

## SUMMARY - ALGORITHM

### Algorithm

#### Step 1: new input

receive  $\underline{u} \in \{0,1\}^N$

build:  $\underline{U} = \underline{1}^m \cdot \underline{u}^T$

binary input vector

binary input matrix  $\begin{bmatrix} 1 & 0 & 0 & 1 & 1 \\ 1 & 0 & 0 & 1 & 1 \\ \vdots & \vdots & \vdots & \vdots & \vdots \end{bmatrix}$

#### Step 2: update output matrix

$$\underline{P} = \underline{X}^1$$

$$\underline{B} = \underline{1} - \underline{1}^m \max_{\text{col}}(\underline{P})$$

$$\underline{C} = \underline{P} + \underline{B}$$

$$\underline{Y} = \underline{C} \cdot \underline{U}$$

represents previous state matrix

burst matrix (with bursting cols)

candidates for active outputs

active outputs

$$\text{or: } \underline{Y} = (\underline{X}^1 + (1 - \max_{\text{col}} \underline{X}^1)) \circ (\underline{1}^m \cdot \underline{U})$$

#### Step 3: state transition = calculate for all $(i,j)$

$$\underline{S}_{ij} = \underline{B}(D_{ij} - \Theta) \quad \text{binary synaptic matrix of cell } ij$$

$$\underline{Z}_{ij} = [\underline{y}(k_{ij1}), \underline{y}(k_{ij2}), \dots, \underline{y}(k_{ijd})] \quad \text{projected output to synaptic connections}$$

$$\underline{Q}_{ij} = \underline{S}_{ij} \circ \underline{Z}_{ij} \quad \text{distal activation matrix}$$

$$\underline{X} = [x_{ij}] = [\|\underline{1}^{\text{DT}} \underline{Q}_{ij}\|_\infty \geq \Theta] \quad \text{new state after transition}$$

#### Step 4: learning

$$\Delta D_{ij} = (S^+ Q_{ij} - S^- Q_{ij}) y_{ij} \quad (S^+ > S^-)$$

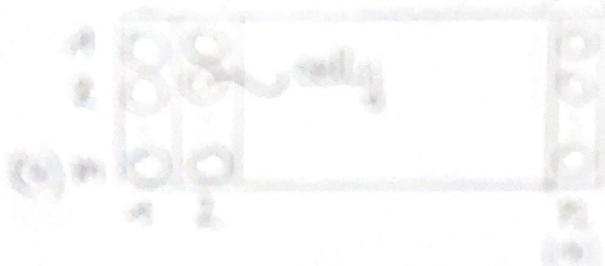
$$D_{ij} = D_{ij} + \Delta D_{ij} \quad \text{update distal weights}$$

\*)  $\circ$  means element wise multiplication

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## HTM SEQUENCE MEMORY

### HTM-Layer



bottom

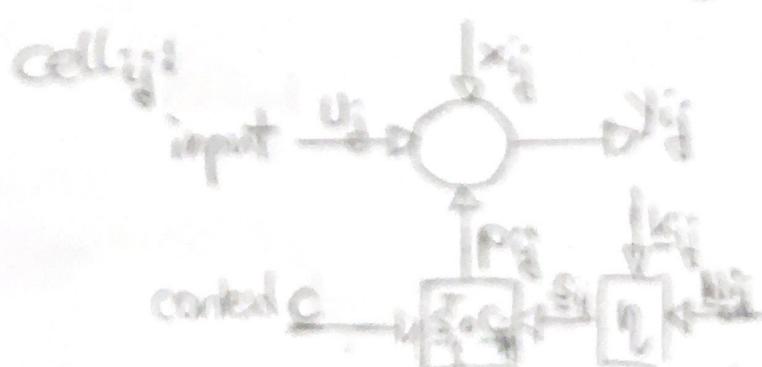
An HTM-layer provides  
a hierarchical structure  
encouraging the cells to

### Definition 3

We use the notation  $B = \{b_i\}$  for the set of memory values, and  $\{b_i\} \times \{x \in \mathbb{R}^n | 0 < x < 1\}$  for the set of inputs. We also write  $b_i \in B$  and  $x \in \mathbb{R}^n$ .

### Cells

Cells (also called HTM-neurons) are the elementary processing units with the following structure:



