y}$  we find that the variance of  $\hat{\beta}$  is

$$\begin{aligned}
\text{Var}(\hat{\beta}) &= \underbrace{\text{Var}\left[(\mathbf{X}^\top \mathbf{X})^{-1} \mathbf{X}^\top\right] \text{Var}(\mathbf{y})}_{0} + \underbrace{\text{Var}\left[(\mathbf{X}^\top \mathbf{X})^{-1} \mathbf{X}^\top\right] (\mathbb{E}(\mathbf{y}))^2 + \text{Var}(\mathbf{y}) (\mathbb{E}\left[(\mathbf{X}^\top \mathbf{X})^{-1} \mathbf{X}^\top\right])^2}_{0}, \\
&= \sigma^2 \left[ (\mathbf{X}^\top \mathbf{X})^{-1} \mathbf{X}^\top \right]^2, \\
&= \sigma^2 \left[ (\mathbf{X}^\top \mathbf{X})^{-1} \mathbf{X}^\top \right] \left[ (\mathbf{X}^\top \mathbf{X})^{-1} \mathbf{X}^\top \right]^\top, \\
&= \sigma^2 \left[ (\mathbf{X}^\top \mathbf{X})^{-1} \mathbf{X}^\top \right] \left[ \mathbf{X} (\mathbf{X}^\top \mathbf{X})^{-1} \right], \\
&= \sigma^2 (\mathbf{X}^\top \mathbf{X})^{-1}.
\end{aligned} \tag{B6}$$

Here we have used that the transpose of  $(\mathbf{X}^\top \mathbf{X})^{-1}$  is itself since it is square and symmetric. mention that sigma is zero in cosmo case, so variance is zero

## 2. MSE expressed in terms of model bias, model variance and noise variance

rewrite to fit report?

Substituting  $\mathbf{y}$  with  $f(\mathbf{x}) + \boldsymbol{\epsilon}$ , and adding and subtracting  $\mathbb{E}[\tilde{\mathbf{y}}]$ , we find that

