The LISSOM Cortical Model

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Problems with SOM retinotopy

The particular model of SOM retinotopy we've been looking at also has other problems:

- There is no known state when the connections from the eye are evenly distributed across a target region; even the initial connections are retinotopic.
- The overall retinotopy is established by axons following gradients of signaling molecules such as Ephrins, though activity may have some role in this process (reviewed in Flanagan 2006; Huberman et al. 2008).

In any case, activity appears to be required for map refinement, and it's interesting that in principle an unfolding process like in the SOM simulation could work.

Problems with SOMs

A Kohonen SOM is very limited as a model of cortical function:

- Picking one winner is valid only for a very small patch with very strong lateral inhibition.
- Full connectivity is possible only for very small cortical networks.
- Lateral interactions are forced to be isotropic, contrary to biological evidence.
- Euclidean distance metric is not clearly relatable to neural firing or synaptic plasticity.

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LISSOM

The LISSOM model (Sirosh & Miikkulainen 1994) was designed to remove some of the artificial limitations and biologically unrealistic features of a SOM:

- Recurrent lateral interactions, instead of global winner
- Specific lateral connections, instead of isotropic neighborhood
- Spatially localized RFs, instead of full connectivity
- Activation by sigmoided dot product, rather than Euclidean distance
- Learning by Hebbian rule

GCAL

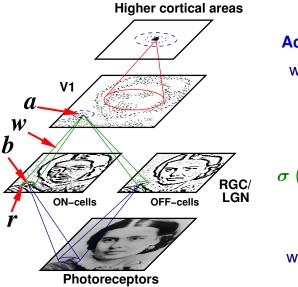
In turn, GCAL (Bednar 2012; Law et al. 2011) was designed to remove some of the artificial limitations and biologically unrealistic features of LISSOM:

- Automatic homeostatic plasticity, instead of hand-adjusted thresholds
- No lateral connection radius shrinking
- Gain control for realistic behavior with contrast (similar to the afferent normalization of CMVC section 8.2.3)

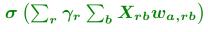
GCAL is otherwise like LISSOM. The CMVC book and older work all focus on LISSOM, but current work uses GCAL; in most cases the distinction is not important.

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HLISSOM Architecture

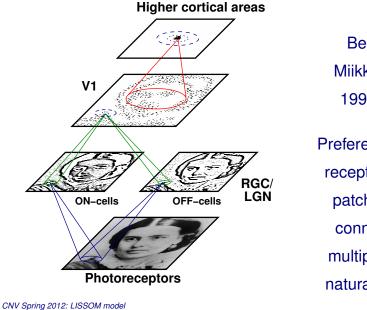


Activity: thresholded weighted sum of all receptive fields $\eta_a =$



Response high
when input matches
weights

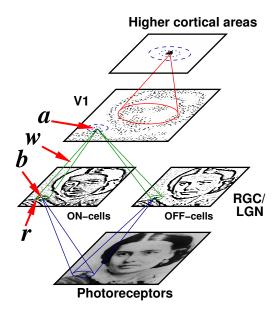
HLISSOM Architecture

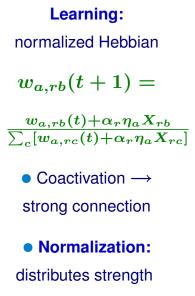


Bednar & Miikkulainen, 1995–2004

Preference maps, receptive fields, patchy lateral connections, multiple areas, natural images

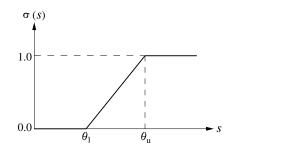
HLISSOM Architecture





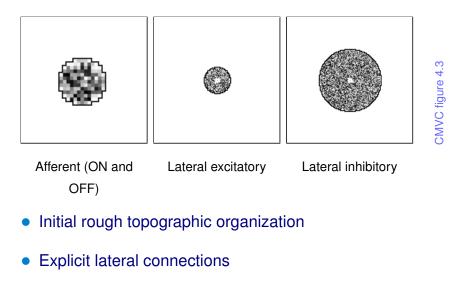
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Neuron activation function $\sigma(s)$

