

Chapter 3

Computational Information– Maximization Models

Some information-maximization models, especially examples of the Independent Component Analysis (ICA), and their importance for the holonomic theory will be presented. Their biological plausibility will be discussed.

3.1 MAXIMAL PRESERVATION OF INFORMATION (“INFOMAX”)

Linsker’s infomax-net. The early “infomax” perception-model by Linsker (1988) was established with an observation that Hebbian adjustment of connections is able to evolve feature-selective neurons which collectively preserve information “as much as possible” during input-to-output processing. Linsker (1988) used a multi-layer network with local feed-forward connections which are Hebb-like – determined by

DOI: 10.4018/978-1-61520-785-5.ch003

covariance. In this case, local means that each neuron in layer $l+1$ receives inputs from neurons in a confined circular region of layer l . The “infomax” ideas used in the classical PCA-like¹ self-organizing net by Linsker (1988, and later works) have later been much developed in direction of ICA.

From second- to higher-order statistics. As has been mentioned in section 1.5, ICA is supposed to be important for cortical image processing because of taking into account the statistics of input data which is also of higher order than the second order.² *Correlation* and *convolution*, for example, are “learning (i.e., memory storage) rules” of *second order*. Such mathematical expressions, used in unsupervised learning models, have been named *Hebbian* (directly or in a generalized sense, e.g. the *covariance* “learning rule” and *PCA*). ICA goes beyond second-order statistics of PCA, because it also processes higher-order terms in a series of statistical quantities – moments or cumulants. This is related with neuronal interactions of higher order, i.e. more than two neurons may interact at the same time because they may be directly connected (e.g., having synapses close to each-other and affecting one another), or their activity may be coupled in another way (e.g., as in coherent oscillations).

In the case of *oscillatory activities* in neurons and subcellular structures including dendritic fields, the relevant variables become local phases and their relations (King *et al.*, 2000). Oscillations are described by functions of complex-valued exponents (the exponent without imaginary unit i is the phase).³

Phases needed for edges. Phase-information is needed for successful approximation of V1-filters which were found to be *localized*, *oriented* and *band-pass* (i.e., selective to structure at different spatial scales). Local angle of orientation is described by local phase. Such filters are needed to trace segments of edges which are themselves oriented. Experiments show that individual *simple* cells of V1 with their specific receptive fields act as such filters, i.e. as selectors or edge-segments by having maximal response to specifically oriented stimuli.

An edge (of an object), as a crucial element of an image, manifests specific relationships among many pixels being encoded in neurons or receptors, not only two (neighboring) ones. Second-order statistics (as in PCA) is sensitive *only to pair-wise relationships*, like correlations encoded in the Hebb rule. Higher-order statistics (as in ICA) is sensitive to multi-neuronal relationships, reflecting *multi-pixel gestalt-structures*, and thus goes beyond the two-pixel (or two-neuron, respectively) relations of PCA. In PCA, gestalts are formed from feature-segments by global organization from local *bilateral* connections (usual simple synapses). In ICA, however, gestalts are formed directly by semi-local *multilateral* connections (complex synaptic structures⁴). This allows ICA to detect edges by *filters* which are essentially more *localized* (*wavelet*-like) than it would be achievable by PCA performing *global* (*Fourier*-like) spatial and spatial-frequency analysis.

25 more pages are available in the full version of this document, which may be purchased using the "Add to Cart" button on the publisher's webpage: www.igi-global.com/chapter/computational-information-maximization-models/50502

Related Content

Particle Swarm Optimization in Bioinformatics, Image Processing, and Computational Linguistics

Badal Soni, Satashree Royand Shiv Warsi (2024). *Research Anthology on Bioinformatics, Genomics, and Computational Biology* (pp. 1270-1292).

www.irma-international.org/chapter/particle-swarm-optimization-bioinformatics-image/342574

An Overview of Graph Indexing and Querying Techniques

Sherif Sakrand Ghazi Al-Naymat (2013). *Bioinformatics: Concepts, Methodologies, Tools, and Applications* (pp. 222-239).

www.irma-international.org/chapter/overview-graph-indexing-querying-techniques/76064

Wave-SOM: A Novel Wavelet-Based Clustering Algorithm for Analysis of Gene Expression Patterns

Andrew Blanchard, Christopher Wolter, David S. McNabband Eitan Gross (2010). *International Journal of Knowledge Discovery in Bioinformatics* (pp. 50-73).

www.irma-international.org/article/wave-som-novel-wavelet-based/45165

Mining Frequent Boolean Expressions: Application to Gene Expression and Regulatory Modeling

Mohammed J. Zaki, Naren Ramakrishnanand Lizhuang Zhao (2010). *International Journal of Knowledge Discovery in Bioinformatics* (pp. 68-96).

www.irma-international.org/article/mining-frequent-boolean-expressions/47097

A Particle Swarm Optimization based Hybrid Recommendation System

Rabi Narayan Beheraand Sujata Dash (2016). *International Journal of Knowledge Discovery in Bioinformatics* (pp. 1-10).

www.irma-international.org/article/a-particle-swarm-optimization-based-hybrid-recommendation-system/172002