

FYS-STK4155 Project 1

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Abstract

Introduction

Data

In this report, we will be working with two separate datasets.

The Franke Function

The first dataset will be given by the *Franke function*, which is defined as follows:

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

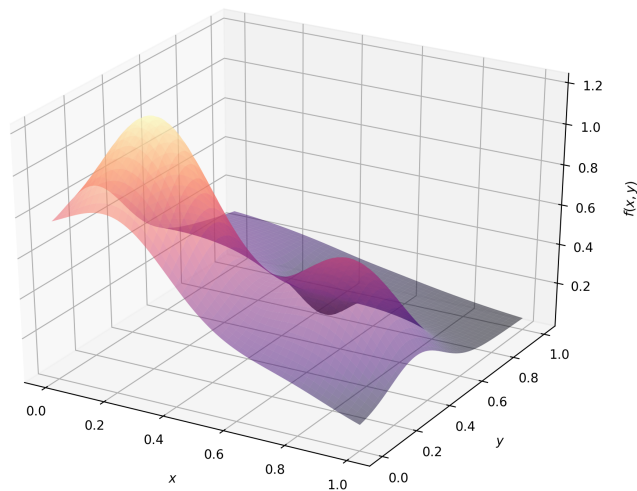


Figure 1: The *Franke function* for x and y values ranging from zero to one.

We will be solving the Franke function for 100 x -values and 100 y -values in the range $[0, 1]$, leaving us with a grid containing a total of 10000 xy coordinate pairs. This leaves us with the values plotted in Figure 1.

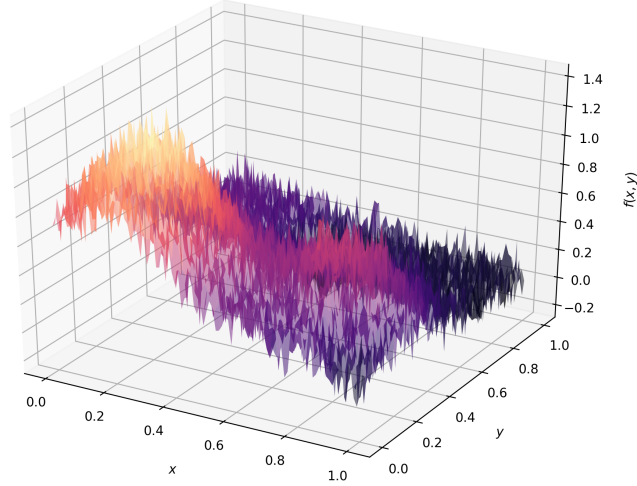


Figure 2: The *Franke function* for x and y values ranging from zero to one, with a Gaussian noise $N(0, 0.01)$

In addition, we will also be adding *Gaussian noise* to each value $f(x, y)$, such that we are left with values as seen in Figure 2.

Møsvatn Austfjell

For our second dataset, we will be using real data taken from the *U.S. Geological Survey* [1] official website <https://earthexplorer.usgs.gov/>. More specifically, we will be using a `.tif` file containing altitude data for a rectangular region of Møsvatn Austfjell shown in Figure 3.

