# Classification

ACTL3143 & ACTL5111 Deep Learning for Actuaries
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### **Lecture Outline**

- TLDR
- Classification
- Stroke Prediction







# Classification models in Keras

If the number of classes is *c*, then:

Target	Output Layer	<b>Loss Function</b>
$\begin{array}{c} \text{Binary} \\ (c=2) \end{array}$	1 neuron with sigmoid activation	Binary Cross-Entropy
Multi-class $(c>2)$	c neurons with softmax activation	Categorical Cross- Entropy





# Optionally output logits

If the number of classes is *c*, then:

Target	Output Layer	<b>Loss Function</b>
$\begin{array}{c} \text{Binary} \\ (c=2) \end{array}$	1 neuron with linear activation	<pre>Binary Cross-Entropy (from_logits=True)</pre>
Multi-class $(c > 2)$	c neurons with linear activation	Categorical Cross- Entropy (from_logits=True)





### Code examples

#### **Binary**

```
model = Sequential([
    # Skipping the earlier layers
    Dense(1, activation="sigmoid")

    ])
model.compile(loss="binary_crossentropy")
```

#### **Binary (logits)**

```
from keras.losses import BinaryCrossentropy
model = Sequential([
    # Skipping the earlier layers
Dense(1, activation="linear")
])
loss = BinaryCrossentropy(from_logits=True)
model.compile(loss=loss)
```

#### **Multi-class**

```
1 model = Sequential([
2  # Skipping the earlier layers
3  Dense(n_classes, activation="softmax")
4 ])
5 model.compile(loss="sparse_categorical_crossentropy")
```

#### **Multi-class (logits)**

```
from keras.losses import SparseCategoricalCrossentropy

model = Sequential([
    # Skipping the earlier layers
    Dense(n_classes, activation="linear")

])

loss = SparseCategoricalCrossentropy(from_logits=True)
model.compile(loss=loss)
```







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### Iris dataset

```
from sklearn.datasets import load_iris
iris = load_iris()
names = ["SepalLength", "SepalWidth", "PetalLength", "PetalWidth"]
features = pd.DataFrame(iris.data, columns=names)
features
```

	SepalLength	SepalWidth	PetalLength	PetalWidth
O	5.1	3.5	1.4	0.2
1	4.9	3.0	1.4	0.2
•••	•••	•••	•••	•••
148	6.2	3.4	5.4	2.3
149	5.9	3.0	5.1	1.8

150 rows × 4 columns





### Target variable

```
1 iris.target_names
array(['setosa', 'versicolor', 'virginica'],
dtype='<U10')</pre>
  1 iris.target[:8]
array([0, 0, 0, 0, 0, 0, 0])
  1 target = iris.target
  2 target = target.reshape(-1, 1)
  3 target[:8]
array([[0],
       [0],
       [0],
       [0],
       [0],
       [0],
       [0],
       [0]])
```

```
classes, counts = np.unique(
             target,
             return_counts=True
  5 print(classes)
  6 print(counts)
[0 1 2]
[50 50 50]
  1 iris.target_names[
       target[[0, 30, 60]]
array([['setosa'],
       ['setosa'],
       ['versicolor']], dtype='<U10')
```





# Split the data into train and test

- 1 X\_train, X\_test, y\_train, y\_test = train\_test\_split(features, target, random\_state=24)
- 2 X\_train

	SepalLength	SepalWidth	PetalLength	PetalWidth
53	5.5	2.3	4.0	1.3
58	6.6	2.9	4.6	1.3
95	5.7	3.0	4.2	1.2
•••	•••	•••	•••	•••
145	6.7	3.0	5.2	2.3
87	6.3	2.3	4.4	1.3
131	7.9	3.8	6.4	2.0

