Research report



BCSE306L - Artificial Intelligence

DIGITAL ASSIGNMENT 2

By: Yana Jain & Tushar Kumar

Reg: 23BAI1292 & 23BAI1121

BreakHis Binary Classification

Basic Swin
Transformer Model

Completed on Mar 8, 2025

Prepared by
Yana Jain
Tushar Kumar

Key insights

- Dataset Processing
- Evaluation of Model
- Comparing with CNN model

Introduction

Swin Transformer (Shifted Window Transformer)

The **Swin Transformer** is a cutting-edge vision transformer developed to enhance image analysis while addressing the computational limitations of traditional **Vision Transformers** (ViT). Unlike ViT, which applies self-attention across the entire image, Swin Transformer introduces a **localized window-based attention mechanism** and a **hierarchical structure**, making it more scalable and effective for processing high-resolution images.

Key Features of Swin Transformer

Hierarchical Feature Learning:

Unlike ViT, which processes images using fixed-size patches, Swin Transformer employs a **multi-scale hierarchical approach**, much like CNNs. This enables it to capture both **fine details** and **broad contextual information**, enhancing its performance in complex image classification tasks.

• Shifted Window Attention:

Traditional transformers compute self-attention across the entire image, leading to high computational costs. Swin Transformer divides images into small, non-overlapping windows, applying self-attention within each. To facilitate information exchange between windows, they are shifted between layers, allowing the model to capture long-range dependencies while maintaining efficiency.

• Improved Computational Efficiency:

By restricting self-attention to **localized windows**, Swin Transformer significantly reduces computational complexity compared to standard transformers. This makes it particularly well-suited for high-resolution images, as it processes smaller regions first before expanding to extract **global features**.

Scalability and Adaptability:

Due to its layered architecture, Swin Transformer can be applied to a wide range of vision tasks, including **image classification**, **object detection**, **and medical imaging**. Its ability to adapt to **different patch sizes and window configurations** makes it more flexible than conventional transformers.

By combining localized attention with hierarchical global feature learning, the Swin Transformer achieves a balance between accuracy and computational efficiency, making it a strong alternative to CNNs and traditional ViTs for vision-related applications.

Both breast tumors benign and malignant can be sorted into different types based on the way the tumoral cells look under the microscope.

Dataset Processing

Why is processing required?

- To ensure compatibility with the model
- To improve model performance
- To handle imbalanced data
- To speed up training

SOURCE CODE

1. sortimages.pu

import os

```
import shutil
import random

# Define source and destination paths
SOURCE_PATH = "/home/yanajain/Downloads/BreaKHis_v1
(2)/histology_slides/breast"
DEST_PATH = "/home/yanajain/Downloads/BreaKHis_Processed"

# Train-validation split ratio
TRAIN_SPLIT = 0.8

# Magnification levels
```

```
MAGNIFICATIONS = ["40X", "100X", "200X", "400X"]
# Categories in dataset
CATEGORIES = ["benign", "malignant"]
# Create train/val directories
for split in ["train", "val"]:
       for category in CATEGORIES:
       for mag in MAGNIFICATIONS:
       os.makedirs(os.path.join(DEST_PATH, split, category, mag),
exist ok=True)
# Function to find images recursively
def get all images(base path, mag level):
       """Finds all images in subdirectories containing a specific
magnification level."""
       image paths = []
       for root, dirs, files in os.walk(base path):
       if mag level in root: # Check if magnification level is in the path
       for file in files:
              if file.endswith((".png", ".jpg", ".jpeg")):
              image paths.append(os.path.join(root, file))
       return image paths
# Process each category (benign/malignant)
for category in CATEGORIES:
       category path = os.path.join(SOURCE PATH, category)
       # Iterate through magnification levels
       for mag in MAGNIFICATIONS:
       images = get all images(category path, mag)
       # Shuffle images for randomness
       random.shuffle(images)
       # Split into train and validation sets
       split index = int(len(images) * TRAIN SPLIT)
       train images = images[:split index]
       val images = images[split index:]
       # Move images to respective directories
       for img path in train images:
       shutil.copy(img_path, os.path.join(DEST_PATH, "train", category,
mag))
```

```
for img_path in val_images:
    shutil.copy(img_path, os.path.join(DEST_PATH, "val", category,
mag))

print("Dataset successfully reorganized!")
```

Here we set 80% dataset for training and 20% dataset for validation

Before processing the data make sure you remember to set up your environment to run the python code.

Steps to execute

- Step 1: Install python 3.11, since 3.12 does not support Tensorflow
- Step 2: In your ubuntu terminal, create a virtual environment and install the required libraries.
- Step 3: In a python script (sortimages.py), organize them according to different magnification levels.
- Step 4: Run for different magnification levels to find the validation accuracy.
- Step 5: As mentioned, consider various window and patch sizes to process the data.

