VAE_log_regression

October 1, 2019

1 Creation of synthetic data for Wisoncsin Breat Cancer data set using a Variational AutoEncoder. Tested using a logistic regression model.

1.1 Aim

To test a a Variational AutoEncoder (VAE) for synthesising data that can be used to train a logistic regression machine learning model.

1.2 Data

Raw data is avilable at:

https://www.kaggle.com/uciml/breast-cancer-wisconsin-data

1.3 Basic methods description

• Create synthetic data by use of a Variational AutoEncoder

Kingma, D.P. and Welling, M. (2013) Auto-encoding variational bayes. arXiv preprint arXiv:1312.6114,2013.

• Train logistic regression model on synthetic data and test against held-back raw data

1.4 Code & results

```
[1]: import numpy as np
  import pandas as pd
  import matplotlib.pyplot as plt

from sklearn.linear_model import LogisticRegression
  from sklearn.model_selection import train_test_split
  from sklearn.decomposition import PCA

# Turn warnings off for notebook publication
  import warnings
  warnings.filterwarnings("ignore")
```

1.4.1 Import Data

```
[2]: def load_data():
        n n n n
        Load Wisconsin Breast Cancer Data Set
        Inputs
        _____
        None
        Returns
        X: NumPy array of X
        y: Numpy array of y
        col_names: column names for X
        11 11 11
        # Load data and drop 'id' column
        data = pd.read_csv('./wisconsin.csv')
        data.drop('id', axis=1, inplace=True)
        # Change 'diagnosis' column to 'malignant', and put in last column place
        malignant = pd.DataFrame()
        data['malignant'] = data['diagnosis'] == 'M'
        data.drop('diagnosis', axis=1, inplace=True)
        # Split data in X and y
        X = data.drop(['malignant'], axis=1)
        y = data['malignant']
        # Get col names and convert to NumPy arrays
       X_col_names = list(X)
        X = X.values
        y = y.values
        return data, X, y, X_col_names
```

1.4.2 Data processing

Split X and y into training and test sets

```
[3]: def split_into_train_test(X, y, test_proportion=0.25):

""""

Randomly split X and y numpy arrays into training and test data sets

Inputs
-----
X and y NumPy arrays
```

```
Returns
-----
X_test, X_train, y_test, y_train Numpy arrays
"""

X_train, X_test, y_train, y_test = \
    train_test_split(X, y, shuffle=True, test_size=test_proportion)

return X_train, X_test, y_train, y_test
```

Standardise data

```
[4]: def standardise_data(X_train, X_test):
    """"
    Standardise training and tets data sets according to mean and standard
    deviation of test set

Inputs
-----
X_train, X_test NumPy arrays

Returns
-----
X_train_std, X_test_std
"""

mu = X_train.mean(axis=0)
std = X_train.std(axis=0)

X_train_std = (X_train - mu) / std
X_test_std = (X_test - mu) / std
return X_train_std, X_test_std
```

