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Think positive: An interpretable neural network for image recognition

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Abstract

The COVID-19 pandemic is an ongoing pandemic and is placing additional burden on healthcare systems around the world. Timely and effectively detecting the virus can help to reduce the spread of the disease. Although, RT-PCR is still a gold standard for COVID-19 testing, deep learning models to identify the virus from medical images can also be helpful in certain circumstances. In particular, in situations when patients undergo routine X-rays and/or CT-scans tests but within a few days of such tests they develop respiratory complications. Deep learning models can also be used for pre-screening prior to RT-PCR testing. However, the transparency/interpretability of the reasoning process of predictions made by such deep learning models is essential. In this paper, we propose an interpretable deep learning model that uses positive reasoning process to make predictions. We trained and tested our model over the dataset of chest CT-scan images of COVID-19 patients, normal people and pneumonia patients. Our model gives the accuracy, precision, recall and F-score equal to 99.48%, 0.99, 0.99 and 0.99, respectively.

Keywords: CT-scan, Prototypes, COVID-19, Pneumonia, Interpretable.

1. Introduction

The pandemic COVID-19 is placing enormous strain on public health systems around the world, and severely affecting the economies of many countries. Although, vaccination is being done for the virus, but the number of the variants of the virus is also increasing. The new variants of the virus can reduce the effectiveness of the vaccines [48]. Therefore, along with vaccination for the virus, detection of the virus is important to reduce the spread of the disease and the development of mutants of the virus. In addition to the prevalent testing technique reverse transcription polymerase chain reaction (RT-PCR), deep learning models can also be helpful in efforts to detect the virus. Most of the deep learning algorithms work as a black-box because their reasoning process for their predictions is not transparent/interpretable. However, the interpretation of the reasoning process of a deep learning model related to a high stake decision is important. There have been cases where
erroneous data fed into the black-box models went unnoticed, due to which wrongful long prison sentences were given (e.g., inmate Glen Rodriguez was denied parole because of wrong COMPAS score) [26] and [47]. The lack of interpretability of the reasoning processes of such deep learning models has become a major issue for whether we can trust predictions that are coming from these models. Therefore, we propose an interpretable deep learning model quasi prototypical part network (Quasi-ProtoPNet), and trained and tested the model over the dataset of chest CT images.

1.1. Related work

In this section, we first discuss those works that are related to our paper because of the interpretability of their reasoning process. Second, we provide a brief summary of the studies that are related to this study as they categorize medical images (chest CT-scan and X-ray images). The models in the second category attempt to distinguish medical images of COVID-19 patients from the medical images of pneumonia patients and normal people, but the models are not necessarily interpretable.

Several approaches have emerged to interpret convolutional neural networks, including post hoc interpretability analysis. Once a neural network performs the classification, post hoc analysis is used to interpret the neural network. Deconvolution [61], saliency visualization [38, 43, 44, 47] and activation maximization [11, 19, 25, 28, 38, 59] are a few examples of post hoc analysis technique. However, these visualization approaches of post hoc analysis do not shed light on the reasoning process with clarity.

Attention-based interpretability is another technique to clarify the reasoning process of the neural networks. The instances of this technique include part-based models [12, 14, 21, 34, 37, 45, 53, 62, 63, 65] and class activation maps (CAM) [64]. In this approach, the aim of a model is to show the patches of an input image that are the focus of its attention; nonetheless, these models do not represent prototypes that resemble the parts of an input image that are the focal points of the models. Recently, a model CXR-specific with class activation maps has also been developed to detect COVID-19 from medical images [32].

Case-based classification techniques that use prototypes [4, 31, 52] or k-nearest neighbours [30, 36, 46] are also related to our work. Throughout this paper, a prototype or a prototypical part will represent a patch of an image. Li et al. [26] have developed a model that uses full image-sized prototypes and requires a decoder for visualizing prototypes. Chen et al. [6] developed a model ProtoPNet which significantly improved on the model developed in [26].

![Figure 1: For a given CT-scan image of a COVID-19 patient, Quasi-ProtoPNet identifies the parts of the image where it thinks that this part of the image is similar to that learned prototype.](image-url)
As shown in Figure 1, ProtoPNet is able to identify different parts of an input image that are similar to different prototypes, and it classifies an image based on the similarity scores. To classify an input image, ProtoPNet finds the Euclidean distance between each latent patch of the input image and the learned prototypes of images from different classes, where prototypes have spatial dimensions $1 \times 1$. The maximum of the inverted distances between a prototype and the patches of the input image is called the similarity score of the prototype. Note that, the smaller the distance, the larger the reciprocal, and there will be only one similarity score for each prototype. A weighted combination of similarity scores is used to determine the logits for different classes and these logits are normalized using Softmax to determine the class of the input image. The weights for the correct class and incorrect class of a training image are set equal to 1 and $-0.5$, respectively. These weights are also called connections of the similarity scores with the classes. The negative weights are assigned to include the negative reasoning process, that is, to reject the incorrect classes. ProtoPNet tries to zero out the negative weights during the training process, and with this assumption of ProtoPNet, a theorem is proved [6, Theorem 1.1]. However, our experiments show that it is hardly possible to zero out the negative connections during the training process after making a negative connection between the similarity scores and incorrect classes.

The models NP-ProtoPNet [42], Gen-ProtoPNet [40] and Ps-ProtoPNet [41] are variations of ProtoPNet, and we refer to these four models collectively the ProtoPNet models or the series of ProtoPNet models. Gen-ProtoPNet model uses a generalized version of the Euclidean distance function, NP-ProtoPNet considers the negative reasoning process and the positive reasoning process equally, and Ps-ProtoPNet model uses the connections between logits and similarity scores as suggested by [41, Theorem 1], and uses the generalized version of the distance function. The theorem [41, Theorem 1] uses a more realistic assumption of fixed negative connections between similarity scores and incorrect classes to find the impact of change in the negative connections on the logits. The impact on the logits is obtained due to the projection of prototypes to the actual patches of training images, that is, the replacement of the prototypes with the patches of the training images. However, the use of fixed negative connections leads to decrease in the logit of correct class and increase in the logit of incorrect classes, consequently the accuracy of Ps-ProtoPNet decreases after the projection of prototypes. In particular, the impact is more severe when the number of classes is small, see [41, Theorem 1]. In summary, each model of the series of ProtoPNet models uses the negative reasoning process along with the positive reasoning process, whereas our model Quasi-ProtoPNet uses only positive reasoning process to categorize images.

