

MONTE CARLO OPTIMIZATION OF NON-LOCAL MEANS DENOISING ALGORITHM

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ABSTRACT. Non-Local Means is a powerful algorithm for image denoising, yet its high computational complexity limits real-time application. This paper replicates the Monte Carlo Non-Local Means Optimization, which utilizes random sampling to accelerate weight calculations without significantly degrading peak signal-to-noise ratio.

Keywords. Denoising, Monte Carlo, Optimization

1. INTRODUCTION

Image noise is a random variation of brightness or color information. In most real-life cases, it can be modeled as additive white Gaussian noise:

$$y = x + \eta \tag{1.1}$$

Where y is the noisy pixel, x the pure pixel and $\eta \sim \mathcal{N}(0, \sigma)$ is the noise.

1.1. Denoising Techniques. Traditional denoising techniques, such as Gaussian smoothing, Median filtering or Local means, operate on the principle of locality, assuming that the true value of the pixel must be similar to the values of its neighbours. This is effective at removing noise, but it also leads to loss of fine textures and blurring of edges. Newer techniques include Convolutional Neural Networks, Wavelet Transforms [8], and Modified Decision-Based Median Filter [9]. Although these methods can achieve impressive results, they often require extensive training data, computational resources, or complex parameter tuning. Non-Local Means (NLM) [3] is remarkable in its simplicity and effectiveness, however its high computational complexity limits real-time application.

1.2. Non-Local Means. The Non-Local Means estimates the true value of the pixel by computing a weighted average of different patches, using a weight function that prioritizes pixels with similar structural patterns. The most similar pixels to a given noisy pixel have no reason to be in its immediate vicinity. For example, patterns such as smooth surfaces, periodic textures or repeated structures can appear in different areas of the image. Although the search space is larger, we are still going to use only a neighbourhood of the pixel to limit the computational cost. As acknowledged by the authors themselves [4], the term “non-local” is somewhat misleading, and a more accurate name would be “*semi-local*”.

Definition 1.1. Suppose Ω is an image domain, and let $p, q \in \Omega$ be two points within the image. Then, the algorithm for denoising is defined as:

$$z(p) = \frac{1}{C(p)} \int_{\Omega} w(p, q) y(q) dq \quad (1.2)$$

where $z(p)$ is the denoised value at point p , $y(q)$ is the noisy value at point q , $w(p, q)$ is the weight function that measures the similarity between patches centered at p and q , and $C(p)$ is a normalizing factor given by:

$$C(p) = \int_{\Omega} w(p, q) dq \quad (1.3)$$

Obviously, in a real-world application we would use a discrete version of this algorithm:

$$z(p) = \frac{1}{C(p)} \sum_{q \in \Omega} w(p, q) y(q) \quad (1.4)$$

where, again, the normalizing factor is given by:

$$C(p) = \sum_{q \in \Omega} w(p, q) \quad (1.5)$$

Given a noisy pixel, the surrounding patch \mathbf{y} will be used to check for similarity with other patches. This object can be flattened into a d-pixel: $\mathbf{y} \in \mathbb{R}^d$. The algorithm requires a set of patches $\mathcal{X} = \{\mathbf{x}_1, \dots, \mathbf{x}_n\}$ that are obtained from reference images. Given this, the discrete version of NLM can be expressed as:

$$z = \frac{\sum_{i=1}^n w_i x_i}{\sum_{i=1}^n w_i} \quad (1.6)$$

where x_i represents the center pixel in the patch \mathbf{x}_i , and the weight w_i measures the similarity between the match \mathbf{y} and \mathbf{x}_i .

A standard choice for the weight function is:

$$w_i = e^{-\|\mathbf{y} - \mathbf{x}_i\|^2 / (2h_r^2)} \quad (1.7)$$

Where h_r is a scalar parameter that controls the decay of the distribution: small values create a fast decay and ensure a focus on stricter similarity between patches, whereas larger values allow different patches to have a greater weight. And $\|\cdot\|$ is the Euclidian norm.

