# Iterative Re-weighted $L_1$ -Norm Principal-Component Analysis<sup>†</sup>

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Abstract-We consider the problem of principal-component analysis of a given set of data samples. When the data set contains faulty measurements/outliers, the performance of classic  $L_2$ principal-component analysis (L2-PCA) deteriorates drastically. Instead,  $L_1$  principal-component analysis ( $L_1$ -PCA) offers outlier resistance due to the  $L_1$ -norm maximization criterion it adopts to compute the principal subspace. In this work, we present an iterative re-weighted  $L_1$ -PCA method (IRW  $L_1$ -PCA) that generates a sequence of  $L_1$ -norm subspaces. In each iteration, the data set comformity of each sample is measured by the  $L_1$ subspace calculated in the previous iteration and used to weigh the data sample before the  $L_1$  subspace update. The approach automatically suppresses outliers in each iteration resulting in increasingly accurate subspace calculation. We provide convergence analysis and compare the proposed algorithm against benchmark algorithms in the literature. Experimental studies demonstrate the superiority of the proposed IRW  $L_1$ -PCA procedure.

Index Terms—Faulty data, feature extraction,  $L_1$ -norm, robust principal component analysis, eigenvector decomposition, outliers.

### I. INTRODUCTION

Principal component analysis (PCA) is a prevalent method for dimensionality reduction and low-rank subspace approximation. Conventional  $L_2$ -norm-based principal component analysis ( $L_2$ -PCA), however, is easily affected by "outlier" values that are numerically distant from the nominal lowrank signal. To deal with the problem of outliers in principalcomponent design there has been a growing interest in robust PCA methods [1]-[7]. In [1]-[4], subspace decomposition is performed under an  $L_1$ -error minimization criterion. In [5], non-negative matrix factorization is performed via Manhattan distance minimization (MahNMF), which robustly estimates the low-rank part and the sparse part of a non-negative matrix. The robust PCA method (RPCA) developed in [6] performs low-rank sparse decomposition by minimizing a weighted sum of the nuclear-norm of the low-rank component and the  $L_1$ -norm of the sparse component. The GoDec algorithm developed in [7] performs low-rank and sparse decomposition as well, by alternatingly solving for the low-rank and sparse components. An accelerated method is proposed in [7] via bilateral random projection (BRP).

Recently, there has been a growing documented effort to calculate robust subspaces by explicit  $L_1$  projection maximization [8]-[11]. The resulting principal components are called  $L_1$ principal components. The work in [8] presented a suboptimal iterative algorithm for the computation of one  $L_1$  principal component and [9] presented an iterative algorithm for suboptimal joint computation of  $d \ge 1$   $L_1$  principal components. In [10], for the first time in the literature, algorithms for exact calculation of  $L_1$  principal components are developed. Later, in [11] an approximate algorithm is developed for fast computation of the  $L_1$  principal components. The  $L_1$ -PCA method has been successfully applied to a wide range of research fields such as direction of arrival (DoA) estimation [12] and robust face recognition [13], [14]. Most recently, compressed-sensed-domain L1-PCA methods were developed for low-rank background scene and sparse foreground moving objects extraction from compressed-sensed surveillance video sequences [15], [16].

Nevertheless, existing  $L_1$ -PCA methods in [8]-[16] adopt "one-shot" processing. That is, for a given data set with potential outliers, the  $L_1$ -PCA algorithm is applied only once to compute from the explicit data the  $L_1$  subspace. For severely contaminated data sets, such one-shot  $L_1$  subspace computation can still be away from the true nominal signal subspace of interest.

In this paper, we propose an iterative re-weighted  $L_1$ -PCA method. Given a fixed data set that potentially contains outliers, the procedure iteratively generates a sequence of improved  $L_1$  subspaces. In each iteration, nominal compliance of each sample is inferred by its position relative to the  $L_1$ subspace calculated in the previous iteration and translated to a "weight." Samples with higher weights tend to be nominal samples and samples with lower weights are more likely to be

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the outliers. Weighted  $L_1$ -PCA calculation is then carried out in which the contribution of outlying samples in the data set is suppressed resulting in an improved  $L_1$ -subspace. The sample weights converge as the iteration number increases and the iterative algorithm terminates when the weights in the current and previous iteration are deemed close enough.

