

A UNIFIED PERSPECTIVE ON ADVANCES OF INDEPENDENT SUBSPACES: BASIC, TEMPORAL, AND LOCAL STRUCTURES

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Abstract:

A general framework of independent subspaces is presented, based on which a number of unsupervised learning topics have been summarized from a unified perspective, featured by different combinations of three basic ingredients. Moreover, advances on these topics are overviewed in three streams, with roadmaps sketched. One consists of studies on the second order independence featured principal component analysis (PCA) and factor analysis (FA), in adaptive and robust implementations as well as with duality and temporal extensions. The other consists of studies on the higher order independence featured independent component analysis (ICA), binary FA, and nonGaussian FA. The third is called mixture based learning that combines the above individual tasks, proportionally or competitively to fulfill a complicated task.

Keywords:

Independence; Subspaces; PCA; MCA; Hebbian learning; Factor analysis (FA); ICA; Temporal FA; Binary FA; nonGaussian FA; Local subspaces; Local FA; Finite mixtures

1. Independent subspaces in a general framework

As shown in Fig.1(a), a sample x is projected to \hat{x} on a subspace with an error $e = x - \hat{x}$. It is nature to minimize the error of using \hat{x} to represent x , with the error e

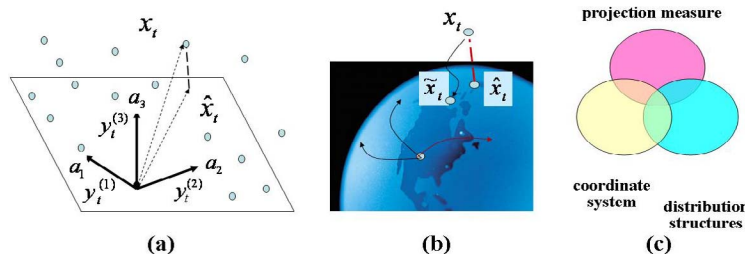


Fig.1 A general framework of independent subspaces

measured by its square length $d = \|e\|^2$ or the Euclidean distance between \hat{x} and x . The minimum is reached when

e is orthogonal to the subspace. Moreover, the subspace can be represented by a linear coordinate system, i.e., spanned by three linear independent unit basis vectors a_1, a_2, a_3 , \hat{x} can be further represented by its projection on each of the basis vectors, resulting in the coordinates $y = [y^{(1)}, y^{(2)}, y^{(3)}]^T$, i.e., $\hat{x} = \sum_j y^{(j)} a_j$ or generally

$$x = \hat{x} + e = Ay + e \tag{1}$$

where $\min E\|e\|^2$ is featured by the following natures:

- a) $Eey^T = 0$, (i.e., not correlated)
- b) e from $G(e|0, \sigma_e^2 I)$, $\sigma_e^2 = \min E\|e\|^2$,
- c) the coordinates in y is reached by an orthogonal transform $y = Wx$.

However, $\min E\|e\|^2$ does not lead to a unique A . Instead, it can consist of any linear independent unit basis vectors. To reduce this indeterminacy, we impose that

- d) a_1, a_2, a_3 are orthonormal basis vectors (i.e., $A^T A = I$), which implies that $Eyy^T = \Lambda$ is a diagonal matrix, i.e., $y^{(1)}, y^{(2)}, y^{(3)}$ are mutually not correlated or independent in a 2nd order statistics sense.

We further move to a general case shown in Fig.1(b). For a meaningful projection \hat{x} on a manifold, there are at least three basic ingredients to be specified. First, the error $e = x - \hat{x}$ needs a measure, based on which a minimum error projection can be implemented. Actually, different measures define different projections. One choice is $d = \|e\|^2$ for a homogeneous medium between \hat{x} and x . For inhomogeneous mediums, other choices can be used. One choice is

$$d = \|e\|_B^2 = e^T \Sigma_e^{-1} e \text{ with } \Sigma_e^{-1} = B^T B, \text{ as if } e \text{ is}$$

mapped to a homogeneous medium by a linear mapping B and measured by Euclidean distance. In Fig.2 this measure is considered, with its special case $d = \|e\|^2$ (i.e., $B=I$) and the degenerated case $e=0$.

To further represent \hat{x} within the manifold, we need the second ingredient, i.e., a coordinate system on the manifold, via either linear vectors in Fig.1(a) or a set of curves in Fig.1(b). In Fig.2, we still start at a linear coordinate system in Fig.1(a). However, minimizing $\|e\|_B^2$ no longer implies the nature a). Instead, it should be explicitly imposed as a condition. Also, the natures b) & c) are modified into

b) e from $G(e|0, \Sigma_e)$,

c) the coordinates y is reached by a linear $y=Wx$, but W is no longer orthogonal.

