

### Hands on Machine Learning for Fluid Dynamics



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# Lecture 4 Tutorial Exercise 1: Regression with Uncertainties

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## Consider Today's dataset



```
# Generate data

n_p=100

np.random.seed(10)

x_s=np.random.uniform(0,10,n_p)

y_s=2*x_s+2+np.random.normal(loc=0,scale=10,size=len(x_s))+x_s**2
```

#### Step 1: Fitting Lines and Parabolas



Build two Python functions to perform the linear regression of the data using the analytic solution we have derived today:

$$\mathbf{y} = \mathbf{X}\mathbf{w}$$

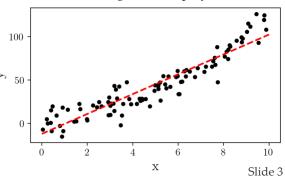
$$\mathbf{X} = \begin{bmatrix} 1 & x_0 \\ | & | \\ 1 & x_{n_p-1} \end{bmatrix} \in \mathbb{R}^{n_p \times 2} \quad \mathbf{w} = \begin{bmatrix} w_0 \\ w_1 \end{bmatrix}$$

return w

Validate your results with numpy's polyfit. Then, build another function that uses a second order polynomial.

$$\mathbf{w}^* = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{y}$$

Note that the exact same approach can be used to fit higher order polynomials.



#### Step 2: Robust (Regularized) Fitting



As we will see more in details in the Lecture 9, it is often a good idea to increase the robustness of the regression by adding a regularization. Let's see it here from a numerical stability point of view.

We are essentially solving the following linear system

$$\mathbf{H}\mathbf{w} = \mathbf{b}$$
 with  $\mathbf{H} = \mathbf{X}^T\mathbf{X}$   $\mathbf{b} = \mathbf{X}^T\mathbf{y}$ 

In a linear system, perturbations in the RHS leads to perturbation in the solution following the classic inequality:

$$\frac{||\delta \mathbf{w}||}{||\mathbf{w}||} \le \underline{\kappa(\mathbf{H})} \frac{||\delta \mathbf{b}||}{||\mathbf{b}||}$$

Condition number of H: ratio of largest to smallest eigenvalue

$$\kappa(\mathbf{H}) = \lambda_M / \lambda_m$$

To regularize means to improve (decrease) the condition number. A simple approach (Ridge regression) consist in adding a small number in the diagonal.

$$\mathbf{H}' = \mathbf{H} + \alpha \mathbf{I}$$

#### Step 2: Robust (Regularized) Fitting



Here are some questions for you

- 1. What is the condition number of the Hessian for the linear and the quadratic problem?
- 2. Derive a formula to link the Ridge coefficient alpha to the largest condition number you tolerate
- 3. Prepare two python functions to compute the weights for the linear and quadratic fit taking as used input a limiting condition number.

```
def Quadratic_s_fit_red(x,y,k_A_l=1000):
    # take input vectors x,y and fit a line
    # with coefficients w_0 and w_1
    n_p=len(x)
    X=np.zeros((n_p,3))
    X[:,0]=
    X[:,1]=
    X[:,2]=
    # Compute the Hessian
    H=X.T@(X)
    # compute the eigenvalues, the largest (1_M) and the smallest (1_m)
    Lambd=np.linalg.eig(H); 1_M=np.max(Lambd[0]); 1_m=np.min(Lambd[0]);
    alpha=
    H_p=H+alpha*np.identity(np.shape(H)[0])
    w=np.linalg.inv(H_p).dot(X.T).dot(y)
    return w
```

Note: check that the regularized Hessian has the condition number you set! (see Python's function np.linalg.cond)

#### Step 3: Uncertainty Analysis



We are interested in analyzing the uncertainties of the regression for both regularized and un-regularized regressions. We follow an ensemble approach to compute uncertainties directly from the posterior population

- 1) Write a function that computes the regression using the analytical tools for n\_e times, using each time 70% of the data for training and 30% of the data for validation.
  - 2) From the n\_e results, compute the average in sample and out of sample errors.
- 3) Then, given an arbitrary grid x\_g, provide the mean prediction and the associated uncertainties using the ensemble statistics:

$$\mu_y(x) = \frac{1}{n_e} \sum_{j=0}^{n_e-1} y_j(x)$$
  $\sigma_{\overline{y}}^2(x) = \frac{1}{n_e} \sum_{j=0}^{n_e-1} (y_j(x) - \mu_y(x))^2$ 

