Error Concealment using 3D Low-Rank Tensor Completion

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1 Introduction

Error concealment is a field of signal processing which deals with corruption and loss of data and aims at reducing the deterioration of the signal.

As data are typically packetised for transmission over a network or storage on a hardware, any data loss or corruption usually results in errors in the form of missing blocks.

This report presents an error concealment algorithm for images and videos. This method reconstructs the corrupted blocks by exploiting the low-rank property of a stack of similar patches. Results obtained by application of this algorithm to movie reconstruction and inpainting are presented here, as well as a comparison with results from another low-rank tensor approximation method.

2 Basics of tensor algebra

Before looking at the error concealment algorithm in details, a few basic tools of tensor algebra need to be defined. This section is based on the paper by Kolda and Bader [3], which should be referred back to for further details.

It is sufficient for our purpose to consider three-dimensional tensors only, but all algebraic concepts presented in this section can be generalised to higher dimensions.

Let $\mathcal{X} \in \mathbb{R}^{I_1 \times I_2 \times I_3}$. The <u>i</u>th matricisation of \mathcal{X} is a reordering of its entries into a matrix $X_{(i)} \in \mathbb{R}^{I_i \times \prod_{j \neq i} I_j}$ such that the indices of \mathcal{X} in the ith dimension become the row indices of $X_{(i)}$ and the indices of \mathcal{X} in the other dimensions are rearranged in lexicographical order to become column indices of X_i .

The i-rank of \mathcal{X} is defined as the rank of its i^{th} matricisation. The <u>multilinear rank</u> of \mathcal{X} is a 3-tuple grouping the i-rank of \mathcal{X} in each of the three dimensions:

$$\operatorname{rank}(\mathcal{X}) := \left[\operatorname{rank}(X_{(1)}), \operatorname{rank}(X_{(2)}), \operatorname{rank}(X_{(3)})\right]$$

The <u>i</u>th-mode product of \mathcal{X} with a matrix $M \in \mathbb{R}^{m \times I_i}$ is a tensor $\mathcal{Y} \in \mathbb{R}^{J_1 \times J_2 \times J_3}$, with $J_i = \overline{I_i}$ for $i \neq i$, and $J_i = m$. \mathcal{Y} is defined by means of the matricisation as follows:

$$\mathcal{Y} = \mathcal{X} \times_i M \iff Y_{(i)} = MX_{(i)}$$

Note that the matricisation operation is a bijection from $\mathbb{R}^{I_1 \times I_2 \times I_3}$ into $\mathbb{R}^{I_i \times \prod_{j \neq i} I_j}$. \mathcal{Y} is thus uniquely determined by any of its matricisation.

The inner product of two tensors $\mathcal{X}, \mathcal{Y} \in \mathbb{R}^{I_1 \times I_2 \times I_3}$ is the sum of the product of their entries, that is to say:

$$<\mathcal{X},\mathcal{Y}>:=\sqrt{\sum_{i_1=1}^{I_1}\sum_{i_2=1}^{I_2}\sum_{i_3=1}^{I_3}x_{i_1,i_2,i_3}y_{i_1,i_2,i_3}}$$
 (1)

The induced <u>norm</u> $\|\mathcal{X}\| := \sqrt{\langle \mathcal{X}, \mathcal{X} \rangle}$ is thus a generalisation of the matrix Frobenius norm to 3 dimensions.

Any tensor \mathcal{X} of multilinear rank $[r_1, r_2, r_3]$ can be decomposed into a core tensor $\mathcal{C} \in \mathbb{R}^{r_1 \times r_2 \times r_3}$ and three orthogonal basis matrices $A_i \in \mathbb{R}^{I_i \times r_i}$, such that

$$\mathcal{X} = \mathcal{C} \times_1 A_1 \times_2 A_2 \times_3 A_3 \tag{2}$$

This is called the <u>Tucker Decomposition</u> and is the basis of the High-Order Singular Value Decomposition (HOSVD) algorithm.

