Appendix A: Principal Component Analysis

Principal Component Analysis (PCA) is almost equivalent to Singular Value Decomposition (SVA), or Karhunen-Loeve expansion. It will be presented first as an important computational method for feature extraction from input-data (Ritter, Martinetz & Schulten, 1992; Haken, 1996; AuxLit 10). To perform PCA, inputpatterns \vec{x}^k are decomposed into a series, i.e. a linear combination of prototypepatterns \vec{w}^r (r = 1, ..., p²):

$$\vec{x}^{k} = \vec{w}^{0} + \sum_{r=1}^{p'} \vec{w}^{r} c_{r}(\vec{x}^{k}) + \vec{R}(\vec{x}^{k})$$
(12.1)

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 $c_r = \vec{w}^r \vec{x}^k$ are the *principal components* of the input-patterns. Their number is p'. Variable c_r could also be treated as the activity-rate of corresponding dominant or cardinal neuron \vec{r} . Index r corresponds to location of a cardinal neuron \vec{r} . At the beginning, all principal components c_r are approximately equal and lower than 1. Then we can talk about potentially-cardinal neurons. Later environmental stimulus gives privilege to one pattern and its c_r increases towards 1 (wins). This means that one neuron becomes actually-cardinal, other neurons get subordinated. We are able to store all inputs \vec{x}^k completely (so, there is no need for data compression) if p = p'. In this "ideal" case index k and index r are equivalent. If, on the other hand, there is p > p', data-compression causes that a higher number (p) of inputs \vec{x}^k is represented by a lower number (p') of cardinal neurons c_r or/and their set of synapses \vec{w}^r (or \vec{w}^r , respectively). c_r -representation is a relatively sparse code; \vec{w}^r is a relatively distributed code.

 \vec{w}^0 specifies the center of weight of the input data distribution $P(\vec{x}^k)$ (\vec{w}^0 are very often represented by mean values – e.g., as in: Bankman in Pribram, 1993, p. 77). Other p' prototype-vectors \vec{w}^r form a so-called eigen-space basis. These basis-vectors, which are typically *mutually orthogonal, coincide with (relatively pictorial) memory representations*, i.e. so-called *eigen-images*. In fact, these p' vectors are eigen-vectors of the *autocorrelation- matrix* C that have *the largest eigen-values* (Ritter et al., 1992):¹

$$\mathbf{C} = \sum_{k=1}^{P} (\vec{x}^{k} - \vec{w}^{0}) \otimes (\vec{x}^{k} - \vec{w}^{0})^{T} P(\vec{x}^{k}).$$
(12.2)

T denotes the transposed vector (i.e., line vector); \otimes denotes the outer or tensor product: $(\vec{a} \otimes \vec{b})_{ij} = a_i b_j$. **C** is an (auto) correlation matrix of the difference-vector $\vec{x}^k - \vec{w}^k$, therefore it is a *covariance matrix*. Typically, mutually orthogonal basis-vectors capture the directions of maximal variance (dispersion) of the data.

Equation (12.1) defines a hyper-plane which passes through the center of weight \vec{w}^0 and is spanned by principal axes along all \vec{w}^r . $\vec{R}(\vec{x}^k)$ is a residual vector which represents a non- vanishing distance from the approximating hyper-plane perpendicular to it. If $\vec{R}(\vec{x}^k)$ would be zero, our approximation with principal eigenvectors \vec{w}^r (prototypes) or, equivalently, with principal components c_r (cardinal neurons corresponding to prototypes) would be perfect.

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