## Q5 ¶

Complete the f objective function in the skeleton code, which computes the objective function for Jlogistic(w). (Hint: you may get numerical overflow when computing the exponential literally, e.g. try e1000 in Numpy. Make sure to read about the log-sum-exp trick and use the numpy function logaddexp to get accurate calculations and to prevent overflow.

```
# numerical overflow check
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt

np.exp(1000)
<ipython-input-1-f582a92b4683>:6: RuntimeWarning: overflow encountered in exp
    np.exp(1000)
```

inf

Complete the fit logistic regression function in the skeleton code using the minimize function from scipy.optimize. Use this function to train a model on the provided data. Make sure to take the appropriate preprocessing steps, such as standardizing the data and adding a column for the bias term.

```
from scipy.optimize import minimize
def fit_logistic_reg(X, y, objective_function, l2_param=1):
          X: 2D numpy array of size (num_instances, num_features)
          y: 1D numpy array of size num_instances objective_function: function returning the value of the objective l2_param: regularization parameter
     Returns:
     optimal_theta: 1D numpy array of size num_features
    # initialize optimal theta
    optimal_theta = np.zeros(X.shape[1])
     # update optimal theta by using minimize function
     optimal_theta = minimize(objective_function, optimal_theta, args = (X, y, l2_param)).x
    return optimal theta
# preprocessing steps: Source from HW2 code
def feature_normalization(train, test):
     # discard features that are constant in the training set
     remove = []
     for i in range(train.shape[1]):
         if len(set(train[:, i])) == 1:
    remove.append(i)
     for i in remove:
          np.delete(train, i, axis=1)
np.delete(test, i, axis=1)
     # min-max scaling
     train_normalized = np.array(train.shape)
test_normalized = np.array(test.shape)
train_max_vals = np.max(train, axis=0)
     train_min_vals = np.min(train, axis=0)
     # use train min-max to transform both train/test dataset
train_normalized = (train - train_min_vals) / (train_max_vals - train_min_vals)
test_normalized = (test - train_min_vals) / (train_max_vals - train_min_vals)
     return train_normalized, test_normalized
# import txt files
# Import tx fites
X_train = np.loadtxt('./X_train.txt', delimiter = ',')
y_train = np.loadtxt('./y_train.txt', delimiter = ',')
X_val = np.loadtxt('./X_val.txt', delimiter = ',')
y_val = np.loadtxt('./y_val.txt', delimiter = ',')
# standardizing the data
X_train_normalized, X_val_normalized = feature_normalization(X_train, X_val)
# adding a column for the bias term
X_train_normalized = np.hstack((X_train_normalized, np.ones((X_train_normalized.shape[0], 1))))
X_{val_normalized} = np.hstack((X_{val_normalized}, np.ones((X_{val_normalized}, shape[0], 1))))
# as we assume outcome space Y = \{-1, 1\}, convert y = 0 values into -1
y_train[y_train == 0] = -1
y_val[y_val == 0] = -1
# Train model
# we need to define f_objective
optimal_theta = fit_logistic_reg(X_train_normalized, y_train, f_objective, l2_param=1)
optimal_theta
          array([ 0.00098731,
          0.001781061)
```

Find the I2 regularization parameter that minimizes the log-likelihood on the validation set. Plot the log-likelihood for different values of the regularization parameter.

```
import matplotlib.pyplot as plt

def neg_log_likelihood(theta, X, y, l2_param):
    # From Q1, we know that n*Rn(w) = NLL(w), in terms of w
    # avg_loss
    avg_loss = (1/X.shape[0]) * sum(np.logaddexp(0, -y*(X @ theta.T )))|
    # return scalar value of objective function
    objective = avg_loss

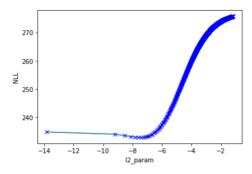
# multiplying by n and return
    return X.shape[0] * objective

# creating nll empty set, and l2 param range
NLL_res = []
l2_param_range = np.arange(0.000001, 0.3, 0.0001)

# for given l2 param range, calculate theta
for l2_param in l2_param_range:
    optimal_theta = fit_logistic_reg(X_train_normalized, y_train, f_objective, l2_param)
    NLL_res.append(neg_log_likelihood(optimal_theta, X_val_normalized, y_val, l2_param))

# plotting the answers
print('Best l2 param:{}, NLL: {}'.format(l2_param_range[np.argmin(NLL_res)], np.min(NLL_res)))
plt.plot(np.log(l2_param_range), NLL_res)
plt.plot(np.log(l2_param_range), NLL_res)
plt.plot(np.log(l2_param_range), NLL_res, 'bx')
plt.xlabel('12_param')
plt.ylabel('NLL')
plt.show()
```

Best l2 param:0.000501, NLL: 232.9126744529017



## **Q8**

[Optional]It seems reasonable to interpret the predictionf(x)= $\phi$ (wTx)=1/(1+e-wTx) as the probability that y = 1, for a randomly drawn pair (x, y). Since we only have a finite sample (and we are regularizing, which will bias things a bit) there is a question of how well "calibrated" our predicted probabilities are. Roughly speaking, we say f(x) is well calibrated if we look at all examples (x,y) for which f(x)  $\approx$  0.7 and we find that close to 70% of those examples have y = 1, as predicted... and then we repeat that for all predicted probabilities in (0, 1). To see how well-calibrated our predicted probabilities are, break the predictions on the validation set into groups based on the predicted probability (you can play with the size of the groups to get a result you think is informative). For each group, examine the percentage of positive labels. You can make a table or graph. Summarize the results. You may get some ideas and references from scikit-learn's discussion.

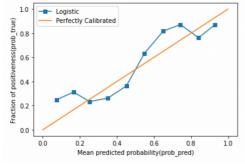
## Ans) LogisticRegression returns well calibrated predictions by default as it directly optimizes Log loss

```
# Best 12 param: 0.000501
from sklearn.calibration import calibration_curve

optimal_theta = fit_logistic_reg(X_train_normalized, y_train, f_objective, l2_param = 0.000501)
y_pred = 1/(1+np.exp(-(X_val_normalized @ optimal_theta.T)))

prob_true, prob_pred = calibration_curve(y_val, y_pred, normalize=False, n_bins=10)

plt.plot(prob_pred, prob_true, marker = 's', label = 'Logistic')
plt.plot([0, 1], [0, 1], label = 'Perfectly Calibrated')
plt.xlabel('Mean predicted probability(prob_pred)')
plt.ylabel('Fraction of positiveness(prob_true)')
plt.legend()
plt.show()
```



## The percentage of positive labels for each group

```
# The percentage of positive labels for each group

for idx, value in enumerate(prob_true):
    print('Group {}) Percentage of positive labels: {:.2f}'.format(idx+1, value))

Group 1) Percentage of positive labels: 0.25

Group 2) Percentage of positive labels: 0.31

Group 3) Percentage of positive labels: 0.23

Group 4) Percentage of positive labels: 0.26

Group 5) Percentage of positive labels: 0.36

Group 6) Percentage of positive labels: 0.63

Group 7) Percentage of positive labels: 0.82

Group 8) Percentage of positive labels: 0.87

Group 9) Percentage of positive labels: 0.87

Group 10) Percentage of positive labels: 0.87
```