1.4.3 Calculate accuracy measures

```
[5]: def calculate_diagnostic_performance(actual, predicted):
    """ Calculate sensitivty and specificty.

Inputs
-----
actual, predted numpy arrays (1 = +ve, 0 = -ve)

Returns
-----
A dictionary of results:
```

```
1) accuracy: proportion of test results that are correct
2) sensitivity: proportion of true +ve identified
3) specificity: proportion of true -ve identified
4) positive likelihood: increased probability of true +ve if test +ve
5) negative likelihood: reduced probability of true +ve if test -ve
6) false positive rate: proportion of false +ves in true -ve patients
7) false negative rate: proportion of false -ves in true +ve patients
8) positive predictive value: chance of true +ve if test +ve
9) negative predictive value: chance of true -ve if test -ve
10) actual positive rate: proportion of actual values that are +ve
11) predicted positive rate: proportion of predicted vales that are +ve
12) recall: same as sensitivity
13) precision: the proportion of predicted +ve that are true +ve
14) f1 = 2 * ((precision * recall) / (precision + recall))
*false positive rate is the percentage of healthy individuals who
incorrectly receive a positive test result
* alse neagtive rate is the percentage of diseased individuals who
incorrectly receive a negative test result
11 11 11
# Calculate results
actual_positives = actual == 1
actual negatives = actual == 0
test_positives = predicted == 1
test_negatives = predicted == 0
test_correct = actual == predicted
accuracy = test_correct.mean()
true_positives = actual_positives & test_positives
false_positives = actual_negatives & test_positives
true_negatives = actual_negatives & test_negatives
sensitivity = true_positives.sum() / actual_positives.sum()
specificity = np.sum(true_negatives) / np.sum(actual_negatives)
positive_likelihood = sensitivity / (1 - specificity)
negative_likelihood = (1 - sensitivity) / specificity
false_postive_rate = 1 - specificity
false negative rate = 1 - sensitivity
positive_predictive_value = true_positives.sum() / test_positives.sum()
negative_predicitive_value = true_negatives.sum() / test_negatives.sum()
actual_positive_rate = actual.mean()
predicted_positive_rate = predicted.mean()
recall = sensitivity
precision = \
    true_positives.sum() / (true_positives.sum() + false_positives.sum())
f1 = 2 * ((precision * recall) / (precision + recall))
```

```
# Add results to dictionary
results = dict()
results['accuracy'] = accuracy
results['sensitivity'] = sensitivity
results['specificity'] = specificity
results['positive_likelihood'] = positive_likelihood
results['negative_likelihood'] = negative_likelihood
results['false postive rate'] = false postive rate
results['false_postive_rate'] = false_postive_rate
results['false_negative_rate'] = false_negative_rate
results['positive_predictive_value'] = positive_predictive_value
results['negative_predicitive_value'] = negative_predicitive_value
results['actual_positive_rate'] = actual_positive_rate
results['predicted_positive_rate'] = predicted_positive_rate
results['recall'] = recall
results['precision'] = precision
results['f1'] = f1
return results
```

1.4.4 Logistic Regression Model

```
[6]: def fit_and_test_logistic_regression_model(X_train, X_test, y_train, y_test):
    """"
    Fit and test logistic regression model.
    Return a dictionary of accuracy measures.
    Calls on `calculate_diagnostic_performance` to calculate results

Inputs
    -----
    X_train, X_test NumPy arrays

Returns
    ------
A dictionary of accuracy results.
    """

# Fit logistic regression model
lr = LogisticRegression(C=0.1)
lr.fit(X_train,y_train)

# Predict tets set labels
y_pred = lr.predict(X_test_std)

# Get accuracy results
```

```
accuracy_results = calculate_diagnostic_performance(y_test, y_pred)
return accuracy_results
```

