

References: Chapter 17 of Geron's book. For 1-Dim plots, Keras tutorial :

<https://www.tensorflow.org/tutorials/generative/autoencoder>

This file trains a VAE with the instances of the normal digits in the training data.

Then, it measures the reconstruction loss for the digits in the test data.

The reconstruction loss for the instances of the abnormal digits in the test data is higher.

A threshold is determined based on the distribution of the reconstruction losses of the normal training data (threshold = mean + 2.5*std of this distribution).

Then, if the reconstruction loss of a digit in the test data is higher than this threshold, it is classified as abnormal.

By comparing with the known labels of test data (with T for normal digit(s) and F for abnormal digit(s)), the confusion matrix and the accuracy is calculated.

```
import sklearn
import tensorflow as tf
from tensorflow import keras
import numpy as np

import matplotlib as mpl
import matplotlib.pyplot as plt
mpl.rc('axes', labelsize=14)
mpl.rc('xtick', labelsize=12)
mpl.rc('ytick', labelsize=12)

from sklearn.metrics import accuracy_score, precision_score, recall_score, confusion_matrix
from sklearn.model_selection import train_test_split
```

Loading the MNIST data and forming arrays of the normal training data, the

- ▼ validation data (normal and abnormal) and the test data (normal and abnormal)

```
#Labels
# 0 T-shirt/top
# 1 Trouser
# 2 Pullover
# 3 Dress
# 4 Coat
# 5 Sandal
# 6 Shirt
# 7 Sneaker
```

```
# 8 Bag
# 9 Ankle boot
```

```
nl1 = 0
nl2 = 2
nl3 = 6
abn1 = 7
abn2 = 7
```

```
(x_train_0, y_train_0), (x_test, y_test) = keras.datasets.fashion_mnist.load_data()

x_train_0 = x_train_0.astype(np.float32) / 255
x_test = x_test.astype(np.float32) / 255

train_size = x_train_0.shape[0] * 9 // 10

x_train, x_valid, y_train, y_valid = train_test_split(x_train_0, y_train_0, train_size = train_size)

normal_data = x_train[(y_train == nl1) | (y_train == nl2) | (y_train == nl3)]           # Normal
normal_labels = y_train[(y_train == nl1) | (y_train == nl2) | (y_train == nl3)]

valid_data = x_valid[(y_valid == abn1) | (y_valid == abn2) | (y_valid == nl1) | (y_valid == nl2)]
valid_labels = y_valid[(y_valid == abn1) | (y_valid == abn2) | (y_valid == nl1) | (y_valid == nl2)]

test_data = x_test[(y_test == abn1) | (y_test == abn2) | (y_test == nl1) | (y_test == nl2)]
test_labels = y_test[(y_test == abn1) | (y_test == abn2) | (y_test == nl1) | (y_test == nl2)]

test_labels_T_F = np.where((test_labels == nl1) | (test_labels == nl2) | (test_labels == nl3))
# Array of T and F, T where test digits are normal and F where test digits are abnormal

valid_labels_T_F = np.where((valid_labels == nl1) | (valid_labels == nl2) | (valid_labels == nl3))
# Array of T and F, T where test digits are normal and F where test digits are abnormal
```

```
Downloading data from https://storage.googleapis.com/tensorflow/tf-keras-datasets/train-32768/29515 [=====] - 0s 0us/step
40960/29515 [=====] - 0s 0us/step
Downloading data from https://storage.googleapis.com/tensorflow/tf-keras-datasets/train-26427392/26421880 [=====] - 0s 0us/step
26435584/26421880 [=====] - 1s 0us/step
Downloading data from https://storage.googleapis.com/tensorflow/tf-keras-datasets/t10k-16384/5148 [=====]
Downloading data from https://storage.googleapis.com/tensorflow/tf-keras-datasets/t10k-4423680/4422102 [=====] - 0s 0us/step
4431872/4422102 [=====] - 0s 0us/step
```



```
normal_data.shape, normal_labels.shape, valid_data.shape, valid_labels.shape, test_data.shape

((16180, 28, 28), (16180,), (2419, 28, 28), (2419,), (4000, 28, 28), (4000,))
```

```
normal_test_data = test_data[(test_labels == nl1) | (test_labels == nl2) | (test_labels == nl3)]
```

```

abnormal_test_data = test_data[(test_labels == abn1) | (test_labels == abn2)]      # The ab
normal_test_labels = test_labels[(test_labels == n11) | (test_labels == n12) | (test_labels =
abnormal_test_labels = test_labels[(test_labels == abn1) | (test_labels == abn2)]      # Their

normal_test_data.shape, abnormal_test_data.shape

((3000, 28, 28), (1000, 28, 28))

normal_valid_data = valid_data[(valid_labels == n11) | (valid_labels == n12) | (valid_labels
abnormal_valid_data = valid_data[(valid_labels == abn1) | (valid_labels == abn2)]      # Th
normal_valid_labels = valid_labels[(valid_labels == n11) | (valid_labels == n12) | (valid_lab
abnormal_valid_labels = valid_labels[(valid_labels == abn1) | (valid_labels == abn2)]      # Th

normal_valid_data.shape, abnormal_valid_data.shape

((1820, 28, 28), (599, 28, 28))

