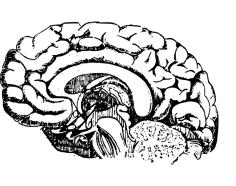
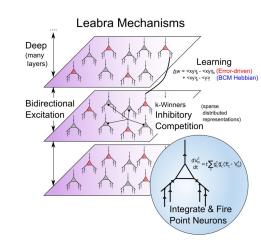
Biologically Based Models of Cortical Cognition



Modeling in Neuroscience

- Project Presentation -



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Contents

- Introduction
- 6 principles for biologically based models of cortical cognition
- Interactions between principles
- LEABRA algorithm
- Experiments
- Conclusion

Introduction

- Number of principles developed for neural-network models of cognition
- Machine learning ideas taken from cognitive neuroscience
- Article provides a set of six principles integrated into a coherent overall framework
 - → Presentation of the paper, the overall framework and some experiments

- 1) Biological Realism
 - Models should be constrained and informed by biological properties of the cortex
 - Computational mechanisms that violate biological properties should not be relied upon

- 2) Distributed Representations
 - Multiple active neuron-like processing units
 - Same unit used in multiple representations
- → Efficiency: combinations of small set of units
- → Robustness: redundancy of items represented by many units
- → Accuracy: representation of graded values
- → Similarity: shared units between similar items

- 3) Inhibitory Competition
 - 20% of cortex neurons are inhibitory interneurons → controls explosion of activations (epilepsy)
 - Allows for sparse distributed representations: general structure of the environment

- 4) Bidirectional Activation Propagation (Interactivity)
 - Both bottom-up and top-down flows
 - Iterative steps through a settling process to stabilize the flows of information

- 5) Error-driven Task Learning
 - Supervised learning: minimize error between desired outcome and network output → back-propagation algorithm
 - Not biologically plausible, except if using interactivity (GeneRec algorithm)
 - Propagates 2 signals and locally take their difference at each unit
 - Expectation phase: expected consequences
 - Outcome phase: experience actual consequences
 - → Teaching signal: difference between expectation and outcome

- 6) Hebbian Model Learning
 - Self-organizing or unsupervised learning
 - Internal representations of statistical structure of the environment
 - Correlational structure → Hebbian
 - Biologically plausible: NMDA-mediated LTP through changes of synaptic strength
 - Co-occurrence of items ⇒ causal relationships (PCA)

Interactions Among the Principles

1) Interactivity and Noise

In an identification scenario

Bidirectional activation propagation

VS

- Exhibit independent contributions from context
- → Because of correlations between neurons

Interactions Among the Principles

- 2) Distributed Representations and Competition
 - Distributed representations with multiple active neurons

 VS
 - Competition that inhibits cooperativity
- → Leads to sparse representation (combinatorial explosion)

KWTA: enforces competition while allowing for sparse distributed representation

1 Not treated within a probabilistic learning framework

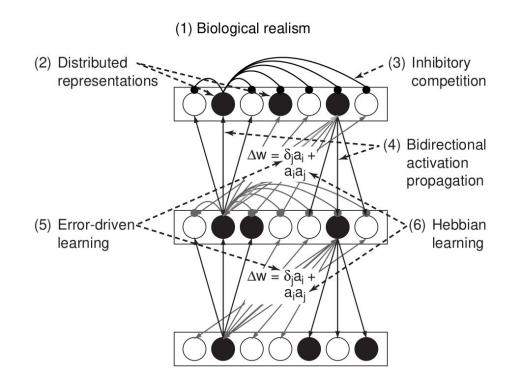
Interactions Among the Principles

- 3) Learning Principles
 - Computational objectives of learning
 - Task learning
 - Model learning
 - Implementational mechanisms
 - Error-driven learning
 - Hebbian learning
- → Quite similar

Only error-driven learning may map arbitrary inputs to outputs.

But may include a Hebbian bias towards representing co-occurrence statistics.

→ Learning in an Error-driven and Associative, Biologically Realistic Algorithm



Point-Neuron Activation Function

$$\frac{dV_m}{dt}(t) = \tau \sum_c g_c(t) \frac{dV_m}{g_c} (E_c - V_m(t))$$

Channels: excitatory input, leak current, inhibitory input

Communication to other cells:
$$y_j(t) = \frac{1}{1 + \frac{1}{\gamma[V_m(t) - \Theta]_+}}$$

K-Winners-Take-All Inhibition (KWTA)

Inhibitory current:
$$g_i = g_{k+1}^{\Theta} + q(g_k^{\Theta} - g_{k+1}^{\Theta}) \quad \text{with} \begin{cases} q & \approx 0.25 \\ g^{\Theta} & = \frac{\sum_{c \neq i} g_c \overline{g_c}(E_c - \Theta)}{\Theta - E_i} \end{cases}$$

Achieves sparse distributed representation.

