



Optimal Hyperbox shrinking in Dendritic Computing applied to Alzheimer's Disease detection in MRI

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Content

- Introduction
- Dendritic Computing
- Experimental results
- Summary and Conclusions

Introduction

Dendritic Computing

- simple and fast
- based on biology
- binary class problems
- based on lattice theory

Introduction

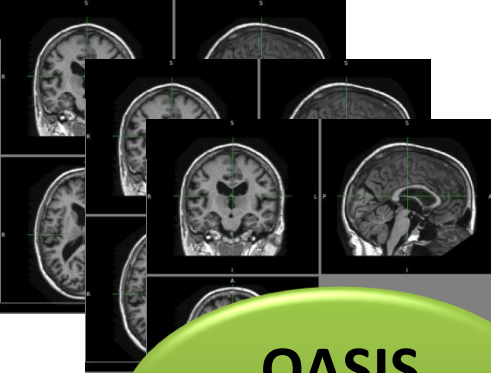
Has been proved to compute a **perfect** approximation to **any data distribution**.

The results of cross-validation experiments give very poor performance: **high sensitivity and very low specificity**.

We attribute this to the fact that the **DC learning algorithm** always tries to **guarantee** the good classification of the class 1 samples.



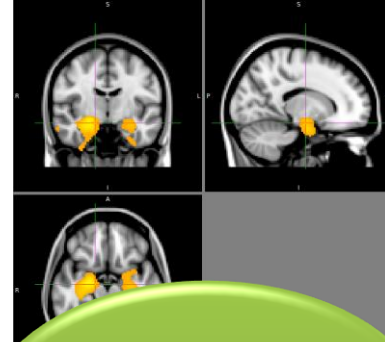
We propose to apply a **reduction factor** on the size of the hyperboxes