```

▼ Building and training the network

```

K = keras.backend
# def rounded_accuracy(y_true, y_pred):
#   return keras.metrics.binary_accuracy(tf.round(y_true), tf.round(y_pred))

# For details please see Geron's book. Uses the reparametrization trick to do stochastic
# sampling from the MVN distribution, while allowing the 2 parallel layers containing the
# means and stds of the MVN distribution for each dimension to be trained via
# backpropogation of the error signal.
class Sampling(keras.layers.Layer):
    def call(self, inputs):
        mean, log_var = inputs
        return K.random_normal(tf.shape(log_var)) * K.exp(log_var / 2) + mean

# For details please see Geron's book.
codings_size = 16    # The number of dimensions of the MVN distribution in the sampling layer

inputs = keras.layers.Input(shape=[28, 28])
z = keras.layers.Flatten()(inputs)
z = keras.layers.Dense(256, activation="selu")(z)
z = keras.layers.Dense(128, activation="selu")(z)
z = keras.layers.Dense(64, activation="selu")(z)

# Parallel layers at the end of the encoder for means
# and standard deviations of the Multivariate Normal (MVN) distribution
# in the dimensions of the coding size (here 32).
codings_mean = keras.layers.Dense(codings_size)(z)

```

```

codings_log_var = keras.layers.Dense(codings_size)(z)

# Sampling layer at the end of the encoder
codings = Sampling()([codings_mean, codings_log_var])
variational_encoder = keras.models.Model(
    inputs=[inputs], outputs=[codings_mean, codings_log_var, codings])

decoder_inputs = keras.layers.Input(shape=[codings_size])
x = keras.layers.Dense(64, activation="selu")(decoder_inputs)
x = keras.layers.Dense(128, activation="selu")(x)
x = keras.layers.Dense(256, activation="selu")(x)
x = keras.layers.Dense(28 * 28, activation="sigmoid")(x)
outputs = keras.layers.Reshape([28, 28])(x)
variational_decoder = keras.models.Model(inputs=[decoder_inputs], outputs=[outputs])

_, _, codings = variational_encoder(inputs)
reconstructions = variational_decoder(codings)
variational_ae = keras.models.Model(inputs=[inputs], outputs=[reconstructions])

# The latent loss function
# latent_loss = -0.5 * K.sum(
#     1 + codings_log_var - K.exp(codings_log_var) - K.square(codings_mean),
#     axis=-1)

# Add the latent loss to the reconstruction loss
# variational_ae.add_loss(K.mean(latent_loss) / 784.)

# For the reconstruction loss binary cross-entropy loss is used.
# For details please see Chapter 17 of Geron's book (Stacked AE and VAE sections)
variational_ae.compile(loss="binary_crossentropy", optimizer="rmsprop")

checkpoint_cb = keras.callbacks.ModelCheckpoint("wo_latent_VAE_model", monitor="val_loss", sa

history = variational_ae.fit(normal_data, normal_data, epochs=100, batch_size=128, callbacks=
                               validation_data=(normal_valid_data, normal_valid_data), shuffle=127/127 [=====] - 1s 8ms/step - loss: 0.3387 - val_loss: 0.3 ▲
Epoch 75/100
127/127 [=====] - 1s 8ms/step - loss: 0.3386 - val_loss: 0.34
Epoch 76/100
121/127 [=====>..] - ETA: 0s - loss: 0.3385INFO:tensorflow:Asse
127/127 [=====] - 3s 24ms/step - loss: 0.3387 - val_loss: 0.
Epoch 77/100
127/127 [=====] - 1s 8ms/step - loss: 0.3385 - val_loss: 0.3
Epoch 78/100
127/127 [=====] - 1s 8ms/step - loss: 0.3385 - val_loss: 0.3
Epoch 79/100
121/127 [=====>..] - ETA: 0s - loss: 0.3384INFO:tensorflow:Asse
127/127 [=====] - 3s 25ms/step - loss: 0.3384 - val_loss: 0.
Epoch 80/100
127/127 [=====] - 1s 8ms/step - loss: 0.3384 - val_loss: 0.3
Epoch 81/100
127/127 [=====] - 1s 8ms/step - loss: 0.3383 - val_loss: 0.3
Epoch 82/100

```

```
[-----] - 1s 8ms/step - loss: 0.3383 - val_loss: 0.3
Epoch 83/100
[-----] - 1s 8ms/step - loss: 0.3382 - val_loss: 0.3
Epoch 84/100
[-----] - 1s 8ms/step - loss: 0.3381 - val_loss: 0.3
Epoch 85/100
[-----] - 1s 8ms/step - loss: 0.3381 - val_loss: 0.3
Epoch 86/100
[-----] - 1s 8ms/step - loss: 0.3381 - val_loss: 0.3
Epoch 87/100
[-----] - 1s 8ms/step - loss: 0.3380 - val_loss: 0.3
Epoch 88/100
[-----] - 1s 8ms/step - loss: 0.3379 - val_loss: 0.3
Epoch 89/100
[-----] - 1s 8ms/step - loss: 0.3379 - val_loss: 0.3
Epoch 90/100
[-----] - 1s 8ms/step - loss: 0.3379 - val_loss: 0.3
Epoch 91/100
[-----] - ETA: 0s - loss: 0.3378INFO:tensorflow:Asse
[-----] - 3s 27ms/step - loss: 0.3378 - val_loss: 0.
Epoch 92/100
[-----] - 1s 8ms/step - loss: 0.3378 - val_loss: 0.3
Epoch 93/100
[-----] - 1s 8ms/step - loss: 0.3377 - val_loss: 0.3
Epoch 94/100
[=====>...] - ETA: 0s - loss: 0.3377INFO:tensorflow:Asse
[-----] - 4s 32ms/step - loss: 0.3377 - val_loss: 0.
Epoch 95/100
[-----] - 1s 8ms/step - loss: 0.3376 - val_loss: 0.3
Epoch 96/100
[-----] - 1s 8ms/step - loss: 0.3376 - val_loss: 0.3
Epoch 97/100
[-----] - 1s 8ms/step - loss: 0.3376 - val_loss: 0.3
Epoch 98/100
[-----] - 1s 8ms/step - loss: 0.3375 - val_loss: 0.3
Epoch 99/100
[-----] - 1s 8ms/step - loss: 0.3375 - val_loss: 0.3
Epoch 100/100
[=====>...] - ETA: 0s - loss: 0.3374INFO:tensorflow:Asse
```

```
plt.plot(history.history["loss"], label="Training Loss")
plt.plot(history.history["val_loss"], label="Validation Loss")
plt.legend()
```

```
<matplotlib.legend.Legend at 0x7f613e639e90>
```



```
model = variational_ae
model.summary(expand_nested=True, show_trainable=True)
```

```
Model: "model_2"
```

