[COM6513] Assignment 2: Topic Classification with a Feedforward Network

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The goal of this assignment is to develop a Feedforward neural network for topic classification.

For that purpose, you will implement:

- Text processing methods for transforming raw text data into input vectors for your network (1 mark)
- · A Feedforward network consisting of:
 - One-hot input layer mapping words into an Embedding weight matrix (1 mark)
 - One hidden layer computing the mean embedding vector of all words in input followed by a ReLU activation function (1 mark)
 - Output layer with a softmax activation. (1 mark)
- The Stochastic Gradient Descent (SGD) algorithm with back-propagation to learn the weights of your Neural network. Your algorithm should:
 - Use (and minimise) the Categorical Cross-entropy loss function (1 mark)
 - Perform a Forward pass to compute intermediate outputs (3 marks)
 - Perform a Backward pass to compute gradients and update all sets of weights (6 marks)
 - Implement and use Dropout after each hidden layer for regularisation (2 marks)
- Discuss how did you choose hyperparameters? You can tune the learning rate (hint: choose small values), embedding size {e.g. 50, 300, 500}, the dropout rate {e.g. 0.2, 0.5} and the learning rate. Please use tables or graphs to show training and validation performance for each hyperparameter combination (2 marks).
- After training a model, plot the learning process (i.e. training and validation loss in each epoch) using a line plot and report accuracy. Does your model overfit, underfit or is about right? (1 mark).
- Re-train your network by using pre-trained embeddings (<u>GloVe (https://nlp.stanford.edu/projects/glove/)</u>) trained on large corpora. Instead of randomly initialising the embedding weights matrix, you should initialise it with the pre-trained weights. During training, you should not update them (i.e. weight freezing) and backprop should stop before computing gradients for updating embedding weights. Report results by performing hyperparameter tuning and plotting the learning process. Do you get better performance? (3 marks).
- Extend you Feedforward network by adding more hidden layers (e.g. one more or two). How does it affect the performance? Note: You need to repeat hyperparameter tuning, but the number of combinations grows exponentially. Therefore, you need to choose a subset of all possible combinations (4 marks)
- Provide well documented and commented code describing all of your choices. In general, you are free to
 make decisions about text processing (e.g. punctuation, numbers, vocabulary size) and hyperparameter
 values. We expect to see justifications and discussion for all of your choices (2 marks).
- Provide efficient solutions by using Numpy arrays when possible. Executing the whole notebook with your code should not take more than 10 minutes on any standard computer (e.g. Intel Core i5 CPU, 8 or 16GB RAM) excluding hyperparameter tuning runs and loading the pretrained vectors. You can find tips in Lab 1 (2 marks).

Data

The data you will use for the task is a subset of the <u>AG News Corpus</u> (http://groups.di.unipi.it/~gulli/AG_corpus_of_news_articles.html) and you can find it in the ./data_topic folder in CSV format:

- data topic/train.csv: contains 2,400 news articles, 800 for each class to be used for training.
- data_topic/dev.csv: contains 150 news articles, 50 for each class to be used for hyperparameter selection and monitoring the training process.
- data_topic/test.csv: contains 900 news articles, 300 for each class to be used for testing.

Pre-trained Embeddings

You can download pre-trained GloVe embeddings trained on Common Crawl (840B tokens, 2.2M vocab, cased, 300d vectors, 2.03 GB download) from here (http://nlp.stanford.edu/data/glove.840B.300d.zip). No need to unzip, the file is large.

Save Memory

To save RAM, when you finish each experiment you can delete the weights of your network using del W followed by Python's garbage collector gc.collect()

Submission Instructions

You should submit a Jupyter Notebook file (assignment2.ipynb) and an exported PDF version (you can do it from Jupyter: File->Download as->PDF via Latex).

You are advised to follow the code structure given in this notebook by completing all given funtions. You can also write any auxilliary/helper functions (and arguments for the functions) that you might need but note that you can provide a full solution without any such functions. Similarly, you can just use only the packages imported below but you are free to use any functionality from the Python Standard Library

(https://docs.python.org/3/library/index.html), NumPy, SciPy (excluding built-in softmax funtcions) and Pandas. You are **not allowed to use any third-party library** such as Scikit-learn (apart from metric functions already provided), NLTK, Spacy, Keras, Pytorch etc.. You should mention if you've used Windows to write and test your code because we mostly use Unix based machines for marking (e.g. Ubuntu, MacOS).

There is no single correct answer on what your accuracy should be, but correct implementations usually achieve F1-scores around 80% or higher. The quality of the analysis of the results is as important as the accuracy itself.

This assignment will be marked out of 30. It is worth 30% of your final grade in the module.

The deadline for this assignment is **23:59 on Mon, 9 May 2022** and it needs to be submitted via Blackboard. Standard departmental penalties for lateness will be applied. We use a range of strategies to **detect unfair** means (https://www.sheffield.ac.uk/ssid/unfair-means/index), including Turnitin which helps detect plagiarism. Use of unfair means would result in getting a failing grade.

In [1]:

```
import pandas as pd
import numpy as np
from collections import Counter
import re
import matplotlib.pyplot as plt
from sklearn.metrics import accuracy_score, precision_score, recall_score, f1_score
import random
from time import localtime, strftime
from scipy.stats import spearmanr, pearsonr
import zipfile
import gc

# fixing random seed for reproducibility
random.seed(16269)
np.random.seed(16269)
```

Transform Raw texts into training and development data

First, you need to load the training, development and test sets from their corresponding CSV files (tip: you can use Pandas dataframes).

In [2]:

```
#import data
df_train = pd.read_csv('./data_topic/train.csv',names = ['label','text'])
df_dev = pd.read_csv('./data_topic/dev.csv',names = ['label','text'])
df_test = pd.read_csv('./data_topic/test.csv',names = ['label','text'])
```

In [3]:

```
df_train
```

Out[3]:

label	text
0 1 Reuters - Venezuelans turned out	early\and in
1 Reuters - South Korean police used	water canno
2 1 Reuters - Thousands of Palestinia	ın\prisoners i
3 1 AFP - Sporadic gunfire and shelling	ıg took place
4 1 AP - Dozens of Rwandan soldiers fle	ew into Suda
395 3 Australia #39;s dominant airline,	Qantas, has
396 3 Reuters - Medtronic Inc. (MDT.N) on	Wednesday\
397 3 SAN FRANCISCO (Reuters) - Google	e Inc. &ItA H
398 3 BHP Billiton, the world #39;s bigg	jest mining c
399 3 Europe's Frustration Grows as Dol	llar Hits Anot

2400 rows × 2 columns

In [4]:

```
#split whole data into content and label (in the required data type)
train_y = df_train.iloc[:,0]
train_x = df_train.iloc[:,1]
train_x = train_x.tolist()
train_y = train_y.values
test_y = df_test.iloc[:,0]
test_x = df_test.iloc[:,1]
test_x = test_x.tolist()
test_y = test_y.values
dev_y = df_dev.iloc[:,0]
dev_x = df_dev.iloc[:,1]
dev_x = dev_x.tolist()
dev_y = dev_y.values
```

In [5]:

```
#change from [1,2,3] to [0,1,2] for fitting the classification model
train_y -= 1
dev_y -= 1
test_y -= 1
```

```
In [6]:
```

```
dev y
```

Out[6]:

Create input representations

To train your Feedforward network, you first need to obtain input representations given a vocabulary. One-hot encoding requires large memory capacity. Therefore, we will instead represent documents as lists of vocabulary indices (each word corresponds to a vocabulary index).

Text Pre-Processing Pipeline

To obtain a vocabulary of words. You should:

- tokenise all texts into a list of unigrams (tip: you can re-use the functions from Assignment 1)
- remove stop words (using the one provided or one of your preference)
- · remove unigrams appearing in less than K documents
- use the remaining to create a vocabulary of the top-N most frequent unigrams in the entire corpus.

In [7]:

```
#use stop words from text processing lecture last semester
stop_words = ['a', 'able', 'about', 'above', 'according', 'accordingly', 'across','ron','co
'also', 'although', 'always', 'am', 'among', 'amongst', 'an', 'and', 'another', 'any', 'any
'have', 'having', 'he', 'hello', 'help', 'hence', 'her', 'here', 'hereafter', 'hereby', 'he
'if', 'ignored', 'immediate', 'in', 'inasmuch', 'inc', 'indeed', 'indicate', 'indicated', '
'may', 'maybe', 'me', 'mean', 'meanwhile', 'merely', 'might', 'more', 'moreover', 'most', 'm
'near', 'nearly', 'necessary', 'need', 'needs', 'neither', 'never', 'nevertheless', 'new',
'old', 'on', 'once', 'one', 'ones', 'only', 'onto', 'or', 'other', 'others', 'otherwise', '
'says', 'second', 'secondly', 'see', 'seeing', 'seem', 'seemed', 'seeming', 'seems', 'seen'
'useful', 'uses', 'using', 'usually', 'uucp', 'v', 'value', 'various', 've', 'very', 'via',
'what', 'whatever', 'when', 'whence', 'whenever', 'where', 'whereafter', 'whereas', 'whereb
```

Unigram extraction from a document

You first need to implement the extract_ngrams function. It takes as input:

- x_raw: a string corresponding to the raw text of a document
- ngram_range: a tuple of two integers denoting the type of ngrams you want to extract, e.g. (1,2) denotes
 extracting unigrams and bigrams.
- token_pattern: a string to be used within a regular expression to extract all tokens. Note that data is already tokenised so you could opt for a simple white space tokenisation.

- stop_words : a list of stop words
- vocab: a given vocabulary. It should be used to extract specific features.

and returns:

a list of all extracted features.

In [8]:

```
def extract ngrams(x raw, ngram range=(1,3), token pattern=r'\b[A-Za-z][A-Za-z]+\b', stop w
    tokenRE = re.compile(token pattern)
    # first extract all unigrams by tokenising
    x_uni = [w for w in tokenRE.findall(str(x_raw).lower(),) if w not in stop_words]
    # this is to store the ngrams to be returned
    x = []
    if ngram_range[0]==1:
        x = x_uni
    # generate n-grams from the available unigrams x uni
    ngrams = []
    for n in range(ngram_range[0], ngram_range[1]+1):
        # ignore unigrams
        if n==1: continue
        # pass a list of lists as an argument for zip
        arg_list = [x_uni]+[x_uni[i:] for i in range(1, n)]
        # extract tuples of n-grams using zip
        # for bigram this should look: list(zip(x uni, x uni[1:]))
        # align each item x[i] in x_uni with the next one x[i+1].
        # Note that x uni and x uni[1:] have different lengths
        # but zip ignores redundant elements at the end of the second list
        # Alternatively, this could be done with for loops
        x_ngram = list(zip(*arg_list))
        ngrams.append(x ngram)
    for n in ngrams:
        for t in n:
            x.append(t)
    if len(vocab)>0:
        x = [w \text{ for } w \text{ in } x \text{ if } w \text{ in } vocab]
    return x
```

Create a vocabulary of n-grams

Then the get_vocab function will be used to (1) create a vocabulary of ngrams; (2) count the document frequencies of ngrams; (3) their raw frequency. It takes as input:

X_raw: a list of strings each corresponding to the raw text of a document

- ngram_range : a tuple of two integers denoting the type of ngrams you want to extract, e.g. (1,2) denotes extracting unigrams and bigrams.
- token_pattern: a string to be used within a regular expression to extract all tokens. Note that data is already tokenised so you could opt for a simple white space tokenisation.
- stop_words : a list of stop words
- min_df: keep ngrams with a minimum document frequency.
- keep topN: keep top-N more frequent ngrams.

and returns:

- vocab: a set of the n-grams that will be used as features.
- df: a Counter (or dict) that contains ngrams as keys and their corresponding document frequency as values.
- ngram_counts : counts of each ngram in vocab

In [9]:

```
def get_vocab(X_raw, ngram_range=(1,3), token_pattern=r'\b[A-Za-z][A-Za-z]+\b',
              min df=0, keep_topN=0, stop_words=[]):
    tokenRE = re.compile(token_pattern)
    df = Counter()
    ngram_counts = Counter()
    vocab = set()
    # iterate through each raw text
    for x in X_raw:
        x_ngram = extract_ngrams(x, ngram_range=ngram_range, token_pattern=token_pattern, s
        #update doc and ngram frequencies
        df.update(list(set(x_ngram)))
        ngram counts.update(x ngram)
    # obtain a vocabulary as a set.
    # Keep elements with doc frequency > minimum doc freq (min_df)
    # Note that df contains all te
    vocab = set([w for w in df if df[w]>=min_df])
    # keep the top N most frequent
    if keep_topN>0:
        vocab = set([w[0] \text{ for } w \text{ in ngram counts.most common(keep topN) if } w[0] \text{ in vocab]})
    return vocab, df, ngram counts
```

Now you should use <code>get_vocab</code> to create your vocabulary and get document and raw frequencies of unigrams:

```
In [10]:
```

```
#get vocab
train_voc_set,train_df,train_ngcount = get_vocab(train_x,ngram_range = (1,1),stop_words=sto
dev_voc_set,dev_df,dev_ngcount = get_vocab(dev_x,ngram_range = (1,1),stop_words=stop_words)
test_voc_set,test_df,test_ngcount = get_vocab(test_x,ngram_range = (1,1),stop_words=stop_words)
```

In [11]:

```
#train_voc_set
```

In [12]:

```
#check the data size
print(len(train_voc_set),len(train_df),len(train_ngcount))
print(len(dev_voc_set),len(dev_df),len(dev_ngcount))
print(len(test_voc_set),len(test_df),len(test_ngcount))
```

```
8596 8596 8596
1629 1629 1629
5672 5672 5672
```

Then, you need to create vocabulary id -> word and word -> vocabulary id dictionaries for reference:

In [13]:

```
#make two dict for id word and word id
voc_id2word = enumerate(train_voc_set)
voc_id2word = dict(voc_id2word)
word2voc_id = {value:key for key,value in voc_id2word.items()}
```

In [14]:

```
word2voc_id
Out[14]:
{'living': 0,
 'inspiration': 1,
 'plotters': 2,
 'grow': 3,
 'brings': 4,
 'engages': 5,
 'quarter': 6,
 'imminent': 7,
 'nationals': 8,
 'honor': 9,
 'air': 10,
 'possibly': 11,
 'dimitris': 12,
 'filings': 13,
 'expectation': 14,
 'lost': 15,
 'breezy': 16,
 'nrimarv': 17.
```

Convert the list of unigrams into a list of vocabulary indices

Storing actual one-hot vectors into memory for all words in the entire data set is prohibitive. Instead, we will store word indices in the vocabulary and look-up the weight matrix. This is equivalent of doing a dot product between an one-hot vector and the weight matrix.

