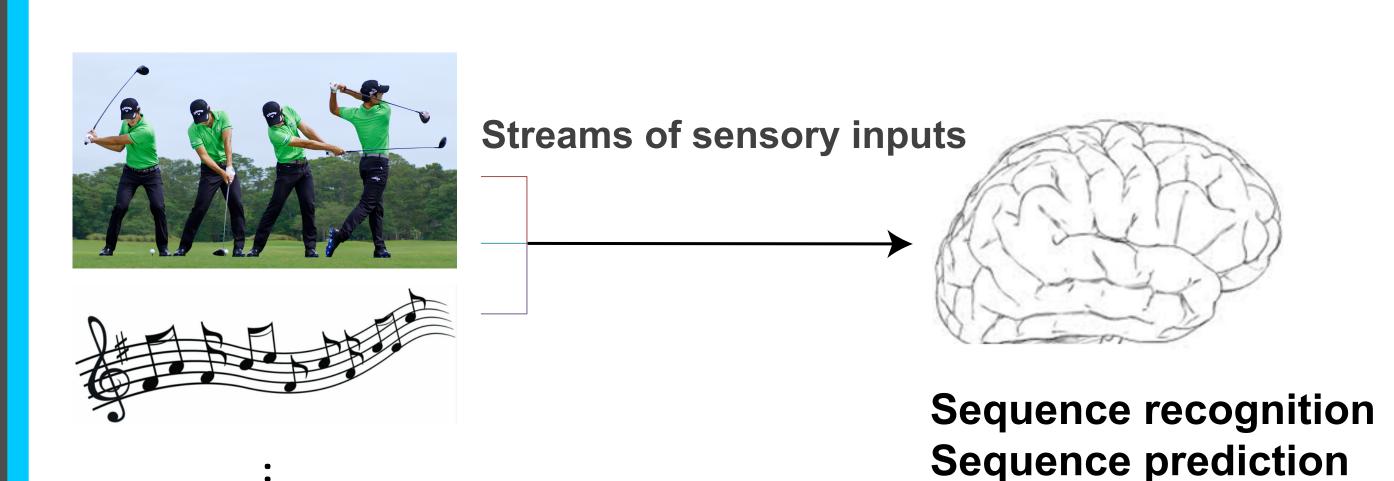


# A theory of sequence memory in the neocortex

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### 1. Sequence learning is ubiquitous in cortex