$$\begin{aligned}
\mathbb{E}[(\mathbf{y} - \tilde{\mathbf{y}})^2] &= \mathbb{E}\left[\underbrace{(f(\mathbf{x}) + \boldsymbol{\epsilon} - \tilde{\mathbf{y}})^2}_{\mathbf{f}}\right], \\
&= \mathbb{E}[(\mathbf{f} + \boldsymbol{\epsilon} - \tilde{\mathbf{y}} + \mathbb{E}[\tilde{\mathbf{y}}] - \mathbb{E}[\tilde{\mathbf{y}}])^2], \\
&= \mathbb{E}\left[\mathbf{f}^2 + \mathbf{f}\boldsymbol{\epsilon} - \mathbf{f}\tilde{\mathbf{y}} + \mathbf{f}\mathbb{E}[\tilde{\mathbf{y}}] - \mathbf{f}\mathbb{E}[\tilde{\mathbf{y}}]\right. \\
&\quad + \boldsymbol{\epsilon}\mathbf{f} + \boldsymbol{\epsilon}^2 - \boldsymbol{\epsilon}\tilde{\mathbf{y}} + \boldsymbol{\epsilon}\mathbb{E}[\tilde{\mathbf{y}}] - \boldsymbol{\epsilon}\mathbb{E}[\tilde{\mathbf{y}}] \\
&\quad - \tilde{\mathbf{y}}\mathbf{f} - \tilde{\mathbf{y}}\boldsymbol{\epsilon} + \tilde{\mathbf{y}}^2 - \tilde{\mathbf{y}}\mathbb{E}[\tilde{\mathbf{y}}] + \tilde{\mathbf{y}}\mathbb{E}[\tilde{\mathbf{y}}] \\
&\quad + \mathbb{E}[\tilde{\mathbf{y}}]\mathbf{f} + \mathbb{E}[\tilde{\mathbf{y}}]\boldsymbol{\epsilon} - \mathbb{E}[\tilde{\mathbf{y}}]\tilde{\mathbf{y}} + (\mathbb{E}[\tilde{\mathbf{y}}])^2 - (\mathbb{E}[\tilde{\mathbf{y}}])^2 \\
&\quad \left. - \mathbb{E}[\tilde{\mathbf{y}}]\mathbf{f} - \mathbb{E}[\tilde{\mathbf{y}}]\boldsymbol{\epsilon} + \mathbb{E}[\tilde{\mathbf{y}}]\tilde{\mathbf{y}} - (\mathbb{E}[\tilde{\mathbf{y}}])^2 + (\mathbb{E}[\tilde{\mathbf{y}}])^2\right], \\
&= \mathbb{E}\left[\mathbf{f}^2 + \mathbf{f}\boldsymbol{\epsilon} + \boldsymbol{\epsilon}\mathbf{f} + \boldsymbol{\epsilon}^2 - \mathbf{f}\mathbb{E}[\tilde{\mathbf{y}}] - \boldsymbol{\epsilon}\mathbb{E}[\tilde{\mathbf{y}}] - \mathbb{E}[\tilde{\mathbf{y}}]\mathbf{f} - \mathbb{E}[\tilde{\mathbf{y}}]\boldsymbol{\epsilon} + (\mathbb{E}[\tilde{\mathbf{y}}])^2\right] \\
&\quad + \mathbb{E}\left[\tilde{\mathbf{y}}^2 - \tilde{\mathbf{y}}\mathbb{E}[\tilde{\mathbf{y}}] - \mathbb{E}[\tilde{\mathbf{y}}]\tilde{\mathbf{y}} + (\mathbb{E}[\tilde{\mathbf{y}}])^2\right] \\
&\quad + \mathbb{E}\left[-\mathbf{f}\tilde{\mathbf{y}} - \boldsymbol{\epsilon}\tilde{\mathbf{y}} + \mathbf{f}\mathbb{E}[\tilde{\mathbf{y}}] + \boldsymbol{\epsilon}\mathbb{E}[\tilde{\mathbf{y}}] - \tilde{\mathbf{y}}\mathbf{f} - \tilde{\mathbf{y}}\boldsymbol{\epsilon} + \tilde{\mathbf{y}}\mathbb{E}[\tilde{\mathbf{y}}] + \mathbb{E}[\tilde{\mathbf{y}}]\mathbf{f} + \mathbb{E}[\tilde{\mathbf{y}}]\boldsymbol{\epsilon} + \mathbb{E}[\tilde{\mathbf{y}}]\tilde{\mathbf{y}} - 2(\mathbb{E}[\tilde{\mathbf{y}}])^2\right].
\end{aligned}$$

Before we move further we may note that the exact function  $f(\mathbf{x})$  generally is not known, and we may therefore assume that our data is a good representation and replace  $\mathbf{f}$  with  $\mathbf{y}$  in the expression above. In practise this  $\mathbf{y}$  is then the part of the data set that we have chosen as test set, while the model is trained with the remaining data set (the training set). Thus, using that  $\mathbb{E}[\mathbb{E}[\mathbf{x}]] = \mathbb{E}[\mathbf{x}]$ ,  $\mathbb{E}[(\mathbb{E}[\mathbf{x}])^2] = (\mathbb{E}[\mathbf{x}])^2$  and  $\mathbb{E}[\mathbf{x}, \mathbf{y}] = \mathbb{E}[\mathbf{x}]\mathbb{E}[\mathbf{y}]$  for any statistically independent  $\mathbf{x}$  and  $\mathbf{y}$ , and that  $\mathbb{E}[\boldsymbol{\epsilon}] = 0$  so that we can remove all first order terms in  $\boldsymbol{\epsilon}$ , we get