OASIS
subjects



VBM
Analysis



SPM



Compute
Feature
Vectors



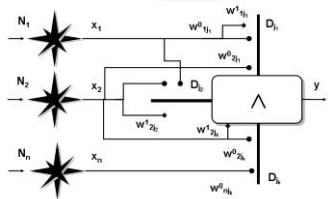
Mean and
standard
Deviation



DC



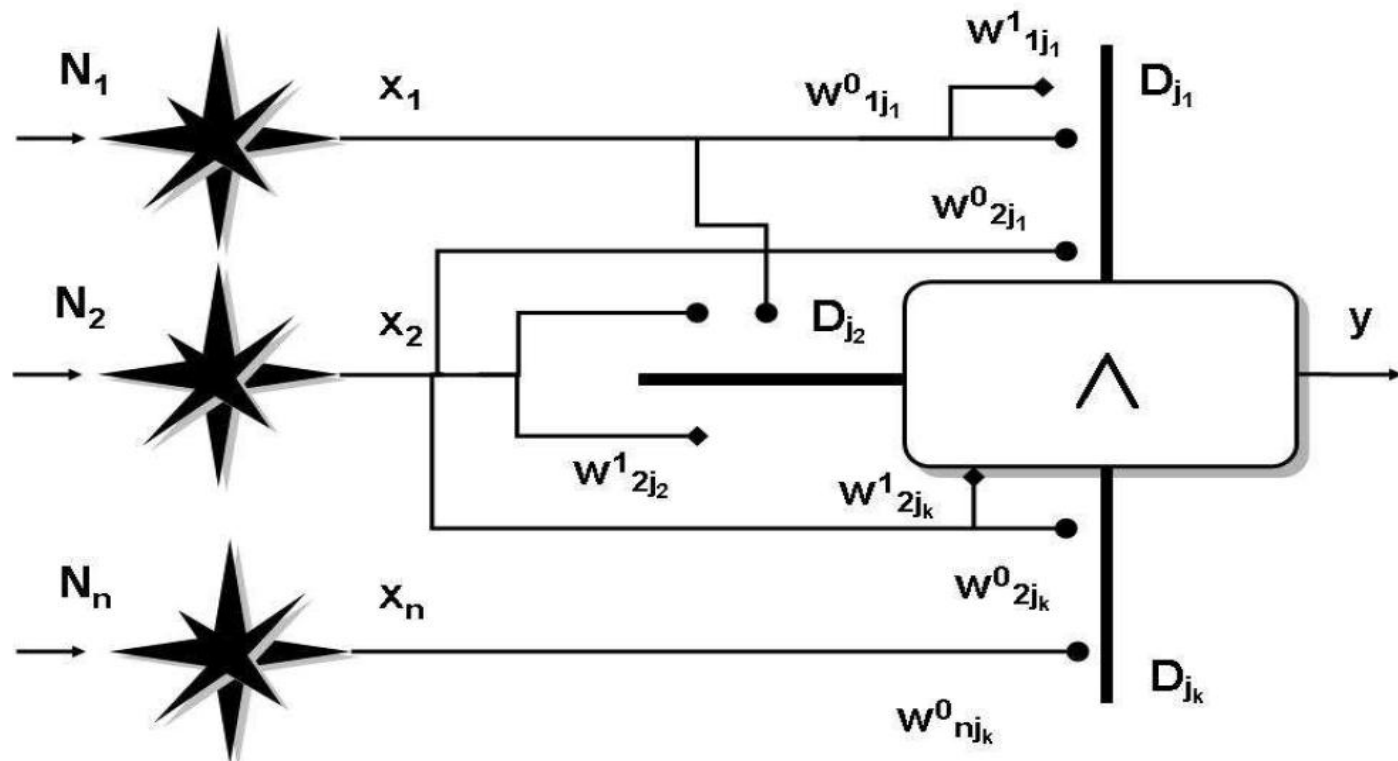
Result



| α | Accuracy | Sensitivity | Specificity |
|-------------|-----------|-------------|-------------|
| 0 | 58 | 94 | 23 |
| 0.5 | 60 | 81 | 40 |
| 0.53 | 59 | 77 | 42 |
| 0.55 | 64 | 85 | 44 |
| 0.57 | 63 | 83 | 43 |
| 0.6 | 62 | 81 | 44 |
| 0.63 | 64 | 83 | 45 |
| 0.65 | 69 | 83 | 54 |
| 0.67 | 64 | 78 | 49 |
| 0.7 | 64 | 79 | 49 |
| 0.73 | 65 | 79 | 52 |
| 0.75 | 65 | 78 | 51 |
| 0.77 | 67 | 78 | 56 |
| 0.8 | 69 | 81 | 56 |
| 0.83 | 66 | 76 | 55 |
| 0.85 | 62 | 73 | 51 |
| 0.87 | 63 | 74 | 52 |
| 0.9 | 63 | 74 | 51 |
| 0.93 | 66 | 74 | 57 |
| 0.95 | 65 | 73 | 57 |
| 0.97 | 61 | 69 | 53 |

Dendritic Computing

Structure of a single output class single layer Dendritic Computing system



Algorithm 1 Dendritic Computing learning based on elimination