The remainder of this paper is organized as follows. In Section II, we introduce necessary background on regular  $L_1$ -PCA. In Section III, the proposed iterative re-weighted  $L_1$ -PCA algorithm is developed. In Section IV, experimental studies are provided to demonstrate the effectiveness of the proposed algorithm. Finally, a few conclusions are drawn in Section V.

## II. BACKGROUND ON $L_1$ -NORM PRINCIPAL-COMPONENT ANALYSIS

Consider N real-valued samples  $\mathbf{x}_1, \mathbf{x}_2, ..., \mathbf{x}_N$  of dimension D that form the  $D \times N$  data matrix

$$\mathbf{X} = [\mathbf{x}_1 \ \mathbf{x}_2 \ \cdots \mathbf{x}_N]. \tag{1}$$

In the common version of the low-rank approximation problem  $(L_2$ -PCA) one seeks to describe (approximate) the data matrix **X** by a rank-r product  $\mathbf{PQ}^T$  where  $\mathbf{P} \in \mathbb{R}^{D \times r}$ ,  $\mathbf{Q} \in \mathbb{R}^{N \times r}$ ,  $r \leq \min\{D, N\}$ . Given the observation data matrix **X**,  $L_2$ -PCA minimizes the sum of the element-wise squared error between the original matrix **X** and its rank-r representation  $\mathbf{PQ}^T$  in the form of Problem  $\mathcal{P}_1^{L_2}$  defined below,

$$\mathcal{P}_{1}^{L_{2}}: (\mathbf{P}_{L_{2}}, \mathbf{Q}_{L_{2}}) = \arg \min_{\substack{\mathbf{P} \in \mathbb{R}^{D \times r}, \mathbf{P}^{T} \mathbf{P} = \mathbf{I}_{r} \\ \mathbf{Q} \in \mathbb{R}^{N \times r}}} \|\mathbf{X} - \mathbf{P}\mathbf{Q}^{T}\|_{2}.$$
(2)

Problem  $\mathcal{P}_1^{L_2}$  is equivalent to

$$\mathcal{P}_{2}^{L_{2}}: \mathbf{P}_{L_{2}} = \arg \max_{\substack{\mathbf{P} \in \mathbb{R}^{D \times r} \\ \mathbf{P}^{T} \mathbf{P} = \mathbf{I}_{r}}} \| \mathbf{X}^{\mathrm{T}} \mathbf{P} \|_{2}$$
(3)

the solution of which is given by the r dominant singular-value left singular vectors of the original data matrix **X**.

By minimizing the sum of squared errors,  $L_2$  principalcomponent calculation becomes sensitive to extreme error value occurrences caused by the presence of outlying samples in the data matrix (samples that are numerically distant from the nominal data, appear only few times in the data matrix and are not to appear under normal system operation upon design). Motivated by this observed drawback of  $L_2$ -subspace signal processing, subspace decomposition approaches that are based on the  $L_1$  norm were proposed for robust low-rank subspace computation. Replacing the  $L_2$ -norm in Problem  $\mathcal{P}_2^{L_2}$  by  $L_1$ norm,  $L_1$ -PCA calculates principal components in the form of

$$\mathcal{P}^{L_1}: \mathbf{P}_{L_1} = \arg \max_{\substack{\mathbf{P} \in \mathbb{R}^{D \times r} \\ \mathbf{P}^T \mathbf{P} = \mathbf{L}}} \| \mathbf{X}^T \mathbf{P} \|_1.$$
(4)