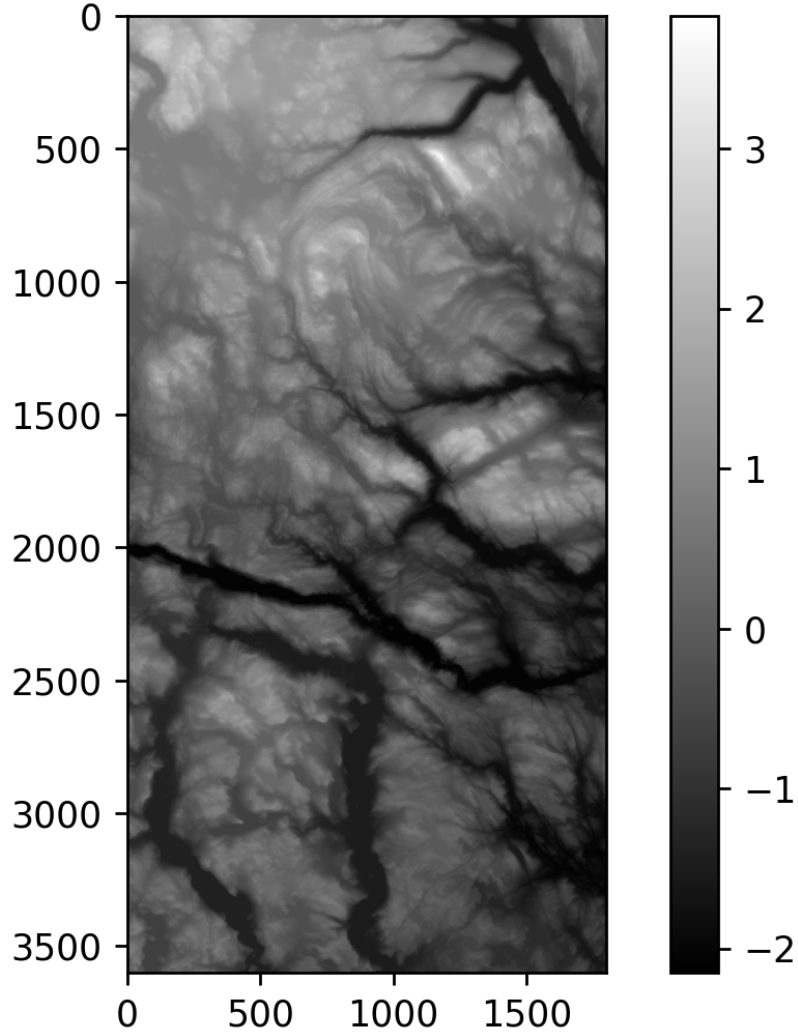


Figure 3: Altitude data for Møsvatn Austfjell, from the USGS website [1].

Method

Generalization of Multidimensional Polynomials

If we wish to construct a p -dimensional polynomial of degree d , we need to know what terms need to be included to give us a completely generalized polynomial. For a 1-D polynomial of second degree, we would have three terms:

$$f(x) = \beta_1 + \beta_2 x + \beta_3 x^2$$

Where β is a vector containing each coefficient. For a 2-D polynomial of second degree, we would have 6 terms:

$$f(x, y) = \beta_1 + \beta_2 x + \beta_3 y + \beta_4 xy + \beta_5 x^2 + \beta_6 y^2$$

And for a 3-D polynomial of second degree, we would have 10 terms:

$$f(x, y, z) = \beta_1 + \beta_2 x + \beta_3 y + \beta_4 z + \beta_5 xy + \beta_6 xz + \beta_7 yz + \beta_8 x^2 + \beta_9 y^2 + \beta_{10} z^2$$

There are many possible combinations of p and d , and the number of terms blows up significantly as these values increase. We can, however, create a general expression [2] for any p and d using summation notation:

$$f(\mathbf{x}) = \sum_{\sum_{j=1}^d i_j \leq p} \left(\beta_{i_1, i_2, \dots, i_d} \prod_{k=1}^d x_k^{i_k} \right) \quad (1)$$

Alternatively, a simple `python` script can be used to find all the terms' exponents by calculating all permutations of the natural numbers from zero to d in sets of length p , then removing all results whose sum is greater than d .

```

1 powers = np.arange(0, degree + 1, 1)
2 powers = np.repeat(powers, p)
3 exponents = list(permutations(powers, p))
4 exponents = np.unique(exponents, axis = 0)
5
6 if p != 1:
7     expo_sum = np.sum(exponents, axis = 1)
8     valid_idx = np.where(np.less_equal(expo_sum, degree))[0]
9     exponents = np.array(exponents, dtype = np.int64)
10    exponents = exponents[valid_idx]
11 else:
12    exponents = np.array(exponents, dtype = np.int64)

```

Ordinary Least-Squares (OLS) Regression

We are given a $p+1$ -dimensional dataset¹ consisting of N datapoints per feature such that:

$$\mathbf{X} = \begin{bmatrix} X_{1,1} & X_{1,2} & \cdots & X_{1,p} \\ X_{2,1} & X_{2,2} & \cdots & X_{2,p} \\ \vdots & \vdots & \ddots & \vdots \\ X_{N,1} & X_{N,2} & \cdots & X_{N,p} \end{bmatrix} \quad \mathbf{y} = \begin{bmatrix} y_1 \\ y_2 \\ \vdots \\ y_N \end{bmatrix} \quad (2)$$

Where the $N \times p$ matrix \mathbf{X} contains the dataset's *input data*, and the N -vector \mathbf{y} contains its *output data*, such that each row in \mathbf{X} corresponds to a single output in \mathbf{y} .

