CNN for TFBS predictions and model interpretability

• Author	資工四甲 U0924028 符絮棻
: Tags	CNN TFBS
∅ 程式碼 來源	<u>Deep-Learning-for-Genomics-/Chapter10/model_interpretability.ipynb at main · PacktPublishing/Deep-Learning-for-Genomics- · GitHub</u>

在調控基因表達和進化的過程中,如DNA複製及RNA轉錄,轉錄因子(TF)與TF結合位點(TFBS)的結合有至關重要的作用 ⇒ 精確類比基因的特異性及尋找TFBS有助於探索細胞表達的機制。

```
TFBS預測問題中,所有基因序列會根據DNA序列中是否能找到TFBS分為兩類,由標籤0或1表示 \begin{cases} 0 & \text{ 陰性 } - \  \  \  \  \end{pmatrix}有與TF的結合位點 \begin{cases} 1 & \text{ 陽性 } - \  \  \  \end{cases}
```

Distribution of Positive and Negative samples

```
In [7]: print(pd, DataFrame(Y). value_counts())

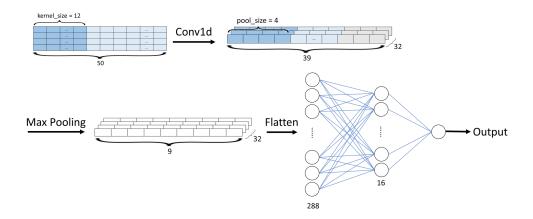
0 1013
1 987
dtype: int64
```

CNN Model

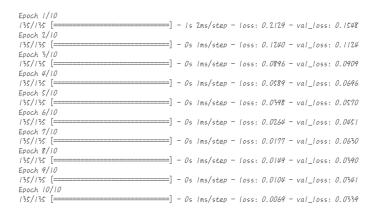
原始模型

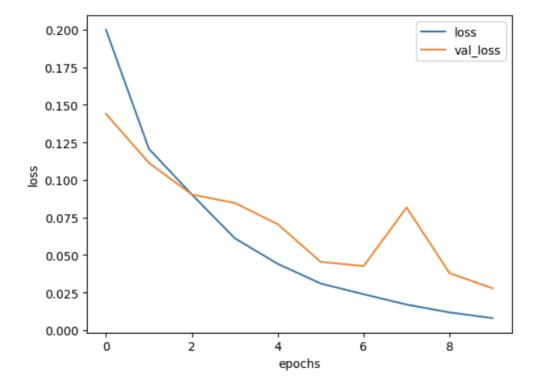
```
In [10]:
          model = Sequential()
          model.add(ConvID(filters=32, kernel_size=12, input_shape=(50, 4)))
          model.add(MaxPoolingID(pool_size=4))
          model.add(Flatten())
          model.add(Dense(16, activation='relu'))
          model.add(Dense(I, activation='sigmoid'))
          model.compile(loss='mean_squared_error', optimizer='adam')
          model.summary()
        Model: "sequential"
         Layer (type)
                                   Output Shape
                                                              Param #
         convld (ConvlD)
                                     (None, 39, 32)
                                                              1568
         max_poolingId (MaxPoolingID (None, 9, 32)
         flatten (Flatten)
                                     (None, 288)
                                     (None, 16)
         dense (Dense)
                                                              4624
         dense_I (Dense)
                                     (None, 1)
```

Total params: 6,209 Trainable params: 6,209 Non-trainable params: 0 由於原始輸出層僅有一個神經元,模型的預測值是一個介於 0 到 1 之間的實數,用於表示是否存在 TFBS 的可能性。



訓練結果:





雖然訓練集及驗證集的損失隨著epoch下降,但驗證集的損失呈現波動狀態,而非穩定下降 ⇒ 猜測模型可能有overfitting 的情況發生。

修改模型

- 1. 使用padding進行填充,提升邊緣資訊的可參考程度
- 2. 修改pool_size,使所有特徵值都能進行max pooling
- 3. 輸出層更改為兩個神經元,使預測結果與實際標籤更相符

