```
[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

import numpy as np
    import cv2
```

```
import cv2
import io
import tensorflow as tf
from PIL import Image
from google.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
    1)
    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
   # Initialize the model
   model = create simple crack model()
   # Assuming the model is already trained, or you can train it here.
    # If you have a pre-trained model saved, you can load it using:
   # model = tf.keras.models.load model('path to model')
   # Upload images
   uploaded = files.upload()
   # Define the detection function
    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 > 0.1 else "No Crack Detected"
        return label, prediction
    # Process uploaded images
    for fname in uploaded.keys():
        # Read using PIL to support all formats, then convert to OpenCV format
        image data = uploaded[fname]
        pil img = Image.open(io.BytesIO(image data)).convert('RGB')
        img array = np.array(pil img)
        img bgr = cv2.cvtColor(img array, cv2.COLOR RGB2BGR)
        # Get prediction and confidence
        label, conf = detect crack(img bgr, model)
        # Display result
        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="lmin")
    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'] == -1]
# ----- 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:



