5.2 Cross-Validation and Regularization

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def rmse(y, y_predicted):

In practical applications the ground truth function will not be given, yet we still have to find a way to balance bias and variance of our models. In order to judge these statistics, one can either keep a disjunct validation set or apply cross-validation if training data is scarce. It is also good practice to keep an additional test set, that is untouched for optimization, in order to judge future performance. The most common method against overfitting is keeping the L2-Norm of the weights as small as possible (also known as Ridge Regression for linear models). This exercise will let you explore the relationship between these concepts.

Task a) Cross-validation and Regularization. * Implement Ridge Regression for polynomial models. Don't use sklearn here! * Implement 3-fold cross-validation and calculate training and validation errors.

Q 5.3.1: What is the effect of the regularization weight on the different evaluation statistics (training, cross-validation and test-error)?

```
In [1]: %%capture
                                %run '5.1 Bias-Variance Analysis.ipynb'
                                # We silently run your solution to the first exercise to have
                                # the previously defined functions and data available.
In [2]: import numpy as np
                               n_folds = 3
                                n_samples = 1
                                 # We draw the points - as before - from a sine curve
                                X_{cv} = np.array([np.random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.uniform(0.1, 0.9, size=(n_samples, 1)) * (-1)**i for i in random.
                                y_cv = np.array([target(X_cv[i]) for i in range(n_folds)])
                                X_{\text{test}} = \text{np.array}([-0.97, -0.4, 0.7, 0.97]).reshape((-1, 1))
                                y_test = target(X_test)
In [3]: %matplotlib inline
                                import matplotlib.pyplot as plt
                                try: import seaborn as sns
                                except ImportError: pass
                                 # Calculate the root mean square error
```

```
# Ridge regression is an extension of linear least squares.
# How do you have to adjust the linear fitting function for this?
# Remember to augment the dataset for polynomial features as before.
def fit_ridge_regression(X, y, regularization_weight):
    verify_shapes(X, y)
    # TODO(Task a):
   pass
# We can use the same method to predict with a ridge regression
# model as in exercise 5.1.
# Bookkeeping
training_errors = np.zeros((n_folds))
validation_errors = np.zeros((n_folds))
test_errors = np.zeros((n_folds))
models = np.zeros((n_folds, poly_degree + 1, 1))
# Here we define our cross-validation function specialized for
# our ridge regression models. The regularization weight is
# the only hyperparameter.
def cross_validate(regularization_weight):
    # Loop over the validation folds
    for i in range(n_folds):
        # Split the data into training and validation set
        # You can use the mask to select the correct training folds
        # X_cv[mask] will give you all folds except for the i-th one
        mask = np.arange(n_folds) != i
        # Remember the corret shape (n_samples, n_features)
        X train = None # TODO
        y_train = None # TODO
        verify_shapes(X_train, y_train)
        X_valid = None # TODO
        y_valid = None # TODO
        verify_shapes(X_valid, y_valid)
        # Train a model and determine the training and validation error
        # Luckily we wrote a function for this
        models[i] = np.nan # TODO
        # Calculate the training, validation and test error
        # for the model trained in this fold
```

return np.sqrt(np.mean(0.5 * (y - y_predicted)**2))

```
# The models can be evaluated with the predict_polynomial function
        # from the first part. We already wrote the rmse error function
        # for comparing it to the ground truth
        training_errors[i] = np.nan # TODO
        validation errors[i] = np.nan # TODO
        test_errors[i] = np.nan # TODO
    # Calculate the statistics for the whole procedure
    # This reduces to taking the mean over the folds
    # The cross-validation error is the measure we're after!
    avg_training_error = np.nan # TODO
    cross_validation_error = np.nan # TODO
    avg_test_error = np.nan # TODO
    return avg_training_error, cross_validation_error, avg_test_error
# TODO: Set the strength of the regularisation here
regularization_weight = 0.0
avg_training_error, cross_validation_error, avg_test_error = \
    cross_validate(regularization_weight)
# Show the statistics for this model class
print('Regularization weight: {:.1e}'.format(regularization_weight))
print('Average training error: {:.3f}'.format(avg_training_error))
print('Cross-validation error: {:.3f}'.format(cross_validation_error))
print('Average test error: {:.3f}'.format(avg_test_error))
# Plotting: For 3 folds, we can still look at each model and how
# well it fits the different data sets.
fix, axes = plt.subplots(n_folds, 1, figsize=(12, 16))
X_{\text{eval}} = \text{np.linspace}(-1, 1, 200).reshape((-1, 1))
y_eval = np.sin(np.pi * X_eval)
for i, axis in enumerate(axes):
    axis.set_title('Fold {:d}'.format(i))
    axis.plot(X_eval, np.sin(np.pi * X_eval), label='Target function', alpha=0.5)
    if not np.any(np.isnan(models[i])):
        axis.plot(X_eval, eval_polynomial(X_eval, models[i]),
                  label='Fitted model', c='m')
   mask = np.arange(n_folds) != i
    axis.scatter(X_cv[mask].flatten(), y_cv[mask].flatten(),
                 label='Training samples', s=30, c='k')
    axis.scatter(X_cv[i].flatten(), y_cv[i].flatten(),
                 label='Validation samples', s=40, c='g')
    axis.scatter(X_test.flatten(), y_test.flatten(),
                 label='Test samples', s=50, c='r')
    axis.set_xlim((-1.1, 1.1))
    axis.set_ylim((-1.5, 1.5))
    axis.legend(loc='best')
```

Regularization weight: 0.0e+00 Average training error: nan Cross-validation error: nan Average test error: nan