Data augmentation is a technique that artificially increases the amount of data used to train deep learning models.

yanajain@yanajain-IdeaPad-3-15IIL05:~\$ python3.11 --version Python 3.11.11

Commands to install python 3.11

sudo apt update && sudo apt upgrade sudo apt install software-properties-common sudo add-apt-repository ppa:deadsnakes/ppa sudo apt update sudo apt install python3.11

```
yanajain@yanajain-IdeaPad-3-15IIL05:-5 python3.11 -version
Python 3.11.11
Python3.11.11
Python3.11
Python3.11.11
Python3.11
Pytho
```

TIMM (Torch Image Models) is an advanced PyTorch library that provides a wide range of pre-trained deep learning models for image classification, object detection, and segmentation. It includes optimized versions of architectures like Swin Transformer, ViTs, and CNNs, facilitating efficient training and fine-tuning on personalized datasets. TIMM is built for versatility, scalability, and high-performance computing, making it well-suited for applications such as breast cancer classification, where accuracy and computational efficiency are paramount. Its integrated optimizations streamline model loading, preprocessing, and evaluation, enabling easy comparison of various architectures within a consolidated framework.

Evaluating the Model

Impact of Patch and Window Sizes in Swin Transformer

1. Model Performance (Accuracy, Loss)

- Smaller patch sizes (32) tend to retain more fine-grained information, leading to better feature extraction and classification accuracy.
- Larger patch sizes (64) reduce spatial resolution, which might lead to lower accuracy for detailed tasks like medical image classification.
- Larger window sizes (128) allow the model to capture more global context but increase computation time.
- Smaller window sizes (64) focus more on local features and might miss long-range dependencies.

2. Computational Efficiency (Memory, Speed)

- Patch Size 32 → Higher memory usage & slower inference due to more patches
- Patch Size 64 → Lower memory usage & faster inference but might lose fine details
- Window Size 64 → Faster computation with more localized feature extraction
- Window Size 128 → More expensive but better global representation

Tuning patch and window sizes during model evaluation directly influences efficiency, accuracy, and feature extraction quality. Swin Transformer achieves a balance between computational cost and accuracy, making it highly effective for breast cancer histopathology analysis.

We applied patch sizes of 32, 64 and window sizes of 64, 128 on all different magnification levels (40x, 100x, 200x, 400x).

SOURCE CODE

swin.pu

import os

import tensorflow as tf

def create model(input shape):

```
model = tf.keras.models.Sequential([
       tf.keras.Input(shape=input shape),
       tf.keras.layers.Conv2D(32, (3, 3), activation='relu'),
       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(512, activation='relu'),
       tf.keras.layers.Dense(1, activation='sigmoid')
       1)
       model.compile(optimizer='adam',
               loss='binary crossentropy',
               metrics=['accuracy'])
       return model
# Helper function to create generators
def create generators(magnification, train or val, target size):
       benign dir = os.path.join(base dir, train or val, 'benign', magnification)
       malignant dir = os.path.join(base dir, train or val, 'malignant', magnification)
       datagen = ImageDataGenerator(rescale=1./255,
                      rotation range=20,
                      width shift range=0.2,
                      height shift range=0.2,
                      shear range=0.2,
                      zoom range=0.2,
                      horizontal flip=True if train or val == 'train' else False)
       combined dir = os.path.join(base dir, train or val, magnification)
       os.makedirs(combined dir, exist ok=True)
       generator = datagen.flow from directory(combined dir,
                              target size=target size,
                              batch size=32,
                              class mode='binary')
       return generator
# Copy images to a combined directory structure
def copy images to combined dir(magnification):
       for train or val in ['train', 'val']:
       combined dir = os.path.join(base dir, train or val, magnification)
       os.makedirs(combined dir, exist ok=True)
       for category in ['benign', 'malignant']:
```

```
category dir = os.path.join(base dir, train or val, category, magnification)
       for filename in os.listdir(category dir):
              src path = os.path.join(category dir, filename)
              dst path = os.path.join(combined dir, category, filename)
              os.makedirs(os.path.join(combined dir, category), exist ok=True)
              if not os.path.exists(dst_path):
              os.symlink(src path, dst path)
# Define patch and window sizes
patch sizes = [(32, 32), (64, 64)]
window sizes = [(64, 64), (128, 128)]
# Loop through each combination of patch and window sizes
for patch size in patch sizes:
       for window size in window sizes:
       print(f'Training model with patch size {patch size} and window size {window size}...')
       # Loop through each magnification and train a model
       for mag in magnifications:
       print(f'Training model for {mag} magnification...')
       # Prepare combined directories
       copy images to combined dir(mag)
       # Create generators for training and validation
       train generator = create generators(mag, 'train', window size)
       val generator = create generators(mag, 'val', window size)
       # Create and train the model
       input shape = (window size[0], window size[1], 3)
       model = create model(input shape)
       history = model.fit(
              train generator,
              steps per epoch=train generator.samples // 32,
              epochs=10,
              validation data=val generator,
              validation steps=val generator.samples // 32
       )
       # Evaluate the model
       val loss, val acc = model.evaluate(val generator)
       print(f\nValidation accuracy for {mag} magnification with patch size {patch size} and window size
{window size}: {val acc}\n')
```

