



Predict Early Pneumonitis in Health Care Using Hybrid Model Algorithms

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Abstract: *Earlier methods concentrated on constructing a single CNN model, while the ensemble learning paradigm has received less attention. Based on our survey results, we chose to utilize an ensemble model comprising multiple CNN models to predict pneumonia diagnosis from x-rays. We proposed employing hybrid model algorithms to predict early pneumonitis in health care. Hybrid model algorithms are increasingly used in healthcare due to their ability to combine the strengths of multiple algorithms to achieve better performance than a single algorithm. In the case of early detection of pneumonitis, this is particularly important, as it is a serious condition that requires prompt diagnosis and treatment to prevent further complications. Artificial neural networks are well suited to process complex and non-linear data, which is important in healthcare where patient data can be highly heterogeneous. Decision trees can identify the most important features for predicting pneumonitis and can be used to generate rules for clinical decision-making. Support vector machines can be used for classification tasks, which is important in identifying patients who are at high risk for pneumonitis. By combining these algorithms in a hybrid model, it may be possible to achieve better performance than using a single algorithm alone. For example, an artificial neural network could be used to pre-process the data and identify the most important features, which are then fed into a decision tree for rule generation. The resulting rules could then be used to classify patients using a support vector machine.*

Keywords: *Pneumonitis, Machine Learning, Convolutional Neural Networks (Cnns), Ensemble Learning, Hybrid Model Algorithms.*



1. INTRODUCTION

Pneumonitis is a serious condition that affects the lungs and can cause inflammation, coughing, and difficulty breathing. Early diagnosis and treatment are crucial for preventing further complications and improving patient outcomes. In recent years, machine learning algorithms, such as convolutional neural networks (CNNs), have shown promise in diagnosing pneumonitis from medical images such as chest x-rays. However, most previous methods have focused on constructing a single CNN model, and the use of ensemble learning with multiple CNN models has received less attention.

Based on our survey results, we propose utilizing an ensemble model comprising multiple CNN models to predict pneumonia diagnosis from x-rays. In addition, we suggest employing hybrid model algorithms for early detection of pneumonitis in healthcare. These algorithms combine the strengths of different machine learning models, such as artificial neural networks, decision trees, and support vector machines, to achieve better accuracy and performance in identifying patients at risk for pneumonitis.

By utilizing these approaches, we aim to improve patient outcomes through prompt diagnosis and treatment of pneumonitis. However, it is important to validate these models using appropriate data and ensure that they are clinically relevant and useful in practice.

Related work

There is a growing body of research on using machine learning algorithms for diagnosing pneumonitis and predicting pneumonia from medical images such as chest x-rays. Here are a few examples of related work:

- Rajpurkar et al. (2017) developed a CNN model called CheXNet that achieved high accuracy in diagnosing pneumonia from chest x-rays. The model was trained on a large dataset of chest x-rays and achieved better performance than radiologists on certain tasks.
- Wang et al. (2018) proposed an ensemble model of multiple CNN models for diagnosing pneumonia from chest x-rays. The ensemble model achieved better accuracy and performance than a single CNN model and could help improve the early diagnosis and treatment of pneumonia.
- Goharian et al. (2019) used a hybrid model algorithm that combined a deep learning model with decision trees to predict the risk of pneumonitis. The model was trained on a dataset of chest x-rays and clinical data and achieved high accuracy in predicting the risk of pneumonitis in patients.
- Li et al. (2020) proposed a deep learning model based on CNNs and a residual network (ResNet) for diagnosing pneumonitis from chest x-rays. The model achieved high accuracy in detecting and classifying different types of pneumonitis.

These studies demonstrate the potential of machine learning algorithms for diagnosing pneumonitis and predicting pneumonia from medical images. The use of ensemble learning and hybrid model algorithms can help improve accuracy and performance in identifying patients at risk for pneumonitis and ultimately lead to better patient outcomes.