Layer (type)	Output Shape	Param #	Trainable
<hr/>			
input_1 (InputLayer)	[(None, 28, 28)]	0	Y
<hr/>			
model (Functional)	[(None, 16), (None, 16), (None, 16)]	244192	Y
<hr/>			
input_1 (InputLayer)	[(None, 28, 28)]	0	Y
flatten (Flatten)	(None, 784)	0	Y
dense (Dense)	(None, 256)	200960	Y
dense_1 (Dense)	(None, 128)	32896	Y
dense_2 (Dense)	(None, 64)	8256	Y
dense_3 (Dense)	(None, 16)	1040	Y
dense_4 (Dense)	(None, 16)	1040	Y
sampling (Sampling)	(None, 16)	0	Y
<hr/>			
model_1 (Functional)	(None, 28, 28)	243920	Y
<hr/>			
input_2 (InputLayer)	[(None, 16)]	0	Y
dense_5 (Dense)	(None, 64)	1088	Y
dense_6 (Dense)	(None, 128)	8320	Y
dense_7 (Dense)	(None, 256)	33024	Y
dense_8 (Dense)	(None, 784)	201488	Y
reshape (Reshape)	(None, 28, 28)	0	Y
<hr/>			
Total params: 488,112			
Trainable params: 488,112			
Non-trainable params: 0			

```

model_encoder = variational_encoder
# model_encoder.summary(expand_nested=True, show_trainable=True)

model_decoder = variational_decoder
# model_decoder.summary(expand_nested=True, show_trainable=True)

model_layers = np.array(model.layers)
n_layers = model_layers.shape[0]
# np.concatenate((np.arange(n_layers).reshape(n_layers,1), model_layers.reshape(n_layers,1)),

```

The original and reconstructed images for the first 30 instances of the normal training data, validation data, normal validation data, abnormal validation data, test data, normal test data, and abnormal test data

```

def plot_image(image):
    plt.imshow(image, cmap="binary")
    plt.axis("off")

def show_reconstructions(model, images, n_images=5):
    reconstructions = model.predict(images[:n_images])
    fig = plt.figure(figsize=(n_images * 1.5, 3))
    for image_index in range(n_images):
        plt.subplot(2, n_images, 1 + image_index)
        plot_image(images[image_index])
        plt.subplot(2, n_images, 1 + n_images + image_index)
        plot_image(reconstructions[image_index])

show_reconstructions(variational_ae, normal_data, 30)
plt.show()

```



```

show_reconstructions(variational_ae, valid_data, 30)
plt.show()

```



```

show_reconstructions(variational_ae, normal_valid_data, 30)

```

```
plt.show()
```



```
show_reconstructions(variational_ae, abnormal_valid_data, 30)
plt.show()
```



```
show_reconstructions(variational_ae, test_data, 30)
plt.show()
```



```
show_reconstructions(variational_ae, normal_test_data, 30)
plt.show()
```



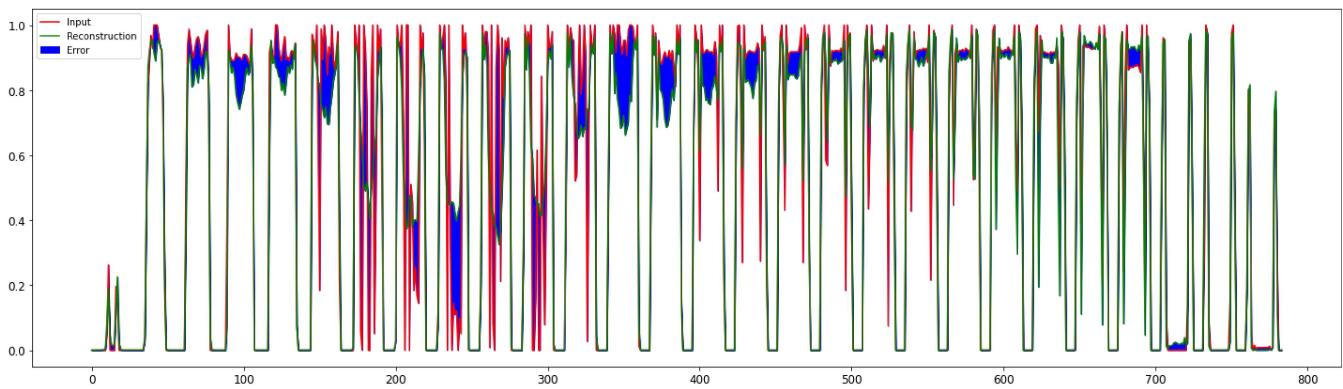
```
show_reconstructions(variational_ae, abnormal_test_data, 30)
plt.show()
```



