

Hybrid Genetic U-Net Algorithm for Medical Segmentation

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Abstract: U-Net based architecture has become the de-facto standard approach for medical image segmentation in recent years. Many researchers have used the original U-Net as a skeleton for suggesting more advanced models such as UNet++ and UNet 3+. This paper seeks to boost the performance of the original U-Net via optimizing its hyperparameters. Rather than changing the architecture itself, we optimize hyperparameters which does not affect the architecture, but affects the performance of the model. For this purpose, we use genetic algorithms. Intensive experiments on medical dataset have been carried out which document a performance gain at a low computation cost. In addition, preliminary results reveal the benefit of the proposed framework for medical image segmentation.

1 INTRODUCTION

Image segmentation models have been gaining traction over the last years (Kheradmandi and Mehranfar, 2022; Iqbal et al., 2022; Lin et al., 2022b). Segmentation models are used in a variety of important fields not only in experimental settings but also in production environments as some of these models are now being applied to real-world applications such as autonomous driving (Wang et al., 2022), and remote sensing (Wu et al., 2022). One of the most notable fields is medical imaging (You et al., 2022; Kheradmandi and Mehranfar, 2022). In fact, these networks have become so useful that Bergen hospital in Norway has begun using them for tumor detection (E-Helse, 2019). The models are used as an assistance tool for doctors, yielding the probability of the patient having a tumor. Our goal is to optimize the U-Net model, which is a wildly used segmentation model. We seek to optimize the hyperparameters of the model using genetic algorithms, further increasing the performance of the model. The work reported in (Ronneberger et al., 2015) introduced U-Net in 2015, and since then, the model has been applied to several domains within deep learning computer vision. U-Net is a successful segmentation model with many successors in medical applications (Lin et al., 2022a). The successors use U-Net as a skeleton, but seek to further improve the model by making minor changes to

the architecture (Fang et al., 2022; Wu et al., 2019). However, none of those successors seeks to improve hyperparameters of the U-Net model itself. Deciding the value of the U-Net hyperparameters may seem arbitrary, as it is extremely difficult to assess the optimal value. In this research work, we propose hyperparameter optimization to enhance and improve the U-Net model by assessing the optimal hyperparameter values. It is an end-to-end framework which uses the genetic algorithm for guiding the training of the U-Net architecture. The main contributions of this research work can be given as follows:

1. We propose a new genetic algorithm which allows exploring the possible combination of the U-Net architecture.
2. We develop new crossover, and mutation operators which intelligently explore the solutions space of the different combinations of the hyperparameter optimization of the U-Net.
3. We test the proposed framework on large data for medical image segmentation. The initial results of the proposed framework are very promising.

The remaining of the paper is presented as follows. Section 2 presents the related work. Section 3 describes the image segmentation problem. Section 4 explains the main components of the proposed framework. Section 5 gives the experimental analysis part, while Section 6 concludes the paper.

2 RELATED WORK

Ultrasound image segmentation has the goal to identify the different labels in a given ultrasound image. In the context of deep learning, the aim is to design efficient models in order to learn the segmentation function. The input of the model is an ultrasound image, and the output will be the label of each pixel in that image. Huang et al. (Huang et al., 2020) proposed a machine learning method for breast ultrasound image segmentation in order to identify tumors. The ultrasound images are first cropped and pre-processed using bilateral filtering, histogram equalization, and pyramid mean shift filtering to remove noise. Simple linear iterative clustering is then performed for grouping the pixel of images into super-pixels. Features are extracted for each super-pixel, where two labels are created, the tumor label if the super-pixel contains a tumor, the normal label, otherwise. The kNN classifier is then performed to classify the pixels located to the super-pixels into tumor or normal. Adjacent tumor super-pixels are finally merged to segment the tumor of the new image. Amiri et al. (Amiri et al., 2020) proposed a two-stage ultrasound image segmentation approach for breast lesion detection. The first use of the U-Net model aims to detect the lesions. The second use of the U-Net model aims to segment the detected lesions. Lee et al. (Lee et al., 2020) introduced the use of channel attention mechanisms to improve CNN performance for breast cancer segmentation in an ultrasound image. Interdependencies of the channels of the image are trained by injecting the statistical feature of each channel features (mean of pixel values) on fully connected layers-based network. The output of this network with the input images are injected into CNN for the segmentation. Wu et al. (Wu et al., 2020) proposed the encoder-decoder deep learning model for thyroid nodule segmentation on ultrasound image data. It contains: i) dense block structure, where any two layers are connected. Batch Normalization is used in order to train this dense block. ii) Atrous spatial pyramid pooling is used for creating contextual multiscale information of input feature map. iii) Model size optimization for reducing the number of parameters learned, where a further 1×1 convolution operation is computed before each convolution layer. The semantic features are obtained from the contextual information, and injected into each layer of the decoder module. The hierarchical feature fusion is also performed to merge the feature maps of the blocks of the decoder module. Zeng et al. (Zeng et al., 2021) proposed a hybrid deep learning architecture for fetal ultrasound image segmentation. A combination of V-Net with attention