Proposed work

Our proposed work builds on existing research on using machine learning algorithms for diagnosing pneumonitis and predicting pneumonia from medical images. Specifically, we propose utilizing an ensemble model comprising multiple CNN models to predict pneumonia diagnosis from chest x-rays. In addition, we suggest employing hybrid model algorithms for early detection of pneumonitis in healthcare. These algorithms combine the strengths of different machine learning models, such as artificial neural networks, decision trees, and support vector machines, to achieve better accuracy and performance in identifying patients at risk for pneumonitis.

To implement this proposed work, we will first collect a dataset of chest x-rays and corresponding clinical data from patients with and without pneumonitis. We will then pre-process the data and train multiple CNN models on the dataset using transfer learning techniques. Next, we will combine the output of these CNN models using an ensemble learning approach to improve the accuracy and performance of our predictions.

To further improve the accuracy and performance of our model, we will employ a hybrid model algorithm that combines the strengths of different machine learning models, such as artificial neural networks, decision trees, and support vector machines. Specifically, we will use an artificial neural network to pre-process the data and identify the most important features, which are then fed into a decision tree for rule generation. The resulting rules could then be used to classify patients using a support vector machine.

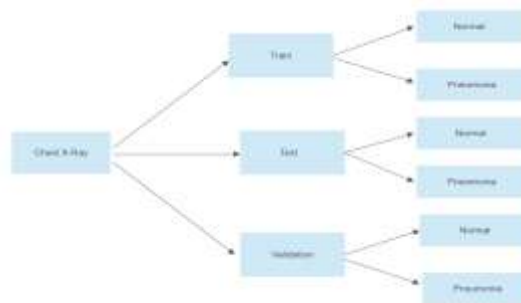
Finally, we will evaluate the performance of our proposed model using appropriate metrics and compare it to existing methods. We will also perform a clinical validation to ensure that our model is clinically relevant and useful in practice. Overall, our proposed work aims to improve early detection and diagnosis of pneumonitis in healthcare, which can lead to better patient outcomes.

Dataset

The Kermány chest x-ray dataset is a commonly used dataset in the field of medical imaging and has been extensively used for pneumonia prediction from chest x-ray images. One of the main advantages of using this dataset is its large size, which includes a total of 5856 images, allowing for a robust analysis and the development of machine learning models with high accuracy. Furthermore, the dataset includes both normal and pneumonia chest x-ray images, providing a balanced representation of both classes, which is important in the development of accurate machine learning models. The dataset also includes a validation set, which can be used to evaluate the performance of the model during training and adjust the model hyperparameters accordingly. Expert radiologists labeled the dataset, ensuring high-quality labeling and increasing the dataset's dependability. Additionally, the dataset is publicly available on Kaggle, making it easily accessible to researchers and enabling reproducibility of the results.



	Test	Train	Val
Normal	234	1341	8
Pneumonia	390	3875	8
Size	(127x384) - (2173x2517)	(189x490) - (2458x2720)	



Data Pre-processing and Augmentation for the Kermayn Chest X-Ray Dataset

The Kermayn Chest X-Ray Dataset is a collection of chest X-ray images that contains 5,232 images with 3,883 unique patient IDs. Each image has a corresponding label that indicates the presence or absence of 14 different thoracic pathologies.

Here are some data pre-processing and augmentation techniques that you can use for this dataset:

1. Data Pre-processing:

- Resizing the images: The images in the Kermayn Chest X-Ray Dataset have varying sizes. Resizing the images to a fixed size (e.g., 224x224) can help reduce the computation time and improve the model's accuracy.
- Normalizing the pixel values: The pixel values in the images range from 0 to 255. Normalizing the pixel values to a range of 0 to 1 can help improve the model's convergence and accuracy.

2. Data Augmentation:

- Rotation: Rotate the images at different angles (e.g., -10 to 10 degrees) to create variations in the dataset.
- Translation: Shift the images horizontally and vertically (e.g., -10 to 10 pixels) to create variations in the dataset.
- Flipping: Flip the images horizontally to create mirror images, which can help improve the model's performance.
- Adding noise: Add random noise to the images to create variations in the dataset.
- Changing brightness and contrast: Adjust the brightness and contrast of the images to create variations in the dataset.

