General information In this kernel I'll work with data from Quora Insincere Questions Classification Competition. This dataset is interesting for NLP researching. We will try to find insincere questions which aren't usefull or are even harmful. I'll do a simple EDA and try an LSTM-CNN model. In []: import numpy as np import pandas as pd import matplotlib.pyplot as plt import seaborn as sns %matplotlib inline from nltk.tokenize import TweetTokenizer import datetime import lightgbm as lgb from scipy import stats from scipy.sparse import hstack, csr matrix from sklearn.model_selection import train test split, cross val score from wordcloud import WordCloud from collections import Counter from nltk.corpus import stopwords from nltk.util import ngrams from sklearn.feature extraction.text import TfidfVectorizer from sklearn.preprocessing import StandardScaler from sklearn.linear_model import LogisticRegression from sklearn.svm import LinearSVC from sklearn.multiclass import OneVsRestClassifier import time pd.set option('max colwidth', 400) from keras.preprocessing.text import Tokenizer from keras.preprocessing.sequence import pad sequences from keras.layers import Dense, Input, LSTM, Embedding, Dropout, Activation, Conv1D, GRU, CuDNNGRU, CuD NNLSTM, BatchNormalization from keras.layers import Bidirectional, GlobalMaxPool1D, MaxPooling1D, Add, Flatten, Masking from keras.layers import GlobalAveragePooling1D, GlobalMaxPooling1D, concatenate, SpatialDropout1D from keras.models import Model, load model from keras import initializers, regularizers, constraints, optimizers, layers, callbacks from keras import backend as K from keras.engine import InputSpec, Layer from keras.optimizers import Adam from keras.callbacks import ModelCheckpoint, TensorBoard, Callback, EarlyStopping, ReduceLROnPlateau from sklearn.preprocessing import OneHotEncoder In []: import os print(os.listdir("../input/embeddings/glove.840B.300d/")) In []: | train = pd.read csv("../input/train.csv") test = pd.read csv("../input/test.csv") sub = pd.read_csv('../input/sample_submission.csv') Data overview This is a kernel competition, where we can't use external data. As a result we can use only train and test datasets as well as embeddings which were provided by organizers. In []: import os print('Available embeddings:', os.listdir("../input/embeddings/")) In []: train["target"].value_counts() We have a serious disbalance - only ~6% of data are positive. No wonder the metric for the competition is f1-score. train.head() In []: In the dataset we have only texts of questions. In []: print('Average word length of questions in train is {0:.0f}.'.format(np.mean(train['question_text'].app ly(lambda x: len(x.split())))) print('Average word length of questions in test is {0:.0f}.'.format(np.mean(test['question text'].apply (**lambda** x: len(x.split())))) In []: print('Max word length of questions in train is {0:.0f}.'.format(np.max(train['question text'].apply(la mbda x: len(x.split())))) print('Max word length of questions in test is {0:.0f}.'.format(np.max(test['question text'].apply(lamb **da** x: len(x.split())))) In []: print('Average character length of questions in train is {0:.0f}.'.format(np.mean(train['question text'].apply(lambda x: len(x))))print('Average character length of questions in test is {0:.0f}.'