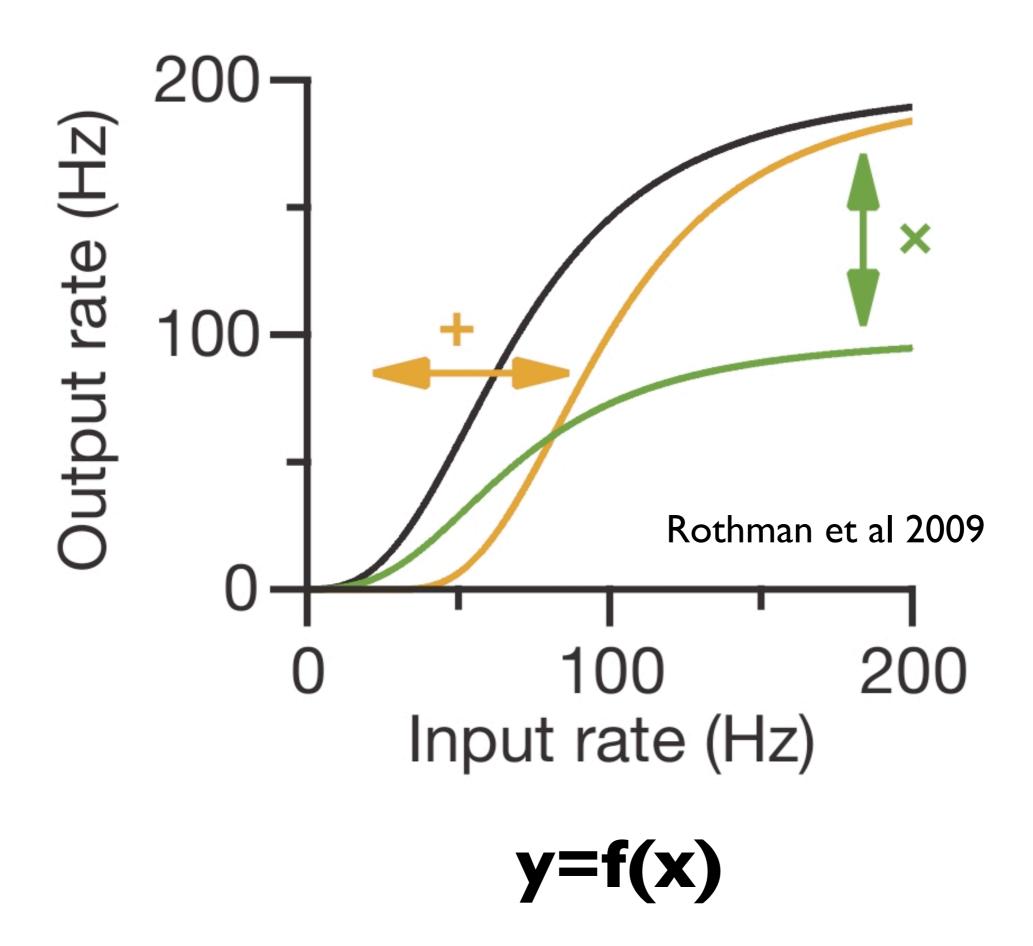
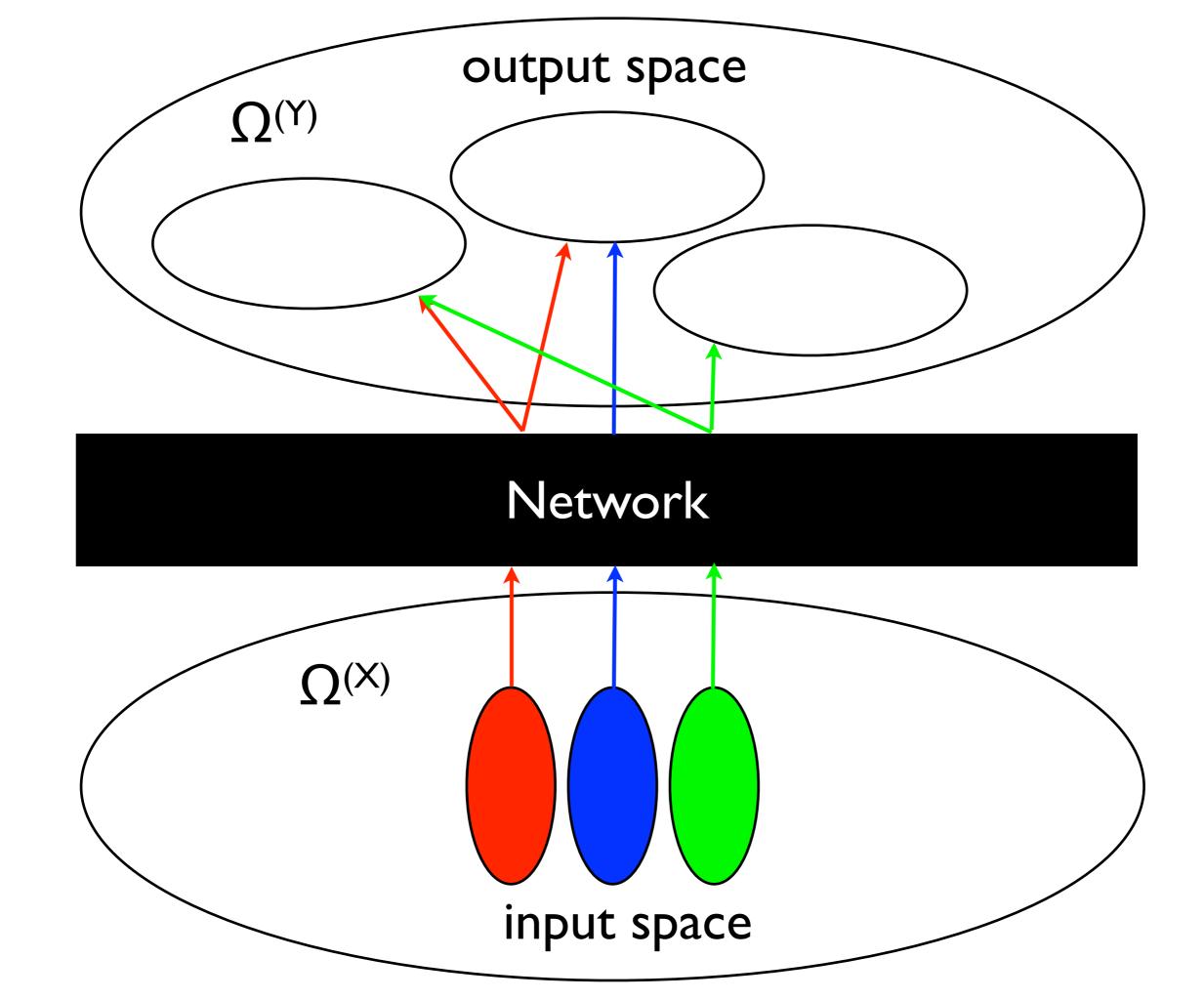