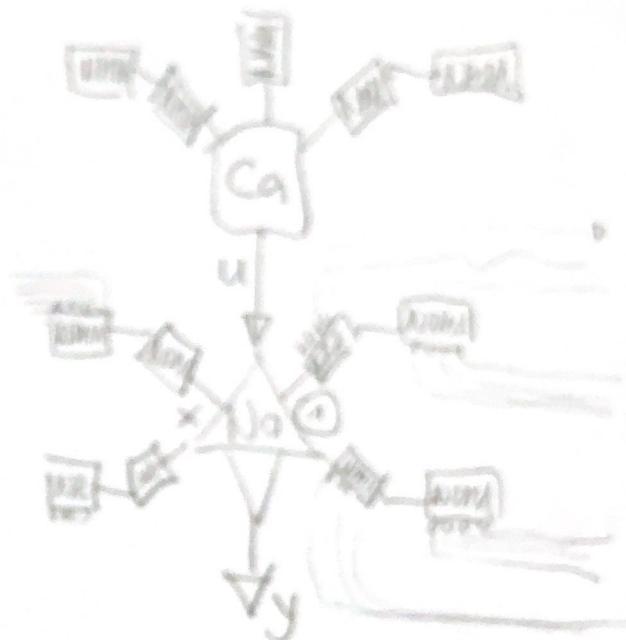
$u_i \in \mathbb{R}^n$   
 $x_i \in \mathbb{R}^n$   
 $y_i \in \mathbb{R}^n$   
 $c \in \mathbb{R}^n$   
 $p_i \in \mathbb{R}^n$   
 $s_i \in \mathbb{R}^n$   
 $k_i \in \mathbb{R}^n$   
 $l_i \in \mathbb{R}^n$   
 $m_i \in \mathbb{R}^n$

There are  $N$  such cells in an HTM-layer.

- $s_i$  is the number of synapses per cell signal and  $d_i$  is the number of total signals
- a cell kernel is frequently called **neocolumn**.
- a cell is also called **HTM-neuron**

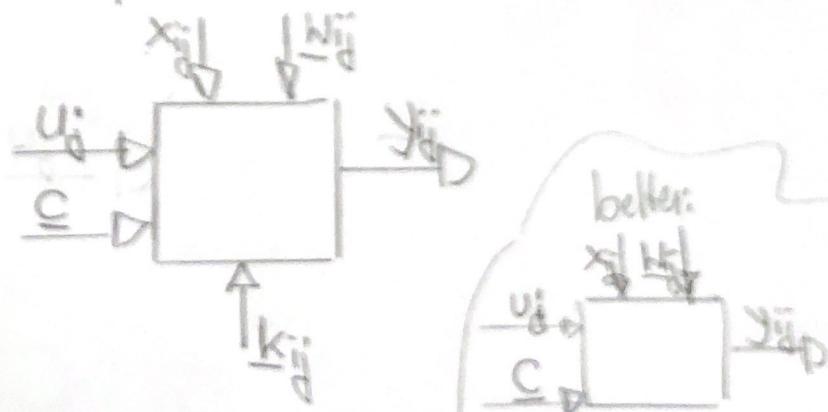
## NEURAL MODEL

Our computational model refers to the following neural model



Larkum et.al. (2009)

- From this neural model our HTM neuron is inspired as a computational block:



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## Units (minicolumns)

The  $m \times n$  cells of an HTM layer are organized as a set of  $n$  units (or minicolumns). Each cell comprises  $m$  cells, indexed by  $(i, j)$ . It is important to realize that all cells of a unit  $j$  have the same common input  $u_j \in [0, 1]$ , and the same context  $c_j$ .

## Vector-Signals/States

For a compact formalism it makes sense to combine scalar signal values to vectors. In this sense we define

$$x_j := [x_{1j}, \dots, x_{mj}]^T \in [0, 1]^m \quad (\text{predictive}) \text{unit state}$$

$$y_j := [y_{1j}, \dots, y_{mj}]^T \in [0, 1]^m \quad \text{unit output}$$

$$p_j := [p_{1j}, \dots, p_{mj}]^T \in \mathbb{N}_0^m \quad \text{prediction vector}$$

$$S_j := [S_{1j}, \dots, S_{mj}] \in [0, 1]^{N \times m} \quad \text{synaptic matrix}$$

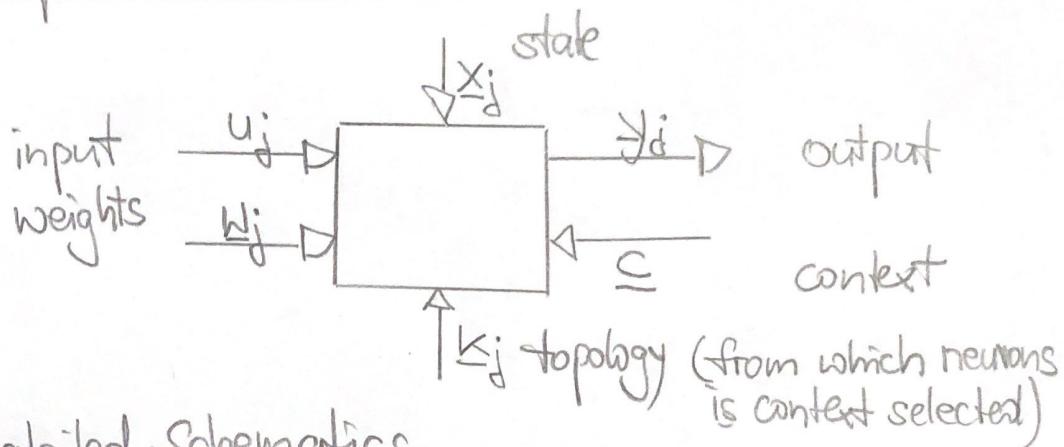
$$k_j := [k_{1j}, \dots, k_{mj}] \in [1..N]^m \quad \text{synaptic index matrix}$$

$$W_j := [W_{1j}, \dots, W_{mj}] \in [0, 1]^{M \times m} \quad \text{synaptic weight matrix}$$