112 rows × 4 columns

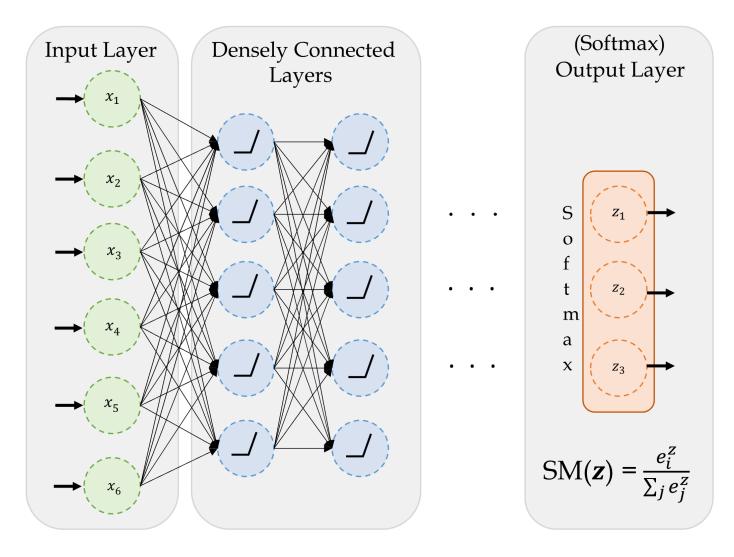
1 X\_test.shape, y\_test.shape

((38, 4), (38, 1))





### A basic classifier network



A basic network for classifying into three categories.





### Create a classifier model

```
1 NUM_FEATURES = len(features.columns)
2 NUM_CATS = len(np.unique(target))
3
4 print("Number of features:", NUM_FEATURES)
5 print("Number of categories:", NUM_CATS)

Number of features: 4
Number of categories: 3
```

#### Make a function to return a Keras model:





### Fit the model

```
1 model = build_model()
2 model.compile("adam", "sparse_categorical_crossentropy")
3
4 model.fit(X_train, y_train, epochs=5, verbose=2);

Epoch 1/5
4/4 - 0s - 93ms/step - loss: 1.3502
Epoch 2/5
4/4 - 0s - 6ms/step - loss: 1.2852
Epoch 3/5
4/4 - 0s - 6ms/step - loss: 1.2337
Epoch 4/5
4/4 - 0s - 6ms/step - loss: 1.1915
Epoch 5/5
4/4 - 0s - 6ms/step - loss: 1.1556
```





### Track accuracy as the model trains

```
1 model = build_model()
2 model.compile("adam", "sparse_categorical_crossentropy", metrics=["accuracy"])
3 model.fit(X_train, y_train, epochs=5, verbose=2);

Epoch 1/5
4/4 - 0s - 91ms/step - accuracy: 0.2946 - loss: 1.3502
Epoch 2/5
4/4 - 0s - 6ms/step - accuracy: 0.3036 - loss: 1.2852
Epoch 3/5
4/4 - 0s - 6ms/step - accuracy: 0.3036 - loss: 1.2337
Epoch 4/5
4/4 - 0s - 6ms/step - accuracy: 0.3304 - loss: 1.1915
Epoch 5/5
4/4 - 0s - 6ms/step - accuracy: 0.3393 - loss: 1.1556
```





### Run a long fit

```
model = build_model()
model.compile("adam", "sparse_categorical_crossentropy", \
metrics=["accuracy"])

time hist = model.fit(X_train, y_train, epochs=500, \
validation_split=0.25, verbose=False)
```

CPU times: user 24.5 s, sys: 3.69 s, total: 28.2 s Wall time: 23.7 s

#### Evaluation now returns both *loss* and *accuracy*.

```
1 model.evaluate(X_test, y_test, verbose=False)
```

[0.09586219489574432, 0.9736841917037964]





### Add early stopping

```
CPU times: user 3.75 s, sys: 530 ms, total: 4.28 s Wall time: 3.61 s Stopped after 68 epochs.
```