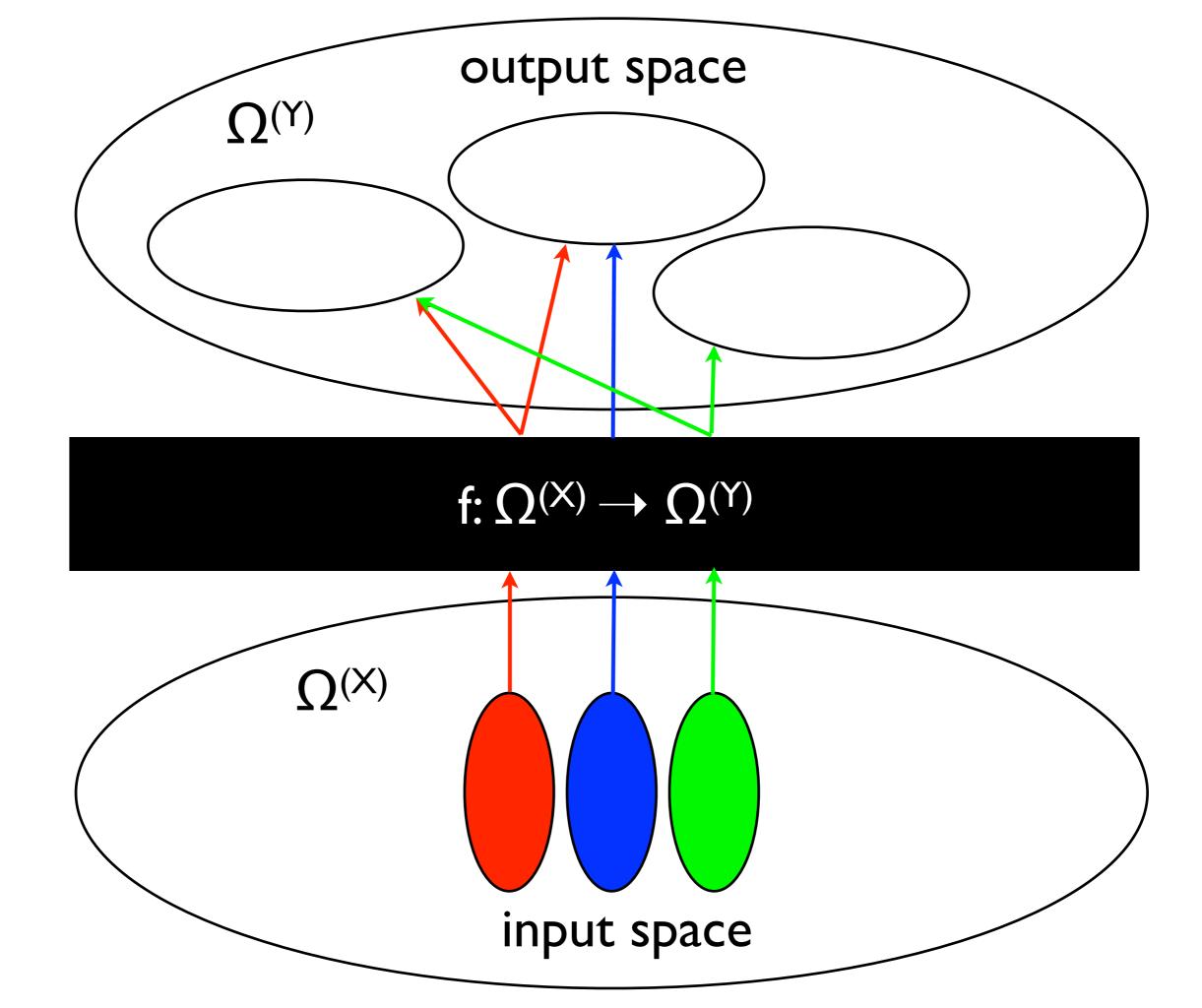
Network Transformations on Code Space