mechanism is carried out in order to reach the better accuracy of the segmentation results. To deal with large range of batch size, global normalization is used instead of batch normalization. A mixed loss function based on the dice similarity coefficient is developed in order to minimize the error ratio. Note that the dice similarity coefficient is determined by the intersection over the union of the ground truth, and the output of the network. Xue et al. (Xue et al., 2021) addressed three issues related to breast lesion ultrasound image segmentation, which are: in homogeneous intensity distributions inside the breast lesion region, ambiguous boundary due to similar appearance between lesion and non-lesion regions, and irregular breast lesion shapes. CNN is first used for multiscale feature maps generation. Each CNN layer is connected to a 1×1 convolutional layer with maxpooling operation for detecting the breast lesion boundaries. The features of all CNN layers are concatenated and combined with spatial-wise, channel-wise blocks for learning the correlation among the generated feature maps, and predict the output image. Liu et al. (Liu et al., 2021) proposed a hybrid deep learning algorithm for detecting prostate cancer using ultrasound images. Feature extraction is performed using the Sobel filter, the features are injected into a RCNN (Regional Convolution Neural Network) for ultrasound image segmentation. Ouahabi et al. (Ouahabi and Taleb-Ahmed, 2021) developed an encode-decoder deep learning model for thyroid segmentation. It adds a new layer which integrates the merits of dense connectivity, dilated convolutions for extracting relevant features, and dealing with varied-size regions, respectively.

The image segmentation models in particular U-Net based architecture require high number of hyper-parameters to be tuned. This research work develops an end-to-end framework based on genetic algorithm, and U-Net for improving the image segmentation process.

3 BACKGROUND ON IMAGE SEGMENTATION

An image is “segmented” or “partitioned” into various groups throughout the image segmentation process. For instance, image segmentation is used to distinguish the speaker from the background in the Zoom call functionality that lets you alter your background. This is but one use case for image segmentation in the real world. Face identification, video surveillance, object detection, medical imaging, and other fields can all benefit from image segmentation.

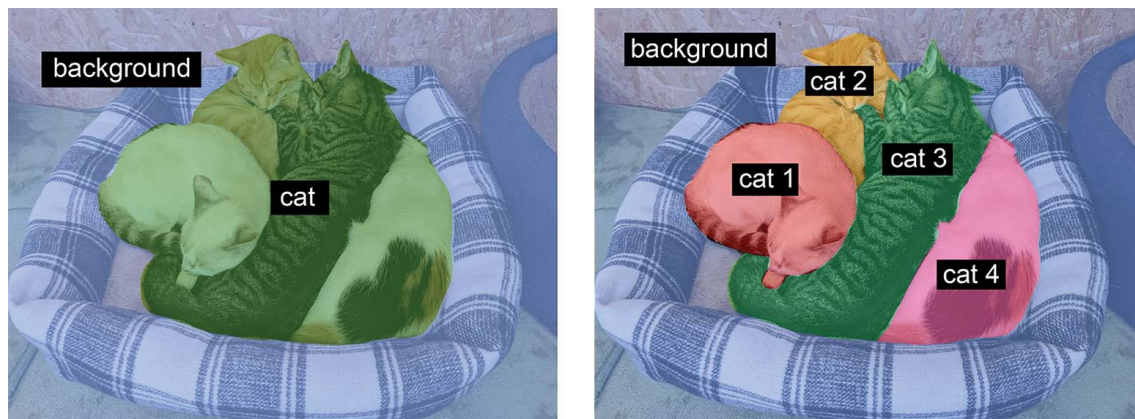


Figure 1: Semantic segmentation vs. instance segmentation (Chollet, 2021).