In order to get rid of the flaws of the ProtoPNet models, especially when the number of classes is small, Quasi-ProtoPNet uses only positive reasoning process by placing zero connection between the similarity scores and incorrect classes. Quasi-ProtoPNet suspends the convex optimization of the last layer to keep the connections constant, where by the suspension of the convex optimization of the last layer means that Quasi-ProtoPNet does not optimize the last layer by freezing all other layers. In addition to the positive reasoning process, Quasi-ProtoPNet uses prototypes of all types of spatial dimensions, that is, rectangular spatial dimensions and square spatial dimensions, whereas ProtoPNet model uses the
prototypes with only square spatial dimensions 1 × 1. Prototypes with large spatial dimensions help our model to classify the images on the basis of objects instead of backgrounds of the objects in the images. However, the optimum spatial dimensions need to be determined to get better accuracy.

To identify an image that has not been previously exposed, humans can compare patches of the image with patches of images of known objects. This type of reasoning is usually used in difficult identification tasks. For example, radiologists may compare suspicious tumors in an X-ray or a CT-scan image with prototype tumor images to diagnose cancer. This type of human reasoning inspired our model where comparison of image parts with learned prototypes is an integral part of the model’s reasoning process. Therefore, our model differentiates between CT-scan images of a COVID-19 patient and CT-scan images of pneumonia patients based on greater similarity between the learned prototypes and the patches of images.

Several non-interpretable networks have been proposed to distinguish chest CT-scan or X-ray images of COVID-19 patients from chest CT-scan or X-ray images of pneumonia patients and normal people, see [1, 2, 5, 7, 8, 9, 10, 15, 17, 22, 23, 24, 29, 33, 32, 60]. Some studies have surveyed the machine learning/deep learning models that classify chest CT-scan images or X-ray images of COVID-19 patients, pneumonia patients and normal people. A survey by Bhattacharya et al. [3] signifies the lack of sufficient and reliable data of the medical images related COVID-19 patients for neural networks, but a model’s reliability depends data. However, we experimented our model over currently publicly available the biggest dataset of the CT-scan images [16]. Few more studies [54, 55, 56, 57, 58] related to multi-view hashing and image retrieval are also worth mentioning.

1.2. Dataset

We choose the dataset [16] of chest CT-scan images of COVID-19 patient, normal people and pneumonia patients to train and test our model. The dataset consists of 143778 training images and 25658 test images. We crop the images using the bounding box information provided with the dataset. Also, we use the information provided with the dataset to segregate the cropped images into three classes Covid, Normal and Pneumonia that contain the images of COVID-19 patients, normal people and pneumonia patients, respectively. We also call these classes first, second and third, and denote them by C, N and P, respectively. The classes C, N and P have 35996, 25496 and 82286 training images, and 12245, 7395 and 6018 test images, respectively. All images have been resized to the dimensions 224 × 224 as required by the base models.

1.3. Contributions

The novelty of our model is that it uses positive reasoning process along with the use of prototypes that can have any type of spatial dimensions, that is, rectangular spatial dimensions and square spatial dimensions. Quasi-ProtoPNet uses an objective function different from the objective function used in the series of ProtoPNet models. The contributions of this paper are summarized below.
Quasi-ProtoPNet uses only the positive reasoning process by maintaining zero connection between the similarity scores and incorrect classes. Quasi-ProtoPNet suspends the convex optimization of the last layer to keep the connections fixed. The suspension of the convex optimization also reduces the training time considerably.

The architecture of Quasi-ProtoPNet helped us to prove a theorem, see Theorem 3.1. The theorem provides the theoretical evidence of the reason of the improvement in the performance of our model over the other ProtoPNet models. We remark that the theorem is not only true for the distance function that we use for our model, but it is also true for any positive-valued function that satisfies the triangular inequality and has appropriate domain.

Quasi-ProtoPNet uses prototypes with both types of spatial dimensions, that is, rectangular spatial dimensions and square spatial dimensions, whereas ProtoPNet model uses prototypes with only square spatial dimensions $1 \times 1$.

The rest of the paper is organized as follows. In Section 2, we provide a detailed information about the architecture of our model, and we explain the training procedure and reasoning process of our model. In Section 3, we provide confusion matrices for our model with different base models, and we compare the performance of our model with the ProtoPNet models and the base models. Also, we show that the improvement in the accuracies given by our model over the accuracies given by the other ProtoPNet models is statistically significant. A graphical comparison of the accuracies is provided. In this section, we also prove a theorem that finds the bounds of the changes in logits due to projection of prototypes on the training images. In Section 4, we talk about the limitations of our model. In Section 5, a brief discussion on our model and the series of ProtoPNet models is provided. Finally, in Section 6, we conclude our work.

2. Method

In this section, we introduce and explain the architecture and the training process of our model Quasi-ProtoPNet in the context of CT-scan images.

2.1. Quasi-ProtoPNet architecture

Quasi-ProtoPNet can be built on convolutional layers of a state-of-the-art base model (baseline), such as: VGG-19 [39], ResNet-34, ResNet-152 [18], DenseNet-121, or DenseNet-161 [20]. As shown in Figure 2, Quasi-ProtoPNet consists of the convolution layers of a base model that are followed by two additional convolutional layers $2 \times 1$ and $1 \times 1$. These convolutional layers are collectively denoted by $L$, and they are followed by a generalized convolutional layer $p_t$ of prototypical parts. The layer $p_t$ is followed by a dense layer $w$ with no bias. The parameters of $L$ and the weight matrix of a dense layer are denoted by $L_{\text{conv}}$ and $w_{\text{init}}$, respectively. The activation functions ReLU and Sigmoid are used for the additional second last convolutional layer and last convolution layer, respectively. Note that, convolutional layers $L$ form a non-interpretable (black-box) part of our model whereas the generalized convolutional layer $p_t$ forms the interpretable (transparent) part of our model.
Figure 2: Quasi-ProtoPNet architecture.

