

# Evolutionary $q$ -Gaussian Radial Basis Functions for Binary-Classification

F. Fernández-Navarro, C. Hervás-Martínez,  
Pedro A. Gutiérrez, M. Cruz-Ramírez and M. Carbonero-Ruz  
pagutierrez@uco.es

Department of Computer Science and Numerical Analysis  
AYRNA Research Group  
University of Córdoba, Spain.



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# Introduction

Different types of neural networks used for classification purposes.

- **Projection** basis functions:
  - Multilayer perceptron neural networks (MLP).
  - Product Unit Neural Networks (PUNNs).
- **Kernel** basis functions:
  - Radial Basis Function (RBF) neural networks.

## Shortcomings of the standard RBF

When dimensionality grows and/or when data is concentrated in boundaries of the  $K$  dimensional space, standard Gaussian basis function lacks its performance.

## Our Proposal

Alternative  $q$ -Gaussian Radial Basis Neural Network obtained by a Hybrid Algorithm (HA).

# Radial Basis Functions Neural Networks

The model of a RBFNN can be described with the following equation (binary classification or regression):

$$f(\mathbf{x}) = \beta_0 + \sum_{i=1}^m \beta_i \cdot \phi_i(d_i(\mathbf{x})) \quad (1)$$

The function  $d_i(\mathbf{x})$  is defined as:

$$d_i(\mathbf{x}) = \frac{\|\mathbf{x} - \mathbf{c}_i\|^2}{r_i^2}, 1 \leq i \leq m \quad (2)$$

# Radial Basis Functions Neural Networks

## Standard RBF (SRBF)

$$\phi_i(d_i(\mathbf{x})) = e^{-d_i(\mathbf{x})} \quad (3)$$

- Very selective response, with high activation for patterns close to the centroid and very small activation for distant patterns.

# Radial Basis Functions Neural Networks

## Cauchy RBF (CRBF)

$$\phi_i(d_i(\mathbf{x})) = \frac{1}{1 + d_i(\mathbf{x})} \quad (4)$$

## Inverse Multiquadratic RBF (IMRBF)

$$\phi_i(d_i(\mathbf{x})) = \frac{1}{(1 + d_i(\mathbf{x}))^{\frac{1}{2}}} \quad (5)$$

- The CRBF and IMRBF have longer tails than the SRBF.  
→ Activation for patterns distant to the centroid of the RBF is bigger than the activation of the SRBF for those patterns.

# $q$ -Gaussian RBFs

## $q$ -Gaussian RBF

$$\phi_i(d_i(\mathbf{x})) = \begin{cases} (1 - (1 - q)d_i(\mathbf{x}))^{\frac{1}{1-q}} & \text{if } (1 - (1 - q)d_i(\mathbf{x})) \geq 0; \\ 0 & \text{Otherwise} \end{cases} \quad (6)$$

- The  $q$ -Gaussian can reproduce different RBFs for different values of the real parameter  $q$ .
  - $q \rightarrow 2$ ;  $q$ -Gaussian  $\equiv$  CRBF.
  - $q \rightarrow 3$ ;  $q$ -Gaussian  $\equiv$  IMRBF.
  - $q \rightarrow 1$ ;  $q$ -Gaussian  $\equiv$  SRBF.
- A small change in the value of  $q$  represents a smooth modification on the shape of the RBF.

# $q$ -Gaussian RBFs

Radial unit activation in one-dimensional space with  $c = 0$  and  $r = 1$  for different RBFs: (a) Gaussian, Cauchy and Inverse Multiquadratic and (b)  $q$ -Gaussian with different values of  $q$

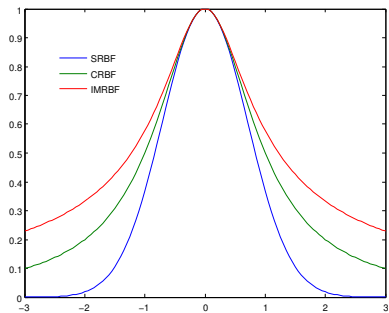


Figure: (a) Alternative RBFs

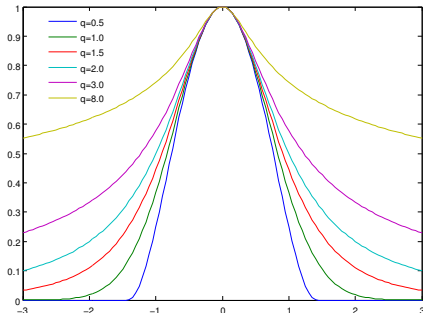


Figure: (b)  $q$ -Gaussian RBF



# $q$ -Gaussian RBFs for binary classification

- **Probabilistic** framework:
  - The activation function of each output node is the softmax function.

$$g(\mathbf{x}) = \frac{\exp f(\mathbf{x})}{1 + \exp f(\mathbf{x})} \quad (7)$$

- It is possible to evaluate the model using the cross-entropy error function, given by:

$$l(g) = -\frac{1}{N} \sum_{n=1}^N [y_n \log f(\mathbf{x}_n) + (1 - y_n) \log(1 - f(\mathbf{x}_n))] \quad (8)$$

# Hybrid Algorithm

- 1: **Hybrid Algorithm:**
- 2: Generate a random population of size  $N$
- 3: **repeat**
- 4:   Calculate the fitness of every individual in the population
- 5:   Rank the individuals with respect to their fitness
- 6:   The best individual is copied into the new population
- 7:   The best 10% of population individuals are replicated and they substitute the worst 10% of individuals
- 8:   Apply parametric mutation to the best  $(p_m)\%$  of individuals
- 9:   Apply structural mutation to the remaining  $(100 - p_m)\%$  of individuals
- 10: **until** the stopping criterion is fulfilled
- 11: Apply *iRprop+* to the best solution obtained by the EA in the last generation.

