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ICA PERMUTATION AMBIGUITY IS FIXED WITH ROUGH GUESSES ON SOURCE KURTOSES

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ICA Permutation Ambiguity is Fixed with Rough Guesses on Source Kurtoses

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Résumé

Ce rapport traite du problème de la séparation aveugle de sources via l'Analyse en Composantes Indépendantes (ICA). Nous montrons qu'une combinaison linéaire des cumulants marginaux d'ordre quatre (kurtosis) des sorties du séparateur fournit une fonction de contraste recevable, sous la condition que les données ont été pré-blanchies spatialement, si les poids de la combinaison sont de même signe que les kurtosis des sources. Si les poids sont égaux aux kurtosis des sources, alors le contraste peut être vu comme un crière d'ajustement de cumulants revenant au maximum de vraisemblance. En outre, si les poids sont distincts et les kurtosis des sources sont distincts, alors l'ambiguïté de permutation, inhérente au problème de l'ICA, disparaît. Les sources peuvent être estimées par ordre de kurtosis croissant. Ceci étend un petit résultat déjà publié, que nous avons ajouté en annexe.

Abstract

The present report addresses the problem of blind source separation via independent component analysis (ICA). We prove that a linear combination of the separator output fourth-order marginal cumulants (kurtoses) is a valid contrast function for ICA under the prewhitening assumption if the weights have the same sign as the actual source kurtoses. If the weights equal the actual source kurtoses, the contrast is a cumulant matching criterion based on the maximum likelihood principle. If the source kurtoses are different and so are the linear combination weights, the contrast eliminates the permutation ambiguity typical to ICA, as the estimated sources are sorted in increasing kurtosis value at the separator output. This extends a previously published result, which we have included in appendix.

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Index Terms

Independent Component Analysis, contrast criteria, kurtosis, Blind Source Separation, Maximum Likelihood.

I. Introduction

One considers the problem of Blind Source Separation (BSS), where a set of N real or complex independent sources, s_n , $1 \le n \le N$, are mixed and observed on N sensors. If data measurements have been spatially prewhitened, it is legitimate to assume that the mixing matrix is unitary, so that the observation model takes the form:

$$\mathbf{x} = \mathbf{Q}\mathbf{s} \tag{1}$$

where Q is a $N \times N$ unitary matrix, and $\mathbf{x} \in \mathbb{C}^N$. The goal is to recover source realizations from the sole observation of realizations of random variable \mathbf{x} . For this purpose, one may estimate the best separating matrix \mathbf{F} so that the output vector

$$y = Fx$$

is equal to the source vector \mathbf{s} up to scale and permutation factors. That is, with the sole assumption that s_n are statistically independent, the best we can hope is to obtain an estimate of \mathbf{s} of the form $\mathbf{y} = \mathbf{\Lambda} \mathbf{P} \mathbf{s}$, where $\mathbf{\Lambda}$ is diagonal invertible and \mathbf{P} is a permutation. This problem is referred to as the Independent Component Analysis (ICA); refer to [1], [2] and references therein.

In [3], it has been proved that a prior knowledge of the source kurtosis signs can fix the permutation ambiguity between sources of different kurtosis signs. As a consequence, it was possible to extract the source of interest in first position if it was the only one to have a positive (resp. negative) kurtosis in the mixture. The resulting computational complexity could hence be reduced, if a special purpose pair-sweeping algorithm is utilized [3].

The present report shows that the permutation ambiguity can be reduced even further if it is known in advance that sources have different kurtoses.

a) Multi-linear relation: The kurtosis μ_i of the separator output y_i is related to those of the observations thanks to the multi-linearity of cumulants. More precisely, we have:

$$\mu_i = \sum_{mnpq} F_{im} F_{ip} F_{ip}^* F_{iq}^* \gamma_{mnpq} \tag{2}$$

where $\mu_i = \text{Cum}\{y_i, y_i, y_i^*, y_i^*\}$ and $\gamma_{mnpq} = \text{Cum}\{x_m, x_n, x_p^*, x_q^*\}$. On the other hand, if we denote G the global filter, i.e. G = FQ, they are also related to source cumulants as:

$$\mu_i = \sum_{n=1}^N |G_{in}|^4 \kappa_n$$

b) First assumption: Denote κ_n the source kurtoses, and assume indices are chosen so that κ_n is non decreasing: $\kappa_{n+1} \geq \kappa_n$, $\forall n$. In addition, assume that the first p sources are known to have a positive kurtosis and the remaining n-p a negative one:

$$\begin{cases} \kappa_n > 0, \forall n : p \le n \le N \\ \kappa_n < 0, \forall n : 1 \le n < p \end{cases}$$
(3)