Although, convolutional layers of any of the base models can be used to construct our model, we provide the explanation of Quasi-ProtoPNet when it is constructed over the convolutional layers of VGG-16. Let \( x \) be an input image. Since the output of the convolutional layers of VGG-16 has depth 512 and spatial dimensions \( 7 \times 7 \), \( L(x) \) has depth 512 and spatial dimensions \( 6 \times 6 \). Note that, the layer \( p_t \) is a vector of prototypical units, and each prototypical unit is a tensor of the shape \( 512 \times h \times w \), where \( 1 \times 1 < h \times w < 6 \times 6 \), that is, \( h \) and \( w \) together are neither equal to 1 nor 6. Suppose \( n \) and \( m \) denote the total number of classes and prototypes for each class, respectively. Let \( P_c = \{ p_{tc} \}_{l=1}^{m} \) be the set of prototypes of a class \( c \) and \( P = \{ P_c \}_{c=1}^{n} \) is set of all prototypes. For our work \( n = 3 \), but we randomly set the hyperparameter \( m = 10 \).

The shapes of \( L(x) \) and \( p_t \) are \( 512 \times 6 \times 6 \) and \( 512 \times h \times w \), where \( h \) and \( w \) lies between 1 and 6 but together they are neither equal to 1 nor 6. Therefore, each prototype can be thought of as a part of \( L(x) \). The model takes into account the spatial relationship between \( L(x) \) and the prototypical parts, and upsamples the part of \( L(x) \) (the part of \( L(x) \) that is at the smallest distance from a prototypical part) to the input image \( x \) to identify the patch on \( x \) that resembles similar to a prototype. The green rectangles in the source images are the parts of the source images from where the prototypes are actually projected. The source image of the prototypes \( p_1^1, p_2^1 \) and \( p_3^1 \) are also shown in the Figure 2. Similar to ProtoPNet (see Section 1.1), Quasi-ProtoPNet computes the similarity scores between an input image and prototypes \( p_1^1 - p_{10}^1, p_3^1 - p_{10}^3 \) and \( p_1^3 - p_{10}^3 \), see Figure 2. The prototype \( p_1^1, p_3^1 \) and \( p_3^3 \) have similarity scores 2.8001, 0.7889 and 1.0233, and the similarity score of \( p_1^1 \) is greater than the other two similarity scores. The complete list of similarity scores obtained from our experiments is given in the matrix \( s_m \), see Section 2.3.

In the dense layer \( w \), the matrices \( w_m \) and \( s_m \) are multiplied to obtain the logits. The logits for the classes \( C, N \) and \( P \) are 38.0688, 10.1137 and 11.1361, respectively. The interpretability/ transparency of our model comes into play when an image is classified into a certain class. Our model is able to tell the reason of the classification of the image to that class, and the reason is that the image has some patches more similar to certain learned prototypes related to that class and it shows those learned prototypes. The learned prototypes
are projected from the training images, so they are actual patched of the training images.

2.2. Training of Quasi-ProtoPNet

Quasi-ProtoPNet uses the generalized version $d$ of the Euclidean distance function, and in this section we show that $d$ is a generalization of the Euclidean distance function. Consider Quasi-ProtoPNet with base model VGG-16. Let $x$ be an input image. Therefore, the shape of $L(x)$ is $512 \times 6 \times 6$ as described in Section 2.1. Let $p$ be any prototype with shape $512 \times h \times w$, where $1 \leq h, w \leq 6$, and $h$ and $w$ together are neither equal to 1 nor 6. The output $O(= L(x))$ of the convolutional layers $L$ has $(7-h)(7-w)$ patches of dimensions $h \times w$. Hence, square of the distance $d(P_{ij}, p)$ between $p$ and $(i,j)$ patch $P_{ij}$ (say) of $O$ is:

$$d^2(P_{ij}, p) = \sum_{l=1}^h \sum_{m=1}^w \sum_{k=1}^{512} \left| O(i+l-1)(j+m-1)k - p_{lmk} \right|^2. \quad (1)$$

Note that, if $p$ has prototypes of spatial dimensions $1 \times 1$, that is, $h = w = 1$, then $d^2(P_{ij}, p) = \sum_{k=1}^{512} \left| O_{ijk} - p_{11k} \right|^2$, which is the square of the Euclidean distance between $p$ and a patch of $O$, where $p_{11k} \approx p_k$. Therefore, the function $d$ is a generalization of the Euclidean distance function. The prototypical unit $p_t$ calculates the following.

$$p_t(O) = \max_{1 \leq i \leq 7-h, 1 \leq j \leq 7-w} \log \left( \frac{d^2(P_{ij}, p) + 1}{d^2(P_{ij}, p) + \epsilon} \right).$$

That is,

$$p_t(L(x)) = \max_{p \in \text{patches}(L(x))} \log \left( \frac{d^2(P, p) + 1}{d^2(P, p) + \epsilon} \right). \quad (2)$$

The Equation (2) exhibits that a prototype $p$ is more similar to the image $x$ if the reciprocal of the distance between $p$ and a latent patch of $x$ is smaller. Quasi-ProtoPNet is trained using the following two steps.

2.2.1. Optimization of all layers before the dense Layer

Let $X = \{x_1 \ldots x_n\}$ and $Y = \{y_1 \ldots y_n\}$ be sets of images and associated labels, respectively, and $D = \{(x_i, y_i) : x_i \in X, y_i \in Y\}$. Then our objective function is:

$$\min_{P, \text{learn}} \frac{1}{n} \sum_{i=1}^n \text{CrosEnt}(h \circ p_t \circ L(x_i), y_i) + \lambda \text{ClstCost}, \quad (3)$$

where ClstCost is given by the equation

$$\text{ClstCost} = \frac{1}{n} \sum_{i=1}^n \min_{j \in P_{ni}} \min_{P \in \text{patches}(L(x_i))} d^2(P, p_j). \quad (4)$$

The Equation (4) discloses that the drop in the cluster cost (ClstCost) leads to the clustering of prototypes around their respective classes. The reduction in cross entropy leads
to better classifications, see the objective function (3). The hyperparameters $\lambda$ is set equal to 0.7. Since $w_m$ is the weight matrix for the dense layer, $w_m^{(i,j)}$ is the weight assigned to the connection between logit of $i$th class and similarity score of $j$th prototype. Therefore, for a class $c$, we put $w_m^{(i,j)} = 1$ for all $j$ with $p_j^c \in P^c$, and for all $p_j^c \notin P^c$ with $c \neq i$, $w_m^{(c,j)} = 0$. The non-negativity of the distance function and optimization of all the layers before the last layer with optimizer SGD help Quasi-ProtoPNet to learn important latent spaces.