$$\begin{aligned}
\mathbb{E}[(\mathbf{y} - \tilde{\mathbf{y}})^2] &= \mathbb{E}\left[\mathbf{y}^2 + \boldsymbol{\epsilon}^2 - \mathbf{y}\mathbb{E}[\tilde{\mathbf{y}}] - \mathbb{E}[\tilde{\mathbf{y}}]\mathbf{y} + (\mathbb{E}[\tilde{\mathbf{y}}])^2\right] \\
&\quad + \mathbb{E}\left[\tilde{\mathbf{y}}^2 - \tilde{\mathbf{y}}\mathbb{E}[\tilde{\mathbf{y}}] - \mathbb{E}[\tilde{\mathbf{y}}]\tilde{\mathbf{y}} + (\mathbb{E}[\tilde{\mathbf{y}}])^2\right] \\
&\quad + \mathbb{E}\left[-\mathbf{y}\tilde{\mathbf{y}} + \mathbf{y}\mathbb{E}[\tilde{\mathbf{y}}] - \tilde{\mathbf{y}}\mathbf{y} + \tilde{\mathbf{y}}\mathbb{E}[\tilde{\mathbf{y}}] + \mathbb{E}[\tilde{\mathbf{y}}]\mathbf{y} + \mathbb{E}[\tilde{\mathbf{y}}]\tilde{\mathbf{y}} - 2(\mathbb{E}[\tilde{\mathbf{y}}])^2\right], \\
&= \mathbb{E}\left[(\mathbf{y} - \mathbb{E}[\tilde{\mathbf{y}}])^2\right] + \mathbb{E}[\boldsymbol{\epsilon}^2] + \mathbb{E}\left[(\tilde{\mathbf{y}} - \mathbb{E}[\tilde{\mathbf{y}}])^2\right] \\
&\quad - \mathbb{E}[\mathbf{y}]\mathbb{E}[\tilde{\mathbf{y}}] + \mathbb{E}[\mathbf{y}]\mathbb{E}[\tilde{\mathbf{y}}] - \mathbb{E}[\tilde{\mathbf{y}}]\mathbb{E}[\mathbf{y}] + \mathbb{E}[\tilde{\mathbf{y}}]\mathbb{E}[\tilde{\mathbf{y}}] + \mathbb{E}[\tilde{\mathbf{y}}]\mathbb{E}[\mathbf{y}] + \mathbb{E}[\mathbf{y}]\mathbb{E}[\tilde{\mathbf{y}}] - 2(\mathbb{E}[\tilde{\mathbf{y}}])^2, \\
&= \text{Bias}[\tilde{\mathbf{y}}] + \text{Var}[\tilde{\mathbf{y}}] + \sigma^2.
\end{aligned} \tag{B7}$$

## Appendix C: Additional Figures

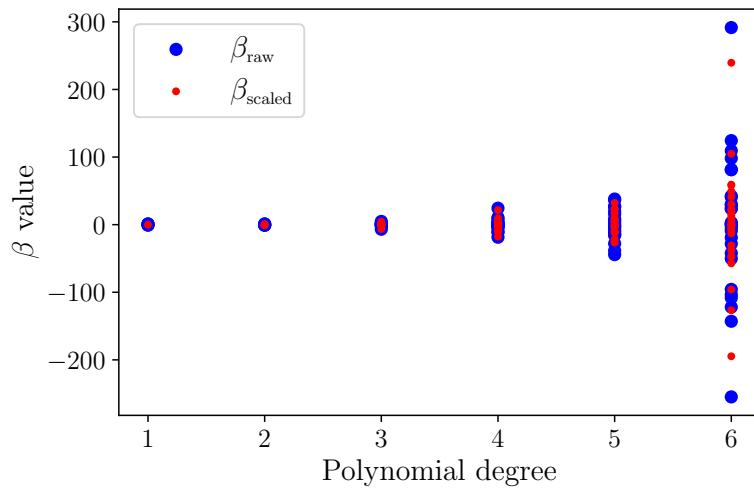


FIG. 16: The coefficients  $\beta$  for the Franke function fit estimated using unscaled (blue) and scaled (red) data, plotted for polynomial degrees in the interval [1, 6].