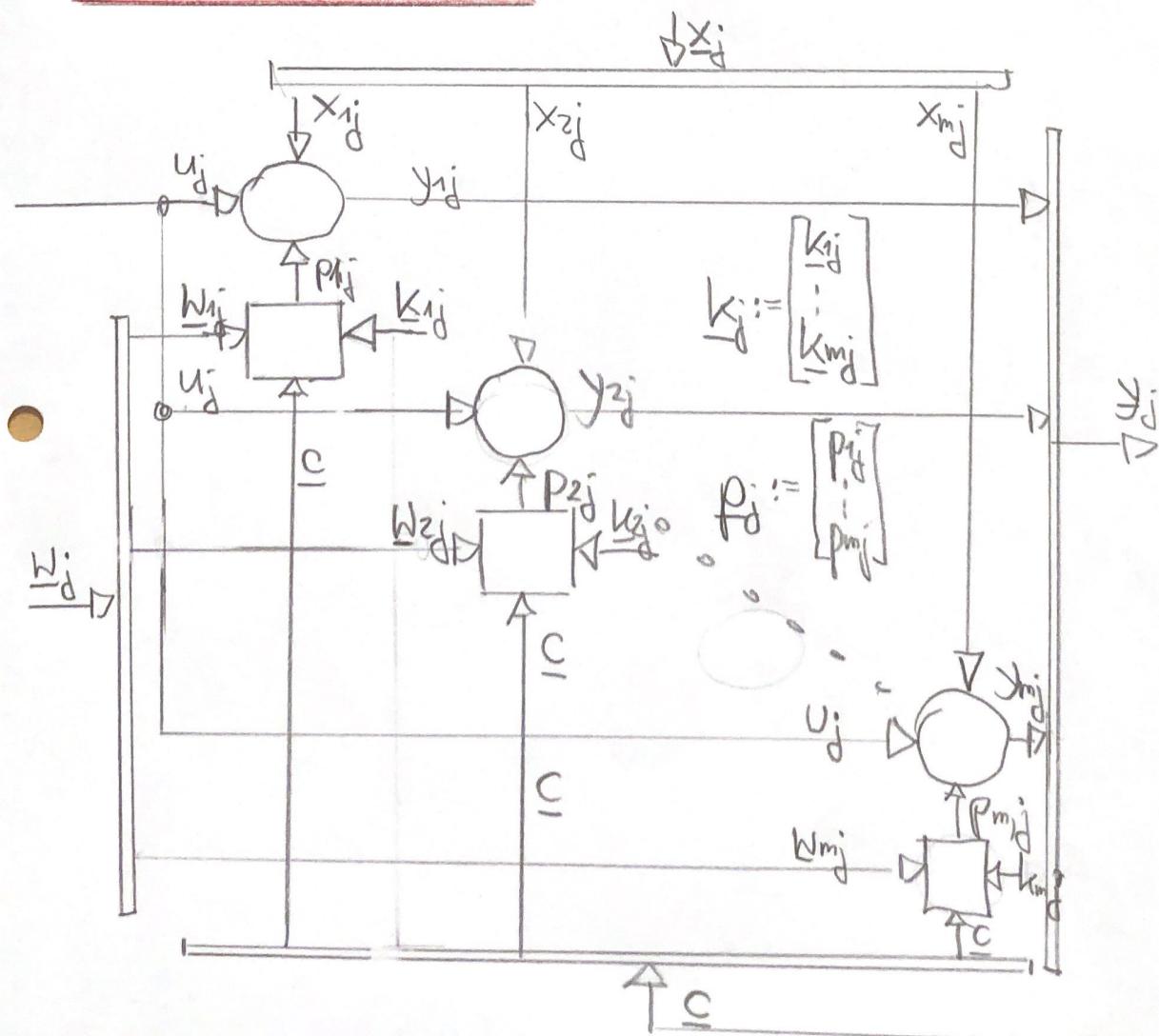
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## SCHEMATICS OF A (CELL) UNIT

### Simplified Schematics



### Detailed Schematics



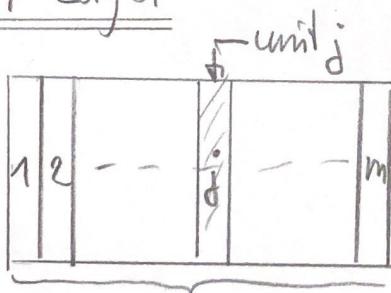
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## HTM-Layer / Recap