First, represent documents in train, dev and test sets as lists of words in the vocabulary:

In [15]:

```
#get the unigram data
uni_train_x = [extract_ngrams(train_x[i],ngram_range=(1,1), stop_words= stop_words) for i i
uni_dev_x = [extract_ngrams(dev_x[i],ngram_range=(1,1), stop_words= stop_words) for i in ra
uni_test_x = [extract_ngrams(test_x[i],ngram_range=(1,1), stop_words= stop_words) for i in
```

In [16]:

```
#uni_train_x[0]
```

Then convert them into lists of indices in the vocabulary:

In [17]:

```
#covert data into indices
train indices x = list(range(len(uni train x)))
dev_indices_x = list(range(len(uni_dev_x)))
test_indices_x = list(range(len(uni_test_x)))
for i in range(len(uni_train_x)):
   temp1 = []
   for j in uni_train_x[i]:
        if j in word2voc_id.keys():
            temp1.append(word2voc_id[j])
   train_indices_x[i] = temp1
for i in range(len(uni_dev_x)):
   temp2 = []
   for j in uni dev x[i]:
        if j in word2voc_id.keys():
            temp2.append(word2voc_id[j])
   dev_indices_x[i] = temp2
for i in range(len(uni test x)):
   temp3 = []
   for j in uni_test_x[i]:
        if j in word2voc id.keys():
            temp3.append(word2voc_id[j])
   test indices x[i] = temp3
```

```
In [18]:
```

```
train_indices_x[0]
Out[18]:
[7949,
 176,
 7922,
 4247,
 3639,
 2828,
 6569,
 1600,
 423,
 4722,
 5400,
 6736,
 6270,
 4796,
 8363.
 5832,
 1013,
 429,
 3586,
 1276,
 6170]
```

Put the labels Y for train, dev and test sets into arrays:

```
In [19]:
```

Network Architecture

Your network should pass each word index into its corresponding embedding by looking-up on the embedding matrix and then compute the first hidden layer \mathbf{h}_1 :

$$\mathbf{h}_1 = \frac{1}{|x|} \sum_i W_i^e, i \in X$$

where |x| is the number of words in the document and W^e is an embedding matrix $|V| \times d$, |V| is the size of the vocabulary and d the embedding size.

Then \mathbf{h}_1 should be passed through a ReLU activation function:

$$\mathbf{a}_1 = relu(\mathbf{h}_1)$$

Finally the hidden layer is passed to the output layer:

$$\mathbf{y} = \operatorname{softmax}(\mathbf{a}_1 W)$$

where W is a matrix $d \times |\mathcal{Y}|$, $|\mathcal{Y}|$ is the number of classes.

During training, a_1 should be multiplied with a dropout mask vector (elementwise) for regularisation before it is passed to the output layer.

You can extend to a deeper architecture by passing a hidden layer to another one:

$$\mathbf{h_i} = \mathbf{a}_{i-1} W_i$$

$$\mathbf{a_i} = relu(\mathbf{h_i})$$

Network Training

First we need to define the parameters of our network by initiliasing the weight matrices. For that purpose, you should implement the network_weights function that takes as input:

- vocab_size : the size of the vocabulary
- embedding_dim: the size of the word embeddings
- hidden_dim: a list of the sizes of any subsequent hidden layers. Empty if there are no hidden layers between the average embedding and the output layer
- num_classes: the number of the classes for the output layer

and returns:

• W: a dictionary mapping from layer index (e.g. 0 for the embedding matrix) to the corresponding weight matrix initialised with small random numbers (hint: use numpy.random.uniform with from -0.1 to 0.1)

Make sure that the dimensionality of each weight matrix is compatible with the previous and next weight matrix, otherwise you won't be able to perform forward and backward passes. Consider also using np.float32 precision to save memory.

In [20]:

In [21]:

```
W = network_weights(vocab_size=5,embedding_dim=10,hidden_dim=[10,2], num_classes=2)
print('W_emb:', W[0].shape)
print('W_out:', W[1].shape)

W_emb: (5, 10)
W_out: (10, 10)
```

In [22]:

```
#check the network_weights work or not
W = network_weights(vocab_size=3,embedding_dim=4,hidden_dim=[3,4], num_classes=2)
W
```

Out[22]:

```
{0: array([[ 0.12224471, -0.0794624 , -0.23009355, 0.29406023],
       [-0.46997094, 0.12204431, -0.0749153, 0.37384233],
       [-0.3974818, 0.3035981, 0.46178967, -0.43050396]],
      dtype=float32),
1: array([[-0.31256202, -0.42996365, -0.2553189]),
       [-0.28818667, -0.15275556, -0.22158206],
       [-0.20424126, 0.4211786, -0.4858057],
       [ 0.45516163, 0.22709407, -0.19921613]], dtype=float32),
2: array([[-0.02515792, 0.37151852, 0.32920408, 0.3873491],
       [0.23128773, -0.08528457, 0.22343004, 0.44983786],
       [-0.23429091, 0.00418209, 0.3896187, 0.01652054]],
      dtype=float32),
3: array([[-0.40125608, -0.16001858],
       [ 0.30069476, 0.21305238],
       [-0.4713758, -0.10194607],
       [ 0.02704137, -0.39511573]], dtype=float32)}
```

Then you need to develop a softmax function (same as in Assignment 1) to be used in the output layer.

It takes as input z (array of real numbers) and returns sig (the softmax of z)

In [23]:

```
def softmax(z):
    soft = np.exp(z- np.max(z))
    soft = soft / soft.sum()
    return soft
```

Now you need to implement the categorical cross entropy loss by slightly modifying the function from Assignment 1 to depend only on the true label y and the class probabilities vector y_preds:

In [24]:

```
def categorical_loss(y, y_preds):
    #print(y_preds)
    y_max = max(y_preds)
    l = (-y) * np.log(y_max)
    return l
```

Then, implement the relu function to introduce non-linearity after each hidden layer of your network (during the forward pass):

$$relu(z_i) = max(z_i, 0)$$

and the relu_derivative function to compute its derivative (used in the backward pass):

```
relu derivative(z_i)=0, if z_i<=0, 1 otherwise.
```

Note that both functions take as input a vector z

Hint use .copy() to avoid in place changes in array z

In [25]:

```
def relu(z):
    a = np.maximum(z,0)
    return a

def relu_derivative(z):
    z_cp = z.copy()
    z_cp[z_cp<=0] =0
    z_cp[z_cp>0] =1
    return z_cp
```

During training you should also apply a dropout mask element-wise after the activation function (i.e. vector of ones with a random percentage set to zero). The dropout_mask function takes as input:

- size: the size of the vector that we want to apply dropout
- dropout_rate: the percentage of elements that will be randomly set to zeros

and returns:

dropout vec: a vector with binary values (0 or 1)

In [26]:

```
def dropout_mask(size, dropout_rate):
    np.random.seed(16269)
    num = size*dropout_rate
    dropout_vec = np.ones(size)
    dropout_ind = np.random.choice(np.arange(size),replace = False,size = int(num))
    dropout_vec[dropout_ind] = 0
    return dropout_vec
```

In [27]:

```
print(dropout_mask(10, 0.2))
print(dropout_mask(10, 0.2))
```

```
[0. 1. 1. 1. 1. 1. 1. 0. 1. 1.]
[0. 1. 1. 1. 1. 1. 1. 0. 1. 1.]
```

Now you need to implement the <code>forward_pass</code> function that passes the input x through the network up to the output layer for computing the probability for each class using the weight matrices in <code>W</code> . The ReLU activation function should be applied on each hidden layer.