Guy Billings
Data Club 02/08/2010





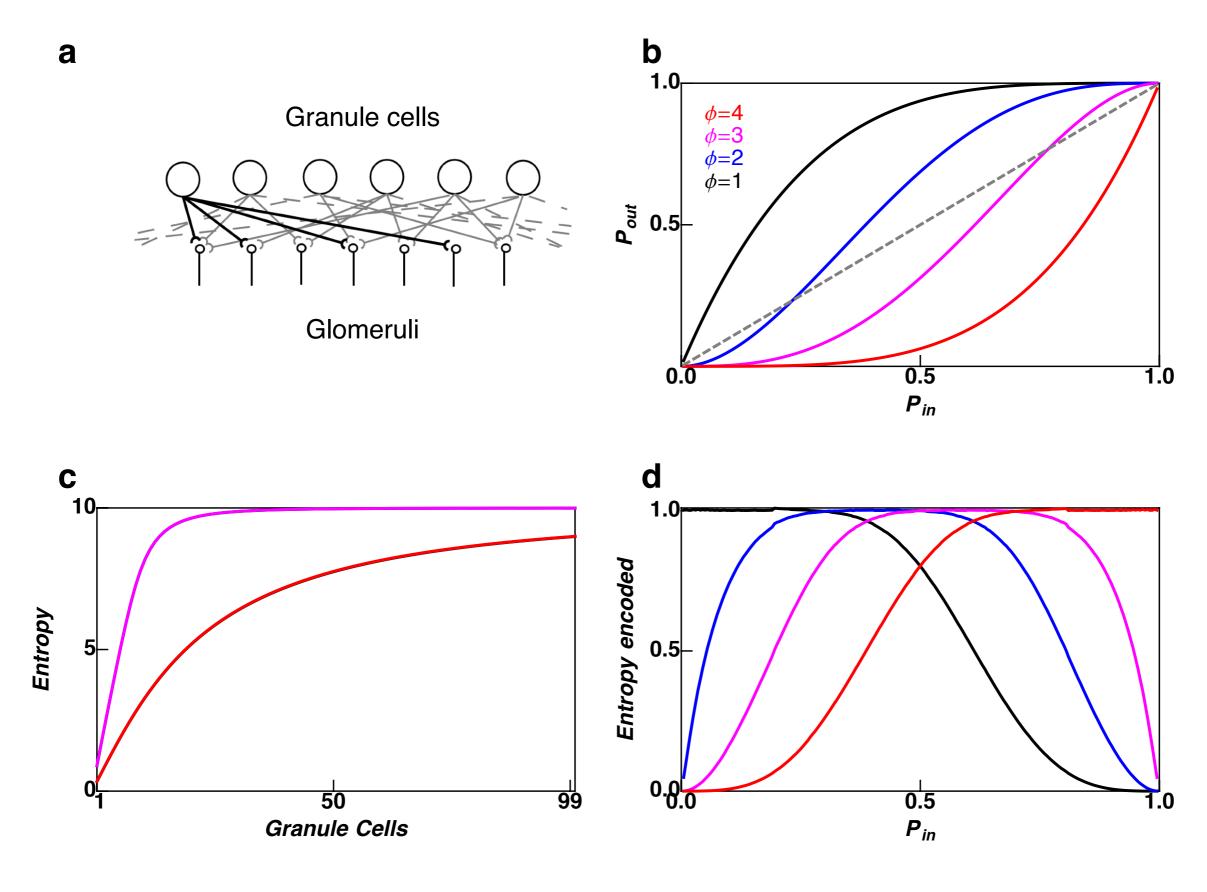


Introduction

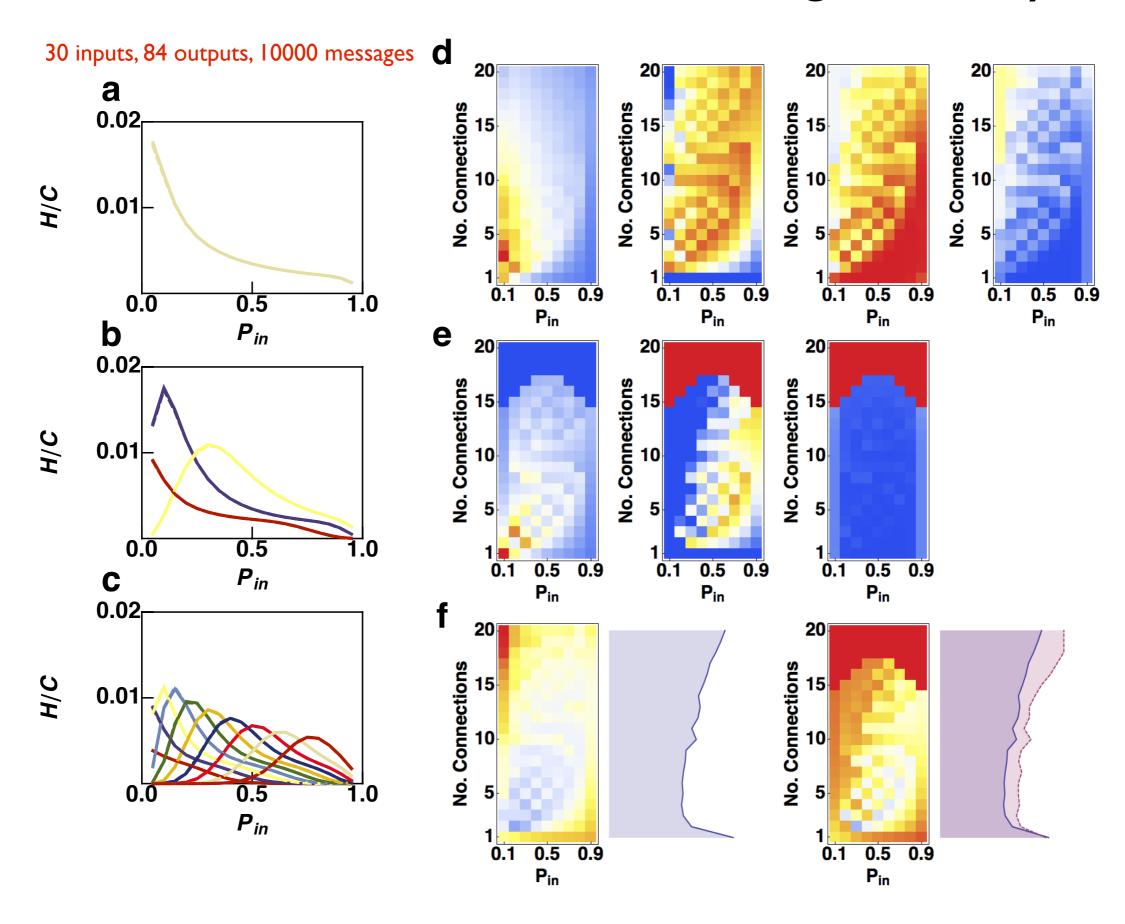