Denote S the set of sources satisfying (3), and Y the set of observations generated by the orthogonal group Q acting on S. Then with these notations we have the following result:

Proposition 1: The optimization criterion $\Psi_p(\mathbf{y})$ defined as:

$$\Psi_{\varepsilon}(\mathbf{Q}) = \sum_{i=1}^{N} \varepsilon_{i} \mu_{i} \tag{4}$$

where $\varepsilon_i = 1$ for $1 \le i \le p$, and $\varepsilon_i = -1$ for $p < i \le N$ is a contrast function over the set of observations $\mathcal{Y} = \mathcal{Q} \cdot \mathcal{S}$.

The proof is given in [3].

c) Second assumption: Now we assume instead that we are given a set of real numbers, α_i , related to the unknown source kurtoses κ_i via an unknown strictly increasing function f passing through the origin: $\alpha_i = f(\kappa_i)$. In other words, we know not only how many positive and negative kurtoses there are, but we also know how many are equal and which ones. For instance, if $\alpha_1 < \alpha_2 \le \alpha_3 < 0 \le \alpha_4$, then it means that $\kappa_1 < \kappa_2 \le \kappa_3 < 0 \le \kappa_4$. Note that because κ_i is non decreasing, so is α_i .

In practice, it often happens that we have enough information to know such an ordering, but not enough to know the source kurtoses with a good accuracy. This lack of accuracy prevents us from resorting to the Maximum Likelihood criterion, and one generally ignores the knowledge of ordering and executes a standard ICA algorithm.

II. NEW OPTIMIZATION CRITERION

The optimization criterion that we propose is a linear combination of output kurtoses, $\sum_i \alpha_i \mu_i$ where $\alpha_i \in \mathbb{R}$ are given and sorted in non decreasing order, and cumulants μ_i are calculated with the help of (2) where γ_{mnpq} are estimated from measurements, $1 \leq i \leq N$. Propositions 2 and 3 rigorously show that it is possible to recover the sources in a pre-assigned order, by maximizing criterion (5) with respect to \mathbf{F} .

Proposition 2: The optimization criterion

$$\Psi_{\alpha}(\mathbf{y}) = \sum_{i=1}^{N} \alpha_i \mu_i \tag{5}$$

is a contrast function over the set of observations $\mathcal{Y} = \mathcal{Q} \cdot \mathcal{S}$.

Lemma 1: Let \mathbf{u} and \mathbf{v} be two vectors of \mathbb{R}^N . Then the permutation \mathbf{P} that maximizes the scalar product $\mathbf{u}^T \mathbf{P} \mathbf{v}$ is that yielding the scalar product value $\sum_i u_{\sigma(i)} v_{\pi(i)}$ where entries $u_{\sigma(i)}$ and $v_{\pi(i)}$ are both sorted in non decreasing order.

Proof: The proof is obvious, and proceeds by contradiction. Assume that for the optimal permutation, u_i are sorted in non decreasing order, but assume that there exist two entries of \mathbf{v} such that $v_k > v_{k+p}$. By construction, we have $(u_{k+p} - u_k)(v_k - v_{k+p}) > 0$. By expanding the product we get $u_{k+p}v_k + u_kv_{k+p} > u_kv_k + u_{k+p}v_{k+p}$, which shows that transposing the two entries of \mathbf{v} increases the scalar product. Hence the permutation was not optimal.

Now let's prove the proposition.

Proof:

Case 1: Distinct α_i 's. By definition (5),

$$\Psi_{\alpha}(\mathbf{y}) \leq \sum_{i} |\alpha_{i}| \left| \sum_{j} G_{ij}^{2} G_{ij}^{2*} \kappa_{j} \right|$$

$$\leq \sum_{ij} |\alpha_{i}| |G_{ij}|^{4} |\kappa_{j}|$$
(6)

Now, recall that ${f G}$ is unitary. Hence $|G_{ij}|^4 \leq |G_{ij}|^2$ for any indices, hence:

$$\Psi_{\alpha}(\mathbf{y}) \le \sum_{ij} |\alpha_i| |G_{ij}|^2 |\kappa_j| \tag{7}$$