The downside of NLM is its high computational complexity. The most expensive step of this algorithm is computing the weights w_i , which leads to $\mathcal{O}(mnd)$ operations, where m is the number of pixels to be denoised, n the number of patches and d the dimensions of the patch. In reality, a study has shown that it's enough to only use a localized window of dimension D around the pixel to be denoised [7], which reduces the complexity to $\mathcal{O}(mD^2d)$. Keep in mind that m and d are also quadratic in nature of the image and patch sizes.

A fun, but not-so-trivial way of improving the computational complexity is to use summed-area tables and the Fast Fourier Transform, which achieves a theoretical complexity of $\mathcal{O}(mD^2\sqrt{d})$. An implementation of this by Jacques

Froment [6], called *Parameter-Free Fast Pixelwise Non-Local Means Denoising*, claims to speed up the process by a factor of 6 to 49.

In reality, both of these implementations, as well as the one below, are easily parallelizable, and can be sped up even further using GPU computing. In 4, we will explore a different approach that uses KD-Trees to accelerate the search for similar patches, at the cost of a slight decrease in the quality of the denoised image.

1.3. Monte Carlo Non-Local Means (MCNLM). This paper focuses on the implementation and effectiveness of a method developed in [5]. The main focus of the approach is to approximate 1.6 by selecting randomly k reference pixel and creating a k -subset of weights $\{w_i\}^k$. The computational complexity of this approach is $\mathcal{O}(mkd)$, which is significantly lower for $k \ll n$.

2. MONTE CARLO NON-LOCAL MEANS

In this section, we discuss the full implementation of the algorithm, the correctness of the approach and some other optimizations that can be added to the method.

2.1. Sampling Patches. There are two ways to extract reference patches. The first one is based on picking them from the original noisy, image, which is called *internal denoising*. The second method is based on having the patches taken from a large database of other images, which is called *external denoising*. The paper focuses on internal denoising.

The Monte Carlo sampling is done on the set $\mathcal{X} = \{x_1, \dots, x_n\}$, considering each reference patch independent. The process is determined by a sequence of random variables $\{I_j\}_{j=1}^n$, where $I_j \sim \text{Bernoulli}(p_j)$. The weight will be sampled only if $I_j = 1$. The vector of these probabilities, $\mathbf{p} := [p_1, \dots, p_n]^T$, is called the *sampling pattern* of the algorithm [5].

The main parameter of this algorithm is ξ , which is the expected value of the random variable which models the ratio between the number of the samples taken and the number of references:

$$S_n = \frac{1}{n} \sum_{j=1}^n I_j \quad (2.1)$$

which has:

$$\xi \stackrel{\text{def}}{=} \mathbb{E}[S_n] = \frac{1}{n} \sum_{j=1}^n \mathbb{E}[I_j] = \frac{1}{n} \sum_{j=1}^n p_j \quad (2.2)$$

So, an important requirement is that:

$$\sum_{j=1}^n = n\xi \quad (2.3)$$

S_n is called the *empirical sampling ratio* and ξ the average sampling ratio [5].

2.2. Algorithm. Given a set of references \mathcal{X} and the sampling ration ξ , we can define the sampling pattern \mathbf{p} in order for it to respect the condition:

$$\sum_{j=1}^n p_j = n\xi \quad (2.4)$$

We can approximate the numerator and the denominator in 1.6 using two random variables:

$$A = \frac{1}{n} \sum_{j=1}^n x_j w_j \frac{I_j}{p_j} \quad \text{and} \quad B = \frac{1}{n} \sum_{j=1}^n w_j \frac{I_j}{p_j} \quad (2.5)$$

To show why we scale by $\frac{1}{p_j}$, we calculate the expected value of the estimators. By linearity of expectation:

$$\mathbb{E}[A] = \frac{1}{n} \sum_{j=1}^n x_j w_j \frac{\mathbb{E}[I_j]}{p_j} = \frac{1}{n} \sum_{j=1}^n x_j w_j \frac{p_j}{p_j} = \frac{1}{n} \sum_{j=1}^n x_j w_j \quad (2.6)$$

and

$$\mathbb{E}[B] = \frac{1}{n} \sum_{j=1}^n w_j \frac{\mathbb{E}[I_j]}{p_j} = \frac{1}{n} \sum_{j=1}^n w_j \quad (2.7)$$