.format(np.mean(test['question text']. apply(lambda x: len(x))))As we can see on average questions in train and test datasets are similar, but there are quite long questions in train dataset. In []: max features = 90000 tk = Tokenizer(lower = True, filters='', num words=max features) full text = list(train['question text'].values) + list(test['question text'].values) tk.fit_on_texts(full_text) In []: | train tokenized = tk.texts to sequences(train['question text'].fillna('missing')) test tokenized = tk.texts to sequences(test['question text'].fillna('missing')) In []: | train['question text'].apply(lambda x: len(x.split())).plot(kind='hist'); plt.yscale('log'); plt.title('Distribution of question text length in characters') We can see that most of the questions are 40 words long or shorter. Let's try having sequence length equal to 70 for now. In []: $\max len = 70$ X train = pad sequences(train tokenized, maxlen = max len) X test = pad sequences(test tokenized, maxlen = max len) In []: embedding_path = "../input/embeddings/glove.840B.300d/glove.840B.300d.txt" #embedding path = "../input/embeddings/paragram 300 s1999/paragram 300 s1999.txt" In []: embed size = 300 In []: def get coefs(word, *arr): return word, np.asarray(arr, dtype='float32') embedding index = dict(get coefs(*o.split(" ")) for o in open(embedding path, encoding='utf-8', errors= all_embs = np.stack(embedding_index.values()) emb_mean,emb_std = all_embs.mean(), all_embs.std() word index = tk.word index nb_words = min(max_features, len(word_index)) embedding_matrix = np.random.normal(emb_mean, emb_std, (nb_words + 1, embed_size)) for word, i in word index.items(): if i >= max features: continue embedding_vector = embedding_index.get(word) if embedding_vector is not None: embedding_matrix[i] = embedding_vector In []: ohe = OneHotEncoder(sparse=False) y ohe = ohe.fit transform(train['target'].values.reshape(-1, 1)) For now I'll use an architecture from my previous kernel in another competition. The architecture in the following: input with embedding; then we have separate "branches" - GRU and LSTM; • each "branch" is processed by two Conv1D layers separately; each Conv1D layer has average and max pooling layers; all pooling layers are concatenated; · two dense layers in the end; In []: def build model(lr=0.0, lr d=0.0, units=0, spatial dr=0.0, kernel size1=3, kernel size2=2, dense units= 128, dr=0.1, conv size=32, epochs=20): file path = "best model.hdf5" check point = ModelCheckpoint(file path, monitor = "val loss", verbose = 1, save best only = True, mode = "min") early stop = EarlyStopping(monitor = "val loss", mode = "min", patience = 3) inp = Input(shape = (max len,)) x = Embedding(max features + 1, embed size, weights = [embedding matrix], trainable = False)(inp) x1 = SpatialDropout1D(spatial dr)(x)x gru = Bidirectional (CuDNNGRU (units, return sequences = True))(x1) x lstm = Bidirectional(CuDNNLSTM(units, return sequences = True))(x1) x conv1 = Conv1D(conv size, kernel size=kernel size1, padding='valid', kernel initializer='he unifo rm')(x gru) avg pool1 gru = GlobalAveragePooling1D()(x conv1) max pool1 gru = GlobalMaxPooling1D()(x conv1) x conv2 = Conv1D(conv size, kernel size=kernel size2, padding='valid', kernel initializer='he unifo rm')(x gru) avg pool2 gru = GlobalAveragePooling1D()(x conv2) max pool2 gru = GlobalMaxPooling1D()(x conv2) x conv3 = Conv1D(conv size, kernel size=kernel size1, padding='valid', kernel initializer='he unifo rm') (x_lstm) avg pool1 lstm = GlobalAveragePooling1D()(x conv3) max pool1 lstm = GlobalMaxPooling1D()(x conv3) x conv4 = Conv1D(conv size, kernel size=kernel size2, padding='valid', kernel initializer='he unifo rm')(x lstm) avg pool2 lstm = GlobalAveragePooling1D()(x conv4) max pool2 lstm = GlobalMaxPooling1D()(x conv4) x = concatenate([avg pool1 gru, max pool1 gru, avg pool2 gru, max pool2 gru, avg pool1 lstm, max pool1 lstm, avg pool2 lstm, max pool2 lstm]) x = BatchNormalization()(x)x = Dropout(dr)(Dense(dense_units, activation='relu') (x)) x = BatchNormalization()(x)x = Dropout(dr)(Dense(int(dense units / 2), activation='relu') (x)) x = Dense(2, activation = "sigmoid")(x)model = Model(inputs = inp, outputs = x) model.compile(loss = "binary crossentropy", optimizer = Adam(lr = lr, decay = lr d), metrics = ["ac curacy"]) model.summary() history = model.fit(X_train, y_ohe, batch_size = 512, epochs = epochs, validation_split=0.1, verbose = 1, callbacks = [check point, early stop]) model = load model(file path) return model In []: | %%time model = build model(lr = 1e-4, lr d = 0, units = 64, spatial dr = 0.5, kernel size1=4, kernel size2=3, dense_units=16, dr=0.1, conv_size=16, epochs=5) In []: | # pred = model.predict(X test, batch size = 1024, verbose = 1) # predictions = np.round(np.argmax(pred, axis=1)).astype(int) # sub['prediction'] = predictions # sub.to_csv("submission.csv", index=False) In []: def build model1(lr=0.0, lr d=0.0, units=0, spatial dr=0.0, kernel size1=3, kernel size2=2, dense units =128, dr=0.1, conv_size=32, epochs=20): file path = "best model.hdf5" check point = ModelCheckpoint(file path, monitor = "val loss", verbose = 1, save best only = True, mode = "min") early_stop = EarlyStopping(monitor = "val_loss", mode = "min", patience = 3) inp = Input(shape = (max len,)) x = Embedding(max features + 1, embed size, weights = [embedding matrix], trainable = False)(inp) x1 = SpatialDropout1D(spatial_dr)(x) x gru = Bidirectional(CuDNNGRU(units, return sequences = **True**))(x1) x_conv1 = Conv1D(conv_size, kernel_size=kernel_size1, padding='valid', kernel_initializer='he_unifo rm')(x gru) avg pool1 gru = GlobalAveragePooling1D()(x conv1) max pool1 gru = GlobalMaxPooling1D()(x conv1) x conv2 = Conv1D(conv size, kernel size=kernel size2, padding='valid', kernel initializer='he unifo rm')(x gru) avg pool2 gru = GlobalAveragePooling1D()(x conv2) max pool2 gru = GlobalMaxPooling1D()(x conv2) x = concatenate([avg_pool1_gru, max_pool1_gru, avg_pool2_gru, max_pool2_gru]) x = BatchNormalization()(x)x = Dropout(dr)(Dense(dense units, activation='relu')(x)) x = BatchNormalization()(x)x = Dropout(dr)(Dense(int(dense_units / 2), activation='relu') (x)) x = Dense(2, activation = "sigmoid")(x)model = Model(inputs = inp, outputs = x) model.compile(loss = "binary_crossentropy", optimizer = Adam(lr = lr, decay = lr_d), metrics = ["ac curacy"]) model.summary() history = model.fit(X train, y ohe, batch size = 512, epochs = epochs, validation split=0.1, verbose = 1, callbacks = [check point, early stop]) model = load_model(file_path) return model In []: # model1 = build model1(lr = 1e-4, lr d = 1e-7, units = 128, spatial dr = 0.3, kernel size1=4, kernel size1=6.3, kernel size1=6.3ze2=3, dense_units=32, dr=0.3, conv_size=32, epochs=5) In []: |#model1 1 = build model1(|1r = 1e-4, |1r d = 1e-7, |2, units = 128, spatial dr = 0.3, kernel size1=4, kernel size2=3, dense_units=32, dr=0.1, conv_size=32, epochs=5) def build model2(lr=0.0, lr d=0.0, units=0, spatial dr=0.0, kernel size1=3, kernel size2=2, dense units =128, dr=0.1, conv size=32, epochs=20): file_path = "best_model.hdf5" check point = ModelCheckpoint(file path, monitor = "val loss", verbose = 1, save best only = True, mode = "min") early stop = EarlyStopping(monitor = "val loss", mode = "min", patience = 3) inp = Input(shape = (max len,)) x = Embedding(max features + 