Yet, the matrix formed with entries $|G_{ij}|^2$ is itself bistochastic since its rows and columns sum up to one. Hence, from the Birkhoff theorem [4], there exist a set of real positive numbers β_{ℓ} such that

$$|G_{ij}|^2 = \sum_{\ell} \beta_{\ell} P_{ij}(\ell)$$
, and $\sum_{\ell} \beta_{\ell} = 1$

where $\mathbf{P}(\ell)$ are permutations matrices. This yields the inequality below

$$\Psi_{\alpha}(\mathbf{y}) \leq \sum_{i,j} |\alpha_i| |\kappa_j| \sum_{\ell} \beta_{\ell} P_{ij}(\ell)$$

The maximum of the right hand side is reached when the convex linear combination reduces to one of its vertex, that is when all β 's are null but one, say $\beta(\ell_o)$. Then from the lemma, $\mathbf{P}(\ell_o)$ precisely relates j and i, so that both $|\alpha_j|$ and $|\kappa_j|$ are sorted in increasing order:

$$\Psi_{\alpha}(\mathbf{y}) \le \sum_{j} |\alpha_{j} \kappa_{j}| = \Psi_{\alpha}(\mathbf{s}) \tag{8}$$

If we have equality, then the same reasoning as in [3] would show that $G = \Lambda P$.

Case 2: Possibly non distinct α_i 's. When α_i 's are not distinct, we can group them by packets of equal values, \mathcal{A}_q . Similarly, values of κ_i can be grouped within the same packets, according to our second assumption. Since permuting indices within a set \mathcal{A}_q does not change the value of the criterion, the proof still holds true.

Proposition 3: If equality holds in (5), then $\mathbf{y} = \mathbf{\Lambda} \mathbf{P} \mathbf{s}$, where permutation \mathbf{P} is equal to the identity matrix for every row i (or column i) for which α_i is distinct from the other α_j 's. In addition, the entries of the diagonal matrix $\mathbf{\Lambda}$ must be of unit modulus.

Proof: Now we shall make use of the fact that not only moduli $|\alpha_i|$ are sorted, but also α_i 's. If equality holds in (8), it means in particular that there exists a permutation **P** such that:

$$\Psi_{\alpha}(\mathbf{y}) = \sum_{ij} \alpha_i P_{ij} \kappa_j = \sum_j \alpha_j \kappa_j = \Psi_{\alpha}(\mathbf{s})$$

From lemma 1, we know that permutation \mathbf{P} is uniquely defined if there is a unique way to sort the κ_n in increasing order. This will be the case if all source kurtoses κ_n are distinct. Should not this be the case, the permutation is not unique: any permutation of indices keeping the order of κ_n non decreasing will still lead to the same maximum of the contrast. The permutation indetermination \mathbf{P} is then formed of diagonal blocks $\mathbf{D}(q)$, whose size corresponds to the number of elements in each set \mathcal{A}_q .

Proposition 2 of [3] may now be seen as a particular case of Proposition 3 above, where coefficients α_i are set to ± 1 .

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A Contrast for Independent Component Analysis with Priors on the Source Kurtosis Signs

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Abstract—A contrast function for Independent Component Analysis (ICA) is presented incorporating the prior knowledge on the sub-Gaussian or super-Gaussian character of the sources as described by their kurtosis signs. The contrast is related to the maximum likelihood principle, reduces the permutation indeterminacy typical of ICA, and proves particularly useful in the direct extraction of a source signal with distinct kurtosis sign. In addition, its numerical maximization can be performed cost-effectively by a Jacobi-like pairwise iteration. Extensions to standardized cumulants of orders other than four are also given.

Index Terms—Blind Source Separation, contrast functions, higher-order statistics, Independent Component Analysis, kurtosis, performance analysis, standardized cumulants.

EDICS: SAS-ICAB (Independent Component Analysis and Blind Source Separation).

I. Introduction

NDEPENDENT Component Analysis (ICA) aims at maximizing the statistical independence between the entries of multivariate data. ICA is the fundamental technique for Blind Source Separation (BSS) in linear mixtures when the sources are assumed mutually independent [1]. The plausibility of the assumption in a wide variety of applications has rapidly made of ICA a reference tool in biomedical engineering, communications and image processing, among many other domains [2], [3], [4].

