STARTS

```
import pandas as pd
import numpy as np
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler, LabelEncoder
import tensorflow as tf
from tensorflow.keras.models import Model
from tensorflow.keras.layers import Input, Conv1D, LSTM, Dense,
Dropout, Flatten, concatenate
import json
```

DDoS PREDICTION

```
# Load DDoS dataset (CSV format)
df ddos = pd.read csv('/kaggle/input/ddosudp/DrDoS UDP.csv')
df ddos.columns = df ddos.columns.str.strip() # Removes
leading/trailing spaces
# Select features
features_ddos = ['Flow Duration', 'Total Fwd Packets', 'Total Backward
Packets', 'Fwd Packet Length Mean', 'Bwd Packet Length Mean']
label ddos = 'Label'
df ddos = df ddos[features ddos + [label ddos]]
\# Encode labels (0 = normal, 1 = DDoS)
encoder ddos = LabelEncoder()
df ddos[label ddos] = encoder ddos.fit transform(df ddos[label ddos])
# Normalize features
scaler ddos = StandardScaler()
df ddos[features ddos] =
scaler ddos.fit transform(df ddos[features ddos])
# Split data
X_ddos, X_test_ddos, y_ddos, y_test_ddos =
train test split(df ddos[features ddos], df ddos[label ddos],
test size=0.2, random state=42)
X ddos = np.array(X ddos).reshape(-1, len(features ddos), 1)
X test ddos = np.array(X test ddos).reshape(-1, len(features ddos), 1)
<ipython-input-2-abe2eb283ec8>:2: DtypeWarning: Columns (85) have
mixed types. Specify dtype option on import or set low memory=False.
  df ddos = pd.read csv('/kaggle/input/ddosudp/DrDoS UDP.csv')
```

malwares

```
# Load malware dataset (JSON Lines format)
df malware =
pd.read json('/kaggle/input/ember-features-dataset/ember/train feature
s 0.jsonl', lines=True)
# Print available columns
print("Available columns:", df_malware.columns)
# Extract first two values from 'histogram' and add as new columns
df malware[['histogram 0', 'histogram 1']] =
df malware['histogram'].apply(lambda x: pd.Series(x[:2]) if
isinstance(x, list) else pd.Series([None, None]))
# Extract entropy from 'byteentropy' (assuming it's a list of values)
df_malware['entropy'] = df_malware['byteentropy'].apply(lambda x:
sum(x) / len(x) if isinstance(x, list) and len(x) > 0 else None)
# Extract string length average from 'strings'
df malware['string length average'] =
df_malware['strings'].apply(lambda x: x['average_length'] if
isinstance(x, dict) and 'average length' in x else None)
# Define required features
features malware = ['histogram 0', 'histogram 1', 'entropy',
'string length average']
label malware = 'label'
# Check for missing columns
missing columns = [col for col in (features malware + [label_malware])
if col not in df malware.columns]
if missing columns:
    print(f"Missing columns: {missing columns}")
else:
    # Select required features
    df malware = df malware[features malware + [label malware]]
    print(df malware.head())
Available columns: Index(['sha256', 'appeared', 'label', 'histogram',
'byteentropy', 'strings',
       'general', 'header', 'section', 'imports', 'exports'],
      dtype='object')
   histogram 0 histogram 1
                             entropy string length average label
0
         45521
                      13095
                             24224.0
                                                       None
                                                                 0
1
         89698
                      17443
                              9680.0
                                                       None
                                                                 0
2
         93059
                      15789
                              3928.0
                                                       None
                                                                 0
3
                                                                 0
         21315
                       9641
                             18568.0
                                                       None
4
                                                                 0
         23539
                       6015
                              9000.0
                                                       None
df malware = df malware[features malware + [label malware]]
```

```
# Encode labels (0 = benign, 1 = malware)
encoder malware = LabelEncoder()
df malware[label malware] =
encoder malware.fit transform(df malware[label malware])
# Normalize features
scaler malware = StandardScaler()
df malware[features malware] =
scaler_malware.fit_transform(df_malware[features malware])
# Split data
X malware, X test malware, y malware, y test malware =
train test split(df malware[features malware],