#### Evaluation on test set:

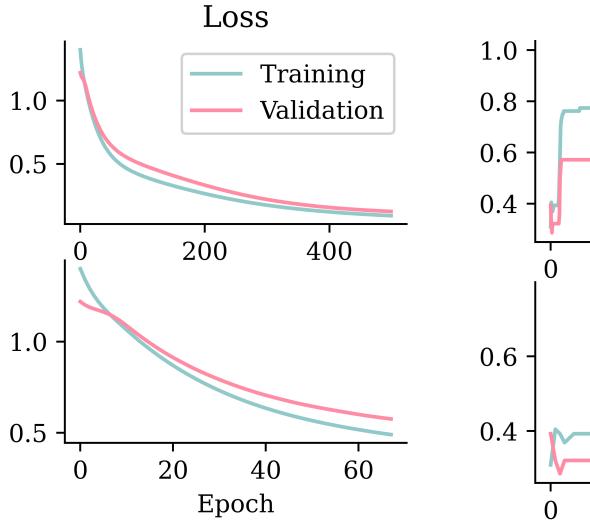
```
1 model.evaluate(X_test, y_test, verbose=False)
```

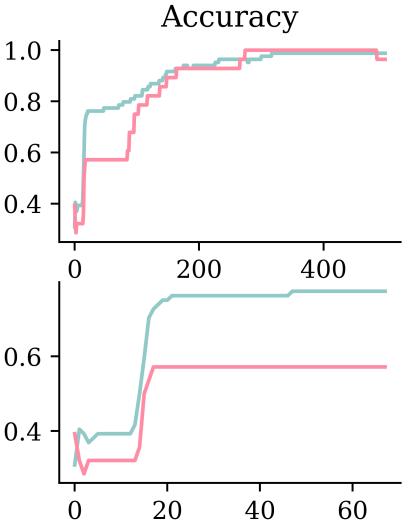
```
[0.9856260418891907, 0.5263158082962036]
```





# Fitting metrics









### What is the softmax activation?

It creates a "probability" vector: Softmax( $\boldsymbol{x}$ ) =  $\frac{\mathbf{e}_i^x}{\sum_j \mathbf{e}_j^x}$ .

#### In NumPy:

```
1 out = np.array([5, -1, 6])
2 (np.exp(out) / np.exp(out).sum()).round(3)
array([0.269, 0.001, 0.731])
```

#### In Keras:

```
1 out = keras.ops.convert_to_tensor([[5.0, -1.0, 6.0]])
2 keras.ops.round(keras.ops.softmax(out), 3)

<tf.Tensor: shape=(1, 3), dtype=float32, numpy=array([[0.269, 0.001, 0.731]], dtype=float32)>
```





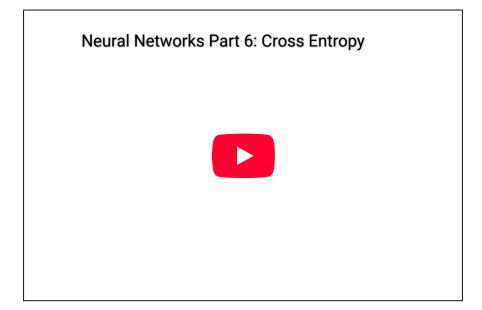
### Prediction using classifiers

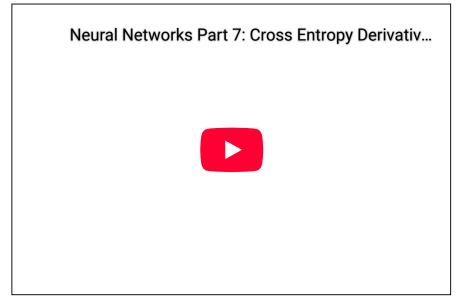
```
1 y_test[:4]
array([[2],
       [2],
       [1],
       [1]])
  1 y pred = model.predict(X test.head(4), verbose=0)
  2 y pred
array([[0.1397096 , 0.5175301 , 0.34276026],
       [0.24611066, 0.44371167, 0.31017768],
       [0.26309973, 0.43174297, 0.30515727],
       [0.259089 , 0.44883674, 0.29207426]], dtype=float32)
  1 # Add 'keepdims=True' to get a column vector.
  2 np.argmax(y_pred, axis=1)
array([1, 1, 1, 1])
  1 iris.target_names[np.argmax(y_pred, axis=1)]
array(['versicolor', 'versicolor', 'versicolor', 'versicolor'],
     dtype='<U10')
```