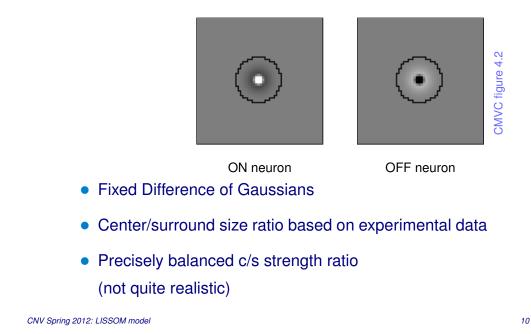


- Piecewise-linear approximation to a sigmoid
- Easy to compute
- Speeds up computation, since most neurons are truly off
- Strongly sensitive to threshold θ_l

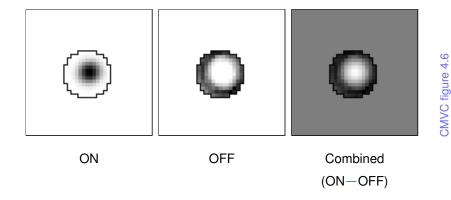
Initial V1 weights



DoG RGC/LGN RFs

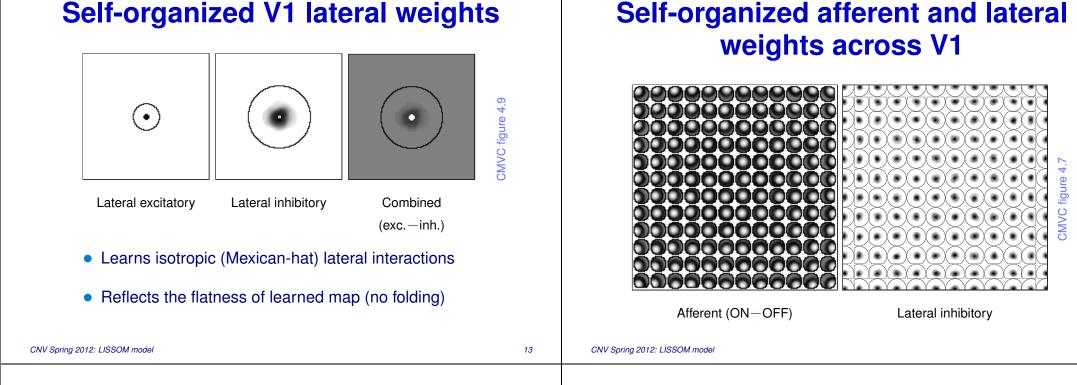


Self-organized V1 afferent weights

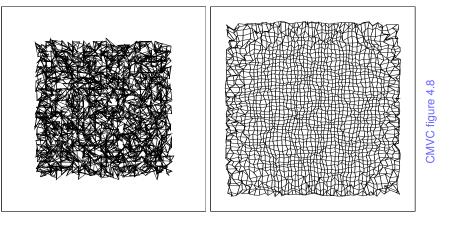


Given isotropic Gaussians, learns isotropic Gaussians

CMVC figure 4.5



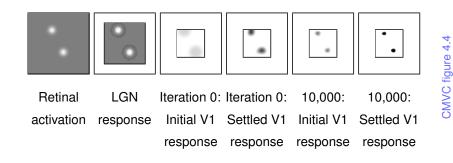
Self-organization of the retinotopic map



Initial disordered map

Final retinotopic map

Retinotopy input and response



- Settling process: Sharpens activity around strongly activated patches
- Multiple winners occur for multiple features on input

Summary

LISSOM: same basic process as a SOM, but:

- More plausible
- More powerful:
 - Multiple winners

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- Specific lateral connections
- More sensitive to parameters
- · More computation and memory intensive

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17

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