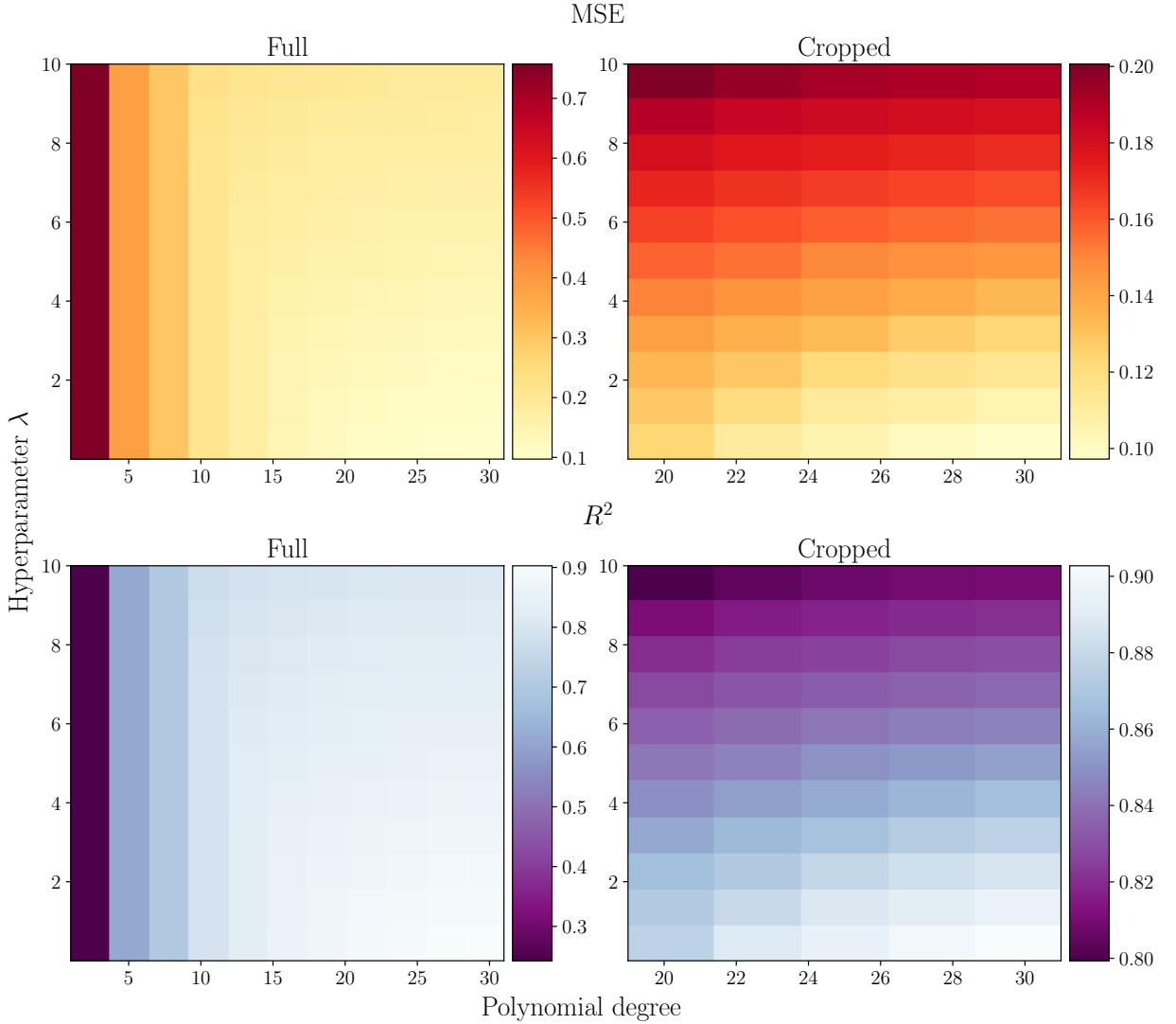


FIG. 17: MSE (upper row) and  $R^2$  (lower row) for Ridge regression evaluated on the cosmological test data and contour plotted as functions of polynomial degree (horizontal axes) and hyperparameter  $\lambda$  (vertical axes). We have used 11 evenly spaced degrees in the interval  $[1, 31]$  and 11 logarithmically spaced values for  $\lambda$  in the interval  $[10^{-5}, 10^{-1}]$ . In the right column we have cropped away degrees smaller than 19 purely for visualization purposes.