# Cross-entropy loss: ELI5



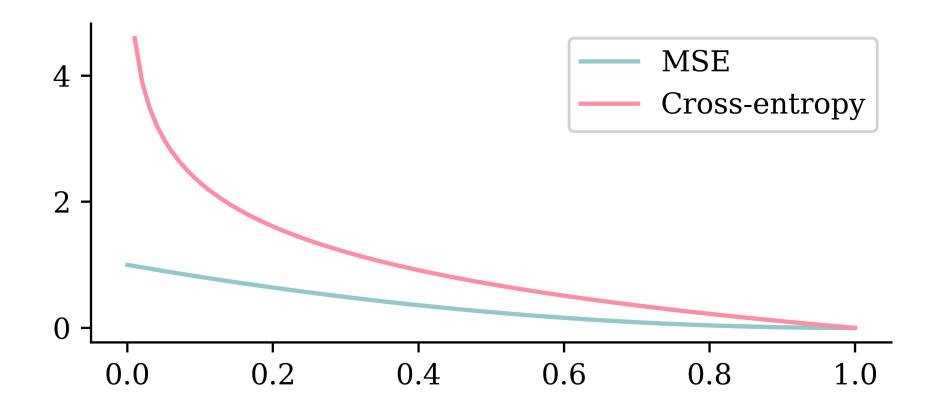






### Why use cross-entropy loss?

```
1  p = np.linspace(0, 1, 100)
2  plt.plot(p, (1 - p) ** 2)
3  plt.plot(p, -np.log(p))
4  plt.legend(["MSE", "Cross-entropy"]);
```







# One-hot encoding

```
from sklearn.preprocessing import OneHotEncoder

enc = OneHotEncoder(sparse_output=False)

y_train_oh = enc.fit_transform(y_train)
y_test_oh = enc.transform(y_test)
```

```
array([[1],
[1],
[1],
[0],
[0]])
```

1 y\_train[:5]

1 y\_train\_oh[:5]

	xo_o	XO_1	XO_2
0	0.0	1.0	0.0
1	0.0	1.0	0.0
2	0.0	1.0	0.0
3	1.0	0.0	0.0
4	1.0	0.0	0.0





### Classifier given one-hot outputs

Create the model (*new loss function*):

```
1 model = build_model()
2 model.compile("adam", "categorical_crossentropy", \
3 metrics=["accuracy"])
```

Fit the model (*new target variables*):

```
1 model.fit(X_train, y_train_oh, epochs=100, verbose=False);
```

Evaluate the model (*new target variables*):

```
1 model.evaluate(X_test, y_test_oh, verbose=False)
```

[0.3470938205718994, 0.9473684430122375]





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# The data

Dataset source: Kaggle Stroke Prediction Dataset.

```
1 data = pd.read_csv("stroke.csv")
2 data.head()
```

	id	gender	age	hypertension	heart_disease	ever_married	work_type	Residenc
0	9046	Male	67.0	0	1	Yes	Private	Urban
1	51676	Female	61.0	0	0	Yes	Self- employed	Rural
2	31112	Male	80.0	О	1	Yes	Private	Rural
3	60182	Female	49.0	О	0	Yes	Private	Urban
4	1665	Female	79.0	1	0	Yes	Self- employed	Rural





### Data description

- 1. id: unique identifier
- 2. gender: "Male", "Female" or "Other"
- 3. age: age of the patient
- 4. hypertension: o or 1 if the patient has hypertension
- 5. heart\_disease: o or 1 if the patient has any heart disease
- 6. ever\_married: "No" or "Yes"

- 8. Residence\_type: "Rural" or "Urban"
- 9. avg\_glucose\_level: average glucose level in blood
- 10. bmi: body mass index
- 11. smoking\_status: "formerly smoked", "never smoked", "smokes" or "Unknown"
- 12. stroke: o or 1 if the patient had a stroke