- x : a list of vocabulary indices each corresponding to a word in the document (input)
- W: a list of weight matrices connecting each part of the network, e.g. for a network with a hidden and an output layer: W[0] is the weight matrix that connects the input to the first hidden layer, W[1] is the weight matrix that connects the hidden layer to the output layer.
- dropout_rate : the dropout rate that is used to generate a random dropout mask vector applied after each hidden layer for regularisation.

and returns:

out_vals: a dictionary of output values from each layer: h (the vector before the activation function), a
 (the resulting vector after passing h from the activation function), its dropout mask vector; and the
 prediction vector (probability for each class) from the output layer.

In [28]:

```
def forward_pass(x, W, dropout_rate=0.2):
    out_vals = {}
    h_{vecs} = []
    a vecs = []
    dropout_vecs = []
    #count the weight of first layer, do relu and dropout function
    w_{emb} = [W[0][index] for index in x]
    h = np.mean(np.array(w_emb),axis=0)
    a = relu(h)
    drop vec = dropout mask(len(a), dropout rate)
    output = a*drop_vec
    h vecs.append(h)
    a_vecs.append(a)
    dropout_vecs.append(drop_vec)
    #other Layers
    for i in range(1,len(W)-1):
        h = np.dot(output,W[i])
        a = relu(h)
        drop_vec = dropout_mask(len(a), dropout_rate)
        output = a*drop_vec
        h vecs.append(h)
        a vecs.append(a)
        dropout vecs.append(drop vec)
    y pre = softmax(np.dot(output,W[len(W)-1]))
    out_vals['h'] = h_vecs
    out vals['a'] = a vecs
    out_vals['dropout_vecs'] = dropout_vecs
    out_vals['y'] = y_pre
    return out vals
```

In [29]:

```
#check is forward_pass function working or not
W = network_weights(vocab_size=100,embedding_dim=20,hidden_dim=[4,2], num_classes=2)

for i in range(len(W)):
    print('Shape W'+str(i), W[i].shape)

print(forward_pass([5,0], W, dropout_rate=0.5))
```

```
Shape W0 (100, 20)
Shape W1 (20, 4)
Shape W2 (4, 2)
Shape W3 (2, 2)
{'h': [array([ 0.17688254, -0.00454105, 0.05746456, -0.00638478, -0.1302145
3,
       -0.04556508, -0.02224372, 0.05884917, -0.2432979, -0.04557189,
        0.06403284, -0.29234985, -0.3149953 , -0.27194694, -0.1704622 ,
        0.06597526, -0.08525288, -0.20908287, -0.18800098, 0.15818825],
      dtype=float32), array([ 0.00223933,  0.03659521, -0.02797977, -0.03967
538]), array([0.00765979, 0.00364189])], 'a': [array([0.17688254, 0.
, 0.05746456, 0.
                        , 0.
                 , 0.
                             , 0.05884917, 0.
       0.
                                         , 0.
       0.06403284, 0.
                             , 0.
                                                     , 0.
                                         , 0.
       0.06597526, 0.
                             , 0.
                                                     , 0.15818825],
      dtype=float32), array([0.00223933, 0.03659521, 0.
]), array([0.00765979, 0.00364189])], 'dropout_vecs': [array([1., 0., 0.,
0., 1., 0., 0., 0., 0., 1., 1., 1., 1., 0., 1., 1.,
       1., 0., 1.]), array([0., 1., 1., 0.]), array([0., 1.])], 'y': array
([0.50009148, 0.49990852])
```

The backward_pass function computes the gradients and updates the weights for each matrix in the network from the output to the input. It takes as input

- x: a list of vocabulary indices each corresponding to a word in the document (input)
- y: the true label
- W: a list of weight matrices connecting each part of the network, e.g. for a network with a hidden and an output layer: W[0] is the weight matrix that connects the input to the first hidden layer, W[1] is the weight matrix that connects the hidden layer to the output layer.
- out vals: a dictionary of output values from a forward pass.
- learning rate: the learning rate for updating the weights.
- freeze emb: boolean value indicating whether the embedding weights will be updated.

and returns:

W: the updated weights of the network.

Hint: the gradients on the output layer are similar to the multiclass logistic regression.

In [30]:

```
def backward_pass(x, y, W, out_vals, lr=0.001, freeze_emb=True):
   #calculate gradient
   grad = out_vals['y'] - (np.arange(len(out_vals['y'])) == y)
   # calculate weights
   out_layer_input = out_vals['a'][-1] * out_vals['dropout_vecs'][-1]
   grad_on_wt = np.outer(grad, out_layer_input)
   grad_on_wt = grad_on_wt.T
   grad = grad.dot(W[len(W)-1].T)
   # update weight
   W[len(W)-1] -= lr * grad_on_wt
   # Update each hidden Layer
   for i in range(len(W) - 2, 1, -1):
        grad *= relu_derivative(out_vals['h'][i])
        # calculate weights
        layer_input = out_vals['a'][i - 1] * out_vals['dropout_vecs'][i - 1]
        grad_on_wt = np.outer(grad, layer_input)
        grad_on_wt = grad_on_wt.T
        grad = grad.dot(W[i].T)
        # update weight
       W[i] -= lr * grad_on_wt
   # Update weights of the initial layer
   if not freeze emb:
        grad *= relu_derivative(out_vals['h'][0])
        W[0][x] -= lr * grad
   return W
```

Finally you need to modify SGD to support back-propagation by using the forward_pass and backward_pass functions.

The SGD function takes as input:

- X_tr: array of training data (vectors)
- Y_tr: labels of X_tr
- W: the weights of the network (dictionary)
- X dev: array of development (i.e. validation) data (vectors)
- Y dev: labels of X dev
- 1r: learning rate
- · dropout: regularisation strength
- · epochs: number of full passes over the training data
- tolerance : stop training if the difference between the current and previous validation loss is smaller than a threshold
- freeze_emb: boolean value indicating whether the embedding weights will be updated (to be used by the backward pass function).