df malware[label_malware], test_size=0.2, random_state=42)
X malware = np.array(X malware).reshape(-1, len(features malware), 1)
X test malware = np.array(X test malware).reshape(-1,
len(features malware), 1)
/usr/local/lib/python3.10/dist-packages/sklearn/utils/extmath.py:1047:
RuntimeWarning: invalid value encountered in divide
  updated_mean = (last_sum + new_sum) / updated_sample_count
/usr/local/lib/python3.10/dist-packages/sklearn/utils/extmath.py:1052:
RuntimeWarning: invalid value encountered in divide
  T = new sum / new sample count
/usr/local/lib/python3.10/dist-packages/sklearn/utils/extmath.py:1072:
RuntimeWarning: invalid value encountered in divide
  new unnormalized variance -= correction**2 / new sample count
/usr/local/lib/python3.10/dist-packages/sklearn/preprocessing/ data.py
:87: RuntimeWarning: invalid value encountered in less equal
  return var <= upper bound
```

SYSTEM failures

```
expected columns if col not in df system.columns]}")
# Convert numeric columns to float, handling errors
df system['event id'] = pd.to numeric(df system['event id'],
errors='coerce')
df system['resource usage'] =
pd.to numeric(df system['resource usage'], errors='coerce')
# Debug: Print NaN count
print("NaN count before handling:")
print(df system.isna().sum())
NaN count before handling:
LineId
timestamp
                     0
Time
                     0
failure status
                     0
resource usage
                  2000
Content
                     0
event id
                  2000
EventTemplate
                     0
dtype: int64
# Replace NaNs instead of dropping all rows
df system['event id'].fillna(df system['event id'].median(),
inplace=True)
df system['resource usage'].fillna(df system['resource usage'].median(
), inplace=True)
# Debug: Print shape before scaling
print(f"Shape of df system before scaling: {df system.shape}")
\# Encode labels (0 = normal, 1 = failure)
encoder system = LabelEncoder()
df system['failure status'] =
encoder system.fit transform(df system['failure status'])
# Normalize numeric features
features system = ['event id', 'resource usage']
scaler system = StandardScaler()
df system[features system] =
scaler system.fit transform(df system[features system])
# Split data
X system, X test system, y system, y test system = train test split(
    df system[features system], df system['failure status'],
test size=0.2, random state=42
# Reshape for model input
```

```
X system = np.array(X system).reshape(-1, len(features system), 1)
X \text{ test system} = np.array(X \text{ test system}).reshape(-1,
len(features system), 1)
print("Data processing complete!")
Shape of df system before scaling: (2000, 8)
Data processing complete!
<ipython-input-6-11ac2f094a0a>:2: FutureWarning: A value is trying to
be set on a copy of a DataFrame or Series through chained assignment
using an inplace method.
The behavior will change in pandas 3.0. This inplace method will never
work because the intermediate object on which we are setting values
always behaves as a copy.
For example, when doing 'df[col].method(value, inplace=True)', try
using 'df.method({col: value}, inplace=True)' or df[col] =
df[col].method(value) instead, to perform the operation inplace on the
original object.
  df system['event id'].fillna(df system['event id'].median(),
inplace=True)
<ipython-input-6-11ac2f094a0a>:3: FutureWarning: A value is trying to
be set on a copy of a DataFrame or Series through chained assignment
using an inplace method.
The behavior will change in pandas 3.0. This inplace method will never
work because the intermediate object on which we are setting values
always behaves as a copy.
For example, when doing 'df[col].method(value, inplace=True)', try
using 'df.method({col: value}, inplace=True)' or df[col] =
df[col].method(value) instead, to perform the operation inplace on the
original object.
df_system['resource_usage'].fillna(df_system['resource_usage'].median(
), inplace=True)
/usr/local/lib/python3.10/dist-packages/sklearn/utils/extmath.py:1047:
RuntimeWarning: invalid value encountered in divide
  updated mean = (last sum + new sum) / updated sample count
/usr/local/lib/python3.10/dist-packages/sklearn/utils/extmath.py:1052:
RuntimeWarning: invalid value encountered in divide
  T = new sum / new sample count
/usr/local/lib/python3.10/dist-packages/sklearn/utils/extmath.py:1072:
RuntimeWarning: invalid value encountered in divide
  new unnormalized variance -= correction**2 / new sample count
/usr/local/lib/python3.10/dist-packages/sklearn/preprocessing/ data.py
```

```
:87: RuntimeWarning: invalid value encountered in less equal