Now, we wish to find a p -dimensional polynomial of degree d which most closely matches our dataset. We will need a design matrix \mathbf{A} ; this will require using the knowledge presented in (1), since a design matrix should contain each polynomial term as an individual column.

Next, we will be using the method of *least squares* [3], whereby we attempt to minimize the *residual sum of squares*:

¹Meaning a set of p input features and 1 output.

$$RSS(\beta) = \sum_{i=1}^N (y_i - \mathbf{A}_i^T \beta)^2$$

Where \mathbf{A}_i represents the i^{th} row in \mathbf{A} , and β is the set of coefficients in the aforementioned polynomial; in matrix form, this can be written more concisely:

$$RSS(\beta) = (\mathbf{y} - \mathbf{A}\beta)^T(\mathbf{y} - \mathbf{A}\beta)$$

To minimize the RSS , we can differentiate it with respect to β and set the right-hand side equal to zero – this allows us to solve for β , which will give us the coefficients to the polynomial that best matches our dataset:

$$\mathbf{A}^T(\mathbf{y} - \mathbf{A}\beta) = 0 \iff \mathbf{A}^T \mathbf{y} = \mathbf{A}^T \mathbf{A} \beta$$

Solving for β then gives us our desired result:

$$\beta = (\mathbf{A}^T \mathbf{A})^{-1} \mathbf{A}^T \mathbf{y} \quad (3)$$

Using the set of coefficients given by (3), we can then match the dataset from (2) as effectively as possible.

Ridge Regression

The solution for β given in (3) can be used without issue in many cases, but if the matrix \mathbf{A} is singular², we run into an issue – namely, we cannot take the inverse of a singular matrix! As a result, we must look to more robust methods; one such method is called *ridge regression*.

The process of obtaining our vector of coefficients β via ridge regression is functionally very similar to that of OLS. The main difference is that we include an extra term in the residual sum of squares:

$$RSS(\beta) = \sum_{i=1}^N (y_i - \mathbf{A}_i^T \beta)^2 + \lambda \sum_{i=1}^N \beta^2$$

In matrix form, this can be rewritten:

$$RSS(\beta) = (\mathbf{y} - \mathbf{A}\beta)^T(\mathbf{y} - \mathbf{A}\beta) + \lambda \beta^T \beta$$

Where λ , known as the *hyperparameter*, is a scalar value. Performing the same process as in the previous subsection, we are left with a solution similar to that in (3):

$$\beta = (\mathbf{A}^T \mathbf{A} + \lambda \mathbf{I})^{-1} \mathbf{A}^T \mathbf{y} \quad (4)$$

In cases where \mathbf{A} is singular, it is therefore possible to make very few changes to the OLS algorithm and still get a good result, one must simply optimize the hyperparameter and find a λ that minimizes the *mean squared error* (yet to be introduced) of our polynomial approximation.

²Meaning that $\det(\mathbf{A}) = 0$

LASSO Regression

The *least absolute shrinkage and selection operator*, commonly abbreviated as *LASSO*, is a method that implements the **L1** norm in place of the **L2** (or Euclidian) norm used in ridge regression. The residual sum of squares is therefore given by:

$$RSS(\beta) = \sum_{i=1}^N (y_i - \mathbf{A}_i^T \beta)^2 + \lambda \sum_{i=1}^N |\beta| \quad (5)$$

Unfortunately, differentiating the above with respect to β will not work as intended, since we cannot take the matrix-form derivative of $\lambda \sum_{i=1}^N |\beta|$. As a result, we must use an iterative *gradient descent* method to minimize the right-hand side of (5).