Since for (cell) unit  $j$  we have the following scalar/  
vector quantities

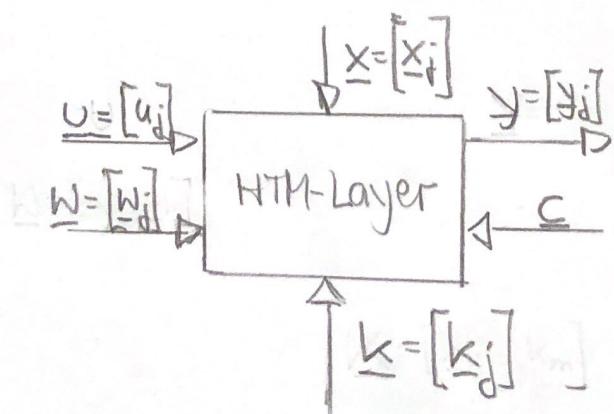
- 1       $u_j \in [0,1]$       input (common for all cells per unit)
- 4       $x_j, y_j \in [0,1]^m$       state and output
- 4       $p_j \in [0:N]^m$       prediction strength
- 10       $c \in [0,1]^{N^*}$       context ( $N^*$  might be greater than  $N$ )
- 40       $S_j \in [0,1]^{m \times N^*}$       synaptic connection matrix
- 12       $k_j \in [0,1]^M$       topology indices
- 48       $w_j \in [0,1]^{Mm}$

### HTM-Layer



$$x := [x_i] = \begin{bmatrix} x_1 \\ \vdots \\ x_n \end{bmatrix}$$

$$y := [y_i] = \begin{bmatrix} y_1 \\ \vdots \\ y_n \end{bmatrix}$$

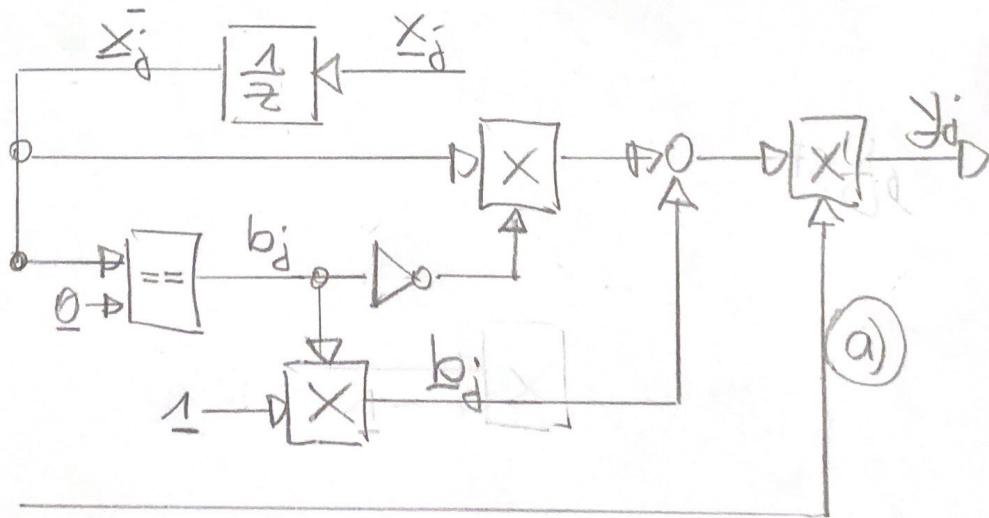


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## Output Activation

- a) Outputs are only active for active input signals
- b) if a cell was predictive in the previous step, then the output is active (given  $u_j = 1$ )
- c) if no cell in the unit was predictive in the previous step, then the output gets active (given  $u_j = 1$ )

## Signal Processing Chart

 $u_j$ 

## Remarks

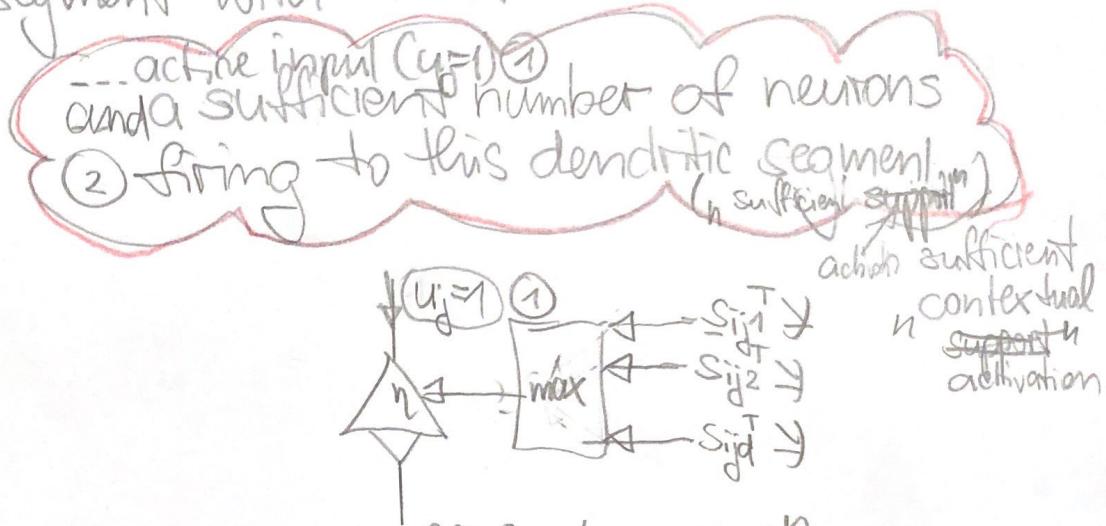
(i) given  $x_j^-$  (previous predictive state) equals  $0$ ,  
 we have  $b_j = 1$ , and  $b_j^- = [1, 1, \dots, 1]$  (burst).