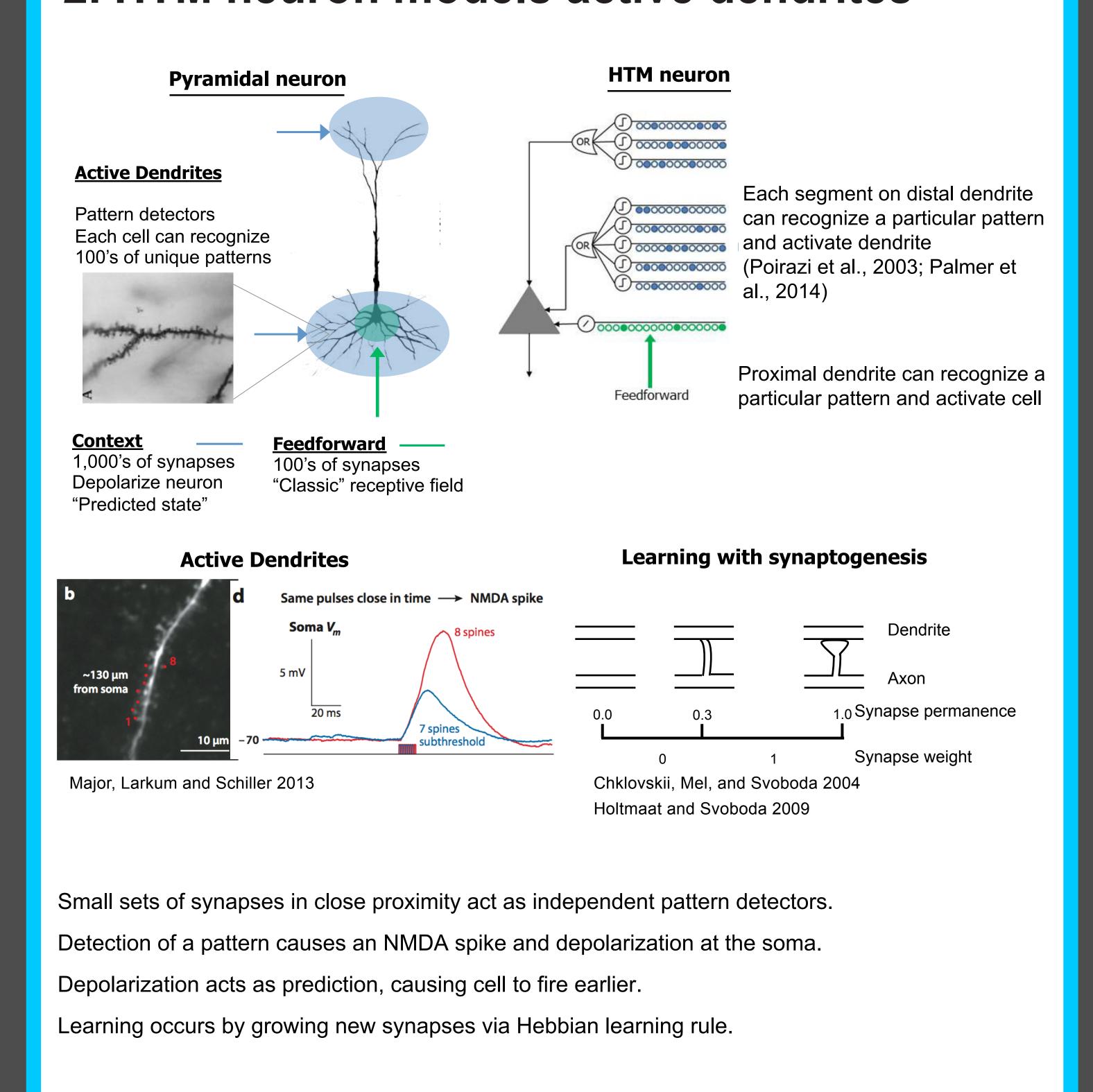
### What is neural mechanism for sequence learning?