2.2.2. Projection of prototypes

Let $x$ be an input image. At the second step, Quasi-ProtoPNet projects the prototypes onto the patches of $x$ that are more similar to the prototypes. That is, a patch of $x$ that is at a smaller distance from a prototype gets projected, and the distance must be at least 93rd percentile of all the inverted distances of the prototype from all the images. For this purpose, Quasi-ProtoPNet makes the following update:

$$p_j^c \leftarrow \arg \min_{\{P: P \in \text{patches}(L(x_i)) \forall i \text{ such that } y_i = c\}} d(P, p_j^c).$$

2.3. Explanation of Quasi-ProtoPNet

In this section, we explain our model with an example of an input image as given in Figure 3.

![Figure 3: The explanation of the reasoning process of the model.](image)

In Figure 3, the image in the first column belongs to the class Covid. In the second column of the figure, the green rectangle on the image are enclosing the patches of the image.
that give the highest similarity score to the prototypes in the third column. In the fourth
column, the rectangles are enclosing the patches on the source images of the prototypes, that
is, the rectangles are pinpointing the patches on the source images from where the prototypes
are projected. In the fifth column, similarity scores between the prototypes and patches of
the test image are displayed. In the sixth column, the connections between similarity scores
and the logits are given. Since the image belongs to the first class $C$, the similarity scores
of the prototypes of the second and third class are assigned zero weight. The entries of the
seventh column are obtained by multiplying similarity scores and class connections, and the
logit (38.0688) for the class $C$ is obtained by adding the entries of the seventh column. The
logit for the class $C$ can also be computed by multiplying the first row of $w_m$ with matrix
$s_m$. The logit for the classes $N$ and $P$ are 10.1137 and 11.1361, respectively, and can be
computed by multiplying second and third row of $w_m$ with matrix $s_m$.

The transpose of the weight matrix $w_m$ and similarity score matrix $s_m$ that we obtain
from our experiments are as follows:

$$w_m^T = \begin{pmatrix} 1 & 0 & 0 \\ 1 & 0 & 0 \\ 1 & 0 & 0 \\ 1 & 0 & 0 \\ 1 & 0 & 0 \\ 1 & 0 & 0 \\ 1 & 0 & 0 \end{pmatrix} \begin{pmatrix} 2.8001 \\ 2.8001 \\ 2.8001 \\ 3.2050 \\ 2.8001 \\ 1.9548 \\ 2.4170 \end{pmatrix}$$

$$s_m = \begin{pmatrix} 2.8001 \\ 2.8001 \\ 2.8001 \\ 3.2050 \\ 2.8001 \\ 1.9548 \\ 2.4170 \\ 1.0003 \\ 1.1130 \\ 1.0463 \\ 1.1354 \\ 1.0233 \end{pmatrix}.$$
3. Results

In this section, we present the metrics given by our model and compare the performance of our model with the performance of the other models.

3.1. The metrics and confusion matrices

Suppose $TP$, $TN$, $FP$ and $FN$ denote the true positives, true negatives, false positives and false negatives for the Covid class. The metrics accuracy, precision, recall and F1-score are [49, 50, 51]:

\[
\text{Accuracy} = \frac{TP + TN}{TP + TN + FP + FN}, \quad \text{Precision} = \frac{TP}{TP + FP}, \quad \text{Recall} = \frac{TP}{TP + FN}, \quad \text{F1-score} = \frac{2 \times \text{Precision} \times \text{Recall}}{\text{Precision} + \text{Recall}}.
\]  

In Figures 4-9, the confusion matrices of Quasi-ProtoPNet with the base models are given. For example, in Figure 4, the confusion matrix $M$ (say) of Quasi-ProtoPNet with baseline VGG-16 is provided. Hence, the numbers $M[0][0]$, $M[1][1] + M[2][2]$, $M[0][1] + M[0][2]$ and $M[1][0] + M[2][0]$ are the $TP$, $TN$, $FP$ and $FN$ of the class Covid. Therefore, by Equations (5) and (6), for Quasi-ProtoPNet, the accuracy, precision, recall and F1-score are equal to 99.05, 0.98, 0.99 and 0.98, respectively.

3.2. The performance comparison of the models

The series of ProtoPNet models are constructed over the convolution layers of the base models. Although, the accuracies of the series of ProtoPNet models and the base models become stabilize prior to 35 epochs (see Section 3.4), but we trained and tested the models for 100 epochs.

The performance comparison in the metrics is provided in Table 1. We see from the third column of the Table 1 that when we build our model on the convolutional layers of VGG-16 then the accuracy, precision, recall and F1-score given by Quasi-ProtoPNet are 99.05, 0.98, 0.99 and 0.98, respectively. The accuracy, precision, recall and F1-score given by the models ProtoPNet,
Table 1: The comparison of performances of the models while experimented over the dataset of CT images