 $\mathbf{P}_{L_1}$  in (4) is likely to be closer to the true nominal rank-r subspace than  $L_2$ -PCA. The r columns of  $\mathbf{P}_{L_1}$  in (4) are the so-called  $r L_1$  principal components that describe the rank-r subspace in which  $\mathbf{X}$  lies. As shown in [10], exact calculation of the  $L_1$  principal components in Problem  $\mathcal{P}^{L_1}$  can be recast

as a combinatorial problem. In short, when the rank of the nominal signal is r = 1, Problem  $\mathcal{P}^{L_1}$  reduces to

$$\mathbf{p}_{L_1} = \arg \max_{\substack{\mathbf{p} \in \mathbb{R}^D \\ \|\mathbf{p}\|_2 = 1}} \|\mathbf{X}^{\mathrm{T}} \mathbf{p}\|_1,$$
(5)

which can be reformulated as

$$\max_{\substack{\mathbf{p}\in\mathbb{R}^{D}\\\|\mathbf{p}\|_{2}=1}} \|\mathbf{X}^{T}\mathbf{p}\|_{1} = \max_{\substack{\mathbf{p}\in\mathbb{R}^{D}\\\|\mathbf{p}\|_{2}=1}} \max_{\substack{\mathbf{b}\in\{\pm1\}^{N}\\\|\mathbf{p}\|_{2}=1}} \mathbf{b}^{T}\mathbf{X}^{T}\mathbf{p}$$
(6)

$$= \max_{\mathbf{b} \in \{\pm 1\}^N} \max_{\substack{\mathbf{p} \in \mathbb{R}^D \\ \|\mathbf{p}\|_2 = 1}} \mathbf{p}^{\mathrm{T}} \mathbf{X} \mathbf{b}$$
(7)

$$= \max_{\mathbf{b} \in \{\pm 1\}^N} \| \mathbf{X} \mathbf{b} \|_2.$$
 (8)

The optimal solution for (8) can be obtained by exhaustive search in the space of the binary antipodal for example, vector **b** with complexity  $O(2^{N-1}DN)$ .

When the rank of the nominal data is r > 1, problem  $\mathcal{P}^{L_1}$  can be rewritten as [10]

$$\max_{\substack{\mathbf{P} \in \mathbb{R}^{D \times r} \\ \mathbf{P}^{\mathrm{T}} \mathbf{P} - \mathbf{L}}} \| \mathbf{X}^{\mathrm{T}} \mathbf{P} \|_{1}$$
(9)

$$= \max_{\substack{\mathbf{P} \in \mathbb{P}^{D \times r} \\ \mathbf{P}^{\mathrm{T}} \mathbf{P} - \mathbf{J}}} \max_{\mathbf{B} \in \{\pm 1\}^{N \times r}} \operatorname{tr}(\mathbf{P}^{\mathrm{T}} \mathbf{X} \mathbf{B})$$
(10)

$$= \max_{\mathbf{B} \in \{\pm 1\}^{N \times r}} \|\mathbf{X}\mathbf{B}\|_* \tag{11}$$

where  $\|\cdot\|_*$  stands for nuclear norm. To find exactly the optimal  $L_1$ -norm projection operator  $\mathbf{P}_{L_1}$  in (9) we can perform the following steps [10].

1) Solve (11) to obtain  $\mathbf{B}_{opt}$ .

2) Perform singular value decomposition (SVD) on  $\mathbf{XB}_{opt} = \mathbf{U} \mathbf{\Sigma} \mathbf{V}^{T}$ .

3) Return  $\mathbf{P}_{L_1} = \mathbf{U}_{:,1:r} \mathbf{V}^{\mathrm{T}}$ .

If we solve (11) by exhaustive search, the overall complexity of the above procedure for finding  $r L_1$  principal components will be  $\mathcal{O}(2^{Nr} \min\{D^2r, Dr^2\})$ . For any fixed data dimension D, a polynomial-time algorithm is developed in [10] to solve optimally (11) with complexity  $\mathcal{O}(N^{\operatorname{rank}(\mathbf{X})r-r+1})$ ,  $\operatorname{rank}(\mathbf{X}) \leq D$ . In [11], a fast approximation algorithm was proposed to solve (11) with complexity  $\mathcal{O}(\min\{ND^2, N^2D\} + N^2(r+2) + ND)$ .