These applications use two-dimensional data in some cases and three-dimensional data in others.

There are two types of image segmentation: *semantic segmentation*, and *instance segmentation*.

- **Semantic segmentation**, where the objective is to assign a category to each pixel in an image. Semantic segmentation aims to classify each tree in a forest image into the appropriate category.
- **Instance segmentation**, This accomplishes the same thing as semantic segmentation but goes a step farther. An instance segmentation of the forest image would then separate the trees into tree 1, tree 2, and so on. Instance segmentation aims to separate items of the same category into a series.

Figure 1 illustrates the distinction between instance and semantic segmentation visually. The terms “semantic segmentation” and “image segmentation” will be used interchangeably. Object segmentation and detection share similarities. Finding the various object classes in a given image is the aim of object detection. The object is marked by object detection with a square frame represented by a bounding box. Object detection just displays the location of an object; it does not identify its shape. Object detection does not meet the criterion for several tasks. For instance, the form of the malignant cell is important when estimating the extent of the disease while trying to identify it.

4 PROPOSED FRAMEWORK

As illustrated in Figure 2, the goal is to find the optimal learning rate, number of epochs, and batch size for the U-Net architecture. All these hyperparameters have a direct impact on the performance of the model without changing the architecture. To optimize the

hyperparameters, we use genetic algorithms. Genetic algorithms (GA) consist of a set of population of individuals, each of which consists of genes. For our task, the genes are the hyperparameters; learning rate, number of epochs, and batch size. The individuals are then tested in the environment. Their genes, or hyperparameters, are applied to the model and the model produces a loss. The individuals are then granted fitness based on the loss, higher loss yields lower fitness, and vice versa. After each individual in the population is tested and has received their fitness, this generation is complete. The last step is for the current population of the next generation. To do so, parents are selected based on their fitness to reproduce, higher fitness yields a higher chance of becoming a parent; survival of the fittest. The next generation’s population is then repopulated by the set of parents, and the process begins anew. Optimally, the GA will converge toward a global optimum where most individuals consist of the optimal hyperparameter values for the model.

1. **Defining the Search Space:** The learning rate, the number of epochs, and the batch size have been specified as the components of the GA’s search space. However, the need to specify and explain the scope of the hyperparameters is still existing. Depending on the restrictions placed by the hyperparameter itself, the range may be quite arbitrary. For instance, the learning rate could only be set to between 0 and 1. It is somewhat wasteful to add such a high learning rate even when it is never utilized.
2. **Selection:** Additionally, we need to choose the individual who will make up the future generation. We applied the tournament-based strategy, which randomly chooses a group of individual from the population. The winner of the “tournament” is chosen to be a parent after the set competes in it. The winner of the competition will be the one