Moreover, impeding to reach the minimum $\|e\|_B^2$, the condition of $A^T A=I$ has to be replaced by a weaker condition:

d) $Eyy^T = \Lambda$ is a diagonal matrix, i.e., components of y becomes uncorrelated.

A re-scaling on components of y will not affect the natures a) & d). That is, there is an indeterminacy of a unknown diagonal matrix D . Also, there is an indeterminacy of a unknown rotation matrix Φ , and an indeterminacy on a specific allocation to its two additive terms in $Exx^T = \Lambda \Lambda^T + \Sigma_e$.

In fact, the above nature d) represents an example of the second ingredient, i.e., how y is distributed within an coordinate system. In Fig.2 we further extend it to an independence of any order statistics. The third ingredient varies from one linear coordinate system to multiple linear coordinate systems at different locations. Alternatively, each subspace can also be represented by another linear coordinate system for its complementary orthogonal subspace. Therefore, different specific types of independent

projection measure	$p(y_t^{(j)} \mu_t^{(j)})$	Gaussian $G(y_t^{(j)} \mu_t^{(j)}, \sigma_j^2)$ or even $G(y_t^{(j)} \mu_t^{(j)}, 1)$	(a) Gram-Charlier expansion (b) Mixture of scalar Gaussians	$y_t^{(j)}$ takes 0 or 1 $\mu_t^{(j)} y_t^{(j)} (1 - \mu_t^{(j)})^{1-y_t^{(j)}}$ $0 \leq \mu_t^{(j)} \leq 1$
$d=0, i.e., e=0, x=Ay$		Decorrelation e.g., whitening $Exx^T = L^T L$ $y = L^{-1}x$	ICA, $y = Wx$, $WA = D\Pi$ D is diagonal Π is permutation	ICA, $y^{(j)} = f(z^{(j)})$, e.g., $f(r) = (1 + e^{-r})^{-1}$ $z = Wx$, $WA = D\Pi$
$d = \ e\ ^2$ $x = Ay + e$ e from $G(e 0, \Sigma_e^2 I)$	$Ux = e$	PCA	NFA & P-ICA $y = Wx$ LMSER-ICA $y^{(j)} = f(z^{(j)})$ $z = Wx$	Binary-LMSER Non-negative PCA
	both	MCA	M-ICA MVNO-ICA	Binary MVNO
		Dual subspaces with each combining parts from both		
$d = e^T B^T B e$ $x = Ay + e$ $e \sim G(x 0, B^T B)$		Factor analysis (FA)	NonGaussian Factor analysis (NFA)	Binary Factor analysis (NFA)

Tab.1 Typical Examples of One Subspace for i.i.d. Samples

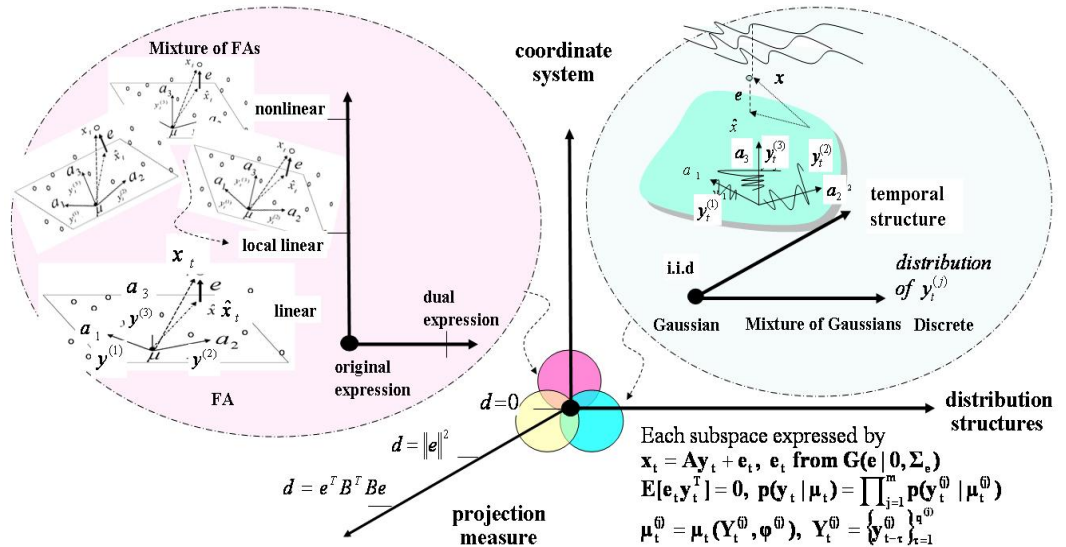


Fig.2 Typical variants of three ingredients

subspaces can be summarized from a unified perspective, featured by different combinations of the three ingredients, as shown in Fig.2. Also, from this perspective, an overview can be made on the past studies.