Behavior generation

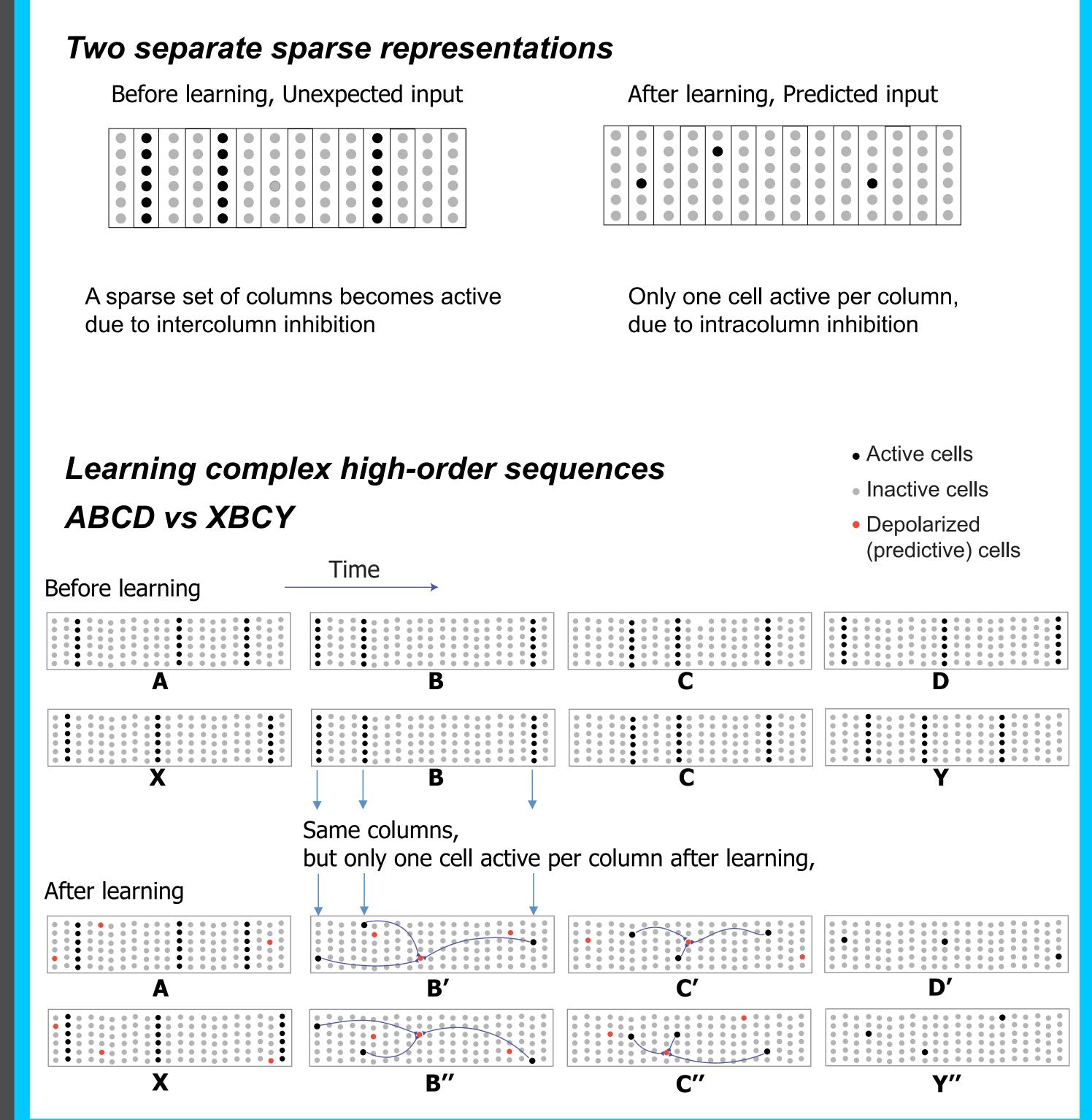
#### HTM Sequence Memory:

- 1. Neurons learn to recognize hundreds of patterns using active dendrites.
- 2. Recognition of patterns act as predictions by depolarizing