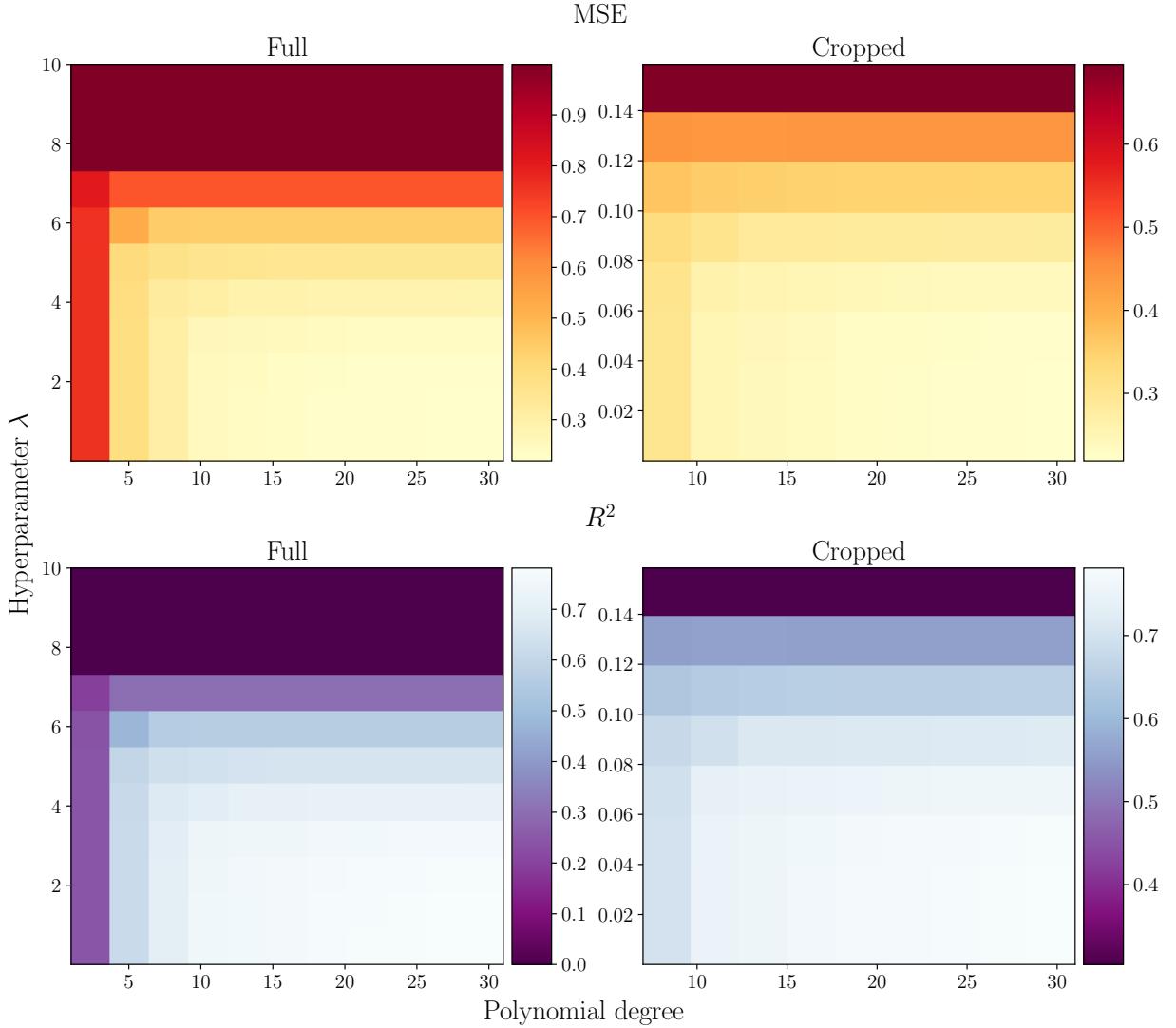


FIG. 18: MSE (upper row) and  $R^2$  (lower row) for Lasso regression evaluated on the cosmological test data and contour plotted as functions of polynomial degree (horizontal axes) and hyperparameter  $\lambda$  (vertical axes). We have used 11 evenly spaced degrees in the interval  $[1, 31]$  and 11 logarithmically spaced values for  $\lambda$  in the interval  $[10^{-5}, 10^{-1}]$ , but in the right column we have cropped away degrees smaller than 19 and the 3 largest  $\lambda$  values.