• print progress : flag for printing the training progress (train/validation loss)

and returns:

- · weights: the weights learned
- training_loss_history: an array with the average losses of the whole training set after each epoch
- validation_loss_history: an array with the average losses of the whole development set after each epoch

In [31]:

```
import time
def SGD(X_tr, Y_tr, W, X_dev=[], Y_dev=[], lr=0.001,
        dropout=0.2, epochs=5, tolerance=0.001, freeze_emb=False,
        print_progress=True):
   #np.random.seed(16269)
   training_loss_history = []
   validation_loss_history = []
   train_docs = list(zip(X_tr, Y_tr))
   dev_docs = list(zip(X_dev,Y_dev))
   for i in range(0,epochs):
        #Randomise the order of each epoch
        np.random.seed(i)
        new_X_tr = np.random.permutation(X_tr)
        np.random.seed(i)
        new_Y_tr = np.random.permutation(Y_tr)
        new_train_docs = list(zip(new_X_tr,new_Y_tr))
        for x_i, y_i in new_train_docs:
            #fo = time.time()
            out_vals = forward_pass(x_i, W, dropout)
            #fo end = time.time()
            #ba = time.time()
            W = backward_pass(x_i, y_i, W, out_vals, lr, freeze_emb)
            #ba end = time.time()
            #print(ba_end - ba, fo_end - fo)
        #train loss
        temp_loss_tr = [categorical_loss(y_i, forward_pass(x_i, W, dropout)['y']) for x_i,
        cur_loss_tr = np.mean(temp_loss_tr)
        # Validation loss
        temp_loss_dev = [categorical_loss(y_i, forward_pass(x_i, W, dropout)['y']) for x_i,
        cur loss dev = np.mean(temp loss dev)
        training_loss_history.append(cur_loss_tr)
        validation_loss_history.append(cur_loss_dev)
        if print progress:
            print(f'Epoch: {i} | Train loss: {cur_loss_tr} | Dev loss: {cur_loss_dev}')
   return W, training_loss_history, validation_loss_history
```

In [32]:

<__array_function__ internals>:5: VisibleDeprecationWarning: Creating an n
darray from ragged nested sequences (which is a list-or-tuple of lists-ortuples-or ndarrays with different lengths or shapes) is deprecated. If you
meant to do this, you must specify 'dtype=object' when creating the ndarra
y.

Now you are ready to train and evaluate your neural net. First, you need to define your network using the network weights function followed by SGD with backprop:

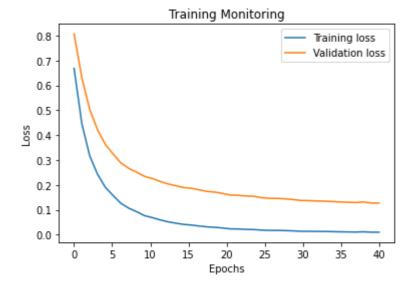
Plot the learning process:

In [33]:

```
x = np.linspace(0,len(loss_tr),len(loss_tr))
y1, y2 = loss_tr, dev_loss

plt.plot(x, y1,label='Training loss')
plt.plot(x, y2, label='Validation loss')

plt.title('Training Monitoring')
plt.xlabel('Epochs')
plt.ylabel('Loss')
plt.legend()
plt.show()
```



According to the graph, it is neither overfitting nor underfitting. It could train more times to get the higher accuracy probably.

Compute accuracy, precision, recall and F1-Score:

In [34]:

Accuracy: 0.8611111111111112
Precision: 0.8627965786817375
Recall: 0.86111111111111
F1-Score: 0.8607497524117416

Discuss how did you choose model hyperparameters?

In [35]:

```
# Firstly We should establish the list to select the parameters
from prettytable import PrettyTable
dim_choice=[100,200,300]
lr_rate = [0.1,0.01,0.05]
dropout_rate = [0.2, 0.4, 0.5]
hyper_result = 0.0
test_result = 0.0
result = list()
dim_his = []
lr his = []
drop his =[]
for dim in range(len(dim choice)):
    for lr in range(len(lr_rate)):
        for drop in range(len(dropout_rate)):
            W = network_weights(vocab_size=len(train_voc_set),embedding_dim=dim_choice[dim]
                                num_classes=3, init_val = 0.1)
            W, loss_tr, dev_loss = SGD(train_indices_x, Y_tr=train_y, W=W, X_dev=dev_indice
                            dropout=dropout_rate[drop],
                            freeze_emb=False,
                            tolerance=0.0001,
                            print_progress = False,
                            epochs=40)
            preds_dev = [np.argmax(forward_pass(x, W, dropout_rate=0.0)['y']) for x,y in zi
            val_acc = accuracy_score(dev_y,preds_dev)
            hyper_result = val_acc
            dim his.append(dim choice[dim])
            lr his.append(lr rate[lr])
            drop_his.append(dropout_rate[drop])
            result.append(hyper_result)
            ind = result.index(max(result))
            #print(hyper_result,dim_choice[dim],lr_rate[lr],dropout_rate[drop])
print(max(result),dim his[ind],lr his[ind],drop his[ind])
(which is a list-or-tuple of lists-or-tuples-or ndarrays with different le
ngths or shapes) is deprecated. If you meant to do this, you must specify
'dtype=object' when creating the ndarray.
  return array(a, dtype, copy=False, order=order)
<__array_function__ internals>:5: VisibleDeprecationWarning: Creating an n
darray from ragged nested sequences (which is a list-or-tuple of lists-or-
tuples-or ndarrays with different lengths or shapes) is deprecated. If you
meant to do this, you must specify 'dtype=object' when creating the ndarra
у.
C:\Users\88690\anaconda3\lib\site-packages\numpy\core\ asarray.py:102: Vis
ibleDeprecationWarning: Creating an ndarray from ragged nested sequences
(which is a list-or-tuple of lists-or-tuples-or ndarrays with different le
ngths or shapes) is deprecated. If you meant to do this, you must specify
'dtype=object' when creating the ndarray.
  return array(a, dtype, copy=False, order=order)
<__array_function__ internals>:5: VisibleDeprecationWarning: Creating an n
darray from ragged nested sequences (which is a list-or-tuple of lists-or-
tuples-or ndarrays with different lengths or shapes) is deprecated. If you
meant to do this, you must specify 'dtype=object' when creating the ndarra
у.
```

In [36]:

```
#print the table of all combinations of parameters
average_Table = PrettyTable(["Accuracy", "embedding_dim", "lr_rate", "dropout_rate"])
for i in range(len(dim_his)):
    average_Table.add_row([result[i],dim_his[i],lr_his[i],drop_his[i]])
print(average_Table)
 0.000000000000000000000
                             200
                                           v. 1
                                                       v. 4
                                                       0.4
  0.86666666666666
                             200
                                          0.1
  0.86666666666666
                             200
                                          0.1
                                                       0.5
         0.9
                             200
                                          0.01
                                                       0.2
         0.9
                             200
                                          0.01
                                                       0.4
         0.9
                             200
                                          0.01
                                                       0.5
         0.88
                                                       0.2
                             200
                                          0.05
                             200
                                          0.05
                                                       0.4
  0.8733333333333333
  0.8733333333333333
                             200
                                          0.05
                                                       0.5
                                                       0.2
         0.88
                             300
                                          0.1
  0.8733333333333333
                             300
                                          0.1
                                                       0.4
  0.86666666666666
                             300
                                          0.1
                                                       0.5
  0.90666666666666
                             300
                                          0.01
                                                       0.2
         0.9
                             300
                                          0.01
                                                       0.4
         0.9
                                                       0.5
                             300
                                          0.01
  0.8733333333333333
                             300
                                          0.05
                                                       0.2
  0.8733333333333333
                             300
                                          0.05
                                                       0.4
  0.873333333333333
                             300
                                          0.05
                                                       0.5
```