In the real-valued case, ICA assumes the following linear model for the observed data vector $\mathbf{x} \in \mathbb{R}^m$:

$$\mathbf{x} = \mathbf{H}\mathbf{s} \tag{1}$$

where $s \in \mathbb{R}^n$ contains the independent components or sources and $\mathbf{H} \in \mathbb{R}^{m \times n}$ represents the mixing matrix, with $m \geq n$. The sources are recovered by maximizing a so-called contrast function measuring the statistical independence between the separator output components [1]. Seminal contrasts such as 'COM1' and 'COM2' originated from cumulant-based approximations (usually at order four) of information-theoretical principles such as maximum likelihood (ML), mutual information and marginal entropy [1], [5]. The hypothesis that the kurtosis (normalized fourth-order marginal cumulant) of all the sources has the same sign allows the definition of

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computationally simpler contrasts [5], [6], but is unable to reduce the ambiguity in the ordering of the recovered sources, or permutation indeterminacy, typical in BSS.

The power of the blind approach lies in its robustness to modelling errors, a feature achieved by making as few assumptions about the problem as possible. However, additional information is often available in practice such as the non-Gaussian character of the sources: that of a digital modulation signal depends on the relative probability of its symbols; the atrial activity signal of an atrial fibrillation electrocardiogram is usually sub-Gaussian or quasi-Gaussian; etc. Separation performance can be considerably improved by capitalizing on this information.

The present contribution puts forward a contrast function that takes into account the prior knowledge about the non-Gaussian character of the sources. The new contrast has optimality properties in the ML sense, is efficiently maximized by Jacobi-like iterations, and alleviates (indeed may totally resolve) the permutation indeterminacy left by blind processing. This latter feature, illustrated in Sec. IV through simulations, has been successfully put into practice, without mathematical proof, on real signals issued from electrocardiography [7], [8].

II. A CONTRAST BASED ON SOURCE KURTOSIS SIGNS

Let us first recall the concept of contrast function. The standardization or whitening (second-order processing) of observation (1) yields another vector $\mathbf{z} = \mathbf{Q}\mathbf{s}$, where \mathbf{Q} is a unitary matrix. The sources can then be recovered by applying a unitary transform $\hat{\mathbf{Q}}$, resulting in the separator output $\mathbf{y} = \hat{\mathbf{Q}}^T\mathbf{z} = \mathbf{G}\mathbf{s}$, where $\mathbf{G} = \hat{\mathbf{Q}}^T\mathbf{Q}$. A function $\Psi(\mathbf{y})$ of the separator-output distribution is an orthogonal contrast for ICA if $\Psi(\mathbf{s}) \geq \Psi(\mathbf{G}\mathbf{s})$, for any orthogonal matrix \mathbf{G} (domination), with equality if and only if \mathbf{G} is a trivial filter

$$G = PD (2)$$

where \mathbf{P} is a permutation and \mathbf{D} a non-singular diagonal matrix (discrimination). Consequently, contrast maximization restores the independent sources at the separator output up to a possible permutation and scaling.

Let κ_i denote the ith-source kurtosis and ε_i its sign, $\varepsilon_i = \mathrm{sign}(\kappa_i), \ 1 \leq i \leq n.$ We assume in the sequel that p sources have positive kurtosis, $\varepsilon_i = 1, \ 1 \leq i \leq p,$ and (n-p) sources have negative kurtosis, $\varepsilon_i = -1, \ p < i \leq n.$ Symbol μ_i represents the kurtosis of the separator's ith output. Proofs for the mathematical results that follow can be found in the Appendix.

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Proposition 1: Criterion

$$\Psi_p(\mathbf{y}) = \sum_{i=1}^n \varepsilon_i \mu_i \tag{3}$$

is a contrast function under the above assumptions.

Remark: The maximum likelihood recovery of the source signals under the whitening constraint is achieved by maximizing the function:

$$\Psi_{\rm ML}(\mathbf{y}) = \sum_{i=1}^{n} \kappa_i \mu_i. \tag{4}$$

This contrast is obtained from an approximation of the Kullback-Leibler divergence based on the Edgeworth expansion of the separator-output probability density function (pdf) truncated at fourth order [6]. If only the source kurtosis signs are known, contrast (4) naturally reduces to (3). Hence, the latter is expected to inherit the optimality features of the approximate ML estimate while reducing the prior information required. The reduced amount of information helps to keep the desirable features of a blind formulation and is capable of partially solving the permutation ambiguity, as shown by Proposition 2 below.

