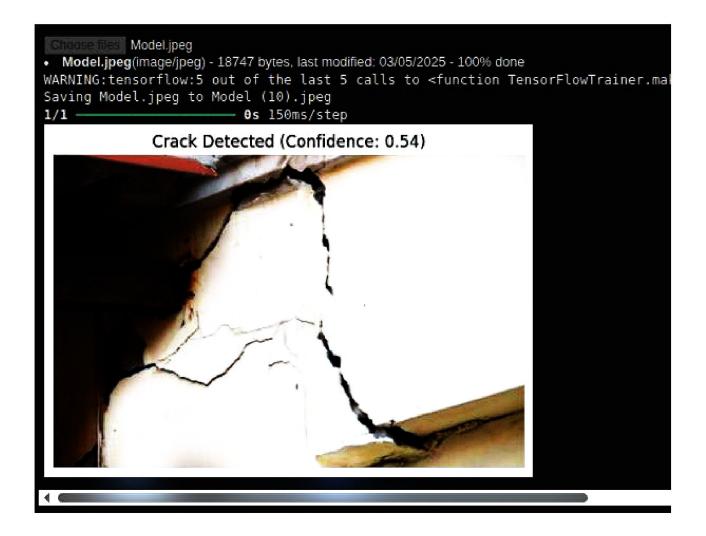
```
[4] !pip install -q tensorflow opency-python matplotlib
[6] import cv2
    import numpy as np
    import tensorflow as tf
    import matplotlib.pyplot as plt
    from tensorflow keras models import load model
    from tensorflow.keras.preprocessing.image import img_to_array
    from google.colab import files
Double-click (or enter) to edit
0
    import numpy as np
    import cv2
    import tensorflow as tf
    from PIL import Image
    from qoogle.colab import files
    import matplotlib.pyplot as plt
    def create simple crack model(input shape=(224, 224, 3)):
        model = tf.keras.Sequential([
            tf.keras.layers.Conv2D(32, (3, 3), activation='relu', input_shape=input_shape),
            tf.keras.layers.MaxPooling2D(2, 2),
            tf.keras.layers.Conv2D(64, (3, 3), activation='relu'),
            tf.keras.layers.MaxPooling2D(2, 2),
            tf.keras.layers.Conv2D(128, (3, 3), activation='relu'),
            tf.keras.layers.MaxPooling2D(2, 2),
            tf.keras.layers.Flatten(),
            tf.keras.layers.Dense(128, activation='relu'),
            tf.keras.layers.Dropout(0.5),
            tf.keras.layers.Dense(1, activation='sigmoid') # Binary classification
        model.compile(optimizer='adam',
                      loss='binary crossentropy',
                      metrics=['accuracy'])
        return model
    model = create simple crack model()
```

```
model.compile(optimizer='adam',
0
                      loss='binary crossentropy',
                      metrics=['accuracy'])
        return model
    model = create simple crack model()
    uploaded = files.upload()
    def detect crack(img array, model):
        img = cv2.resize(img array, (224, 224))
        img = img.astype('float32') / 255.0
        img = np.expand dims(img, axis=0)
        prediction = model.predict(img)[0][0]
        label = "Crack Detected" if prediction > θ.1 else "No Crack Detected"
        return label, prediction
    for fname in uploaded.keys():
        image data = uploaded[fname]
        pil imq = Image.open(io.BytesIO(image data)).convert('RGB')
        img array = np.array(pil img)
        imq bgr = cv2.cvtColor(img array, cv2.COLOR RGB2BGR)
        label, conf = detect crack(img bgr, model)
        plt.imshow(pil img)
        plt.title(f'{label} (Confidence: {conf:.2f})')
        plt.axis('off')
        plt.show()
```

## output:



```
import pandas as pd
      import numpy as np
      import matplotlib.pyplot as plt
      import seaborn as sns
      from sklearn.ensemble import IsolationForest
      from sklearn.preprocessing import StandardScaler
      import os
      from datetime import datetime
      # ----- Setup Paths -----
      BASE DIR = "/content"
      RAW_DIR = os.path.join(BASE_DIR, "raw")
      os.makedirs(RAW_DIR, exist_ok=True)
      CSV_PATH = os.path.join(RAW_DIR, "shm_data.csv")
      CHUNK SIZE = 500
      MAX CHUNKS = 2 # Limit for concise output
      # ----- Generate Synthetic Data -----
      def generate_data(filepath, rows=1000):
          timestamps = pd.date range(start="2023-01-01", periods=rows, freq="1min")
          df = pd.DataFrame({
               "timestamp": timestamps,
               "strain": np.random.normal(50, 10, rows),
               "vibration": np.random.normal(0.3, 0.1, rows),
               "displacement": np.random.normal(5, 2, rows),
               "temperature": np.random.normal(30, 3, rows),
          })
          anomalies = np.random.choice(rows, size=20, replace=False)
          df.loc[anomalies, 'strain'] += np.random.normal(80, 15, len(anomalies))
df.loc[anomalies, 'vibration'] += np.random.normal(1.5, 0.3, len(anomalies))
          df.to_csv(filepath, index=False)
          print(f"Data saved at {filepath}")
# ----- Read Data in Chunks -----
def read_chunks(path, chunk_size):
    return pd.read_csv(path, chunksize=chunk_size, parse_dates=["timestamp"])
# ----- Detect Anomalies -----
def detect_anomalies(df):
    features = ["strain", "vibration", "displacement", "temperature"]
    scaler = StandardScaler()
    X_scaled = scaler.fit_transform(df[features])
    model = IsolationForest(contamination=0.02, random_state=42)
    df['anomaly'] = model.fit_predict(X_scaled)
    return df[df['anomaly'] ==
# ----- Plot One Graph Only ------
def plot_anomalies(df, anomalies, chunk_num):
    plt.figure(figsize=(12, 5))
    sns.lineplot(x='timestamp', y='strain', data=df, label='Strain')
    if not anomalies.empty:
        sns.scatterplot(x='timestamp', y='strain', data=anomalies, color='red', label='Anomaly')
    plt.title(f"Chunk {chunk num} - Strain with Anomalies")
    plt.xticks(rotation=45)
    plt.tight_layout()
    plt.show()
# ----- Main SHM Function ------
def monitor_shm():
    if not os.path.exists(CSV_PATH):
       generate_data(CSV_PATH)
    for i, chunk in enumerate(read chunks(CSV PATH, CHUNK SIZE)):
       if i >= MAX_CHUNKS:
           break
        anomalies = detect_anomalies(chunk)
       print(f"Chunk {i+1}: {len(anomalies)}) anomalies detected")
       plot_anomalies(chunk, anomalies, i+1)
monitor shm()
```

## **Output:**