Case study: Binary model of the granule cell layer

Spike train space

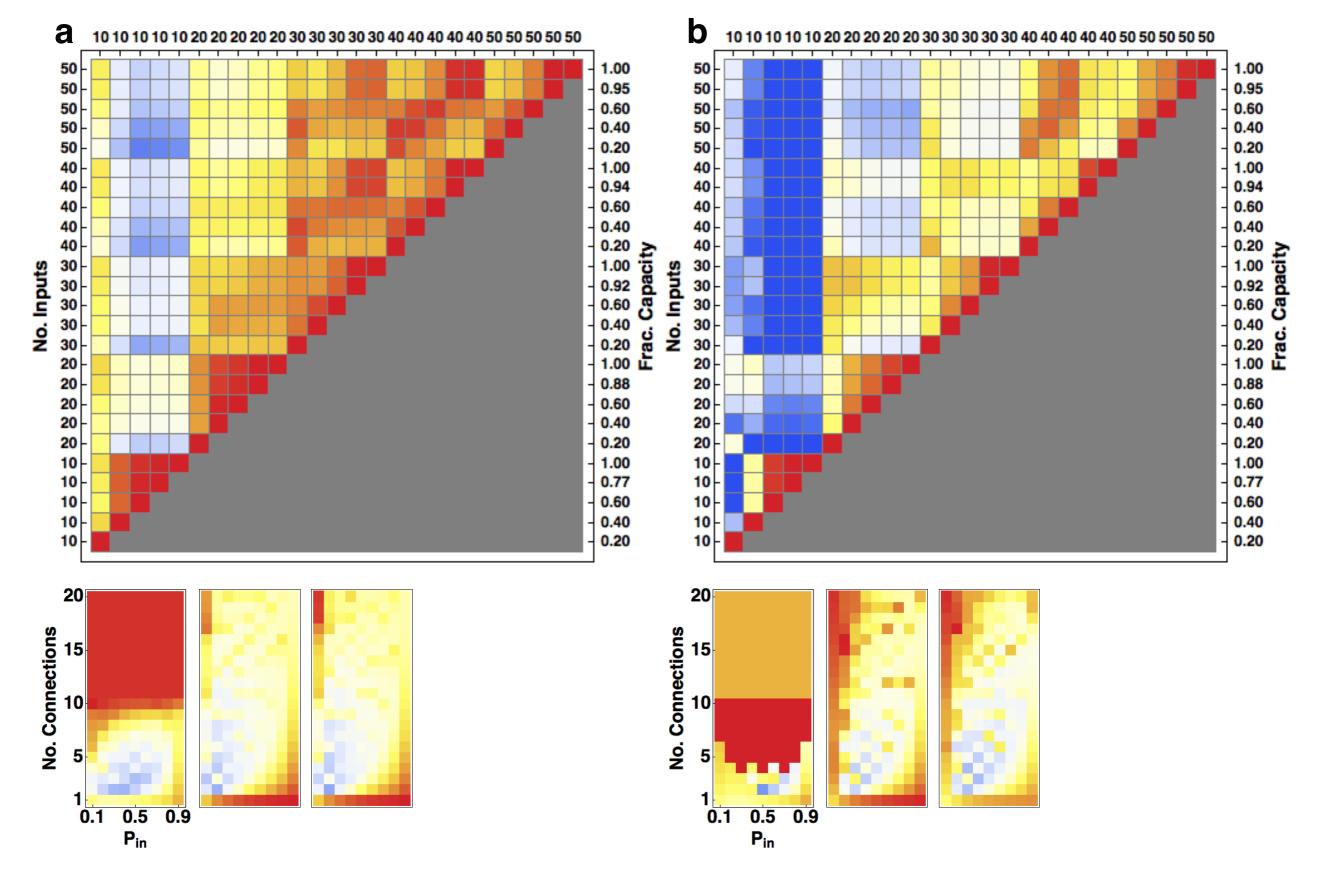
Simple model of the cerebellar granule cell layer