So, A and B are *unbiased estimators* of the true numerator and denominator. In the end, we will approximate the result z by another random variable:

$$Z = \frac{A}{B} = \frac{\sum_{j=1}^n x_j w_j \frac{I_j}{p_j}}{\sum_{j=1}^n w_j \frac{I_j}{p_j}} \quad (2.8)$$

However,

$$\mathbb{E}[Z] = \mathbb{E}\left[\frac{A}{B}\right] \neq \frac{\mathbb{E}[A]}{\mathbb{E}[B]} = z \quad (2.9)$$

so, Z is a *biased estimator* of z . However, section 3.2 proves that as $n \rightarrow \infty$, the deviation $|Z - z|$ drops exponentially. This shows that, for a larger number of patches, the more efficient MCNLM gives results comparable to that of the simple NLM.

Algorithm 1 Monte Carlo NLM Denoising

Require: Noisy patch \mathbf{y} , reference set \mathcal{X} , sampling pattern \mathbf{p}

Ensure: Denoised pixel z

```
1: for  $j$  in  $1 \dots n$  do
2:   Generate random variable  $I_j \sim \text{Bernoulli}(p_j)$ 
3:   if  $I_j = 1$  then
4:     Compute  $w_j$ 
5:   end if
6: end for
7: Compute  $A = \frac{1}{n} \sum_{j=1}^n w_j x_j \frac{I_j}{p_j}$ 
8: Compute  $B = \frac{1}{n} \sum_{j=1}^n w_j \frac{I_j}{p_j}$ 
9: return  $Z = \frac{A}{B}$ 
```

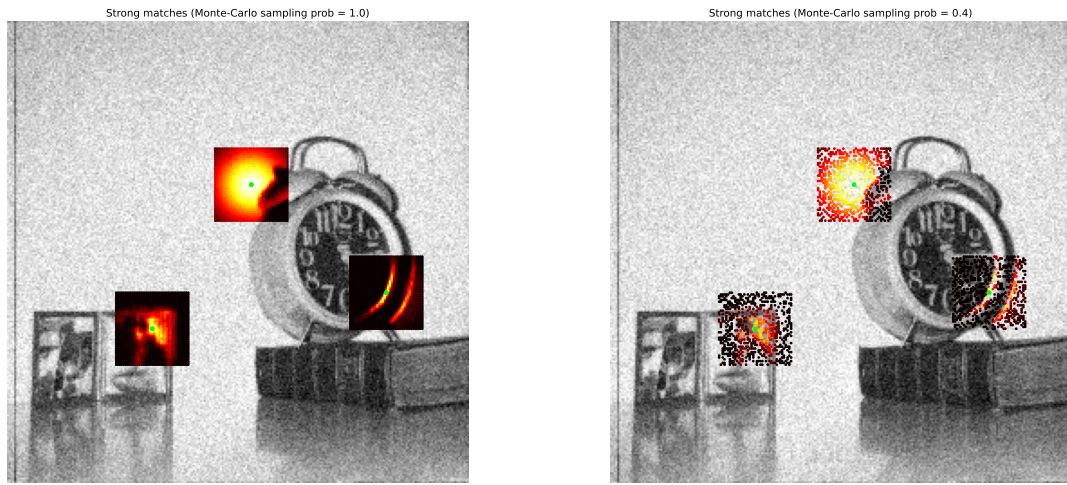
2.3. Spatial Sampling. We must also consider that locality matters in images. In a picture of a starry night, for example, a bright star might look like noise if compared to the rest of the dark sky. We can integrate this insight with the ingenious structural handling of MCNLM by incorporating a spatial distance parameter into the weighting function:

$$w_j = w_j^s \cdot w_j^r \quad (2.10)$$

where w_j^r is the weight mentioned at 1.7 and w_j^s is a spatial weight:

$$w_j^s = e^{-(d_j^2)^2 / (2h_s^2)} \cdot \mathbb{I}\{d_\infty^j \leq \rho\} \quad (2.11)$$

where d_j^2 is the Euclidean distance between the noisy patch and the reference patch we compare to and d_∞^j is the Chebyshev distance. The indicator function \mathbb{I} is used to determine a spatial search window of width ρ .