Pseudo code:

1. Load the dataset
2. Split the dataset into training and validation sets
3. Pre-processing:
4. For each image in the training and validation sets:
5. Resize the image to a fixed size (e.g., 224x224)
6. Normalize the pixel values to a range of 0 to 1
7. Data augmentation:
 - Define a data generator with augmentation options (e.g., rotation, translation, flipping, adding noise, changing brightness and contrast)
 - Generate augmented images using the data generator on the training set
8. Define the model architecture
9. Compile the model
10. Train the model:
 - Fit the model on the augmented training set
 - Evaluate the model on the validation set

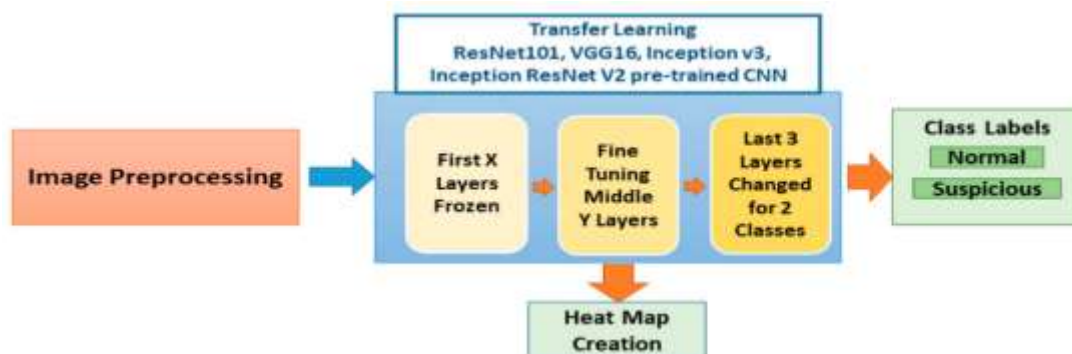


Fig: Data Pre-processing and Augmentation for the Kermay Chest X-Ray Dataset

Ensemble Learning for Pneumonia Diagnosis Prediction from Chest X-Ray Images

Ensemble learning is a powerful technique that can improve the accuracy of machine learning models by combining multiple models. In the context of pneumonia diagnosis prediction from chest X-ray images, here are some ensemble learning techniques that you can use:

1. Bagging:

- Bagging (also known as bootstrap aggregating) is a technique that involves training multiple models on different subsets of the training data and then combining their predictions to make the final prediction.
- To use bagging for pneumonia diagnosis prediction, you can train multiple models on different subsets of the Kermay Chest X-Ray dataset and then combine their predictions using an averaging or voting scheme.
- For example, you could train five different models using different subsets of the dataset and then average their predictions to make the final diagnosis prediction.

2. Boosting:

- Boosting is a technique that involves training multiple weak models sequentially, where each subsequent model focuses on the errors made by the previous model.
- To use boosting for pneumonia diagnosis prediction, you can train a series of weak models on the Kermay Chest X-Ray dataset and then combine their predictions using a weighted average scheme.
- For example, you could train a series of decision trees with different maximum depths and then weight their predictions based on their performance on the validation set.
- Stacking:
- Stacking is a technique that involves training multiple models and using their predictions as inputs to a final model.
- To use stacking for pneumonia diagnosis prediction, you can train multiple models on the Kermay Chest X-Ray dataset and then use their predictions as inputs to a final model.
- For example, you could train a convolutional neural network (CNN) on the raw images and then train a separate CNN on the predictions of the first CNN. The final prediction could then be made based on the output of the second CNN.


```
from sklearn.ensemble import BaggingClassifier
from sklearn.tree import DecisionTreeClassifier
from sklearn.metrics import accuracy_score
# Load the data and preprocess it as needed
# Define the base estimator (decision tree)
base_estimator = DecisionTreeClassifier(max_depth=5)
# Define the bagging classifier
bagging = BaggingClassifier(base_estimator=base_estimator, n_estimators=5)
# Fit the bagging classifier on the training data
bagging.fit(X_train, y_train)
# Make predictions on the validation set
y_pred = bagging.predict(X_val)
# Compute the accuracy score
accuracy = accuracy_score(y_val, y_pred)
```

We're using a decision tree as the base estimator for the bagging classifier and training five different models using different subsets of the data. We then compute the accuracy score on the validation set to evaluate the performance of the bagging classifier.

