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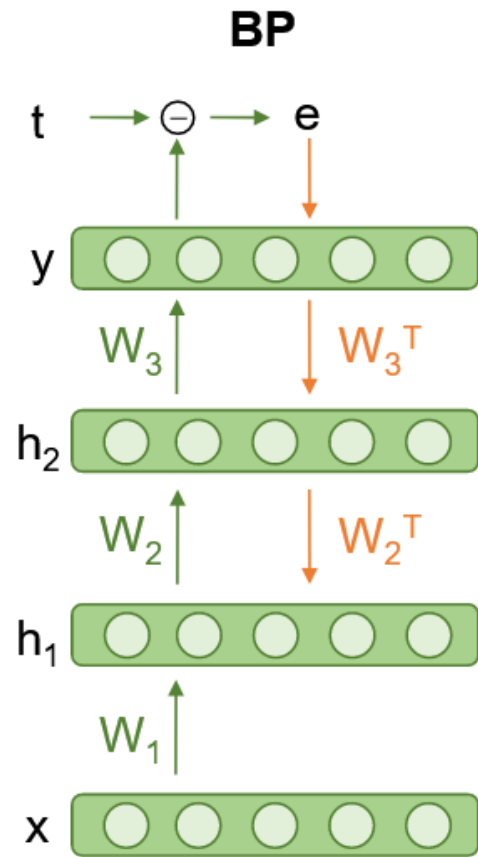
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Error-driven Input Modulation: Solving the Credit Assignment Problem without a Backward Pass

Giorgia Dellaferrera & Gabriel Kreiman

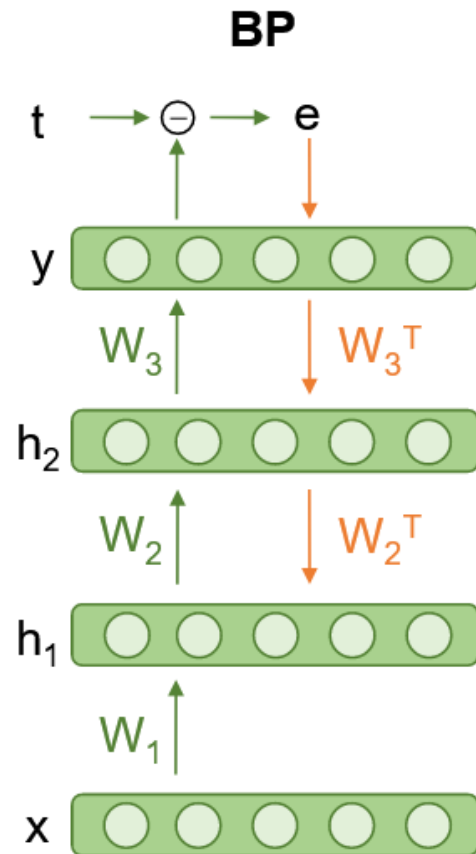
ICML 2022

Backpropagation of the error is not biologically plausible



Rumelhart et al., 1995

Backpropagation of the error is not biologically plausible



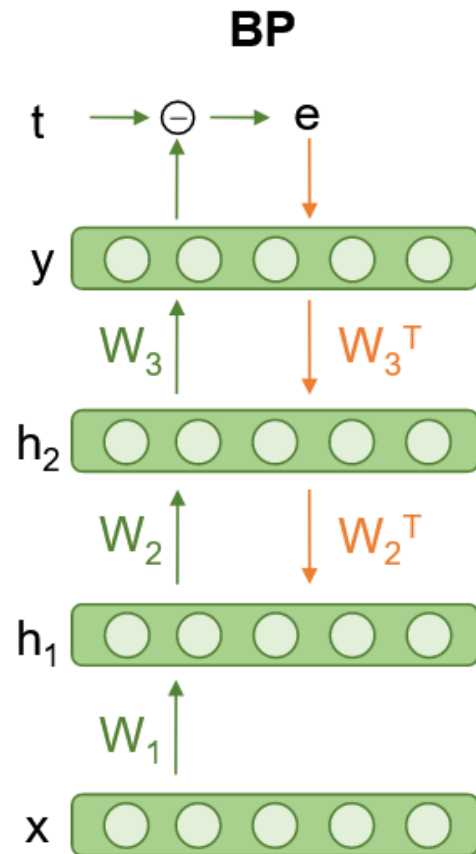
» Weight transport problem

- Symmetric weights for forward and backward computation

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Lillicrap et al., 2020

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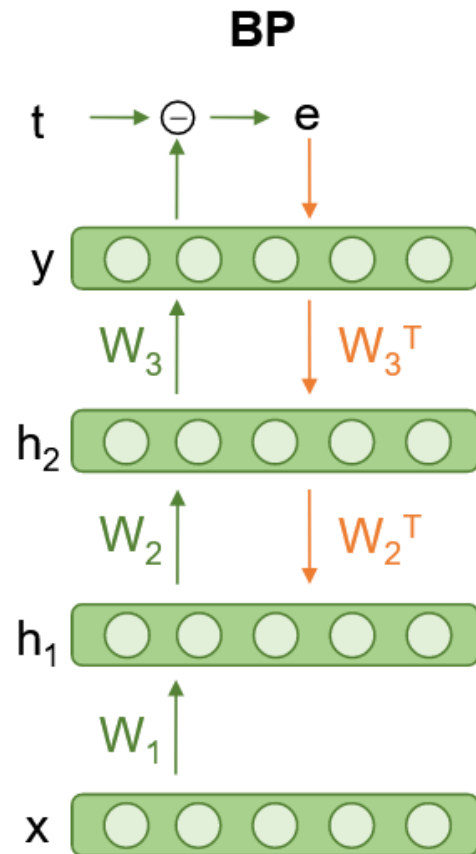
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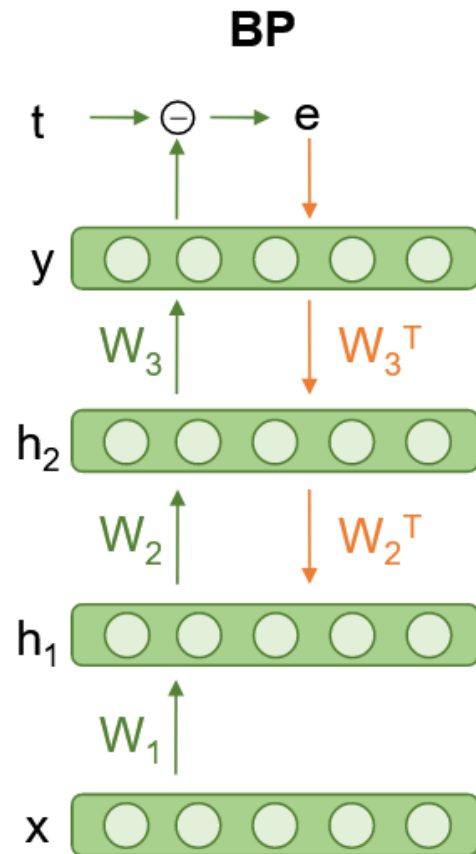
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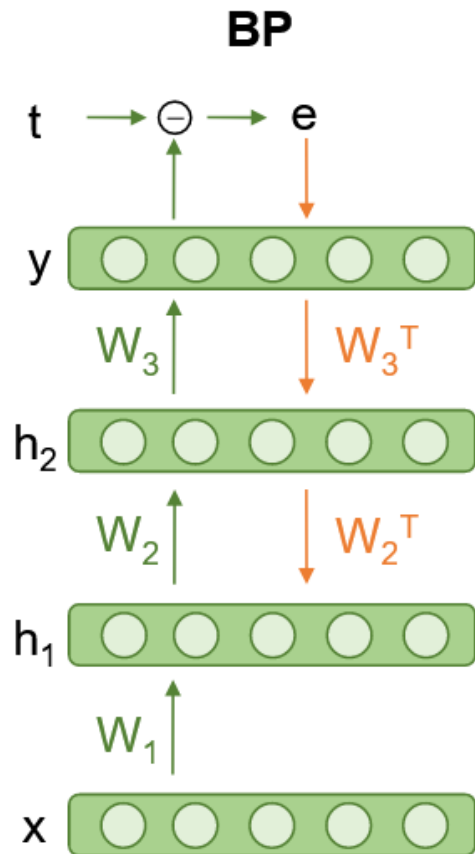
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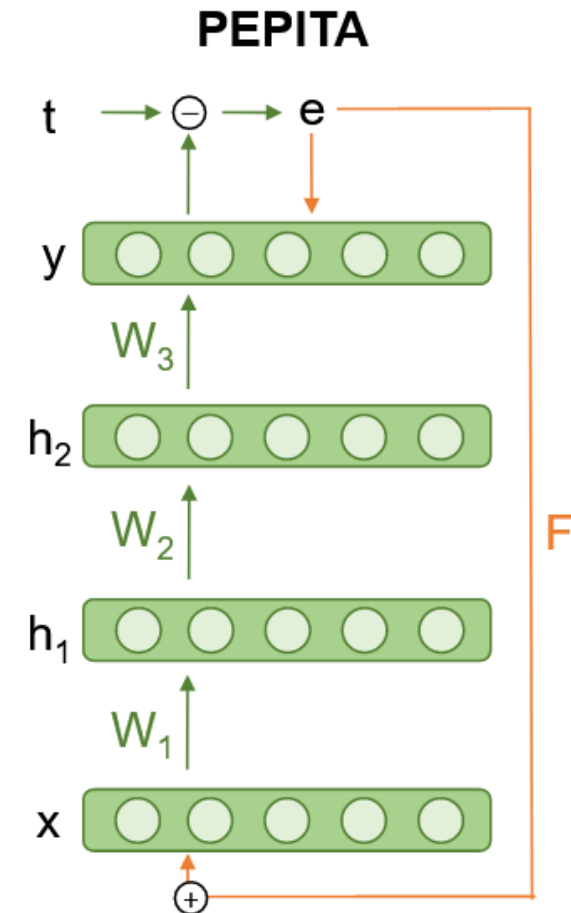


Alternative Training Schemes

Lillicrap et al., 2020

The PEPITA learning rule for Fully Connected Neural Networks

- » **PEPITA** = Present the Error to Perturb the Input To modulate Activity
 - Substitutes the standard Forward+Backward scheme with **two Forward Passes**



The PEPITA learning rule for Fully Connected Neural Networks

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 - Substitutes the standard Forward+Backward scheme with **two Forward Passes**

Algorithm 1 Implementation of PEPITA

Given: Input (x) and label ($target$)

#standard forward pass

$$h_0 = x$$

for $\ell = 1, \dots, L$

$$h_\ell = \sigma_\ell(W_\ell h_{\ell-1})$$

$$e = h_L - target$$

#modulated forward pass

$$h_0^{err} = x + Fe$$

for $\ell = 1, \dots, L$

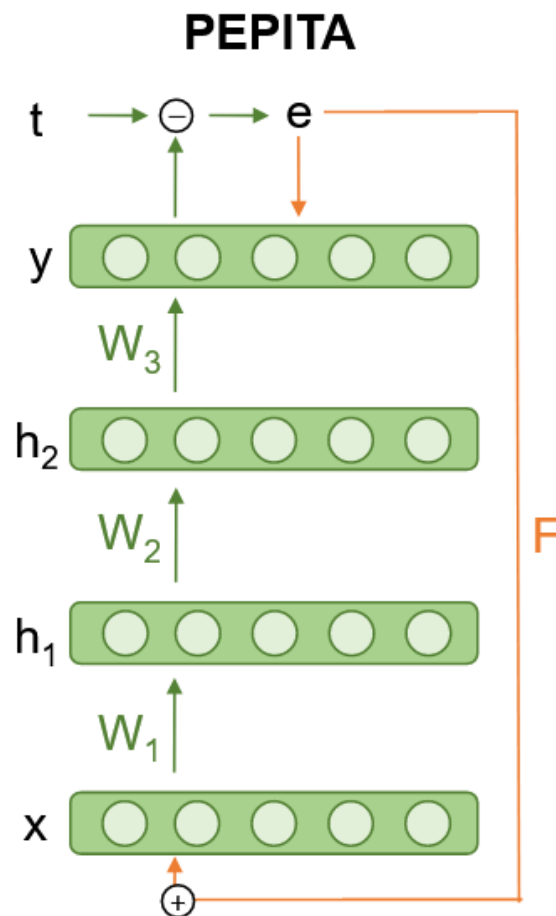
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if $\ell < L$:

$$\Delta W_\ell = (h_\ell - h_\ell^{err}) \cdot (h_{\ell-1}^{err})^T$$

else:

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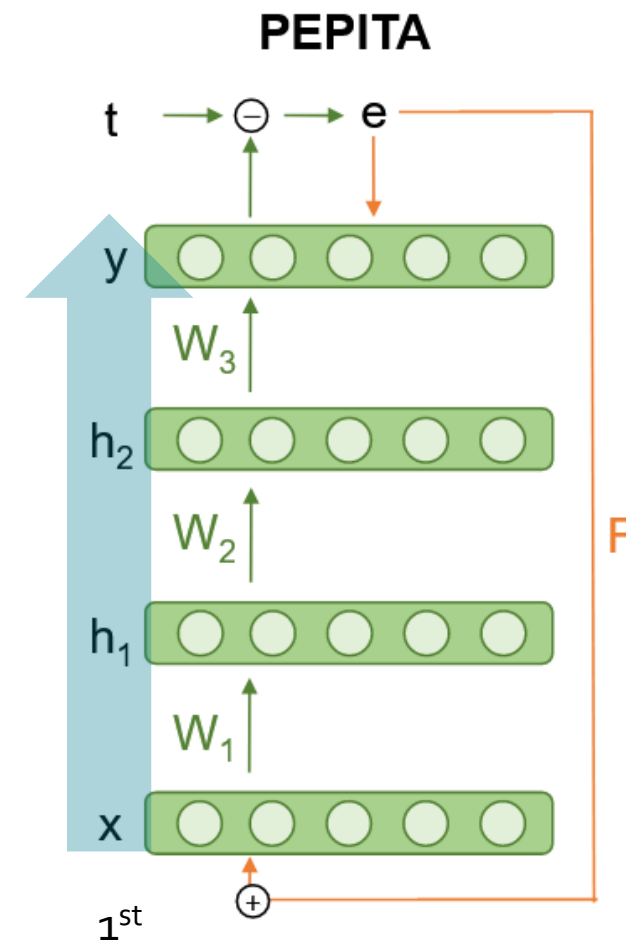
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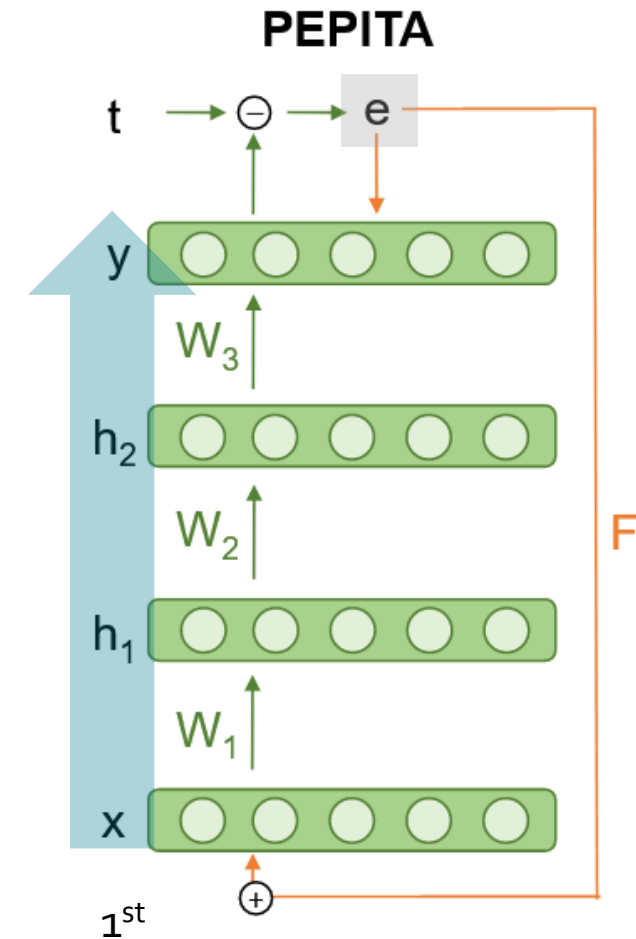
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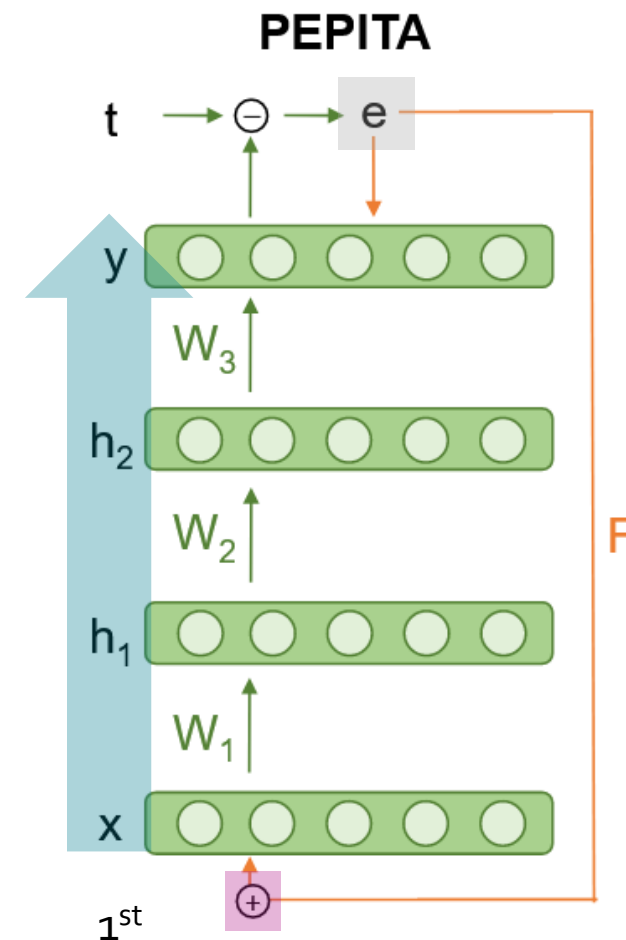
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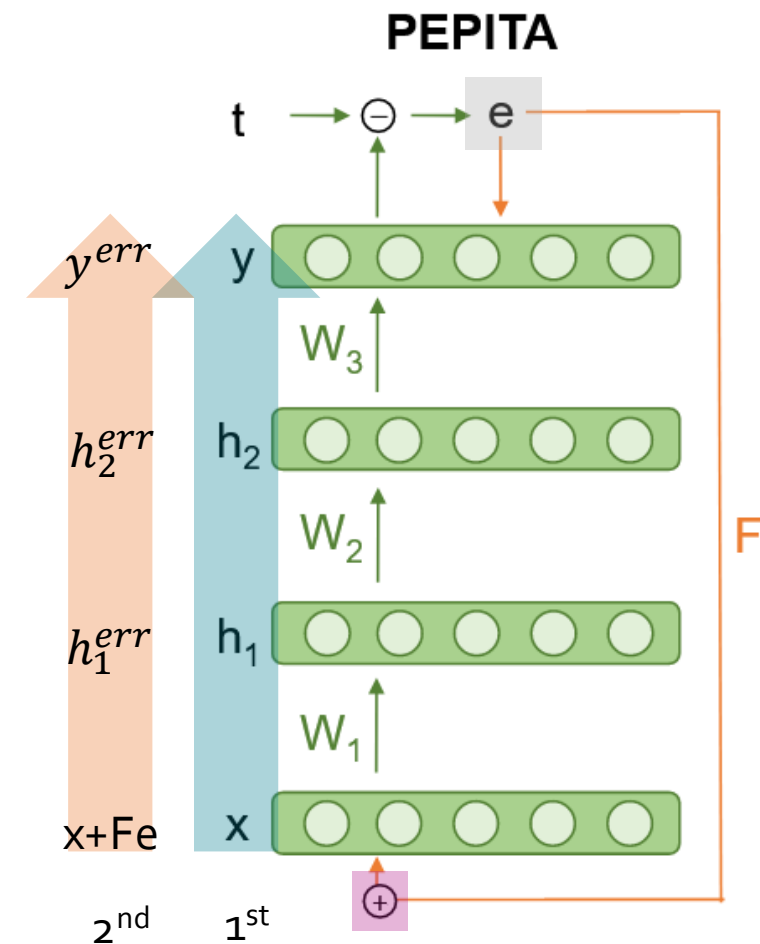
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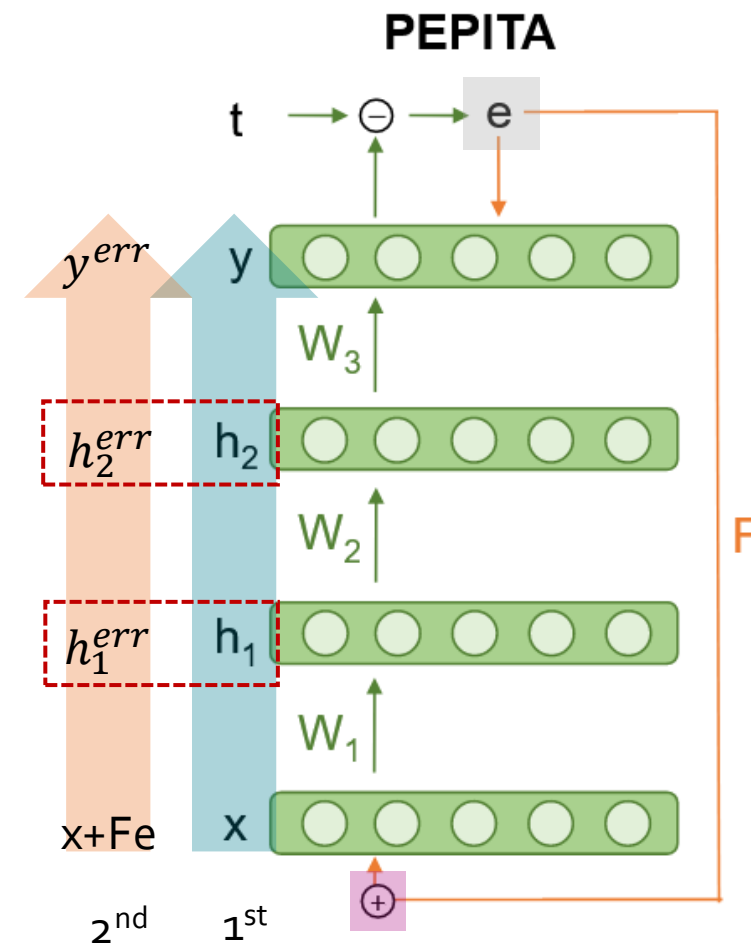
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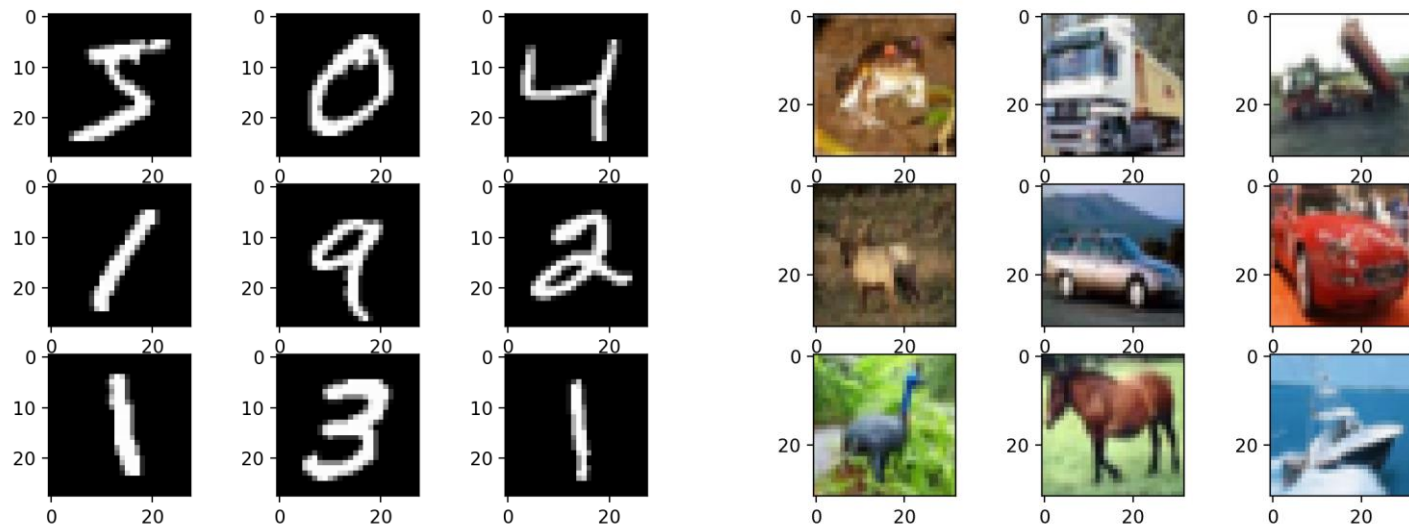
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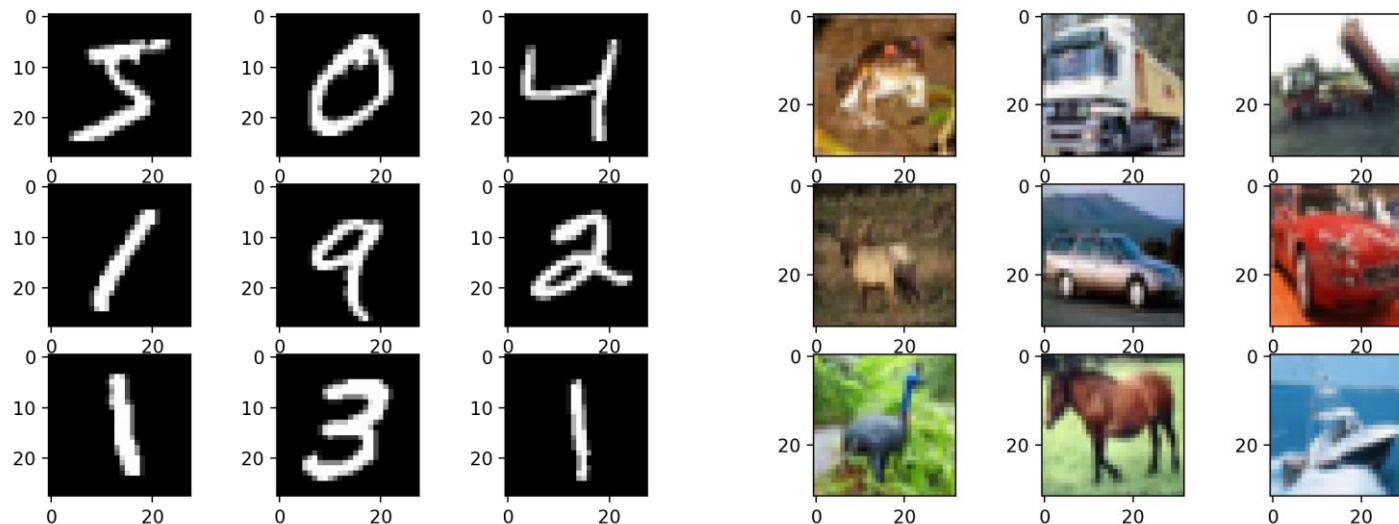


Testing PEPITA on image classification tasks - experimental results



	FULLY CONNECTED MODELS			CONVOLUTIONAL MODELS		
	MNIST	CIFAR10	CIFAR100	MNIST	CIFAR10	CIFAR100
BP	98.63±0.03	55.27±0.32	27.58±0.09	98.86±0.04	64.99±0.32	34.20±0.20
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DRTP	95.10±0.10	45.89±0.16	18.32±0.18	97.32±0.25	50.53±0.81	20.14±0.68
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Testing PEPITA on image classification tasks - experimental results