(A) Weight values in search window

(B) Weights for sampled center pixels

FIGURE 1. Main caption describing both images

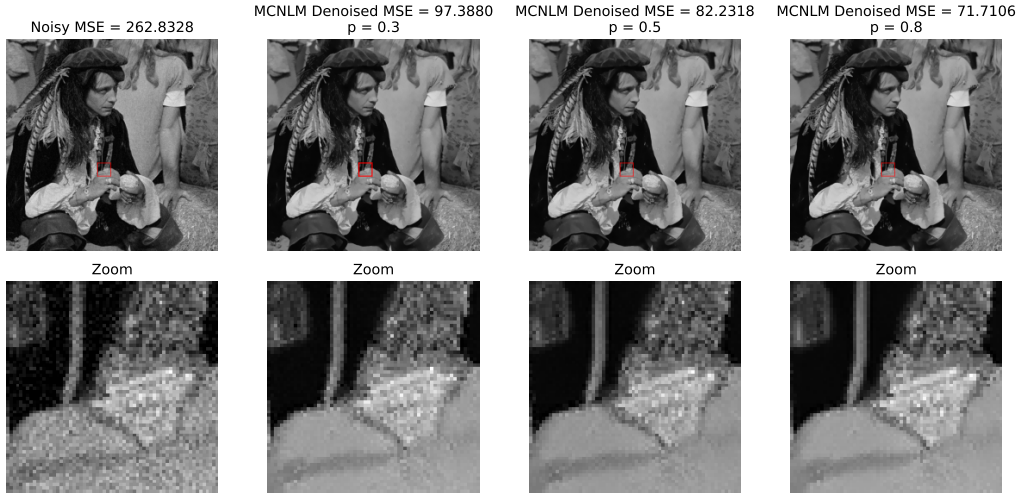


FIGURE 2. MCNLM with $\xi \in \{0.3, 0.5, 0.8\}$

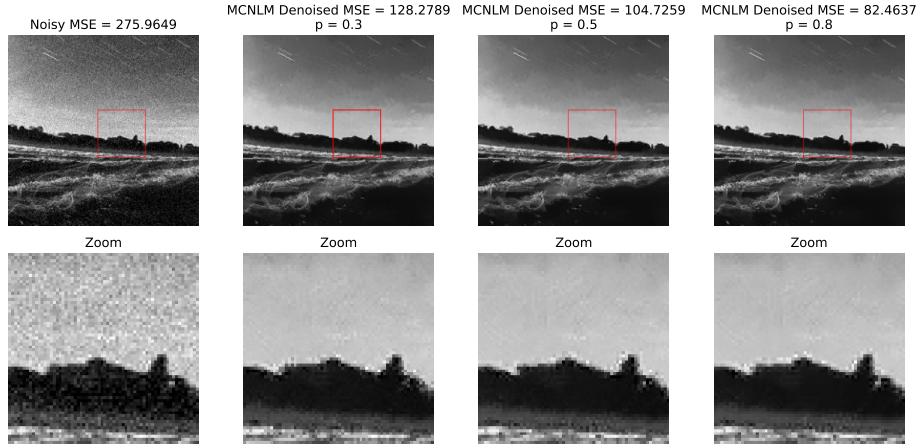


FIGURE 3. MCNLM with $\xi \in \{0.3, 0.5, 0.8\}$

Figure 4 visualizes the weight function within a search window centered on a noisy pixel, where brighter values indicate higher weight magnitudes. This illustrates how the method preserves structure: pixels with similar patches exert a stronger influence on the weighted average, ensuring that structural details are retained in the denoised output.

2.4. Results. In the figures 7, 3 we can see how the MCNLM algorithm behaves on an 1024×1024 image ($\approx 10^6$ pixels), with Gaussian noise with $\sigma = 17/255$ (taken into account that we normalize our pixels to have values in the interval $[0, 1]$) and zero mean. The comparison is done on 5×5 patches, and on a 20×20 search window around the noisy pixel. The sampling pattern is chosen as uniform, with $\mathbf{p} = \{\xi, \dots, \xi\}$. It is obvious that an increase in the sampling ratio only provides a marginally better result.