1-Dim plot of pixels of the first normal test data

```
reconstructions_nl_test = variational_ae.predict(normal_test_data)
```

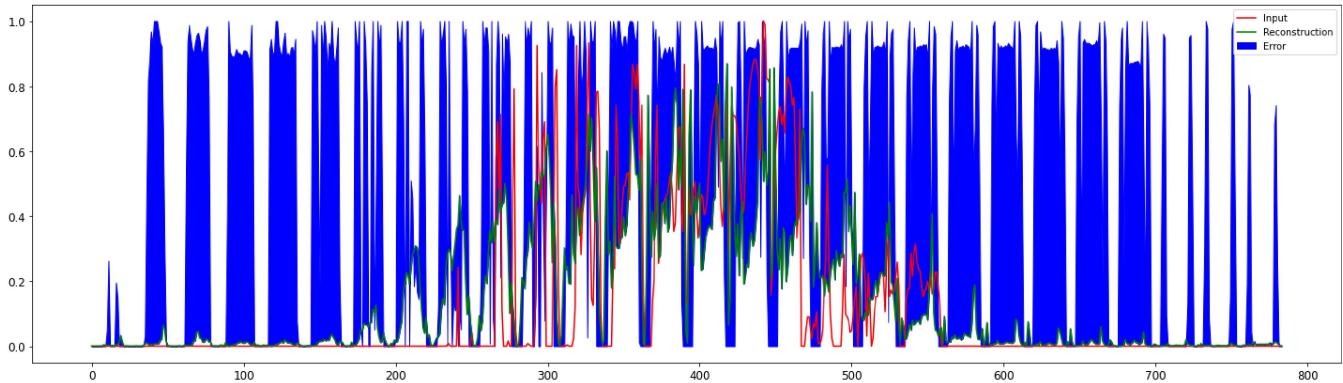
```
plt.figure(figsize=(25,7))
plt.plot(normal_test_data[0].ravel(), 'r')
plt.plot(reconstructions_nl_test[0].ravel(), 'g')
plt.fill_between(np.arange(28*28), reconstructions_nl_test[0].ravel(), normal_test_data[0].ravel())
plt.legend(labels=["Input", "Reconstruction", "Error"])
plt.show()
```



1-Dim plot of pixels of the first abnormal test data

```
reconstructions_abn_test = variational_ae.predict(abnormal_test_data)
```

```
plt.figure(figsize=(25,7))
plt.plot(abnormal_test_data[0].ravel(), 'r')
plt.plot(reconstructions_abn_test[0].ravel(), 'g')
plt.fill_between(np.arange(28*28), reconstructions_abn_test[0].ravel(), normal_test_data[0].ravel(), color='blue')
plt.legend(labels=["Input", "Reconstruction", "Error"])
plt.show()
```

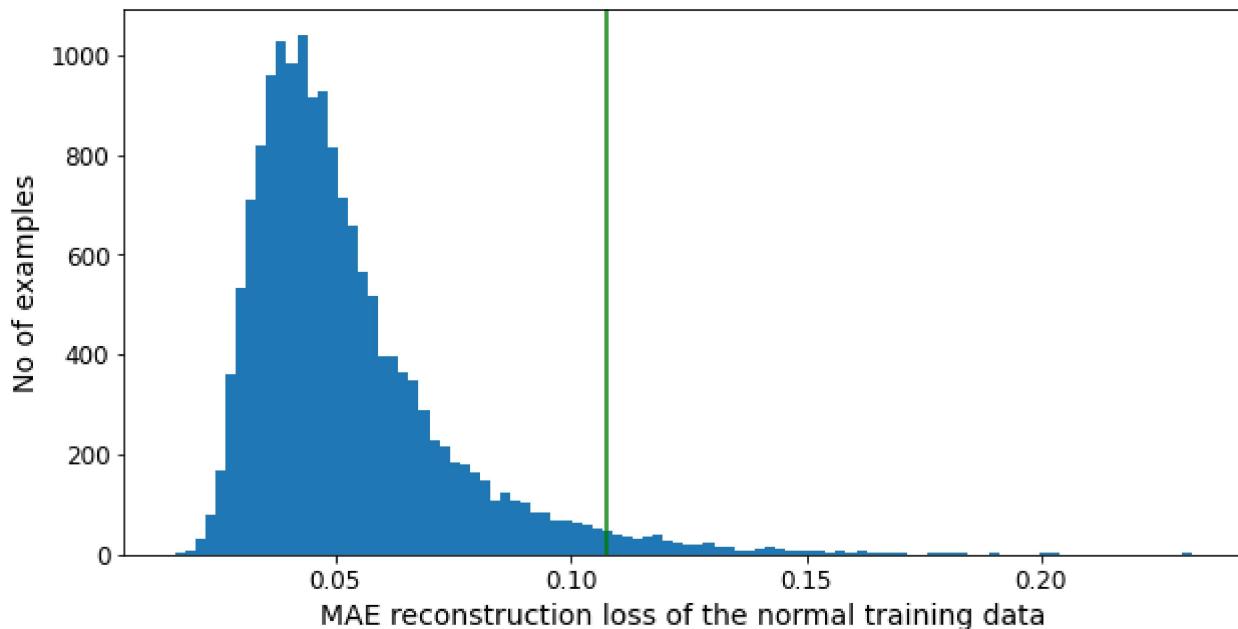


▼ Distributions of the reconstruction losses and the calculation of the threshold.