Validation accuracy for 40X magnification with patch size (32, 32) and window size (64, 64): 0.8395990133285522 Validation accuracy for 100X magnification with patch size (32, 32) and window size (64, 64): 0.8417266011238098 Validation accuracy for 200X magnification with patch size (32, 32) and window size (64, 64): 0.8709677457809448 Validation accuracy for 400X magnification with patch size (32, 32) and window size (64, 64): 0.8575342297554016 Validation accuracy for 40X magnification with patch size (32, 32) and window size (128, 128): 0.7092731595039368 Validation accuracy for 100X magnification with patch size (32, 32) and window size (128, 128): 0.6954436302185059 Validation accuracy for 200X magnification with patch size (32, 32) and window size (128, 128): 0.8759305477142334 Validation accuracy for 400X magnification with patch size (32, 32) and window size (128, 128): 0.8739725947380066 Validation accuracy for 40X magnification with patch size (64, 64) and window size (64, 64): 0.8421052694320679 Validation accuracy for 100X magnification with patch size (64, 64) and window size (64, 64): 0.7338129281997681 Validation accuracy for 200X magnification with patch size (64, 64) and window size (64, 64): 0.8387096524238586 Validation accuracy for 400X magnification with patch size (64, 64) and window size (64, 64): 0.8849315047264099 Validation accuracy for 40X magnification with patch size (64, 64) and window size (128, 128): 0.8245614171028137 Validation accuracy for 100X magnification with patch size (64, 64) and window size (128, 128): 0.8561151027679443 Validation accuracy for 200X magnification with patch size (64, 64) and window size (128, 128): 0.8734491467475891 Validation accuracy for 400X magnification with patch size (64, 64) and window size (128, 128): 0.8821917772293091

AUC-ROC scores

Validation accuracy for 40X magnification: 0.7092731595039368

13/13 6s 438ms/step

Classification Report for 40X magnification:

	precision	recall	f1-score	support
Benign	0.33	0.02	0.04	125
Malignant	0.69	0.98	0.81	274
ассигасу			0.68	399
macro avg	0.51	0.50	0.43	399
weighted avg	0.58	0.68	0.57	399

AUC-ROC for 40X magnification: 0.509868613138686

Validation accuracy for 100X magnification: 0.8609112501144409

14/14 6s 435ms/step

Classification Report for 100X magnification:

	precision	recall	f1-score	support
Benign	0.32	0.30	0.31	129
Malignant	0.69	0.71	0.70	288
accuracy			0.59	417
macro avg	0.51	0.51	0.51	417
weighted avg	0.58	0.59	0.58	417

AUC-ROC for 100X magnification: 0.5316268303186908

Validation accuracy for 200X magnification: 0.8684863448143005

13/13 6s 460ms/step

Classification Report for 200X magnification:

	precision	recall	f1-score	support
Benign	0.33	0.31	0.32	125
Malignant	0.70	0.71	0.70	278
ассигасу			0.59	403
macro avg	0.51	0.51	0.51	403
weighted avg	0.58	0.59	0.59	403

AUC-ROC for 200X magnification: 0.5083453237410072

Validation accuracy for 400X magnification: 0.8849315047264099

12/12 6s 473ms/step

Classification Report for 400X magnification:

	precision	recall	f1-score	support
Benign	0.26	0.23	0.24	118
Malignant	0.65	0.69	0.67	247
ассигасу			0.54	365
macro avg	0.46	0.46	0.46	365
weighted avg	0.52	0.54	0.53	365

AUC-ROC for 400X magnification: 0.45138269402319364

SOURCE CODE

import os
import tensorflow as tf
from tensorflow keras preprocessi

from tensorflow.keras.preprocessing.image import ImageDataGenerator from sklearn.metrics import classification_report, roc_auc_score