3. STOCHASTIC APPROXIMATION OF NON-LOCAL MEANS

We yield the question whether the Monte-Carlo variant of our algorithm is indeed reliable and approaches the full Non-Local Means solution. For this we need to analyze the approximation error $|Z - z|$.

To understand our approximation, we will study the empirical sampling ratio S_n . The law of large numbers states that the empirical mean S_n of a large number of indepent random variable is close to the true mean.

3.1. Probability Bounds. For any error bound $\varepsilon > 0$ and any sampling pattern p with $0 < p_j \leq 1$ and $\sum_{j=1}^n p_j = \varepsilon$ we want to study the probability:

$$\mathbb{P}(|S_n - \mathbb{E}[S_n]| > \varepsilon) \quad (3.1)$$

From the Cebyshev inequality we know that:

$$\mathbb{P}(|S_n - \mathbb{E}[S_n]| > \varepsilon) < \frac{\text{Var}[I_1]}{n\varepsilon^2}, \forall \varepsilon > 0 \quad (3.2)$$

This bound provided by the Cebyshev inequality above is too loose, only providing a linear descent of the error probability as $n \rightarrow \infty$.

We can use the Bernstein inequality to provide a much tighter exponential bound. The following is true for a sequence of n Bernoulli i.i.d. variables:

$$\mathbb{P}(|S_n - \mathbb{E}[S_n]| > \varepsilon) \leq \exp\left(-\frac{n\varepsilon^2}{2(\xi(1-\xi) + \varepsilon/3)}\right) \quad (3.3)$$

where $S_n = \frac{1}{n} \sum_{j=1}^n I_j$.

3.2. General Error Probability Bound. To rigorously quantify the reliability of the Monte Carlo approximation, we refer to the concentration bound derived in [5]. For a sample size n and an error tolerance $\varepsilon > 0$, the probability that the Monte Carlo estimate Z deviates from the exact NLM value z is bounded by:

$$\begin{aligned} \mathbb{P}(|Z - z| > \varepsilon) &\leq \exp\{-n\xi\} \\ &+ 2 \exp\left\{\frac{-n(\mu_B\varepsilon)^2}{2\left(\frac{1}{n} \sum_{j=1}^n \alpha_j^2 \left(\frac{1-p_j}{p_j}\right) + \frac{(\mu_B\varepsilon)M_\alpha}{6}\right)}\right\} \end{aligned} \quad (3.4)$$

3.2.1. Simplification for Uniform Sampling. While Equation 3.4 accounts for variable sampling probabilities, our implementation utilizes a uniform sampling pattern where $p_j = p$ for all patches. Furthermore we have $\mu_B \leq 1$.

Under these experimental conditions, the bound simplifies significantly. Neglecting the trivial failure term $\exp\{-n\xi\}$ (which approaches zero for our window size of $n = 441$), the failure probability becomes:

$$\mathbb{P}(|Z - z| > \varepsilon) \leq 2 \exp\left(\frac{-n\varepsilon^2}{2\left(\rho \sum_{j=1}^n \alpha_j^2 + \varepsilon/6\right)}\right) \quad (3.5)$$

where $n = 441$ is the total pixels in the 21×21 search window, $\rho = (1 - p)/p$ is the sampling penalty factor and $\alpha_j = \frac{w_j}{\sum_{k=1}^n w_k}$ is the normalized weight of the patch j .

3.2.2. Derivation of Numerical Bounds. To obtain a concrete probability bound, we will replace the values of the variables in Equation 3.5 with the parameters from our experimental setup. The parameters are, as follows, $n = 441$, for a 21×21 search window, $\rho = (1 - 0.5)/0.5 = 1$. We select the desired error tolerance of $\varepsilon = 0.05$.

The remaining term, $\sum_{j=1}^n \alpha_j^2$, is non-trivial as it depends on the image content. We can, however, derive bounds for two limiting cases:

Case 1: Homogeneous Regions. In flat image regions (e.g., sky or smooth walls), the NLM weights are distributed nearly uniformly across the search window.

$$w_j \approx w \quad \forall j \implies \alpha_j \approx \frac{1}{n}$$

Substituting this into the variance term:

$$\sum_{j=1}^n \alpha_j^2 \approx \sum_{j=1}^n \left(\frac{1}{n}\right)^2 = n \cdot \frac{1}{n^2} = \frac{1}{441} \approx 0.0023$$

From 3.5, we calculate that the resulting probability is $\mathbb{P}(|Z - z| > \varepsilon) < 10^{-20}$, confirming that random sampling is statistically safe in smooth regions.