Then (given  $u_j = 1$ ) the output becomes  $y_j = [1, 1, 1, 1]$

(ii) given  $x_j^- \neq 0$  we have  $b_j = 0$  and  $y_j$  gets  $x_j^-$  (given  $u_j = 1$ )

## STATE TRANSITION

In every cell (HTM neuron) the updated state is 1, iff the cell has at least one distal segment with

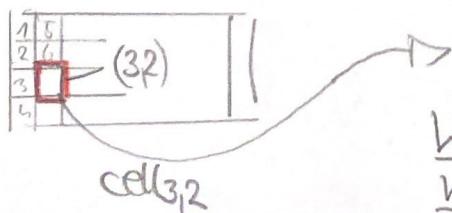


What means „sufficient support“

=> it means that the total sum of synaptic connections to activated cells exceed threshold  $n$

### Example

Layer:  $4 \times 10$  ( $N=40$  cells) with 2 distal segments a 6 potential connections;



$$\begin{matrix} & \begin{matrix} 2 & 7 & 18 & 32 & 35 & 39 \end{matrix}^T \\ \begin{matrix} 2 \\ 7 \end{matrix} & \left[ \begin{matrix} W_{321} \\ W_{322} \end{matrix} \right] \\ & \begin{matrix} 18 & 16 & 22 & 33 & 35 \end{matrix}^T \end{matrix} \Rightarrow W_{32} = \begin{bmatrix} 0.7 \\ 0.8 \\ 0.25 \\ 0.31 \\ 0.12 \\ 0.2 \end{bmatrix}$$

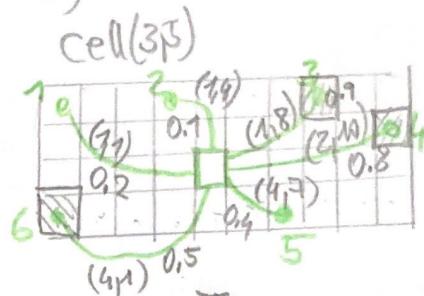
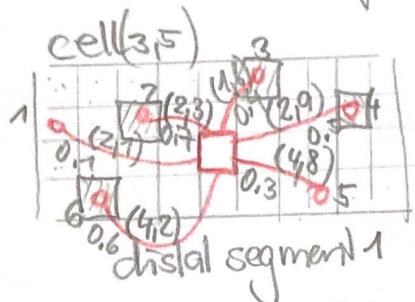
$$\begin{aligned} K_{321} &= [2, 7, 18, 32, 35, 39] \\ W_{321} &= [0.7, 0.8, 0.25, 0.31, 0.12, 0.2] \\ K_{322} &= [18, 16, 22, 33, 35] \\ W_{322} &= [ ] \end{aligned}$$

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## DISTAL ACTIVATION

The distal segments are characterized by a weight matrix  $\underline{W}_{ij}$  and a connection index matrix  $K_{ij}$ .

Let  $\sigma_e(i,j)$  be the index mapping function from  $(i,j) \rightarrow k$ , given by  $\sigma_e(i,j) = (m-1)j + i$ . We could have for example ( $s=6, d=2$ )



$$K_{351} = [\sigma_e(21), \sigma_e(23), \sigma_e(16), \sigma_e(29), \sigma_e(48), \sigma_e(42)]^T = \\ [2, 10, 21, 34, 32, 8]^T$$

$$K_{352} = [\sigma_e(11), \sigma_e(14), \sigma_e(18), \sigma_e(210), \sigma_e(47), \sigma_e(41)]^T = \\ [1, 13, 25, 38, 28, 4]^T$$

$$\underline{W}_{351} = [0.1, 0.7, 0.9, 0.5, 0.3, 0.6]^T$$

$$\underline{W}_{352} = [0.2, 0.1, 0.9, 0.8, 0.4, 0.5]^T$$

This leads us to the synaptic connection matrices

$$S_{351} = \begin{bmatrix} 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \end{bmatrix} = 8(K_{351} \otimes \underline{W}_{351} - \eta) \quad \left\{ \begin{array}{l} \text{with} \\ \eta = 0.5 \\ (\text{in our example}) \end{array} \right.$$

$$S_{352} = \begin{bmatrix} 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \end{bmatrix} = 5(K_{352} \otimes \underline{W}_{352} - \eta)$$

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## DISTAL ACTIVATION VECTOR

Given a cell  $(ij)$  with  $d$  distal segments given by  $\underline{K}_{ij} = [k_{ij1}, \dots, k_{ijd}]$  and  $\underline{W}_{ij} = [w_{ij1}, \dots, w_{ijd}]$  the distal activation is defined as

$$\underline{d}_{ij} = (\underline{W}_{ij} \underline{S}_{ij}^T \circ \underline{Y})$$

$$\underline{S}_{ij} := [S_{ij1}, \dots, S_{ijd}]$$

$$\underline{S}_{ijk} := S_{ijk}(:, i)$$

### Example

Given  $\underline{Y} = Y(m \times n) = \begin{bmatrix} 1 & 0 & 0 & 1 & 0 & 1 & 0 & 0 & 0 \\ 1 & 0 & 0 & 1 & 0 & 1 & 0 & 0 & 1 \\ 1 & 0 & 0 & 1 & 0 & 1 & 0 & 0 & 1 \\ 1 & 0 & 0 & 1 & 0 & 1 & 0 & 0 & 0 \\ 1 & 0 & 0 & 1 & 0 & 1 & 0 & 0 & 0 \end{bmatrix}$

