

An Optimized Solution: Deep Residual Learning for ECG Signal Classification

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Abstract

Time matters the most in medical diagnosis. Therefore, ECG has been extensively used in diagnosis and precise identification of several cardiac conditions. As continuation to the previous work, which concerned about feature extraction and classification of ECG signals applying CNN on distinct datasets, this paper presents residual learning model that makes training of networks easier and substantially deeper since the deeper neural networks are difficult to train. The proposed solution is classified into two types, one with no residual connection (Basic CNN) and the other analyzing the repetition of residual connections (Residual CNN) for displaying the variation in performance level of each model. The proposed model offers extensive observational proof indicating that residual model is an optimized framework that shows better results of accuracy from considerably increased depth. From the performance of the (Deep Residual CNN) we also see that we can better exploit these residual connections by making the model much deeper. Such a model can be conveniently used for constant monitoring of ECG in real time environment.

Keywords

Basic CNN, Residual CNN, Convolution Layer, Max Pool Block

Introduction

Deep Learning, to know the technique for cardiac arrhythmia (17 training) detection primarily depends on lengthy-length electrocardiography (ECG) sign analysis. Prevention of Cardiovascular disorder is major role of any medical device as about 50 million humans are liable to coronary heart disorder within the international. Despite the fact that automatic analysis of the ECG sign may be very famous, present-day strategies are not super. research is based totally on 1000 ECG sign fragments from the MIT - BIH Arrhythmia database for one lead (MLII) from 45 ladies and men. a way based totally on the assessment of 10-s ECG signal fragments (now not a single QRS complicated) is applied (on common, thirteen instances fewer classifications/evaluation). A whole quit-to-cease shape changed into designed in place of the hand-crafted characteristic extraction and choice applied in conventional strategies. Deep 1D-CNN performed a popularity widespread accuracy of 17 cardiac arrhythmia problems (lessons) at a stage of 91.33% and category time according to single9 pattern of 0.015 s.

In spite of the fact that convolutional neural organizations (CNNs) might be utilized to order electrocardiogram (ECG) beats in the examination most recent cardiovascular disorder, ECG alarms are commonly prepared as one-dimensional alerts at the same time as CNN's are better perfect to a multidimensional pattern or photograph reputation packages. on this have a look at, the morphology and rhythm modern-day heartbeats are fused into a dimensional data vector for subsequent processing via CNNs that encompass adaptive modern-day charge and biased dropout methods. The outcomes exhibit that the CNN model is effective for detecting abnormal heartbeats or arrhythmias via computerized feature extraction. When the version became examined at the MIT-BIH arrhythmia database & PTBD database, the version did better performance than different state-of-the-art methods for five and eight heartbeat categories (the common accuracy become 99.1% and 97%). especially, the system shown higher performance in terms modern-day the sensitivity and wonderful predictive rate for V beats by extra than 4.3% and 5.4%, respectively, and also for S beats by using extra than 22.6% and 25.9%, respectively, while compared to current algorithms. it's far predicted that the technique might be appropriate for implementation on portable devices for the e-domestic health monitoring contemporary cardiovascular disease. In this paper, we propose Deep Residual CNN which might be the optimal solution when compared to Vaseline CNN and Residual CNN in terms of Accuracy, F1 score and other basic parameters with respect to datasets individually.

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Background and Related Work

Class of Electrocardiogram (ECG) alerts performs a enormous role within the identity of the functioning of the coronary heart.[1] This painting pertains with the ECG signals, wherein the classifier is evolved for identity of normal or odd situations of the heart. The uncooked ECG signals are gathered from an internet database (www.physioNet.org) for category. The raw ECG signal is pre-processed for noise elimination, and the frequency spectrum is analyzed to evaluate uncooked and denoised ECG sign. Attributes (P, Q, R, S, T time periods) from denoised ECG signal is analyzed and categorized the usage of Convolution Neural network (CNN). The paper reports a type method to distinguish ECG signals from the MIT-BIH database (arrhythmia database, arrhythmia p-wave annotations, atrial traumatic inflammation). The CNN analyses the deviation among nominal tiers of attributes (amplitude and time c language) and classifies between the abnormality and regular ECG wave. This painting offers an easy technique for deciphering ECG related situation for the clinician and helps clinical practitioners to make diagnostic selections.

