# IMPROVISED NATURE INSPIRED DEEP BELIEF NETWORK & ITS APPLICATION IN HEART DISEASE PREDICTION

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UNDER THE GUIDANCE OF PROF. ANUPAM SHUKLA & DR. RITU TIWARI PRESENTED BY:

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- Contents
- Introduction
- Motivation
- Background
- 6 Aim
- 6 Objective
- Methodology
- Results and Discussions
- Onclusions and Future Scope

# Introduction

- **Earlier**, the use of computer was to build knowledge from medical experts.
- This time consuming process depends on medical experts' subjective opinions.
- Machine learning techniques have been developed to gain knowledge automatically from raw data.
- Here, an improvised deep belief network is presented for the diagnosis of heart disease.

# DEEP LEARNING

- Training data through multiple neuron layers.
- It allows to solve **complex** problems.
- This creates multiple layers of abstraction.
- Deep learning achieves desired results in many application— Example: Recognition of complex patterns by using many layers-first layer can recognize simple edges, second layer can recognize the figures like rectangle and triangle and higher layers can be used to recognize complex shapes.
- Here, deep belief network, a part of deep learning is used

# DEEP BELIEF NETWORK

- DBN = RBM + NN
- Multi-layer belief network
- Multiple RBM layers stacked together to form DBN
- Topmost layer is a perceptron layer which classifies the data which have come through RBM layers
- Greedy layer-wise training in which most optimized trained value of one hidden layer is passed on to next layer

# **RBM**

- Boltzmann machine are bidirectional graph used to model the probability distribution
- Graph is bipartite
- Consists of hidden layer and visible layer with weight distribution between them
- extract features and reconstruct input

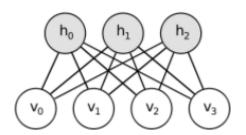


Figure: Structure of RBM<sup>1</sup>



# PROBABILITY DISTRIBUTION BY RBM

### STEP 1:

**Energy function**  $E(v,h)^2$  defining energy distribution for the network is:

$$E(v,h) = h'Wv + b'v + c'h$$
 (1)

h,v: activations of hidden and visible layer

b,c: biases of hidden and visible layer

W: weight matrix of RBM

### STEP 2:

Joint probability distribution of RBM:

$$p(\mathbf{v}, \mathbf{h}) = \frac{1}{N} e^{E(\mathbf{v}, \mathbf{h})} \tag{2}$$

<sup>&</sup>lt;sup>2</sup>Bengio, Y., Lamblin, P., Popovici, D., Larochelle, H. et al.: 2007, Greedy layer-wise training of deep networks, Advances in neuralinformation processing systems 19, 153

# STEP 3:

# Conditional probability distribution of RBM

The conditional probabilities associated with the joint probability distribution are denoted by Q(h|v) and P(v|h).

$$P(v_k = 1|h) = sigm(-b_k - \sum_j (W_{jk} * h_j))$$
 (3)

$$Q(h_j = 1|v) = sigm(-c_j - \sum_k (W_{jk} * v_k))$$
 (4)

Using above, P and Q sampling on RBM unit is carried out.



# STEP 4 AND 5:

**STEP 4:** We calculate the log likelihood of the input observations

$$log(P(v_0)) = log(\sum_{h} P(v_0, h)) = log(\sum_{h} (e^-E(v_0, h))) - log(\sum_{h, v} (e^-E(v, h)))$$

Here L is the learning rate

**STEP 5:** By applying gradient descent we minimize the negative log likelihood of  $v_0$ 

# Training of DBN

- 1 X be a feature **input matrix**
- 2 Training of RBM on X to obtain is done to have its weight matrix W
- 3 X is **transformed** by RBM and a new data X' is formed.
- 4 Repeating this procedure with X ← X' for the next pair of layers till the two top layers of network are reached.
- 5 **Fine-tuning** of all the parameters of this deep architecture with **supervised training**.