Training set $T = \left\{ \left(\mathbf{x}^\xi, c_\xi \right) \mid \mathbf{x}^\xi \in \mathbb{R}^n, c_\xi \in \{0, 1\}; \xi = 1, \dots, m \right\}$, $C_1 = \{ \xi : c_\xi = 1 \}$, $C_0 = \{ \xi : c_\xi = 0 \}$

1. Initialize $j = 1$, $I_j = \{1, \dots, n\}$, $P_j = \{1, \dots, m\}$, $L_{ij} = \{0, 1\}$,

$$w_{ij}^1 = - \bigwedge_{c_\xi=1} x_i^\xi; w_{ij}^0 = - \bigvee_{c_\xi=1} x_i^\xi, \forall i \in I$$

Construct a basic hyperbox

2. Compute response of the current dendrite D_j , with $p_j = (-1)^{\text{sgn}(j-1)}$:

$$\tau_j(\mathbf{x}^\xi) = p_j \bigwedge_{i \in I_j} \bigwedge_{l \in L_{ij}} (-1)^{1-l} (x_i^\xi + w_{ij}^l), \forall \xi \in P_j.$$

3. Compute the total response of the neuron:

$$\tau(\mathbf{x}^\xi) = \bigwedge_{k=1}^j \tau_k(\mathbf{x}^\xi); \xi = 1, \dots, m.$$

4. If $\forall \xi \left(f(\tau(\mathbf{x}^\xi)) = c_\xi \right)$ the algorithm stops here with perfect classification of the training set.