1, embed size, weights = [embedding matrix], trainable = False)(inp) x1 = SpatialDropout1D(spatial_dr)(x) x_gru = Bidirectional(CuDNNGRU(units * 2, return_sequences = True))(x1) x gru = Bidirectional(CuDNNGRU(units, return sequences = **True**))(x gru) x_conv1 = Conv1D(conv_size, kernel_size=kernel_size1, padding='valid', kernel_initializer='he_unifo avg pool1 gru = GlobalAveragePooling1D()(x conv1) max pool1 gru = GlobalMaxPooling1D()(x conv1) x conv2 = Conv1D(conv size, kernel size=kernel size2, padding='valid', kernel initializer='he unifo rm')(x gru) avg pool2 gru = GlobalAveragePooling1D()(x conv2) max pool2 gru = GlobalMaxPooling1D()(x conv2) x = concatenate([avg pool1 gru, max pool1 gru, avg pool2 gru, max pool2 gru]) x = BatchNormalization()(x)x = Dropout(dr)(Dense(dense_units, activation='relu') (x)) x = BatchNormalization()(x)x = Dropout(dr)(Dense(int(dense units / 2), activation='relu') (x)) x = Dense(2, activation = "sigmoid")(x) model = Model(inputs = inp, outputs = x)model.compile(loss = "binary_crossentropy", optimizer = Adam(lr = lr, decay = lr_d), metrics = ["ac curacy"]) model.summary() history = model.fit(X train, y ohe, batch size = 512, epochs = epochs, validation split=0.1, verbose = 1, callbacks = [check point, early stop]) model = load model(file path) return model In []: #%%time #model2 = build model2(lr = 1e-4, lr d = 1e-7, units = 256, spatial dr = 0.3, kernel size1=4, kernel si ze2=3, dense units=32, dr=0.1, conv size=32, epochs=5) In []: |#model3| = build model2(lr = 1e-3, lr d = 1e-7, units = 256, spatial dr = 0.3, kernel size1=4, kerneze2=3, dense units=32, dr=0.1, conv size=16, epochs=5) In []: | %%time model4 = build model2(lr = 1e-4, lr d = 1e-7, units = 64, spatial dr = 0.3, kernel size1=4, kernel size 2=3, dense units=32, dr=0.1, conv size=8, epochs=5) In []:|#model5 = build model2(lr = 1e-4, lr d = 1e-7, units = 256, spatial dr = 0.1, kernel size1=4, kernel si ze2=3, dense units=32, dr=0.1, conv size=16, epochs=5) Model with attention https://github.com/Diyago/ML-DLscripts/blob/9e161a96580efa9993805ca28f610df72fe36406/DEEP%20LEARNING/LSTM%20RNN/Sentiment%20analysis%20LSTM%20wth In []: class Attention(Layer): def __init__(self, step_dim, W regularizer=None, b regularizer=None, W constraint=None, b constraint=None, bias=True, **kwargs): 11 11 11 Keras Layer that implements an Attention mechanism for temporal data. Supports Masking. Follows the work of Raffel et al. [https://arxiv.org/abs/1512.08756] # Input shape 3D tensor with shape: `(samples, steps, features)`. # Output shape 2D tensor with shape: `(samples, features)`. Just put it on top of an RNN Layer (GRU/LSTM/SimpleRNN) with return sequences=True. The dimensions are inferred based on the output shape of the RNN. Example: model.add(LSTM(64, return sequences=True)) model.add(Attention()) self.supports masking = True #self.init = initializations.get('glorot uniform') self.init = initializers.get('glorot uniform') self.W_regularizer = regularizers.get(W_regularizer) self.b_regularizer = regularizers.get(b_regularizer) self.W_constraint = constraints.get(W_constraint) self.b constraint = constraints.get(b constraint) self.bias = bias self.step dim = step dim self.features dim = 0super(Attention, self). init (**kwargs) def build(self, input shape): assert len(input shape) == 3 self.W = self.add weight((input shape[-1],),initializer=self.init, name='{}_W'.format(self.name), regularizer=self.W regularizer, constraint=self.W constraint) self.features_dim = input_shape[-1] if self.bias: self.b = self.add weight((input