Case 2: Structured Regions. In textured regions or near edges, the weights concentrate on a small subset of patches that have a high similarity to the patch we want to denoise. We estimate this by defining an *effective neighbor count* k_{eff} . If the algorithm finds only $k_{\text{eff}} \approx 10$ strong matches in the window (a conservative estimate for detailed texture), the normalized weights approximate $\alpha_j \approx 1/10$ for the matches and 0 otherwise.

$$\sum_{j=1}^n \alpha_j^2 \approx \sum_{j=1}^{10} \left(\frac{1}{10}\right)^2 = 10 \cdot \frac{1}{100} = 0.1$$

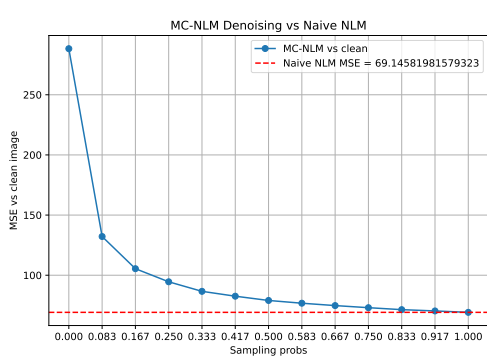
Using this estimate in 3.5:

$$\text{Exponent} = \frac{-441 \cdot (0.05)^2}{2(1 \cdot 0.1 + 0.05/6)} \approx \frac{-1.1025}{0.216} \approx -5.1$$

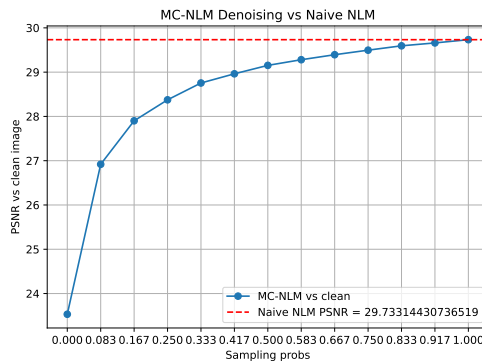
therefore

$$\mathbb{P}(|Z - z| > \varepsilon) \leq 2e^{-5.1} \approx 0.012$$

This shows that even in worst-case scenarios, the probability of the Monte Carlo estimate deviating by more than 5% is strictly bounded below 1.2%.



(A) MSE across different sampling probabilities



(B) PSNR across different sampling probabilities

FIGURE 4. Monte-carlo converge results

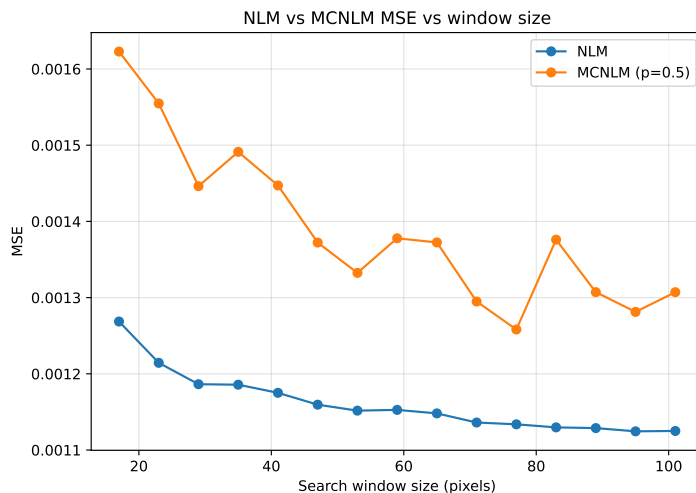


FIGURE 5. NLM vs MCNLM mse for different window sizes

4. KD-TREE ACCELERATED NON-LOCAL MEANS

This section describes a possible improvement of the basic Non-Local Means algorithm that would improve its theoretical time complexity by using a KD-Tree data structure to speed up the finding of similar patches, at the cost of slight accuracy loss. While the theoretical time complexity of this approach is better, the naive algorithm is laughably simple to parallelize, meanwhile this approach does have some bottlenecks in the query phase that are harder to do so; as such, the actual runtime improvement may be less than expected. However, it is still an interesting approach to explore and implement, and can be used as a middle ground between the naive NLM and more heuristic approaches like Monte Carlo NLM.