## III. PROPOSED ITERATIVE RE-WEIGHTED $L_1$ PRINCIPAL COMPONENT ANALYSIS

The regular  $L_1$ -PCA problem in (4) seeks a rank-r subspace from the data matrix  $\mathbf{X} \in \mathbb{R}^{D \times N}$  by one-shot calculation. Although the adopted  $L_1$ -norm maximization is less affected by outliers compared to  $L_2$ -norm maximization in  $L_2$ -PCA in (3), the produced  $L_1$  subspace  $\mathbf{P}_{L_1}$  can still be away from the true nominal signal low-rank subspace. In this section, we propose an iterative method that generates a sequence of improved  $L_1$  subspaces for the same data matrix  $\mathbf{X}$ .

## A. Algorithm

We consider the calculation of r principal components  $\mathbf{P}_{L_1} \in \mathbb{R}^{D \times r}$ , D > r > 1. Initially, the direct  $L_1$  subspace is computed via (4) and denoted by  $\mathbf{P}_{L_1}^{(0)}$ . Next, the distance of each sample  $\mathbf{x}_n$  from subspace  $\mathbf{P}_{L_1}^{(0)}$  is defined as the  $L_2$  error between  $\mathbf{x}_n$  and its rank-r surrogate

$$d_n^{(1)} = \|\mathbf{x}_n - \mathbf{P}_{L_1}^{(0)} \mathbf{P}_{L_1}^{(0)^{\mathrm{T}}} \mathbf{x}_n\|_2, \quad n = 1, ..., N.$$
(12)

We expect large  $d_n^{(1)}$  if  $\mathbf{x}_n$  is an "outlier" and small  $d_n^{(1)}$  if  $\mathbf{x}_n$  is a nominal sample. Therefore, the nominal-likeness (weight) of each sample can be measured as the reciprocal of its  $L_2$  distance from the subspace, i.e.,

$$w_n^{(1)} = (d_n^{(1)})^{-1}, \quad n = 1, ..., N,$$
 (13)

followed by normalization,

$$\widetilde{w}_{n}^{(1)} = \frac{w_{n}^{(1)}}{\sum\limits_{n=1}^{N} w_{n}^{(1)}}, \quad n = 1, ..., N.$$
(14)

When computing the  $L_1$  subspace, data samples with larger nominal-likeness (weight) should contribute more and samples with smaller nominal-likeness (weight) should be suppressed such that the resulting calculated  $L_1$  subspace is more accurate. In this direction, we propose that each data sample  $\mathbf{x}_n$  is weighed by  $\tilde{w}_n^{(1)}$ . We form a weight matrix

$$\widetilde{\mathbf{W}}^{(1)} = \begin{bmatrix} \widetilde{w}_1^{(1)} & 0 & 0 & \cdots \\ 0 & \widetilde{w}_2^{(1)} & 0 & \cdots \\ \vdots & & \\ 0 & 0 & \cdots & \widetilde{w}_N^{(1)} \end{bmatrix}$$
(15)

and update the  $L_1$  subspace by

$$\mathbf{P}_{L_1}^{(1)} = \arg \max_{\substack{P \in \mathbb{R}^{D \times r} \\ \mathbf{P}^T \mathbf{P} = \mathbf{I}_r}} \| (\mathbf{X} \widetilde{\mathbf{W}}^{(1)})^{\mathrm{T}} \mathbf{P} \|_1.$$
(16)

Generalizing, in the (k + 1)th iteration new weights are computed using the  $L_1$  subspace  $\mathbf{P}_{L_1}^{(k)}$  computed at the kth iteration, i.e.