the cell without generating an immediate action potential.
- 3. A network of neurons with active dendrites forms a powerful sequence memory

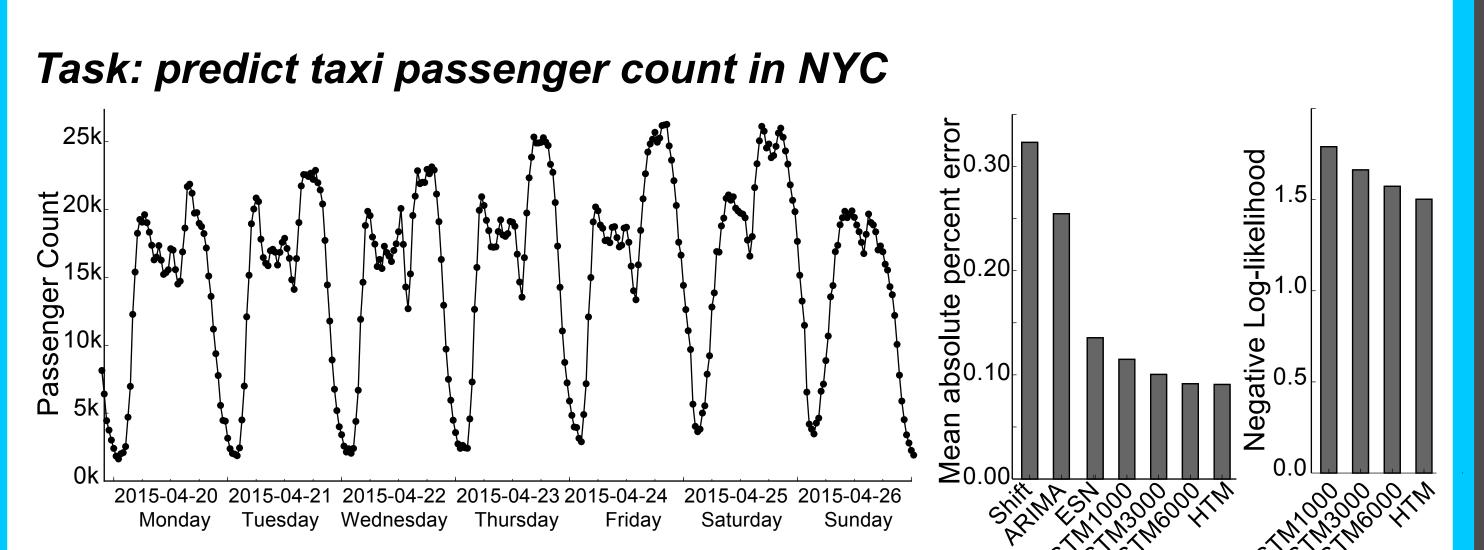
### 2. HTM neuron models active dendrites