Note that a Tucker decomposition is not unique. Indeed, if \mathcal{X} can be decomposed as in Equation 2 then for any invertible $V_1 \in \mathbb{R}^{r_1 \times r_1}, V_2 \in \mathbb{R}^{r_2 \times r_2}, V_3 \in \mathbb{R}^{r_3 \times r_3}$, the following decomposition holds as well:

$$\mathcal{X} = \tilde{\mathcal{C}} \times_1 \tilde{A}_1 \times_2 \tilde{A}_2 \times_3 \tilde{A}_3$$

where $\tilde{\mathcal{C}} = \mathcal{C} \times_1 V_1 \times_2 V_2 \times_3 V_3$ and $\tilde{A}_i = A_i V_i^{-1}$.

HOSVD aims at computing a rank- $[R_1, R_2, R_3]$ approximation of a tensor \mathcal{X} of higher rank in the Tucker format. This is done by matricising \mathcal{X} in each dimension i = 1, 2, 3 and using the R_i principal left singular vectors as basis A_i of the Tucker decomposition. The pseudo-code for the HOSVD procedure is given in Algorithm 1.

Algorithm 1 Higher-Order Singular Value Decomposition

- 1: **procedure** HOSVD $(\mathcal{X}, R_1, R_2, R_3)$
- 2: **for** i = 1, 2, 3 **do**
- 3: $A_i \leftarrow R_i$ leading left singular vectors of $X_{(i)}$
- 4: $\mathcal{C} \leftarrow \mathcal{X} \times_1 A_1^{\top} \times_2 A_2^{\top} \times_3 A_3^{\top}$
- 5: **return** C, A_1, A_2, A_3

This procedure can be viewed as a generalisation of the matrix Singular Value Decomposition (SVD) as a truncated SVD is performed on each matricisation of \mathcal{X} to obtain a lower rank approximation. However, in contrast to the 2-dimensional case, the HOSVD does not lead to the best rank- $[R_1, R_2, R_3]$ approximation of \mathcal{X} . Let us denote the subspace of all 3-dimensional tensors of rank $r = [R_1, R_2, R_3]$ by $\mathcal{M}_r := \{\mathcal{T} \in \mathbb{R}^{I_1 \times I_2 \times I_3} \mid \operatorname{rank}(\mathcal{T}) = [R_1, R_2, R_3]\}$ and let $P_{\mathcal{M}_r}\mathcal{X}$ be a best approximation of \mathcal{X} in the subspace \mathcal{M}_r relatively to the norm defined above. If $P_{HO}\mathcal{X}$ represents the result of the HOSVD procedure applied to \mathcal{X} then the following results holds [2]:

$$\|\mathcal{X} - P_{HO}\mathcal{X}\| \leqslant \sqrt{3}\|\mathcal{X} - P_{\mathcal{M}_r}\mathcal{X}\|$$

This means that a best low-rank approximation of \mathcal{X} is only better by a factor $\frac{1}{\sqrt{3}}$ than what is computed by means of the HOSVD procedure.

3 Statement of the problem

The description is made here for the case of movie reconstruction. Image inpainting can be viewed as a particular case of movie reconstruction where the number of frames is equal to 1

Firstly, a Macro-Block (MB) grid is defined for each frame. The size of a MB, denoted by N, is typically 8×8 or 16×16 pixels and the MB grid is shifted by half a block size in the

vertical and horizontal direction of the frame. A given MB therefore typically overlaps with 8 other MBs from the same frame (see Figure 1) and a natural division of each MB into 4 sub-blocks arises.



Figure 1: A given MB (grey) typically overlaps with 8 other MBs (red, yellow, blue and green contour) from the same frame.

3.1 Tensor construction via selection of similar patches

In each frame of the movie separately, all MBs containing corrupted entries are put in a list and sorted relatively to the number of clean (i.e. non-corrupted) pixels they contain. We call P^0 a corrupted MB and Ω the set of its clean and concealed pixels. P^0 s with a larger set Ω will be reconstructed first.

Once a P^0 is chosen for reconstruction of its missing entries, similar MBs that are fully clean will be looked for in a some defined reference frames. We call these similar MBs P^i and seek to select the K-1 P^i s which best match P^0 . The best matching P^i is defined as the one which minimises the error

$$||P_{\Omega}^{0} - P_{\Omega}^{i}||_{1} = \sum_{j} |P_{j}^{0} - P_{j}^{i}|$$
(3)

where j is an index that loops over all pixels of Ω . All MBs are rescaled to have the same L_2 -norm as P^0 (on the set Ω only) before the matching error is computed.