$$d_n^{(k+1)} = \|\mathbf{x}_n - \mathbf{P}_{L_1}^{(k)} \mathbf{P}_{L_1}^{(k)^{\mathrm{T}}} \mathbf{x}_n\|_2, 1 \le n \le N,$$
(17)  
$$w^{(k+1)} = (d^{(k+1)})^{-1}$$
(18)

$$w_n^{(k+1)} = (u_n^{(k+1)}),$$
 (18)  
 $\widetilde{w}_n^{(k+1)} = w_n^{(k+1)}$  (19)

$$\hat{w}_{n}^{(k+1)} = \frac{n}{\sum_{n=1}^{N} w_{n}^{(k+1)}},$$
(19)

$$\widetilde{\mathbf{W}}^{(k+1)} = \begin{bmatrix} \widetilde{w}_1^{(k+1)} & 0 & 0 & \cdots \\ 0 & \widetilde{w}_2^{(k+1)} & 0 & \cdots \\ \vdots & & \\ 0 & 0 & \cdots & \widetilde{w}_N^{(k+1)} \end{bmatrix}.$$
(20)

Subsequently, the  $L_1$  subspace at the (k + 1)th iteration is updated to

$$\mathbf{P}_{L_1}^{(k+1)} = \arg \max_{\substack{P \in \mathbb{R}^{D \times r} \\ \mathbf{P}^T \mathbf{P} = \mathbf{I}_r}} \| (\mathbf{X} \widetilde{\mathbf{W}}^{(k+1)})^{\mathrm{T}} \mathbf{P} \|_1.$$
(21)

#### B. Convergence Analysis

To guarantee a convergent weight sequence for practical algorithmic implementation, we modify the weight update formula as follows. In the (k+1)th iteration, we first compute the  $\ell_2$  error (distance) for each sample as in (17). Then, we define

$$u_n^{(k+1)} = (d_n^{(k+1)})^{-1}$$
(22)

and update the weight based on  $u_n^{(k+1)}$  by

$$w_n^{(k+1)} = \begin{cases} w_n^{(k)}(1-\beta^k), \text{ if } u_n^{(k+1)} < w_n^{(k)}(1-\beta^k), \\ u_n^{(k+1)}, \text{ if } w_n^{(k)}(1-\beta^k) \le u_n^{(k+1)} \le w_n^{(k)}(1+\beta^k) \\ w_n^{(k)}(1+\beta^k), \text{ if } u_n^{(k+1)} > w_n^{(k)}(1+\beta^k) \end{cases}$$

where  $0 < \beta < 1$  is a pre-defined parameter. Intuitively, we avoid updating the weights too aggressively by restricting the new weight  $w_n^{(k+1)}$  to be within a small neighborhood of the weight in the previous iteration  $w_n^{(k)}$ . The size of the neighborhood depends on  $\beta$ . Subsequently,  $w_n^{(k+1)}$  is normalized as in (19), followed by weight matrix construction in (20). The convergence of the weight sequence can be verified by

$$\lim_{k \to \infty} \beta^k = 0, \tag{23}$$

$$\lim_{k \to \infty} (w_n^{(k+1)} - w_n^{(k)}) = 0,$$
 (24)

$$\lim_{k \to \infty} (\widetilde{w}_n^{(k+1)} - \widetilde{w}_n^{(k)}) = 0.$$
<sup>(25)</sup>

### C. Stopping Criterion

In implementing the proposed iterative algorithm, we exit the algorithm when the difference between the weight vectors at the kth and (k + 1)th iteration is smaller than a predefined threshold  $\epsilon > 0$ , that is,

$$\|\mathbf{w}^{(k+1)} - \mathbf{w}^{(k)}\| < \epsilon, \tag{26}$$

where  $\mathbf{w}^{(k)} = [w_1^{(k)}, w_2^{(k)}, \cdots, w_N^{(k)}]^T$  and  $\mathbf{w}^{(k+1)} = [w_1^{(k+1)}, w_2^{(k+1)}, \cdots, w_N^{(k+1)}]^T$ .