Using uniform level of inhibition ⇒ no more than k units active at a time

- Error-driven Learning
- → Symmetric version of the biologically plausible GeneRec algorithm

$$\Delta w_{ij}^{E} = x_{i}^{+} y_{j}^{+} - x_{i}^{-} y_{j}^{-}$$

2 phases:

- Expectation phase (-)
- Outcome phase (+)

Hebbian Learning

Variant of Oja normalization:
$$\Delta w_{ij}^H = x_i y_j - y_j w_{ij} = y_j (x_i - w_{ij})$$

 \rightarrow Compute expected value of sender conditional on receiver: $w_{ij}^H \approx \langle x_i | y_j \rangle$

Total update:
$$\Delta w_{ij} \propto \Delta w_{ij}^E + \lambda \Delta w_{ij}^H$$

Experiments

- Two neurons
- Pattern association
- Iris dataset classification

Two Neurons

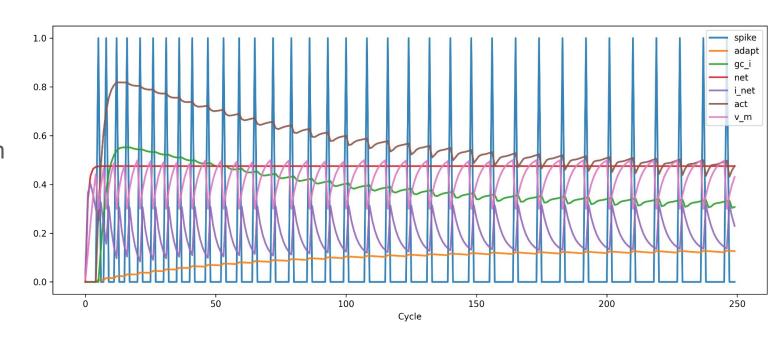
• Weight: 0.5

Input activation: 1

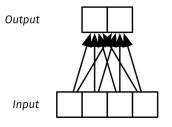
• 1 cycle: 1ms

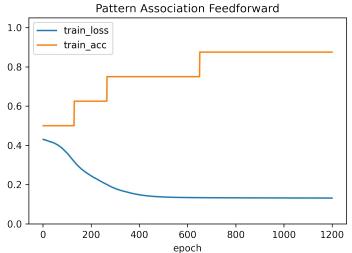
V_m > thresh

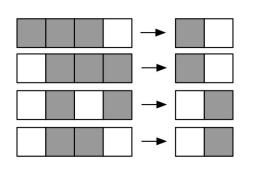
⇒ spike

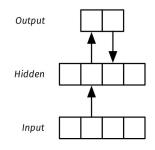


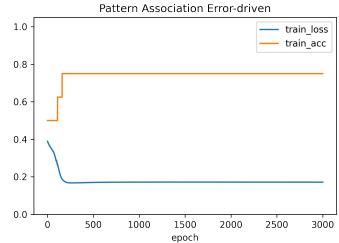
Pattern Association





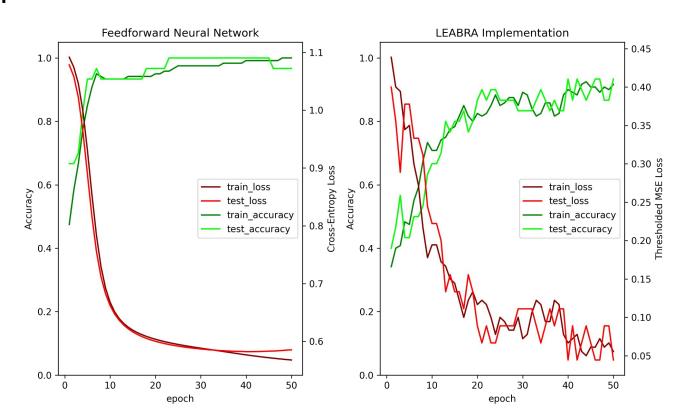






Iris Classification

- 40 input units
- 23 hidden units
- 3 output units



References

- [1] Randall C. O'Reilly. "Six principles for biologically based computational models of cortical cognition". In: Trends in Cognitive Sciences 2.11 (1998).
- [2] R. C. O'Reilly. "Biologically Plausible Error-Driven Learning Using Local Activation Differences: The Generalized Recirculation Algorithm". In:Neural Computation 8.5 (1996).
- [3] Erkki Oja. "Simplified neuron model as a principal component analyzer". In: Journal of Mathematical Biology 15.3 (1982).
- [4] C. Daniel Greenidge, Noam Miller, and Kenneth A. Norman. "Leabra7: a Python package for modeling recurrent, biologically-realistic neural networks". In: CoRRabs/1809.04166 (2018).

Conclusion

- Presented 6 principles
- Can be integrated into a coherent framework
- LEABRA does learn
- Training is unstable
- Analysis is hard (no probabilistic framework / sparsity / complex equations)
- Great potential for applications in ML and further research in cognitive neuroscience