Architecture:
1 hidden layer +
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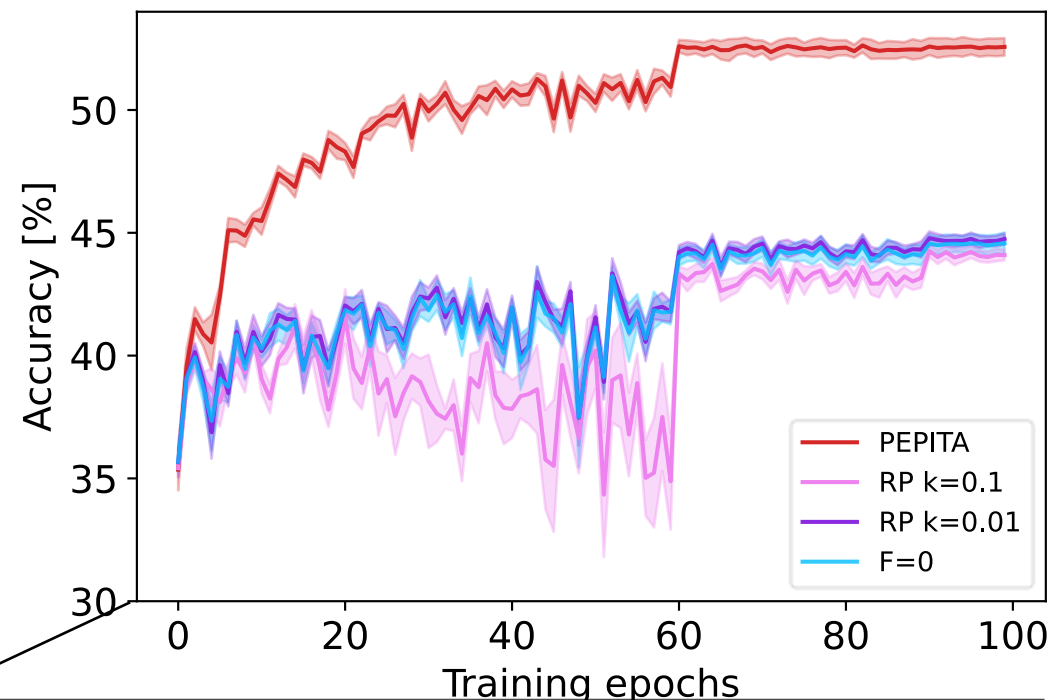
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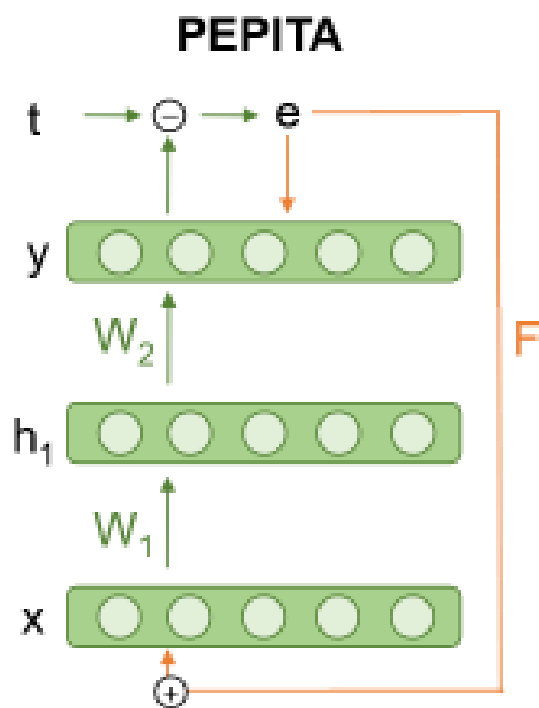


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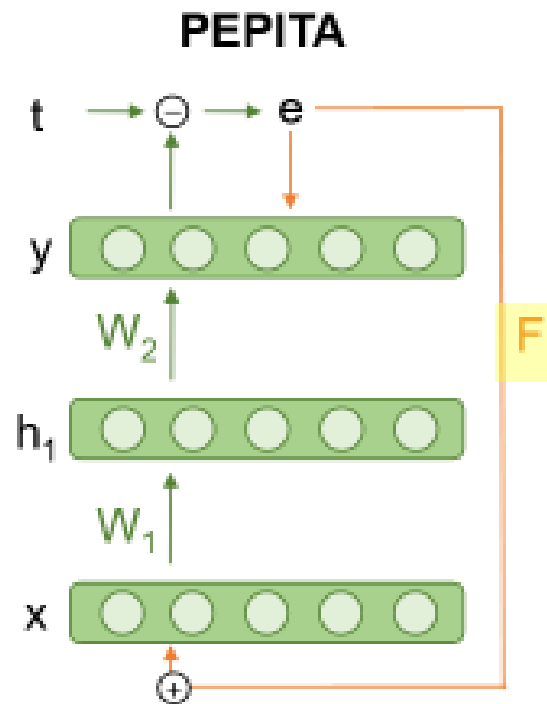
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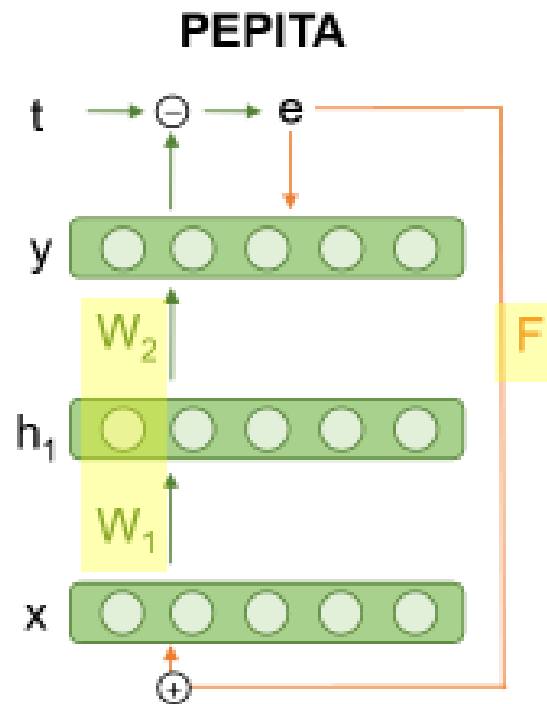
- Angle between
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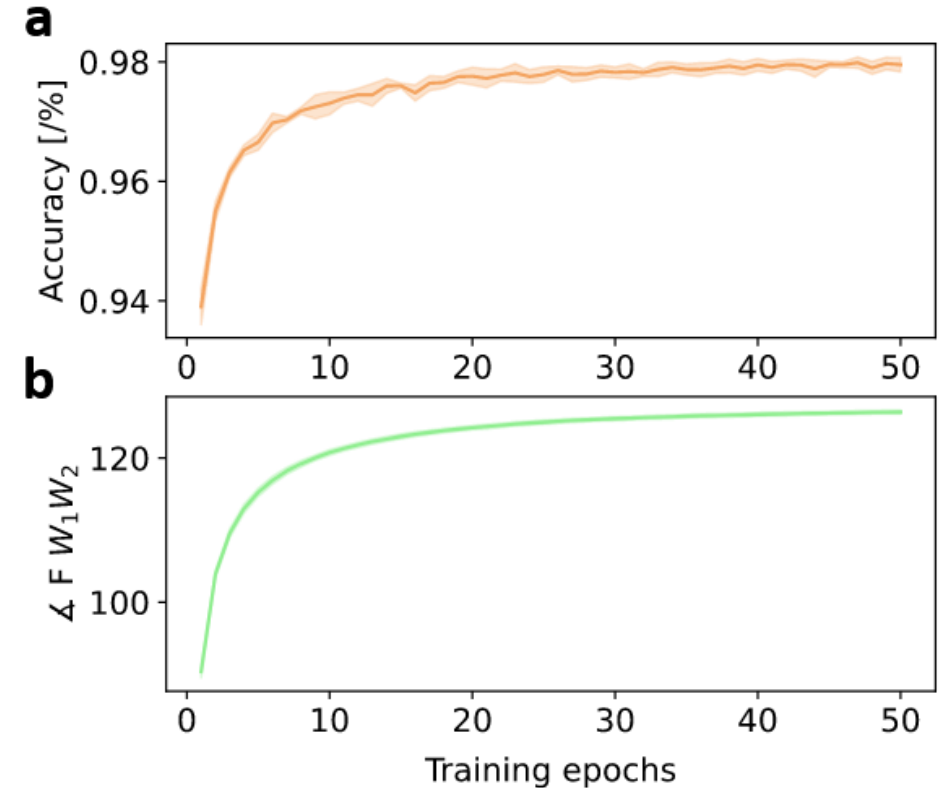
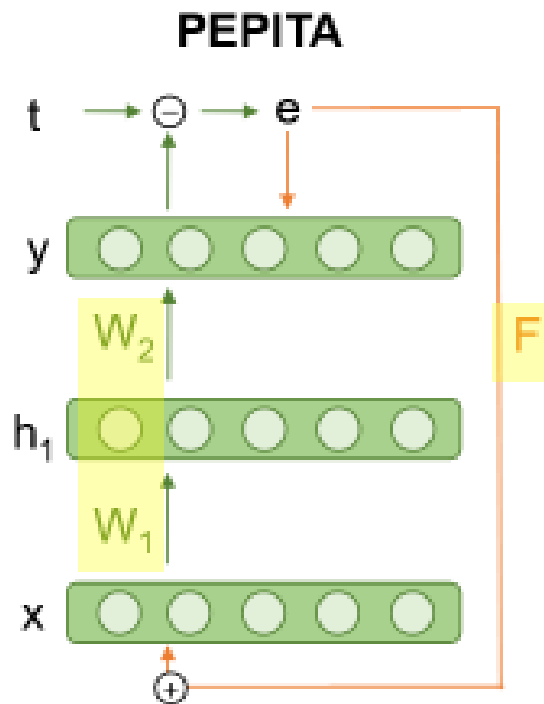
- Angle between
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Why it works: soft-antialignment

» Soft-antialignment

- Angle between
 - projection matrix F and
 - product between the forward weight matrices
- Evolution during learning \rightarrow soft antialignment
- Analytically proven for one-hidden layer linear network



Summary and Outlook

» PEPITA

- Is a novel training scheme relying only on **forward computations**
- Solves weight transport, freezing of neural activity, non-local weight updates and backward locking
- Achieves performance on-par with FA on simple image classification tasks

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» PEPITA

- Is a novel training scheme relying only on **forward computations**
- Solves weight transport, freezing of neural activity, non-local weight updates and backward locking
- Achieves performance on-par with FA on simple image classification tasks

» Challenges

- Performance **does not improve with depth**
- Different non-linearities
- Residual connection
- Training the F matrix

*Thank you for
your attention!*

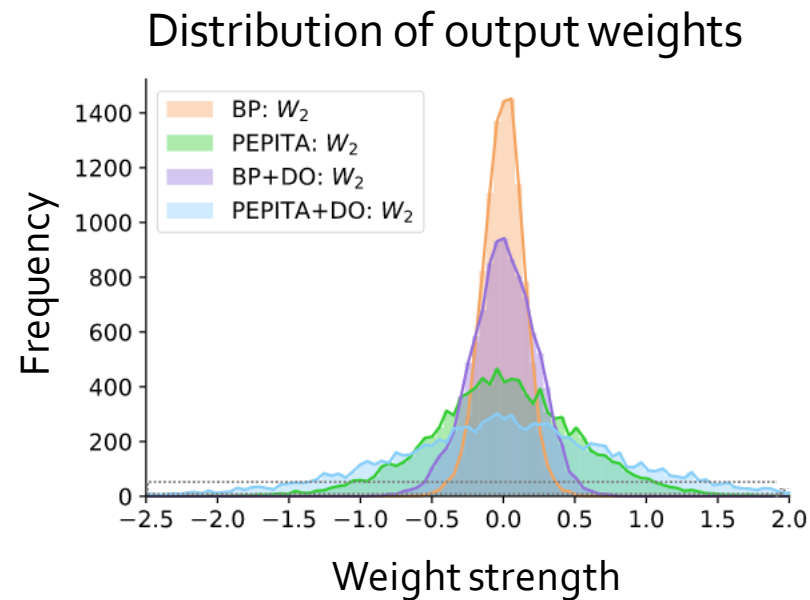
- » Questions?
- » Ideas?
- » Suggestions?

 gde@zurich.ibm.com

Weight distribution after training

» Wider weight distribution

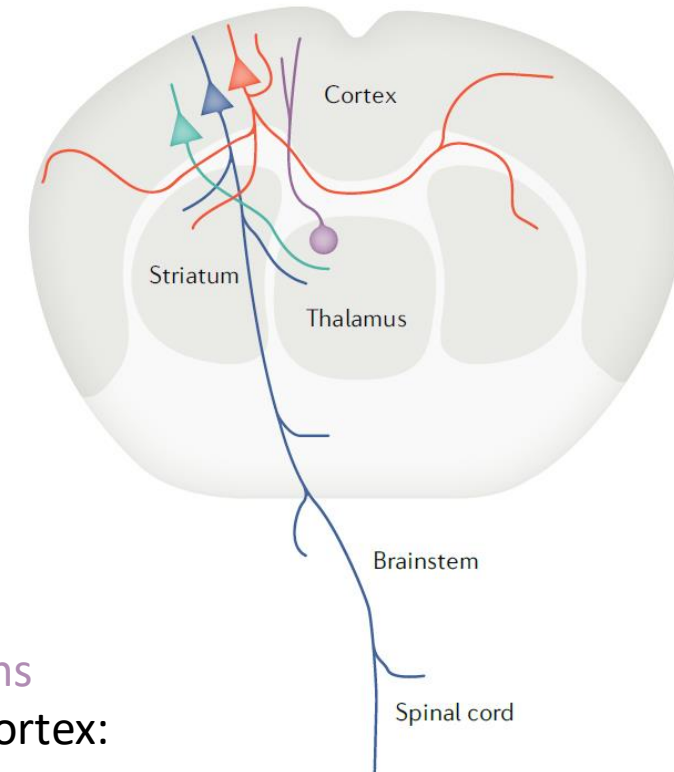
- PEPITA learns different solutions compared to BP



Analysis of PEPITA from a biological standpoint

PEPITA solves the BP's issues of biological plausibility, but it introduces additional elements:

- » Projection of the error onto the input through a fixed random matrix
 - Reminiscent of cortico-thalamo-cortical loops



In the thalamus:

- Thalamocortical (TC) neurons

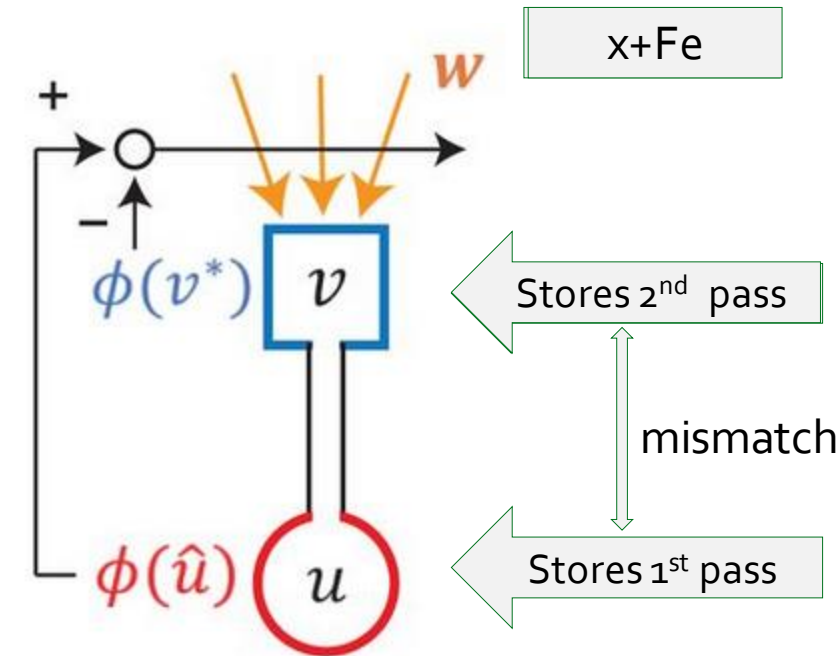
Excitatory neurons in the neocortex:

- Intratelencephalic (IT)
- Pyramidal tract (PT)
- Corticothalamic (CT) neurons

Analysis of PEPITA from a biological standpoint

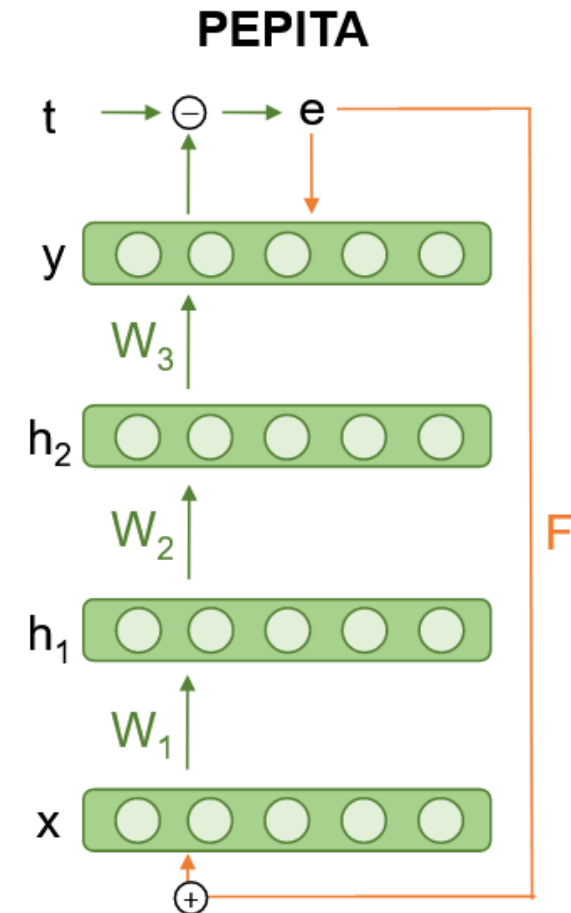
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- » Projection of the error onto the input through a fixed random matrix
 - Reminiscent of cortico-thalamo-cortical loops
- » Storing of the activation of the *Standard pass* until the *Modulated pass*
 - Can be implemented in biological neurons through mismatch between dendritic and somatic activity

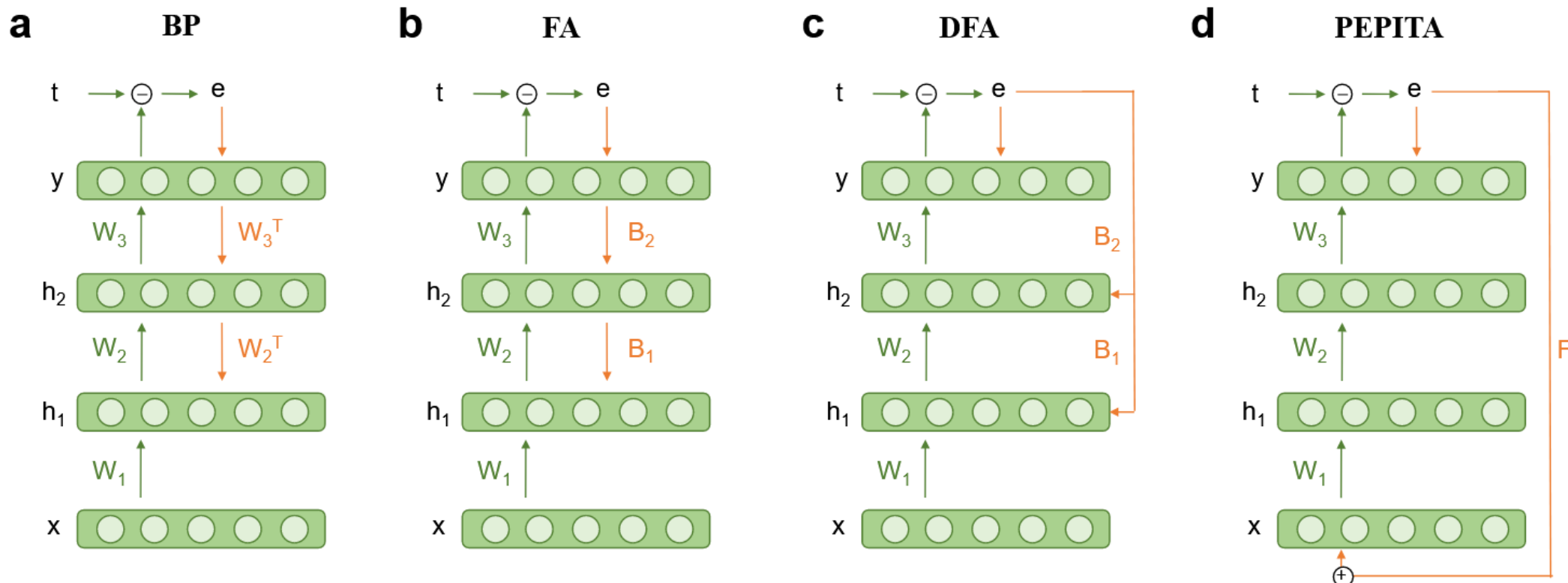


The PEPITA learning rule

- » **PEPITA** = Present the Error to Perturb the Input To modulate Activity
- » Substitutes the standard Forward+Backward scheme with **two Forward Passes**
 - *Standard Forward pass* → same as for standard algorithms
 - *Modulated Forward pass* → input is modulated by the error
- » F = **projection matrix** to add the error onto the input
- » Update relies **on difference of activations** between *Standard* and *Modulated pass*



PEPITA solves the biologically implausible aspects of BP



$$\Delta W_\ell = -(W_{\ell+1}^T \delta a_{\ell+1}) \odot f'(a_\ell) h_{\ell-1}^T$$

$$-(B_\ell^T \delta a_{\ell+1}) \odot f'(a_\ell) h_{\ell-1}^T$$

$$-(B_\ell^T e) \odot f'(a_\ell) h_{\ell-1}^T$$

$$(h_\ell - h_\ell^{err}) \cdot (h_{\ell-1}^{err})^T$$

WEIGHT-TRANSPORT-FREE

✗

✓

✓

✓

LOCAL RULE

✗

✗

✗

✓

FREEZING OF ACTIVITY

✗

✗

✗

✓

UPDATE-UNLOCKED

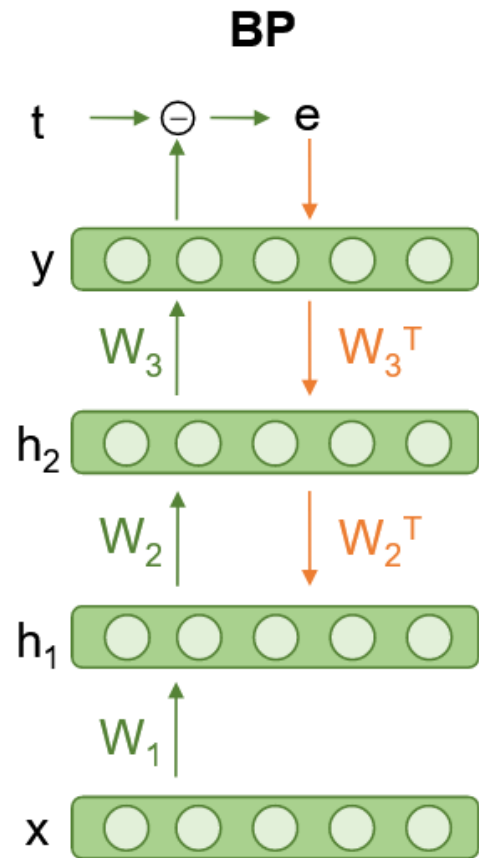
✗

✗

PARTIALLY

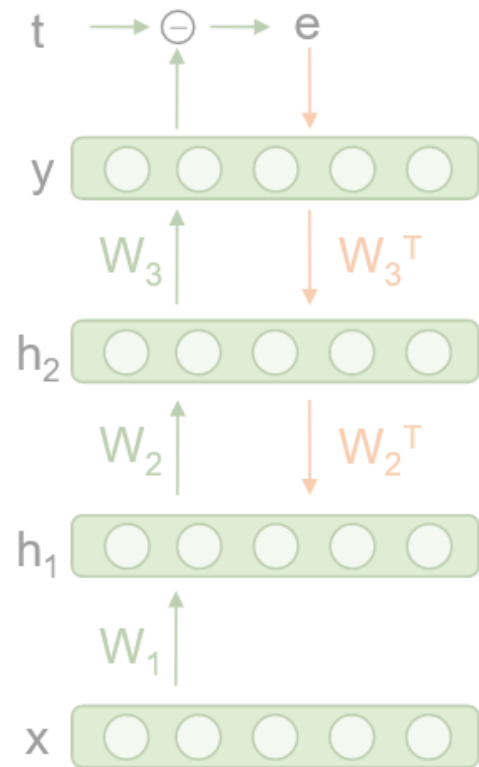
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Alternatives to BP: relaxing symmetry requirements



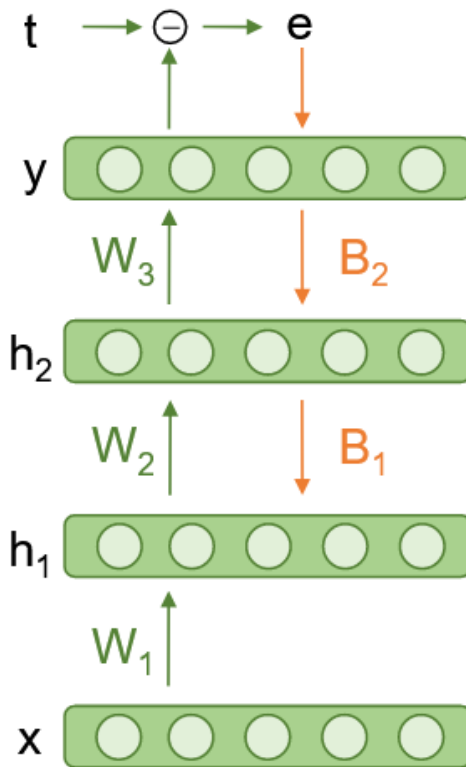
Alternatives to BP: relaxing symmetry requirements

BP



Rumelhart et al., 1995

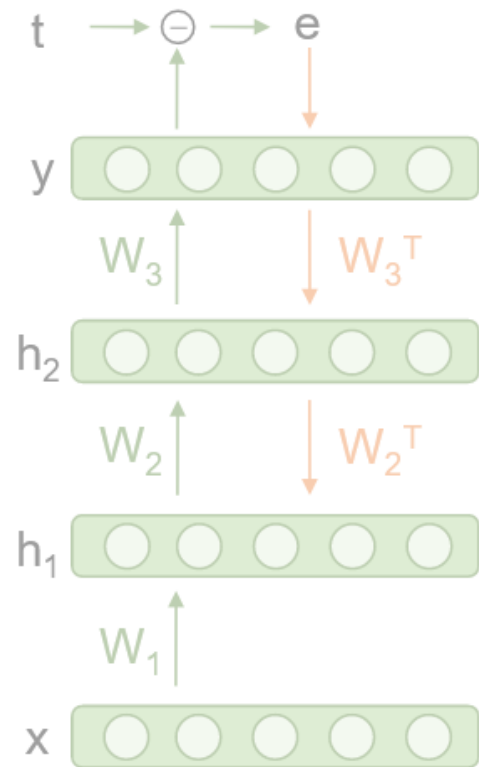
FA



Lillicrap et al., 2016

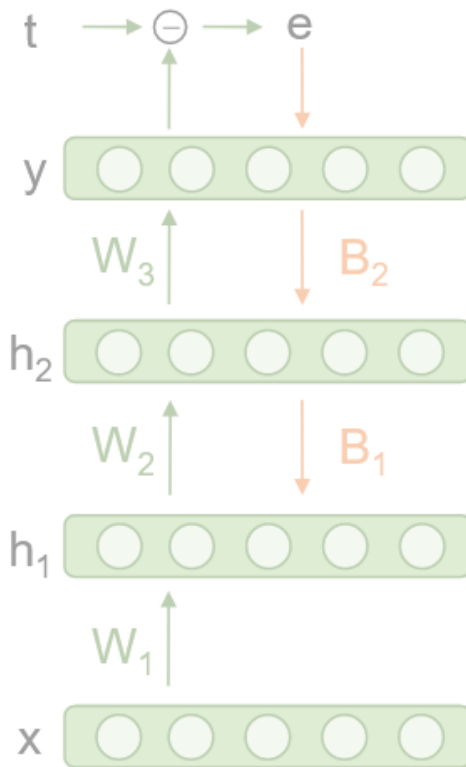
Alternatives to BP: relaxing symmetry requirements

BP



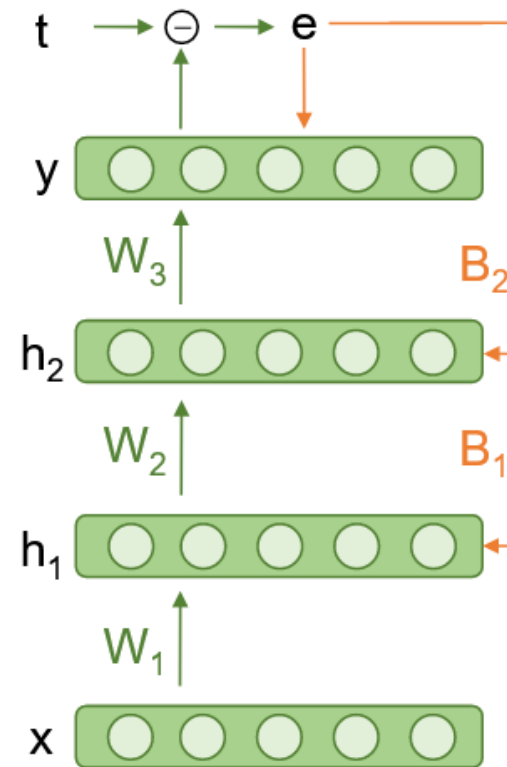
Rumelhart et al., 1995

FA



Lillicrap et al., 2016

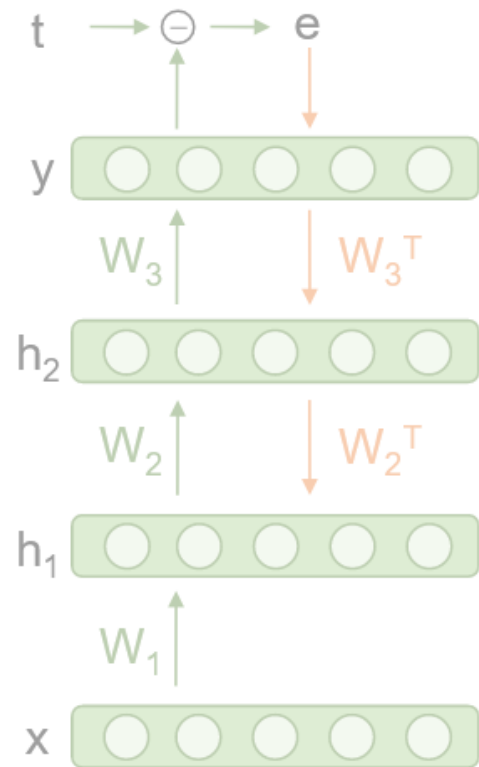
DFA



A. Nokland, 2016

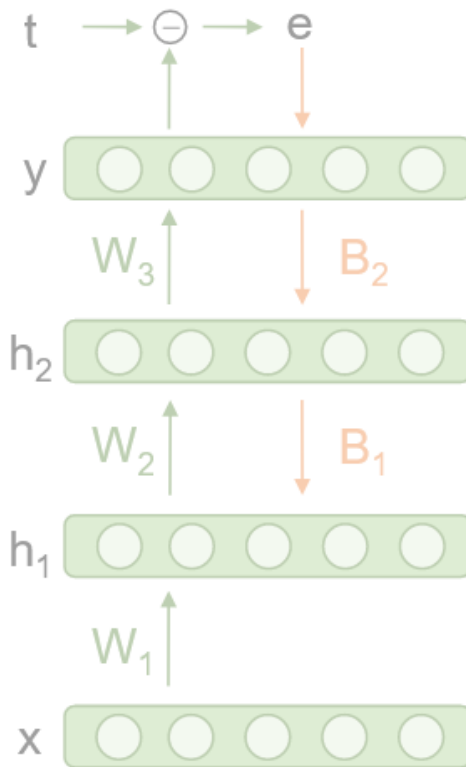
Alternatives to BP: relaxing symmetry requirements