shape[1],), initializer='zero', name='{}_b'.format(self.name), regularizer=self.b regularizer, constraint=self.b constraint) else: self.b = None self.built = True def compute_mask(self, input, input_mask=None): # do not pass the mask to the next layers return None def call(self, x, mask=None): # eij = K.dot(x, self.W) TF backend doesn't support it # features dim = self.W.shape[0] # step_dim = x._keras_shape[1] features dim = self.features dim step_dim = self.step_dim eij = K.reshape(K.dot(K.reshape(x, (-1, features dim)), K.reshape(self.W, (features dim, 1))), (-1, step dim))if self.bias: eij += self.b eij = K.tanh(eij) a = K.exp(eij)# apply mask after the exp. will be re-normalized next if mask is not None: # Cast the mask to floatX to avoid float64 upcasting in theano a *= K.cast(mask, K.floatx()) # in some cases especially in the early stages of training the sum may be almost zero a /= K.cast(K.sum(a, axis=1, keepdims=True) + K.epsilon(), K.floatx()) a = K.expand dims(a)weighted input = x * a#print weigthted input.shape return K.sum(weighted input, axis=1) def compute output shape(self, input shape): #return input_shape[0], input_shape[-1] return input_shape[0], self.features_dim **Attention** In []: def build model3(lr=0.0, lr d=0.0, units=0, spatial dr=0.0, dense units=128, dr=0.1, use attention=True inp = Input(shape = (max len,)) x = Embedding(max features + 1, embed size, weights = [embedding matrix], trainable = False)(inp) x1 = SpatialDropout1D(spatial dr)(x)x gru = Bidirectional(CuDNNGRU(units * 2, return sequences = True))(x1) if use attention: x att = Attention(max len)(x gru) x = Dropout(dr)(Dense(dense units, activation='relu') (x att)) else: x att = Flatten() (x gru) x = Dropout(dr)(Dense(dense units, activation='relu') (x att)) x = BatchNormalization()(x)#x = Dropout(dr)(Dense(int(dense units / 2), activation='relu')(x))x = Dense(2, activation = "sigmoid")(x)model = Model(inputs = inp, outputs = x) model.compile(loss = "binary_crossentropy", optimizer = Adam(lr = lr, decay = lr_d), metrics = ["ac curacy"]) #model.summary() #history = model.fit(X train, y ohe, batch size = 512, epochs = epochs, validation split=0.1, verbose = 1, callbacks = [check point, early stop]) #model = load model(file path) return model In []: %%time file path = "best model.hdf5" check point = ModelCheckpoint(file path, monitor = "val loss", verbose = 1, save best only = True, mode = "min") early stop = EarlyStopping(monitor = "val loss", mode = "min", patience = 3) model6 = build model3(lr = 1e-3, lr d = 1e-7, units = 64, spatial dr = 0.3, dense units=16, dr=0.1, useattention=**True**) history = model6.fit(X_train, y_ohe, batch_size = 512, epochs = 10, validation_split=0.1, verbose = 1, callbacks = [check point, early stop]) In []: | # #%%time # file path = "best model.hdf5" # check point = ModelCheckpoint(file path, monitor = "val loss", verbose = 1, save best only = True, mode = "min") # early stop = EarlyStopping(monitor = "val loss", mode = "min", patience = 3) # model6 1 = build model3(lr = 1e-3, lr d = 1e-7, units = 64, spatial dr = 0.3, dense units=16, dr=0.1, use_attention=False) # history = model6 1.fit(X train, y ohe, batch size = 512, epochs = 5, validation split=0.1, verbose = 1, callbacks = [check point, early stop]) One branch In []: def build model4(lr=0.0, lr d=0.0, units=0, spatial dr=0.0, kernel size1=3, kernel size2=2, dense units =128, dr=0.1, conv size=32, epochs=20): file path = "best model.hdf5" check point = ModelCheckpoint(file path, monitor = "val loss", verbose = 1, save best