2. Independent subspaces of 2nd order independence

We start at a subspace for samples of independently and identically distributed (i.i.d.), as shown in Tab.1 and the two bottom dimensions in Fig.2. Thus, there is no temporal structure, we have $Y_t^{(j)}$ empty and $\mu_t^{(j)} = \mu^{(j)}$.

For a Gaussian $p(y_t^{(j)} | \mu^{(j)})$, we encounter factor

links among studies and on topics missing in the existing surveys and textbooks, to the author's best knowledge.

As shown on Roadmap A, this stream originated from over 100 years ago. The first adaptive learning one is Oja 1st-PC rule [66] that finds the first principal component (PC) without explicitly estimating the sample covariance Σ . Extended to find multi-PCs, one way is featured by an asymmetrical or a sequential implementation of one 1st-PC rule, but suffering error-accumulation. Readers are referred to [5,6,67,76,96] for overview. The other way is updating the weights of W symmetrically, e.g., Oja subspace rule [65]. Further studies are made in the following branches:

MCA, dual subspace, and TLS fitting Advocating to use not only a multi-PCs based subspace, but also its complementary part, i.e., minor components (MCs) that correspond the smallest eigenvalues of Σ , Xu, Krzyzak & Oja in 1991 suggests a dual pattern recognition with the first adaptive 1st-MC learning rule proposed via eqn.(11a) in [119]. Also, Minor component analysis (MCA) was firstly named by Xu, Oja & Suen in [116] and used for a total least square (TLS) curve fitting implemented by the above eqn.(11a) that finds the 1st-MC. Not only further progresses have been made on finding multi-MCs [62,63], but also this topic has been brought to the signal processing literature by Gao, Ahmad & Swamy [32] that was motivated by a visit of Gao to Xu's office where Xu introduced him the result of [116]. Thereafter, adaptive MCA learning for TLS filtering becomes a popular topic in the signal processing literature, e.g., see [24,30,58,60]. Also, efforts are made on performing either of PCA and MCA by simply switching the updating sign with a normalization as originally suggested in [119]. Since a PCA learning that converges correctly may become unstable or diverging after sign switching, studies have been made to examine the existing PCA rules on whether they remain stable after sign switching, while attempting to avoid its division computing for normalization. The jobs are quite tedious and need heavy mathematical analyses of ODE stability (e.g., Chen & Amari, [16]). The other line is turning an optimization of a PCA cost into a stable optimization of an induced cost for MCA. One example that turns the LMSER cost into one for a subspace spanned by MCs is given in [111]. Generally, for a cost $\min_W J(W)$ for PCA by its gradient descending $\Delta W = -\eta \nabla J(W)$, after switching sign we have the updating $\Delta W = \eta \nabla J(W)$ in one of the following three situations:

- Becoming divergence if $J(W)$ has no upper bound;
- Get a wrong solution if $J(W)$ is upper bounded but a maximum is reached at a non-orthogonal W ;

- Get the MCA solution if $J(W)$ is upper bounded and a maximum is reached at the MCA solution.

LMSER learning and Subspace tracking In Xu (1991)[118], a new adaptive PCA rule is derived from the gradient $\nabla_{E_2(W)}$ for a least mean square error reconstruction (LMSER). In [118], the first proof on global convergence of Oja subspace rule was provided, which was previously regarded as difficult. Further comparative studies were made on Oja rule and the LMSER rule, e.g., in [14,15,47,48,54,71,72], and shown both mathematically and experimentally that LMSER improves Oja rule in both performance and converging speed. Two years after [118], Yang (1993) uses this $E_2(W)$ via a recursive least square method for signal subspace tracking [120], then followed by others in the signal processing literature [30,55]. Alternatively, Xu in 1994 also pointed out that PCA and subspace analysis can also be performed by several other theories or cost functions [111,112]. Recently in [25,28], Fiori analyzed the algebraic and geometric properties of one among them, called relative uncertainty theory (RUT). Moreover, the NIC criterion for subspace tracking [58] is actually a special case of RUT, which can be observed by comparing eqn.(20) in [58] with equation of ρ_e at the end of Sec.III.B in [111].