3 connections maximises coding efficiency



Scaling of encoding



Granule cell layer encoding

3 connections optimises energy efficiency of encoding

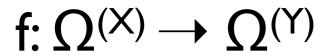
Overall, having a small number of connections (<10) minimises the performance distance between the network and the perfect network

Primary determinant of the nature of encoding (lossy/lossless, sparsity) is thresholding (tonic inhibition)

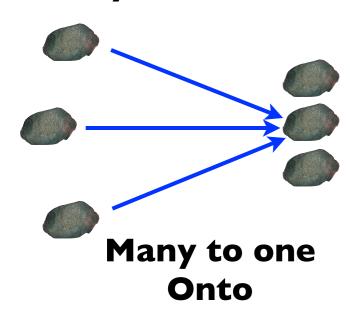
Further questions:

What is the optimal distribution? Inclusion of metric pattern separation Effect of correlations

NTCS perspective



Lossy



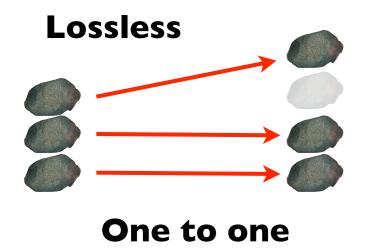
Output space: Discrete topology

Define: pre-images of open sets as open

Input space is not separable

Metric space not possible (pseudo-metric space)

Efficient (compression?)



Not onto

Output space: Discrete topology

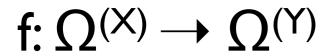
Define: pre-images of open sets as open

Input space is separable (has discrete topology)

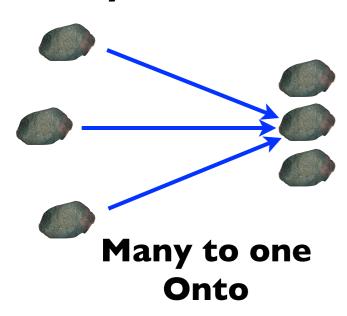
Can be metrisized

Costly (separation?)

NTCS perspective



Lossy



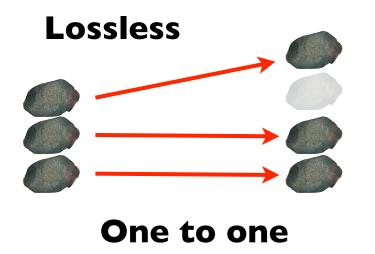
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Output space: Discrete topology

Define: pre-images of open sets as open

Input space is separable (has discrete topology)

Can be metrisized

Costly (separation?)

In analogy to the transfer function: Statements about the transformation are restricted Restrictions directly related to our **choice of code space** (domain)

The obvious choice of code space....

SPIKETRAIN SPACE!

SPIKETRAIN SPACE!

But this introduces extreme case of 'curse of dimensionality'

SPIKETRAIN SPACE!

But this introduces extreme case of 'curse of dimensionality'

Need a way of controlling the complexity of the problem

Ways forward

Let $\Omega^{(X)}$ be 'spike train space'. Embed $\Omega^{(X)}$ in an 'ambient' space (\mathbb{R}^n).

I) Define a metric Embedded spike train space Clustering/dimensionality-reduction/other analysis based on chosen metric (Victor Purpura 1996, van Rossum 2001, Houghton 2010)

2) 'Connect the dots'
Simplical complex
Computational topology to find Homology group of the complex (i.e. its 'shape')
(Carlsson 2008, Singh et al 2008)

Ways forward

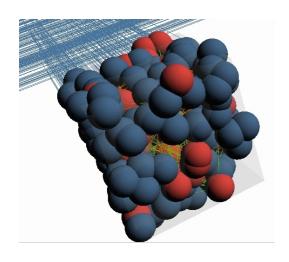
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 (Victor Purpura 1996, Houghton 2010)
- 2) 'Connect the dots'
 Simplical complex
 Computational topology to find Homology group of the complex (i.e. its 'shape')
 (Carlsson 2008, Singh et al 2008)

30 glomeruli, 84 granule cells

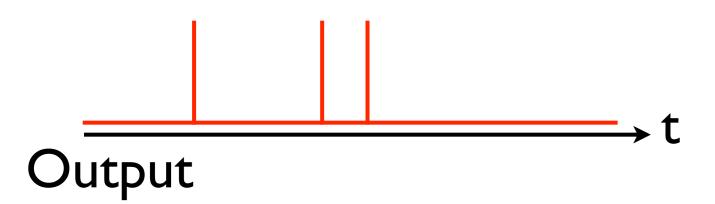
Input patterns: Inputs randomly activated (fire at 30Hz) with prob. 0. I

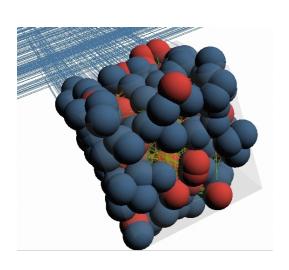
10 patterns, each with 10 repetitions0Hz otherwise