4) For each regression, store the resulting weights. Then look at the distribution (histogram) of each weight.

```
def Boot_Strap_Model_2_esamble(X,y,x_g,n_e=500,tp=0.3):
      poot Strap model_Z_esamble(X,y,x_g,n_e=>ue,tp=0.3):
Pop_Y=np.zeros((len(x_g),n_e)) # posterior population
mu_y=np.zeros(len(x_g)) # mean prediction
Unc_y=np.zeros(len(x_g)) # Uncertainty in the prediction
J_i_mean=np.zeros(n_e) # in sample error
J_o_mean=np.zeros(n_e) # out of sample error
w_e=np.zeros((3,n_e)) # Output distribution of weights
# Take back the x vector
     # Loop over the ensamble
for j in range(n_e):
# Split the
         xs, xss, ys, yss = train_test_split(x,y, test_size=tp)
         # Fit the weights
        w_s=
# Assign vectors to the distributions
        # Assign vectors to the distributions

w_e[:,j]=w_s

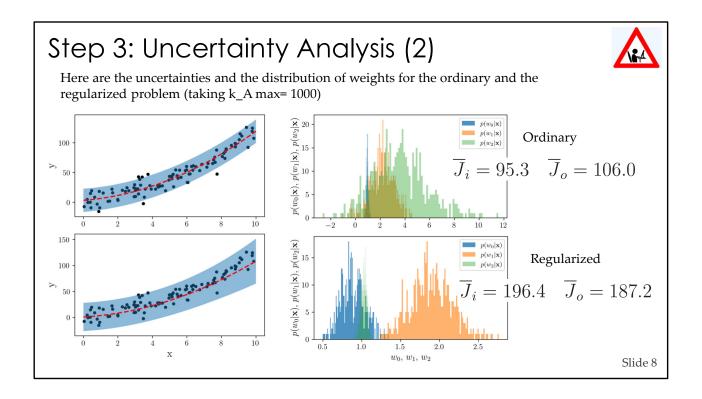
# Make in-sample prediction------

y_p_s=

# In-sample error
         J_i_mean[j]=
         # Make out-of sample prediction (and errors)
        y_p_ss=
# Out of sample error
         J_o_mean[j]=
        # Fill the population matrix Pop_Y[:,j]=
       \ensuremath{\text{\#}} Compute the mean and the uncertainty
       mu_y=
Var_y=
Var_Y_model=
       Var_y_tot=
Unc_y=1.96*np.sqrt(Var_y_tot)
       return mu_y, Unc_y, J_i_mean, J_o_mean, w_e
```

Here's a possible structure The general idea is:

- (1) Initialize all the vectors that will be exported and the matrices that will be filled
- (2) Loop over the ensemble. Use scikit learn's function train\_test\_split
- (3) Train the regression on the training data and compute performances on both training and test data
- (4) Compute the ensemble statistics using python's function np.mean and np.std



#### Step 4: Batch Gradient Descent



Write a python function to implement the mini-batch gradient descent, taking as inputs a function to compute costs and gradients, a decaying learning rate, a given number of epochs and a batch size.

First, you should write the following functions

```
def cost(w,X,y):
return J
```

```
def grad(w,X,y):
return Nabla_J
```

Then you should aim for a function of this form:

```
def Batch_GD(cost,grad,w0,X,y,eta_0,decay,n_epochs,n_batch):
    return w_opt, w_evolution, Err_SGD
```

```
def Batch_GD(cost,grad,w0,X,y,eta_0,decay,n_epochs,n_batch):
     # Prepare the loop per epoch
     n_iter=
     # number of points and features
     n_p,n_f=np.shape(X)
     # Initialize batch sample Design Matrix
     X b=
     # Current estimate of w
     # Initialize the weight evolution and the error evolution 
Err_SGD=np.zeros(n_iter); #Err_SGD[0]=cost(w,X,y) 
w_evolution=np.zeros((n_f,n_iter)); #w_evolution[:,0]=w0
     for j in range(n_iter):
    # Select randomly some data points for the batch
    # Note that replace=False means that there is no repetition
        # Construct the matrix X_b
X_b=X[Indices,:]; y_b=y[Indices]
# Get the current cost
        Err_SGD[j]=cost(w,X_b,y_b)
        #Get the gradient
Nabla_J_w=grad(w,X_b,y_b)
# Compute the learning rate
        eta=eta_0/(1+decay*j)
        # Weght update
        # Store the result in the history
        w_evolution[:,j]=w
        Mex='Iteration: {:d}; Epoch: {:d}; Cost: {:.3f}; Grad_abs: {:.3f}'\
    .format(j,j//n_batch,Err_SGD[j],np.linalg.norm(Nabla_J_w))
      w_opt=w # Final result on the weight
     return w_opt, w_evolution, Err_SGD
```



Here's a possible structure

The general idea is:

- (1) Take some random indices for the batch selection
  - (2) Sample the data
- (3) Evaluate cost and gradients
  - (4) Update the current estimate of the weights