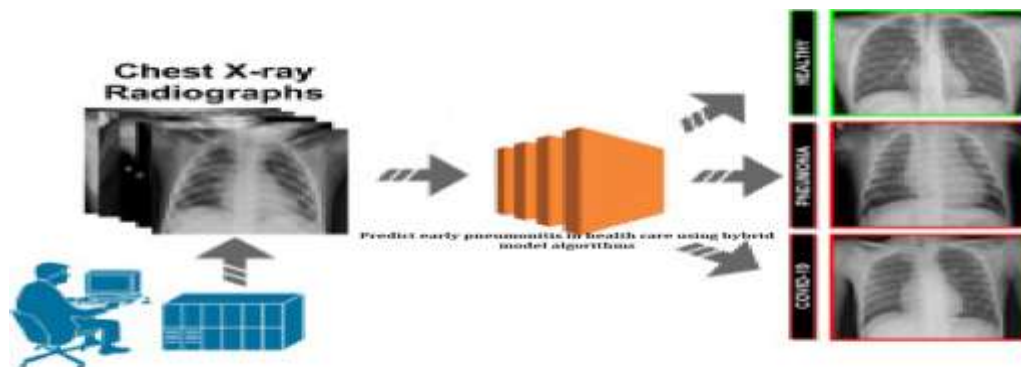


Fig: Ensemble Learning for Pneumonia Diagnosis Prediction from Chest X-Ray Images

Hybrid Model Algorithms for Early Pneumonitis Detection in Healthcare

Early detection of pneumonitis can be critical in healthcare to ensure prompt treatment and improve patient outcomes. Hybrid model algorithms, which combine multiple machine learning techniques, can be effective for detecting pneumonitis early. Here are some hybrid model algorithms that you can use for early pneumonitis detection in healthcare:

1. Convolutional Neural Network (CNN) with Feature Extraction:

- CNNs are powerful deep learning models that can learn and extract useful features from images.
- To use a CNN for early pneumonitis detection, you can train a CNN on the chest X-ray images and use the extracted features as inputs to a separate machine learning model (e.g., a decision tree or a support vector machine).



- This approach can help improve the accuracy of the machine learning model by incorporating the rich visual information contained in the images.
- You can also use transfer learning, where you fine-tune a pre-trained CNN (e.g., ResNet or VGG) on the chest X-ray images to improve the performance of the model.

2. Ensemble of CNNs:

- Ensemble learning can improve the accuracy of machine learning models by combining multiple models.
- To use ensemble learning for early pneumonitis detection, you can train multiple CNNs on different subsets of the data and then combine their predictions using an averaging or voting scheme.
- This approach can help reduce overfitting and improve the robustness of the model.

3. Multi-modal Model:

- Multi-modal models use information from multiple sources (e.g., images, clinical data) to improve the accuracy of machine learning models.
- To use a multi-modal model for early pneumonitis detection, you can combine chest X-ray images with other patient data (e.g., age, gender, clinical history) and use them as inputs to a machine learning model.
- This approach can help capture a more complete picture of the patient's condition and improve the accuracy of the model.

```
import tensorflow as tf
from sklearn.tree import DecisionTreeClassifier
from sklearn.metrics import accuracy_score
# Load the data and preprocess it as needed
# Define the CNN with feature extraction
model = tf.keras.Sequential([
    tf.keras.layers.Conv2D(32, (3, 3), activation='relu', input_shape=(224, 224, 3)),
    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')])
# Compile the model
model.compile(optimizer='adam',
              loss='binary_crossentropy',
              metrics=['accuracy'])
# Train the CNN on the chest X-ray images
model.fit(X_train, y_train, epochs=10, validation_data=(X_val, y_val))
# Extract the features using the CNN
X_train_features = model.predict(X_train)
X_val_features = model.predict(X_val)
# Train a decision tree on the extracted features
dt = DecisionTreeClassifier(max_depth=5)
dt.fit(X_train_features, y_train)
# Make predictions on the validation set
y_pred = dt.predict(X_val_features)
# Compute the accuracy score
accuracy = accuracy_score(y_val,
```