5. Create a new dendrite $j = j + 1$, $I_j = I' = X = E = H = \emptyset$, $D = C_1$

6. Select \mathbf{x}^γ such that $c_\gamma = 0$ and $f(\tau(\mathbf{x}^\gamma)) = 1$.

7. $\mu = \bigwedge_{\xi \neq \gamma} \left\{ \bigvee_{i=1}^n |x_i^\gamma - x_i^\xi| : \xi \in D \right\}$.

8. $I' = \left\{ i : |x_i^\gamma - x_i^\xi| = \mu, \xi \in D \right\}$; $X = \left\{ (i, x_i^\xi) : |x_i^\gamma - x_i^\xi| = \mu, \xi \in D \right\}$.

9. $\forall (i, x_i^\xi) \in X$

a. if $x_i^\gamma > x_i^\xi$ then $w_{ij}^1 = -(x_i^\xi + \alpha \cdot \mu)$, $E_{ij} = \{1\}$

b. if $x_i^\gamma < x_i^\xi$ then $w_{ij}^0 = -(x_i^\xi - \alpha \cdot \mu)$, $H_{ij} = \{0\}$

10. $I_j = I_j \cup I'$; $L_{ij} = E_{ij} \cup H_{ij}$

11. $D' = \left\{ \xi \in D : \forall i \in I_j, -w_{ij}^1 < x_i^\xi < -w_{ij}^0 \right\}$. If $D' = \emptyset$ then goto step 2, else $D = D'$ goto step 7.

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Hard-limiter function

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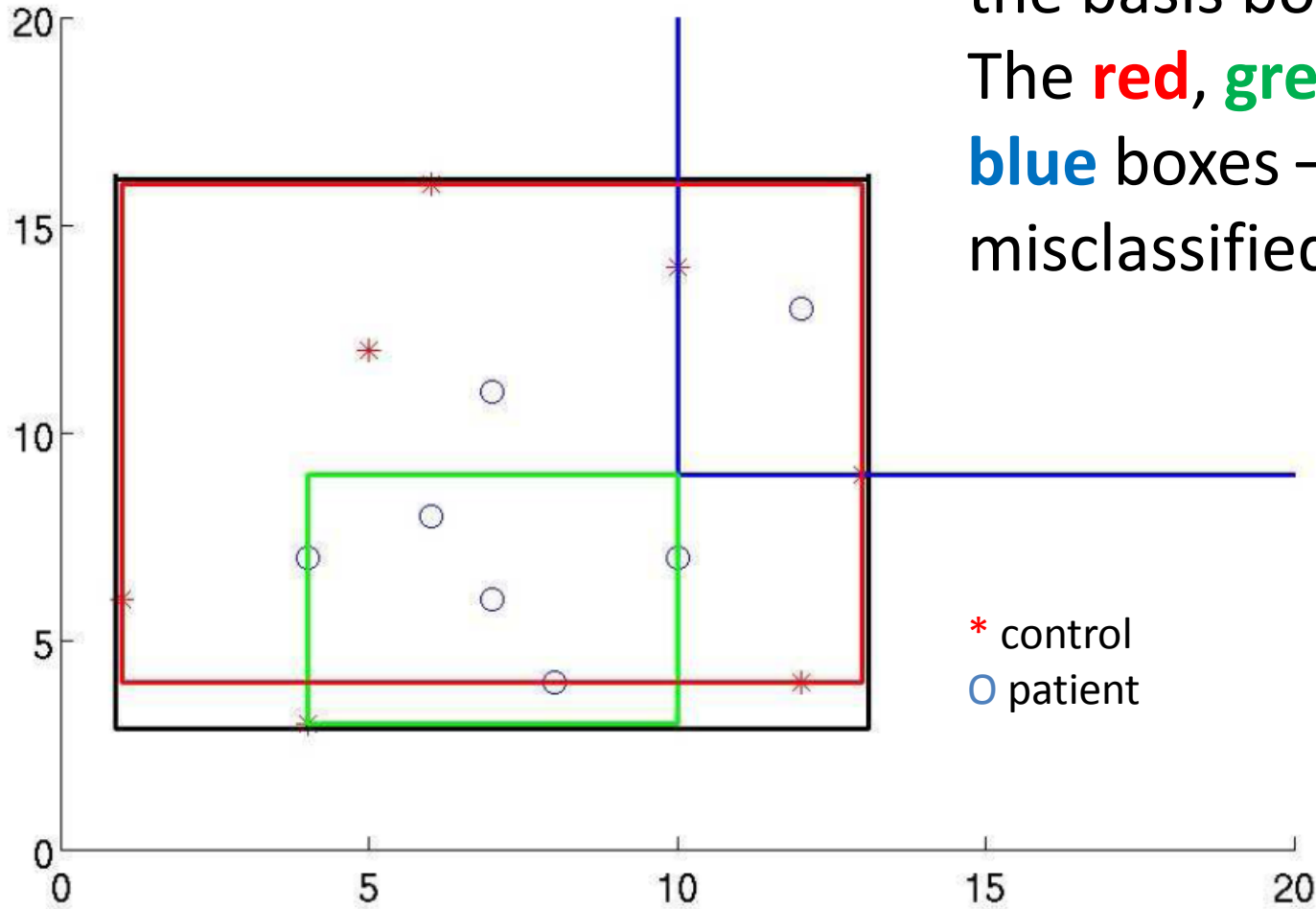
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Adding of dendrites to remove misclassified patterns of class 0 that fall inside this hyperbox