only = True, mode = "min") early stop = EarlyStopping(monitor = "val loss", mode = "min", patience = 3) inp = Input(shape = (max len,)) x = Embedding(max features + 1, embed size, weights = [embedding matrix], trainable = False)(inp) x1 = SpatialDropout1D(spatial dr)(x)x gru = Bidirectional (CuDNNGRU (units, return sequences = True))(x1) x conv1 = Conv1D(conv size, kernel size=kernel size1, padding='valid', kernel initializer='he unifo avg pool1 gru = GlobalAveragePooling1D()(x conv1) max pool1 gru = GlobalMaxPooling1D()(x conv1) x = concatenate([avg pool1 gru, max pool1 gru]) x = BatchNormalization()(x)x = Dropout(dr)(Dense(dense units, activation='relu') (x)) x = BatchNormalization()(x)#x = Dropout(dr)(Dense(int(dense units / 2), activation='relu')(x))x = Dense(2, activation = "sigmoid")(x)model = Model(inputs = inp, outputs = x) model.compile(loss = "binary_crossentropy", optimizer = Adam(lr = lr, decay = lr_d), metrics = ["ac curacy"]) model.summary() history = model.fit(X train, y ohe, batch size = 512, epochs = epochs, validation split=0.1, verbose = 1, callbacks = [check point, early stop]) model = load model(file path) return model In []: | %%time model7 = build model4(lr = 1e-4, lr d = 1e-7, units = 64, spatial dr = 0.3, kernel size1=3, dense units =32, dr=0.1, conv size=8, epochs=5) In []: |#model8| = build model4(lr = 1e-4, lr d = 1e-7, units = 128, spatial dr = 0.3, kernel size1=4, dense units=32, dr=0.1, conv size=8, epochs=5) Masking In []: | def build model5(lr=0.0, lr d=0.0, units=0, spatial dr=0.0, kernel size1=3, kernel size2=2, dense units =128, dr=0.1, conv size=32, epochs=20): file path = "best model.hdf5" check point = ModelCheckpoint(file path, monitor = "val loss", verbose = 1, save best only = True, mode = "min") early stop = EarlyStopping(monitor = "val loss", mode = "min", patience = 3) reduce lr = ReduceLROnPlateau(monitor='val loss', factor=0.1, patience=2, min lr=0.001) inp = Input(shape = (max len,)) x = Embedding(max features + 1, embed size, weights = [embedding matrix], trainable = False)(inp) x1 = SpatialDropout1D(spatial dr)(x)x m = Masking()(x1)x gru = LSTM(units)(x m)x = BatchNormalization()(x gru)x = Dropout(dr)(Dense(dense_units, activation='relu') (x)) x = BatchNormalization()(x)#x = Dropout(dr)(Dense(int(dense units / 2), activation='relu')(x))x = Dense(2, activation = "sigmoid")(x)model = Model(inputs = inp, outputs = x) model.compile(loss = "binary_crossentropy", optimizer = Adam(lr = lr, decay = lr_d), metrics = ["ac curacy"]) model.summary() history = model.fit(X_train, y_ohe, batch_size = 512, epochs = epochs, validation split=0.1, verbose = 1, callbacks = [check point, early stop, reduce lr]) model = load model(file path) return model In []: # model9 = build model5 (lr = 1e-4, lr d = 1e-7, units = 128, spatial dr = 0.3, kernel size1=4, dense units=32, dr=0.1, conv size=8, epochs=10) In []: | pred1 = model.predict(X test, batch size = 1024, verbose = 1) pred4 = model4.predict(X test, batch size = 1024, verbose = 1) pred += pred4 pred2 = model7.predict(X_test, batch_size = 1024, verbose = 1) pred += pred2 # pred3 = model9.predict(X test, batch size = 1024, verbose = 1) # pred += pred3 pred4 = model6.predict(X_test, batch_size = 1024, verbose = 1) pred += pred4 # pred5 = model7.predict(X test, batch size = 1024, verbose = 1) # pred += pred5 pred = pred / 4In []: #pred = model9.predict(X test, batch size = 1024, verbose = 1) predictions = np.round(np.argmax(pred, axis=1)).astype(int) sub['prediction'] = predictions sub.to_csv("submission.csv", index=False)