### Split the data

First, look for missing values.

```
1 number_missing = data.isna().sum()
  2 number missing[number missing > 0]
bmi
       201
dtype: int64
  1 features = data.drop(["id", "stroke"], axis=1)
  2 target = data["stroke"]
  3
  4 X_main, X_test, y_main, y_test = train_test_split(
         features, target, test_size=0.2, random_state=7)
  6 X_train, X_val, y_train, y_val = train_test_split(
         X_main, y_main, test_size=0.25, random_state=12)
  8
  9 X train.shape, X val.shape, X test.shape
((3066, 10), (1022, 10), (1022, 10))
```



### What values do we see in the data?

```
1 X_train["gender"].value_counts()
                                                    1 X_train["work_type"].value_counts()
gender
                                                 work_type
Female
                                                 Private
          1802
                                                                   1754
Male
          1264
                                                 Self-employed
                                                                   490
Name: count, dtype: int64
                                                 children
                                                                   419
                                                 Govt_job
                                                                    390
  1 X_train["ever_married"].value_counts()
                                                 Never worked
                                                                     13
                                                 Name: count, dtype: int64
ever married
                                                    1 X_train["smoking_status"].value_counts()
Yes
       2007
No
       1059
Name: count, dtype: int64
                                                 smoking_status
                                                 never smoked
                                                                     1130
  1 X_train["Residence_type"].value_counts()
                                                 Unknown
                                                                      944
                                                 formerly smoked
                                                                      522
Residence_type
                                                 smokes
                                                                     470
Urban
         1536
                                                 Name: count, dtype: int64
Rural
         1530
```



Name: count, dtype: int64



# Preprocess columns individually

- 1. Take categorical columns  $\hookrightarrow$  one-hot vectors
- 2. binary columns  $\hookrightarrow$  do nothing
- 3. continuous columns  $\hookrightarrow$  impute NaNs & standardise.





### Scikit-learn column transformer

```
from sklearn.pipeline import make pipeline
   cat_vars = ["gender", "ever_married", "Residence_type",
       "work type", "smoking status"]
 6 ct = make column transformer(
     (OneHotEncoder(sparse output=False, handle unknown="ignore"), cat vars),
     ("passthrough", ["hypertension", "heart disease"]),
     remainder=make pipeline(SimpleImputer(), StandardScaler()),
     verbose feature names out=False
10
11 )
12
13 X_train_ct = ct.fit_transform(X_train)
14 X_val_ct = ct.transform(X_val)
15 X test ct = ct.transform(X test)
16
   for name, X in zip(("train", "val", "test"), (X_train_ct, X_val_ct, X_test_ct)):
       num_na = X.isna().sum().sum()
18
       print(f"The {name} set has shape {X.shape} & with {num na} NAs.")
19
```

The train set has shape (3066, 20) & with 0 NAs. The val set has shape (1022, 20) & with 0 NAs. The test set has shape (1022, 20) & with 0 NAs.



### Handling unseen categories

```
1 X_train["gender"].value_counts()

gender
Female 1802
Male 1264
Name: count, dtype: int64
```

```
ind = np.argmax(X_val["gender"] = "Othe
X_val.iloc[ind-1:ind+3][["gender"]]
```

gender

	genuer
4970	Male
3116	Other
4140	Male
2505	Female

<pre>1 X_val["gender"].value_counts()</pre>
gender Female 615
Male 406 Other 1
Name: count, dtype: int64

1	gender_cols	= X_val_ct[["gender_Female",	
2	gender_cols	<pre>.iloc[ind-1:ind+3]</pre>	

	gender_Female	gender_
4970	0.0	1.0
3116	0.0	0.0
4140	0.0	1.0
2505	1.0	0.0





### Setup a binary classification model

```
def create_model(seed=42):
    random.seed(seed)
    model = Sequential()
    model.add(Input(X_train_ct.shape[1:]))
    model.add(Dense(32, "leaky_relu"))
    model.add(Dense(16, "leaky_relu"))
    model.add(Dense(1, "sigmoid"))
    return model
model = create_model()
model.summary()
```