Note that K needs to be defined as a user input, but in case less than K-1 "good matching" MBs are actually available in the reference frames, then only $K_{eff} - 1 < K - 1$ MBs are effectively selected, in order to avoid taking into account random MBs which would pervert the results. In my particular implementation, any MB with an error more then twice bigger than the best matching P^i was discarded from the selection.

It is important here to realise that, generally, the higher the number of reference frames for the selection of P^i s, the better the matching with P^0 and thus the better the reconstruction. However, the scanning of reference frames for detection of matching MBs is a very time-consuming process. The number of reference frames should therefore be a trade-off between computation time and quality of the results.

 P^0 and all the P^i s are then gathered into a 3-dimensional tensor \mathcal{X} of size $N \times N \times K_{eff}$, where P^0 is the first slice of the tensor. Later in this report, K will be used in place of K_{eff} for the sake of simplicity, but it should be remembered that this K might be smaller than the value given as user input.

3.2 Low-rank property of the tensor

Ideally, if the objects in the movie were not moving and the camera was keeping the same view, then \mathcal{X} would be composed of K times the same patch. It could therefore be represented

as a product between a 2D patch P and a constant vector 1 in the 3^{rd} dimension. Allowing some small sparse noise \mathcal{E} , it would write:

$$\mathcal{X} = \mathcal{X}_l + \mathcal{E} = P \times_3 1 + \mathcal{E} \tag{4}$$

After decomposition of \mathcal{X}_l in the Tucker format

$$\mathcal{X}_l = \mathcal{C} \times_1 A_1 \times_2 A_2 \times_3 A_3$$

and comparison with Equation 4 we can write

$$P = \mathcal{C} \times_1 A_1 \times_2 A_2$$
 and $A_3 = 1$

In practice, \mathcal{X}_l is not composed of K exactly identical patches and its rank can thus not be 1. It is however reasonable to assume that it still has a small 3-rank. In this light, two different algorithms have been used to find a good low-rank approximation of \mathcal{X} .

4 Description of the algorithms for low-rank approximation

4.1 Alternating Least Squares Algorithm (ALS)

We call now $\bar{\Omega}$ the set of missing entries in P^0 .

The Alternating Least Square algorithm presented in the paper by Nguyen, Dao and Tran [1] starts by filling $P_{\bar{\Omega}}^0$ in the first slice of \mathcal{X} with the mean values of the P^i s at these locations (see Algorithm 2, line 2).

Algorithm 2 Alternating Least Square

```
1: Form \mathcal{X} from P^0, ..., K-1

2: (\mathcal{X}(:,:,1))_{\bar{\Omega}} = (\frac{1}{K-1} \sum_{i=1}^{K-1} P^i)_{\bar{\Omega}}

3: Choose mode ranks [R_1, R_2, R_3] and tolerance \sigma_{rel}, \sigma_{it}

4: Initialise A_1 \in \mathbb{R}^{N \times R_1}, A_2 \in \mathbb{R}^{N \times R_2}, A_3 \in \mathbb{R}^{K \times R_3} randomly.

5: A^3(:,1) = [1, ..., 1]^{\top}/K

6: for i = 1, 2, 3 do

7: \mathcal{Y} = \mathcal{X}

8: for j \neq i do

9: \mathcal{Y} = \mathcal{Y} \times_j A_j^{\top}

10: Y_i \leftarrow \text{unfold } \mathcal{Y} \text{ in mode } i

11: A_i = \text{first } R_i \text{ principal components of } Y_i

12: \mathcal{C} = \mathcal{X} \times_1 A_1^{\top} \times_2 A_2^{\top} \times_3 A_3^{\top}

13: \mathcal{X}_l = \mathcal{C} \times_1 A_1 \times_2 A_2 \times_3 A_3

14: If convergence is reached, stop; else return to Step 5

15: Recover missing area in P^0: (P^0)_{\bar{\Omega}} = (\mathcal{X}_l(:,:,1))_{\bar{\Omega}}
```

While the multilinear rank R_3 is supposed to be small (it was set to 3 in the following examples), no assumption can be made on R_1 and R_2 . These two variables are thus generally kept to their maximal possible value N.

The ALS procedure is a modified version of the HOSVD presented in Algorithm 1. It is an iterative process which updates the basis matrices A_i one by one, keeping the other two

constant. In this way, A_3 can be enforced at each iteration to fulfil an additional condition. Indeed, in the ideal case described in subsection 3.2, A_3 was a constant vector 1. Since we look for a solution which is known to be close to the ideal case, convergence can be helped by forcing the first vector of A_3 to be constant and normalised to 1.