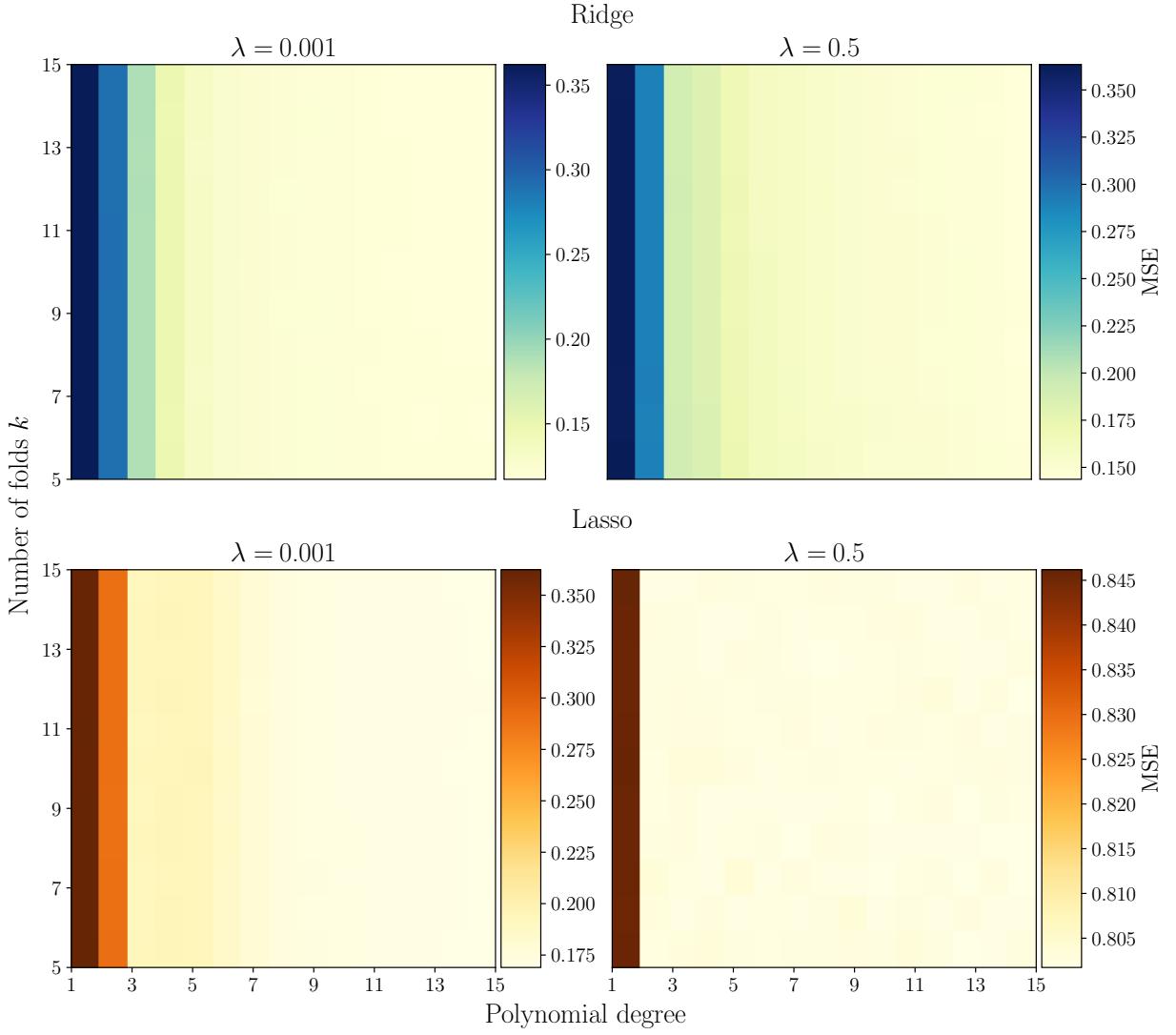


FIG. 19: MSE evaluated on Franke test data as function of polynomial degrees in the interval  $[1, 15]$  and number of folds  $k \in [5, 15]$  used in cross-validation, for Ridge (upper row) and Lasso (lower row) regression. We