Principal subspace vs multi-PCs Oja subspace rule reaches a principal subspace but not truly the multi-PCs due to a unknown rotation, while it is experimentally demonstrated by Xu in 1991 that the converged rows of W approximate the multi-PCs well by adding a sigmoid function $s(r)$ [118]. Worked at Harvard by the late summer 1991, Xu got aware of Brockett (1991)[11] and extended the Brockett flow of $n \times n$ orthogonal matrices to that of $n \times n_1$ orthogonal matrices with $n > n_1$, from which two learning rules for truly the multi-PCs are obtained from modifying the LMSER rule and Oja subspace rule accordingly. The two rules were included as eqns (13)&(14) in Xu [115] that was submitted in 1991, independently and differently from that of Oja [63]. In [83], Tanaka unifies these rules into an expression controlled by one parameter and makes a comparative study on them as well as the rules in [16], with eqn(14) in [115] shown to be the most promising one. In addition, the multi-PCs were also shown to be adaptively learned by several other theories or costs (Xu, 1994b) [112].

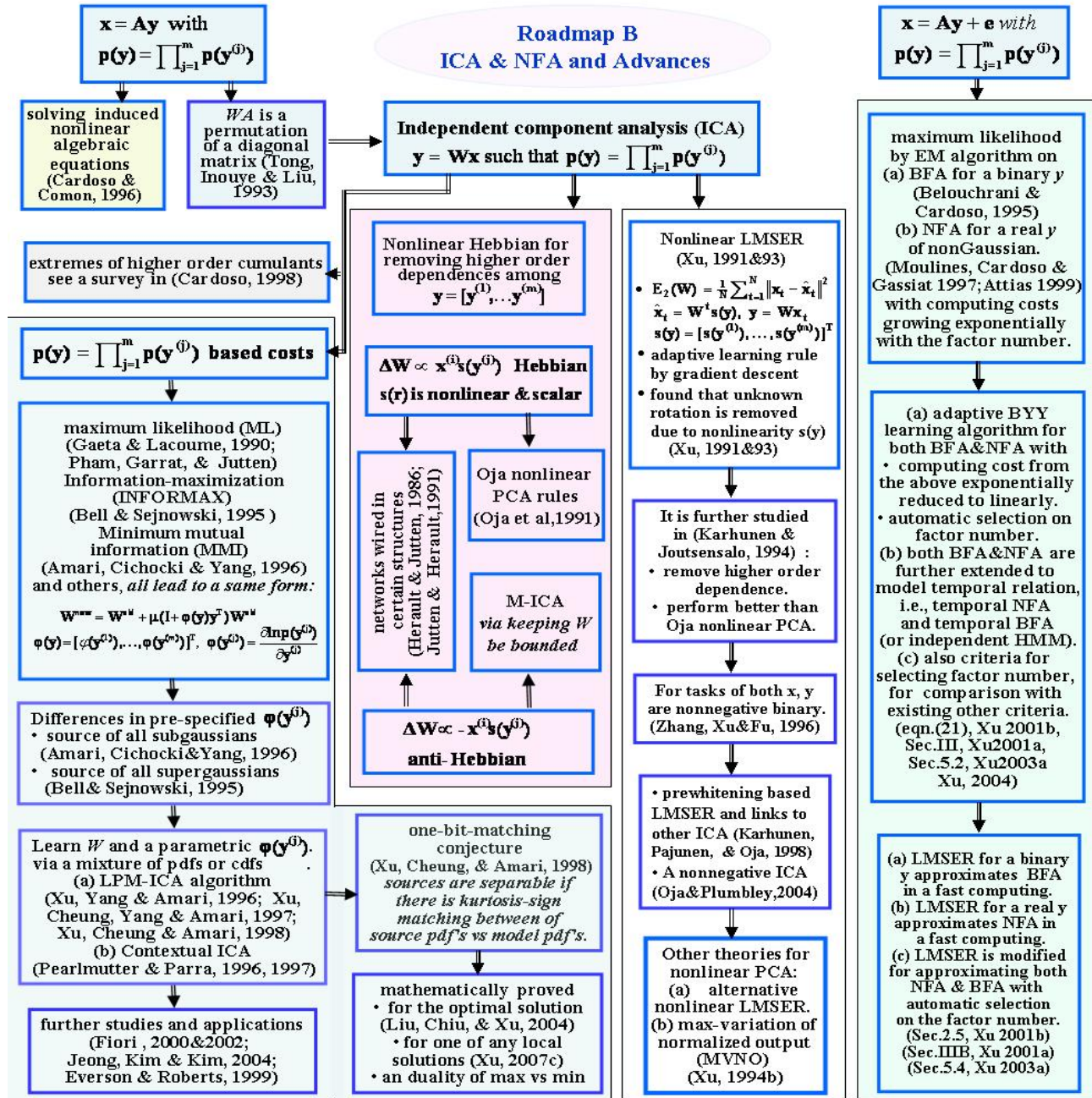
Adaptive robust PCA In the robust statistics literature, robust PCA was proposed to resist outliers via a robust