Algorithm 1 The LASSO algorithm, over the course of 500 iterations.

```

1:  $z = \sum_i A_i^2$ 
2:  $i = 0$ 
3: while  $i \leq 500$  do
4:    $i = i + 1$ 
5:    $j = 0$ 
6:   while  $j \leq p$  do
7:      $\hat{y} = \sum_{k \neq j} \beta A_{*,k}$ 
8:      $\rho = \sum_k A_{*,k}(\mathbf{y} - \hat{\mathbf{y}})$ 
9:     if  $\rho < -\lambda/2$  then
10:       $\beta_j = (\rho + \lambda/2)/z_j$ 
11:     else if  $\rho > \lambda/2$  then
12:       $\beta_j = (\rho - \lambda/2)/z_j$ 
13:     else
14:       $\beta_j = 0$ 
15:     end if
16:   end while
17: end while

```

Mean Squared Error

To get a measure of success with respect to the implemented method and parameters, we can calculate the mean difference in the squares of each measured output y_i and their respective predicted outputs \hat{y}_i :

$$MSE(\mathbf{y}, \hat{\mathbf{y}}) = \frac{1}{N} \sum_{i=1}^N (y_i - \hat{y}_i)^2 = \mathbb{E}[(\mathbf{y} - \hat{\mathbf{y}})^2] \quad (6)$$

The lower the *MSE*, the closer the polynomial approximation is to the original dataset. If it is too low, however, we run the risk of overfitting our dataset, which is not desirable either – fortunately, this not an issue within the scope of this report.

R² Score

Another measure of success is the *coefficient of determination*, colloquially known as the R^2 score, is given by the following expression:

$$R^2 = 1 - \frac{\sum_{i=1}^N (y_i - \hat{y}_i)^2}{\sum_{i=1}^N (y_i - \bar{y}_i)^2} \quad (7)$$

The closer R^2 is to one, the closer the polynomial approximation is to the input/output dataset, although a perfect score can once again arise due to overfitting just as in the case of the MSE .

Bias-Variance Tradeoff

Before we continue, we can decompose the range of outputs \mathbf{y} as follows:

$$\mathbf{y}(\mathbf{X}) = f(\mathbf{X}) + N(0, \sigma) \quad (8)$$

Where $f(\mathbf{X})$ represents the *actual* function used to generate the dataset, and $N(0, \sigma)$ is a Gaussian noise with a standard deviation of σ .

As a regression model increases in complexity³, it so happens that the *variance* of a prediction increases. Variance is defined as follows:

$$\text{Var}(\mathbf{y}) = \frac{1}{n} \sum_{i=1}^n (y_i - \mathbb{E}[\mathbf{y}])^2 \quad (9)$$

On the other hand, we have that the *bias* of the prediction decreases as the complexity increases. We define the bias as:

$$\text{Bias}(\mathbf{y}) = \frac{1}{n} \sum_{i=1}^n (f_i - \mathbb{E}[\mathbf{y}])^2 \quad (10)$$

Note that in order to calculate the bias, we need to know the original function f used to generate \mathbf{y} .

Interestingly enough, taking the sum of (9) and (10) as well as σ^2 will yield the *mean squared error*. We will show this to be the case in the following subsection.

Derivation

We wish to show that:

$$\mathbb{E}[(\mathbf{y} - \hat{\mathbf{y}})^2] = \frac{1}{n} \sum_{i=1}^n (f_i - \mathbb{E}[\hat{\mathbf{y}}])^2 + \frac{1}{n} \sum_{i=1}^n (\hat{y}_i - \mathbb{E}[\hat{\mathbf{y}}])^2 + \sigma^2 \quad (11)$$

We begin by rewriting the MSE into summation notation, decomposing the terms as defined in (8), and adding/subtracting a term $\mathbb{E}[\hat{\mathbf{y}}]$:

³For a polynomial regression, this would refer to its *degree*.

$$\begin{aligned}\mathbb{E}[(\mathbf{y} - \hat{\mathbf{y}})^2] &= \frac{1}{n} \sum_{i=1}^n (y_i - \hat{y}_i)^2 = \frac{1}{n} \sum_{i=1}^n (f_i + \varepsilon - \hat{y}_i)^2 \\ &= \frac{1}{n} \sum_{i=1}^n (f_i + \varepsilon - \hat{y}_i + \mathbb{E}[\hat{\mathbf{y}}] - \mathbb{E}[\hat{\mathbf{y}}])^2\end{aligned}$$