| Base      | Metric   | Quasi-ProtoPNet | Ps-ProtoPNet [41] | Gen-ProtoPNet [40] | NP-ProtoPNet [42] | ProtoPNet [6] | Base only |
|-----------|----------|-----------------|-------------------|-------------------|-------------------|---------------|-----------|
| VGG-16    | accuracy | 99.05           | 98.83             | 95.85             | 98.23             | 90.84         | 99.03     |
|           | precision| 0.98            | 0.96              | 0.93              | 0.95              | 0.93          | 0.98      |
|           | recall   | 0.99            | 0.98              | 0.95              | 0.94              | 0.91          | 0.99      |
|           | F1-score | 0.98            | 0.97              | 0.94              | 0.94              | 0.90          | 0.98      |
| VGG-19    | accuracy | 99.15           | 98.53             | 98.17             | 98.23             | 96.54         | 98.71     |
|           | precision| 0.98            | 0.97              | 0.95              | 0.91              | 0.93          | 0.98      |
|           | recall   | 0.99            | 0.99              | 0.99              | 0.96              | 0.95          | 0.99      |
|           | F1-score | 0.98            | 0.98              | 0.97              | 0.93              | 0.94          | 0.98      |
| ResNet-34 | accuracy | 99.29 ± 0.04    | 98.97 ± 0.12      | 98.40 ± 0.07      | 98.45 ± 0.06      | 97.05 ± 0.04  | 99.24 ± 0.10 |
|           | precision| 0.99            | 0.97              | 0.96              | 0.96              | 0.95          | 0.99      |
|           | recall   | 0.99            | 0.99              | 0.99              | 0.99              | 0.96          | 0.99      |
|           | F1-score | 0.99            | 0.98              | 0.97              | 0.97              | 0.96          | 0.99      |
| ResNet-152| accuracy | 99.26 ± 0.05    | 98.85 ± 0.09      | 99.90 ± 0.09      | 98.48 ± 0.06      | 88.20 ± 0.08  | 99.40 ± 0.05 |
|           | precision| 0.98            | 0.97              | 0.93              | 0.99              | 0.87          | 0.99      |
|           | recall   | 0.99            | 0.98              | 0.93              | 0.99              | 0.87          | 0.99      |
|           | F1-score | 0.98            | 0.97              | 0.93              | 0.99              | 0.87          | 0.99      |
| DenseNet-121 | accuracy | 99.44 ± 0.04  | 99.24 ± 0.02      | 99.97 ± 0.02      | 98.83 ± 0.02      | 98.48 ± 0.02  | 99.32 ± 0.03 |
|           | precision| 0.99            | 0.98              | 0.98              | 0.99              | 0.98          | 0.99      |
|           | recall   | 0.99            | 0.99              | 0.98              | 0.98              | 0.98          | 0.99      |
|           | F1-score | 0.99            | 0.98              | 0.98              | 0.98              | 0.98          | 0.99      |
| DenseNet-161 | accuracy | 99.37 ± 0.02  | 99.02 ± 0.02      | 98.87 ± 0.02      | 98.88 ± 0.02      | 98.76 ± 0.02  | 99.41 ± 0.07 |
|           | precision| 0.98            | 0.96              | 0.98              | 0.97              | 0.97          | 0.99      |
|           | recall   | 0.99            | 0.99              | 0.98              | 0.99              | 0.99          | 0.99      |
|           | F1-score | 0.99            | 0.97              | 0.98              | 0.97              | 0.98          | 0.99      |

NP-ProtoPNet, Gen-ProtoPNet, Ps-ProtoPNet with base model VGG-16, and the base model itself (Base only) are 90.84, 0.89, 0.91 and 0.90; 98.23, 0.93, 0.95 and 0.94; 95.85, 0.93, 0.95 and 0.94; 98.83, 0.96, 0.98 and 0.97; and 99.03, 0.98, 0.99 and 0.98, respectively. The highest accuracies obtained with different base models are in bold. Moreover, we see from the Table 1 that accuracies given by Quasi-ProtoPNet are even better than the accuracies given by the base models when Quasi-ProtoPNet is constructed over the convolutional layers of VGG-16, VGG-19 and DenseNet-121. Furthermore, the highest accuracy (99.48%) achieved by Quasi-ProtoPNet with base model DenseNet-121 is equal to the highest accuracy (99.48%) achieved by the non-interpretable model DenseNet-161.

In addition to achieving excellent accuracy, Quasi-ProtoPNet can explain why an input image is classified into a certain class, whereas such explanations are not possible with black-box models.
Table 2: The p-values obtained with the test of hypothesis for system of two proportions (accuracies) between our proposed model and each of the other models

| Base          | Ps-ProtoPNet [41] | Gen-ProtoPNet [40] | NP-ProtoPNet [42] | ProtoPNet [6] | Base only |
|---------------|-------------------|--------------------|-------------------|---------------|-----------|
| VGG-16        | 0.00755           | 0.00002            | 0.00002           | 0.00002       | 0.40905   |
| VGG-19        | 0.00002           | 0.00002            | 0.00002           | 0.00002       | 0.00002   |
| ResNet-34     | 0.00005           | 0.00002            | 0.00002           | 0.00002       | 0.44828   |
| ResNet-152    | 0.00002           | 0.00002            | 0.00002           | 0.00002       | 0.02169   |
| DenseNet-121  | 0.00480           | 0.00002            | 0.00002           | 0.00002       | 0.03836   |
| DenseNet-161  | 0.00002           | 0.00002            | 0.00002           | 0.00002       | 0.08692   |

That is, our model exhibits some prototypes from the image class that are similar to some patches of the classified image. In other words, if an image is classified to a certain class then it must have some patches similar to the prototypes of that class. The model also gives prototypes that can be manually compared with some patches of the classified image to know why a certain class has been assigned to the image.

3.3. The test of hypothesis for the accuracies

Since an accuracy is the proportion of correctly classified images among all the test images, the test of hypothesis concerning system of two proportions can be applied to determine whether the differences between the accuracies are statistical significant. Let \( n_d \) be the size of test dataset. Let \( x_1 \) and \( x_2 \) be the number of images correctly classified by models 1 and 2, respectively. Let \( \tilde{p}_1 = x_1/n_d \) and \( \tilde{p}_2 = x_2/n_d \). The statistic for the test concerning difference between two proportions (accuracies) is as follows [35]:

\[
Z = \frac{\tilde{p}_1 - \tilde{p}_2}{\sqrt{\tilde{p}(1 - \tilde{p})/n_d}}, \quad \text{where } \tilde{p} = (x_1 + x_2)/2n_d.
\]  

(7)

Suppose the models 1 and 2 give the accuracies \( p_1 \) and \( p_2 \). Then, our hypothesis:

- \( H_0 : (p_1 - p_2) = 0 \) (null hypothesis)
- \( H_a : (p_1 - p_2) \neq 0 \) (alternative hypothesis)

Let the level of confidence (\( \alpha \)) be 0.05. Therefore, to reject the null hypothesis, the \( p \)-value must be less than 0.025 because we have two-tailed hypothesis. Suppose \( p_1 \) represents the accuracy given by Quasi-ProtoPNet and the accuracies given by the other models are represented by \( p_2 \). The values of test statistic \( Z \) are obtained by the above formula, see Equation (7). We use the standard normal table to obtain the associated \( p \)-values, and list the \( p \)-values in the Table 2.