have used  $\lambda = 0.001$  and  $\lambda = 0.5$  in the left and right columns, respectively.

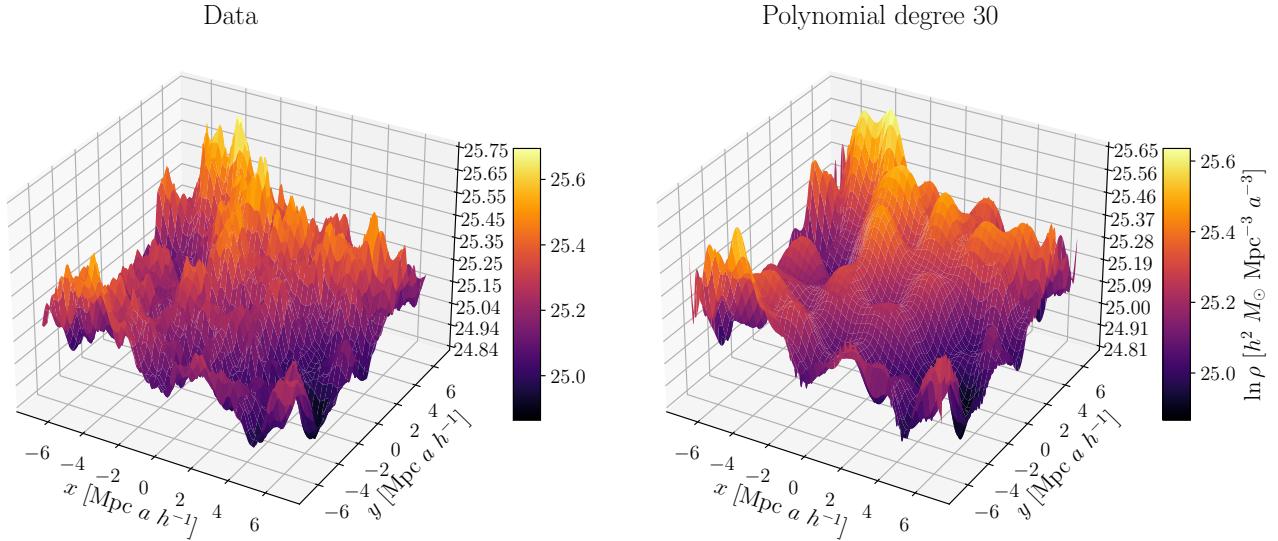


FIG. 20: Surface plots of the cosmological simulation data (left) and the OLS fit for polynomial degree 30 (right).