We get:

$$\underline{d}_{351} = \underline{S}_{ij1}^T \underline{Y} = 3, \quad \underline{d}_{352} = \underline{S}_{ij2}^T \underline{Y} = 1$$

$$\Rightarrow \underline{d}_{35} = \begin{bmatrix} 3 \\ 1 \end{bmatrix}$$

which means

For distal segment 1 there are 3 feeding cells to synaptic connections, and for distal segment 2 there is only 1 feeding cell, given the activated output vector  $\underline{Y}$ .

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## DISTAL ACTIVATION MATRIX AND STATE TRANSITION

### Distal Activation Matrix

$$D := [d_{ij} \ d_{ij-1} \ d_{ij}] \quad \text{with } d_{ij} = d_{ij} \text{ given by eqn 1}$$

The distal activation  $p \in [0, 1]^N$  is defined as

$$P = \max_{1 \leq i \leq N} D^i$$

$\max(\cdot)$  means  
Find the maximum  
element of each row  
has to be taken

### Example

$$D = \begin{bmatrix} \dots & 3 & \dots \\ \dots & 1 & \dots \\ \vdots & & \vdots \\ \ddots & & \ddots \end{bmatrix} \Rightarrow \max_{1 \leq i \leq N} D^i = \begin{bmatrix} 3 \\ \vdots \\ 1 \end{bmatrix} = D (\max_{1 \leq i \leq N} D^i \geq 2) = \begin{bmatrix} 1 \\ 1 \\ 1 \end{bmatrix}$$

### State transition

The new state  $x$  is calculated

$$x = (p \geq \Theta)$$

precisely:  $x^+ = (p^t \geq \Theta)$

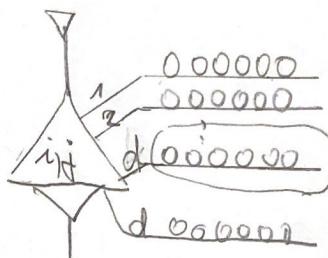
Example:  $\Theta = 2$

$$D = \begin{bmatrix} \dots & 3 & \dots \\ \dots & 1 & \dots \\ \vdots & & \vdots \\ \ddots & & \ddots \end{bmatrix}, p = \max_{1 \leq i \leq N} D^i = \begin{bmatrix} 3 \\ \vdots \\ 1 \end{bmatrix} = D \xrightarrow{\Theta=2} \begin{bmatrix} 1 \\ 1 \\ 1 \end{bmatrix}$$

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## PROJECTED OUTPUT VECTOR

Let us focus on cell  $(ij)$  with  $n_d$  distal segments indexed from  $1 \dots n_d$  each having  $s$  synapses



distal segment  $d \Rightarrow w_{ijd}, k_{ijd}$

Let  $\underline{k}_{ijd} = \underline{\gamma} = [\gamma_1 \dots \gamma_s]^T$   $\underline{\gamma}$ -index vector under focus

Then  $\underline{\gamma}$  sets a focus on the cell output vector  $\underline{y}$  given by the  $s$  indices  $\gamma_1 \dots \gamma_s$ . This is visualized below.

$$\underline{y}^T = [y_{\gamma_1} \ y_{\gamma_2} \ y_{\gamma_3} \ \dots \ y_{\gamma_s}]$$

$\gamma_1 \quad \gamma_2 \quad \gamma_3 \quad \dots \quad \gamma_s$

$\uparrow \quad \uparrow \quad \uparrow \quad \dots \quad \uparrow$

$\gamma_1 \quad \gamma_2 \quad \gamma_3 \quad \dots \quad \gamma_s$

The projected vector  $p_{\underline{\gamma}} = [y_{\gamma_1} \ y_{\gamma_2} \ y_{\gamma_3} \ \dots \ y_{\gamma_s}]^T$  is the <sup>concept</sup> summary of the focused elements of  $\underline{y}$  given by  $\underline{\gamma}$ , and can be calculated as a projection

$$p_{\underline{\gamma}} = P(\underline{\gamma}) \cdot \underline{y} \quad P \in \{0,1\}^{s \times N}$$

where

$$P(\underline{\gamma}) := \begin{bmatrix} \underline{e}_{\gamma_1}^T \\ \underline{e}_{\gamma_2}^T \\ \vdots \\ \underline{e}_{\gamma_s}^T \end{bmatrix}$$

and  $\underline{e}_i^T$  being the  $i$ -th row of the  $N \times N$  unit matrix  $I^{N \times N}$

## DISTAL ACTIVATION $Q_{ij}$

In each distal segment  $j$  of cell  $(i,j)$   
we have

$$q_{ijd} := \underbrace{\sigma(w_{ijd})}_{=S_{ijd}} \circ \underbrace{P(k_{ijd})}_{=Z_{ijd}} \gamma \quad \sigma(x) := \begin{cases} 1 & x \geq 0 \\ 0 & x < 0 \end{cases}$$

as the activation vector for the distal segment  
and

$$Q_{ij} := [q_{ij1} \dots q_{ijd}]$$

as the "distal activation matrix".