The procedure is considered to have reached convergence at iteration n if one of these two conditions is fulfilled:

1. The low-rank tensor $\mathcal{X}_{l}^{(n)}$ has become close enough to \mathcal{X} , i.e.

$$\epsilon_{rel}^{(n)} := \|\mathcal{X}_l^{(n)} - \mathcal{X}\| / \|\mathcal{X}\| < \sigma_{rel}$$

2. The change made during in 1 iteration has become negligible, i.e.

$$|\epsilon_{rel}^{(n)} - \epsilon_{rel}^{(n-1)}| < \sigma_{it}$$

4.2 GeomCG

The second algorithm, GeomCG, is the one presented in the paper written by Kressner, Steinlechner, Vandereycken [2]. Their implementation of the algorithm is available on their website [4] and is the one which was used here.

The aim of GeomCG is to minimise the cost function

$$f_{\mathcal{X}}(\mathcal{Y}) := \|P_{\Omega}\mathcal{Y} - P_{\Omega}\mathcal{X}\|^2$$

under the constraint $\mathcal{Y} \in \mathcal{M}_r$.

The algorithm is based on the fact that \mathcal{M}_r is a smooth subspace of $\mathbb{R}^{I_1 \times I_2 \times I_3}$ and that, together with the inner product of Equation 1, it forms a Riemannian manifold. An extension of the non-linear Conjugate Gradient method is used to minimise the cost function $f_{\mathcal{X}}(\mathcal{Y}_k)$. The direction of optimisation and the step size are computed by projecting the euclidian gradient of $f_{\mathcal{X}}(\mathcal{Y}_k)$ into the tangent plan of \mathcal{M}_r at position \mathcal{X} . After each optimisation the resulting tensor is retracted back onto \mathcal{M}_r , which ensures that the low-rank condition on \mathcal{Y} remains satisfied.

5 Results

In the following, p denotes the corruption ratio, i.e. the proportion of corrupted pixels in every frame. The corruption of sub-blocks was done randomly in each frame separately.

The original bus movie was downloaded from [5] in CIF format and was preprocessed so as to reduce the size of the problem and the computation time. The number of pixels in every frame was reduced by a factor 16 using cubic interpolation, and then cropped so as to keep only 151×196 pixels per frame.

5.1 Reconstruction of the Bus Sequence

Firstly, the bus movie was corrupted by removing sub-blocks of size $n \times n = N/2 \times N/2 = 4 \times 4$ with a corruption ratio p = 15%. The 100 first frames were then reconstructed by using ALS with the following parameters:

• MB size: N = 8

• Rank of \mathcal{X}_l tensor: r = [N, N, 3]

• Number of reference frames: $2 \cdot 5 + 1 = 11$ (the frame being reconstructed itself + 5 nearest previous frames + 5 nearest following frames)

Figure 2 shows the result obtained in the first frame. See how most of the reconstructed patches fit well in the image. The sequence of all 100 reconstructed frames can be viewed in the file recoveredBusN8.avi in annex.





Figure 2: Left: 1^{st} frame of the corrupted sequence. Right: 1^{st} frame of the reconstructed sequence.

All further results were obtained with a MB size N=16 to reduce the computation time.

5.1.1 Importance of initial guess

In order to test the importance of the initialisation of $P_{\bar{\Omega}}^0$ (line 2 of Algorithm 2), the same computations were carried on the bus sequence, but, this time, with a random initial guess rather than the average over similar MBs. Typical results of the reconstruction in both cases are displayed in Figure 3 and Figure 4.

Notice on Figure 3 that the MB reconstructed by means of low-rank approximation is only slightly better than the simple average computed initially. The reconstructed MB is smoother but remains very close to the initial guess.

Figure 4 shows that the quality of the result depends strongly on the initialisation. This is due to the fact that ALS only computes a local optimisation around the initial guess.

By selecting a higher number of P^i in the tensor building step (bigger K), the effect of the initial guess should decrease. This would however probably lead to the selection of poor matches and degrade the quality of the concealment.

It should be noticed that in both cases the error stagnates after one iteration of ALS only.

