

Spectral Graph Theory and High Performance  
Computing  
Master Thesis

David Wobrock `david.wobrock@gmail.com`

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# Chapter 1

## Introduction

### 1.1 Background

The talk by Milanfar [1], working at Google Research, about using the Graph Laplacian Operator for Image Processing purposes awakes curiosity.

Indeed, Milanfar reports that these techniques to build filters are used on smartphones, which means it is done in a reasonable time with limited computational resources. Over 2 billion photos are shared daily on social media [1], with very high resolutions and filters are often applied on them.

### 1.2 Objective

The objective of this thesis is to, in a first place, understand and implement some image processing algorithms shown by Milanfar's work on 2D images. We want to observe the performance and results of these algorithms to be validate the usage of intrinsic spectral properties of an image. For instance, the spectrum of the graph associated to an image can serve for image segmentation. Applied in a fine-grained manner, this segmentation could lead to image compression. More classic algorithm can also be achieved with the described approach, such as image denoising, deblurring, sharpening, etc.

The next step includes extending these techniques on 3D images. Given the results, new scalable processing approaches for 3D images could be explored. These also include standard processing filters such as denoising and sharpening, but also segmentation and compression techniques.

These approaches, even though faster than other filters, still require computations which can grow fast in the 3D case. That is why we want to apply the algorithm on high performance architectures in order to achieve reasonable computation times.

## 1.3 Related work

In this section we will explore three topics concerning this project. First of all, we will dive into spectral graph theory. Then, we will deal with 2D image processing, focusing three applications: denoising, sharpening and image segmentation. For each of them, we show a classical algorithm, a new spectral approach and then a quick comparison of them. Finally, we present what has been done in 3D image processing so far using spectral graph theory, essentially using the graph laplacian operator.

### 1.3.1 Spectral graph theory

Spectral graph theory has a long history starting with matrix theory and linear algebra that were used to analyse adjacency matrices of graphs. It consists in studying the properties of graphs in relation to the eigenvalues and eigenvectors of the adjacency or Laplacian matrix of the graph. The eigenvalues of such a matrix are called the spectrum of the graph. The second-smallest eigenvalue has been called “algebraic connectivity of a graph” by Fiedler [2], and is therefore also known as *Fiedler value*.

This field is very broad and the eigen-decomposition of graphs is used in a lot of areas. First, it was applied to chemistry because eigenvalues can be associated with the stability of molecules. Spectral graph theory has a many other applications such as graph coloring, graph isomorphism testing, random walks and graph partitioning among others.

One of the most complete works about spectral graph theory is [3] by Fan Chung. This monograph exposes many properties of graphs, the power of the spectrum and how spectral graph theory links the discrete world with the continuous one.

**Laplacian matrix** Since the adjacency matrix of a graph only holds basic information about it, we usually augment it to the Laplacian matrix. Multiple definitions of the Laplacian matrix are given in [3] and each hold different properties. The most common ones are the normalised Laplacian and the Random walk Laplacian.

**Cheeger’s inequality** One of the most fundamental theorems of spectral graph theory concerns the Cheeger’s inequality and Cheeger constant. It approximates the sparsest cut of a graph with the second eigenvalue of its Laplacian.

The Cheeger constant measures the degree of “bottleneck” of a graph, useful for constructing well-connected graphs. Considering a graph  $G$  of  $n$  vertices, the Cheeger constant is defined as

$$h(G) = \min_{0 < |S| \leq \frac{n}{2}} \frac{|\partial S|}{|S|},$$

where  $S$  is a subset of the vertices of  $G$  and  $\partial S$  is the *edge boundary* of  $S$  to have all edges with exactly one endpoint in  $S$ , or formally

$$\partial S = \{u, v \in E(G) : u \in S \text{ and } v \notin S\},$$

with  $E(G)$  the edges of graph  $G$ .