$Q_{ij}$  gives an overview  
over the activation  
potentials arriving  
at the synapses

### Example

consider:  $W_{ij} = \begin{bmatrix} 1 & 2 \\ 5 & 8 \\ 8 & 7 \\ 3 & 4 \\ 7 & 0 \\ 2 & 6 \end{bmatrix} \cdot \frac{1}{10}$ ,  $K_{ij} = \begin{bmatrix} 1 & 2 \\ 3 & 4 \\ 5 & 6 \\ 6 & 7 \\ 8 & 9 \end{bmatrix} = [k_{ij1} \ k_{ij2}]^T$

Let:  $y = [0 \ 1 \ 1 \ 0 \ 0 \ 1 \ 0 \ 1 \ 1 \ 0 \ 0 \ 1 \ 0 \dots]^T$

Then:

$$P_{ij1} = [e_1, e_3, e_5, e_6, e_8, e_9]^T y = [0 \ 1 \ 0 \ 1 \ 1 \ 1]^T$$

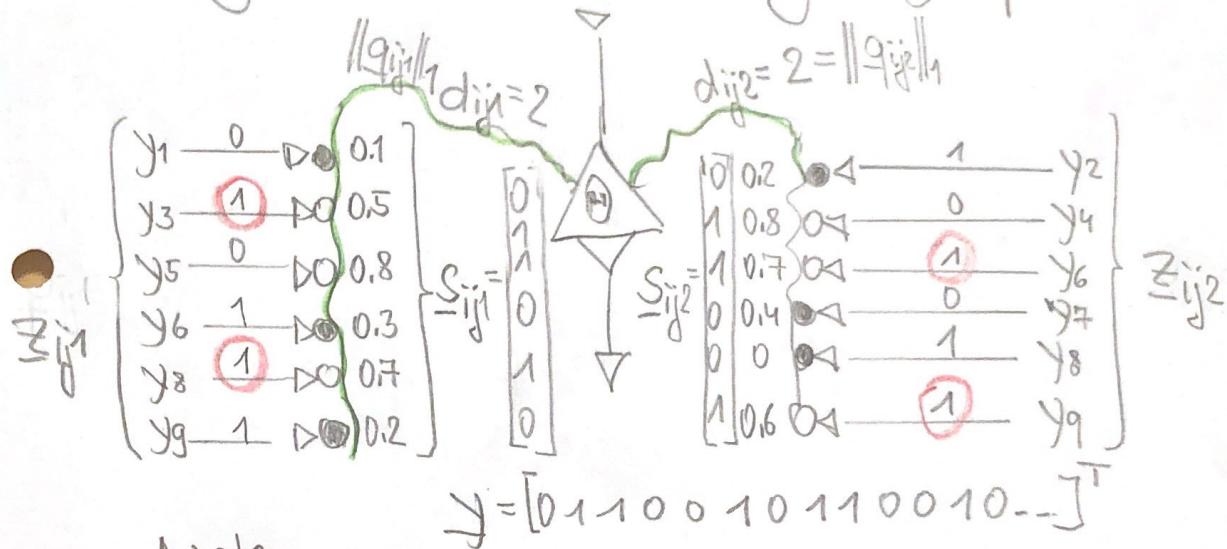
$$P_{ij2} = [e_2, e_4, e_7, e_9, e_8, e_9]^T y = [1 \ 0 \ 1 \ 0 \ 1 \ 1]^T$$

$$q_{ij1} = \sigma([158372]/10) \circ P_{ij1} = [0 \ 1 \ 0 \ 0 \ 1 \ 0]^T$$

$$q_{ij2} = \sigma([287406]/10) \circ P_{ij2} = [0 \ 0 \ 1 \ 0 \ 0 \ 1]^T \Rightarrow Q_{ij} = [0 \ 1 \ 0 \ 0 \ 1 \ 0]^T$$

## WHAT IS DISTAL ACTIVATION

Consider cell  $(i, j)$ , which has  $d$  distal segments, each containing  $s$  synapses



### Note

- Synapses with weight  $< \eta$  ( $\eta$  - synaptic threshold) are considered as not connected (black circles) and have as such no contribution