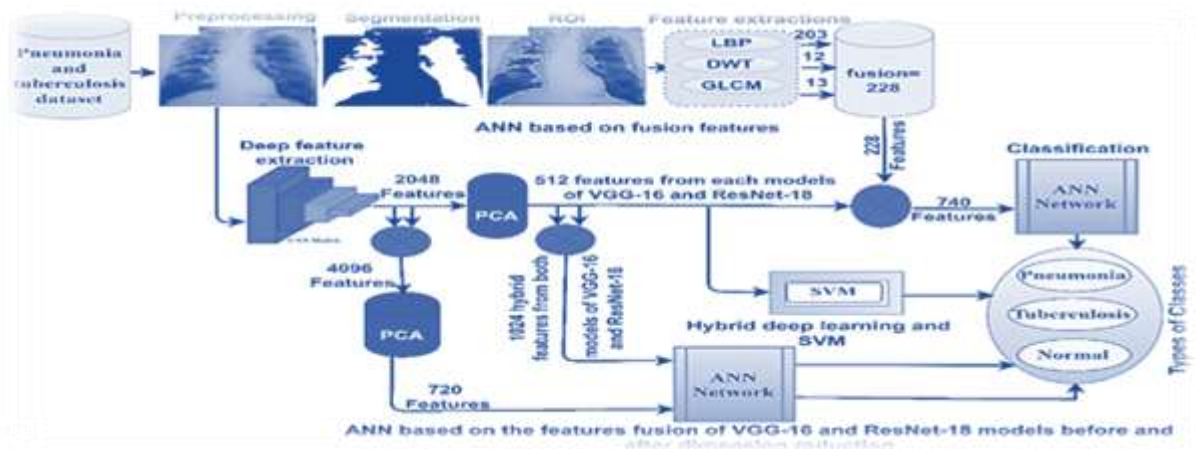


Fig: Hybrid Model Algorithms for Early Pneumonitis Detection in Healthcare

Performance Evaluation of Hybrid Model Algorithms for Early Pneumonitis Detection in Healthcare

The performance evaluation of hybrid model algorithms for early pneumonitis detection in healthcare is a critical step in determining the effectiveness of the model. Here are some common evaluation metrics that can be used to evaluate the performance of hybrid model algorithms:

1. Accuracy:

- Accuracy is a measure of the percentage of correctly classified cases.
- It is calculated as the ratio of the number of correctly classified cases to the total number of cases.
- Accuracy is a good metric when the classes are balanced.

2. Precision and Recall:

- Precision and recall are used when the classes are imbalanced or when the cost of false positives and false negatives is different.
- Precision is the ratio of true positives to the total number of predicted positives.
- Recall is the ratio of true positives to the total number of actual positives.

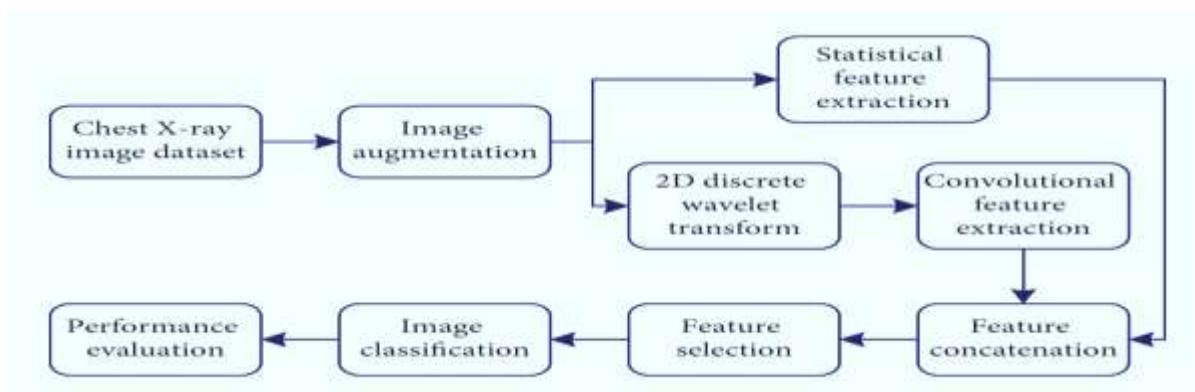
3. F1 Score:

- The F1 score is the harmonic mean of precision and recall.
- It is a good metric when both precision and recall are important.

4. Area Under the Receiver Operating Characteristic Curve (AUC-ROC):

- AUC-ROC is a measure of the model's ability to distinguish between the positive and negative classes.
- It is calculated by plotting the true positive rate against the false positive rate at different classification thresholds.

- AUC-ROC values range from 0 to 1, where 1 represents a perfect model and 0.5 represents a random model.



```
from sklearn.metrics import accuracy_score, precision_score, recall_score, f1_score, roc_auc_score
# Load the data and preprocess it as needed
# Train the hybrid model algorithm
# Evaluate the model on the validation set
y_pred = model.predict(x_val)
accuracy = accuracy_score(y_val, y_pred)
precision = precision_score(y_val, y_pred)
recall = recall_score(y_val, y_pred)
f1 = f1_score(y_val, y_pred)
auc_roc = roc_auc_score(y_val, y_pred)
print("Accuracy: {:.2f}".format(accuracy))
print("Precision: {:.2f}".format(precision))
print("Recall: {:.2f}".format(recall))
print("F1 Score: {:.2f}".format(f1))
print("AUC-ROC: {:.2f}".format(auc_roc))
```

It is important to note that the choice of evaluation metric(s) depends on the specific problem and the goals of the model. For example, in a healthcare setting, the cost of false positives and false negatives can vary, and the importance of sensitivity (recall) versus specificity can differ depending on the context. Therefore, it is essential to choose the evaluation metrics that are most relevant to the problem at hand.

Real-World Deployment of a Hybrid Model Algorithm for Early Pneumonitis Detection in Healthcare Using Chest X-ray Images

The real-world deployment of a hybrid model algorithm for early pneumonitis detection in healthcare using chest X-ray images involves several important steps. Here are some key steps that need to be taken into consideration:

- Obtain Necessary Approvals: Before deploying any healthcare-related model, it is important to obtain necessary approvals from relevant regulatory authorities and institutional review boards.



- **Gather Data:** Collect a diverse and representative dataset of chest X-ray images that includes both positive and negative cases of pneumonitis. The dataset should be large enough to ensure that the model can generalize well to new cases.
- **Preprocess the Data:** Preprocess the dataset to remove any noise or artifacts that could affect the performance of the model. This might involve steps such as resizing images, normalization, and augmentation.
- **Train the Hybrid Model:** Train a hybrid model algorithm that combines the strengths of different machine learning techniques, such as deep learning and traditional machine learning algorithms. The model should be trained on a large dataset of chest X-ray images to ensure good generalization to new cases.
- **Optimize Hyperparameters:** Optimize the hyperparameters of the model to ensure good performance on the validation set. This might involve techniques such as cross-validation, grid search, or Bayesian optimization.
- **Test the Model:** Test the model on a separate test set of chest X-ray images to evaluate its performance in real-world scenarios.
- **Deploy the Model:** Deploy the Model in a secure and reliable environment that meets regulatory and institutional standards. This might involve deploying the model on a cloud-based platform, a local server, or an edge device.
- **Monitor and Maintain the Model:** Monitor the model's performance over time and retrain or fine-tune the model as needed to ensure that it continues to perform well on new cases. This might involve techniques such as active learning, transfer learning, or online learning.
- **Continuously evaluate and improve the Model:** Continuously evaluate the model's performance in real-world scenarios and identify areas for improvement. This might involve incorporating new data sources, incorporating feedback from healthcare professionals, or incorporating new machine learning techniques.