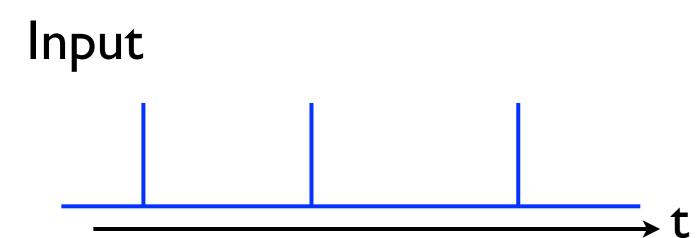


Output patterns: Resulting spike trains from granule cells Weights adjusted to match experiment

Embed $\Omega^{(x)}$ in an 'ambient' space (\mathbb{R}^n)



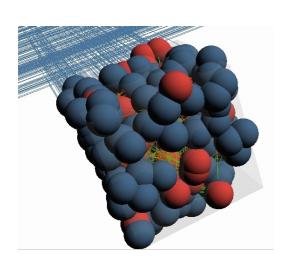




Embed $\Omega^{(x)}$ in an 'ambient' space (\mathbb{R}^n)

(vR metric) 5ms t

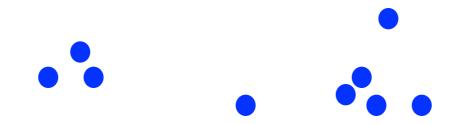
Output



Input

Embed $\Omega^{(x)}$ in an 'ambient' space (\mathbb{R}^n) (vR metric) <u>5ms</u>← Output 8.0 Time (ms) 0.6 50 0.4 0.2 Input 100 3 4 5 6 7 Cell index 8

Embed $\Omega^{(x)}$ in an 'ambient' space (\mathbb{R}^n) (vR metric) 5ms Output Each 'point' in the space represents activity of all cells within a particular time segment 8.0 Time (ms) 0.6 50 0.4 0.2 Input 100 2 4 5 6 7 Cell index 3 8

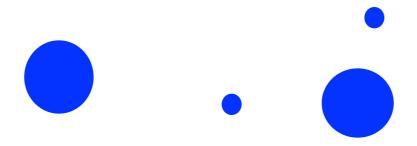






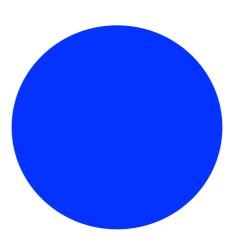




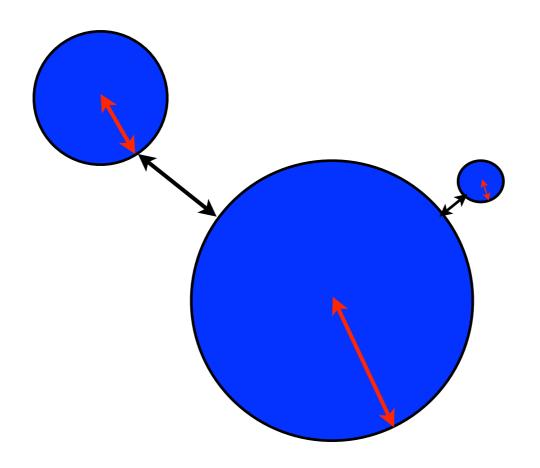




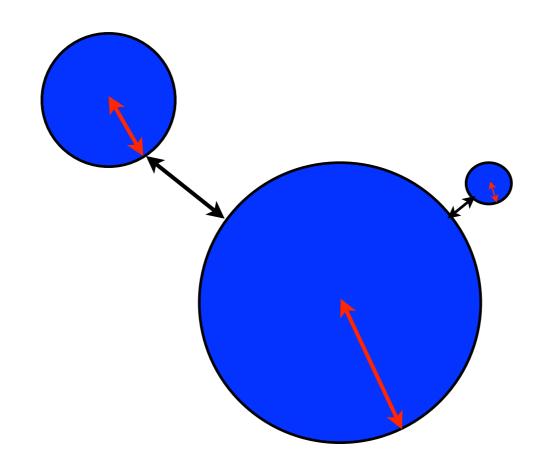




Clusters data into groups by merging successive closest points

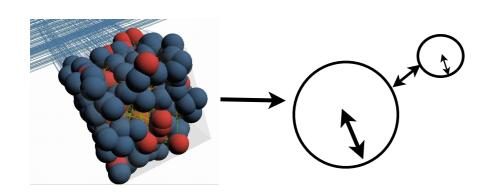


Clusters data into groups by merging successive closest points



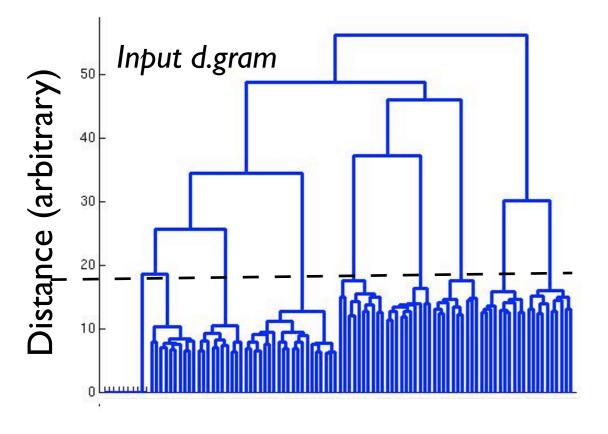
Use 'ward's linkage' which clusters on the basis of squared distance