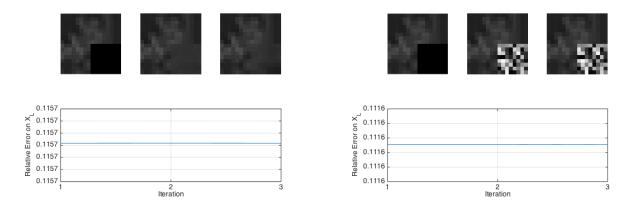


Figure 3: Corrupted MB (left), initialisation of $P_{\bar{\Omega}}^0$ using educated guess (middle), reconstructed MB (right)

Figure 4: Corrupted MB (left), random initialisation of $P_{\bar{\Omega}}^0$ (middle), poorly reconstructed MB (right)

5.1.2 Effect of the corruption ratio

The effect of p on the quality of the results was then investigated. Observe on Figure 5 how the reconstruction of the 15^{th} frame becomes relatively bad for $p \ge 35\%$. This is due to the fact that it becomes harder to find clean similar patches in the set of reference frames when p is higher. The tensor \mathcal{X} thus has to be formed from MBs that do not fit well with the MB being reconstructed.

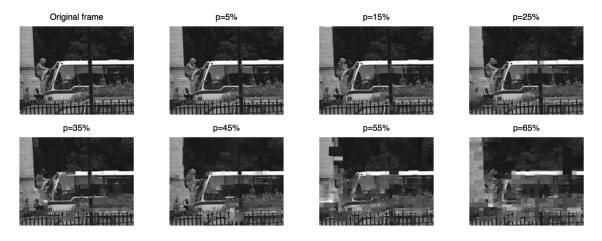


Figure 5: The quality of the reconstruction depends on the percentage of corrupted pixels

In each frame, every reconstructed MB was compared with the original one, and the relative error in Frobenius norm was computed. Figure 6 shows the average of this error over each frame for different values of p. Even though the error is not constant over different frames, it can be noticed that generally the smaller p, the smaller the error. Moreover the rate of growth of the error seems to increase with p.

In order to estimate the strength of the error growth, the values displayed in Figure 6 were averaged over the 30 frames for each p. Figure 7 shows that the rate of growth of the error is faster than exponential since the curve does not form a straight line in semilogarithmic

scale.

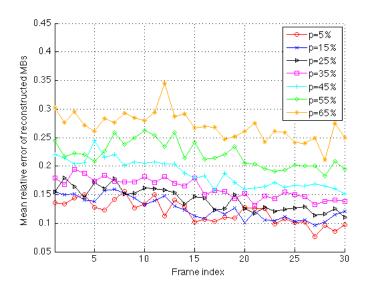
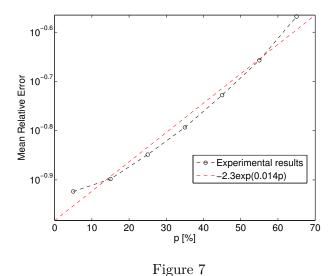


Figure 6



5.1.3 Reconstruction with GeomCG

The whole procedure was kept the same as with ALS, except for the low-rank approximation of the \mathcal{X} tensor, which was performed with GeomCG. As initial guess, GeomCG was provided with the result for \mathcal{X}_l given at line 13 in Algorithm 2 after a single iteration.

It was noticed that the GeomCG algorithm could not provide acceptable results when the mode-3 rank R_3 was set too high. The values of the reconstructed pixels were then out of range.

For this reason, the computations with GeomCG, as well as with ALS, were done with the following multilinear rank: $r = [R_1, R_2, R_3] = [N, N, 1]$.

See on Figure 8 that the results returned by both algorithm are equivalent. This can be explained by the fact that ALS converges in a single iteration, and that this result was provided to GeomCG as initial guess. GeomCG therefore does not have much to optimise from the result of ALS.

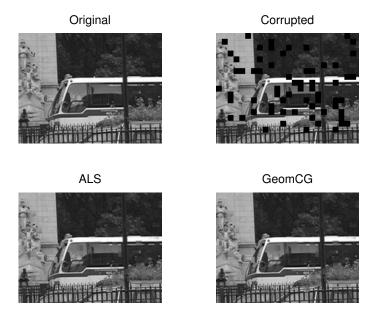


Figure 8: Same results are obtained with ALS and GeomCG (here frame 20)

To be convinced that the results were actually the same, the error has been plotted in Figure 9.