1.4.5 Synthetic Data Method - Variational AutoEncoder

```
[7]: def sampling(args):
        n n n
        Reparameterization trick by sampling from an isotropic unit Gaussian.
        Instead of sampling from Q(z|X), sample epsilon = N(0,I)
        z = z_{mean} + sqrt(var) * epsilon
        # Arguments
            args (tensor): mean and log of variance of Q(z|X)
        # Returns
            z (tensor): sampled latent vector
        import tensorflow
        from tensorflow.keras import backend as K
        z_mean, z_log_var = args
        batch = K.shape(z_mean)[0]
        dim = K.int_shape(z_mean)[1]
        \# by default, random_normal has mean = 0 and std = 1.0
        epsilon = K.random_normal(shape=(batch, dim))
        sample = z_{mean} + K.exp(0.5 * z_{log_var}) * epsilon
        return sample
[8]: def make_synthetic_data_vae(X_original, y_original,
                                 batch_size=256,
                                latent_dim=8,
                                 epochs=10000,
                                learning_rate=2e-5,
                                dropout=0.25,
                                number_of_samples=1000):
        11 11 11
        Synthetic data generation.
        Calls on `get_principal_component_model` for PCA model
        If number of components not defined then the function sets it to the number
          of features in X
        Inputs
```

```
original_data: X, y numpy arrays
number_of_samples: number of synthetic samples to generate
n_components: number of principal components to use for data synthesis
Returns
X_synthetic: NumPy array
y_synthetic: NumPy array
11 11 11
import tensorflow
from tensorflow.keras import layers
from tensorflow.keras.models import Model
from tensorflow.keras.optimizers import Adam
from tensorflow.keras import backend as K
from tensorflow.keras.losses import mean_squared_error
# Standardise X
mean = X_original.mean(axis=0)
std = X_original.mean(axis=0)
X_std = (X_original - mean) / std
# network parameters
input_shape = X_original.shape[1]
intermediate_dim = X_original.shape[1]
# Split the training data into positive and negative
mask = y_original == 1
X_train_pos = X_std[mask]
mask = y_original == 0
X_train_neg = X_std[mask]
# Set up list for positive and negative synthetic data sets
synthetic_X_sets = []
# Run fir twice: once for positive label examples, the other for negative
for training_set in [X_train_pos, X_train_neg]:
    # Clear Tensorflow
    K.clear_session()
    # VAE model = encoder + decoder
    # build encoder model
    inputs = layers.Input(shape=input_shape, name='encoder_input')
    encode_dense_1 = layers.Dense(
        intermediate_dim, activation='relu')(inputs)
```

```
dropout_encoder_layer_1 = layers.Dropout(dropout)(encode_dense_1)
encode_dense_2 = layers.Dense(
    intermediate_dim, activation='relu')(dropout_encoder_layer_1)
z_mean = layers.Dense(latent_dim, name='z_mean')(encode_dense_2)
z_log_var = layers.Dense(latent_dim, name='z_log_var')(encode_dense_2)
# use reparameterization trick to push the sampling out as input
# note that "output_shape" isn't necessary with the TensorFlow backend
z = layers.Lambda(
    sampling, output_shape=(latent_dim,), name='z')([z_mean, z_log_var])
# instantiate encoder model
encoder = Model(inputs, [z_mean, z_log_var, z], name='encoder')
# build decoder model
latent_inputs = layers.Input(shape=(latent_dim,), name='z sampling')
decode_dense_1 = layers.Dense(
    intermediate_dim, activation='relu')(latent_inputs)
dropout_decoder_layer_1 = layers.Dropout(dropout)(decode_dense_1)
decode_dense_2 = layers.Dense(
    intermediate_dim, activation='relu')(dropout_decoder_layer_1)
outputs = layers.Dense(input_shape)(decode_dense_2)
# instantiate decoder model
decoder = Model(latent_inputs, outputs, name='decoder')
# instantiate VAE model
outputs = decoder(encoder(inputs)[2])
vae = Model(inputs, outputs, name='vae_mlp')
# Train the autoencoder
optimizer = Adam(lr=learning_rate)
# VAE loss = mse_loss or xent_loss + kl_loss
vae.compile(optimizer=optimizer, loss = mean_squared_error)
# Train the autoencoder
vae.fit(training_set, training_set,
```

```
batch_size = batch_size,
            shuffle = True,
            epochs = epochs,
            verbose=0)
    # Produce synthetic data
    z_new = np.random.normal(size = (number_of_samples, latent_dim))
    reconst = decoder.predict(np.array(z_new))
    reconst = mean + (reconst * std)
    synthetic_X_sets.append(reconst)
    # Clear models
    K.clear session()
    del encoder
    del decoder
    del vae
# Combine data
# Combine positive and negative and shuffle rows
X_synthetic = np.concatenate(
        (synthetic_X_sets[0], synthetic_X_sets[1]), axis=0)
y_synthetic_pos = np.ones((number_of_samples, 1))
y_synthetic_neg = np.zeros((number_of_samples, 1))
y_synthetic = np.concatenate((y_synthetic_pos, y_synthetic_neg), axis=0)
# Randomise order of X, y
synthetic = np.concatenate((X_synthetic, y_synthetic), axis=1)
shuffle_index = np.random.permutation(np.arange(X_synthetic.shape[0]))
synthetic = synthetic[shuffle_index]
X_synthetic = synthetic[:,0:-1]
y_synthetic = synthetic[:,-1]
return X_synthetic, y_synthetic
```

1.4.6 Main code

```
[9]: # Load data
original_data, X, y, X_col_names = load_data()

# Set up results DataFrame
results = pd.DataFrame()
```

Fitting classification model to raw data

```
[10]: # Set number of replicate runs
number_of_runs = 30
```