estimator on Σ [78,22]. First in 1992 [117] and then given in [117], Xu & Yuille generalized the rules of Oja, LMSER, and MCA into the robust adaptive learning by statistical physics, related to the M-estimators [40]. Also, PCA costs in [111] are extended to robust versions in Tab.2 of [112]. Thereafter, efforts have been further made, including its use in computer vision, e.g., [9,21,45,52].

On Roadmap A, another branch consists of advances

on FA that is equivalent to PCA in the special case $\Sigma_e = \sigma_e^2 I$ (shown firstly [3] in 1956, revisited in [85,102]).

In the past decade, there is a renewed interest on FA, not only the EM algorithm for FA [74] in is brought to PCA [75], but also adaptive EM algorithm and other advances are developed in help of the BYY harmony learning [96, 99,100,102,104], as listed on Roadmap A.



3. Independent subspaces of higher order independence

In Fig.2 we further extend $p(\mathbf{y}_t^{(i)} | \boldsymbol{\mu}_t^{(i)})$ to nonGaussian, with more constraint than a diagonal $E\mathbf{y}\mathbf{y}^T = \Lambda$ imposed by $\prod_{j=1}^m p(\mathbf{y}_t^{(j)} | \boldsymbol{\mu}_t^{(j)})$. We start at the degenerated case $e=0$, shown at the left-upper corner on Roadmap B. The problem is solving $\mathbf{x}=\mathbf{A}\mathbf{y}$ from samples of \mathbf{x} and a constraint on higher order independence among components of \mathbf{y} . One way to handle is solving induced nonlinear algebraic equations, with details referred to a survey paper [12]. From $\mathbf{x}=\mathbf{A}\mathbf{y}^*$ with at most one component of \mathbf{y}^* being Gaussian, it has been showed in [87] that “ $\mathbf{y}=\mathbf{W}\mathbf{x}$ becomes component-wise independent” means “ $\mathbf{y}=\mathbf{W}\mathbf{x}$ recovers \mathbf{y}^* up to scales and a permutation of components”. Thus, the problem can be turned into independent component analysis (ICA), which was further tackled in four branches, as shown on Roadmap B.

One is seeking extremes of the higher order cumulants of y [13]. The other branch is featured by nonlinear extension of Hebbian learning. One example combines Hebbian and anti-Hebbian rules in a network with nonlinear neurons [37,44]. Another example is Oja nonlinear PCA rules by adding nonlinearity into Hebbian learning [64].

The third branch optimizes a cost or principle that directly aims at the independence in a product of component densities. Interestingly, different aspects lead to a same updating equation shown on Roadmap B, with difference coming from pre-specifying the nonlinearity of $\phi(\mathbf{y}^{(j)})$. As a result, one works well when the source components of \mathbf{y}^* are all subgaussians [2] while the other works well when the source components of \mathbf{y}^* are all supergaussians [7]. The problem is solved by learning jointly \mathbf{W} and $\phi(\mathbf{y}^{(j)})$ via a parametric model [68, 106, 107, 108]. Also, it has been found that a rough estimate of each source pdf is enough for source separation, which leads to one-bit-matching conjecture that is mathematically proved to be true first in a weak sense that the global optimization is reached [51] and then in a strong sense that anyone of local optimal solutions is reached [94], also applicable to partial separation by a partial matching and a duality of maximization and minimization.

The last branch for ICA is featured by Nonlinear LMSER (Xu, 1991&93)[116,118], with details on Roadmap B. Additionally, clarifications are here made on two confusions. One relates to an omission of the origin of Nonlinear LMSER in [48], which has already been clarified in [41,47,71,72] that clearly spell out that both the nonlinear $E_2(\mathbf{W})$ and its adaptive gradient rule were firstly

proposed in (Xu, 1991)[118]. The second comes from that ICA is usually regarded as a counterpart of PCA [44]. As already stated in [96,100], it is inappropriate and yields confusion. It follows from the first row of Tab.1 that ICA by $\mathbf{y}=\mathbf{W}\mathbf{x}$ is an extension of decorrelation analysis, including any combinations of PCs and MCs, while it follows from the second row of Tab.1 that the counterpart of MCA is minor ICA (M-ICA) that is performed via nonlinear anti-Hebbian, minimizing cumulants, minimizing MVNO (See Roadmap B); while the counterpart of PCA is principal ICA (P-ICA) that is reached by nonlinear Hebbian, maximizing cumulants, and nonlinear LMSER.