Dendritic Computing

Synthetic 2D dataset.



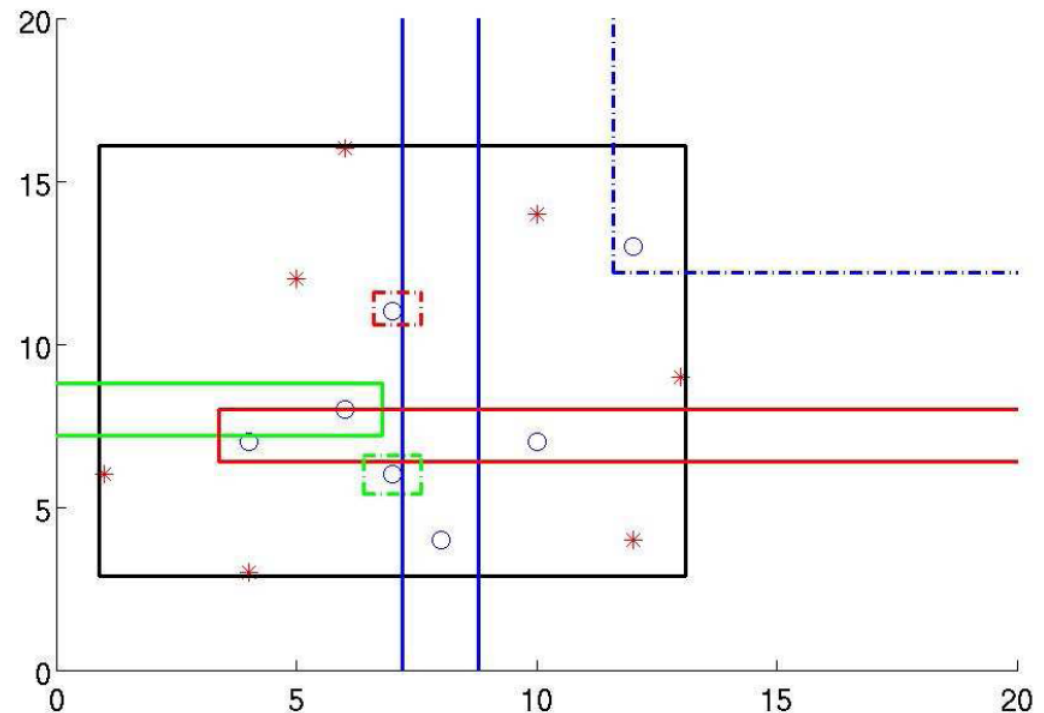
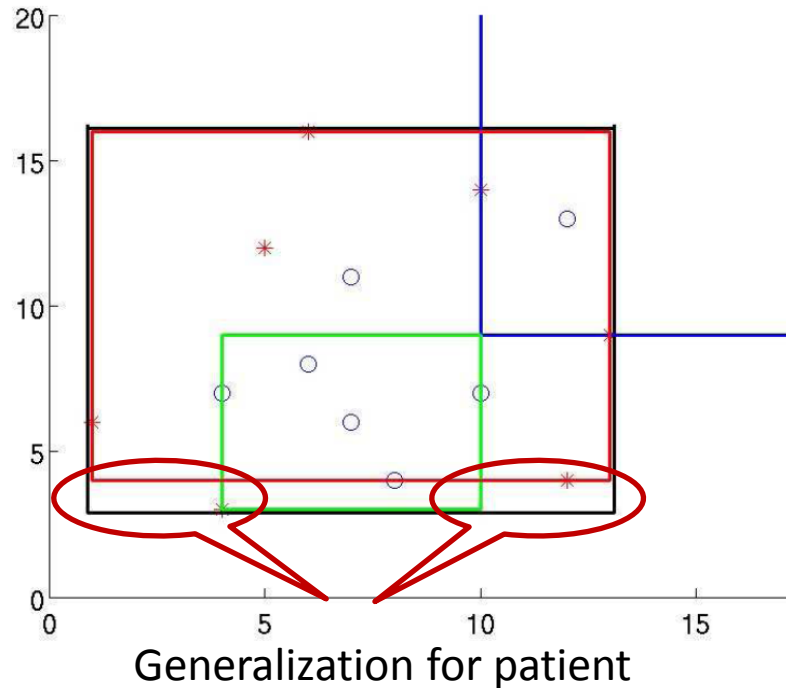
The **black** hyperbox is the basis box.