Next, we set $a \equiv f_i - \mathbb{E}[\hat{\mathbf{y}}]$ and $b \equiv \hat{y}_i - \mathbb{E}[\hat{\mathbf{y}}]$ and expand:

$$\frac{1}{n} \sum_{i=1}^n (a - b + \varepsilon)^2 = \frac{1}{n} \sum_{i=1}^n (a^2 - 2ab + b^2 - 2b\varepsilon + \varepsilon^2 + 2a\varepsilon)$$

The next few steps are messy, and require lots of algebraic manipulation:

$$\begin{aligned}& \frac{1}{n} \sum_{i=1}^n (f_i - \mathbb{E}[\hat{\mathbf{y}}])^2 + \frac{1}{n} \sum_{i=1}^n (\varepsilon^2) + \frac{1}{n} \sum_{i=1}^n (\hat{y}_i - \mathbb{E}[\hat{\mathbf{y}}])^2 - \frac{2}{n} \sum_{i=1}^n \varepsilon(\hat{y}_i - \mathbb{E}[\hat{\mathbf{y}}]) + \\ & \frac{2}{n} \sum_{i=1}^n \varepsilon(f_i - \mathbb{E}[\hat{\mathbf{y}}]) - \frac{2}{n} \sum_{i=1}^n (f_i - \mathbb{E}[\hat{\mathbf{y}}])(\hat{y}_i - \mathbb{E}[\hat{\mathbf{y}}]) = \\ & \frac{1}{n} \sum_{i=1}^n (f_i - \mathbb{E}[\hat{\mathbf{y}}])^2 + \frac{1}{n} \sum_{i=1}^n (\hat{y}_i - \mathbb{E}[\hat{\mathbf{y}}])^2 + \sigma^2 - \mathbb{E}[\varepsilon] \frac{2}{n} \sum_{i=1}^n (\hat{y}_i - \mathbb{E}[\hat{\mathbf{y}}]) + \\ & \mathbb{E}[\varepsilon] \frac{2}{n} \sum_{i=1}^n (f_i - \mathbb{E}[\hat{\mathbf{y}}]) - \frac{2}{n} \sum_{i=1}^n (f_i - \mathbb{E}[\hat{\mathbf{y}}])(\hat{y}_i - \mathbb{E}[\hat{\mathbf{y}}])\end{aligned}$$

Finally, we see that our original assumption given by (11) is correct.

$$= \frac{1}{n} \sum_{i=1}^n (f_i - \mathbb{E}[\hat{\mathbf{y}}])^2 + \frac{1}{n} \sum_{i=1}^n (\hat{y}_i - \mathbb{E}[\hat{\mathbf{y}}])^2 + \sigma^2 \quad \square$$

Where $\frac{1}{n} \sum_{i=1}^n (f_i - \mathbb{E}[\hat{\mathbf{y}}])$ is clearly the *bias* and $\frac{1}{n} \sum_{i=1}^n (\hat{y}_i - \mathbb{E}[\hat{\mathbf{y}}])^2$ is the *variance*.

Cross-Validation

So far, all our methods involving validation⁴ may have involved the separation of our data into a training and testing set, whereby the vector of coefficients is calculated using the *training* set, and the validation is performed on the *testing* set that remains. There is however one more way to obtain a clearer picture of how well a model works: *k-fold* cross validation.

In short, once the training and testing set have been separated, we can choose a value for k . Next, we divide the training set into k equally sized parts⁵. The next step is to be performed k times; here we take $k - 1$ of the parts and combine them into a temporary training set, and leave the last part as our testing set, and we perform validation on the testing set, and save the

⁴This refers to calculating the *MSE*, variance, bias, and so on.

⁵We can have slightly unequal-sized parts without it being an issue, if the size of the dataset doesn't divide perfectly into k .

values. During each iteration, we must shuffle our parts such that each step has a unique training-testing split.

Finally, we can take all the calculated MSE values, among others, and take their average. This leaves us with a well-rounded result *without* the need for more input data!

Results

Discussion

Appendix

References

- [1] “Earthexplorer.”
- [2] M. (<https://math.stackexchange.com/users/58320/macavity>), “What is the general form of a polynomial of degree n and with m variables?.” Mathematics Stack Exchange. URL:<https://math.stackexchange.com/q/2482654> (version: 2017-10-21).
- [3] T. Hastie, R. Tibshirani, and J. H. Friedman, *The elements of statistical learning: data mining, inference, and prediction*. Springer, 2013.