The danger of tampering exists for classic patron recognition strategies based totally mostly on biometrics collectively with face and fingerprint.[2] Presently, research on patron reputation using biometric signs and symptoms which incorporates electrocardiogram (ECG), electroencephalogram (EEG), and electromyogram (EMG) has been actively carried out to triumph over this hassle. We herein suggest a purchaser popularity method utilizing a deep learning approach based totally mostly on ensemble networks after transforming ECG signs into -dimensional (2d) pix. A preprocessing approach for one-dimensional ECG indicators is carried out to remove noise or distortion; in the end, they're projected onto a second picture region and transformed into photo records. For the set of regulations, they designed deep mastering-based ensemble networks to decorate the degraded performance arising from overfitting in a single community. Our experimental consequences showcase that the ensemble networks display off an accuracy that is 1.7% better than that of the unmarried community. Mainly, the overall performance of the ensemble networks is as a good deal as 13% better in comparison to the unmarried network that degrades the popularity rate via manner of displaying similar abilities between training. The electrocardiogram (ECG) has been considerably used in the diagnosis of a coronary heart sickness which

The electrocardiogram (ECO) has been considerably used in the diagnosis of a coronary heart stekness which incorporates arrhythmia because of its simplicity and non-invasive nature.[3] Arrhythmia can be classified into many sorts, collectively with existence-threatening and non-lifestyles-threatening. Accurate detection of arrhythmic kinds can effectively prevent coronary heart sickness and reduce mortality. Strategies on this observe, specific deep learning approach for type of cardiac arrhythmia regular with the deep residual network (ResNet) is presented. We developed a 31-layer one-dimensional (1D) residual convolutional neural network. The set of rules consists of 4 residual blocks, every of which includes 3 1D convolution layers, 3 batch normalization (BP) layers, 3 rectified linear unit (ReLU) layers, and an "identification shortcut connections" form. Similarly, we suggest the use of 2-lead ECG signs in combination with deep learning techniques to routinely pick out five particular styles of heartbeats. They've acquired a median accuracy, sensitivity and incredible predictivity of 99.06%, 93.21% and 96.76% respectively for single-lead ECG heartbeats. Within the 2-lead datasets, the consequences display that the deep ResNet version has excessive classification average performance, accomplishing an accuracy of 99.38%, the sensitivity of 94.54%, and specificity of 98.14%.

ECG class is vital to the diagnosis of present-day cutting-edge cardiovascular sickness.[4] This paper develops a robust and accurate set state-of-the-art rules for computerized detection contemporary coronary heart arrhythmias from ECG indicators recorded with one lead. A unique version based totally on the convolutional neural community is proposed to extract low-level and immoderate-level talents state-of-the-art brief-term ECG. Similarly, statistics-Theoretic Metric today's is applied as a very last type version to enhance the discrimination capabilities brand new the community trained capabilities. The experimental consequences over the MIT-BIH arrhythmia database display that the model achieves a comparable performance with the maximum present-day the state-of-the-art strategies and information-Theoretic Metric brand new modern similarly enhance the general performance. Besides the best accuracy finished, the proposed method balances unique standards.

Atrial traumatic irritation is the maximum commonplace persistent shape of arrhythmia.[5] A way primarily based totally on wavelet remodel combined with a deep convolutional neural network is applied for the automated type of electrocardiograms. For the reason that ECG sign is without difficulty inferred, the ECG sign is decomposed into 9 varieties of sub-indicators with unique frequency scales through using wavelet function, and then wavelet reconstruction is performed after segmented filtering to remove the have an impact on of noise. A 24-layer convolution neural network is used to extract the hierarchical abilities through convolution kernels of different sizes, and in the end, the SoftMax classifier is used to classify them. The approach of the ECG information set furnished by way of the 2017 PhysioNet/CINC venture. After cross-validation, this technique can achieve 87.1% accuracy and the F1 score is 86.46%. Properly timed prediction of cardiovascular illnesses with the assist of a pc-aided diagnosis device minimizes the mortality rate of cardiac ailment sufferers. Cardiac arrhythmia detection is one of the toughest obligations, because of the reality the versions of electrocardiogram (ECG) sign are very small, which cannot be detected via human eyes.



Figure 1: A Flowchart of ECG Signal Preprocessing Level

Methodology & Proposed Solution Baseline CNN

This model has no residual connections and includes five primary blocks, a final convolution and then a totally related component. each main block consists of 2 Convolutional layers and a MaxPool block. Then it is fed into a completely linked portion of the community that finally ends up with a softmax layer for prediction.