More complex approaches that are based on this idea have been presented by Adams et al. in the SIGGRAPH 2009 paper [1], that explores the idea of a *Gaussian KD-Tree*, and by Brox et al. [2] who introduced the notion of *Cluster*

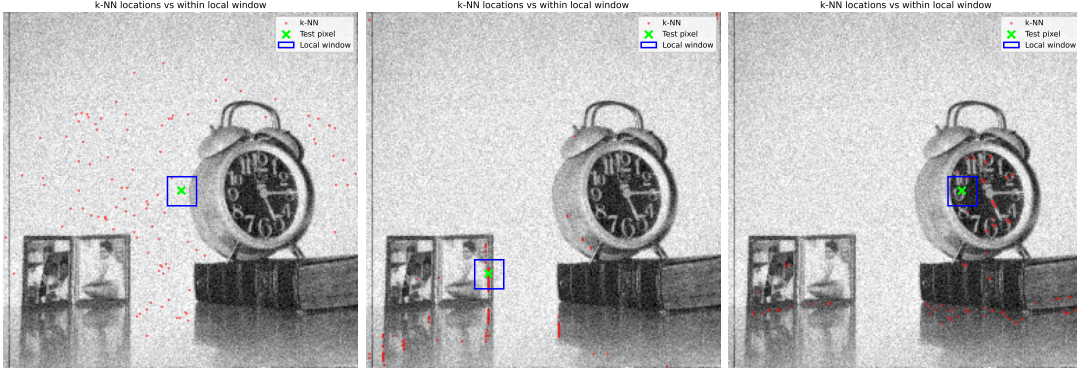


FIGURE 6. MCLNM vs KD-Tree NLM patch selection comparison

Trees. The approach proposed by Adams et al., is a more advanced data structure that combines KD-Trees with Gaussian Mixture Models to further speed up the search for similar patches. This approach is more difficult to implement, but can provide even better performance, since the original authors implemented this using a GPU with CUDA support.

4.1. Algorithm Description. Unlike the Monte Carlo NLM algorithm, which randomly samples patches from the search window, this approach builds a KD-Tree from all the patches in the search window, and then queries it for the K nearest neighbors of the current patch. This way, we can find the most similar patches more efficiently than by brute-force searching through all patches in the search window. More specifically, for each pixel p in the image, we extract its patch $B(p, f)$ and consider a point in $\mathbb{R}^{(2f+1)^2}$ corresponding to the flattened patch. This $(2f+1)^2$ -dimensional point is then inserted into the KD-Tree. After building the KD-Tree for all patches in the search window, we can query it for the K nearest neighbors of the patch $B(p, f)$, and use these patches to compute the weights and denoise the pixel p . The weight computation and denoising steps remain the same as in the original NLM algorithm, but we only consider the K nearest neighbors instead of all patches in the search window. Below is a high-level pseudocode of the proposed idea.

For RGB images, we would need to also consider the color channels, resulting in points in $\mathbb{R}^{3(2f+1)^2}$. However, as before, we will focus on grayscale images for simplicity.

Algorithm 2 KD-Tree Accelerated Non-Local Means Denoising

Require: Noisy image Y , patch half-size f , filtering parameter h , number of neighbors K