In particular, when convolutional layers of VGG-16 are used to construct the models, we get the \( p \)-values from the accuracy given by Quasi-ProtoPNet along with the accuracies given by Ps-ProtoPNet, Gen-ProtoPNet, NP-ProtoPNet, ProtoPNet and VGG-16 equal to 0.00755, 0.00002, 0.00002, 0.00002 and 0.40905, respectively. The null hypothesis for all the \( p \)-values that correspond to the series of ProtoPNet models got rejected, because the \( p \)-values are less than 0.025, see the Table 2. Therefore, the accuracies given by Quasi-ProtoPNet with different base models are statistically significantly (with 95% confidence) better than the accuracies given by the ProtoPNet
models. However, the p-values given in the last column of Table 2 corresponding to the base models VGG-16, ResNet-34, ResNet-152, DenseNet-121 and DenseNet-161 are greater than 0.025. So, the accuracies given by these base models are not significantly different from the accuracies given by our model.

Figure 10: Quasi-ProtoPNet with VGG-16.

Figure 11: Quasi-ProtoPNet with VGG-19.

Figure 12: Quasi-ProtoPNet with ResNet-34.

Figure 13: Quasi-ProtoPNet with ResNet-152.

Figure 14: Quasi-ProtoPNet with DenseNet-121.

Figure 15: Quasi-ProtoPNet with DenseNet-161.
3.4. The graphical comparison of the accuracies

In the Figures 10-15, graphical comparison of the accuracies given by Quasi-ProtoPNet and the other models is provided. Although, the accuracies given the models become stable before 35 epochs, the models are trained and tested for 100 epochs over the dataset [16], and the graphical comparisons of the accuracies are provided over 50 epochs.

The Figure 10 provides a comparison of the accuracies given by the models when they are constructed over the convolutional layers of VGG-16. Although, it is difficult to see the difference between the accuracies in the Figures 10-15, the difference is clear before the models stabilize.

3.5. The effect of the projection of prototypes

In this section, we prove a theorem similar to [6, Theorem 2.1]. The theorem [6, Theorem 2.1] assumes that the negative connections between similarity scores and incorrect classes can be made equal to zero during the training process. As mentioned in Section 1.1, our experiments show that it is hardly possible to make the negative connections zero during the training process. However, we do not need to make this assumption because our model uses only positive reasoning process, and the suspension of the convex optimization of the last layer of our model keeps the connection between similarity scores and incorrect classes zero. Furthermore, [6, Theorem 2.1] is proved with the Euclidean distance function, whereas our theorem is neither restricted to the Euclidean distance function nor to its generalized version \(d\), but the distance function can be replaced with any positive-valued function that satisfies the triangular inequality and has an appropriate domain. However, we present the theorem with a hemimetric, a distance function more general than the distance function \(d\).

Theorem 3.1. Let \(f\) be a hemimetric. Suppose \(f\) and the distance function \(d\) have the same domain, and \(f^2\) denotes the square of \(f\). Let \(h \circ p \circ L\) be a Quasi-ProtoPNet. For a class \(k\), let \(a^k_i\) and \(b^k_i\) be the values of \(i\)-th prototype for class \(k\) after the projection of \(p^k_i\) and before the projection of \(p^k_i\), respectively. Let \(x\) be an input image that is correctly classified by Quasi-ProtoPNet before the projection, and \(k\) be the correct class label of \(x\). Let \(O^k_i\) be the patch of \(L(x)\) closest to \(a^k_i\). Suppose there exists some \(\delta\) with \(0 < \delta < 1\) such that:

1. for all \(k' \neq k\) and \(i \in \{1, \ldots, m_k\}\), we have \(f(a^k_i, b^k_i) \leq \theta f(O^k_i, b^k_i) - \sqrt{\epsilon}\), where \(\theta = \min(\sqrt{1+\delta-1}, \frac{1}{\sqrt{2-\delta}})\) and \(\epsilon\) is given by \(p_i(L(x)) = \max_{p \in \text{patches}(L(x))} \log \left( \frac{f^2(P, p) + 1}{\epsilon} \right)\);

2. for all \(i \in \{1, \ldots, m_k\}\), we have \(f(a^k_i, b^k_i) \leq (\sqrt{1+\delta} - 1) f(O^k_i, b^k_i)\) and \(f(O^k_i, b^k_i) \leq \sqrt{1-\delta}\).

Then after projection,

1. the output logit \(\Delta_k\) (say) for the correct class \(k\) can decrease at most by \(m \log(1+\delta)(2-\delta)\), that is, \(\Delta_k \geq -m \log(1+\delta)(2-\delta)\);

2. the output logit \(\Delta_{k'}\) (say) for incorrect classes \(k'\) can increase at most by \(m \log(1+\delta)(2-\delta)\), that is, \(\Delta_{k'} \leq m \log(1+\delta)(2-\delta)\).