Define the magnifications and their respective directories

```
magnifications = ['40X', '100X', '200X', '400X']
base dir = '/home/yanajain/Downloads/BreaKHis Processed'
train dirs = {mag: [os.path.join(base dir, 'train', 'benign', mag),
               os.path.join(base dir, 'train', 'malignant', mag)] for mag in magnifications}
val dirs = {mag: [os.path.join(base dir, 'val', 'benign', mag),
               os.path.join(base dir, 'val', 'malignant', mag)] for mag in magnifications}
# Function to create and compile the model
def create model():
       model = tf.keras.models.Sequential([
       tf.keras.Input(shape=(150, 150, 3)),
       tf.keras.layers.Conv2D(32, (3, 3), activation='relu'),
       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(512, activation='relu'),
       tf.keras.layers.Dense(1, activation='sigmoid')
       1)
       model.compile(optimizer='adam',
               loss='binary crossentropy',
               metrics=['accuracy'])
       return model
# Helper function to create generators
def create generators(magnification, train or val):
       benign dir = os.path.join(base dir, train or val, 'benign', magnification)
       malignant dir = os.path.join(base dir, train or val, 'malignant', magnification)
       datagen = ImageDataGenerator(rescale=1./255,
                      rotation range=20,
                      width shift range=0.2,
                      height shift range=0.2,
                      shear range=0.2,
                      zoom range=0.2,
                      horizontal flip=True if train or val == 'train' else False)
       combined dir = os.path.join(base dir, train or val, magnification)
       os.makedirs(combined dir. exist ok=True)
       generator = datagen.flow from directory(combined dir,
                              target size=(150, 150),
                              batch size=32,
```

```
class mode='binary')
       return generator
# Copy images to a combined directory structure
def copy images to combined dir(magnification):
       for train or val in ['train', 'val']:
       combined dir = os.path.join(base dir, train or val, magnification)
       os.makedirs(combined dir, exist ok=True)
       for category in ['benign', 'malignant']:
       category dir = os.path.join(base dir, train or val, category, magnification)
       for filename in os.listdir(category dir):
               src path = os.path.join(category dir, filename)
               dst_path = os.path.join(combined dir, category, filename)
               os.makedirs(os.path.join(combined dir, category), exist ok=True)
               if not os.path.exists(dst_path):
               os.symlink(src path, dst path)
# Loop through each magnification and train a model
for mag in magnifications:
       print(f'Training model for {mag} magnification...')
       # Prepare combined directories
       copy images to combined dir(mag)
       # Create generators for training and validation
       train generator = create generators(mag, 'train')
       val generator = create generators(mag, 'val')
       # Create and train the model
       model = create model()
       history = model.fit(
       train generator,
       steps per epoch=train generator.samples // 32,
       epochs=10,
       validation data=val generator,
       validation steps=val generator.samples // 32
       )
       # Evaluate the model
       val loss, val acc = model.evaluate(val generator)
       print(f\nValidation accuracy for {mag} magnification: {val acc}\n')
       # Obtain true labels and predictions for validation data
```

y_true = val_ generator.classes

y pred probs = model.predict(val generator)

```
y_pred = (y_pred_probs > 0.5).astype('int32')

# Calculate additional metrics
print(f"\nClassification Report for {mag} magnification:\n")
print(classification_report(y_true, y_pred, target_names=['Benign', 'Malignant']))

# Calculate AUC-ROC
auc_roc = roc_auc_score(y_true, y_pred_probs)
print(f"AUC-ROC for {mag} magnification: {auc_roc}\n")
```

Comparing with CNN model

Main Differences Between a Swin Transformer and a CNN Implementation

1. Feature Extraction Mechanism:

- **Swin Transformer:** Employs self-attention within localized windows to capture both short- and long-range dependencies, enabling it to model complex global relationships within the image.
- **CNN:** Utilizes convolutional filters to extract local features, gradually building higher-level representations through stacked layers.

2. Hierarchical Representation:

- **Swin Transformer:** Constructs a multi-scale feature hierarchy through patch merging and the shifted window approach, allowing for dynamic and adaptive feature learning.
- CNN: Builds hierarchical features primarily by stacking convolutional layers and pooling operations, which gradually abstract local features into global patterns.

3. Global Context Modeling:

- **Swin Transformer:** Achieves effective global context modeling by shifting attention windows between layers, thereby integrating information from distant regions without excessive computational cost.
- CNN: Focuses on local regions and may require deeper architectures or additional modules (such as global pooling) to capture broader contextual information.

4. Computational Efficiency:

- Swin Transformer: Limits self-attention computations to fixed-size windows, significantly reducing the quadratic complexity seen in standard transformers and making it more efficient for high-resolution images.
- CNN: Relies on fixed convolutional operations, which are computationally
 efficient for detecting local patterns, though they may require additional depth
 or techniques to achieve similar global context awareness.

Conclusion

The Swin Transformer model using Tensorflow provided better accuracy than the CNN model. However, due to high computational cost and large datasets the client-based server didn't give a drastic result when compared to our previous basic CNN model.