For MIT-I	BIH dataset	t:	For	For PTBDB dataset:			
Layer (type)	Output Shape	Param #	Layer (type)	Output Shape	Param #		
input_1 (InputLayer)	[(None, 187, 1)]	0	input_1 (InputLayer)	[(None, 187, 1)]	0		
conv1d (Conv1D)	(None, 183, 16)	96	convld (Conv1D)	(None, 183, 16)	96		
convld_1 (Conv1D)	(None, 179, 16)	1296	convid_1 (ConviD)	(None, 179, 16)	1296		
max_pooling1d (MaxPooling1D)	(None, 89, 16)	0	max_pooling1d (MaxPooling1D)	(None, 89, 16)	0		
iropout (Dropout)	(None, 89, 16)	0	dropout (Dropout)	(None, 89, 16)	0		
conv1d_2 (Conv1D)	(None, 87, 32)	1568	convld_2 (ConvlD)	(None, 87, 32)	1568		
conv1d_3 (Conv1D)	(None, 85, 32)	3104	convid_3 (ConviD)	(None, 85, 32)	3104		
max_pooling1d_1 (MaxPooling1	(None, 42, 32)	9	max_pooling1d_1 (MaxPooling1	(None, 42, 32)	0		
ropout_1 (Dropout)	(None, 42, 32)	0	dropout_1 (Dropout)	(None, 42, 32)	0		
onv1d_4 (Conv1D)	(None, 40, 32)	3104	convld_4 (ConvlD)	(None, 40, 32)	3104		
onv1d_5 (Conv1D)	(None, 38, 32)	3104	convid_5 (ConviD)	(None, 38, 32)	3104		
max_pooling1d_2 (MaxPooling1	(None, 19, 32)	0	<pre>max_pooling1d_2 (MaxPooling1</pre>	(None, 19, 32)	0		
Iropout_2 (Dropout)	(None, 19, 32)	0	dropout_2 (Dropout)	(None, 19, 32)	0		
conv1d_6 (Conv1D)	(None, 17, 256)	24832	convld_6 (Conv1D)	(None, 17, 256)	24832		
inal_conv (Conv1D)	(None, 15, 256)	196864	<pre>final_conv (Conv1D)</pre>	(None, 15, 256)	196864		
lobal_max_pooling1d (Global	(None, 256)	0	global_max_pooling1d (Global	(None, 256)	0		
ropout_3 (Dropout)	(None, 256)	0	dropout_3 (Dropout)	(None, 256)	0		
lense_1 (Dense)	(None, 64)	16448	dense_1 (Dense)	(None, 64)	16448		
lense_2 (Dense)	(None, 64)	4160	dense_2 (Dense)	(None, 64)	4160		
lense_3_mitbih (Dense)	(None, 5)	325	dense_3_ptbdb (Dense)	(None, 1)	65		
Fotal params: 254,901 Frainable params: 254,901 Vonstrainable params: 0			Total params: 254,641 Trainable params: 254,641 Non-trainable params: 0				

Figure 2: Model for Baseline CNN MIT-BIH dataset Figure 3: Model for Baseline CNN PTBDB dataset

Residual CNN

A residual block has a 3 x 3 convolution layer accompanied via a batch normalization layer and a ReLU activation feature. that is again continued by means of a three x three convolution layer and a batch normalization layer. The pass connection basically skips each those layers and adds immediately before the ReLU activation characteristic. Such residual blocks are repeated to shape a residual network.

For MIT-BIH dataset:			
aven (Even)	Outout Shaoa	00000 B	Connected to
put 1 (InputLayer)	[(None, 187, 1)]	0	
wy1d (Conv10)	(None, 183, 32)	192	input 1(8)(8)
nyld 1 (ConylD)	(None, 183, 32)	5152	
invid 2 (ConviD)	(None, 183, 32)		convid 1[8][8]
			conv1d[0][0]
tivation (Activation)	(None, 183, 32)		
x nonlineld (MaxPonlinelD)	(None, 91, 32)		activation(0)(0)
nyld 3 (Conv1D)	(None, 91, 32)	5152	max poplingid[9][9]
1v1d 4 (Conv10)	(None, 91, 82)		convid 3[8][8]
			max_pooling1d[0][0]
testion 1 (Activation)	(None 91 32)		Contro_4[0][0]
<pre>x ecoliog1d 1 (MaxBooliog10)</pre>	(1000 45 23)		
avid 5 (Convin)	(None, 45, 32)		any poplingid ligital
wid 6 (Convin)	(10000 45 22)	6163	
1_2 (Add)	(None, 45, 32)	0	max_pooling1d_1[0][0]
			convia_e[e][e]
ivation_2 (Activation)	(None, 45, 32)		700 ⁻ 5[6][6]
x_pooling1d_2 (MaxPooling1D)	(None, 22, 32)		activation_2[8][8]
ivid_7 (convib)	(None, 22, 32)	5152	max_poolingid_2[u][u]
nv1d_8 (Conv1D)	(None, 22, 32)	5152	conv1d_7[0][0]
4_3 (Add)	(None, 22, 32)		max_pooling1d_2[0][0] conv1d_8[0][0]
tivation_3 (Activation)	(None, 22, 32)		add_3[0][0]
<pre>x_pooling1d_3 (MaxPooling1D)</pre>	(None, 11, 32)		activation_3[0][0]
nv1d_9 (Conv1D)	(None, 11, 32)	\$152	max_poolingid_3[0][0]
mv1d_10 (Conv10)	(None, 11, 32)	5152	conv1d_9[0][0]
d_4 (Add)	(None, 11, 32)		max_poolingid_3[0][0]
tivation_4 (Activation)	(None, 11, 32)	6	add_4[0][0]
x_pooling1d_4 (MaxPooling1D)	(None, 5, 32)		activation_4[0][0]
nse_3_mitbih (Dense)	(None, 5)		dense_2[0][0]
stal params: 282,853 rainable params: 282,853 on-trainable params: 0			