Actually, the concept ‘principal’ involves $e=\mathbf{x}-\mathbf{A}\mathbf{y}\neq 0$, i.e., eq.(1) or the model at the right-upper corner on Roadmap B. It extends FA to the cases beyond that y is from a Gaussian, e.g., binary FA (BFA) if y is binary, and nonGaussian FA (NFA) if y is real but not from a Gaussian, as shown by the third row in Tab.1. Details are referred to the leftmost column on Roadmap B. Particularly, similar to that FA becomes equivalent to PCA in the special case $\Sigma_e = \sigma_e^2 I$, both BFA and NFA perform P-ICA at this special case too. As discussed in [96], by using a nonlinear map $y_i^{(j)} = s(z_i^{(j)})$ and $\mathbf{z}=\mathbf{W}\mathbf{x}$ to avoid expensive computing cost for y , the nonlinear LMSER implements either an approximate BFA with a Bernoulli $p(\mathbf{y}^{(j)})$ in probability $p_j = \frac{1}{N} \sum_{i=1}^N s(z_i^{(j)})$ or a real factor NFA with $p(\mathbf{y}^{(j)})$ being a pseudo uniform distribution on $[0,+\infty)$ or $(-\infty,+\infty)$ in help of BYY harmony learning [96], from which further results have been obtained too. Moreover, when $p(\mathbf{y}^{(j)})$ is a Bernoulli or on $[0,+\infty)$, nonlinear LMSER relates to nonnegative ICA [71,72].

In the cases with $e=\mathbf{x}-\mathbf{A}\mathbf{y}\neq 0$, another critical point is that it is impossible for a linear $\mathbf{y}=\mathbf{W}\mathbf{x}$ to lead to the minimum error $E\|\mathbf{e}\|_B^2$ or the maximum likelihood. Instead, a nonlinear map $\mathbf{y}=\mathbf{f}(\mathbf{x})$ is needed, which incurs open theoretical issues that will be addressed in the last section. From the aspect of algorithms, seeking an appropriate $\mathbf{y}=\mathbf{f}(\mathbf{x})$ incurs expensive computing costs too. Further structures should be imposed on either or both of $\mathbf{f}(\mathbf{x})$ and $p(\mathbf{y}_t^{(i)} | \boldsymbol{\mu}_t^{(i)})$. Nonlinear LMSER provides an approximate implementation with a post-sigmoid structure for $\mathbf{f}(\mathbf{x})$, which saves computing costs considerably and is experimentally shown to work well [48,71,72]. However, no quantitative analysis has been made on how this approximation affects performance yet. The EM algorithm in [4,61] implements the maximum likelihood learning by

considering $p(\mathbf{y}_t^{(i)} | \boldsymbol{\mu}^{(i)})$ via a Gaussian mixture, but suffering a computing cost that increases exponentially with the dimension m of \mathbf{y} . Also, the BYY harmony learning [99,100] considers $p(\mathbf{y}_t^{(i)} | \boldsymbol{\mu}^{(i)})$ in a Gaussian mixture or other parametric models, and a preliminary experimental study showed that it gets a performance similar to the above EM algorithm based maximum likelihood learning but with a computing cost linearly increasing with m [95]. Further comparative studies on the three approaches are needed both theoretically and experimentally.

4. Extensions to temporal and localized structures

We further move to observe the first row of Tab.2 that extends the third row in Tab.1 after considering temporal structures. A typical one is embedded in $p(\mathbf{y}_t^{(i)} | \boldsymbol{\mu}^{(i)})$

with $\boldsymbol{\mu}_t^{(i)} = \boldsymbol{\mu}^{(i)}(\mathbf{Y}_t^{(i)}, \boldsymbol{\varphi}_j)$, $\mathbf{Y}_t^{(i)} = \{\mathbf{y}_{t-\tau}^{(i)}\}_{\tau=1}^q$, e.g., a linear regression $\boldsymbol{\mu}_t^{(i)} = \sum_{\tau=1}^q \beta_{\tau}^{(j)} \mathbf{y}_{t-\tau}^{(i)}$. Information can be carried over time in two ways. One is via computing $\boldsymbol{\mu}_t^{(i)}$ by the regression, with learning on $\boldsymbol{\varphi}_j$ made in help of the gradient with respect to $\boldsymbol{\varphi}_j$ by the chain rule through the regression. The second way is via computing the integral $\int p(\mathbf{y}_t^{(i)} | \boldsymbol{\mu}_t^{(i)}) p(\mathbf{Y}_t^{(i)}) d\mathbf{Y}_t^{(i)}$ and getting the gradient with respect to $\boldsymbol{\varphi}_j$ by the chain rule through the integral. Details are referred to [95,96,99].