Cheeger's inequality defines a bound and relationship on the smallest positive eigenvalue of the Laplacian matrix  $\mathcal{L}$  such as

$$\lambda_1(\mathcal{L}) \geq \frac{h^2(\mathcal{L})}{4}.$$

When the graph  $G$  is  $d$ -regular, thanks to [4], we also have an inequality between  $h(G)$  and the second-smallest eigenvalue  $\lambda_2$  such as

$$\frac{1}{2}(d - \lambda_2) \leq h(G) \leq \sqrt{2d(d - \lambda_2)}.$$

### 1.3.2 2D image processing

#### Denoising

**Background** Even with high quality cameras, denoising and improving a taken picture remains important. The two main issues that have to be addressed by denoising are blur and noise. The effect of blur is internal to cameras since the number of samples of the continuous signal is limited and it has to hold the Shannon-Nyquist theorem [5]. Noise comes from the light acquisition system that fluctuates in relation to the amount of incoming photons.

To model these problems, we can formulate the deficient image as,

$$y = z + e,$$

where  $e$  is the noise vector of variance  $\sigma^2$ ,  $z$  the clean signal vector and  $y$  the noisy picture.

What we want is a high-performance denoiser, capable of scaling up in relation to increasing the image size and keeping reasonable performances. The output image should come as close as possible to the clean image. As an important matter, it is now accepted that images contain a lot of redundancy. This means that, in a natural image, every small enough window has many similar windows in the same image.

**Traditional, patch-based methods** The image denoising algorithms review proposed by [5] suggests that the NL-means algorithm, compared to other reviewed methods, comes closest to the original image when applied to a noisy image. This algorithm takes advantage of the redundancy of natural images and for a given pixel  $i$ , predicts its value by using the pixels in its neighbourhood.

In [6], the authors propose the BM3D algorithm, a denoising strategy based on grouping similar 2D fragments of the image into 3D data arrays. Then,

collaborative filtering is performed on these groups and return 2D estimates of all grouped blocks. This algorithm exposed state-of-the-art performance in terms of denoising at that time. The results are still one of the best for a reasonable computational cost.

**Global filter** In the last couple of years, global image denoising filters came up, based on spectral decompositions [7]. This approach considers the image as a complete graph, where the filter value of each pixel is approximated by all pixels in the image. We define the approximated clean image  $\hat{z}$  by,

$$\hat{z} = Wy,$$

where  $W$  is our data-dependent global filter, a  $n \times n$  matrix,  $n$  the number of pixels in the picture.  $W$  is computed from the graph affinity matrix<sup>1</sup>  $K$ , also of size  $n \times n$ , such as

$$K = \mathcal{K}_{ij},$$

where  $\mathcal{K}$  is a similarity function between two pixels  $i$  and  $j$ . As the size of  $K$  can grow very large, we sample the image and compute an approximated  $\hat{K}$  on this subset using the Nyström extension.  $K$  can in fact be approximated through its eigenvalues and eigenvectors. Knowing that the eigenvalues of  $K$  decay very fast [1], the first eigenvectors of the subset of pixels are enough to compute the approximated  $\hat{K}$ .

Generally, as proposed in [7] and [8], to improve the denoising performance of global filters, pre-filtering techniques are used. It is proposed to first apply a NL-means algorithm to the image to reduce the noise, but to still compute the global filter on the noisy input and apply the filter to the pre-filtered image.

**Comparison between patch-based and global methods** As [8] suggests, global filter methods have the possibility to converge to a perfect reconstruction of the clean image, which seems to be impossible for techniques like BM3D. Global filtering also seems promising for creating more practical image processing algorithms.

The thesis [9] proposes a normalised iterative denoising algorithm which is patch-based. The work reports that this technique has slightly better results than the global filter but essentially has a better runtime performance.

## Sharpening

**Background** Sharpening consists basically in a high-pass filter which will magnify high frequency details [9]. The two main issues again with this approach are that noise often displays high frequency attributes, which will be amplified by the filter. Secondly, the phenomenon of overshooting and undershooting, which appears when sharpening already sharp parts (edges for example), will exhibit unpleasant artifacts.

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<sup>1</sup>Also called kernel matrix or similarity matrix

Modeling the problem is the same as for denoising,  $\hat{z} = Fy$ ,  $F$  being our filter.