BP



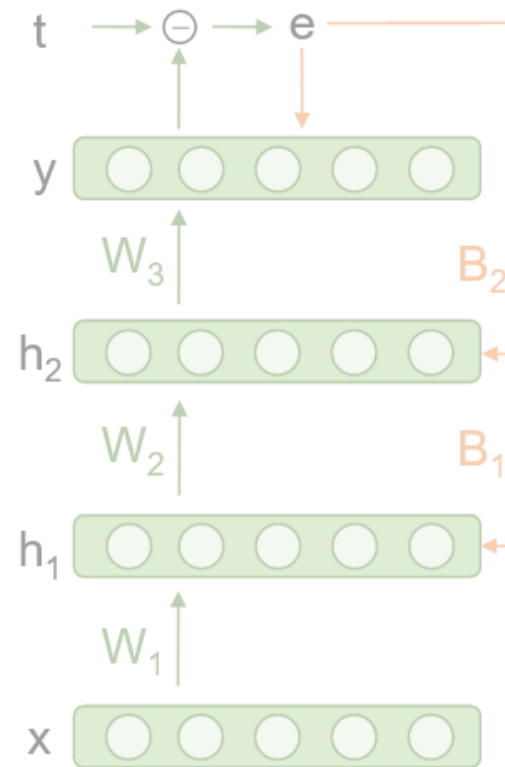
Rumelhart et al., 1995

FA



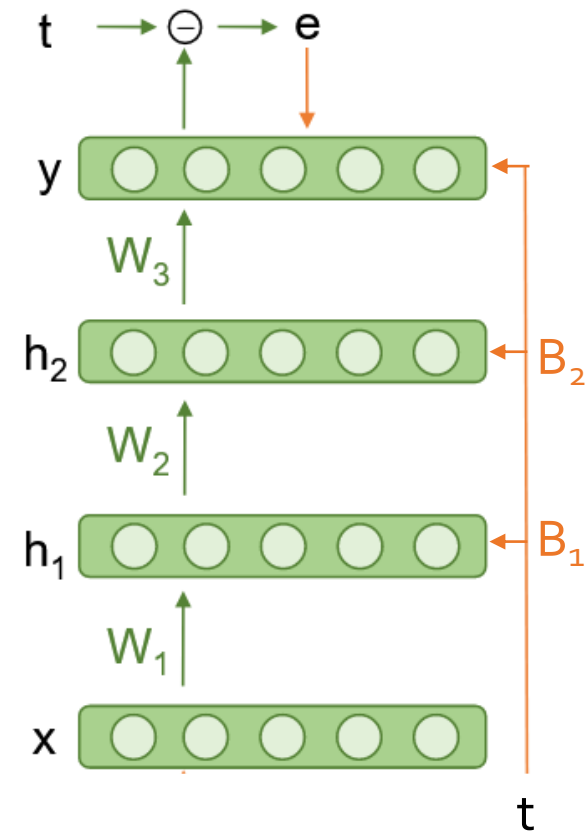
Lillicrap et al., 2016

DFA



A. Nokland, 2016

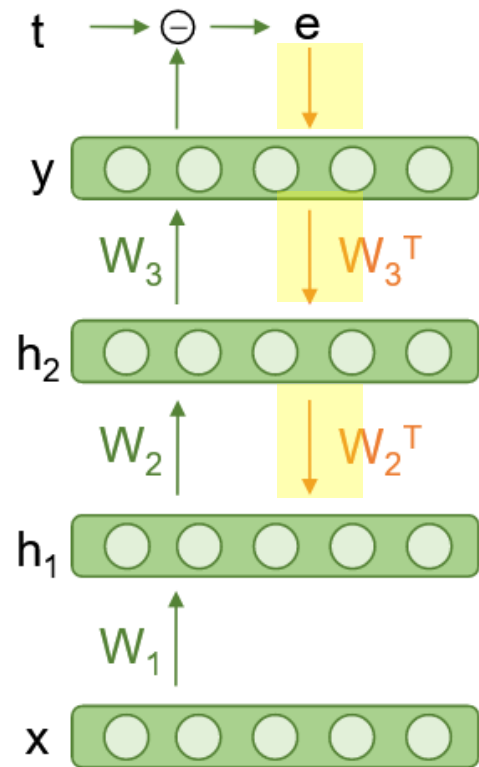
DRTP



Frenkel et al., 2019

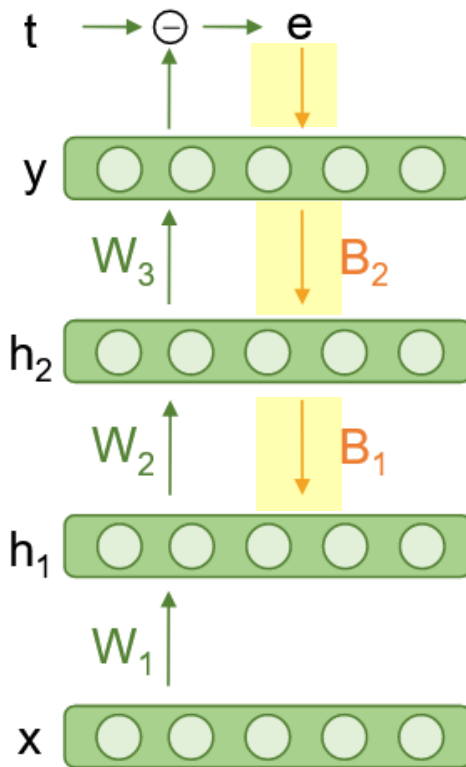
Alternatives to BP: relaxing symmetry requirements

BP



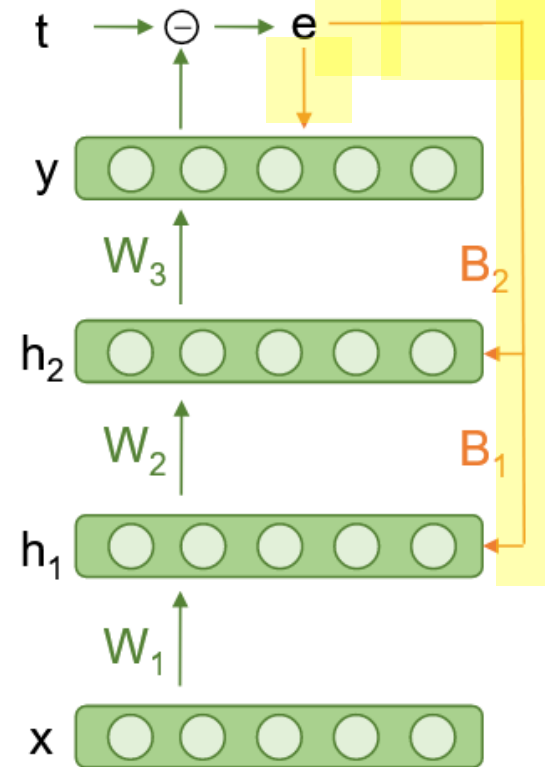
Rumelhart et al., 1995

FA



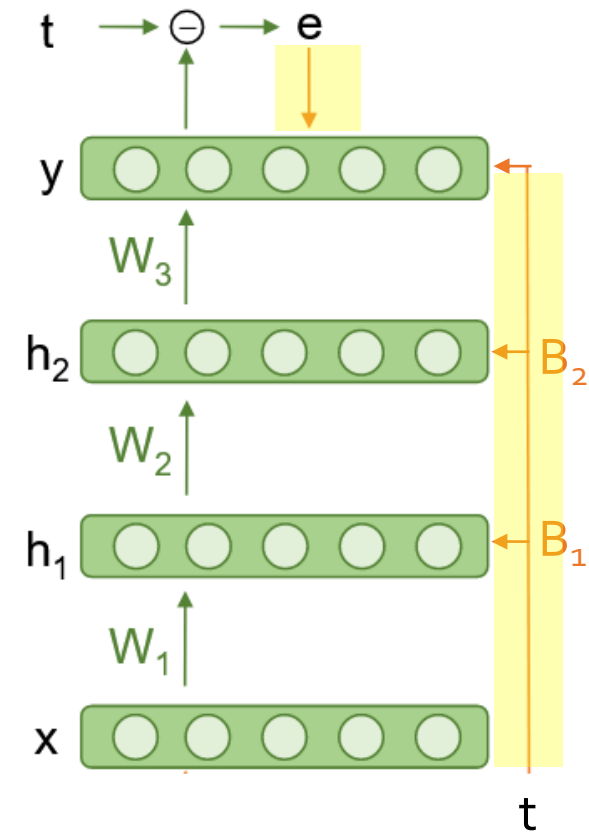
Lillicrap et al., 2016

DFA



A. Nokland, 2016

DRTP



Frenkel et al., 2019

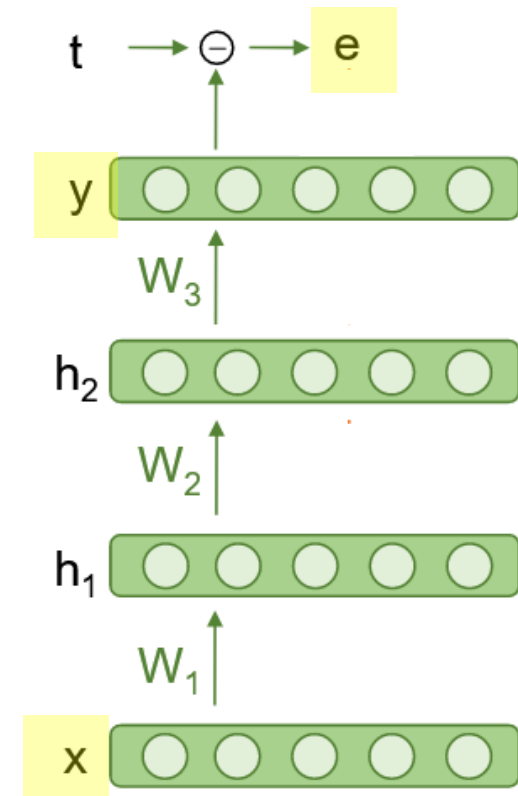
The backpropagation algorithm

» Forward pass

- Network's response to input
- Error function $e = y - t$
- Weight updates proportional to its negative gradient

» Backward pass

- Error signal flows backward through the network
- Computed recursively via the chain rule
- Update phase



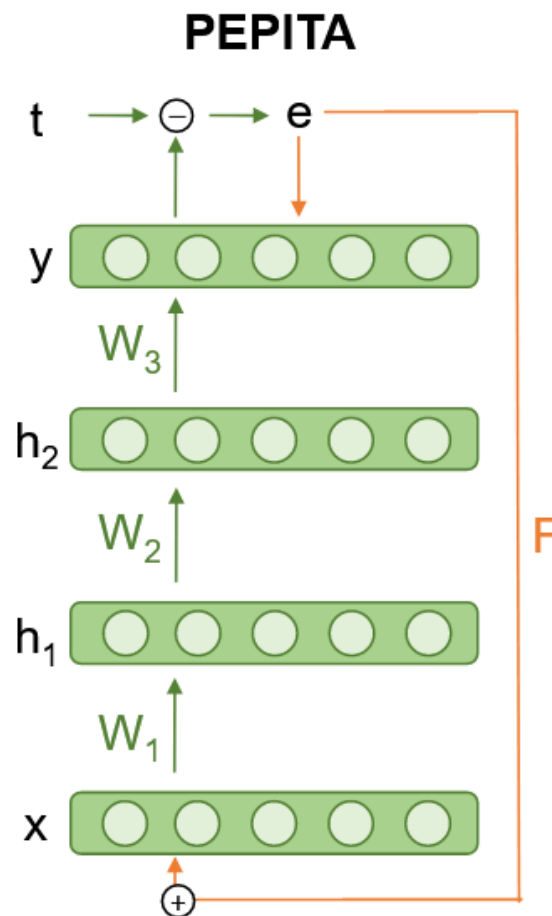
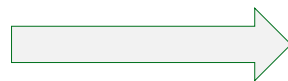
The Backward Pass

The backward pass implies:

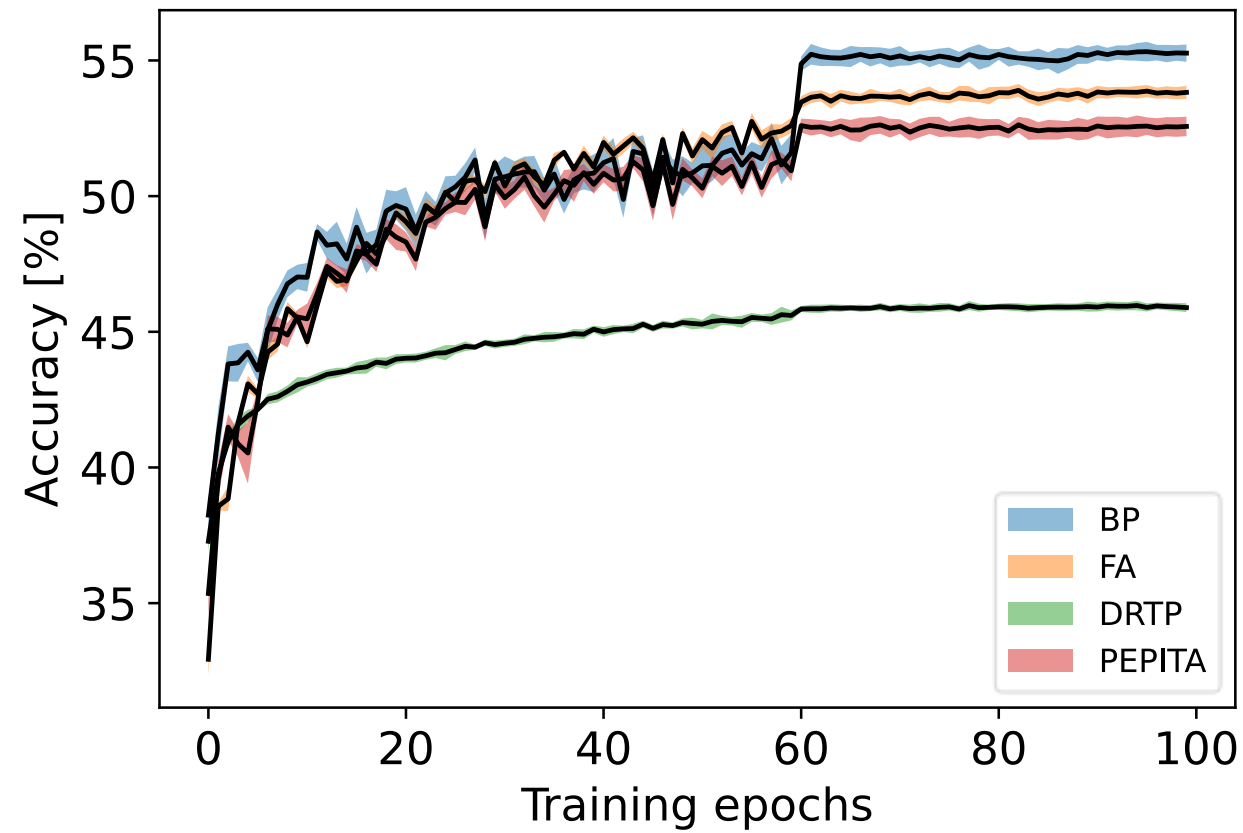
- » Non-locality
- » Frozen activity
- » At least partial update locking



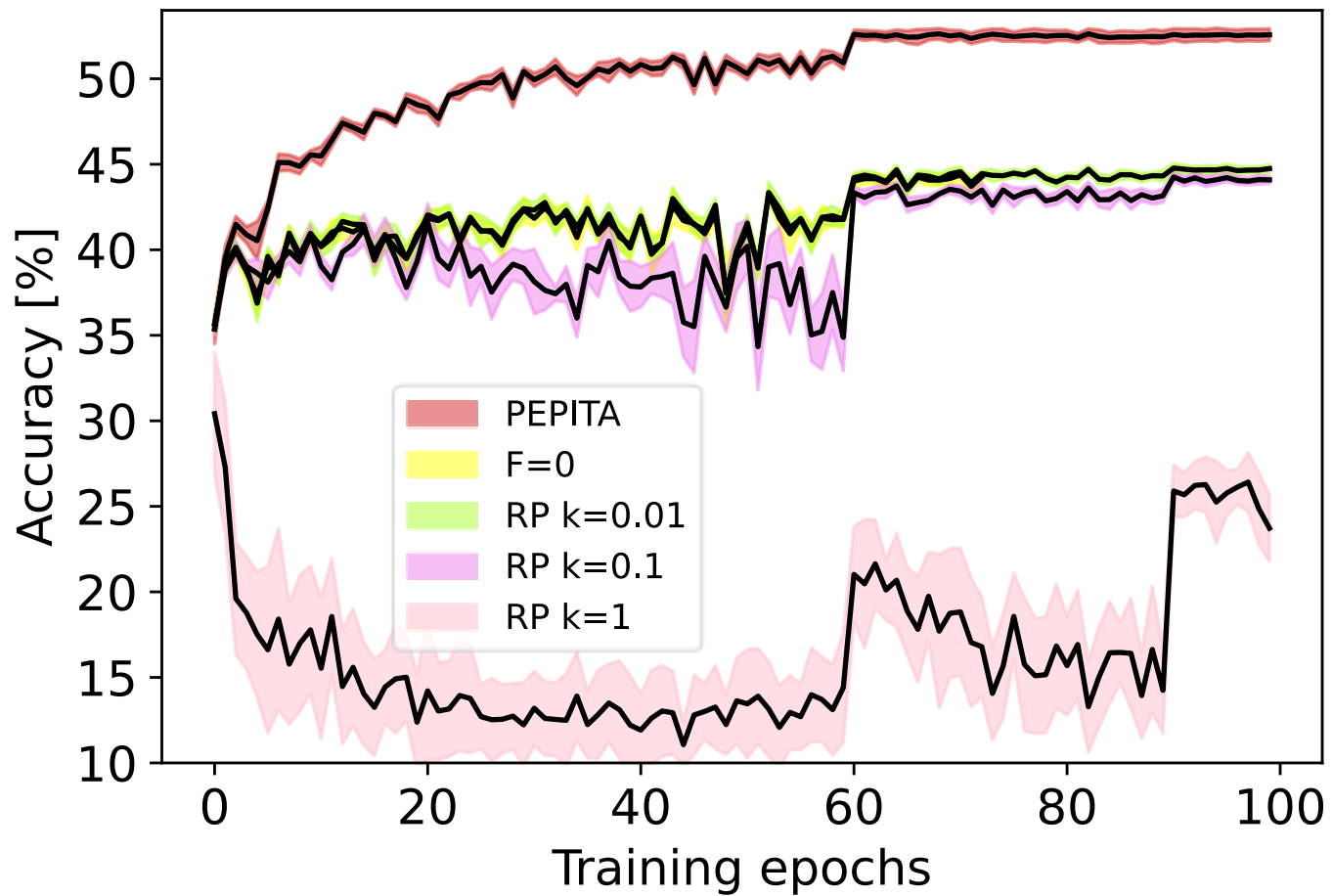
Remove the backward pass



Test curves on CIFAR-10

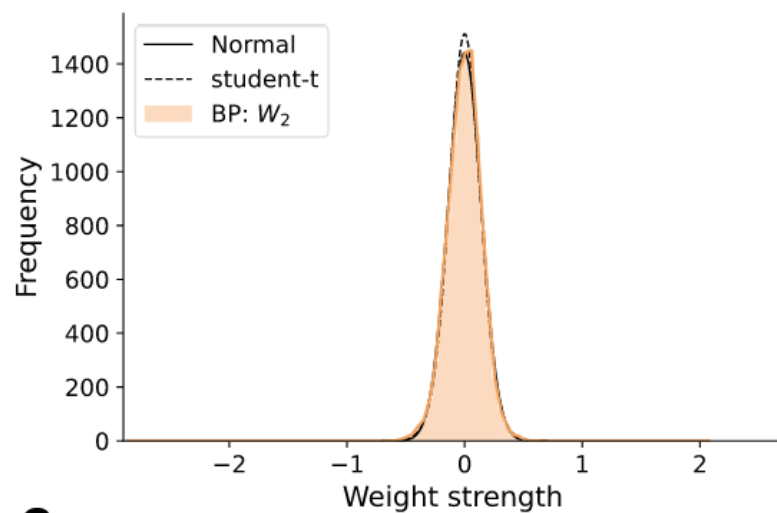


Error-based modulation is key for good performance

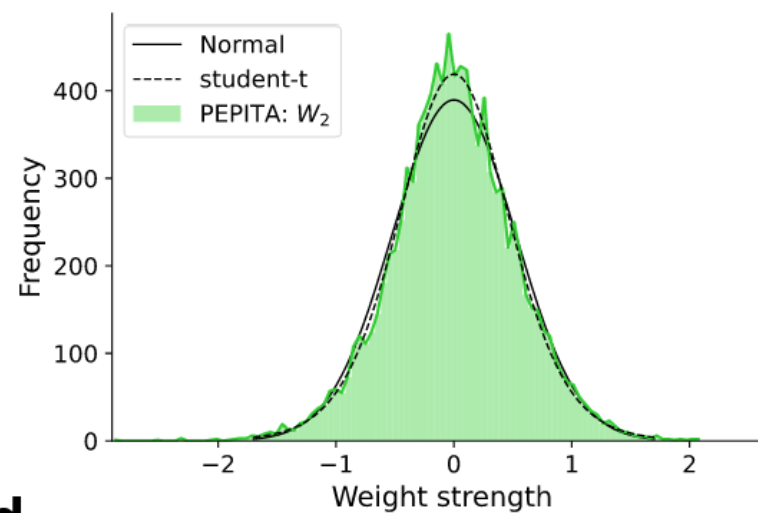


Weight distribution – heavy tailed

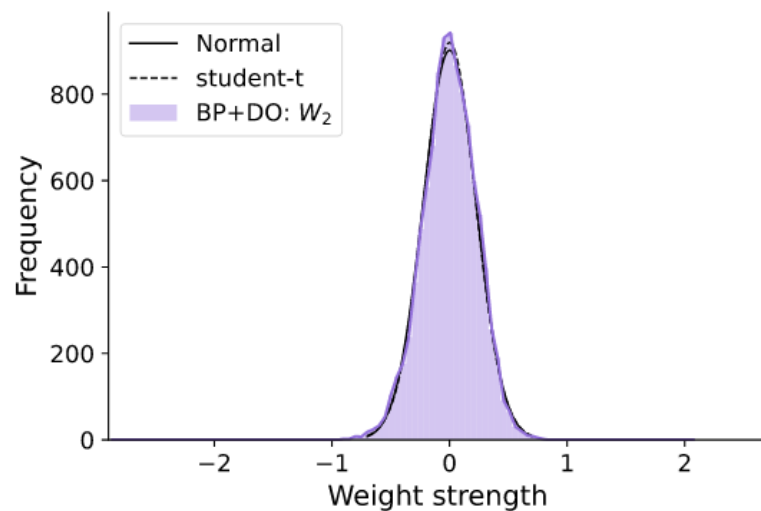
a



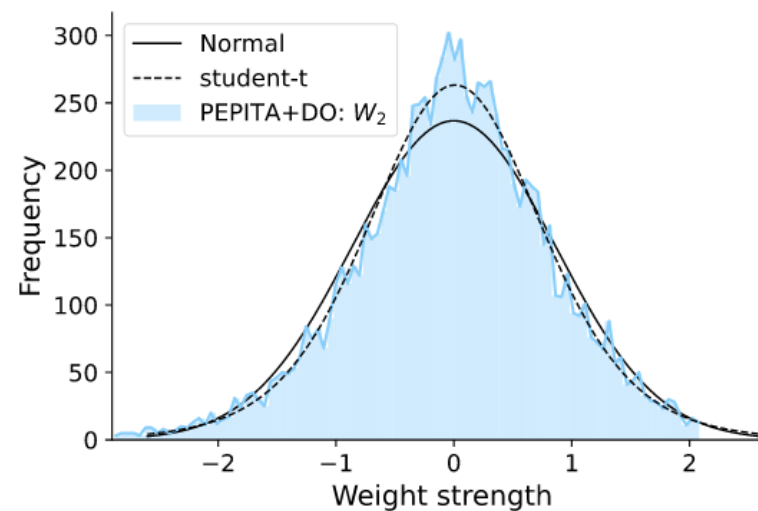
b



c



d

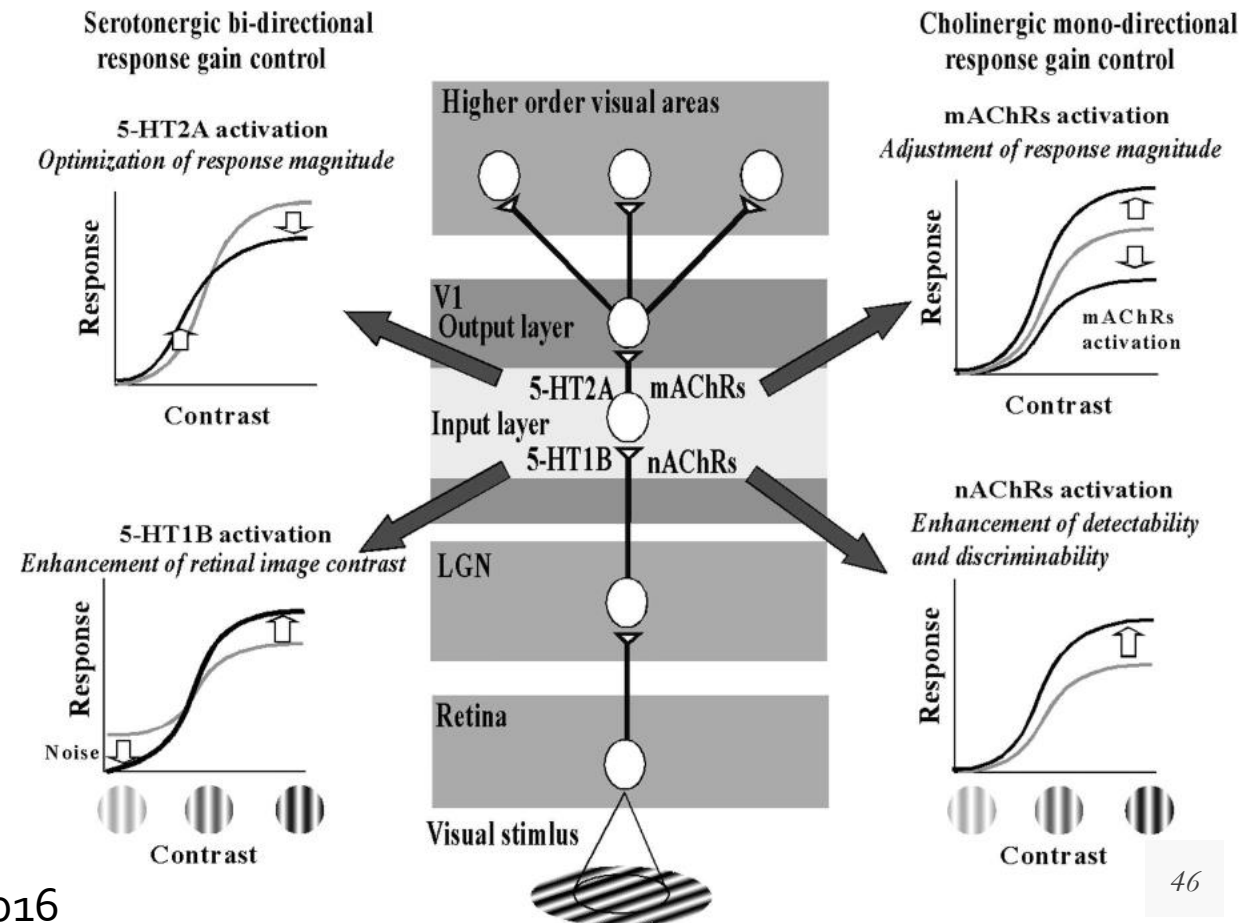


Analysis of PEPITA from a biological standpoint

PEPITA solves the BP's issues of biological plausibility, but it introduces additional elements:

» Projection of the error onto the input through a fixed random matrix

- Global neuromodulatory signals modulate activity in V1



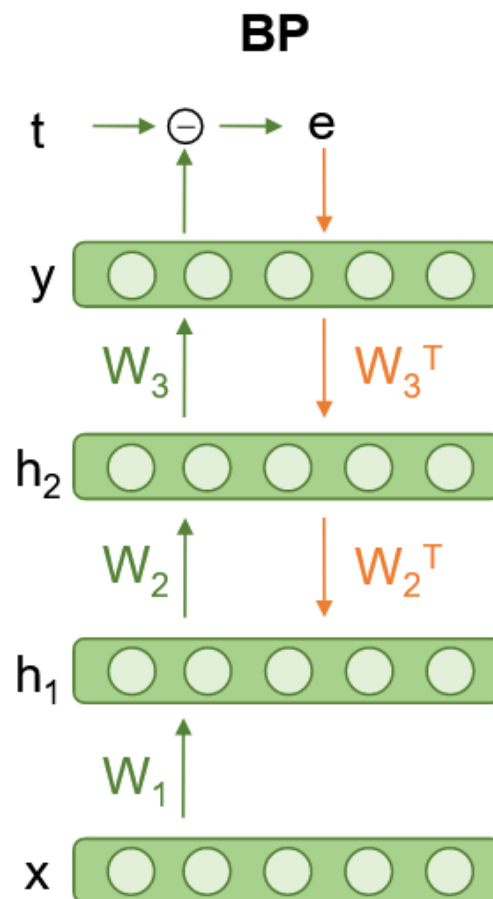
Alternatives to BP: Sign symmetry

» Asymmetric backpropagation

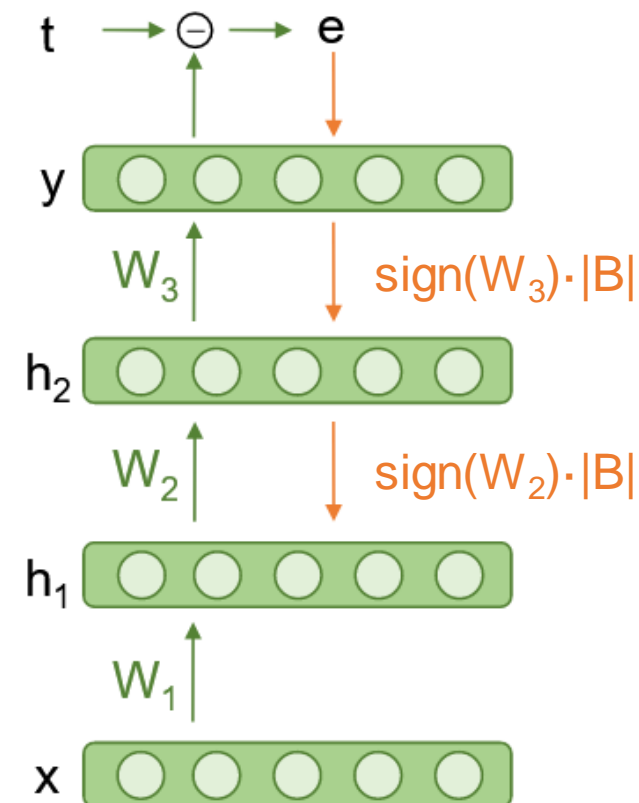
- Sign-concordant Feedback

» Relax weight symmetry requirement

- the magnitudes of feedback weights do not matter to performance
- the signs of feedback weights do matter — the more concordant signs between feedforward and their corresponding feedback connection

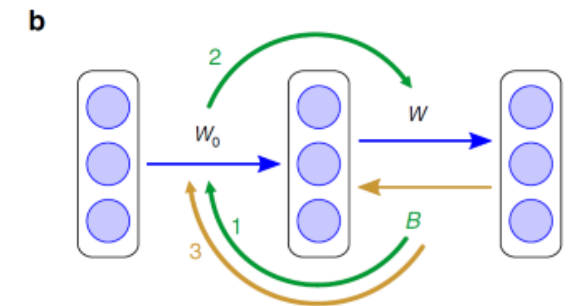
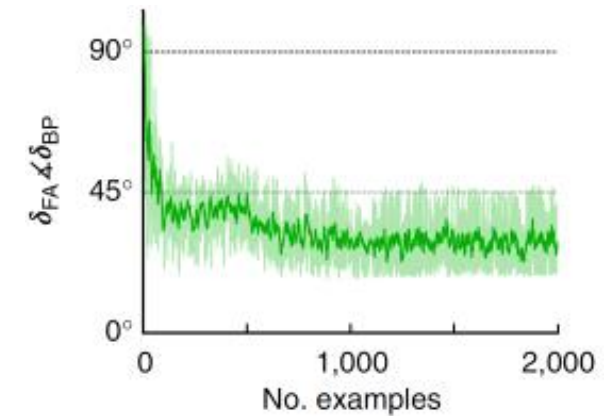
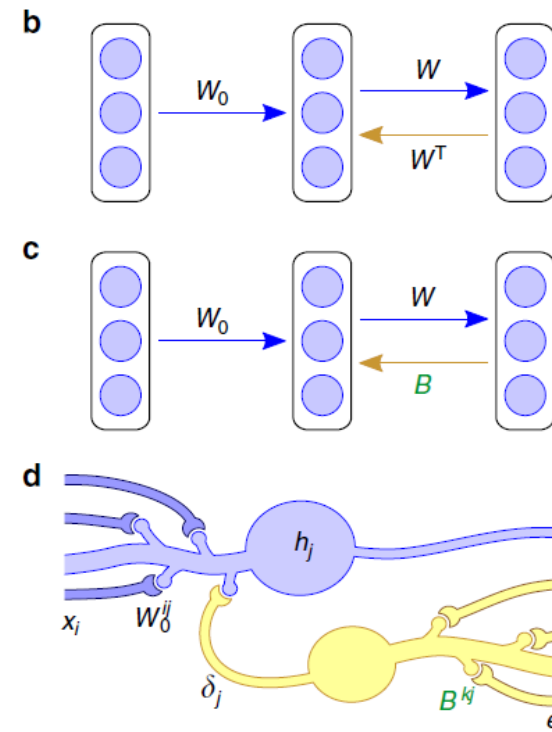


Sign-concordant Feedback



Alternatives to BP: Feedback Alignment

- » Precise symmetric connectivity between connected layers is not required to obtain quick learning
- » **Replaces W^T with a matrix of fixed random weights B**
 - Each neuron in the hidden layer receives a random projection of the error vector
 - Avoids all transport of synaptic weight information
- » **The circuit learns by encouraging a soft alignment of W with B^T**
 - The angle between modulator vectors prescribed by feedback alignment and backprop decreases
 - As W aligns with B^T , B begins to act like W^T , sending useful teaching signals to the hidden units



Alternatives to BP: Direct and Indirect Feedback Alignment

» The FA principle is used for training hidden layers more independently from the rest of the network

» **Feedback path disconnected from the forward path**

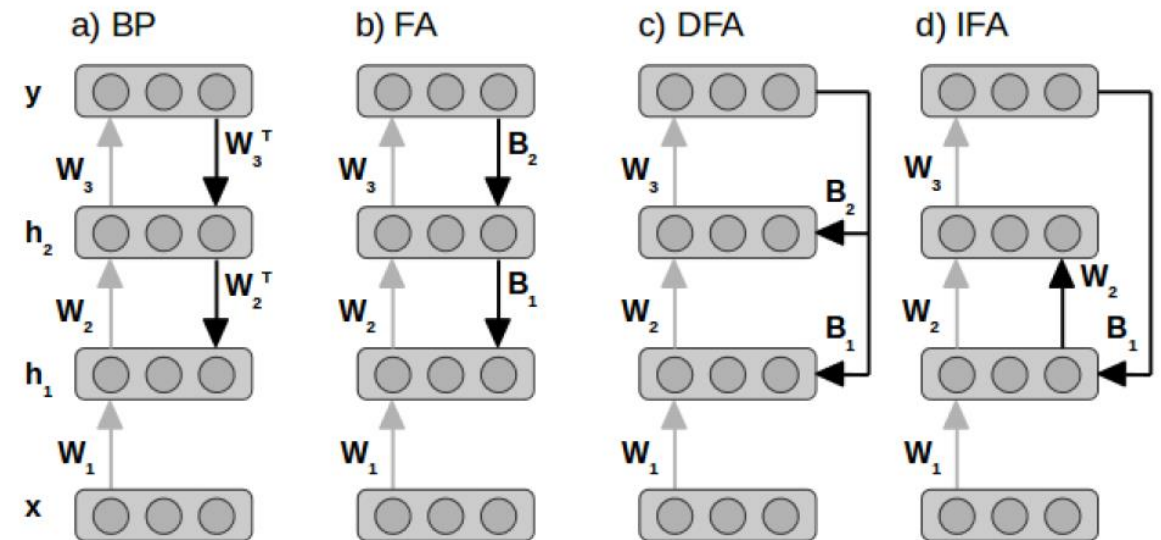
- Possibility that the error in the feedback layer is represented by neurons not participating in the forward pass
- layer is no longer reciprocally connected to the layer above

» **DFA**

- direct feedback path to each hidden layer

» **IFA**

- direct feedback path connecting to the first hidden layer
- then visiting every layer on its way forward



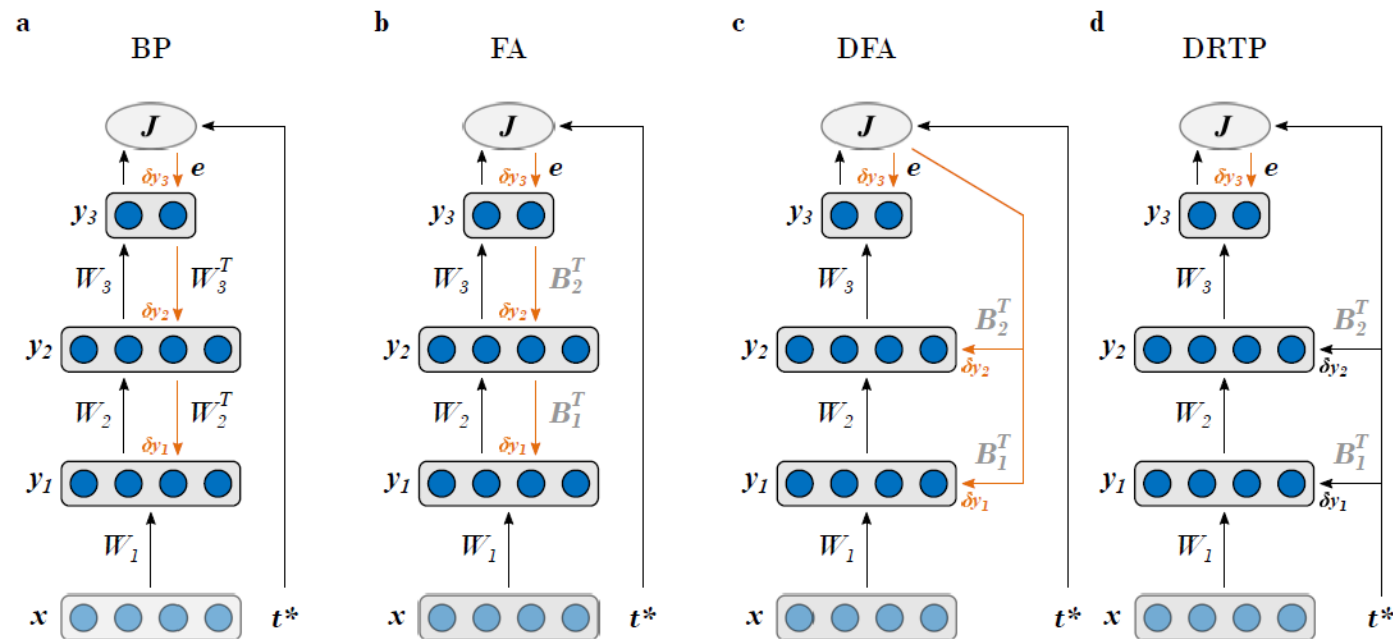
MODEL	BP	FA	DFA
7x240 Tanh	2.16 ± 0.13%	2.20 ± 0.13% (0.02%)	2.32 ± 0.15% (0.03%)
100x240 Tanh			3.92 ± 0.09% (0.12%)
1x800 Tanh	1.59 ± 0.04%	1.68 ± 0.05%	1.68 ± 0.05%
2x800 Tanh	1.60 ± 0.06%	1.64 ± 0.03%	1.74 ± 0.08%
3x800 Tanh	1.75 ± 0.05%	1.66 ± 0.09%	1.70 ± 0.04%
4x800 Tanh	1.92 ± 0.11%	1.70 ± 0.04%	1.83 ± 0.07% (0.02%)
2x800 Logistic	1.67 ± 0.03%	1.82 ± 0.10%	1.75 ± 0.04%
2x800 ReLU	1.48 ± 0.06%	1.74 ± 0.10%	1.70 ± 0.06%
2x800 Tanh + DO	1.26 ± 0.03% (0.18%)	1.53 ± 0.03% (0.18%)	1.45 ± 0.07% (0.24%)
2x800 Tanh + ADV	1.01 ± 0.08%	1.14 ± 0.03%	1.02 ± 0.05% (0.12%)

Test error on MNIST

A. Nokland, 2016

Alternatives to BP: Direct Random Target Propagation

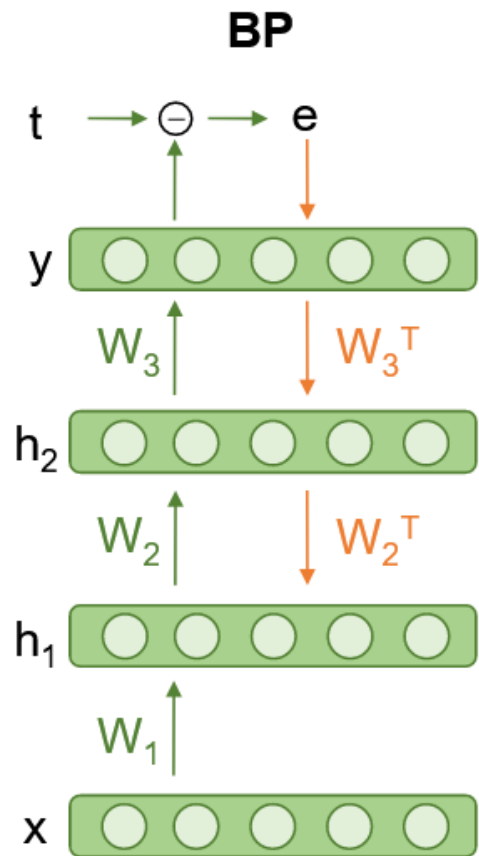
- » The error sign provides useful modulatory signals to multi-layer networks
 - Targets (i.e. one-hot-encoded labels) used in place of the output error
 - Targets are projected onto the hidden layers
- » Fully solves both the weight transport and the update locking problems
- » BUT: lower performance than BP, FA, DFA