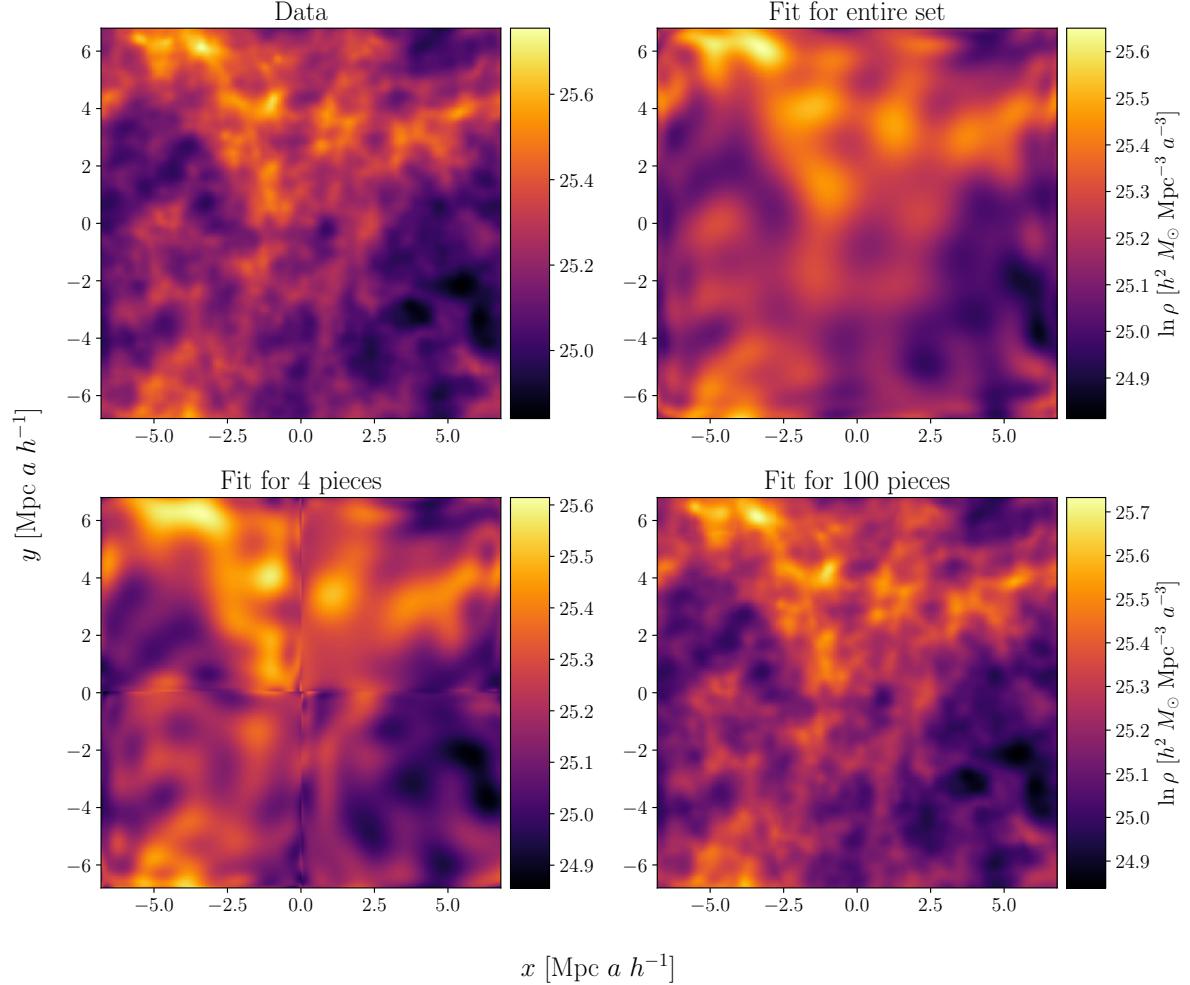


FIG. 21: Contour plots of the cosmological simulation data (upper left) and the OLS fits for the entire data (upper right), 4 separate pieces stitched back together (lower left) and 100 separate pieces stitched back together (lower right). We have used a polynomial degree of 30 for all fits.