**Classical Difference of Gaussian (DoG) operator** This technique consists of computing the difference between two gaussian kernels, which will produce a range of different kernels with various frequencies. Indeed, DoG involves subtracting a blurred version of the input image to another, less blurred version of this original image [10]. A blurred image can be obtained by convolving the input image with a gaussian kernel. This gaussian blurring actually removes high-frequency spatial information. So the idea is that by subtracting this blurred image from a less blurred one, we will keep the high-frequency information, like a high-pass filter<sup>2</sup>, which results in image sharpening.

**Structure-aware sharpening** We know from its definition [11] that the smoothing filter  $W$  is a symmetric and doubly stochastic matrix<sup>3</sup>. So the largest eigenvalue  $\lambda_1 = 1$  shows that it has low pass filter characteristics. With the definition of the Laplacian, in relation to the matrix  $W$  such as  $\mathcal{L} = I - W$ , we can consider that it is a data-adaptive high-pass filter [9]. So we can define a data-adaptive unsharp mask filter as

$$F_1 = I + \beta(I - W),$$

with  $\beta > 0$ . This is basically a weighted high-pass filtered version of the input image [1].

But with this approach, the noise amplifying and overshoot problems remain. The proposed solution in [11] applies first a smoothing filter to the image, then the unsharp mask filter and finally the same smoothing filter again. The smoothing and sharpening filters use possibly two different kernels matrices as basis. The first smoothing filter aims to reduce the noise, while we still avoid over-smoothing. The second and final smoothing controls the effect of the amplified noise and the overshoot artifacts. This gives us the filter

$$F = W_1(I + \beta(I - W_2))W_1,$$

where  $W_1$  and  $W_2$  are constructed from the affinity matrices  $K_1$  and  $K_2$ , which can differ in relation to the parameters of their respective kernel function.

**Comparison** As [11] suggests, the structure-aware sharpening is data-adaptive and more noise robust than a classical DoG filter. Even though, both approaches use a similar concept. The proposed filter is actually a data-derived DoG-based filter. We can rewrite the proposed filter as

$$F = (1 + \beta)W_1^2 - \beta W_1 W_2 W_1,$$

where we can actually see the difference between two versions of the input image.

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<sup>2</sup>More precisely, like a band-pass filter here

<sup>3</sup>Depending on the used Laplacian definition. For example the one obtained by ‘‘Sinkhorn’’ iterations [12]

## Image Segmentation

**Background** To recognise patterns, find groups and cut images is a similar problem to graph partitioning. As is known, most formulations of partitioning a graph result in a NP-hard problem. So it will be our goal to approximate quickly good partitions instead of looking for an optimal solution.

A graph  $G = (V, E)$  is partitioned into two disjoint sets  $A$  and  $B$  such as  $A \cup B = V$  and  $A \cap B = \emptyset$  by removing edges between  $A$  and  $B$ . One can compute the value of such a *cut* with the degree of dissimilarity between  $A$  and  $B$ , which is calculated from the total weight of the removed edges

$$cut(A, B) = \sum_{u \in A, v \in B} w(u, v).$$

**Minimum cut** The most common way of partitioning a graph is the *minimum cut*. This problem consists in looking for the optimal bipartition of a graph, which is one that minimises the *cut* value. The *minimum cut* problem is well-studied and has efficient algorithms for solving it, even though it has an exponential amount of partitions that can be explored.

Some work on image segmentation using the *minimum cut* has been done [13] [14] [15]. But the results can only be qualified as proposals for recognised regions, such does [14] and would need further optimisation to be the final groups. For instance, [13] introduced a cut criterion using *minimum cut* on a graph but it was observed that it favors finding too small components and [15] tries to optimise the latter.

However, this problem still remains and *minimum cut* can produce bad groups on certain cases. Additionnaly, these cases often appear in natural images, where it is more important to group a large area of the background before small details.

**Normalised cut** To improve this state, [16] proposed, instead of looking at the total weigh of the edges connecting two groups, to look at a fraction of the total edge connections to all nodes. This is the *normalised cut*, or *Ncut*. The disassociation measure can be formulated as

$$Ncut(A, B) = \frac{cut(A, B)}{assoc(A, V)} + \frac{cut(A, B)}{assoc(B, V)},$$

where  $assoc(A, V) = \sum_{u \in A, t \in V} w(u, t)$ , the total connection from the nodes in  $A$  to all nodes in the graph.