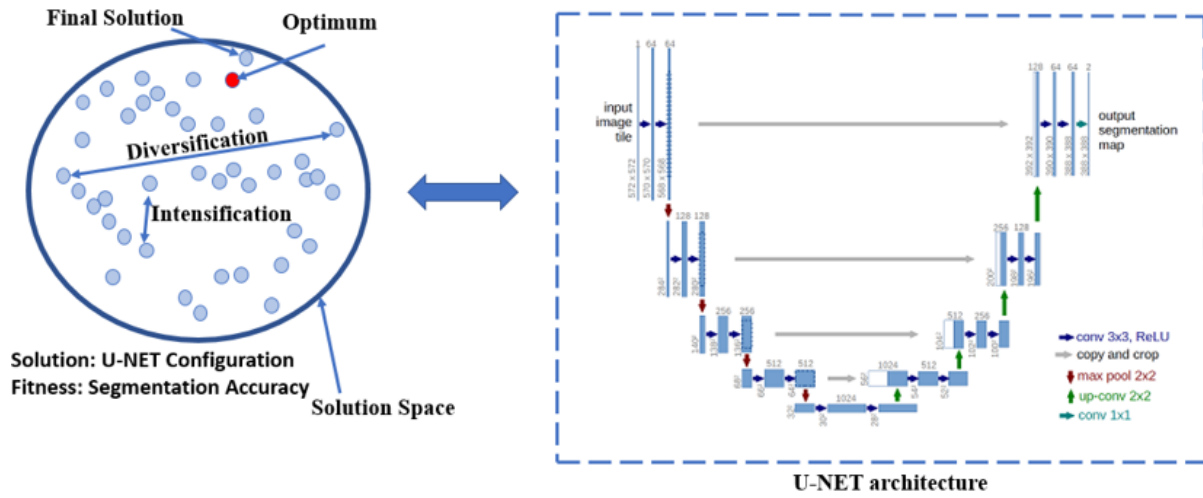


Figure 2: The proposed framework that combines both the genetic algorithm, and the U-Net architecture to improve the medical segmentation process (U-NET diagram is retrieved from (Ronneberger et al., 2015)).

with the highest level of fitness. Therefore, the outcome is somewhat predetermined. The competitors in the competition are not subjected to any new tests. The selection process for tournaments includes some built-in exclusions for the very lowest performers. For example, in a population of 100 individuals, where five individuals are competing, the four less performing individuals can never be chosen as parents, given these values. There is no tournament set in the populace where these four individuals can prevail. Figure 3 illustrates a selection tournament example that is applied to the suggested framework.

3. **Crossover:** The chosen parents must reproduce and give creation to new individuals after submitting an application for tournament selection and locating the set of parents. The new individual is made up of a combination of the DNA from the parents. We can transform the values to binary representations or use the genes as their values while executing crossover. Successful parents receive the crossover process, but it has the potential to alter the order of the many genes. Every possible gene combination is included in the search space, and the crossover process aids in the systematic exploration of the various combinations. Utilizing a binary format allows for even more profound investigation of the crossover process because it gives each gene's value the opportunity to be changed.
4. **Mutation:** There is a possibility that a newly produced individual will become mutated. The individual's genes are changed via mutation. By changing the gene's value, mutation might occur

to one or many genes. Within the limitations of the gene, alternation causes the gene to be randomized. New genes are added to the population through mutation. The modified individual bearing the new genes will be quickly wiped out of the population if these genes are bad, that is, if they score a poor fitness. If the genes are sound, however, the GA will continue to use the newly discovered genes and procreate the population.

5 PERFORMANCE EVALUATION

Table 1: The table contains the best individual after 50 generations, given the different mutation rates.

Mutation Rate	IoU Performance
5%	0.7274
10%	0.7273
15%	0.7465
20%	0.7278
25%	0.7289
30%	0.7366
35%	0.7274
40%	0.7276
45%	0.7272

We apply our solution to the ultrasound nerve dataset. As with most medical imaging datasets, the data is imbalanced. The dataset contains 5600 images where all the images are labeled. We use 90% of images to train a new model with each individual hyperparameters, and we use 10% of images for testing the model. The evaluation of the proposed framework is calculated using the Intersection-over-Union (IoU), (equa-