The **red**, **green** and **blue** boxes – to remove misclassified controls

* control
○ patient

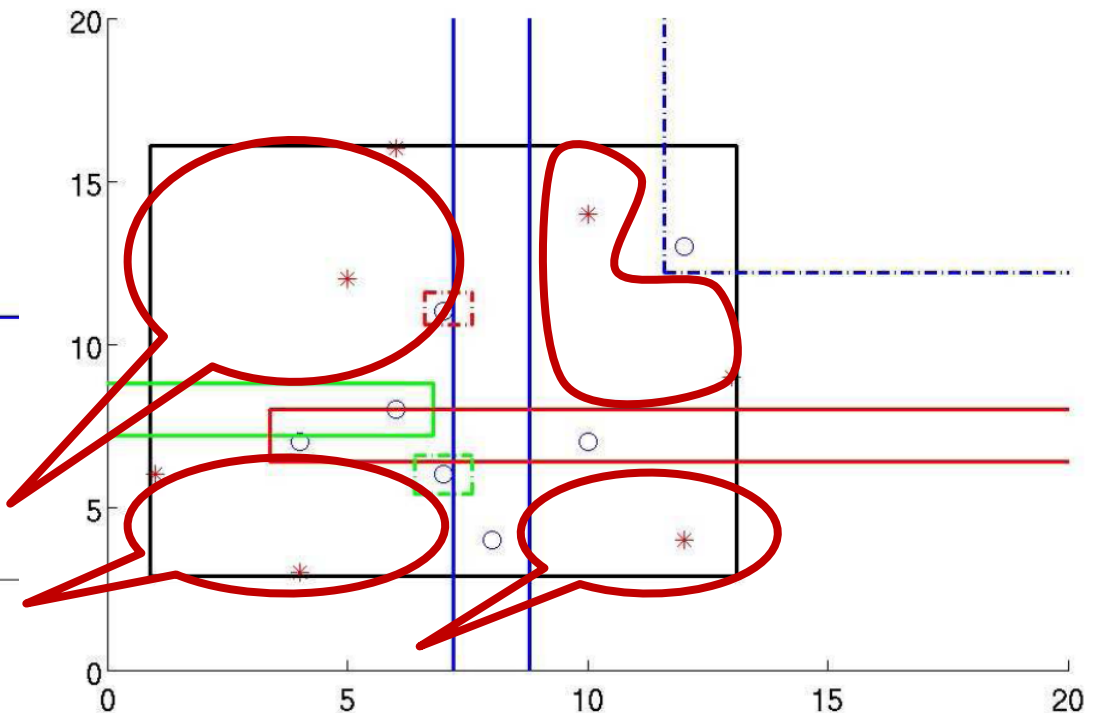
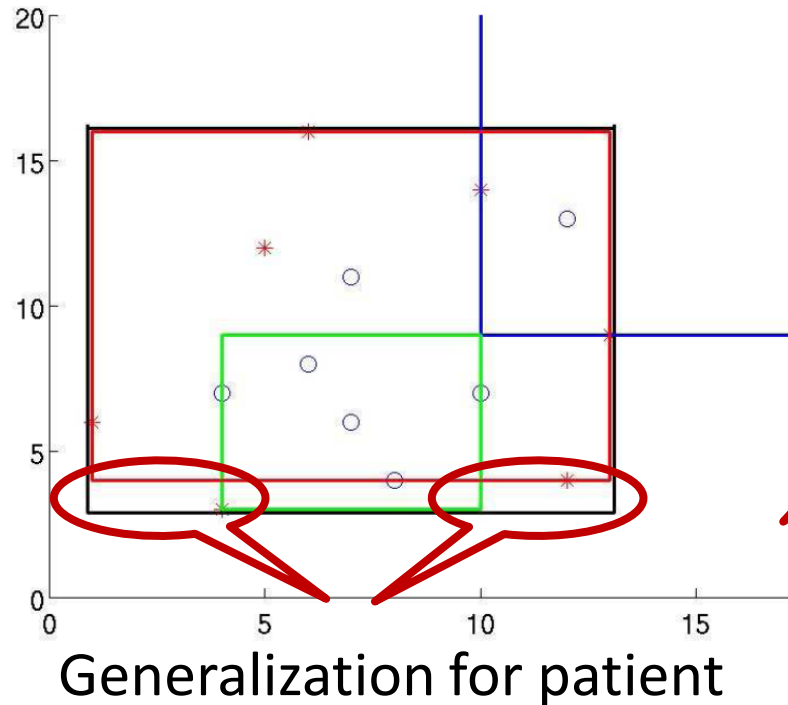
Dendritic Computing

- To establish specificity and sensitivity we propose shrinking the boundaries of the hyperbox.
- Exclude the region occupied by a misclassified item of control



Dendritic Computing

- To balance specificity and sensitivity we propose shrinking the boundaries of the hyperbox corresponding to each dendrite
- Exclude the region occupied by a misclassified item of control



Results on Alzheimer Detection

| α | Accuracy | Sensitivity | Specificity |
|-------------|-----------|-------------|-------------|
| 0 | 58 | 94 | 23 |
| 0.5 | 60 | 81 | 40 |
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First row – **baseline DC**