The goal is still to minimise the *Ncut* value, but when *cut* is small for isolated pixels, the fraction will be greater because the similarity *assoc* between this one pixel and all pixels will be small. This fraction's denominator will be bigger if the similarity between the selected pixels and all pixels is more important, which makes the fraction smaller.



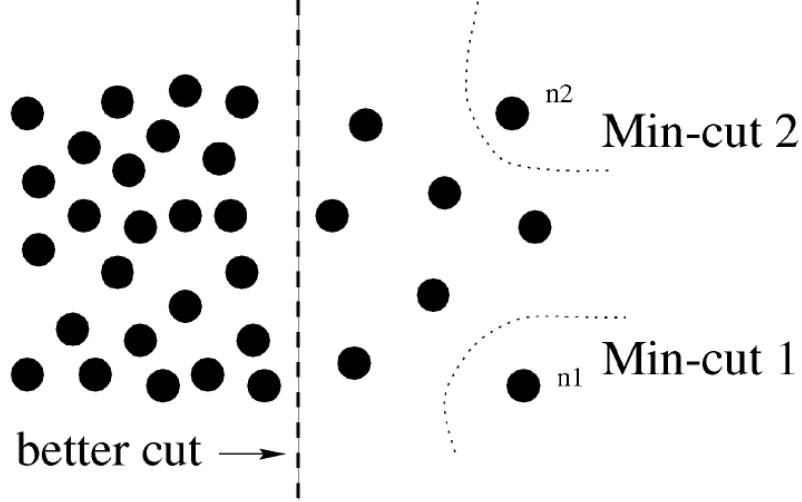


Figure 1.1: Bad partitioning from *minimum cut* in a particular case [16]

As it follows, we can define the normalised association measure within groups, such as

$$Nassoc(A, B) = \frac{assoc(A, A)}{assoc(A, V)} + \frac{assoc(B, B)}{assoc(B, V)},$$

where  $assoc(A, A)$  is the total weights of edges inside group  $A$ . This measure quantifies how tightly the nodes of a group are connected to each other. Deriving the  $Ncut$  equation, we can define that

$$Ncut(A, B) = 2 - Nassoc(A, B).$$

It was shown by [17] and [16] that minimising the  $Ncut$  exactly is a NP-complete problem. However, [16] proposes a method to approximate discrete solutions efficiently for the *normalised cut* problem.

Indeed, the proposed algorithm starts by computing the associated similarities for graph  $G = (V, E)$  for the image and store it in the matrices  $K$  and  $D$ .

Then, solve  $(D - K)x = \lambda Dx$  for the eigenvectors with the smallest eigenvalues. The way this equation is derived is explicitly shown in [16]. The eigenvector with the second smallest is used to bipartition the graph by finding the best splitting point to minimise  $Ncut$ . After that, it is possible to use the other eigenvectors to bipartition the graph even more. Once this is done, we can recursively apply the same algorithm on the segmented groups, stoping when the maximum number of groups is reached or the partitions are not good enough depending on an empirical threshold.

**Edge separators and spectral rounding** Introduced by [18], edge separators of a graph are created by iteratively reweighting edges until the graph disconnects into a certain number of groups. At each iteration, only a small number of eigenvectors are computed to reweight the edges. This spectral rounding method produces directly discrete solutions, and seems to compare nicely against the *Ncut* approximation by producing more human-like results [19].

### 1.3.3 3D image processing

In general, [20] sets a very good overview of most variants and applications of spectral processing.

## 1.4 Delimitations

The number of investigated algorithms and variants depend on the pre-study and objectives that make the most sense for 3D images. It is obviously also limited by the amount of time given to the degree project.

## Chapter 2

# 2D Image Processing using the Graph Laplacian Operator

### 2.1 Theoretical basis

Multiple image processing filters can be built by Graph Laplacians. As Milanfar mentions in [1], smoothing, deblurring, sharpening, dehazing, and other filters can be created. Laplacians can also be used for compression artifact removal, low-light imaging and image segmentation.