Shown in the second row of Tab.2 are extensions of third row in Tab.1 to multiple subspaces at different locations and thus under the name of local sth. Studies on this stream are summarized on Roadmap C, where a key point is how to allocate a sample x_t to different coordinate systems. There are two typical ways. One is made during implementing the EM algorithm for a maximum likelihood (ML) learning or a Bayesian learning. The other is made via competition, including the classic competitive learning, and the rival penalized competitive learning (RPCL) and the one in the Bayesian Ying Yang (BYY) harmony learning [92,93].

$\mathbf{d} = \mathbf{e}^T \mathbf{B}^T \mathbf{B} \mathbf{e}$ $\mathbf{x} = \mathbf{A} \mathbf{y} + \mathbf{e}$	$p(\mathbf{y}_t^{(i)} \boldsymbol{\mu}_t^{(i)})$ Gaussian $G(\mathbf{y}_t^{(i)} \boldsymbol{\mu}_t^{(i)}, \boldsymbol{\sigma}_t^2)$ or even $G(\mathbf{y}_t^{(i)} \boldsymbol{\mu}_t^{(i)}, \mathbf{1})$	(a) Gram-Charlier expansion (b) Mixture of scalar Gaussians	$\mathbf{y}_t^{(i)}$ takes 0 or 1 $\boldsymbol{\mu}_t^{(i) \text{ y}^{(i)}} (\mathbf{1} - \boldsymbol{\mu}_t^{(i)})^{1 - \mathbf{y}^{(i)}}$ $0 \leq \boldsymbol{\mu}_t^{(i)} \leq 1$
From i.i.d. samples to temporal samples $\boldsymbol{\mu}_t^{(i)} = \boldsymbol{\mu}^{(i)}(\mathbf{y}_t^{(i)}, \boldsymbol{\varphi}_j)$ $\mathbf{y}_t^{(i)} = \{\mathbf{y}_{t-\tau}^{(i)}\}_{\tau=1}^q$	Temporal factor analysis (TFA)	Temporal nonGaussian Factor analysis (T-NFA) Temporal LMSER (T-LMSER) $\mathbf{y}^{(i)} = f(\mathbf{z}^{(i)}), \mathbf{z} = \mathbf{W} \mathbf{x}$	Temporal binary factor analysis (T-BFA) or independent HMM binary T-LMSER
From one subspace to multiple subspaces	Local FA (LFA) Local PCA/PSA	Local NFA LOCAL LMSER	Local BFA Local LMSER
Both from i.i.d. to temporal from one to multiple	Local TFA	Local T-NFA Local T-LMSER	Local T-BFA Local independent HMM Local T-LMSER

Tab.2 Typical examples with temporal and localized structures

5. Subspace number and subspace dimension

Roadmap C Several Types of Mixtures of Subspaces

models		$p(\mathbf{y})$ is Gaussian		$p(\mathbf{y})$ is nonGaussian	
learning		PCA/FA/TFA	MCA/Surface fitting	ICA	NFA/BFA/LMSER and temporal extensions
Competitive Learning	Classic	Local PCA (Sec.3.2, Xu 1995) (Kambhatla & Leen, 1997) PCA competitive learning (Lopez-Rubio, et al 2004)	Local MCA by MML (Sec.4.1, Xu 1995)	Competitive ICA (eqn.37, Xu 2001b) Competitive Temporal ICA (eqn. 88, Xu 2001a)	the degenerated cases from the ones below
	RPCL & BYY	Local PCA, Local FA, Local LMSER (eqns.14&16, Xu 1998b) (Sec.4.3, Xu 2001b) Local FA (Sec.3.3.2, Xu, 2007a) Local TFA (Xu, 2004)	Local MCA by MML (eqn.15, Xu, 1998b) (Sec.4.2, Xu, 2001b)	Improved competitive ICA (Sec.4, Xu 2002)	Local BFA, NFA, LMSER (Sec.4, Xu 2002) (eqns.43&44, Xu 2001b) temporal extensions (Xu, 2004)
with automatic selection on the number of subspaces or both of the number and dimensions of subspaces.					
EM Algorithm	ML	Local PCA by a simplified EM (Sec.V(B)(D), Xu 1994c) Mixtures of FA (Ghahramani & Hinton, 1996) Mixtures of probabilistic PCA (Tipping & Bishop, 1999)	Local MCA by a simplified EM (Sec.V(C)(D), Xu 1994e) MCA Co-integration (Xu & Leung, 1998) Probabilistic MCA (Williams & Agakov, 2002)	ICA mixture (Lee, Lewicki, & Sejnowski, 2000)	One possible way is getting extension from (Moulines, Cardoso & Gassiat 1997; Attias 1999) but with much expensive computing costs
	Bayesian	Variational Mixture (Ghahramani & Beal, 2000; Utsugi & Kumagai, 2001)		Variational Mixture (Choudrey & Roberts 2003)	