# PARTICLE SWARM OPTIMISATION

- PSO<sup>3</sup> was introduced by James Kennedy and Russel Eberhart in 1995
- Used for optimising a non-linear function
- Based on information sharing between the agents in the search space.
- Velocity and position of particle:

$$U_i(k+1) = U_i(k) + c_1 r_1(P_i - X_i(k)) + c_2 r_2(P_g - X_i(k))$$
 (6)

$$X_i(k+1) = X_i(k) + U_i(k+1)$$
 (7)

<sup>&</sup>lt;sup>3</sup>Eberhart, R. C., Kennedy, J. et al.: 1995, A new optimizer using particle swarm theory, Proceedings of the sixth international symposium on micro machine and human science, Vol. 1, New York, NY, pp. 3943

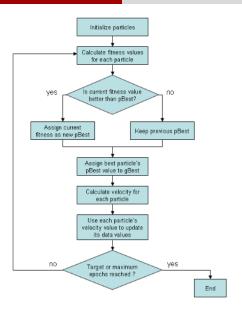


Figure: Flowchart of PSO

# **MOTIVATION**

- Clinical decisions are often taken on the basis of doctors perception and experience rather than on the knowledge rich data masked in the database. Every doctor is not evenly expert in every sub specialty and they are in several places a scarce resource.
- Therefore, automatic, precise, accurate and fast classification of medical heart disease data is today's need.

# BACKGROUND

- Data mining UCI repository dataset<sup>4</sup>
- Paper<sup>5</sup> describes deep belief nets as probabilistic generative models that are composed of multiple layers of stochastic latent variables (also called "feature detectors" or "hidden units").

(MNIST dataset , CD , SVM , Limitations , large amount of time)

(WIVIST dataset, CD, SVW, Elimitations, large amount of time

<sup>&</sup>lt;sup>4</sup>eart data set, U.: uci, Uci heart data set

<sup>&</sup>lt;sup>5</sup>Hinton, G.: 2011, Deep belief nets, Encyclopedia of Machine Learning, Springer, pp.

- In this paper<sup>6</sup>. PSO is compared with the other algorithms and it was found that it needs fewer parameters. The paper concludes with the fact that more research can be made on PSO by improving the topology of particle swarm, by blending with other intelligent optimization algorithm.
- The best accuracy achieved on:

Cleveland: 90% using SVM+GA [9] (Bhatia,2008)

Hungarian: 92% using k-NN [12] (Setiawan, 2009)

<sup>&</sup>lt;sup>6</sup>Bai, Q.: 2010, Analysis of particle swarm optimization algorithm, Computer andinformation science3(1), 180

Table 1 PSO publications and citations by year

Year	Google Scholar Hits	IEEE Xplore Publications
1995	<b>(</b> 28)	(2)
1996	<b>(</b> 14)	(0)
1997	<b>■</b> (24)	(3)
1998	$\blacksquare (49)$	<b>(</b> 4)
1999	$\blacksquare (62)$	<b>(</b> 7)
2000	$\blacksquare (78)$	<b>(</b> 8)
2001	$\blacksquare (118)$	<b>(11)</b>
2002	(256)	<b>(</b> 46)
2003	(373)	(82)
2004	(850)	(181)
2005	(1210)	(279)
2006	(1230)	(462)

<sup>&</sup>lt;sup>7</sup>Kennedy, J.: 2011, Particle swarm optimization, Encyclopedia of machine learning, and

# RESEARCH GAP

- There are many hyper parameters to tune like number of layers, number of hidden units, learning rate and many more. The eciency of function is determined by these parameters therefore it is necessary to optimize all these parameters to get accurate results
- To optimize all these input parameters its takes a considerable amount of time. Therefore there is a need of some algorithm which can optimize these parameters in limited time giving better performance than simple DBN.

# PROBLEM STATEMENT

- On time diagnosis of heart disease is necessary for patient from a cardiologist in an ideal situation. In case of emergencies we need some fast and efficient alternatives.
- The large number of parameters required by deep belief network like regularization and dimensions of neuron and RBM layer have a non-linear dependency which causes difficulty in training it.

# AIM

 To construct a classifier using deep belief network and further optimize it by nature inspired particle swarm optimization

# **OBJECTIVE**

- To predict heart disease risk using deep belief network optimized using PSO
- **Comparison** of results with existing works done on the chosen heart disease dataset

# Data set description

# UCI repository dataset<sup>8</sup>

Cleveland clinical foundation Database

303 records of patients

Each record has 13 attributes

The output is either 0,1,2,3 or 4

1,2,3 or 4: patient has heart disease(based on severity 1-4)

0 : normal person

Hungarian Database

It has 294 records. Each record has 13 attributes. The output to each record is either 0 or 1. 0 indicates normal person and 1 indicates abnormal person.