As it is known, an image filter consists of a function which outputs one pixel, taking all pixels as input and applying weights to them. We can write this as

$$z_i = \sum_j W_{ij} y_j,$$

$z_i$  being the output pixel,  $W_{ij}$  the weight and  $y_j$  all input pixels. This means that we have a vector of weights for each pixel.

So, as a practical notation, we can say that, with  $W$  the matrix of weights and  $y$  the input image as a vector,

$$z = Wy.$$

Now we want to represent the image as a graph. Each pixel is a node and has edges to multiple other nodes. We can arbitrarily define how the pixels connect to each other, so we can say that the graph is complete, each node connects to all other nodes. But we can weigh the edges to measure the similarity between pixels.

Affinity, or similarity, is a subjective term which we can also define as we need. Pixels can be similar if they are spatially close or if they have the same color, or both. These similarities give us the so-called affinity matrix  $K$ .

By extending this affinity matrix, we obtain the Graph Laplacian  $\mathcal{L}$ . And we want to build the filter  $W$ , containing the weights, from the Laplacian.

According to [21], to build a filter from the Laplacian, we have

$$W = I - \mathcal{L},$$

and so reciprocally

$$\mathcal{L} = I - W.$$

The Graph Laplacian has multiple definitions, but the simplest is the unnormalised one:

$$\mathcal{L}_U = D - K,$$

with  $D$  a positive definite diagonal matrix with the normalising factors such as  $D_{jj} = \text{diag}\{\sum_i K_{ij}\}$  along its diagonal.  $D$  corresponds in fact to the degrees of each node in graph theory terminology. Other definitions of the Laplacian are shown later in 2.3.3.

Interestingly, we can define from [7] that, by approximating  $W$  by a symmetric, positive definite, double stochastic matrix, this symmetric  $W$  can be computed thanks to this eigen-decomposition

$$W = VSV^T,$$

$V$  being the eigenvectors and  $S$  the eigenvalues as a diagonal matrix.

Our global filter can be expressed as

$$z = Wy = VSV^Ty.$$

This will be useful since the filter matrix  $W$  becomes huge very quickly and we will need a way to approximate it. Indeed,  $W$  is a square matrix with  $n * n$  elements,  $n$  the number of pixels in the picture.

**Approximation by Nyström Extension** To compute the filter, the first matrix that is needed is the affinity matrix  $K$ . Both matrices have the same size and are computationally expensive.

That is why we approximate the affinity matrix by sampling the picture. We sample  $p$  pixels, such as  $p \ll n$ .

Different sampling techniques exist and are shown in 2.3.1.

Once sampled, we compute  $K_A$  and  $K_B$ , which are parts of  $K$  such as

$$K = \begin{bmatrix} K_A & K_B \\ K_B^T & K_C \end{bmatrix}.$$

$K_A$  represents the affinity matrix of size  $p \times p$  between the sample pixels, whereas  $K_B$  is the affinity matrix of size  $p \times m$ , such as  $m = n - p$ , between the sample pixels and the remaining pixels.

These matrices will be computed thanks to a kernel function (or affinity function). This can be the bilateral filter, non-local means or another, as shown in 2.3.2.

Once computed, we can approximate the eigenvectors  $\Phi$  and eigenvalues  $\Pi$  of  $K$  thanks to the eigen-decomposition of  $K_A$ .

As stated in [7], we can define the decomposition of  $K_A$

$$K_A = \Phi_A \Pi_A \Phi_A^T$$

and the approximation of  $K$  such as

$$\tilde{K} = \tilde{\Phi} \Pi_A \tilde{\Phi}^T$$

and finally the approximation of the  $p$  first eigenvectors of  $K$

$$\tilde{\Phi} = \begin{bmatrix} \Phi_A \\ K_B^T \Phi_A \Pi_A^{-1} \end{bmatrix}$$

From this eigen-decomposition approximation of  $K$ , we can then approximate the eigenvectors and eigenvalues of  $W$ .

One might wonder how an approximation of the first elements of the eigen-decomposition can be enough to approximate the whole matrix. In fact, the eigenvalues decay quickly as shown in [1] and [22].