Another important problem is how to determine the number k of subspaces and the dimension m_ℓ of each subspace. It is usually referred under the name model selection. A classic way is implementing a two-phase procedure. First, a number of candidates are learned by enumerating k and m_ℓ . Second, a criterion is used to select the best among the candidates. An example is using AIC for one coordinate system based FA [10]. Other choices of criteria include BIC or equivalently MDL, the cross validation, etc. However, such a two-phase implementation is computationally very extensive and thus impractical. Moreover, when the sample

size is finite and k , m_ℓ are not too small, the values of criterion are difficult to be estimated accurately, which makes the performance degenerate considerably.

Alternative solutions have been sought. One type includes incremental approaches, e.g., as m_ℓ increases to $m_\ell + 1$, learning is made incrementally with the parts already learned kept or partially adjusted such that redundant computing can be saved. However, it usually leads to a suboptimal performance because not only newly added parameters have to be learned, but also the old parameter set have to be relearned. Oppositely, we can decrease from m_ℓ to $m_\ell - 1$. Still we can not simply discard those extra parameters, i.e., all the parameters in the case $m_\ell - 1$ have to be re-learned.

Another direction is seeking model selection made automatically during learning. That is, in a model with k and m_ℓ initially set to be large enough such that the correct one is included, learning will not only determine parameters but also automatically shrink k and m_ℓ down to appropriate ones. One such an effort is RPCL [114], which can make k determined automatically during learning. The other and better choice is the BYY harmony learning that is a general approach applicable to various statistical learning tasks with automatic model selection. Readers are referred to [92,93] for an adaptive algorithm to implement local FA with automatic determination of k and m_ℓ , and for a systematic comparative experimental study on a large number of simulated data sets and several real data sets from UCI repository, in comparison with AIC, CAIC, BIC/MDL, CV as well as two incremental approaches [34,80]. Both performances and computing times are compared, and it can be observed that the BYY harmony learning algorithm outperforms the counterparts considerably. Readers are referred to [92,93,95,97,98,99,100] for the BYY harmony learning based criteria and algorithms for the rest cases in Tabs 1 & 2.

4. Concluding remarks

Studies of three closely related unsupervised learning streams have been overviewed in an extensive scope and from a rather systematic perspective. A general framework of independent subspaces is presented, from which a number of learning topics are summarized via its features of choosing and combining the three basic ingredients.

There are already extensive studies on the cases with independence in a sense of second order statistics. Also, there are extensive studies on ICA with noise free (i.e, $e=0$).

Trends move towards the cases with components of y being mutually independent in higher order statistics and with noise $e \neq 0$. Though a few algorithms are available, further comparative studies on them are needed both theoretically and experimentally. Even more importantly, there are still a number of open issues, some of which are listed below:

- Which part of unknown parameters in $x = Ay + e$ can be determined uniquely? which part is indeterminable and how it can be improved?
- Under what conditions, the independence of $\prod_{j=1}^m p(y_t^{(j)} | \mu_t^{(j)})$ can be ensured conceptually? How can it be further achieved by a learning algorithm?
- When and under what condition, $\hat{x} = Ay$ can be said to be the best reconstruction of x ?
- Under what condition, both ensuring $\prod_{j=1}^m p(y_t^{(j)} | \mu_t^{(j)})$ and the best reconstruction of x by $\hat{x} = Ay$ can be achieved jointly? If not, how to trade off the two?
- What is the best nonlinear map $y = f(x)$ in term of $p(y_t^{(j)} | \mu_t^{(j)})$ and e ? Can the best be obtained analytically or via an effective computing?

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