In [37]:

C:\Users\88690\anaconda3\lib\site-packages\numpy\core_asarray.py:102: Vis ibleDeprecationWarning: Creating an ndarray from ragged nested sequences (which is a list-or-tuple of lists-or-tuples-or ndarrays with different le ngths or shapes) is deprecated. If you meant to do this, you must specify 'dtype=object' when creating the ndarray.

return array(a, dtype, copy=False, order=order)

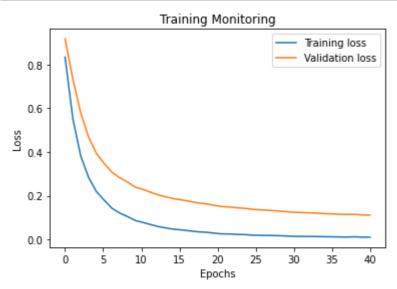
<__array_function__ internals>:5: VisibleDeprecationWarning: Creating an n
darray from ragged nested sequences (which is a list-or-tuple of lists-ortuples-or ndarrays with different lengths or shapes) is deprecated. If you
meant to do this, you must specify 'dtype=object' when creating the ndarra
y.

In [38]:

```
#plot the training loss and val loss with using best parameters
x = np.linspace(0,len(loss_tr),len(loss_tr))
y1, y2 = loss_tr, dev_loss

plt.plot(x, y1,label='Training loss')
plt.plot(x, y2, label='Validation loss')

plt.title('Training Monitoring')
plt.xlabel('Epochs')
plt.ylabel('Loss')
plt.legend()
plt.show()
```



This model is well, neither overfitting nor underfitting. Could increase epochs to train more times

In [39]:

Accuracy: 0.8588888888888889 Precision: 0.8619866910203213 Recall: 0.858888888888889 F1-Score: 0.8585911905628026

We could find out that accuracy is a little bit lower than the previous one which the parameters chosen randomly. The reason is that there are many same accuracy when we use validation to find the best parameters, and we choose randomly from one of the sets of the parameters.

Use Pre-trained Embeddings

Now re-train the network using GloVe pre-trained embeddings. You need to modify the backward_pass function above to stop computing gradients and updating weights of the embedding matrix.

Use the function below to obtain the embedding martix for your vocabulary. Generally, that should work without any problem. If you get errors, you can modify it.

In [40]:

```
def get_glove_embeddings(f_zip, f_txt, word2id, emb_size=300):
    w_emb = np.zeros((len(word2id), emb_size))
    with zipfile.ZipFile(f_zip) as z:
        with z.open(f_txt) as f:
        for line in f:
            line = line.decode('utf-8')
            word = line.split()[0]

        if word in train_voc_set:
            emb = np.array(line.strip('\n').split()[1:]).astype(np.float32)
            w_emb[word2id[word]] +=emb
    return w_emb
```

In [41]:

```
w_glove = get_glove_embeddings("glove.840B.300d.zip","glove.840B.300d.txt",word2voc_id)
```

First, initialise the weights of your network using the network_weights function. Second, replace the weights of the embedding matrix with w_glove . Finally, train the network by freezing the embedding weights:

In [42]:

```
Epoch: 0 | Train loss: 0.9046048299881092 | Dev loss: 0.8721254910692425
Epoch: 1 | Train loss: 0.8862225808382912 | Dev loss: 0.855000323019056
Epoch: 2 | Train loss: 0.8624540794587242 | Dev loss: 0.8358953980042821
Epoch: 3 | Train loss: 0.8272456805670378 | Dev loss: 0.8000549095154829
Epoch: 4 | Train loss: 0.7997430045066902 | Dev loss: 0.7785728351115514
Epoch: 5 | Train loss: 0.7692634053738563 | Dev loss: 0.7522427314756711
Epoch: 6 | Train loss: 0.7313739775435604 | Dev loss: 0.7142393658895781
Epoch: 7 | Train loss: 0.7004890602177151 | Dev loss: 0.6866705619824497
Epoch: 8 | Train loss: 0.6792048702656869 | Dev loss: 0.6665691649869395
Epoch: 9 | Train loss: 0.6528442550999817 | Dev loss: 0.6440841954963159
Epoch: 10 | Train loss: 0.6391529304793625 | Dev loss: 0.632417938097807
Epoch: 11 | Train loss: 0.6058533596235515 | Dev loss: 0.6006359834723779
Epoch: 12 | Train loss: 0.590246720274034 | Dev loss: 0.5873136111516857
Epoch: 13 | Train loss: 0.5755612689392734 | Dev loss: 0.573922717718161
Epoch: 14 | Train loss: 0.5562470321908112 | Dev loss: 0.5553715905281894
Epoch: 15 | Train loss: 0.5464131946607462 | Dev loss: 0.5463636181586466
```

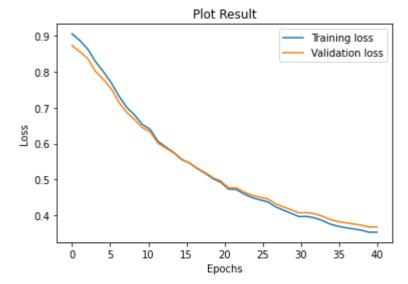
In [43]:

In [44]:

```
# Now we could plot result of the training
x = np.linspace(0,len(loss_tr),len(loss_tr))
y1, y2 = loss_tr, dev_loss

plt.plot(x, y1,label='Training loss')
plt.plot(x, y2, label='Validation loss')

plt.title('Plot Result')
plt.xlabel('Epochs')
plt.ylabel('Loss')
plt.legend()
plt.show()
```



Two loss lines are quire close, it a little bit overfitting. Guessing the reason is we choose parameters manually. Below, we will use validation set to find out the best set of parameters.

Discuss how did you choose model hyperparameters?