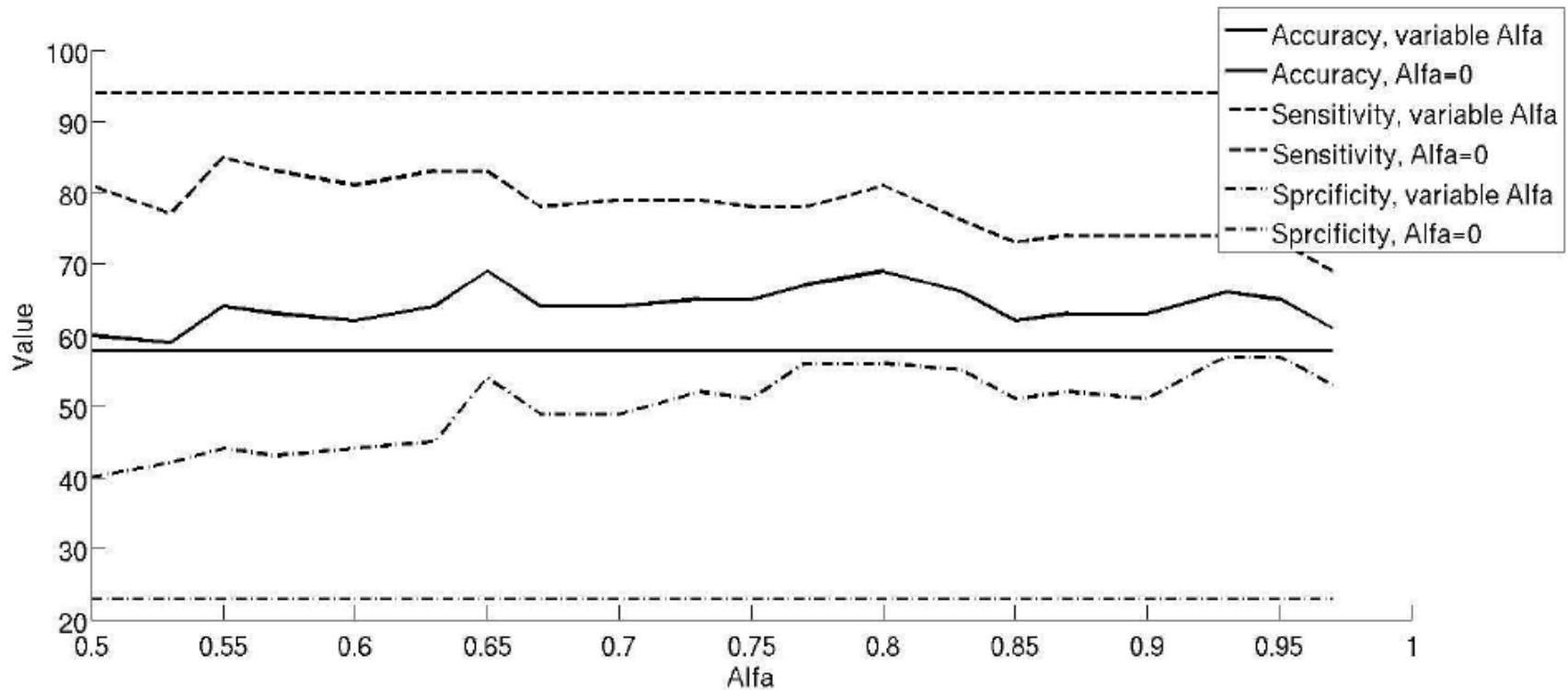
We find Shrinkage parameter of the box $\alpha \in [0, 1)$.

For each shrinkage parameter we have performed 10 –fold cross validation



The best result:
Sensitivity worse
Specificity **increase** – the best balance which gives the best Accuracy

Results on Alzheimer Detection



- **Specificity** (at the bottom) – control classification:
For 0.5 increase very fast and still increase
- **Sensitivity** (at the top) – patient classification:
Shows the better results for baseline DC
- But the **Accuracy** (in the middle)
Shows the best result for $\alpha=0.8$

Conclusions

We found empirically, performing **cross-validation** on an **Alzheimer's Disease database** of features computed from **MRI** scans, that a single layer neuron model endowed with **Dendritic Computing** has poor generalization capabilities.

The model shows **high sensitivity** but **poor specificity**. In this paper we have proposed a **simple change** in the learning algorithm

- that produces a significant **increase** in performance in terms of accuracy,
- obtaining a better trade-off between sensitivity and specificity.

Thank you for your attention!

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