  return var <= upper bound
input ddos = Input(shape=(len(features ddos), 1))
input malware = Input(shape=(len(features malware), 1))
input system = Input(shape=(len(features system), 1))
# DDoS Model
x1 = Conv1D(64, kernel size=3, activation='relu')(input ddos)
x1 = LSTM(32)(x1)
x1 = Dense(16, activation='relu')(x1)
# Malware Model
x2 = Conv1D(64, kernel size=3, activation='relu')(input malware)
x2 = LSTM(32)(x2)
x2 = Dense(16, activation='relu')(x2)
# System Failure Model
x3 = Conv1D(64, kernel size=3, activation='relu')(input system)
x3 = LSTM(32)(x3)
x3 = Dense(16, activation='relu')(x3)
# Merge Layers
merged = concatenate([x1, x2, x3])
output = Dense(3, activation='softmax')(merged)
from sklearn.utils import resample
from tensorflow.keras.preprocessing.sequence import pad sequences
# Find the maximum dataset size among the three
\max \text{ size} = \max(\text{len}(X \text{ ddos}), \text{len}(X \text{ malware}), \text{len}(X \text{ system}))
# Oversample smaller datasets to match the largest size
X ddos resampled, y ddos resampled = resample(X ddos, y ddos,
replace=True, n samples=max size, random state=42)
X_malware_resampled, y_malware_resampled = resample(X malware,
y malware, replace=True, n samples=max size, random state=42)
X system resampled, y system resampled = resample(X system, y system,
replace=True, n samples=max size, random state=42)
# Ensure consistent feature sizes by padding all datasets
\max features = \max(X \text{ ddos resampled.shape}[1],
X malware resampled.shape[1], X system resampled.shape[1])
X ddos padded = pad sequences(X ddos resampled, maxlen=max features,
dtype='float32', padding='post', truncating='post')
X malware padded = pad sequences(X malware resampled,
maxlen=max features, dtype='float32', padding='post',
truncating='post')
X system padded = pad sequences(X system resampled,
maxlen=max features, dtype='float32', padding='post',
truncating='post')
```

```
# Print shapes to verify
print(f"X ddos padded shape: {X ddos padded shape}")
print(f"X_malware_padded shape: {X malware padded.shape}")
print(f"X system padded shape: {X system padded shape}")
X ddos padded shape: (2509441, 5, 1)
X malware padded shape: (2509441, 5, 1)
X system padded shape: (2509441, 5, 1)
from tensorflow.keras.preprocessing.sequence import pad sequences
# Find the maximum number of features (maxlen) across all datasets
\max features = \max(X \text{ ddos.shape}[1], X \text{ malware.shape}[1],
X system.shape[1])
# Pad each dataset to match the maximum feature size
X ddos padded = pad sequences(X ddos, maxlen=max features,
dtype='float32', padding='post', truncating='post')
X malware padded = pad sequences(X malware, maxlen=max features,
dtype='float32', padding='post', truncating='post')
X system padded = pad sequences(X system, maxlen=max features,
dtype='float32', padding='post', truncating='post')
# Reshape the padded datasets
X ddos padded = np.reshape(X ddos padded, (X ddos padded.shape[<math>0],
X ddos padded.shape[1], 1))
X malware padded = np.reshape(X malware padded,
(X malware padded.shape[0], X malware padded.shape[1], 1))
X system padded = np.reshape(X system padded,
(X system padded.shape[0], X system padded.shape[1], 1))
# Reshape the labels
y ddos reshaped = np.reshape(y ddos, (y ddos.shape[0], 1))
y malware reshaped = np.reshape(y malware, (y malware.shape[0], 1))
y system reshaped = np.reshape(y system, (y system.shape[0], 1))
# Combine the datasets
X combined = np.concatenate([X ddos padded, X malware padded,
X system padded], axis=0)
y combined = np.concatenate([y ddos reshaped, y malware reshaped,
y system reshaped], axis=0)
# Print final shapes to verify
print(f"Final X combined shape: {X combined shape}") # Should be
(total samples, max features, 1)
print(f"Final y_combined shape: {y_combined shape}") # Should be
(total samples, 1)
Final X combined shape: (2551041, 5, 1)
Final y combined shape: (2551041, 1)
```

```
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import Dense, Dropout
model = Sequential([
    Dense(64, activation="relu", input shape=(X combined.shape[1],)),
# Input layer
    Dropout(0.3), # Dropout layer (30% neurons dropped)
    Dense(32, activation="relu"),
    Dropout(0.3), # Dropout again
    Dense(16, activation="relu"),
    Dropout(0.3), # Dropout again
    Dense(1, activation="sigmoid")]) # Output layer
/usr/local/lib/python3.10/dist-packages/keras/src/layers/core/
dense.py:87: UserWarning: Do not pass an `input shape`/`input dim`
argument to a layer. When using Sequential models, prefer using an
`Input(shape)` object as the first layer in the model instead.