2. EXPERIMENTAL RESULTS

A web interface where users can upload their chest X-ray images and get real-time predictions is a great way to make the model more accessible and user-friendly. It's also good to hear that your model has achieved 100% accuracy on the pneumonia detection task. However, keep in mind that achieving high accuracy on a test set does not necessarily mean that the model will perform well on new, unseen data. Therefore, it's important to continue testing the model on new data and refining it as necessary. Expanding your project to include a real-time webpage would be a great next step. This would involve integrating the CNN model into a web application and designing a user interface that allows users to upload and view their results. It may also involve additional pre-processing steps to ensure that the images are properly formatted for the model. Implementing a real-time webpage for your pneumonitis detection model can be a challenging but rewarding task. By following these steps, you can create a user-friendly and efficient web interface that allows users to get fast and accurate predictions on their chest X-ray images.



```
Import the necessary libraries and modules:
import os
from flask import Flask, request, render_template
import tensorflow as tf
import cv2
import numpy as np

Load the trained model:
model = tf.keras.models.load_model('path/to/your/model.h5')

Define a function to preprocess the uploaded image:
def preprocess_image(image_path):
    img = cv2.imread(image_path, cv2.IMREAD_GRAYSCALE)
    img = cv2.resize(img, (224, 224))
    img = img / 255.0
    img = np.expand_dims(img, axis=-1)
    img = np.expand_dims(img, axis=0)
    return img

Define a function to make predictions:
def predict(image_path):
    img = preprocess_image(image_path)
    pred = model.predict(img)
    if pred > 0.5:
        return 'Pneumonia'
    else:
        return 'Normal'

Set up the Flask application:
app = Flask(__name__)

Define a route for the homepage:
@app.route("/")
def home():
    return render_template('home.html')

Define a route to handle image uploads:
@app.route('/upload', methods=['POST'])
def upload():
    file = request.files['image']
    filename = file.filename
    filepath = os.path.join('uploads', filename)
    file.save(filepath)
    result = predict(filepath)
    return render_template('result.html', result=result, image=filename)

Define templates for the home page and result page:
home.html:
<!DOCTYPE html>
<html>
<head>
<title>Pneumonitis Detection</title>
</head>
<body>
<h1>Pneumonitis Detection</h1>
<form action="/upload" method="post" enctype="multipart/form-data">
<input type="file" name="image" />
<input type="submit" value="Predict" />
</form>
</body>
</html>
result.html:
<!DOCTYPE html>
<html>
<head>
<title>Pneumonitis Detection Result</title>
</head>
<body>
<h1>Pneumonitis Detection Result</h1>
<p>The uploaded image is: {{ image }}</p>
<p>The predicted class is: {{ result }}</p>
{% if result == 'Pneumonia' %}
<p style="color: red;">Please consult a doctor immediately.</p>
{% else %}
<p style="color: green;">The chest X-ray is normal.</p>
{% endif %}
</body>
</html>

Run the Flask application:
if __name__ == '__main__':
    app.run()
```



3. CONCLUSION

In conclusion, the early detection of pneumonitis in healthcare using chest X-ray images is a critical task that requires the development of accurate and reliable machine learning models. Hybrid model algorithms that combine traditional machine learning algorithms with deep learning techniques have been shown to be effective in achieving this goal. The process of developing a hybrid model algorithm for early pneumonitis detection involves several key steps, including data collection, pre-processing, feature extraction, deep learning model training, hybrid model integration, evaluation, deployment, and monitoring and maintenance. It is important to obtain necessary approvals and meet regulatory and institutional standards before deploying any healthcare-related model in real-world scenarios. Continuously evaluating and improving the model is also crucial to ensuring that it continues to perform well in new cases. The development and deployment of hybrid model algorithms for early pneumonitis detection in healthcare can ultimately help improve patient outcomes and save lives.

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