### 3. HTM network model for sequence learning

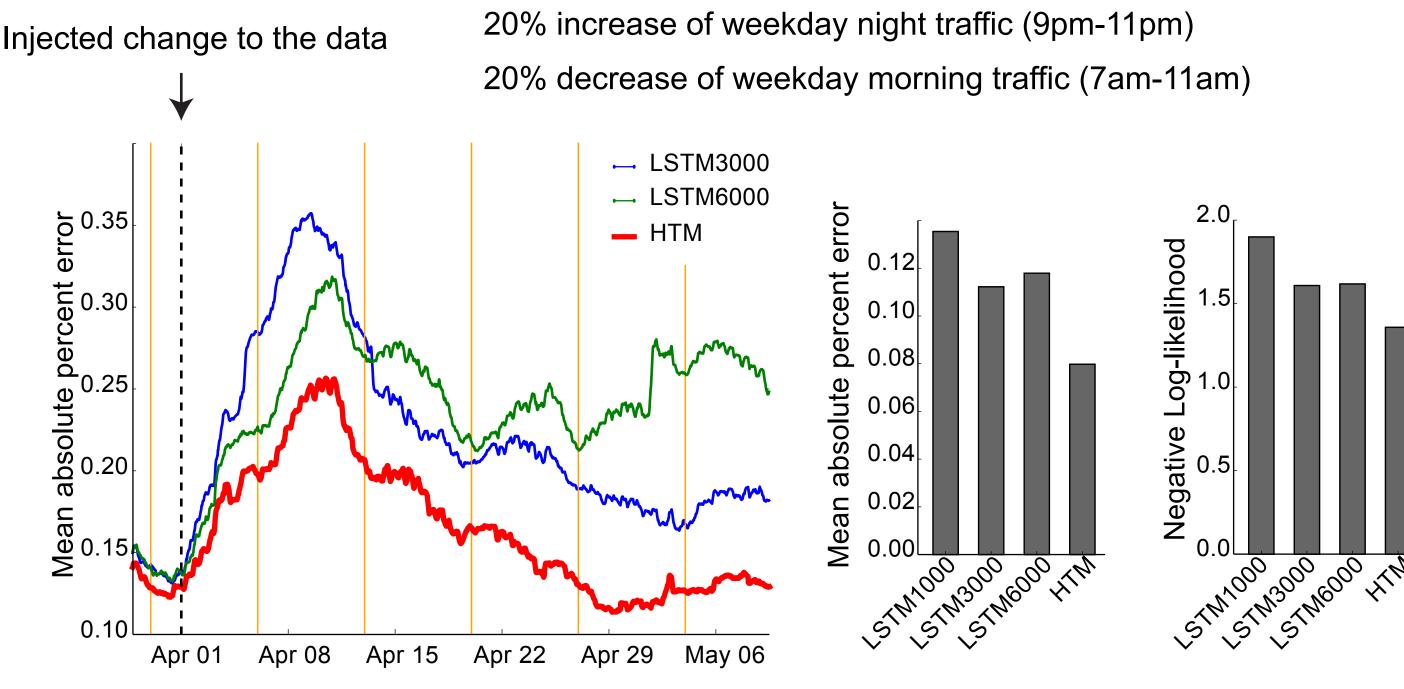


### 4. Works well on real-world problems



HTM has comparable performance to state-of-the-art algorithms

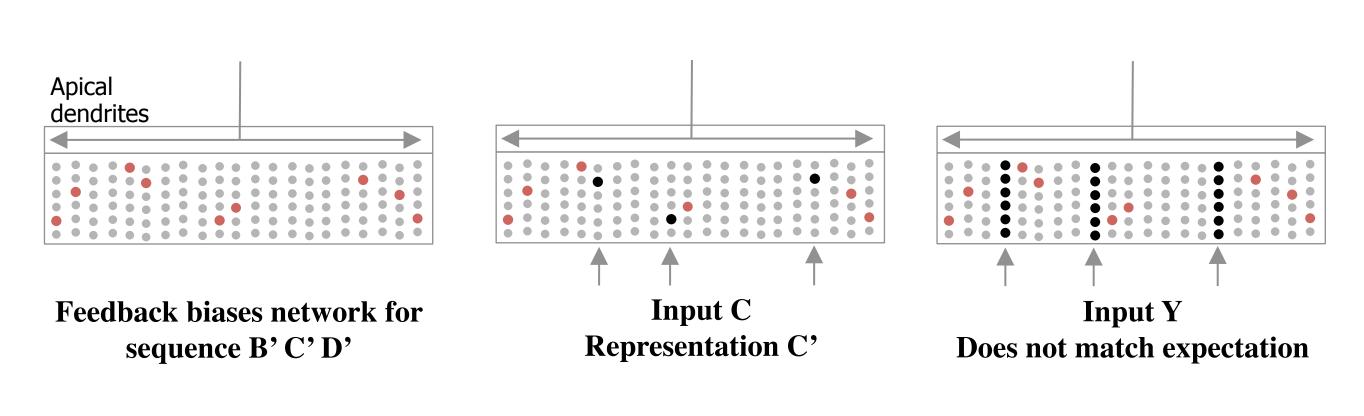
Data sources: http://www.nyc.gov/html/tlc/html/about/trip\_record\_data.shtml



HTM quickly adapts to changes due to its ability to learn continuously

### Apical inputs predict entire sequences

It has been speculated that feedback connections implement expectation or bias (Lamme et al., 1998). Our neuron model suggests a mechanism for top-down expectation in the cortex.



Feedback to the apical dendrites can predict multiple elements simultaneously. New feedforward input will be intepreted as part of the predicted sequence.

### Learning and activation rules

#### **Activation rules:**

Select the top 2% of columns with strongest inputs on proximal dendrite as active columns

If any cell in an active column is predicted, only the predicted cells fire

If no cell in an active column is predicted, all cells in the column fire

#### Unsupervised Hebbian-like learning rules:

If a depolarized cell becomes active subsequently, its active dendritic segment will be reinforced

If a depolarized cell does not become active, we apply a small decay to active segments of that cell