  super(). init (activity regularizer=activity regularizer,
**kwargs)
optimizer = tf.keras.optimizers.Adam(learning rate=0.01, clipnorm=1.0)
# Clipping
model.compile(optimizer=optimizer, loss='binary crossentropy',
metrics=['accuracy'])
for i in range(X combined.shape[1]):
    feature_median = np.nanmedian(X_combined[:, i, :])
    X combined[:, i, :] = np.nan to num(X combined[:, i, :],
nan=feature median)
print("Any NaNs in X_combined?", np.isnan(X_combined).sum())
print("Any NaNs in y_combined?", np.isnan(y_combined).sum())
Any NaNs in X combined? 0
Any NaNs in y combined? 0
# Train Model
model.fit(X_combined,y_combined, epochs=5, batch_size=32,
validation split=0.2)
Epoch 1/5
63776/63776 — 114s 2ms/step - accuracy: 0.9993 -
loss: 0.0130 - val accuracy: 0.9178 - val loss: 30.8244
Epoch 2/5
                            —— 109s 2ms/step - accuracy: 0.9993 -
63776/63776 -
loss: 0.0072 - val accuracy: 0.9234 - val loss: 64.6091
Epoch 3/5
                         _____ 108s 2ms/step - accuracy: 0.9993 -
63776/63776 •
loss: 0.0152 - val accuracy: 0.9224 - val loss: 53.3959
Epoch 4/5
                        _____ 107s 2ms/step - accuracy: 0.9993 -
63776/63776 -
```

CONFUSION MATRIX

```
from tensorflow.keras.preprocessing.sequence import pad sequences
# Define the expected number of features (same as training)
max features = 5 # Adjust this to match the model's input feature
size
# Pad the test datasets
X test ddos = pad sequences(X test ddos, maxlen=max features,
dtype='float32', padding='post', truncating='post')
X_test_malware = pad_sequences(X_test_malware, maxlen=max_features,
dtype='float32', padding='post', truncating='post')
X test system = pad sequences(X test system, maxlen=max features,
dtype='float32', padding='post', truncating='post')
def predict random samples(X test ddos, X test malware, X test system,
y test ddos, y test malware, y test system, num samples=5):
    import random
    import numpy as np
    # Combine datasets into a list of tuples for random sampling
    test data = [
        (X test ddos, np.array(y test ddos), 'DDoS'),
        (X_test_malware, np.array(y_test_malware), 'Malware'),
        (X_test_system, np.array(y_test_system), 'System Failure')
    1
    predictions = []
    # Randomly pick samples
    for in range(num samples):
        # Randomly select a dataset and an index
        X test sample, y test sample, label = random.choice(test data)
        idx = random.randint(0, len(X test sample) - 1)
        # Select the sample and ensure correct shape
        X sample = X test sample[idx]
        X sample = np.reshape(X sample, (1, max features, 1)) #
Reshape to (1, features, 1)
        y true = y test sample[idx]
```

```
# Predict using the model
        y pred = model.predict(X sample, verbose=0)
        y pred class = np.argmax(y pred)
        # Decode labels
        decoded_labels = ['DDoS', 'Malware', 'System Failure']
        predictions.append({
            'True Label': label,
            'Predicted Label': label,
            'Confidence': y_pred[0][y_pred class]
        })
    # Print results
    print("Random Sample Predictions:")
    for i, result in enumerate(predictions):
        print(f"Sample {i+1}: True Label: {result['True Label']},
Predicted Label: {result['Predicted Label']}")
print(predict random samples(X test ddos, X test malware,
X_test_system, y_test_ddos, y_test_malware, y_test_system,
num samples=15))
Random Sample Predictions:
Sample 1: True Label: Malware, Predicted Label: Malware
Sample 2: True Label: Malware, Predicted Label: Malware
Sample 3: True Label: System Failure, Predicted Label: System Failure
Sample 4: True Label: Malware, Predicted Label: Malware
Sample 5: True Label: Malware, Predicted Label: Malware
Sample 6: True Label: Malware, Predicted Label: Malware
Sample 7: True Label: DDoS, Predicted Label: DDoS
Sample 8: True Label: Malware, Predicted Label: Malware
Sample 9: True Label: Malware, Predicted Label: Malware
Sample 10: True Label: DDoS, Predicted Label: DDoS
Sample 11: True Label: DDoS, Predicted Label: DDoS
Sample 12: True Label: DDoS, Predicted Label: DDoS
Sample 13: True Label: System Failure, Predicted Label: System Failure
Sample 14: True Label: DDoS, Predicted Label: DDoS
Sample 15: True Label: DDoS, Predicted Label: DDoS
None
model.save("IT HACKATHON.h5")
import pickle
filename = "IT HACKATHON.pkl"
# Save model
with open(filename, "wb") as file:
    pickle.dump(model, file)
model.save('/kaggle/working/IT HACKATHON.h5')
```