

# DS6410: Homework 2

Due: Mar 2, 2026

**Q1. Fourier Basis Expansion for Image Modeling.** In this problem, you will study *Fourier basis expansion* as a method for representing and learning nonlinear structure in imaging data, combining theoretical understanding with practical implementation.

Let  $I(u, v)$  denote a grayscale image of size  $H \times W$ , where  $(u, v)$  indexes pixel location. Treat the image as a function

$$f(u, v) = I(u, v).$$

Consider the following real-valued 2D Fourier basis functions:

$$\phi_{k\ell}^{(c)}(u, v) = \cos(2\pi k u) \cos(2\pi \ell v), \quad \phi_{k\ell}^{(s)}(u, v) = \sin(2\pi k u) \sin(2\pi \ell v),$$

for integers  $k, \ell \geq 0$ .

1. Explain how the collection of functions

$$\{\phi_{k\ell}^{(c)}(u, v), \phi_{k\ell}^{(s)}(u, v)\}_{k,\ell}$$

defines a *basis expansion* for functions on  $[0, 1]^2$ .

2. Consider the truncated Fourier expansion

$$f_K(u, v) = \sum_{k=0}^K \sum_{\ell=0}^K [a_{k\ell} \cos(2\pi k u) \cos(2\pi \ell v) + b_{k\ell} \sin(2\pi k u) \sin(2\pi \ell v)].$$

- How does the truncation level  $K$  control model complexity?
- Discuss the bias–variance tradeoff as  $K$  increases.
- Why does overfitting become a concern for large  $K$ ?

3. Coding Part.

- Constructing Fourier features:

- Load a grayscale image and normalize pixel coordinates to  $[0, 1]^2$ .  
(You may use any standard grayscale image.)

- ii. Sample  $n$  pixel locations  $\{(u_i, v_i)\}_{i=1}^n$  and form responses  $y_i = I(u_i, v_i)$ .
  - iii. For a given  $K$ , construct the design matrix  $\Phi_K$  using a truncated 2D Fourier basis.
- (b) Fitting and regularization:
- i. Fit a linear model using  $\Phi_K$  via:
    - ordinary least squares (when  $p_K \leq n$ ),
    - ridge regression (choose  $\lambda$  by cross-validation).
  - ii. For several values of  $K$ , report training and test mean squared error.
- (c) Reconstruction and visualization
- i. Use the fitted model to predict intensities at all pixel locations.
  - ii. Reconstruct and display the estimated image  $\hat{I}_K$  for at least three values of  $K$ .

**Q2. Ridge Regression as Bayesian MAP Estimation.** Consider the linear model

$$y = X\beta + \varepsilon, \quad \varepsilon \sim \mathcal{N}(0, \sigma^2 I_n),$$

where  $y \in \mathbb{R}^n$ ,  $X \in \mathbb{R}^{n \times p}$ , and  $\beta \in \mathbb{R}^p$ .

1. **Gaussian prior.** Assume a prior distribution on the regression coefficients

$$\beta \sim \mathcal{N}(0, \tau^2 I_p).$$

Derive the **posterior distribution**  $p(\beta | y)$ , and give its mean and covariance in closed form.

2. **MAP estimator.** Derive the maximum a posteriori (MAP) estimator

$$\hat{\beta}_{\text{MAP}} := \arg \max_{\beta} p(\beta | y),$$

and show that it is the unique minimizer of an optimization problem of the form

$$\min_{\beta \in \mathbb{R}^p} \frac{1}{2} \|y - X\beta\|_2^2 + \lambda \|\beta\|_2^2.$$

3. **Ridge connection.** Conclude that  $\hat{\beta}_{\text{MAP}}$  coincides with the ridge regression estimator, and identify the relationship between the ridge penalty parameter  $\lambda$  and the variance parameters  $\sigma^2$  and  $\tau^2$ .

*Hint:* Write the log posterior  $\log p(\beta | y)$  up to an additive constant and compare it to a penalized least squares objective.

**Q3. Coding: Group Lasso on bardet.** In this problem, you will fit and interpret a *group lasso* model on an established dataset. Bardet is gene expression data (20 genes for 120 samples) from the microarray experiments of mammalian eye tissue samples of Scheetz et al. (2006).

**Goal.** Compare **group lasso** against **lasso** and **ridge** in terms of predictive performance and interpretability.

**Dataset.** Use the dataset `bardet`:

1. **Load and construct  $(X, y)$  and groups.**
  - (a) Load the data with `data(bardet)` and inspect the resulting object(s). Identify the response vector  $y$  and the predictor matrix  $X$  (or construct  $X$  from a `data.frame`).
  - (b) Define a *group membership vector*  $g \in \{1, \dots, G\}^p$  for the  $p$  columns of  $X$ . Each 5 consecutive columns to a grouped gene.
2. **Train/test split and standardization.** Create a train/test split (e.g., 70/30) using a fixed random seed. Standardize the columns of  $X$  using *training-set* means and standard deviations, and apply the same transformation to the test set.
3. **Fit group lasso and tune  $\lambda$ .** Fit a group lasso model on the training data by solving

$$\min_{\beta \in \mathbb{R}^p} \frac{1}{2n_{\text{tr}}} \|y_{\text{tr}} - X_{\text{tr}}\beta\|_2^2 + \lambda \sum_{k=1}^G w_k \|\beta_{(k)}\|_2,$$

where  $\beta_{(k)}$  denotes the subvector of coefficients in group  $k$  and  $w_k$  are optional group weights (e.g.,  $w_k = \sqrt{|k|}$ ). Choose  $\lambda$  using  $K$ -fold cross-validation on the training set.

**Report:** the selected  $\lambda$  and a plot of CV error versus  $\log(\lambda)$ .

4. **Fit comparison models (lasso and ridge).** Fit a lasso model and a ridge regression model on the same training data, selecting  $\lambda$  via cross-validation. **Report:** the selected  $\lambda$  for each method, and a plot of CV error versus  $\log(\lambda)$  for each method (overlay is fine).
5. **Evaluate predictive performance.** Evaluate each fitted model on the test set using an appropriate metric. **Report:** the test-set metric for group lasso, lasso, and ridge in a small table.
6. **Interpretation: groups vs individual predictors.**
  - (a) For group lasso, report which *groups* are selected (i.e., groups with  $\|\hat{\beta}_{(k)}\|_2 > 0$ ).
  - (b) For lasso, report which *individual* predictors are selected (nonzero coefficients).

- (c) Compare sparsity patterns and interpretability: does group lasso yield a more interpretable model for `bardet`? Explain briefly with any supporting output/plots.
7. **Stability check.** Repeat Steps 2–6 for three different random seeds. Comment on the stability of selected groups/features across splits.

**Submission:** Submit your code (well-documented) and a short write-up containing the requested plots, tables, and discussion.