Distribution of the reconstruction losses of the normal training data

```
reconstructions = variational_ae.predict(normal_data)
train_loss = tf.keras.losses.mae(reconstructions.reshape(-1, 784), normal_data.reshape(-1, 784))
plt.figure(figsize=(10,5))
plt.hist(train_loss[None,:], bins=100)
threshold1 = np.mean(train_loss) + 2.5*np.std(train_loss)
```

```
plt.axvline(threshold1,c='g')
plt.xlabel("MAE reconstruction loss of the normal training data")
plt.ylabel("No of examples")
plt.show()
```



```
mean1 = np.mean(train_loss)
std1 = np.std(train_loss)
print("Mean: ", mean1)
print("Std: ", std1)
```

```
Mean:  0.0527785
Std:  0.021893192
```

```
threshold_train_mean_2_5_std = np.mean(train_loss) + 2.5*np.std(train_loss)
print("Threshold based on the mean of the training data MAE reconstruction losses + 2.5 std:
```

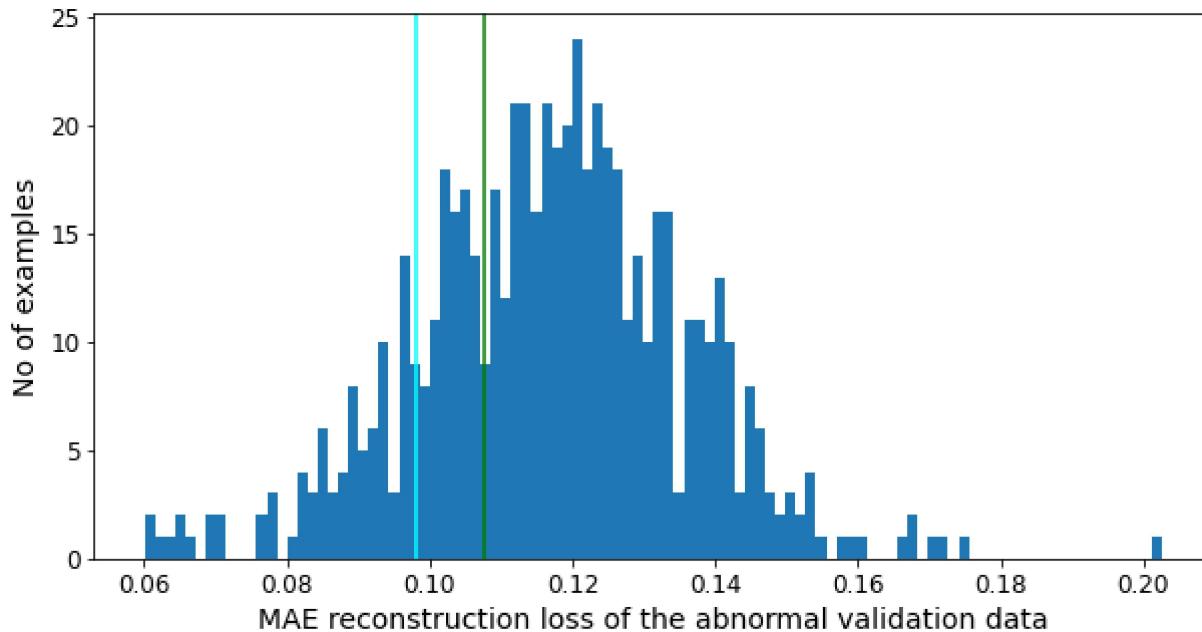
```
Threshold based on the mean of the training data MAE reconstruction losses + 2.5 std:  €
```

```
threshold1 = threshold_train_mean_2_5_std
```

Distribution of the reconstruction losses of the abnormal validation data

```
reconstructions = variational_ae.predict(abnormal_valid_data)
abn_valid_loss = tf.keras.losses.mae(reconstructions.reshape(-1,784), abnormal_valid_data.reshape(-1,784))
plt.figure(figsize=(10,5))
plt.hist(abn_valid_loss[None, :], bins=100)
threshold2 = np.mean(abn_valid_loss) - np.std(abn_valid_loss)
plt.axvline(threshold2,c='cyan')
plt.axvline(threshold1,c='g')
plt.xlabel("MAE reconstruction loss of the abnormal validation data")
```

```
plt.ylabel("No of examples")
plt.show()
```



```
abnormal_valid_mean_loss = np.mean(abn_valid_loss)
```

```
abnormal_valid_mean_loss , np.std(abn_valid_loss)
```