Remark: Reference [9] addresses the so-called one-bit matching conjecture, whereby the sources can be separated if there exists a one-to-one correspondence between the kurtosis signs of the sources and those resulting from the truncated Gram-Charlier expansion of their pdf's. A function obtained in [9] bears certain resemblance to contrast (3) but the proof of the conjecture is cumbersome and valid only when the source skewness (standardized third-order cumulant) is null. We prove in the Appendix that function (3) is a contrast for all orders $r \geq 3$, of which Proposition 1 is just a particular case for r = 4.

Proposition 2: Trivial filters associated with contrast (3) are of the form (2), where

$$\mathbf{P} = \left(\begin{array}{cc} \mathbf{P}_1 & \mathbf{0} \\ \mathbf{0} & \mathbf{P}_2 \end{array} \right). \tag{5}$$

 \mathbf{P}_1 and \mathbf{P}_2 being permutation matrices of size $p \times p$ and $(n-p) \times (n-p)$, respectively, and \mathbf{D} made up of unit-norm diagonal entries.

Remark: Sources with positive kurtosis are extracted separately from sources with negative kurtosis by contrast (3), provided that parameter p is known. In particular, a source of interest can be recovered without permutation ambiguity if its kurtosis sign is different from all the others'. The Appendix shows that contrast (3) enjoys this source ordering property for standardized cumulants of even order $r \geq 4$.

III. CONTRAST OPTIMIZATION

The Jacobi-like pairwise iteration technique originally proposed in [1] can also be used to optimize contrast function (3). The function is maximized for each signal pair in turn over several sweeps until convergence. Let us assume that we are processing pair $\mathbf{z}_{12} = [z_1, z_2]^T$, the result being

readily adapted to other pairs by a simple change of indices. The corresponding two-signal separator output is given by $\mathbf{y}_{12} = \hat{\mathbf{Q}}^T \mathbf{z}_{12}$, where $\hat{\mathbf{Q}}$ is a Givens rotation that can be parameterized as

$$\hat{\mathbf{Q}}(\theta) = \frac{1}{\sqrt{1+t^2}} \begin{pmatrix} 1 & -t \\ t & 1 \end{pmatrix} \tag{6}$$

with $t = \tan \theta$. The associated pairwise contrast is $\Psi(\mathbf{y}_{12}) = \varepsilon_1 \mu_1 + \varepsilon_2 \mu_2$. By virtue of the multilinearity property of cumulants, this function can easily be expressed in terms of the unknown t and the 4th-order cumulants of \mathbf{z}_{12} , denoted as $c_{ij} = \operatorname{Cum}_{ij}(z_1, z_2)$, with (i+j) = 4 (using Kendall's notation). The stationary points of $\Psi(\mathbf{y}_{12})$ are then found to be the solutions to the quartic equation:

$$a_3t^4 + 2(a_2 - 2a_4)t^3 + 3(a_1 - a_3)t^2 + 2(2a_0 - a_2)t - a_1 = 0$$
 (7)

where $a_0=(\varepsilon_1c_{40}+\varepsilon_2c_{04}),\ a_1=4(\varepsilon_1c_{31}-\varepsilon_2c_{13}),\ a_2=6(\varepsilon_1+\varepsilon_2)c_{22},\ a_3=4(\varepsilon_1c_{13}-\varepsilon_2c_{31}),\ and\ a_4=(\varepsilon_1c_{04}+\varepsilon_2c_{40}).$ The above quartic can be solved by radicals (Ferrari's formula) at a cost that can be considered negligible compared to the cumulant computation. The solutions can also be simply expressed in terms of the extended ML (EML) estimator of [10] if $\varepsilon_1=\varepsilon_2$ or the alternative EML (AEML) estimator of [11] if $\varepsilon_1\neq\varepsilon_2$. Typically, about $O(\sqrt{n})$ sweeps over all signal pairs are required for convergence, as suggested in [1]. However, as a by-product of Proposition 2, the extraction of a source of interest with distinct (e.g., positive) kurtosis sign can be carried out by sweeping the contrast over pairs \mathbf{z}_{1j} only, with $\varepsilon_1=1,\,\varepsilon_j=-1,\,$ for $2\leq j\leq n$. After convergence, the desired source will appear at the first entry of the separator output vector.