In [45]:

```
from prettytable import PrettyTable
lr_rate = [0.01, 0.001, 0.003]
dropout_rate = [0.2, 0.4, 0.5]
hyper result = 0.0
test_result = 0.0
result = list()
lr_his = []
drop_his =[]
for lr in range(len(lr_rate)):
   for drop in range(len(dropout rate)):
        W = network_weights(vocab_size=len(train_voc_set),embedding_dim=300,hidden_dim=[],
                            num classes=3, init val = 0.1)
       W[0] = w_glove
       W, loss tr, dev_loss = SGD(train_indices_x, Y_tr=train_y, W=W, X_dev=dev_indices_x,
                        dropout=dropout rate[drop],
                        freeze emb=True,
                        tolerance=0.0001,
                        print_progress = False,
                        epochs=40)
        preds_dev = [np.argmax(forward_pass(x, W, dropout_rate=0.0)['y']) for x,y in zip(de
        val acc = accuracy score(dev y,preds dev)
        hyper_result = val_acc
        lr_his.append(lr_rate[lr])
        drop_his.append(dropout_rate[drop])
        result.append(hyper result)
        ind = result.index(max(result))
        #print(hyper_result,dim_choice[dim],lr_rate[lr],dropout_rate[drop])
print(max(result), lr his[ind], drop his[ind])
totebepreedetonmarning, eredetng an nadrray from ragged neseed sequences
(which is a list-or-tuple of lists-or-tuples-or ndarrays with different le
ngths or shapes) is deprecated. If you meant to do this, you must specify
'dtype=object' when creating the ndarray.
  return array(a, dtype, copy=False, order=order)
<__array_function__ internals>:5: VisibleDeprecationWarning: Creating an n
darray from ragged nested sequences (which is a list-or-tuple of lists-or-
tuples-or ndarrays with different lengths or shapes) is deprecated. If you
meant to do this, you must specify 'dtype=object' when creating the ndarra
у.
C:\Users\88690\anaconda3\lib\site-packages\numpy\core\ asarray.py:102: Vis
ibleDeprecationWarning: Creating an ndarray from ragged nested sequences
(which is a list-or-tuple of lists-or-tuples-or ndarrays with different le
ngths or shapes) is deprecated. If you meant to do this, you must specify
'dtype=object' when creating the ndarray.
  return array(a, dtype, copy=False, order=order)
<__array_function__ internals>:5: VisibleDeprecationWarning: Creating an n
darray from ragged nested sequences (which is a list-or-tuple of lists-or-
tuples-or ndarrays with different lengths or shapes) is deprecated. If you
meant to do this, you must specify 'dtype=object' when creating the ndarra
```

In [46]:

```
emb_Table = PrettyTable(["Accuracy", "lr_rate", "dropout_rate"])
for i in range(len(result)):
    emb_Table.add_row([result[i],lr_his[i],drop_his[i]])
print(emb_Table)
```

lr_rate	dropout_rate
0.01	0.2
0.01	0.4
0.01	0.5
0.001	0.2
0.001	0.4
0.001	0.5
0.003	0.2
0.003	0.4
0.003	0.5
	0.01 0.01 0.001 0.001 0.001 0.003 0.003

In [47]:

```
Shape W0 (8596, 300)
Shape W1 (300, 3)
```

C:\Users\88690\anaconda3\lib\site-packages\numpy\core_asarray.py:102: Vis ibleDeprecationWarning: Creating an ndarray from ragged nested sequences (which is a list-or-tuple of lists-or-tuples-or ndarrays with different le ngths or shapes) is deprecated. If you meant to do this, you must specify 'dtype=object' when creating the ndarray.

```
return array(a, dtype, copy=False, order=order)
```

<__array_function__ internals>:5: VisibleDeprecationWarning: Creating an n
darray from ragged nested sequences (which is a list-or-tuple of lists-ortuples-or ndarrays with different lengths or shapes) is deprecated. If you
meant to do this, you must specify 'dtype=object' when creating the ndarra
y.

In [48]:

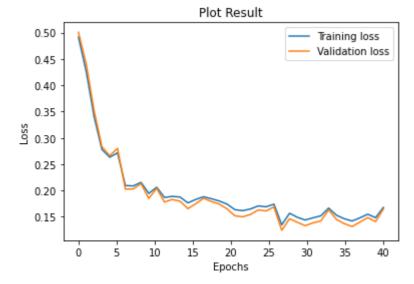
Accuracy: 0.886666666666667 Precision: 0.887098394913521 Recall: 0.886666666666667 F1-Score: 0.8868245713460235

In [49]:

```
# Now we could plot result of the training
x = np.linspace(0,len(loss_tr),len(loss_tr))
y1, y2 = loss_tr, dev_loss

plt.plot(x, y1,label='Training loss')
plt.plot(x, y2, label='Validation loss')

plt.title('Plot Result')
plt.xlabel('Epochs')
plt.ylabel('Loss')
plt.legend()
plt.show()
```



Extend to support deeper architectures

Extend the network to support back-propagation for more hidden layers. You need to modify the backward_pass function above to compute gradients and update the weights between intermediate hidden layers. Finally, train and evaluate a network with a deeper architecture. Do deeper architectures increase performance?

In [85]:

In [86]:

C:\Users\88690\anaconda3\lib\site-packages\numpy\core_asarray.py:102: Vis ibleDeprecationWarning: Creating an ndarray from ragged nested sequences (which is a list-or-tuple of lists-or-tuples-or ndarrays with different le ngths or shapes) is deprecated. If you meant to do this, you must specify 'dtype=object' when creating the ndarray.

return array(a, dtype, copy=False, order=order)

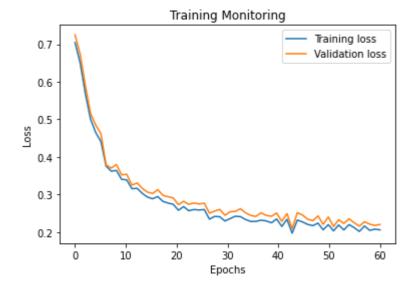
<__array_function__ internals>:5: VisibleDeprecationWarning: Creating an n
darray from ragged nested sequences (which is a list-or-tuple of lists-ortuples-or ndarrays with different lengths or shapes) is deprecated. If you
meant to do this, you must specify 'dtype=object' when creating the ndarra
y.

In [87]:

```
x = np.linspace(0,len(loss_tr),len(loss_tr))
y1, y2 = loss_tr, dev_loss

plt.plot(x, y1,label='Training loss')
plt.plot(x, y2, label='Validation loss')

plt.title('Training Monitoring')
plt.xlabel('Epochs')
plt.ylabel('Loss')
plt.legend()
plt.show()
```



In [88]:

```
preds_te = [np.argmax(forward_pass(x, W, dropout_rate=0.0)['y']) for x,y in zip(test_indice)
print('Accuracy:', accuracy_score(test_y,preds_te))
print('Precision:', precision_score(test_y,preds_te,average='macro'))
print('Recall:', recall_score(test_y,preds_te,average='macro'))
print('F1-Score:', f1_score(test_y,preds_te,average='macro'))
```

Accuracy: 0.77555555555556 Precision: 0.7851723233255484 Recall: 0.7755555555556 F1-Score: 0.7735548372159694 Using manual parameter, so the accuracy looks quite low

Discuss how did you choose model hyperparameters?