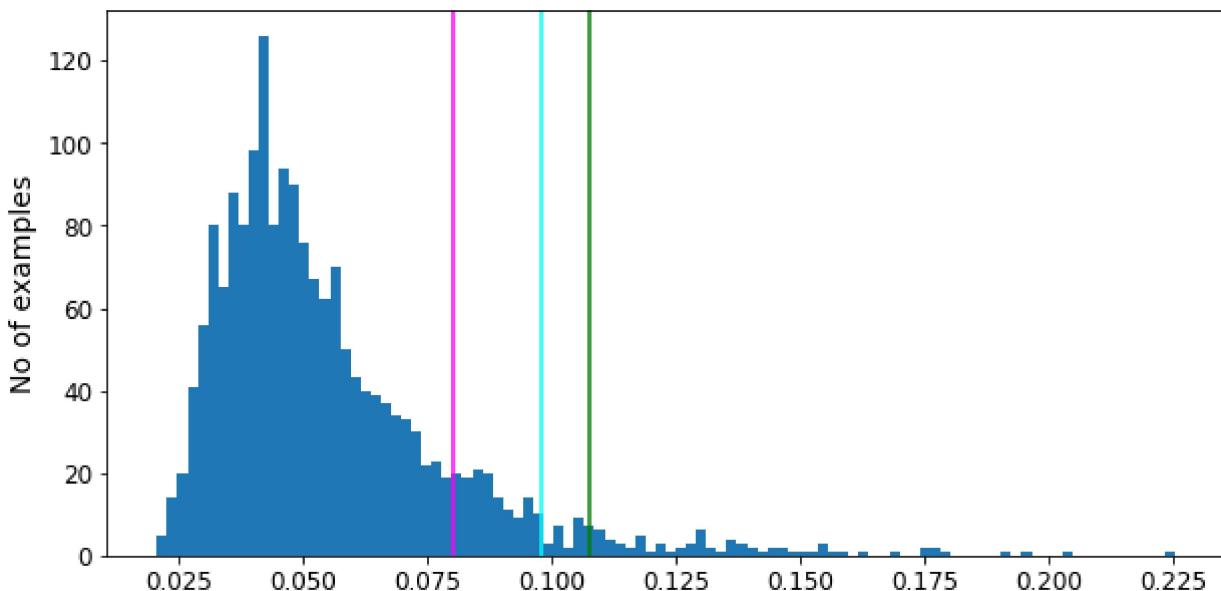
```
(0.11706579, 0.01891169)
```

```
threshold2 = abnormal_valid_mean_loss - np.std(abn_valid_loss)
print("Threshold2: ", threshold2)
```

```
Threshold2: 0.0981541
```

Distribution of the reconstruction losses of the normal validation data

```
reconstructions = variational_ae.predict(normal_valid_data)
nl_valid_loss = tf.keras.losses.mae(reconstructions.reshape(-1,784), normal_valid_data.reshape(-1,784))
plt.figure(figsize=(10,5))
plt.hist(nl_valid_loss[None, :], bins=100)
threshold3 = np.mean(nl_valid_loss) + np.std(nl_valid_loss)
plt.axvline(threshold3, c='magenta')
plt.axvline(threshold2, c='cyan')
plt.axvline(threshold1, c='g')
plt.xlabel("MAE reconstruction loss of the normal validation data")
plt.ylabel("No of examples")
plt.show()
```



```
normal_valid_mean_loss = np.mean(nl_valid_loss)
```

```
normal_valid_mean_loss , np.std(nl_valid_loss)
```

```
(0.05539579, 0.02476868)
```

```
threshold3 = normal_valid_mean_loss + np.std(nl_valid_loss)
print("Threshold3: ", threshold3)
```

```
Threshold3: 0.08016447
```

Calculation of a preliminary threshold based on $(\text{threshold2} + \text{threshold3}) / 2$ = Average of (mean + std of the distribution of the reconstruction losses of the normal validation data) and (mean - std of the distribution of the reconstruction losses of the abnormal validation data)

```
Avg_of_threshold_2_3 = (threshold2 + threshold3)/2
print("Average of threshold 2 and 3: ", Avg_of_threshold_2_3)
```

```
Average of threshold 2 and 3: 0.0891592800617218
```

```
threshold4 = Avg_of_threshold_2_3
```

Calculation of the threshold that gives the best accuracy on the validation data and set this as the threshold.

```
def predict(model, data, threshold):
    reconstructions = model.predict(data)
    loss = tf.keras.losses.mae(reconstructions.reshape(-1, 784), data.reshape(-1, 784))
```

```

return tf.math.less(loss, threshold)

increment = (abnormal_valid_mean_loss - normal_valid_mean_loss)/100
thresholds = np.arange(normal_valid_mean_loss, abnormal_valid_mean_loss, increment)
thrs_size = thresholds.shape[0]
accuracies = np.zeros(thrs_size)
for i in range(thrs_size):
    preds = predict(variational_ae, valid_data, thresholds[i])
    accuracies[i] = accuracy_score(preds, valid_labels_T_F)
argmax = np.argmax(accuracies)
valid_data_best_threshold = thresholds[argmax]
print("The best threshold based on validation data: ", valid_data_best_threshold)

The best threshold based on validation data:  0.09609798803925529

thr_acc = np.zeros((thrs_size, 2))
thr_acc[:, 0] = thresholds
thr_acc[:, 1] = accuracies
thr_acc[argmax-2:argmax+3]

array([[0.09486459,  0.92269533],
       [0.09548129,  0.92269533],
       [0.09609799,  0.92393551],
       [0.09671469,  0.92145515],
       [0.09733139,  0.92062836]])

```

```
threshold5 = valid_data_best_threshold
```

```
threshold = threshold5
```

Distribution of the reconstruction losses of all the validation data (normal and abnormal)

The blue line is threshold4 (= the average of threshold3 [magenta] and threshold2 [cyan]).