IV. NUMERICAL EXPERIMENTS

The contrast is tested on synthetic random unitary mixtures of n=10 binary signals composed of 1000 samples. Sources kurtosis values of either $\kappa=2$ (super-Gaussian) or $\kappa=-2$ (sub-Gaussian) are obtained by setting the probability of the two states in the binary distribution accordingly [12]. The error

$$E = \frac{1}{2n(n-1)} \left[\sum_{i=1}^{n} \left(\sum_{j=1}^{n} \frac{|G_{ij}|}{\max_{k} |G_{ik}|} - 1 \right) + \sum_{j=1}^{n} \left(\sum_{i=1}^{n} \frac{|G_{ij}|}{\max_{k} |G_{kj}|} - 1 \right) \right]$$
(8)

is used as a separation performance criterion [13], [4]. The error is always positive, and zero if and only if matrix G is a trivial filter of the form (2). Error values are averaged over 250 independent realizations of the sources and the mixing matrix. Three contrasts are considered: 'COM2' [1] (\triangle marker); 'COM1+' and 'COM1-', which correspond to the contrast of [5] assuming that all sources have positive and negative kurtosis, respectively (+ and × markers, resp.); and function (3), which we refer to as 'kurtosis sign priors (KSP)' contrast (\bigcirc marker). For each tested contrast, we carry out $5(1 + |\sqrt{n}|)$ sweeps over all signal pairs.

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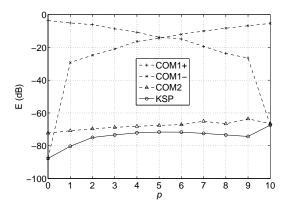


Fig. 1. Source separation performance of ICA contrasts as a function of the number of positive-kurtosis sources p. The KSP method employs the correct value of p.

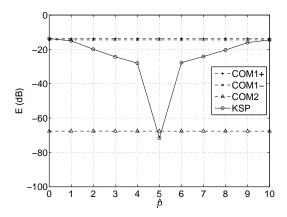


Fig. 2. Source separation performance of ICA contrasts as a function of the estimated number of positive-kurtosis sources \hat{p} . The correct value is p = 5.

Fig. 1 shows the performance variation as a function of the number p of sources with positive kurtosis, where p is assumed to be perfectly known a priori. As expected, COM1+ and COM1- fail to perform the separation except when all sources have the same kurtosis sign. KSP outperforms the other contrasts.

The robustness of contrast (3) to a mismatch in the prior information is analyzed in Fig. 2, where \hat{p} sources are assumed to have positive kurtosis while, actually, p=5. KSP's separation performance degrades as the available knowledge becomes less accurate.

Finally, we set p=1 and aim at the single source with positive kurtosis through the extraction procedure described at the end of Sec. III. Fig. 3 plots the average interference-to-signal ratio (ISR) for the estimation of the first source, defined as

$$ISR = 1 - \frac{|G_{11}|^2}{\sum_{j=1}^n |G_{1j}|^2}$$

as a function of the sweep number. This result illustrates the ability of the KSP contrast (3) to extract a source of known kurtosis sign from a mixture where all other sources have the opposite sign, without having to separate the whole mixture and resolve the permutation ambiguity after separation.

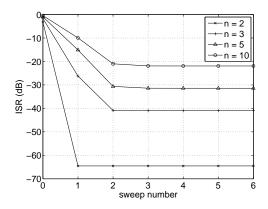


Fig. 3. Source extraction performance of the KSP contrast (3) for different mixture sizes.

V. CONCLUSIONS

An orthogonal contrast for ICA has been proposed which takes into account the non-Gaussian character of the source signals as measured by the sign of their fourth-order marginal cumulants (kurtosis). The contrast is linked to an approximate ML principle, and is able to separate the independent sources into two groups, depending on their kurtosis sign, thus partially solving the permutation ambiguity usually associated with ICA. The iterative pairwise maximization of the proposed contrast can be carried out at low complexity by closed-form solutions. As opposed to alternative fully blind techniques, the new contrast is particularly suited to the direct extraction of a source with known kurtosis sign distinct from the others'. The principle extends to higher-order cumulants other than kurtosis, as proved in the Appendix.

APPENDIX A

Proof of Proposition 1: The following proof generalizes the result of Proposition 1 to rth-order cumulants, with $r \geq 3$. Accordingly, in the sequel κ_i and μ_i denote the standardized rth-order cumulant of source s_i and output y_i , respectively, whereas $\varepsilon_i = \mathrm{sign}(\kappa_i)$.