Proof. For any class \(c\), let \(G_c(x, \{p^c_i\}_{i=1}^m)\) be the output logit for input image \(x\), where \(\{p^c_i\}_{i=1}^m\) denote the prototypes of class \(c\). The connection between similarity score and incorrect classes is
Therefore,
\[ G_c(x, \{p_i^c\}_{i=1}^m) = \sum_{i=1}^m \log \left( \frac{f^2(O_i^c, p_i^c) + 1}{f^2(O_i^c, p_i^c) + \epsilon} \right). \]

Let \( \Delta_c \) be the difference between the output logit of class \( c \) after the projection and before the projection of prototypes. Suppose \( G_c(x, \{a_i^c\}_{i=1}^m) \) and \( G_c(x, \{b_i^c\}_{i=1}^m) \) denote the logits after the projection and before the projection, respectively. Therefore, we have
\[ \Delta_c = G_c(x, \{a_i^c\}_{i=1}^m) - G_c(x, \{b_i^c\}_{i=1}^m) = \sum_{i=1}^m \log \left( \frac{f^2(O_i^c, a_i^c) + 1}{f^2(O_i^c, a_i^c) + \epsilon} \right). \]

Assume,
\[ \Psi_i = \frac{f^2(O_i^c, a_i^c) + 1}{f^2(O_i^c, b_i^c) + 1} \times \frac{f^2(O_i^c, b_i^c) + \epsilon}{f^2(O_i^c, a_i^c) + \epsilon}. \]

Therefore,
\[ \Delta_c = \sum_{i=1}^m \log \Psi_i. \]

First, to prove 1, that is, to find the lower bound of \( \Delta_c \); assume \( c = k \) in the above equations (9) and (10), where \( k \) is the correct class of \( x \).

From the inequality given in the assumption 2, we have
\[ \frac{f^2(O_i^k, a_i^k) + 1}{f^2(O_i^k, b_i^k) + 1} \geq \frac{1}{f^2(O_i^k, b_i^k) + 1} \geq \frac{1}{2 - \delta}. \]

Using the triangular inequality, we have
\[ \frac{f^2(O_i^k, b_i^k) + \epsilon}{f^2(O_i^k, a_i^k) + \epsilon} \geq \frac{f^2(O_i^k, b_i^k) + \epsilon}{(f(O_i^k, b_i^k) + f(a_i^k, b_i^k))^2 + \epsilon}. \]

By the assumption 2, we have
\[ f(a_i^k, b_i^k) \leq (\sqrt{1 + \delta} - 1)f(O_i^k, b_i^k), \] that is,
\[ f(a_i^k, b_i^k) + f(O_i^k, b_i^k) \leq f(O_i^k, b_i^k)\sqrt{1 + \delta}. \]

Square inequality (13) and add \( \epsilon \) to the result, we obtain
\[ (f(a_i^k, b_i^k) + f(O_i^k, b_i^k))^2 + \epsilon \leq (1 + \delta)f^2(O_i^k, b_i^k) + \epsilon \leq (1 + \delta)(f^2(O_i^k, b_i^k) + \epsilon). \]

On rearranging inequality (14), we have
\[ \frac{f^2(O_i^k, b_i^k) + \epsilon}{(f(a_i^k, b_i^k) + f(O_i^k, b_i^k))^2 + \epsilon} \geq (1 + \delta). \]

By inequalities (12) and (15), we have
\[ \frac{f^2(O_i^k, b_i^k) + \epsilon}{f^2(O_i^k, a_i^k) + \epsilon} \geq (1 + \delta). \]
Therefore, by equations (11) and (16), we have
\[ \Psi_k^i = \frac{f^2(O_i^k, a^k_i) + 1}{f^2(O_i^k, b^k_i) + 1} \times \frac{f^2(O_i^k, b^k_i) + \epsilon}{f^2(O_i^k, a^k_i) + \epsilon} \geq \frac{1}{(1 + \delta)(2 - \delta)}. \]  
(17)

Hence, by the equations (8) and (17), we have
\[ \Delta_k \geq \sum_{i=1}^{m} \log \left( \frac{1}{(1 + \delta)(2 - \delta)} \right), \]  
that is, \( \Delta_k \geq -m \log(1 + \delta)(2 - \delta). \)

Second, to prove 2, that is, to find the upper bound of \( \Delta_k^c \), assume \( c = k' \) in the above equations (9) and (10), where \( k' \) is the incorrect class of \( x \).

By the triangle inequality,
\[ \frac{f^2(O_i^{k'}, a_k^{k'}) + 1}{f^2(O_i^{k'}, b_k^{k'}) + 1} \leq \frac{(f(O_i^{k'}, b_k^{k'}) + f(a_k^{k'}, b_k^{k'}))^2 + 1}{f^2(O_i^{k'}, b_k^{k'}) + 1}. \]  
(18)

The assumption 1 gives:
\[ f(a_k^{k'}, b_k^{k'}) \leq (\sqrt{1 + \delta} - 1)f(O_i^{k'}, b_k^{k'}) - \sqrt{\epsilon} \leq (\sqrt{1 + \delta} - 1)f(O_i^{k'}, b_k^{k'}). \]  
(19)

By the inequality (19), we have
\[ (f(O_i^{k'}, b_k^{k'}) + f(a_k^{k'}, b_k^{k'}))^2 \leq (f(O_i^{k'}, b_k^{k'}) + (\sqrt{1 + \delta} - 1)f(O_i^{k'}, b_k^{k'}))^2 \]
\[ = ((\sqrt{1 + \delta}f(O_i^{k'}, b_k^{k'}))^2 = (1 + \delta) f^2(O_i^{k'}, b_k^{k'}). \]
(20)

The inequality (20) gives:
\[ \frac{f(O_i^{k'}, b_k^{k'}) + f(a_k^{k'}, b_k^{k'}) + 1}{f(O_i^{k'}, b_k^{k'}) + 1} \leq \frac{(1 + \delta) f(O_i^{k'}, b_k^{k'}) + 1}{f(O_i^{k'}, b_k^{k'}) + 1} \]
\[ \leq \frac{(1 + \delta) f(O_i^{k'}, b_k^{k'}) + 1 + \delta}{f(O_i^{k'}, b_k^{k'}) + 1} = 1 + \delta. \]  
(21)

From the inequalities (18) and (21), we have
\[ \frac{f^2(O_i^{k'}, a_k^{k'}) + 1}{f^2(O_i^{k'}, b_k^{k'}) + 1} \leq 1 + \delta. \]  
(22)

Again, by the triangle inequality, we have
\[ f(O_i^{k'}, a_k^{k'}) \geq f(O_i^{k'}, b_k^{k'}) - f(a_k^{k'}, b_k^{k'}). \]  
(23)