#### Model: "sequential 5"

Layer (type)	Output Shape	Param #
dense_10 (Dense)	(None, 32)	672
dense_11 (Dense)	(None, 16)	528
dense_12 (Dense)	(None, 1)	17

**Total params:** 1,217 (4.75 KB) Trainable params: 1,217 (4.75 KB) Non-trainable params: 0 (0.00 B)







### Add metrics, compile, and fit

```
model = create_model()
     pr auc = keras.metrics.AUC(curve="PR", name="pr auc")
    model.compile(optimizer="adam", loss="binary_crossentropy",
         metrics=[pr_auc, "accuracy", "auc"])
     es = EarlyStopping(patience=50, restore best weights=True,
         monitor="val_pr_auc", verbose=1)
     model.fit(X_train_ct, y_train, callbacks=[es], epochs=1_000, verbose=0,
       validation data=(X val ct, y val));
Epoch 65: early stopping
Restoring model weights from the end of the best epoch: 15.
  1 model.evaluate(X val ct, y val, verbose=
[0.14444079995155334,
0.13122102618217468,
0.9589040875434875,
0.8215014934539795]
```





### Overweight the minority class

```
model = create_model()

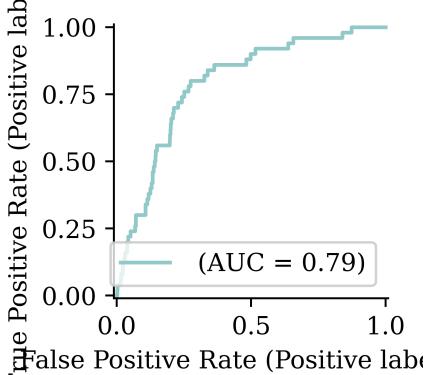
pr_auc = keras.metrics.AUC(curve="PR", name="pr_auc")
model.compile(optimizer="adam", loss="binary_crossentropy",
metrics=[pr_auc, "accuracy", "auc"])

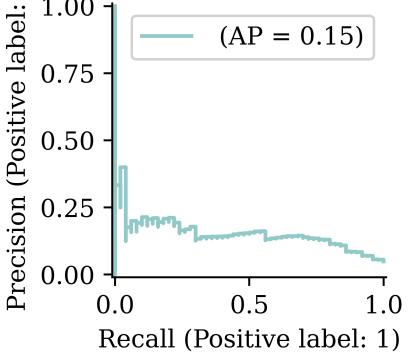
es = EarlyStopping(patience=50, restore_best_weights=True,
monitor="val_pr_auc", verbose=1)
model.fit(X_train_ct, y_train.to_numpy(), callbacks=[es], epochs=1_000, verbose=0,
validation_data=(X_val_ct, y_val), class_weight={0: 1, 1: 10});
```

Epoch 74: early stopping
Restoring model weights from the end of the best epoch: 24.



### Classification Metrics





```
1 y_pred_stroke = y_pred > 0.5
2 confusion_matrix(y_test, y_pred_stroke)
```

array([[792, 180],

1 y\_pred\_stroke = y\_pred > 0.3
2 confusion\_matrix(y\_test, y\_pred\_stroke)

array([[662, 310],



# Package Versions

1 from watermark import watermark
2 print(watermark(python=True, packages="keras,matplotlib,numpy,pandas,seaborn,scipy,torch

Python implementation: CPython Python version : 3.11.11 IPython version : 8.32.0

keras : 3.8.0
matplotlib: 3.10.0
numpy : 1.26.4
pandas : 2.2.3
seaborn : 0.13.2
scipy : 1.13.1

torch : 2.5.1+cu124

tensorflow: 2.18.0 tf\_keras : 2.18.0





# Glossary

- accuracy
- classification problem
- confusion matrix
- cross-entropy loss
- metrics
- sigmoid activation function
- sofmax activation