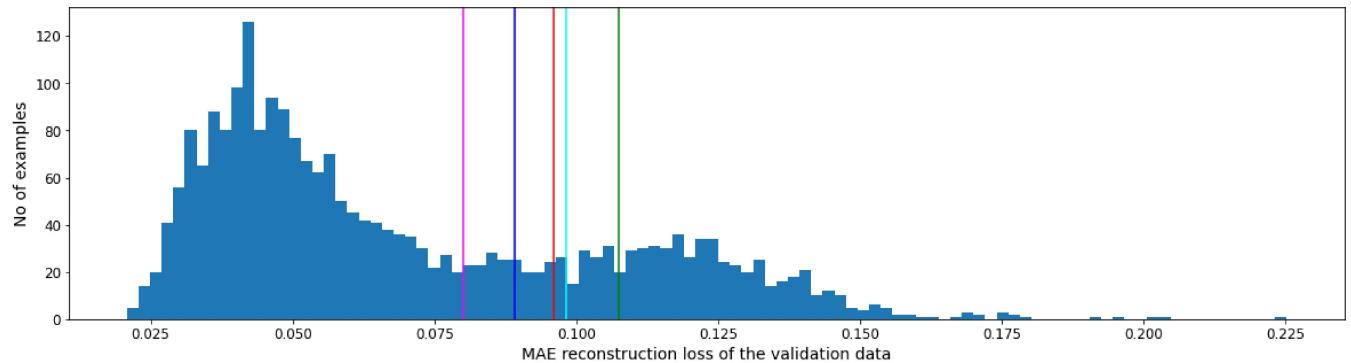
The red line is the threshold that gives the best accuracy for the validation data.

```

reconstructions = variational_ae.predict(valid_data)
valid_loss = tf.keras.losses.mae(reconstructions.reshape(-1,784), valid_data.reshape(-1,784))
plt.figure(figsize=(20,5))
plt.hist(valid_loss[None, :], bins=100)
plt.axvline(threshold, c='r')
plt.axvline(threshold4, c='b')
plt.axvline(threshold2, c='cyan')
plt.axvline(threshold3, c='magenta')
plt.axvline(threshold1, c='green')

```

```
plt.xlabel("MAE reconstruction loss of the validation data")
plt.ylabel("No of examples")
plt.show()
```



▼ Distribution of the reconstruction losses of the test data (normal and abnormal)

The blue line is threshold4 (= the average of threshold3 [magenta] and threshold2 [cyan]).

The red line is the threshold that gives the best accuracy for the validation data.

```
reconstructions = variational_ae.predict(test_data)
test_loss = tf.keras.losses.mae(reconstructions.reshape(-1,784), test_data.reshape(-1,784))
plt.figure(figsize=(20,5))
plt.hist(test_loss[None, :], bins=100)
plt.axvline(threshold, c='r')
plt.axvline(threshold4, c='b')
plt.axvline(threshold2, c='cyan')
plt.axvline(threshold3, c='magenta')
plt.axvline(threshold1, c='green')
plt.xlabel("MAE reconstruction loss of the test data")
plt.ylabel("No of examples")
plt.show()
```

Mean and standard deviation of reconstruction losses for normal and abnormal test data

```
reconstructions = variational_ae.predict(normal_test_data)
nl_test_loss = tf.keras.losses.mae(reconstructions.reshape(-1,784), normal_test_data.reshape(
np.mean(nl_test_loss) , np.std(nl_test_loss))

(0.055375244, 0.023500977)

reconstructions = variational_ae.predict(abnormal_test_data)
abn_test_loss = tf.keras.losses.mae(reconstructions.reshape(-1,784), abnormal_test_data.resha
np.mean(abn_test_loss) , np.std(abn_test_loss)

(0.117089696, 0.018401166)
```

Calculation of the accuracy and the confusion matrix on the test data with threshold set based on the best threshold from the validation data

```
# def predict(model, data, threshold):
#     reconstructions = model.predict(data)
#     loss = tf.keras.losses.mae(reconstructions.reshape(-1, 784), data.reshape(-1, 784))
#     return tf.math.less(loss, threshold)
mean_nl_test_loss = np.mean(nl_test_loss)
std_nl_test_loss = np.std(nl_test_loss)

mean_abn_test_loss = np.mean(abn_test_loss)
std_abn_test_loss = np.std(abn_test_loss)

def print_stats(predictions, labels):
    cf = confusion_matrix(labels, predictions)

    print("mean_nl_test_loss: {}".format(mean_nl_test_loss))
    print("std_nl_test_loss: {}".format(std_nl_test_loss))
    print("mean_abn_test_loss: {}".format(mean_abn_test_loss))
    print("std_abn_test_loss: {}".format(std_abn_test_loss))

    print("Confusion Matrix: \n prediction: F T ")
    print("      {} {}".format(preds[preds == False].shape[0], preds[preds == True].sh
print(" label: F [[{} {}] {}] {}".format(cf[0,0], cf[0,1], test_labels_T_F[test_labels_T_
print("      T [{} {}]] {}".format(cf[1,0], cf[1,1], test_labels_T_F[test_labels_T_
print("Accuracy = {}".format(accuracy_score(labels, predictions)))
print("Precision = {}".format(precision_score(labels, predictions)))
print("Recall = {}".format(recall_score(labels, predictions)))

preds = predict(variational_ae, test_data, threshold)
```

```
print_stats(preds, test_labels_T_F)

mean_nl_test_loss: 0.055375244468450546
std_nl_test_loss: 0.023500977084040642
mean_abn_test_loss: 0.11708969622850418
std_abn_test_loss: 0.018401166424155235
Confusion Matrix:
prediction: F      T
              1074   2926
label: F    [[883   117]    1000
             [191   2809]]   3000
Accuracy = 0.923
Precision = 0.9600136705399863
Recall = 0.9363333333333334
```

```
print(confusion_matrix(test_labels_T_F, preds))
```

```
[[ 883  117]
 [ 191 2809]]
```

Extra accuracy info

Just informative. Please record the above accuracy.