#### IV. APPLICATIONS AND EXPERIMENTAL STUDIES

In this section, we assess the effectiveness of the proposed iterative re-weighted  $L_1$ -PCA (IRW  $L_1$ -PCA) algorithm through two experiments: (i) Dimensionality reduction of a 2-dimensional Gaussian data set (artificial data), and (ii) video surveillance (field data foreground extraction).

### A. Dimensionality Reduction

We generate a nominal data set  $\mathbf{X}_{D\times N}$  of N = 30two-dimensional (D = 2) observation points drawn from the Gaussian distribution  $\mathcal{N}\left(\mathbf{0}_2, \begin{bmatrix} 10.5 & 13\\ 13 & 30 \end{bmatrix}\right)$  as shown in Fig. 1. We assume that our data matrix is corrupted by four additional outlier measurements,  $\mathbf{o}_1, \mathbf{o}_2, \mathbf{o}_3, \mathbf{o}_4$ , shown in the bottom right corner of Fig. 1. For the corrupted data matrix  $\mathbf{X}^{\text{CRPT}} = [\mathbf{X}, \mathbf{o}_1, \mathbf{o}_2, \mathbf{o}_3, \mathbf{o}_4]$ , we calculate and plot in Fig. 1 the rank-1 subspace by  $L_2$ -PCA, regular  $L_1$ -PCA, as well as the proposed IRW  $L_1$ -PCA method with number of

iterations k = 1, 4, 8, 15, 21 ( $\beta = 0.9$ ). For reference purposes, we also plot the true nominal-data maximum-variance direction, i.e., the dominant eigenvector of the covariance matrix 10.5 13 . We observe that the proposed IRW  $L_1$ -PCA 1330 approach offers better estimation of the principal component than the  $L_2$  and regular  $L_1$ -PCA methods [10]. As the number of iterations increases from k = 1 to k = 21, the IRW  $L_1$  principal component comes closer to the true rank-1 subspace. The algorithm converges empirically at k = 21. To quantify the impact of the outliers, we generate 1000 new independent evaluation data points from  $\mathcal{N}\left(\mathbf{0}_{2}, \begin{bmatrix} 10.5 & 13\\ 13 & 30 \end{bmatrix}\right)$ . In Fig. 2, we estimate the mean-square-fit-error (MSFE)  $\mathbf{E}\{\|\mathbf{x} - \mathbf{p}\mathbf{p}^{\mathrm{T}}\mathbf{x}\|_{2}^{2}\}$ 1000 by  $\frac{1}{1000} \sum_{i=1}^{1000} \|\mathbf{x}_i - \mathbf{p}\mathbf{p}^{\mathrm{T}}\mathbf{x}_i\|_2^2$  for  $\mathbf{p}_{L_2}(\mathbf{X}^{\mathrm{CRPT}})$ ,  $\mathbf{p}_{L_1}(\mathbf{X}^{\mathrm{CRPT}})$ , and IRW  $\mathbf{p}_{L_1}(\mathbf{X}^{\mathrm{CRPT}})$ . Again, for reference purposes, we plot the MSFE curve for the true nominal-data maximum-variance direction. We observe that the MSFE value of the proposed IRW  $L_1$  component decreases rapidly as the iteration number increases and converges toward the minimum at k = 21.



Fig. 1. Training data matrix  $\mathbf{X}_{2\times 30}$  corrupted by four outlier points in bottom right with calculated rank-1  $L_2$ , regular  $L_1$ , and IRW  $L_1$ principal components at iteration k = 1, 4, 8, 15, and 21.