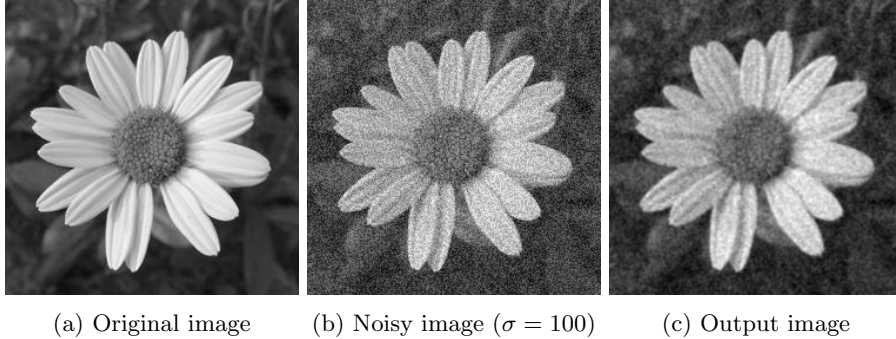
As the experiment below shows, the eigenvalues decrease quickly which means that the whole matrix is mainly defined by the first elements.

## 2.2 First implementation

**Details** The first implementation of this flow is aimed to denoise the input image. This represent an early stage of the internship. The sample pixels are selected in a spatially uniform manner. It uses the NLM from [5] as kernel method to compute the pixels similarity. The Laplacian matrix is defined through the Sinkhorn iterative algorithm [12], where the resulting approximated eigenvectors require to be orthogonalised, which can be done in one step, as discussed in [23]. We finally apply our approximated filter to the noisy image.

The implementation is generally greatly inspired by the MatLab implementation of [7].

**Results** The results of this very first experiment are not yet satisfying as we can observe visually:



We observe that the output picture is still quite noisy and blurry, we can certainly do better. But still, as a first result, this is better than nothing.

## 2.3 Algorithm variations

### 2.3.1 Sampling method

The sample requires to represent only less than 1% of the pixels of the image. To achieve this, we can use different approaches. The chosen method is decisive for the Nyström method.

**Random sampling (RS)** most common and simple sampling scheme, but no deterministic guarantee of the output quality. Can produce good results for images with poor resolution, but with a huge amount of data, random sampling is limited because it cannot reflect the structure of the data set [24].

**K-means sampling (KS)** associate to each pixel a 5-D space (R, G, B, X, Y) and divide the pixels into K clusters (K centers). These clusters are a good sampling scheme for images with simple and uniform backgrounds [25] [26].

**Uniform spatially sampling** the uniformity of the sample gives good results for image sampling because of the spatial correlation of pixels. This method remains simple but effective [7].

**Incremental sampling (INS)** is an adaptive sampling scheme, meaning that it select points according to the similarity, so that we can have an approximate optimal rank-k subspace of the original image [24].

**Mean-shift segmentation-based sampling** this scheme performs good for complex backgrounds. The method consists in over-segmenting the image into  $n$  regions and only one pixel of each region will be sampled using the spatially closest pixel to the center of the region given a formula in [25].

### 2.3.2 Affinity function

The kernel function  $K_{ij}$  measures the similarity between the pixel  $y_i$  and  $y_j$ .

#### Listing

**Spatial Gaussian Kernel** takes only into account the spatial distance between two pixels [1].

**Photometric Gaussian Kernel** considers the intensity and color similarity of the pixels [1].

**Bilateral Kernel** one of the most used kernel which smooths images by a nonlinear combination of the spatial and photometric gaussian kernels [1] [7].

**Non-Local Means (NLM)** is similar to the bilateral kernel, a data-dependent filter, except that the photometric affinity is captured patch-wise [7].