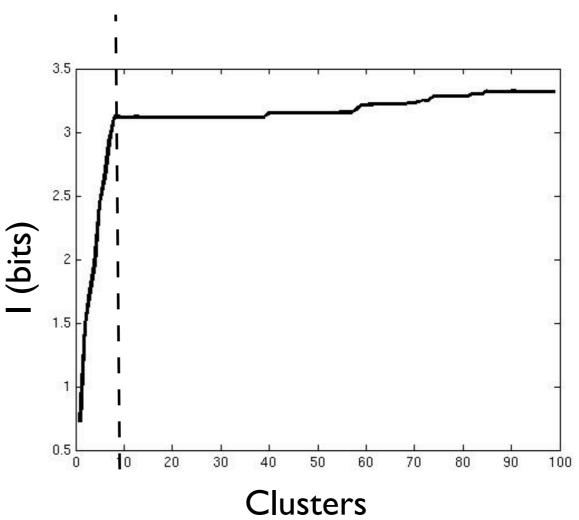
Hierarchical clustering gives variable resolution information measure



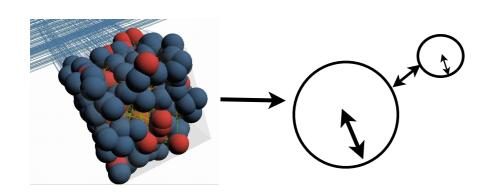
$$I(d,c) = \sum_{p \in P} \sum_{c \in C} p(d,c) \log\left[\frac{p(d,c)}{p(d)p(c)}\right]$$

Source entropy=3.32 bits



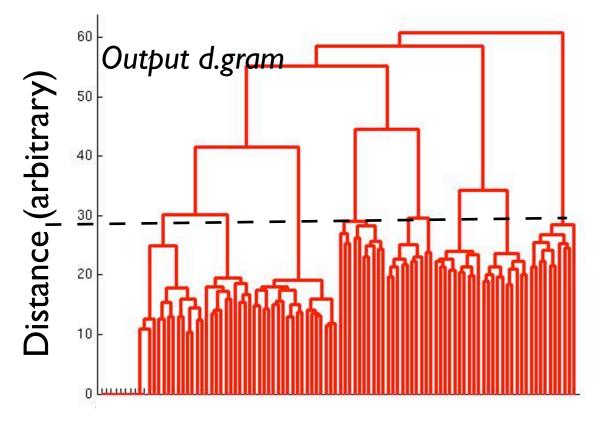


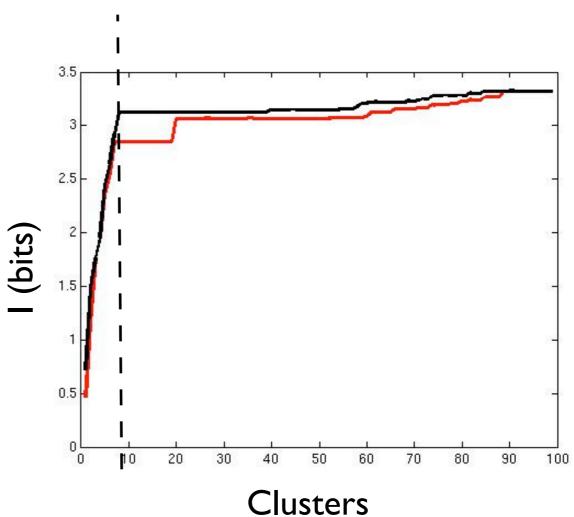
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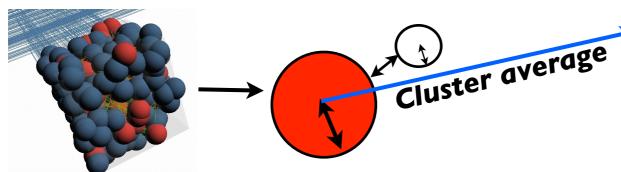
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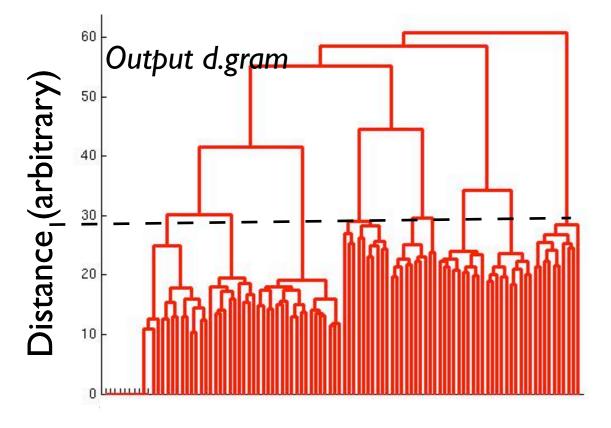