<sup>&</sup>lt;sup>8</sup>Lichman, M.: 2013, UCI machine learning repository.http:archive.ics.uci.edu/ml

### Methodology

Attribute	Description	Value Description
age	Age	Numerical
sex	Sex	1 if male, 0 if female
ср	Chest pain type	1 to 4
trestbps	Resting systolic blood	Numerical
	pressure	
chol	Serum chole-	Numerical
	strol(mg/dl)	
fbs	Fasting blood sugar	1 if yes and 0 if no
restecg	Resting cardigraphic	0 to 2
	results	
thalach	Maximum heart rate	Numerical
	achieved	
exang	Exercise induced	1 if yes and 0 if no
	angina	
oldpeak	ST depression induced	Numerical
	by excercise relative to	
	rest	
slope	The slope of the peak	1 upsloping and 2 flat
	exercise ST segment	
ca	Number of major ves-	Numerical
	sels	
thal		3,6 or 7
num	Output Attribute	0 or 1

Figure: Summary of disease attributes<sup>9</sup>

# PSEUDO CODE OF DBN

```
RBMupdate(v[0], epsilon, W, b, c)
```

```
for all hidden units i
2.
         compute Q(h[0][i]=1|v[0])# sigmoid(b[i]+sum_i(W[i][i]*v[0][i]))
3.
        sample h[0][i] from Q(h[0][i]=1|v[0])
4.
     for all visible units j:
5.
        compute P(v[1][i] = 1 | h[0]) \# sigmoid(c[i] + sum_i(W[i][i])^* h[0][i]))
6.
        sample v[1][j] from P(v[1][j] = 1|h[0])
7.
     for all hidden units i:
8.
        compute Q(h[1][i] = 1|v[1])#sigmoid(b[i]+sum_i(W[i][i]*v[1][i]))
     W += epsilon * (h[0] * v[0]' - Q(h[1][.] = 1 | v[1]) * v[1]')
9.
      a += epsilon * (h[0] - Q(h[1][.] = 1 | v[1]))
10.
      b += epsilon * (v[0] - v[1])
11.
```

# PreTrainSupervisedDBN(Z, epsilon, L, n, W, b, V, c)

```
let X the marginal over the input part of Z
1.
2.
    initialize b[0]=0
3.
    for l=1 to L-1
        initialize W[i]=0, b[i]=0
4.
5.
        while not stopping criterion:
6.
           sample g[0]=x from X
7.
           for i=1 to l-1:
8.
              sample g[i] from Q(g[i]|g[i-1])
9.
        RBMupdate(g[l-1], epsilon, W[l], b[l], b[l-1])
10.
      initialize W[L]=0, b[L]=0, V=0, c=0
11.
      while not stopping criterion:
12.
         sample (g[0]=x, y) from Z
13.
         for i=1 to I-1:
14.
            sample g[i] from Q(g[i]|g[i-1])
         RBMupdate((g[L-1],y), epsilon, (W[L],V'), b[L], (b[L-1],c))
15.
```

# DBNSupervisedFineTuning(Z, C, epsilon\_C, L, n, W, b, V,c)

Define the network output function f(x) = V \* x + c where x is the data transformed through rbm layers

Iteratively minimize the expected value of C(f(x),y)

for pairs (x,y) sampled from Z by tuning parameters W, b, V, c

This can be done by stochastic gradient descent with learning rate epsilon\_C,

using appropriate stopping criterion such as early stopping on validation set

# TrainSupervisedDBN(Z, C, epsilon\_CD, epsilon\_C, L, n, W, b, V):

- 1. let X the marginal over the input part of Z
- 2. PreTrainSupervisedDBN(X, epsilon\_CD, L, n, W, b)
- 3. DBNSupervisedFineTuning(Z, C, epsilon\_C, L, n, W, b, V, c

### Deep Belief Network

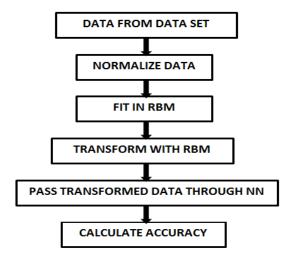


Figure: Deep belief network structure

# PSEUDO CODE FOR PSO

```
For each particle
        Initialize particle
3.
     Do until maximum iterations or minimum error criteria
4.
        For each particle
5.
            Calculate Data fitness value
6.
            If the fitness value is better than pBest
7.
               Set pBest = current fitness value
8.
           if pbest is better than gbest
9.
               Set\ gBest = pBest
10.
         For each particle
11.
                Calculate particle velocity
12.
```