Figure 4: Model for Residual CNN MIT-BIH dataset Figure 5: Model for Residual CNN PTBDB dataset

Deep Residual CNN

Currently, convolutional neural network (CNN)-based totally methods accomplished superb overall performance in ECG signals denoising. significantly, CNN with deeper and thinner structures is bendier to obtain the ECG sign info. but, direct stacking a few current networks is hard to gain satisfactory denoising performance. the principle structure brand new DRCNN is the residual block comprises of two convolutional layers, made of skip connections without batch normalization operation. The input ECG signal were transferred to the hidden layers by the less effective pass connection without any delay. Although it takes part in reducing the route length contemporary gradient transfer allowing the gradient switch to mitigate the vanishing gradient issue through a brief route. When compared to the various state of art algorithms, DRCNN shows effective progress in denoising based on the experimental consequences tested. We calculate f1 score, accuracy score, AUROC score, AUPRC score and get the Confusion Matrix of DRCNN and show that it's better than Baseline CNN and normal Residual Learning.

The model relies on repeated 1D convolutions to build lower dimensional representations of the output. The structure's main feature is the repetition of residual blocks. These are made up of two Convolutional layers, a residual connection from the input to the block, ReLU and Maxpooling.

The dimensionality reduction is carried out using MaxPool layers, and each Conv layer maintains the same number of filters (32), which is somewhat unusual.

The residual networks aid in backpropagation of the gradient, however, this model is perhaps not quite deep enough to warrant these, as shown by the performance of the Baseline CNN, which is essentially this model without the residual connections.

From the performance of the Deep Residual CNN, we also see that we can better exploit these residual connections by making the model much deeper.

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For MIT-BIH Dataset:

For PTBDB Dataset:



Figure 6: Model for Deep Residual CNN MIT-BIH Dataset

Figure 7: Model for Deep Residual CNN PTBDB Dataset

Results and Analysis Baseline CNN For MIT-BIH dataset:

> Test f1 score : 0.9173432846885803 Test accuracy score : 0.9850173579389732 Confusion Matrix : [[18033 52 24 3 6] [101 444 10 0 1] [48 2 1380 13 5] [22 0 20 120 0] [18 0 3 0 1587]]

For PTBDB dataset:

Test f1 score : 0.9902820573595639 Test accuracy score : 0.9859154929577465 AUC-ROC score : 0.9859154929577465 AUPRC score : 0.9925274848085369 Confusion Matrix : [[781 28] [13 2089]]

```
Residual CNN
For MIT-BIH dataset:
```

Test	f1	score	: 0.92	230	023698	36939288	
Test	acc	uracy	score	:	0.987	1185821304587	7
[[180	45	35	28		4	6]	
[85	461	9		0	1]	
[32	1	1396		16	3]	
[30	0	13		119	0]	
[18	0	1		0	1589]]	

For PTBDB dataset:

Test f1 score : 0.9832263269055602 Test accuracy score : 0.9866025420817588 AUROC score : 0.9866025420817588 AUPRC score : 0.9929940444947872 Confusion Matrix : [[783 26] [13 2089]]

Deep Residual CNN For MIT-BIH dataset:

> Test f1 score : 0.9222344281630622 Test accuracy score : 0.9877124063584871 Confusion Matrix : [[18066 20 20 6 6] [98 445 11 1 1] 27 4 1397 17 [3] 23 1 15 122 1] [12 0 3 0 1593]] [

For PTBDB dataset:

Test f1 score : 0.9892641926193407 Test accuracy score : 0.9914118859498454 AUROC score : 0.9914118859498454 AUPRC score : 0.9954446209642036 Confusion Matrix: [[792 17] [8 2094]]

Conclusion and Future Work

The proposed work includes evaluation of Baseline CNN, Deep Residual CNN and calculated f1 score (weighted average of Precision and Recall) dataset accuracy score, AUC-ROC score, AUPRC score & Confusion Matrix and checked the performance variation in each for MIT-BIH & PTBDB datasets respectively. For future scope, we can check the variation in performance level for RNN (Recurrent Neural Network) and DR-RNN (Deep Residual Recurrent Neural Network).

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