In [89]:

```
# Firstly We should establish the list to select the parameters
from prettytable import PrettyTable
hdim_choice=[[200],[100,50],[200,50]]
lr_rate = [0.01,0.001,0.005]
dropout_rate = [0.2,0.4,0.5]
hyper_result = 0.0
test_result = 0.0
result = list()
hdim_his = []
lr his = []
drop_his =[]
for hdim in range(len(hdim_choice)):
    for lr in range(len(lr_rate)):
        for drop in range(len(dropout_rate)):
            W = network_weights(vocab_size=len(train_voc_set),embedding_dim=300,hidden_dim=
            W[0] = w_glove
            W, loss_tr, dev_loss = SGD(train_indices_x, Y_tr=train_y, W=W, X_dev=dev_indice
                            dropout=dropout_rate[drop],
                            freeze_emb=True,
                            tolerance=0.0001,
                            print_progress = False,
                            epochs=50)
            preds_dev = [np.argmax(forward_pass(x, W, dropout_rate=0.0)['y']) for x,y in zi
            val_acc = accuracy_score(dev_y,preds_dev)
            hyper_result = val_acc
            hdim_his.append(hdim_choice[hdim])
            lr_his.append(lr_rate[lr])
            drop_his.append(dropout_rate[drop])
            result.append(hyper_result)
            ind = result.index(max(result))
            #print(hyper_result,dim_choice[dim],lr_rate[lr],dropout_rate[drop])
print(max(result),hdim_his[ind],lr_his[ind],drop_his[ind])
  recurri urruy (u, ucype, copy russe, oruci oruci,
<__array_function__ internals>:5: VisibleDeprecationWarning: Creating an n
darray from ragged nested sequences (which is a list-or-tuple of lists-or-
tuples-or ndarrays with different lengths or shapes) is deprecated. If you
meant to do this, you must specify 'dtype=object' when creating the ndarra
у.
C:\Users\88690\anaconda3\lib\site-packages\numpy\core\_asarray.py:102: Vis
ibleDeprecationWarning: Creating an ndarray from ragged nested sequences
(which is a list-or-tuple of lists-or-tuples-or ndarrays with different le
ngths or shapes) is deprecated. If you meant to do this, you must specify
'dtype=object' when creating the ndarray.
  return array(a, dtype, copy=False, order=order)
<__array_function__ internals>:5: VisibleDeprecationWarning: Creating an n
darray from ragged nested sequences (which is a list-or-tuple of lists-or-
tuples-or ndarrays with different lengths or shapes) is deprecated. If you
meant to do this, you must specify 'dtype=object' when creating the ndarra
C:\Users\88690\anaconda3\lib\site-packages\numpy\core\_asarray.py:102: Vis
ibleDeprecationWarning: Creating an ndarray from ragged nested sequences
(which is a list-or-tuple of lists-or-tuples-or ndarrays with different le
```

In [90]:

```
hidden_Table = PrettyTable(["Accuracy", "hdim", "lr_rate", "dropout_rate"])
for i in range(len(hdim_his)):
    hidden_Table.add_row([result[i],hdim_his[i],lr_his[i],drop_his[i]])
print(hidden_Table)
```

+ Accuracy	+ hdim	 lr_rate	dropout_rate
+	+	+	+
0.7	[200]	0.01	0.2
0.746666666666666	[200]	0.01	0.4
0.666666666666666	[200]	0.01	0.5
0.78	[200]	0.001	0.2
0.7733333333333333	[200]	0.001	0.4
0.72	[200]	0.001	0.5
0.706666666666666	[200]	0.005	0.2
0.746666666666666	[200]	0.005	0.4
0.706666666666666	[200]	0.005	0.5
0.60666666666666	[100, 50]	0.01	0.2
0.533333333333333	[100, 50]	0.01	0.4
0.3866666666666666	[100, 50]	0.01	0.5
0.71333333333333334	[100, 50]	0.001	0.2
0.646666666666666	[100, 50]	0.001	0.4
0.4933333333333333	[100, 50]	0.001	0.5
0.6933333333333334	[100, 50]	0.005	0.2
0.62	[100, 50]	0.005	0.4
0.42	[100, 50]	0.005	0.5
0.40666666666666	[200, 50]	0.01	0.2
0.393333333333333	[200, 50]	0.01	0.4
0.3533333333333333	[200, 50]	0.01	0.5
0.8	[200, 50]	0.001	0.2
0.66	[200, 50]	0.001	0.4
0.546666666666666	[200, 50]	0.001	0.5
0.64	[200, 50]	0.005	0.2
0.546666666666666	[200, 50]	0.005	0.4
0.41333333333333333	[200, 50]	0.005	0.5

In [91]:

```
#use best parameters to train model
W = network_weights(vocab_size=len(train_voc_set),embedding_dim=300,hidden_dim=hdim_his[ind
W[0] = w_glove
for i in range(len(W)):
    print('Shape W'+str(i), W[i].shape)
W, loss_tr, dev_loss = SGD(train_indices_x, Y_tr=train_y,
                            W=W.
                             X_dev=dev_indices_x,
                             Y dev=dev y,
                             lr=lr_his[ind],
                             dropout=drop_his[ind],
                             freeze_emb=True,
                             tolerance=0.0001,
                             epochs=50)
Epocii. 12 | 114111 1033. 0.20/0111/30/330<del>1</del>3/ | DCV 1033. 0.2110110/<del>11</del>033
Epoch: 13 | Train loss: 0.18401984100297528 | Dev loss: 0.1910326302465990
Epoch: 14 | Train loss: 0.15608280133177393 | Dev loss: 0.1694495314915581
Epoch: 15 | Train loss: 0.1585296901535762 | Dev loss: 0.1735731317679735
Epoch: 16 | Train loss: 0.17357361265567023 | Dev loss: 0.1843862458325776
Epoch: 17 | Train loss: 0.1689096068225143 | Dev loss: 0.1820939831906187
Epoch: 18 | Train loss: 0.1845781325580215 | Dev loss: 0.19366855206727324
Epoch: 19 | Train loss: 0.187484538543126 | Dev loss: 0.19845834296444215
Epoch: 20 | Train loss: 0.142257529597178 | Dev loss: 0.1593739136507278
Epoch: 21 | Train loss: 0.13397894044796793 | Dev loss: 0.1499274382145753
Epoch: 22 | Train loss: 0.13609049846507595 | Dev loss: 0.1539676378808554
Epoch: 23 | Train loss: 0.1750883819973416 | Dev loss: 0.18635389255631912
Epoch: 24 | Train loss: 0.15845126512427843 | Dev loss: 0.1744797852464582
Epoch: 25 | Train loss: 0.15879847175281941 | Dev loss: 0.1771930314972779
Epoch: 26 | Train loss: 0.09310445800059576 | Dev loss: 0.0937442776944847
```

In [92]:

```
x = np.linspace(0,len(loss_tr),len(loss_tr))
y1, y2 = loss_tr, dev_loss

plt.plot(x, y1,label='Training loss')
plt.plot(x, y2, label='Validation loss')

plt.title('Training Monitoring')
plt.xlabel('Epochs')
plt.ylabel('Loss')
plt.legend()
plt.show()
```



In [93]:

Accuracy: 0.7977777777778 Precision: 0.8099701165161451 Recall: 0.7977777777778 F1-Score: 0.7950477027476411

Full Results

Add your final results here:

Model	Precision	Recall	F1-Score	Accuracy
Average Embedding	0.8588888888888889	0.8619866910203213	0.8588888888888889	0.8585911905628026
Average Embedding (Pre- trained)	0.88666666666666667	0.887098394913521	0.8866666666666666667	0.8868245713460235
Average Embedding (Pretrained) + X hidden layers	0.79777777777778	0.8099701165161451	0.7977777777778	0.7950477027476411

Please discuss why your best performing model is better than the rest.

The last model which is the one with pre-embedding and hidden layers is tge worst. The model with pre-trained embedding is better than average embedding according to the accuracy. For pre-trained embedding, we use tge GloVe model, this model could help to do co-courrence count. With hidden layers could consume more computation resources. I'll say at first, I didn't put the seed in the function, and the accuracy of the last model could reach 87.5%. However, once I put the seed in it, the accuracy always around 80%. This issue makes me confuse. I'll try to find out the reason in my future work.

In []:		