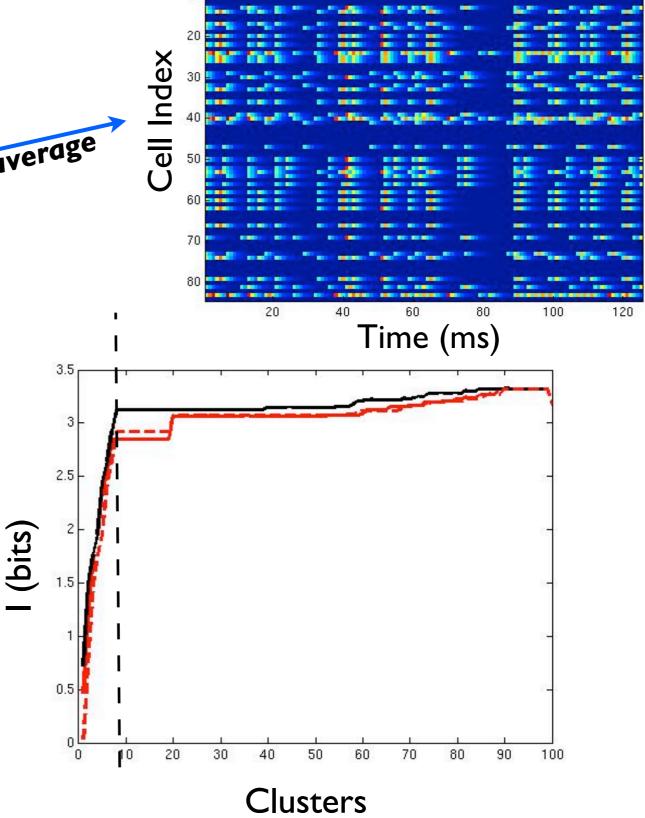


Building a 'decoder'



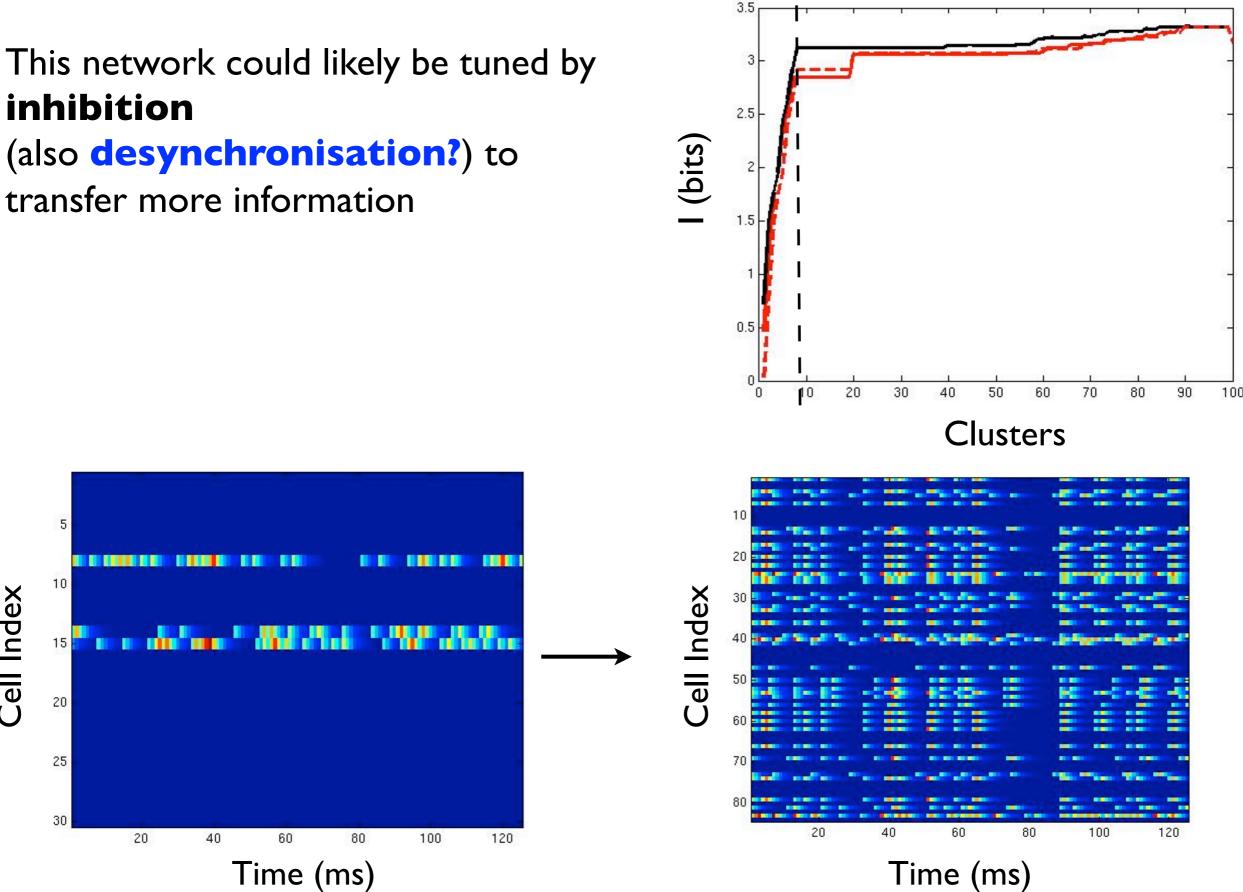
Source entropy=3.32 bits





inhibition (also desynchronisation?) to

Cell Index



Summary

Application of metric techniques to spike train space allows information theoretic analysis

Clustering allows identification of putative transformations

This can be used to repeat the previous granule cell study but with a more realistic model

Further work

Not clear how to choose the space/clustering method - ideally build a 'neurometric'

Application of topology - removing the need to define a metric