The contribution according to connected synapses is represented by the distal activation vectors

$$\text{distal segment 1: } \mathbf{q}_{ij1} = \sum_j s_{ij1} \mathbf{z}_{ij1} = \begin{bmatrix} 0 \\ 1 \\ 1 \\ 0 \\ 0 \\ 1 \\ 1 \\ 0 \\ 1 \end{bmatrix} \begin{bmatrix} 0 \\ 1 \\ 0 \\ 1 \\ 0 \\ 1 \\ 1 \\ 0 \\ 1 \end{bmatrix} = \begin{bmatrix} 0 \\ 1 \\ 0 \\ 0 \\ 0 \\ 1 \\ 1 \\ 0 \\ 1 \end{bmatrix}$$

$$\text{distal segment 2: } \mathbf{q}_{ij2} = \sum_j s_{ij2} \mathbf{z}_{ij2} = \begin{bmatrix} 0 \\ 1 \\ 1 \\ 0 \\ 0 \\ 1 \\ 0 \\ 1 \\ 1 \end{bmatrix} \begin{bmatrix} 1 \\ 0 \\ 0 \\ 1 \\ 0 \\ 1 \\ 1 \\ 0 \\ 1 \end{bmatrix} = \begin{bmatrix} 0 \\ 1 \\ 0 \\ 0 \\ 0 \\ 1 \\ 0 \\ 0 \\ 1 \end{bmatrix}$$

$$\text{thus } \mathbf{Q}_{ij} = [\mathbf{q}_{ij1} \ \mathbf{q}_{ij2}] = \begin{bmatrix} 0 & 0 \\ 1 & 0 \\ 0 & 1 \\ 0 & 0 \\ 1 & 0 \\ 0 & 0 \\ 1 & 1 \\ 0 & 1 \\ 1 & 0 \end{bmatrix}$$

distal activation  
 { in segment 1 ↑  
 { in segment 2 ↑

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## LEARNING

If a cell is correctly predicted, it means that it has at least one distal segment which has sufficient distal activation, i.e.

$$\|q_{ijd}\|_1 \geq \Theta$$

$$\|q_{ijd}\|_1 = S_{jd}^T Z_{jd} \geq \Theta$$

Thus we calculate

$$d_{ij} := \begin{bmatrix} \|q_{ij1}\|_1 \\ \|q_{ij2}\|_1 \\ \|q_{ijd}\|_1 \end{bmatrix} = Q_{ij}^T \underline{1}$$

Example

$$d_{ij} = \underbrace{\begin{bmatrix} 0 & 1 & 0 & 0 & 1 & 0 \\ 0 & 0 & 1 & 0 & 0 & 1 \end{bmatrix}}_{Q_{ij}^T} \begin{bmatrix} 1 \\ 1 \\ 1 \end{bmatrix} = \begin{bmatrix} 2 \\ 2 \end{bmatrix}$$

Learning Rule:

Any distal segment of an active cell ( $y_{ij} = 1$ ) is reinforced by  $\delta^+$

$$\Delta w_{ij} = \delta^+ Q_{ij}^T - \delta^- \underline{1}^{s \times d}, \quad \delta^+ > \delta^-$$

## Again: Output Calculation

Let  $\underline{P} = \underline{x} = [p_1, \dots, p_m] \in \{0,1\}^{m \times n}$

$\Rightarrow$  define a basis matrix  $\underline{B} = [b_1, \dots, b_m]$

with  $b_j = \begin{cases} 0 & \text{for } p_i + 0 \\ 1 & \text{for } p_i = 0 \end{cases}$

$$\underline{B} = 1 - \underline{I}_{\text{col}}(\underline{P})$$

"zeros" at those columns which are zero in  $\underline{P}$  (so it complements)

Then the output matrix  $\underline{Y}$  calculates:

$$\underline{Y} = (\underline{P} + \underline{B}) \cdot (\underline{1}^m \cdot \underline{u}^T)$$

### Example

$$\underline{P} = \begin{bmatrix} 0 & 0 & 1 & 0 & 0 & 1 & 0 & 1 & 1 \\ 1 & 0 & 1 & 1 & 0 & 0 & 1 & 1 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 \\ 0 & 1 & 0 & 1 & 0 & 1 & 0 & 1 & 0 \end{bmatrix}$$

$$\underline{u} = [0 \ 0 \ 0 \ 0 \ 0 \ 1 \ 1 \ 0 \ 0]^T$$

$$\underline{B} = \begin{bmatrix} 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 1 \\ 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 \end{bmatrix} = 1 - \underbrace{\begin{bmatrix} 1 \\ 1 \\ 1 \\ 1 \end{bmatrix} \begin{bmatrix} 1 & 1 & 1 & 1 & 0 & 1 & 1 & 1 \end{bmatrix}^T}_{\text{col}(\underline{P})}$$

$$\underline{P} + \underline{B} = \begin{bmatrix} 1 & 0 & 1 & 0 & 1 & 1 & 0 & 1 & 1 \\ 1 & 0 & 1 & 1 & 0 & 1 & 0 & 1 & 1 \\ 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 1 \\ 0 & 1 & 0 & 1 & 0 & 1 & 0 & 1 & 0 \end{bmatrix}$$

$$\underline{Y} = \begin{bmatrix} 1 & 0 & 1 & 0 & 1 & 1 & 1 & 0 & 0 \\ 0 & 0 & 1 & 0 & 0 & 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \end{bmatrix} = (\underline{P} + \underline{B}) \cdot (\underline{1}^m \cdot \underline{u}^T)$$

mask according  
to input  $\underline{u}$