The assumption 1 implies \( f(O_i^{k'}, b_k^{k'}) - f(a_k^{k'}, b_k^{k'}) > 0 \). Therefore, by the inequality (23), we have
\[ \frac{f^2(O_i^{k'}, b_k^{k'}) + \epsilon}{f^2(O_i^{k'}, a_k^{k'}) + \epsilon} \leq \frac{f^2(O_i^{k'}, b_k^{k'}) + \epsilon}{f(O_i^{k'}, b_k^{k'}) - f(a_k^{k'}, b_k^{k'}) + \epsilon} \leq \left( \frac{f(O_i^{k'}, b_k^{k'}) + \sqrt{\epsilon}}{f(O_i^{k'}, b_k^{k'}) - f(a_k^{k'}, b_k^{k'})} \right)^2. \]  
(24)
Again, by the assumption 1, we have
\[ f(a_k^n_i, b_k^n_i) \leq \left(1 - \frac{1}{\sqrt{2} - \delta}\right) f(O_k^n_i, b_k^n_i) - \sqrt{\epsilon}. \]

On simplifying the above inequality, we obtain
\[ \frac{1}{\sqrt{2} - \delta} f(O_k^n_i, b_k^n_i) + \sqrt{\epsilon} \leq f(O_k^n_i, b_k^n_i) - f(a_k^n_i, b_k^n_i). \]

Therefore,
\[ \frac{1}{\sqrt{2} - \delta} f(O_k^n_i, b_k^n_i) + \sqrt{\epsilon} \leq \frac{1}{\sqrt{2} - \delta} f(O_k^n_i, b_k^n_i) + \sqrt{\epsilon} \leq f(O_k^n_i, b_k^n_i) - f(a_k^n_i, b_k^n_i). \] (25)

By the inequality 25, we have
\[ \frac{f(O_k^n_i, b_k^n_i) + \sqrt{\epsilon}}{f(O_k^n_i, b_k^n_i)} \leq \frac{\sqrt{2} - \delta}{\sqrt{2} - \delta}. \] (26)

On combining the inequalities (24) and (26), we obtain
\[ \frac{f(O_k^n_i, b_k^n_i) + \epsilon}{f(O_k^n_i, a_k^n_i) + \epsilon} \leq (\sqrt{2} - \delta)^2 = 2 - \delta. \] (27)

On combining the inequalities (23) and (27), we have
\[ \Psi'_i = \frac{f^2(O_k^n_i, a_k^n_i) + 1}{f^2(O_k^n_i, b_k^n_i) + 1} \times \frac{f^2(O_k^n_i, b_k^n_i) + \epsilon}{f^2(O_k^n_i, a_k^n_i) + \epsilon} \leq (1 + \delta)(2 - \delta). \] (28)

Therefore, by equation 10, and inequality 28, we have
\[ \Delta'_i \leq \sum_{i=1}^{m} \log(1 + \delta)(2 - \delta) \leq m \log(1 + \delta)(2 - \delta). \] (29)

Hence, \( \Delta'_k \leq m \log(1 + \delta)(2 - \delta). \)

### 4. Limitations

As mentioned in the Section 1.1, Quasi-ProtoPNet gives better performance than the series of ProtoPNet models when classification is to be made over only a few classes. As the number of classes grows bigger, we cannot guarantee that the performance of our model will be better than the performance of ProtoPNet and Ps-ProtoPNet. However, there are many cases similar to the case of CT-scan images as discussed in this paper when we need to classify images over only a few classes. Therefore, our model can be really useful for such situations.
5. Discussion

Quasi-ProtoPNet model suspends the convex optimization of the last layer to keep the connections constant and it uses the objective function that accommodates only the positive reasoning process. Also, the suspension reduced the training time of our model. Quasi-ProtoPNet is closely related to the series of other ProtoPNet models, but strikingly different from them due to its reasoning process for the classifications. Quasi-ProtoPNet uses the positive reasoning process whereas other ProtoPNet models use the negative reasoning process along with the positive reasoning process that leads to decrease in their accuracy, especially when number of classes is small. In particular, our model can be useful during this pandemic when deadly mutants of coronavirus (e.g. omicron variant) are being identified.

6. Conclusions

The use of positive reasoning process along with the use of prototypes with rectangular spatial dimensions and square spatial dimensions helped our model to improve its performance over the series of the other ProtoPNet models. Moreover, as observed in Section 3.2, Quasi-ProtoPNet gives the highest accuracy (99.48%) when DenseNet-121 is used as the base model, and the highest accuracy given by Quasi-ProtoPNet is equal to the highest accuracy (99.48%) given by the non-interpretable model DenseNet-161.

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Highlights

Regina, Canada, December 6, 2021.

• Quasi-ProtoPNet uses the positive reasoning process that is inspired from reasoning process that an intelligent student uses to solve a multiple choice question.

• The performance of our model is on par with the performance of the state-of-the-art non-interpretable models.

• The theorem provides the evidence of better performance of our model, see Theorem 3.1.

• Quasi-ProtoPNet uses prototypes with both types of spatial dimensions, that is, rectangular spatial dimensions and square spatial dimensions.

• Our model keeps constant connections between similarity scores and logits. The suspension of the convex optimization of the last layer also reduces the training time considerably.

Sincerely yours,

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Think positive: An interpretable neural network for image recognition

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Abstract

The COVID-19 pandemic is an ongoing pandemic and is placing additional burden on healthcare systems around the world. Timely and effectively detecting the virus can help to reduce the spread of the disease. Although, RT-PCR is still a gold standard for COVID-19 testing, deep learning models to identify the virus from medical images can also be helpful in certain circumstances. However, the transparency/interpretability of the reasoning process of predictions made by such deep learning models is essential. In this paper, we propose an interpretable deep learning model that uses positive reasoning process to make predictions. We trained and tested our model over the dataset of chest CT-scan images of COVID-19 patients, normal people and pneumonia patients. Our model gives the accuracy, precision, recall and F-score equal to 99.48%, 0.99, 0.99 and 0.99, respectively.

Keywords: CT-scan, Prototypes, COVID-19, Pneumonia, Interpretable.

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Conflicts of Interest Statement

Regina, Canada, December 6, 2021.

The author declares no conflict of interest.

Sincerely yours,

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