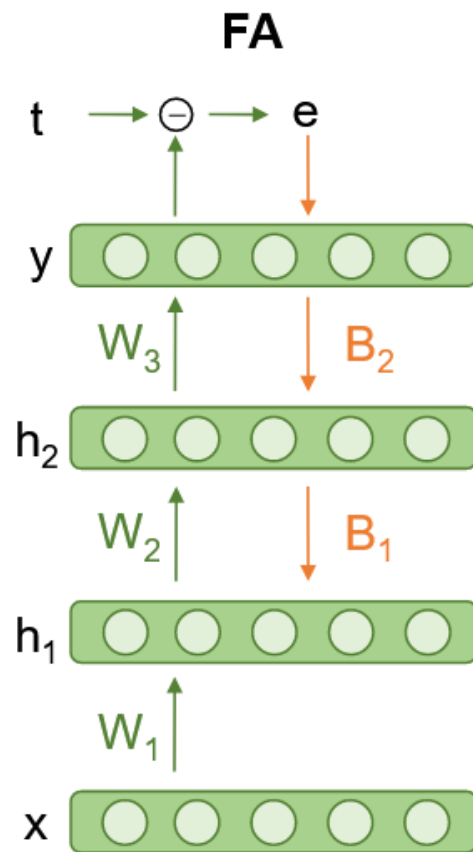
Network		BP	FA	DFA	DRTP
FC1-500	DO 0.0	1.72±0.08%	1.92±0.08%	2.59±0.11%	4.58±0.12%
	DO 0.1	1.55±0.03%	1.66±0.06%	2.17±0.10%	4.65±0.13%
	DO 0.25	1.64±0.06%	1.73±0.05%	2.32±0.08%	5.36±0.11%

Test error on MNIST

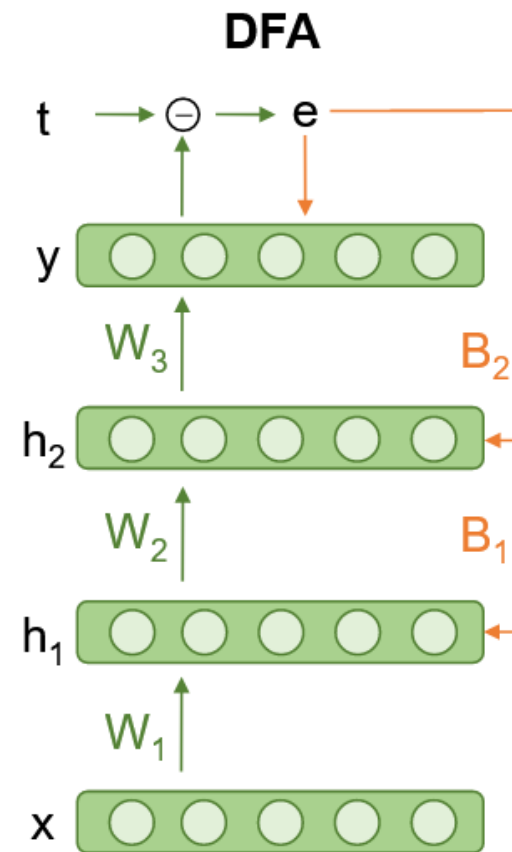
Training without a backward path: modulating the input through the error



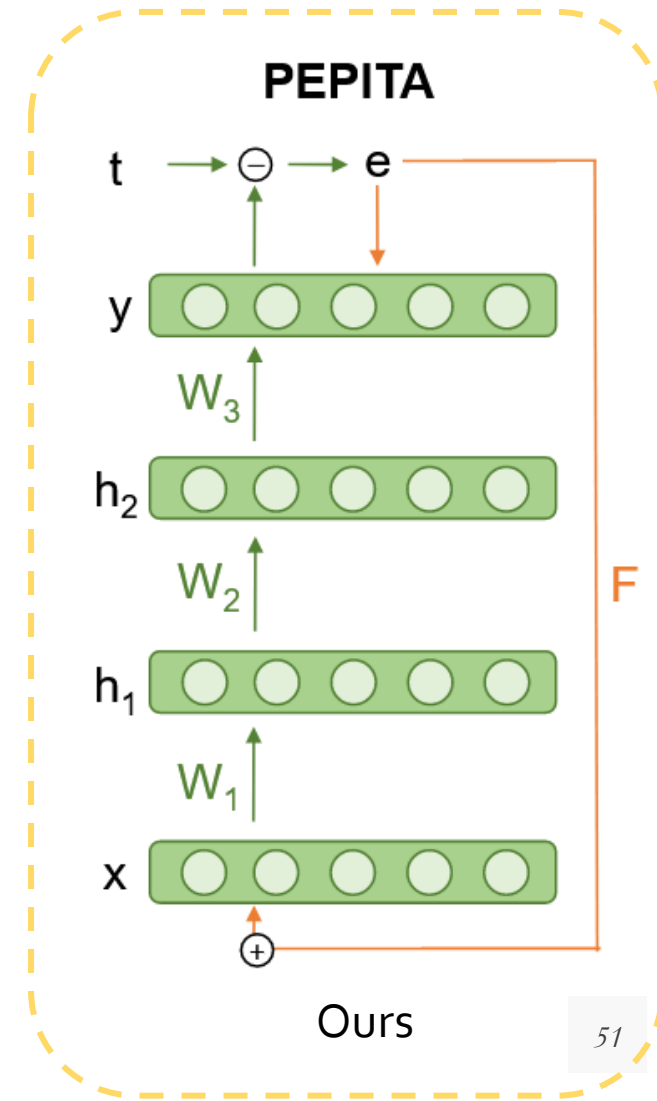
Rumelhart et al., 1995



Lillicrap et al., 2016



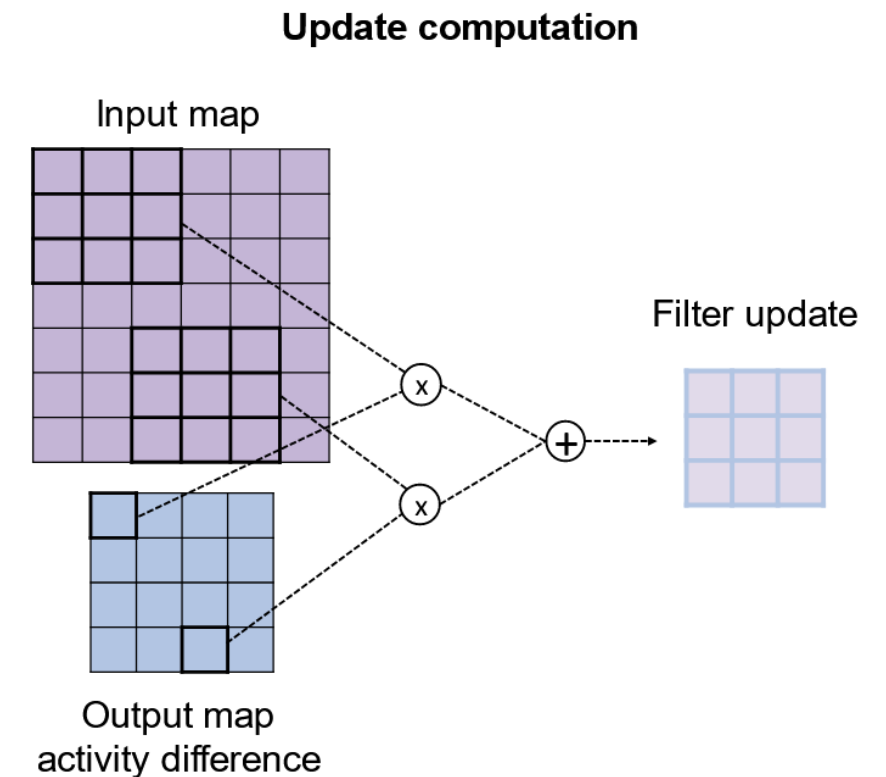
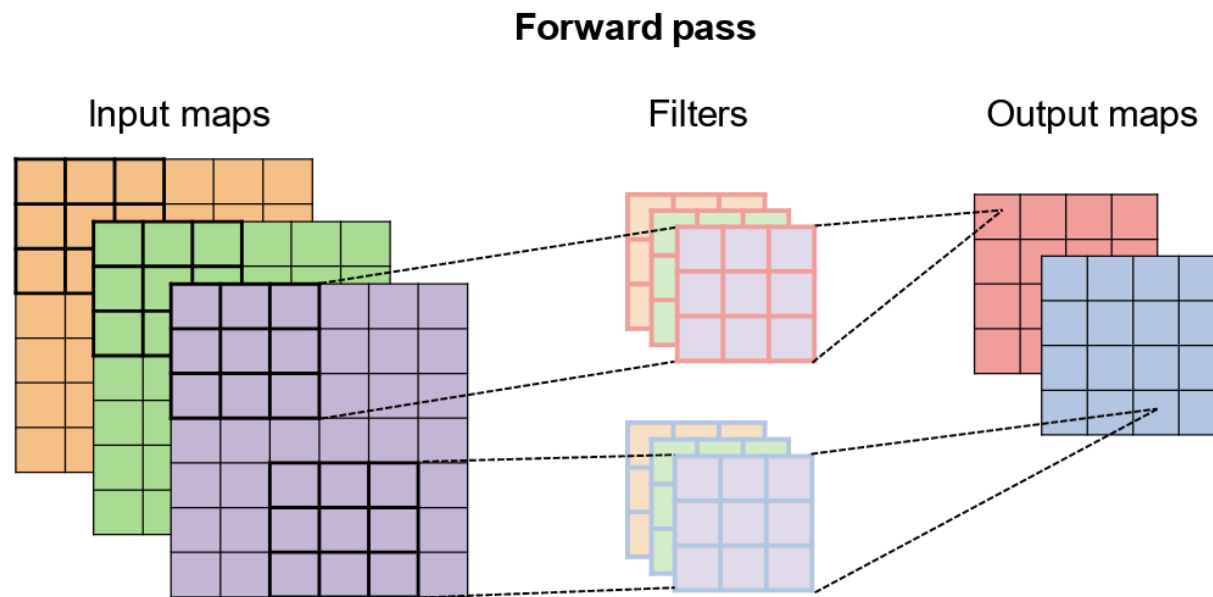
Nokland, 2016



Ours

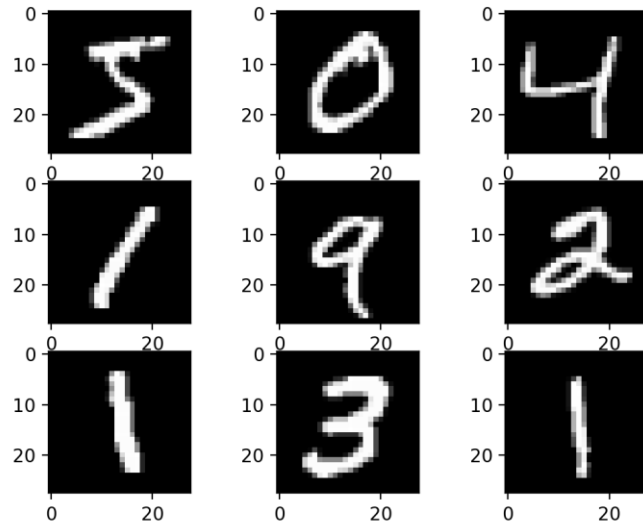
The PEPITA learning rule for Convolutional Neural Networks

- » Same approach with *Standard* and *Modulated pass*
- » Takes into account *weight sharing* of convolutional layers
- » Each filter is updated based on the contributions of each *input-map-region* – *output-map-element* pair

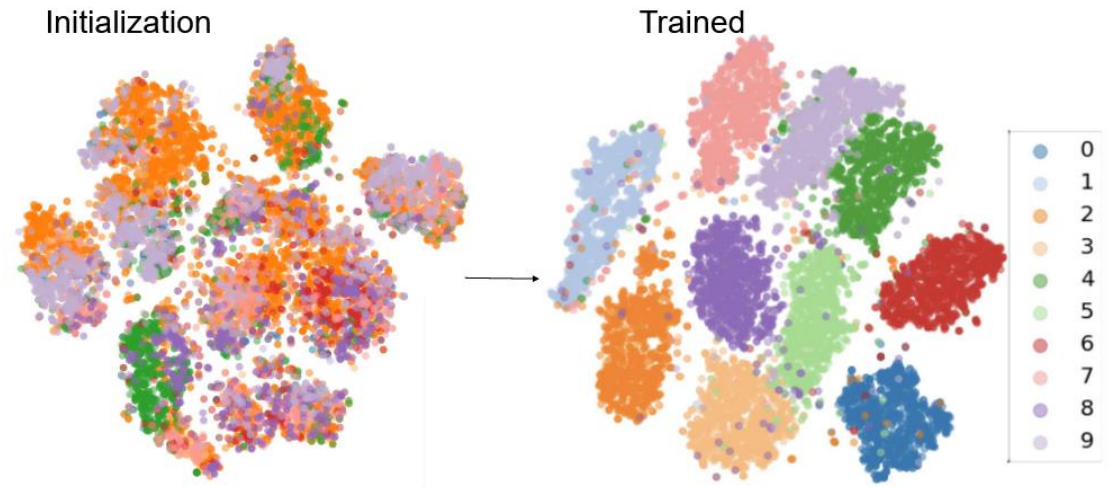


Testing PEPITA on image classification tasks - experimental results

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- » PEP
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	FULLY CONNECTED MODELS			CONVOLUTIONAL MODELS		
	MNIST	CIFAR10	CIFAR100	MNIST	CIFAR10	CIFAR100
BP	98.63±0.03	55.27±0.32	27.58±0.09	98.86±0.04	64.99±0.32	34.20±0.20
FA	98.42±0.07	53.82±0.24	24.61±0.28	98.50±0.06	57.51±0.57	27.15±0.53
DRTP	95.10±0.10	45.89±0.16	18.32±0.18	97.32±0.25	50.53±0.81	20.14±0.68
PEPITA	98.01±0.09	52.57±0.36	24.91±0.22	98.29±0.13	56.33±1.35	27.56±0.60

Acknowledgements



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*Harvard Medical
School*