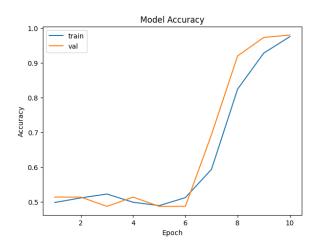
```
#標籤做one hot encoding
labels = pd.read_csv('labels.txt')
Labels = np.array(labels).reshape(-1)
print(Labels[:5])
iecd = iec.fit_transform(Labels[:])
iecd = np.array(iecd).reshape(-1, 1)
ohed = ohe.fit_transform(iecd)
Y = np.stack(ohed.toarray())
```

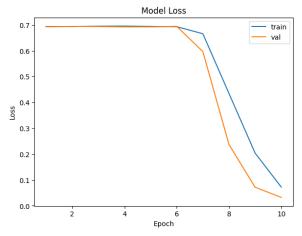
- 4. 增加模型的深度
- 5. 設定dropout ratio, 避免overfitting

```
dropout = 0.2
model = Sequential()
model.add(Conv1D(filters=32, kernel_size=12, input_shape=(50, 4), padding='same'))
model.add(MaxPooling1D(pool_size=5))
model.add(Flatten())
model.add(Dense(256, activation='relu'))
model.add(Dropout(dropout))
model.add(Dense(128, activation='relu'))
model.add(Dropout(dropout))
model.add(Dense(64, activation='relu'))
model.add(Dropout(dropout))
model.add(Dense(32, activation='relu'))
model.add(Dropout(dropout))
model.add(Dense(16, activation='relu'))
model.add(Dense(16, activation='relu'))
model.add(Dense(8, activation='relu'))
model.add(Dense(8, activation='relu'))
model.add(Dense(4, activation='relu'))
model.add(Dense(4, activation='relu'))
model.add(Dense(2, activation='softmax'))
model.compile(loss='mean_squared_error', optimizer='adam')
model.summary()
```

訓練結果:

```
Epoch 1/10
135/135 [==
Epoch 2/10
                                      ===] - 2s 7ms/step - 1oss: 0.6938 - accuracy: 0.4978 - val_loss: 0.6927 - val_accuracy: 0.5133
                                   =====] 1 6ms/step - loss: 0.6935 - accuracy: 0.5111 - val_loss: 0.6940 - val_accuracy: 0.5133
135/135 [==
                                    =====] - 1s 5ms/step - loss: 0.6948 - accuracy: 0.5222 - val_loss: 0.6934 - val_accuracy: 0.4867
135/135 [=
Epoch 4/10
135/135 [==
                                        =] - 1s 5ms/step - 1oss: 0.6952 - accuracy: 0.4985 - val_loss: 0.6922 - val_accuracy: 0.5133
Epoch 5/10
135/135 [=
                                           - 1s 5ms/step - 1oss: 0.6940 - accuracy: 0.4889 - val_loss: 0.6922 - val_accuracy: 0.4867
Epoch 6/10
135/135 [=
Epoch 7/10
                                           - 1s 6ms/step - loss: 0.6929 - accuracy: 0.5119 - val_loss: 0.6943 - val_accuracy: 0.4867
135/135 [==
                                  ======] - 1s 6ms/step - 1oss: 0.6657 - accuracy: 0.5933 - val_loss: 0.5966 - val_accuracy: 0.6933
                                    ====] - 1s 8ms/step - loss: 0.4345 - accuracy: 0.8244 - val_loss: 0.2372 - val_accuracy: 0.9200
135/135 [==
Epoch 9/10
135/135 [==
                                      ===] - 1s 8ms/step - loss: 0.2039 - accuracy: 0.9281 - val_loss: 0.0717 - val_accuracy: 0.9733
Epoch 10/10
                                   :====] - 1s 8ms/step - 1oss: 0.0726 - accuracy: 0.9756 - val_loss: 0.0326 - val_accuracy: 0.9800
```





Evaluating the Model

- roc_curve → 在不同閾值設定下,分類模型的真陽性率(True Positive Rate, TPR)和假陽性率(False Positive Rate, FPR)之間的關係。
- AUC \rightarrow 考慮了不同閾值下的真陽性率及偽陽性率,並將其合成一個數值,通常取值範圍在0.5~1之間。當AUC愈大,模型的預測能力越強。
- AUPRC (精確率-召回率曲線) → 關注模型的精確率及召回率之間的權衡。

使用測試集資料評估模型的預測能力:

原始

AUC 0.9971483042022461 AUPRC 0.9969029074117326 修改後

AUC 0. 9903474903474904 AUPRC 0. 9796747967479674