Moreover, the error of the reconstruction computed with ALS and $R_3 = 3$ was compared to both ALS and GeomCG with $R_3 = 1$. Note that all results are the same here.

This means that a rank-1 approximation of \mathcal{X} already captures most of its structure. The average of the three first singular values $\sigma_1, \sigma_2, \sigma_3$ of $\mathcal{X}_{(3)}$ (where, in the first slice of \mathcal{X} , $P_{\bar{\Omega}}^0$ was replaced by the pixels of the original image) was computed over all tensors \mathcal{X} . This lead to the following results:

$$\sigma_1 = 18.1367, \sigma_2 = 1.7986, \sigma_3 = 1.2682$$

and thus

$$\frac{\sigma_2}{\sigma_1} = 9.9\%$$
 and $\frac{\sigma_3}{\sigma_1} = 7\%$

Therefore, even though the 2^{nd} and 3^{rd} singular values of $\mathcal{X}_{(3)}$ are not negligible in comparison to σ_1 , the results of ALS do not seem to improve significantly when R_3 is set to 3.

Finally GeomCG was provided, as a test, with a random initial guess instead of the result returned by one iteration of ALS. The result of a typical frame can be seen in Figure 10. Observe how the reconstruction of some MBs is completely out of range. By decreasing R_1

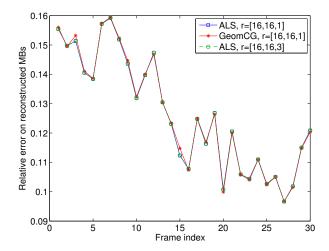


Figure 9: The error is the same with ALS and GeomCG

and R_2 , the number of wrongly reconstructed MBs decreases as well. However, even with $R_1 = R_2 = N/2$, $R_3 = 1$, most frames reconstructed with GeomCG still contained at least one MB that was out of range. Recall that decreasing R_1 and R_2 forces the reconstructed MB to have a lower-rank, even though the original MB might have been full-rank. Too low R_1 and R_3 therefore damage the quality of the reconstruction.



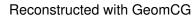




Figure 10: GeomCG returns poor results when provided with a random initial guess (here frame 20)

5.2 Inpainting

The images used for this inpainting experiment were downloaded from [6] and [7]. The number of pixels was reduced by spatial interpolation for the sake of rapidity.

In both cases (see Figure 11 and Figure 12), a portion of the image was deleted by hand, and then reconstructed with a MB size N=8.

The aim with the first picture was to test the ability of the ALS algorithm to reconstruct a regular texture. It can be observed that the horizontal stripes are very well recovered as a prolongation of the stripes at the boundary. In the opposite, internal vertical white stripes which are not a prolongation from anything at the boundary are not recovered at all. More precisely, instead of reconstructing clear vertical lines at defined locations, the algorithm

distributes ghosts of these white lines in the whole reconstructed portion. This tends to make the reconstruction slightly lighter than the original image.







Figure 11: Left: Original image. Middle: Deletion of a portion of texture. Right: Reconstruction of the texture

The aim with the picture of the snowy park was to make the bank disappear by deleting all of its pixels and reconstructing them from the rest of the image. It should be noticed in Figure 12 that the right side of the reconstructed portion is lighter than the left part, and that the transition between both parts is relatively abrupt. This is an illustration of the fact that the sub-blocks are reconstructed from the boundary towards the center. The reconstruction from each side is independent from the other side, and the matching of different parts as they join can only be done on a short range (size of a MB). This can lead to relatively big gradients in the reconstructed image, when the set of contiguous fully corrupted MB is too wide.







Figure 12: Left: Original image. Middle: Deletion of the bank. Right: Reconstruction without the bank.

6 Conclusion and suggestion for further research

The ALS algorithm which has been implemented for this report shows some successful results of movie reconstruction and inpainting.

The final reconstruction depends strongly on the selection of similar MBs in the set of reference frames. This selection step is actually the bottleneck of the algorithm in term of computation time. It would therefore by interesting to find a way to optimise the search of matheing MBs.

Finally, the size of the MBs is a parameter which is left free for the user to tune. It is however often not clear what value would be optimal for a given problem, and the smaller N does not always mean the better the reconstruction. It would therefore be useful to characterise the optimal N depending on the image resolution or the shape of the corrupted regions.

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