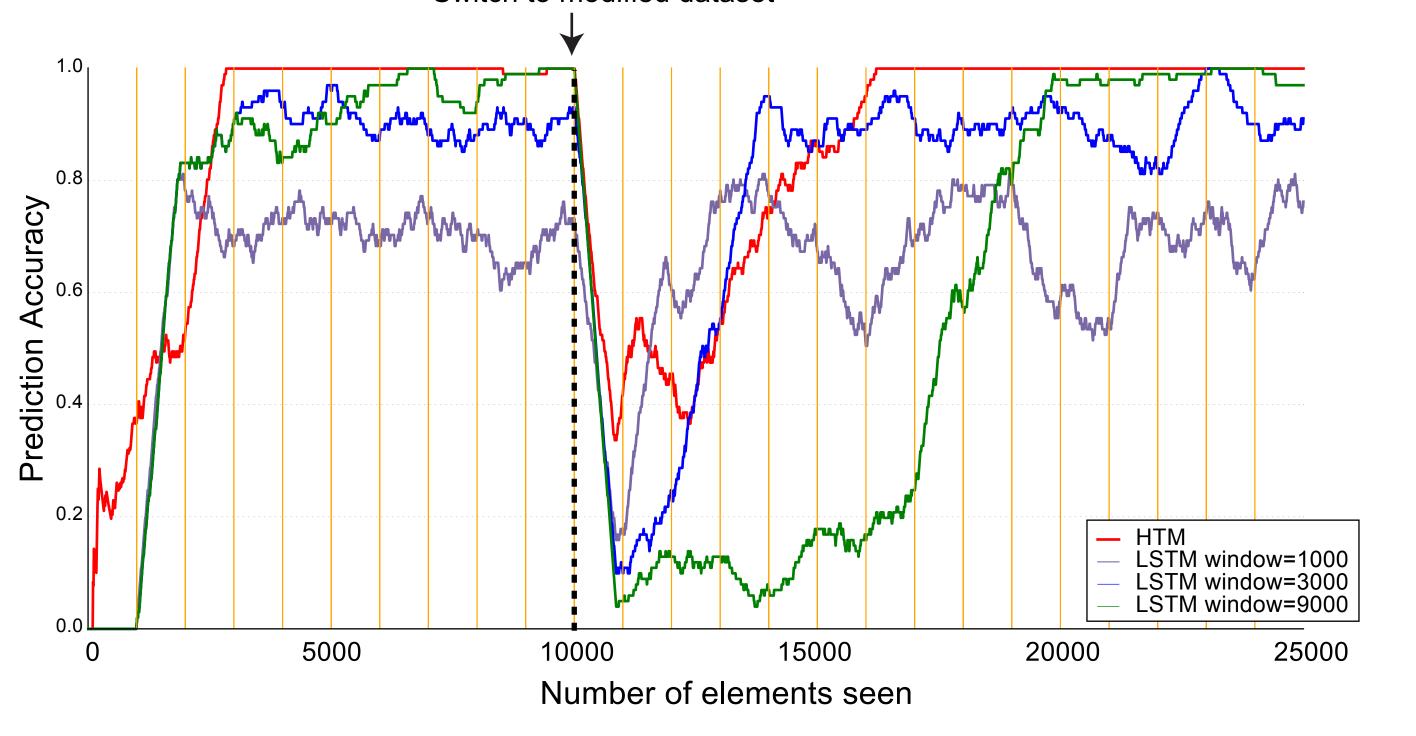
If no cell in an active column is predicted, the cell with the most activated segment gets reinforced

# Continuous learning from sequence streams

Task: sequence prediction with streams of high-order sequences

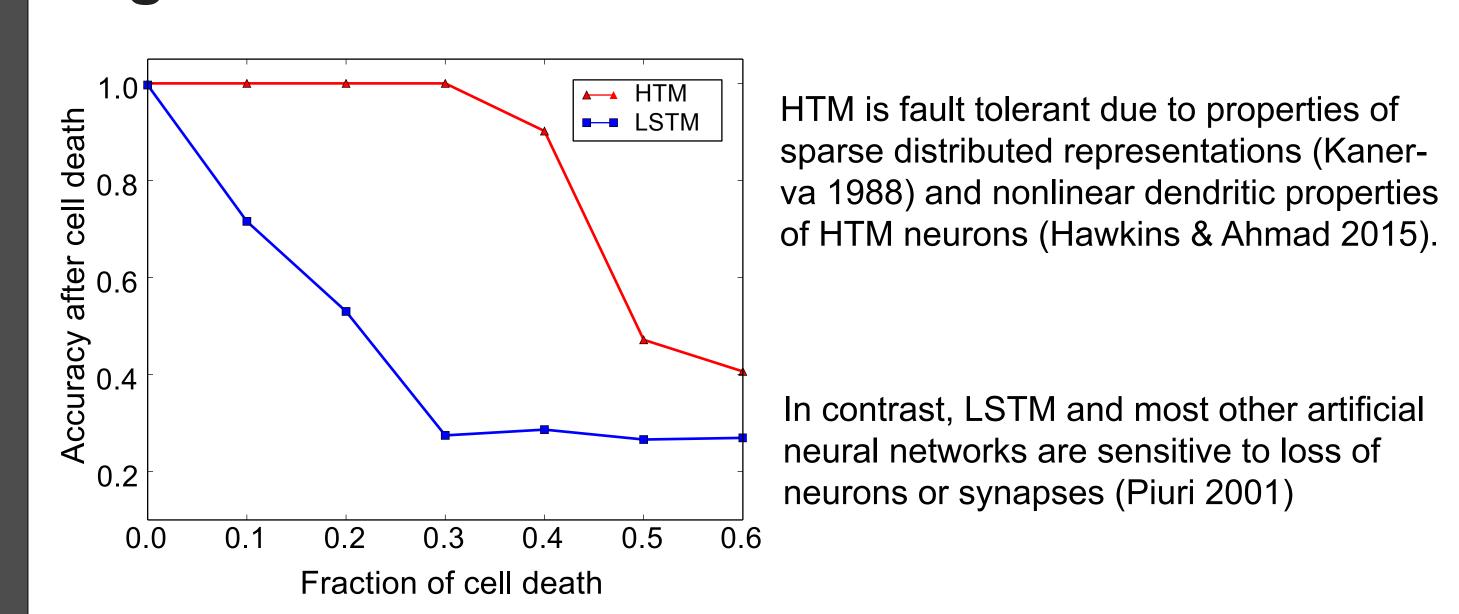
Check prediction at the end of each sequence