By the multilinearity property of cumulants, we have $\mu_i = \sum_{j=1}^n G_{ij}^r \kappa_j$, where $G_{ij} = [\mathbf{G}]_{ij}$. Hence:

$$\Psi_p(\mathbf{y}) = \sum_{i=1}^n \varepsilon_i \sum_{j=1}^n G_{ij}^r \kappa_j.$$

The triangular inequality yields

$$\Psi_p(\mathbf{y}) \le \sum_{i=1}^n \sum_{j=1}^n |G_{ij}|^r |\kappa_j| \le \sum_{i=1}^n \sum_{j=1}^n |G_{ij}|^2 |\kappa_j|$$

where the right-hand side term stems from the fact that $r \geq 3$ and the orthonormality of matrix G, which can be expressed as $\sum_i |G_{ij}|^2 = 1$. Invoking again this property, we obtain

$$\Psi_p(\mathbf{y}) \le \sum_{j=1}^n |\kappa_j| = \sum_{j=1}^n \varepsilon_j \kappa_j = \Psi_p(\mathbf{s}).$$

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This proves the domination. Now if the equality $\Psi_p(\mathbf{y}) = \Psi_p(\mathbf{s})$ holds, we must have

$$\sum_{i=1}^{n} \sum_{j=1}^{n} \left[|G_{ij}|^2 - |G_{ij}|^r \right] |\kappa_j| = 0.$$

Yet all the terms in the sums are positive and thus they must all vanish. In other words, $|G_{ij}|^2 - |G_{ij}|^r = 0, \forall i, j$, with $r \geq 3$, which can occur only if $|G_{ij}| \in \{0,1\}$. Because \mathbf{G} is orthonormal, it must then have only one nonzero element in every row and column. Hence, \mathbf{G} is of the form (2), with $D_i = [\mathbf{D}]_{ii} = \pm 1$. This proves the discrimination property. Function $\Psi_p(\mathbf{y})$ is thus a contrast for ICA.

Proof of Proposition 2: This proof extends the validity of Proposition 2 to any even order $r \geq 4$. As seen above, equality $\Psi_p(\mathbf{y}) = \Psi_p(\mathbf{s})$ holds if and only if

$$\sum_{i=1}^{n} \varepsilon_i \sum_{j=1}^{n} G_{ij}^r \kappa_j = \sum_{j=1}^{n} \varepsilon_j \kappa_j.$$

Because $D_j = \pm 1$ and **P** is a permutation, we have that $G_{ij}^r = P_{ij}$, with $P_{ij} = [\mathbf{P}]_{ij}$, as r is even. Also, $\varepsilon_i^2 = 1$ and $\varepsilon_j \kappa_j = |\kappa_j|$, so that

$$\sum_{i=1}^{n} \left[1 - \sum_{i=1}^{n} \varepsilon_i P_{ij} \varepsilon_j \right] |\kappa_j| = 0 .$$

Yet, since all the terms in the sum are positive, they must individually vanish, yielding the relation

$$\sum_{i=1}^{n} \varepsilon_i P_{ij} \epsilon_j = 1, \quad \forall j.$$

Now, by splitting the sum into two parts, we are able to replace ε_i by its value, yielding $\sum_{i=1}^p P_{ij}\varepsilon_j - \sum_{i=p+1}^n P_{ij}\varepsilon_j = 1$. Let us distinguish between the cases $j \leq p$ and j > p, and take into account the fact that, for any permutation, $\sum_{i=1}^n P_{ij} = 1$. Then:

$$\begin{cases} 1 - 2 \sum_{i=p+1}^{n} P_{ij} = 1 & \forall j \leq p \\ 1 - 2 \sum_{i=1}^{p} P_{ij} = 1 & \forall j > p \end{cases}.$$

The first equality yields, for any $j \leq p$, $\sum_{i=p+1}^{n} P_{ij} = 0$. That is, by positivity, $P_{ij} = 0$. Thus, the $(n-p) \times p$ bottom left block of \mathbf{P} is null. Analogously, we see that for any j > p, $\sum_{i=1}^{p} P_{ij} = 0$, and thus the $p \times (n-p)$ top right block of \mathbf{P} must also be null. Consequently, the permutation matrix takes indeed the form (5).

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