Accuracy on the test data with threshold set based on (threshold2 + threshold3) / 2 = Average of (mean + std of the distribution of the reconstruction losses of the normal validation data) and (mean - std of the distribution of the reconstruction losses of the abnormal validation data)

```
preds = predict(variational_ae, test_data, Avg_of_threshold_2_3)
print_stats(preds, test_labels_T_F)
```

```
mean_nl_test_loss: 0.055375244468450546
std_nl_test_loss: 0.023500977084040642
mean_abn_test_loss: 0.11708969622850418
std_abn_test_loss: 0.018401166424155235
Confusion Matrix:
prediction: F      T
              1178   2822
label: F    [[929   71]    1000
             [249  2751]]   3000
Accuracy = 0.92
Precision = 0.9748405386250886
Recall = 0.917
```

Accuracy on the test data with threshold set based on the mean of the training data MAE

```

preds = predict(variational_ae, test_data, threshold_train_mean_2_5_std)
print_stats(preds, test_labels_T_F)

mean_nl_test_loss: 0.055375244468450546
std_nl_test_loss: 0.023500977084040642
mean_abn_test_loss: 0.11708969622850418
std_abn_test_loss: 0.018401166424155235
Confusion Matrix:
prediction: F      T
            843    3157
label: F   [[728    272]    1000
            T   [115    2885]]   3000
Accuracy = 0.90325
Precision = 0.91384225530567
Recall = 0.9616666666666667

```

Extra Info

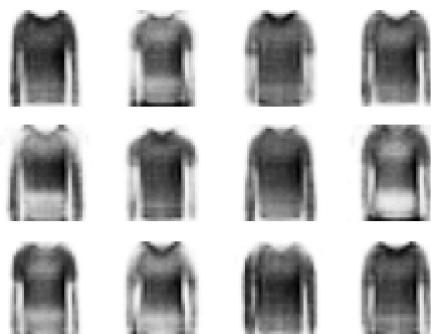
Giving the VAE codings (please see book) (Just informative, not the goal here)

```

def plot_multiple_images(images, n_cols=None):
    n_cols = n_cols or len(images)
    n_rows = (len(images) - 1) // n_cols + 1
    if images.shape[-1] == 1:
        images = np.squeeze(images, axis=-1)
    plt.figure(figsize=(n_cols, n_rows))
    for index, image in enumerate(images):
        plt.subplot(n_rows, n_cols, index + 1)
        plt.imshow(image, cmap="binary")
        plt.axis("off")

codings = tf.random.normal(shape=[12, codings_size])
images = variational_decoder(codings).numpy()
plot_multiple_images(images, 4)
# save_fig("vae_generated_images_plot", tight_layout=False)

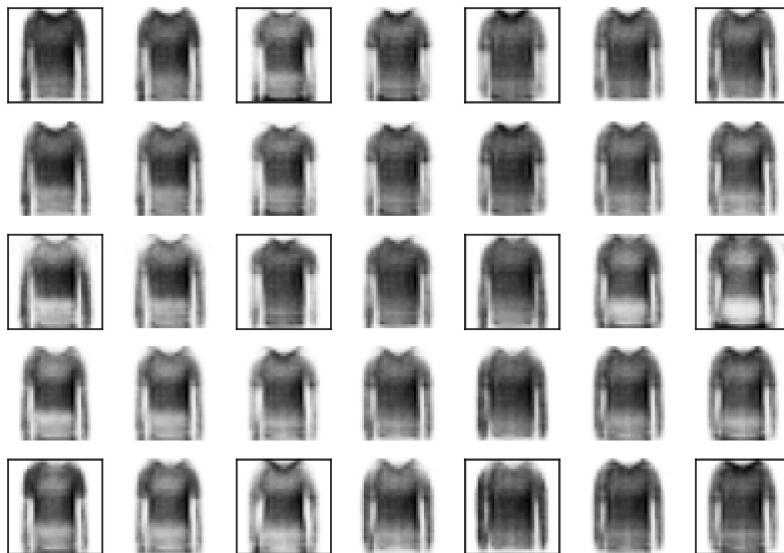
```



```
codings_grid = tf.reshape(codings, [1, 3, 4, codings_size])
```

```
larger_grid = tf.image.resize(codings_grid, size=[5, 7])
interpolated_codings = tf.reshape(larger_grid, [-1, codings_size])
images = variational_decoder(interpolated_codings).numpy()

plt.figure(figsize=(7, 5))
for index, image in enumerate(images):
    plt.subplot(5, 7, index + 1)
    if index%7==0 and index//7==0:
        plt.gca().get_xaxis().set_visible(False)
        plt.gca().get_yaxis().set_visible(False)
    else:
        plt.axis("off")
    plt.imshow(image, cmap="binary")
# save_fig("semantic_interpolation_plot", tight_layout=False)
```



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