Example sequences:
G, I, H, E, C, D, A
B, I, H, E, C, D, F
A, J, H, I, F, D, E, B
C, J, H, I, F, D, E, G
C, J, H, I, F, D, E, G
Switch to modified dataset



HTM learns continuously, no batch training required. HTM is more robust and recovers more quickly.

## High fault tolerance to neuron death



# References

Poirazi P, Brannon T, Mel BW (2003) Neuron 37:989–999.
Chklovskii, D. B., Mel, B. W., and Svoboda, K. (2004). Nature 431, 782–8.
Hawkins, J., and Ahmad, S. (2015). arXiv:1511.00083 [q-bio.NC]
Holtmaat A, Svoboda K (2009) Nat Rev Neurosci 10:647–658.
Kanerva, P, Sparse Distributed Memory. The MIT Press, 1988
Lamme, V. A., Supèr, H., and Spekreijse, H. (1998). Curr. Opin. Neurobiol. 8, 529–35.
Losonczy, A., Makara, J. K., and Magee, J. C. (2008). Nature 452, 436–41.
Major, G., Larkum, M. E., and Schiller, J. (2013). Annu. Rev. Neurosci. 36, 1–24.
Piuri, V., J Parallel Distr Com., vol. 61, pp. 18-48.
Smith SL, Smith IT, Branco T, Häusser M (2013) Nature 503:115–120.
Vinje, W. E., and Gallant, J. L. (2002). J. Neurosci. 22, 2904–2915.

### Acknowledgements

We thank Scott Purdy, Alex Lavin, Matthew Larkum, Ahmed El Hady for helpful discussions.

### Our code is open source

We believe in open research and full transparency. Numenta's research and algorithm code is part of the open-source project Numenta Platform for Intelligent Computing (NuPIC). As a fast growing project, NuPIC currently has more than 4,000 followers and more than 1000 forks on Github.

### 5. Summary

HTM exhibits many desirable features for sequence learning:

- Unsupervised learning
- Quickly adapts to changes in data
- Learns high-order structure in sequences
- Robust and fault tolerant
- Makes multiple simultaneous predictions
- Works well on real-world problems
- Accurate biological model

### 6. Testable predictions

- 1) Sparser activations during a predictable sensory stream. (Vinje & Gallant 2002)
- 2) Unanticipated inputs lead to a burst of activity correlated vertically within mini-columns.
- 3) Neighboring mini-columns will not be correlated.
- 4) Predicted cells need fast inhibition to inhibit nearby cells within mini-column.
- 5) For predictable stimuli, dendritic NMDA spikes will be much more frequent than somatic action potentials. (Smith et al., 2013)
- 6) Strong LTP in distal dendrites requires bAP and NMDA spike (Losonczy et al., 2008)
- 7) Weak LTP (in the absence of NMDA spikes) in dendritic segments if a cluster of synapses become active followed by a bAP.
- 8) Localized weak LTD when an NMDA spike is not followed by a bAP.

Further implementation detail can be found at https://github.com/numenta/nupic