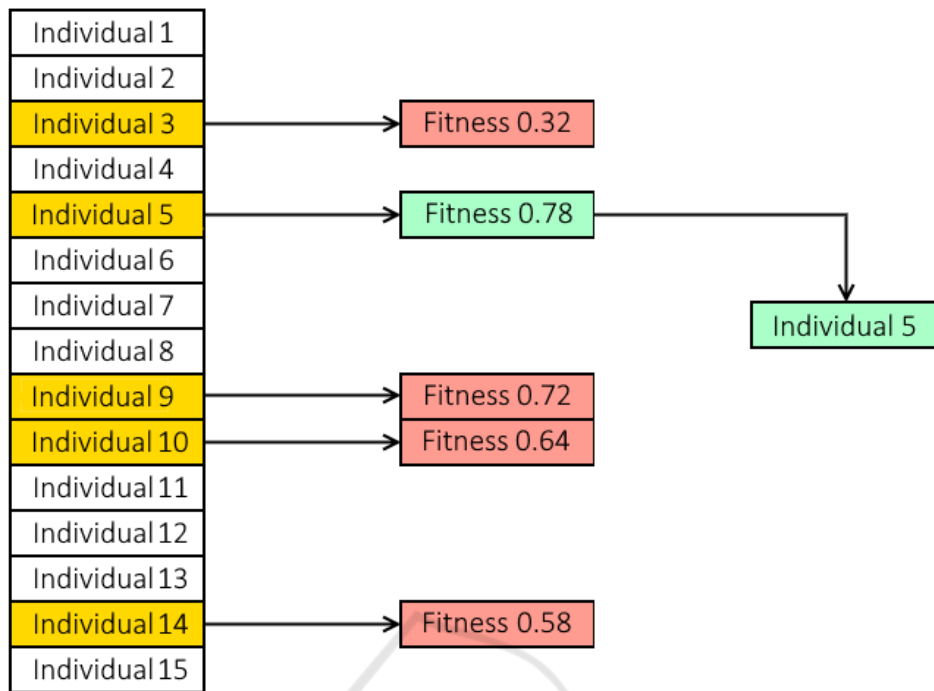


Figure 3: A population containing 15 individuals where tournament selection is applied. At random, 5 individuals are randomly chosen to compete in the tournament. The winner of the tournament is based on the predetermined fitness of the individual. In this case, individual 5 wins the tournament and is selected to be a parent.

Table 2: Comparison of the proposed solution with UNet algorithm.

% Images	UNet	Proposed Solution
10%	0.7104	0.7239
20%	0.7329	0.7692
30%	0.7567	0.7985
50%	0.7859	0.8003
80%	0.7971	0.8120
100%	0.8001	0.8431

tion 1).

$$IoU(U, V) = \frac{|U \cap V|}{|U \cup V|} \quad (1)$$

IoU is a method of measuring the overlapping labels. It measures the overlapping true and false labels, then divides it by the union of the labels. As expected, this heavily punishes the model by falsely predicting a nerve in the wrong spot. But in addition, it also punishes the model if it were to only predict false labels (no nerve) for every input. Encouraging the model to actually find the nerves and not stall the learning. We run a set of short tests to evaluate the optimal mutation rate. These tests consist of a training set of 100 images from the nerve dataset, and the epoch range is set between 0 and 1. Due to the low amount of data used, and the low number of epochs, we evaluate from the training loss, and not the validation loss. We

run these tests to find the mutation rate we want to use and to see if there are any discrepancies between the different mutation rates. Table 1 shows the different range of mutation rates and the performance. The performance is calculated by the dice loss functions. Regardless of the mutation rate, there is no large different in the performance. The 15% mutation rate is slightly above the rest, this may be random. Regardless, we will use 15% mutation rate for our initial experiment. In the following experiments, we will use the hyperparameters of the top individual found by the genetic algorithm. Table 2 compares the results of the proposed solution with the UNet algorithm. By varying the number of images from 10% to 100%, the proposed solution outperforms the UNet algorithm in terms of IoU. For instance, when training 100% of images, the IoU of UNet is only 0.8001%, where the IoU of the proposed solution is 0.8431%. These results are achieved thanks to the hyperparameter optimization technique used to find the best parameters of the UNet model. The last experiment aims to visualize some results of the developed model. Figure 4 shows three images; the input, the hand annotated nerve cell, and the predicted segmentation. The results indicate a low gap between the ground truth and the predicted segmentation of the proposed model. These results are achieved thanks to the strategy used in the segmentation process, where an efficient hyperparameter