### B. Video Surveillance

Consider a sequence of surveillance video frames  $\mathbf{X}_t \in \mathbb{R}^{m \times n}$  with frame resolution of  $m \times n$  pixels and time index t = 1, ..., N. For a surveillance video sequence, the background scene is usually static and the objective is to extract foreground moving objects. In our experiment, we perform block-by-block IRW  $L_1$ -PCA for low-rank background modeling and foreground extraction. We divide each frame  $\mathbf{X}_t$  into J blocks  $\mathbf{X}_t^j \in \mathbb{R}^{m_b \times n_b}$ , j = 1, ..., J. We let  $\mathbf{x}_t^j \in \mathbb{R}^D$ ,  $D = m_b n_b$ , represent vectorization of  $\mathbf{X}_t^j$  via column concatenation. For each sequence of co-located blocks,  $\mathbf{x}_t^j$ , t = 1, ..., N, it is likely that the moving objects appear only in a few of these blocks, therefore we can model the



Fig. 2. Mean square-fit-error of  $L_2$ , regular  $L_1$ , and IRW  $L_1$  principal components calculated from corrupted training data set  $\mathbf{X}^{CRPT}$  versus iteration index k.

static background scene as a low-rank component  $\mathbf{z}_t^j$  and the foreground moving objects as an outlying component  $\mathbf{s}_t^j$ . That is,

$$\mathbf{x}_{t}^{j} = \mathbf{z}_{t}^{j} + \mathbf{s}_{t}^{j}, \ t = 1, ..., N.$$
 (27)

In matrix form representation of the *j*th block across N frames,  $\mathbf{X}^{j} \triangleq [\mathbf{x}_{1}^{j}, ..., \mathbf{x}_{N}^{j}] \in \mathbb{R}^{D \times N}$  and

$$\mathbf{X}^j = \mathbf{Z}^j + \mathbf{S}^j. \tag{28}$$

To extract the low-rank background information, we carry out IRW  $L_1$ -PCA on  $\mathbf{X}^j$  and obtain the rank-1  $L_1$  subspace  $\mathbf{p}_{L_1}^j$  at convergence. Afterwards, the background blocks can be approximated by  $\widehat{\mathbf{Z}}^j = \mathbf{p}_{L_1}^j \mathbf{p}_{L_1}^{j^T} \mathbf{X}^j$  and the foreground blocks can be extracted as  $\widehat{\mathbf{S}}^j = \mathbf{X}^j - \widehat{\mathbf{Z}}^j$ , j = 1, ..., J.

We test the method on the *Airport* video sequence with 96 frames, each of  $144 \times 176$  pixels. We process N = 8 successive frames at a time. To mitigate the "blockiness" artifact, we divide each frame into J = 370 overlapping blocks of size  $26 \times 32$  and apply the proposed IRW  $L_1$ -PCA method independently to each group of co-located blocks across 8 frames. The final background and foreground scenes are obtained by averaging the extracted background pixels (as well as the foreground pixels) for which multiple results are available.

Fig. 3 displays the background and foreground extracted at multiple distinct time slots t = 5, 7, 66, 67 with r = 1 principal component by the proposed IRW  $L_1$ -PCA, regular  $L_1$ -PCA [10], and the robust PCA method of [6]. The results show that both the regular  $L_1$ -PCA and robust PCA method suffer from severe "ghost" presence in the estimated background scene, which results in problematic foreground extraction. In contrast, IRW  $L_1$ -PCA significantly mitigates the "ghost" effect in the estimated background and offers a much clearer foreground scene.



Fig. 3. Airport sequence: Original frame [row (i)] of time slot t = 5, 7, 66, and 67; proposed IRW  $L_1$ -PCA reconstructed background and moving objects [rows (ii) and (iii)]; robust PCA [6] reconstructed background and moving objects [rows (iv) and (v)]; regular  $L_1$ -PCA [10] reconstructed background and moving objects [rows (vi) and (vii)].

#### V. CONCLUSION

In this work, we proposed an iterative re-weighted  $L_1$  principal-component analysis algorithm to compute principal subspaces from data sets that may contain outliers. Instead of computing a "one-shot"  $L_1$  subspace, the proposed procedure iteratively computes a sequence of  $L_1$  subspaces. In every iteration, each data sample is weighed according to compliance to nominal data behavior measured by the  $L_1$  subspace computed in the previous iteration. We evaluated the effectiveness of the proposed IRW  $L_1$ -PCA method experimentally and the results showed significantly better performance than regular  $L_1$ -PCA and state-of-the-art robust PCA methods.

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