Ensure: Denoised image Z

```
1:  $\mathcal{P} \leftarrow$  Extract all patches  $B(p, f)$  from image  $Y$  and flatten them into vectors  
   in  $\mathbb{R}^{(2f+1)^2}$   
2:  $\text{KD} \leftarrow \text{BuildKDTree}(\mathcal{P})$   
3: for each pixel  $p$  in image  $Y$  do  
4:    $B_p \leftarrow$  Extract and flatten patch  $B(p, f)$   
5:    $\mathcal{N}_p \leftarrow \text{KD.Query}(B_p, K)$   
6:    $Z(p) \leftarrow 0, W \leftarrow 0$   
7:   for each patch  $B(q, f)$  in  $\mathcal{N}_p$  do  
8:      $d \leftarrow \text{ComputeDistance}(B_p, B(q, f))$   
9:      $w \leftarrow \exp\left(-\frac{d^2}{h^2}\right)$   
10:     $Z(p) \leftarrow Z(p) + w \cdot Y(q)$   
11:     $W \leftarrow W + w$   
12:   end for  
13:    $Z(p) \leftarrow Z(p)/W$   
14: end for  
15: return  $Z$ 
```

4.2. Experimental Results. We compared the two approaches (Monte Carlo NLM and KD-Tree NLM) with the `clock` image from the **SIPI Image Database**, corrupted with Gaussian noise with $\sigma = 17$. We can see that both approaches have very similar performance in terms of PSNR and MSE, even though they look pretty different. The Monte Carlo approach works very well with large uniform spaces, but struggles near rough edges, while the KD-Tree approach can outperform it in these areas, but looks more noisy in large uniform areas. This is to be expected, since the Monte Carlo approach is more likely to be able to find good patches in large similar areas, and struggle in very dense, high-frequency ones. The KD-Tree approach, on the other hand, is not restricted by a fixed window search area, and can find good patches even in high-frequency areas, as long as they are present somewhere in the image. However, the noise in the background can be attributed to the fact that it's behaving too good — since there are many patches similar to the noisy one, it's likely to pick the ones with similar noise patterns, resulting in less effective denoising.

5. NOISE ESTIMATION AND ALGORITHM IMPLEMENTATION

In the above algorithm it is assumed that the noise standard deviation σ is known. However, in real-life applications this is not always the case. There are multiple methods for estimating this value, but we are going to focus on a simple one based on the Fast Fourier Transform. The idea is that, in the frequency domain, the high-frequency components are more likely to represent noise rather

than actual image details. By analyzing these components, we can estimate the noise level. A pseudo-code of the algorithm is given below.

Algorithm 3 Gaussian Noise Standard Deviation Estimation using FFT

Require: Input image

- 1: $F \leftarrow \text{FFT}(\text{image})$
 - 2: Shift the zero-frequency component F to the center of the spectrum
 - 3: Define a high-frequency mask M that selects the high-frequency components
 - 4: $H \leftarrow F \cdot M$
 - 5: $\text{noise} \leftarrow \text{Inverse FFT Shift of } H$
 - 6: Keep only the real part of noise and remove the mean.
 - 7: **return** $\sigma \leftarrow \text{std}(\text{noise})$
-

While this obviously doesn't yield perfect results like already knowing the noise level, it provides a reasonable estimate that can be used in the denoising process. We can see that the estimated noise standard deviation is close to the actual value, which is sufficient for our denoising algorithm to perform effectively. However, the blurring effect can start to be noticeable when we compare the two images side by side.

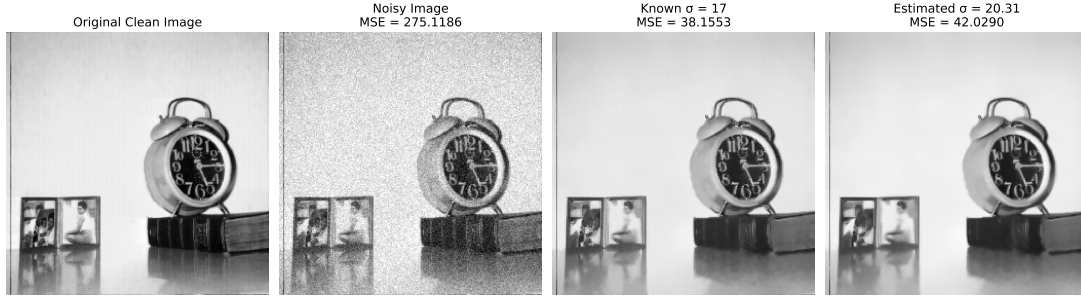


FIGURE 7. Comparison between using a known σ and an estimated one

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