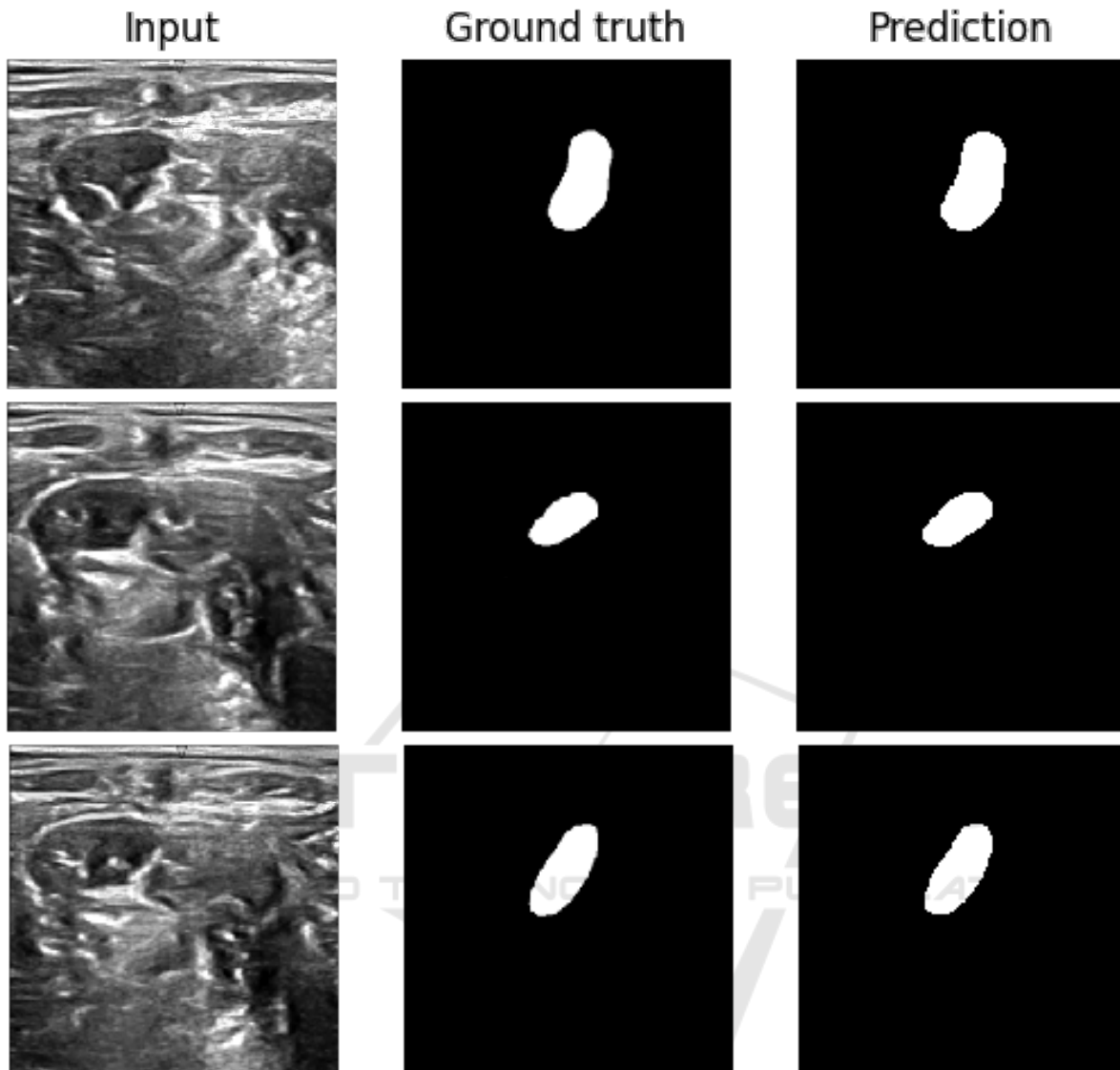


Figure 4: Ultrasound images of the neck as input on the left side, center shows the hand annotated nerve, right side shows the prediction by the model. These are some of the best results.

optimization is running to retrieve the optimal parameters of the designed model. These results demonstrate the applicability of the developed model in real settings to help the practitioners and doctors for medical decision-making.

6 CONCLUSION

Although U-Net model shows a great behavior for solving segmentation problem in medical applications, some limitations remain unsolved. In this paper, we solved the hyperparameter optimization is-

sue by developing an end-to-end intelligent framework which combines the genetic algorithm with the U-Net architecture to achieve the optimal accuracy in training complex medical data. We used genetic algorithms to optimize the hyperparameters of the UNet architecture for medical segmentation. The results reveal the superiority of the developed model compared to the UNet model. As future perspective, we aim to explore other evolutionary algorithms such as particle swarm optimization, Ant Colony, and mimetic algorithm in order to speed the convergence to the optimum. Exploring evolving learning, and in particular the NEAT optimization, is also in our future agenda.

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