Use gbest and Velocity to update particle data

# PSEUDO CODE FOR PROPOSED ALGORITHM

# ModifiedTrainSupervisedDBN (Z, C, epsilon\_CD, epsilon\_C, L, n, W, b, V)

- 1. let X the marginal over the input part of Z
- 2. PreTrainSupervisedDBN (X, epsilon\_CD, L, n, W, b)
- 3. DBNSupervisedFineTuning (Z, C, epsilon\_C, L, n, W, b, V, c)
- 4. Calculate accuracy of DBN with a testing data using W
- 5. return accuracy

# PSEUDO CODE FOR PROPOSED ALGORITHM

```
\label{lem:modifiedTrainSupervisedDBNwithPSO} ModifiedTrainSupervisedDBNwithPSO(Z,C,epsilonCD,epsilonC,L,W,b,v,range\_n,swarmsize,iter)
```

```
1. let vector s denote the swarm position of the particles
```

- 2. vector v denotes the velocity vector of the particles
- 3.  $initialize global_best = -infinity$
- 4. initialize personal\_best = -infinity
- 5. for i<swarmsize
- 6. initialize s[i] in range\_n
- 7. increment i
- 8. while (iterations < iter)
- 9. for i<swarmsize
- 10.  $fitness\_val = modifiedTrainSupervisedDBN(Z, C, epsilonCD, epsilonC, L, s[i], W, b, V)$

# PSEUDO CODE FOR PROPOSED ALGORITHM

```
11.
             if fitness_val>particle_best
12.
                set particle_best = fitness_val
13.
              if particle_best > global_best
14.
                set global_best=particle_best
15.
             for i < swarmsize
16.
                calculate the particle velocity for the particle i
17.
                update s[i] according to v[i]
18.
                use global best to update particle position
19.
          return global_best that is the best dimension for the deep belief
        network
```

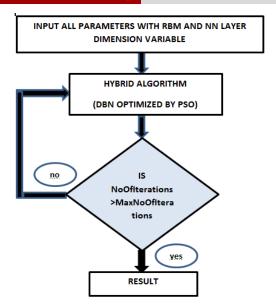


Figure: Proposed hybrid algorithm

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# RESULTS

Table: Comparison between existing & proposed algo-Cleveland

Year	Algorithm	Accuracy
2000[13]	Active SVM	87%
2008[9]	SVM+GA	90%
2009[12]	FDSS	87%
Proposed	DBN	88%
Proposed	DBN+PSO	91.18%

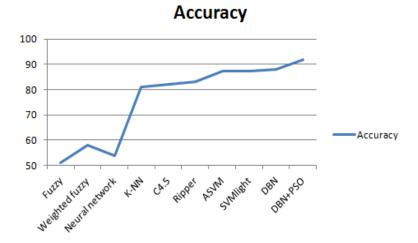


Figure: Accuracy of different algorithms on cleveland dataset

# RESULTS

# Table: Comparison between existing & proposed algo-HUNGARIAN

Algorithm	Accuracy
Fuzzy[11]	36
NN[13]	46.417
Weighted Fuzzy [13]	50.83
MLP-ANN [12]	54
C4.5 [12]	66
Ripper [12]	68
K-NN [12]	92
Deep belief network	91.33
DBN+PSO(proposed Stategy)	94.88

# RESULTS

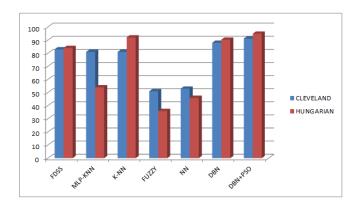


Figure: Comparison between algorithms

# **DISCUSSIONS**

- Comparison of proposed strategy with DBN
- Comparison of proposed strategy with existing algorithms applied on the same data set
- The algorithm used i.e. PSO for optimizing deep belief network is giving better results than simple deep belief network algorithm.
- The proposed algorithm gave better results than work done on the existing algorithm
- This is because we have taken into account the optimization of variables in our hybrid algorithm .

# Conclusions

- Machine learning algorithms like DBN can be used to make the diagnosis fast and less costly.
- The proposed strategy of hybrid DBN algorithm with PSO successfully classified the heart disease type from the input data set provided.
- Mathematically simple[10] PSO can optimize multiple parameters of DBN at a time giving the most efficient results.
- It has been predicted in a very accurate manner whether a person has heart disease or not reducing the dependency on traditional techniques.

# FUTURE SCOPE

- (i) This work can be extended by using other nature inspired algorithms like firefly, Ant colony Optimization(ACO), Flower pollination.
- (ii) Also the system can be made for real time data. This will then take data in real time as well as predict or classify in real time whether a person has the disease or not.
- (iii) Larger datasets can be managed using big data concepts.

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