Figure: Hybrid Algorithm (HA) framework

# Hybrid Algorithm (Main characteristics I)

- **Error and Fitness Functions.**
  - $l(g)$  as the error function.
  - $A(g) = \frac{1}{1+l(g)}$ , where  $0 < A(g) \leq 1$  as the fitness measure.
- **Initialization of the Population.** 5, 000 random RBFNNs:
  - $k$ -means algorithm for different values of  $k$ ,  $k \in [M_{min}, M_{max}]$ , where  $M_{min}$  and  $M_{max}$  are parameters of the algorithm.
  - Widths ( $r_i$ ) of the RBFNNs  $\rightarrow$  geometric mean of the distance to the two nearest neighbours.
  - $q_i \rightarrow$  values close to 1 (SRBF).
  - Then we select the best 500 RBFNNs, and we evolve them.

## Hybrid Algorithm (Main characteristics II)

- **Structural Mutation.** There are four different structural mutations: hidden node addition, hidden node deletion, connection addition and connection deletion. If the structural mutator adds a new node in the RBFNN, the  $q$  parameter is assigned to a value in the interval  $[0.75, 1.25]$ .
- **Parametric Mutation.** Centre, Radii and  $q$  Mutation and Output-to-Hidden Node Connection Mutations → adding a Gaussian noise.
- **iRprop+.** We have carried out the adaptation of the *iRprop+* local improvement procedure to the softmax activation function and the cross-entropy error function.

# Experimental Design

- Eleven binary classification datasets taken from the UCI repository.
- The performance of each method has been evaluated using the correct classification rate ( $C$ ) in the generalization set.
- The experimental design was conducted using a 10-fold cross-validation procedure, with 10 repetitions per each fold.
- Comparison of the results obtained to:
  - SRBF.
  - CRBF.
  - IMRBF.

# Statistical Results (Mean $\pm$ Standard Deviation)

	Method( $C_G(\%)$ )			
	SRBF	CRBF	IMRBF	$q$ -Gaussian
Labor	91.33 $\pm$ 12.09	<b>95.00 <math>\pm</math> 11.24</b>	91.66 $\pm$ 8.78	<i>93.33 <math>\pm</math> 11.65</i>
Promoters	75.54 $\pm$ 13.56	80.18 $\pm$ 6.66	<i>81.09 <math>\pm</math> 8.69</i>	<b>84.00 <math>\pm</math> 6.15</b>
Hepatitis	<b>86.33 <math>\pm</math> 8.09</b>	83.16 $\pm$ 7.15	85.12 $\pm$ 7.52	<i>85.30 <math>\pm</math> 7.54</i>
Sonar	<b>78.38 <math>\pm</math> 9.03</b>	74.09 $\pm$ 10.20	76.02 $\pm$ 11.16	<i>76.04 <math>\pm</math> 13.56</i>
Heart	81.85 $\pm$ 8.97	83.70 $\pm$ 8.76	<b>84.81 <math>\pm</math> 8.45</b>	<i>84.07 <math>\pm</math> 7.20</i>
BreastC	72.04 $\pm$ 6.39	71.35 $\pm$ 8.00	<b>73.10 <math>\pm</math> 6.39</b>	<i>73.06 <math>\pm</math> 6.77</i>
Heart-C	85.44 $\pm$ 3.83	85.45 $\pm$ 5.59	<i>85.77 <math>\pm</math> 3.05</i>	<b>85.79 <math>\pm</math> 5.20</b>
Liver	<i>68.41 <math>\pm</math> 5.15</i>	65.23 $\pm$ 8.23	65.52 $\pm$ 6.31	<b>71.30 <math>\pm</math> 6.50</b>
Vote	<b>96.32 <math>\pm</math> 3.97</b>	95.39 $\pm$ 3.59	94.94 $\pm$ 2.36	<i>96.08 <math>\pm</math> 3.45</i>
Card	86.08 $\pm$ 3.14	<i>86.52 <math>\pm</math> 3.55</i>	85.94 $\pm$ 3.80	<b>87.87 <math>\pm</math> 0.37</b>
German	74.80 $\pm$ 3.82	<i>74.90 <math>\pm</math> 3.17</i>	74.40 $\pm$ 2.50	<b>75.25 <math>\pm</math> 2.98</b>
$\overline{C_G}(\%)$	81.50	81.36	<i>81.67</i>	<b>82.91</b>
$\overline{R}$	2.72	2.99	2.72	<b>1.54</b>
$p$ -Value	<b>0.03</b>	<b>0.00</b>	<b>0.03</b>	-
$\alpha'_{Hommel}$	0.10	0.03	0.05	-

The best result is in bold face and the second best result in italics

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# Conclusions

## Conclusions

- The models proposed,  $q$ -Gaussian Radial Basis Functions as transfer functions, are a viable alternative for obtaining more accurate binary classifications.
- These models have been designed with a HA constructed specifically for taking into account the characteristics of this kernel model.

## Future research

- To study other alternative RBFs (Generalized RBFs).
- To consider multi-class problems.

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