**Locally Adaptive Regression Kernel (LARK)** uses the geodesic distance based on estimated gradients [12] [27].

### 2.3.3 Graph Laplacian

Graph Laplacian has multiple possible definitions and each has its own properties. A good summary can be found in [1]. A Graph Laplacian can be symmetric which is important for eigen-decomposition of the matrix. It can have a DC eigenvector, which means that the Laplacian has to give 0 if we apply it to a constant image. This is also useful to have. And the spectral range, corresponding to the range of the eigenvalues, is important because we will use the filters derived from the Laplacian multiple times, and if the eigenvalues are not between 0 and 1, then the filters tend to be unstable. With  $K$  being the affinity matrix,  $d_i = \sum_j K_{ij}$  and  $D = \text{diag}\{d_i\}$ :

Laplacian Name	Formula	Symmetric	DC eigenvector	Spectral Range
Un-normalised	$D - K$	Yes	Yes	$[0, n]$
Normalised	$I - D^{-1/2} K D^{-1/2}$	Yes	No	$[0, 2]$
Random Walk	$I - D^{-1} K$	No	Yes	$[0, 1]$
“Sinkhorn” [12]	$I - C^{-1/2} K C^{-1/2}$	Yes	Yes	$[0, 1]$
Re-normalised	$\alpha(D - K), \alpha = \mathcal{O}(n^{-1})$	Yes	Yes	$[0, 1]$

Generally, it is a good practice to stick to one definition of the Laplacian.

## 2.4 Algorithm variations examples

To illustrate the impact of the affinity function, here are some examples of the affinity matrix for certain pixels. The more a pixel is colored in red, the more

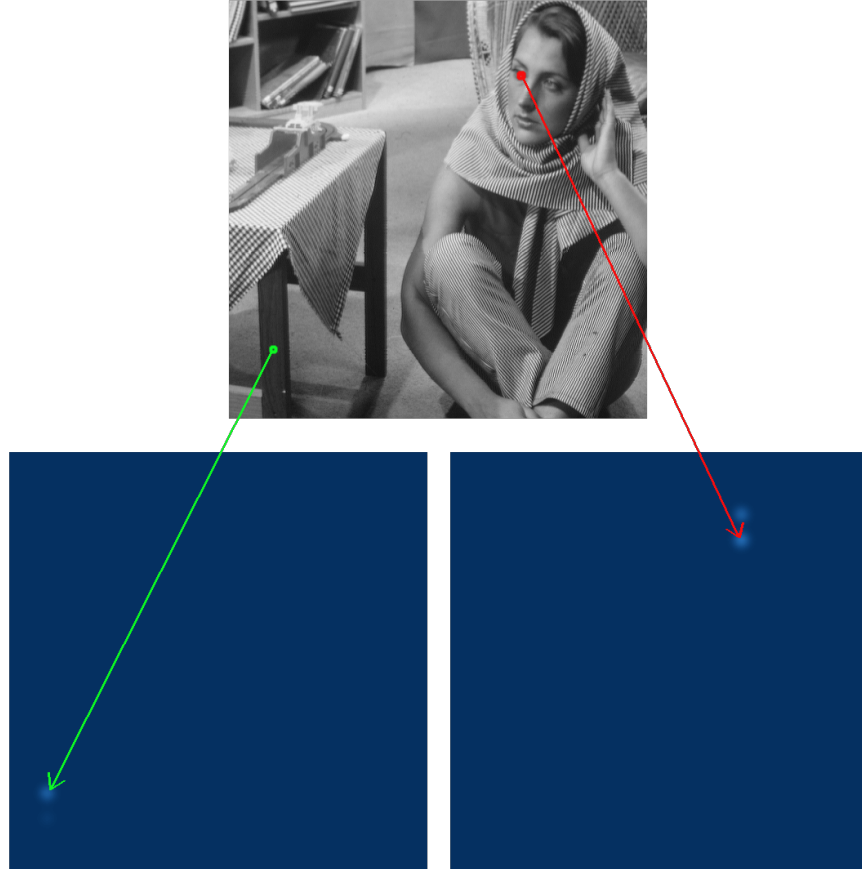
similar it is to the selected pixel, with respect to the chosen function. A blue colored pixel is dissimilar to the considered pixel.

To generate these examples, we use an image of the famous Barbara of dimension 512x512 pixels (grayscale image). Generating the affinity matrix takes approximately 120 seconds on an Intel i5 processor. We use a spatially uniform sampling technique and select 0.1% of the pixels. We show two affinity matrices for each techniques, the first one is of a pixel on the table leg and the second on Barbara's eye.

**Spatial Gaussian Kernel** The formula of this kernel is

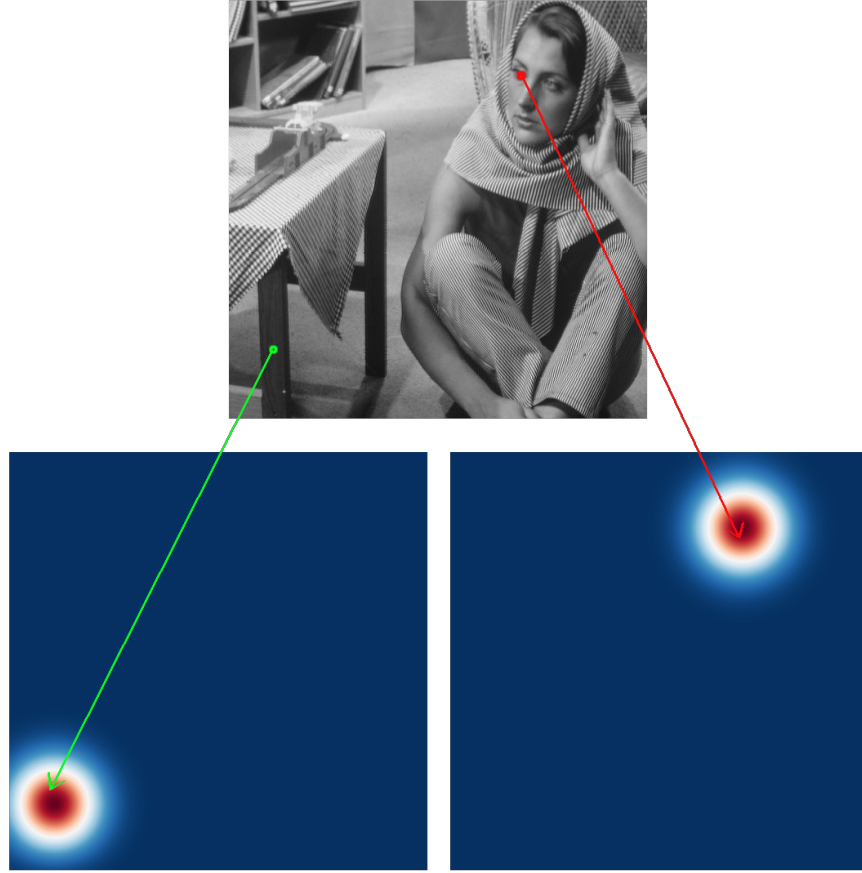
$$K(x_i, x_j) = \exp\left(-\frac{\|x_i - x_j\|^2}{h_x^2}\right).$$

Affinity matrices with  $h_x = 10$ :



Affinity matrices with  $h_x = 50$ :



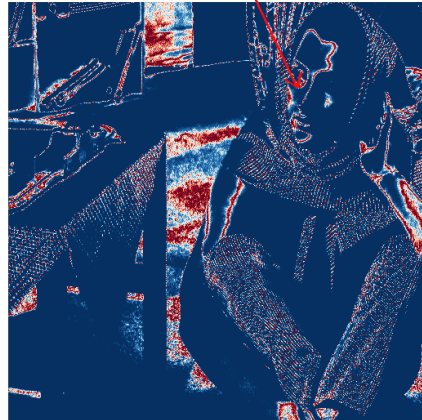
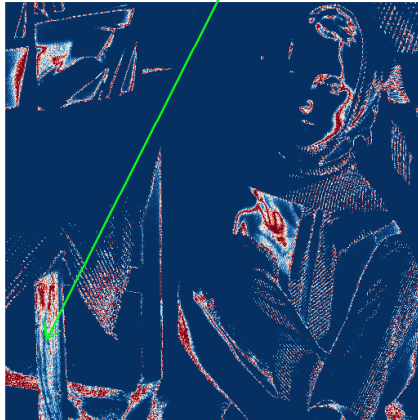


As we can see, the parameter is influencing on the normalisation of the values and gaussian standard deviation. The bigger it is, the more tolerant the spatial distance computation will be.

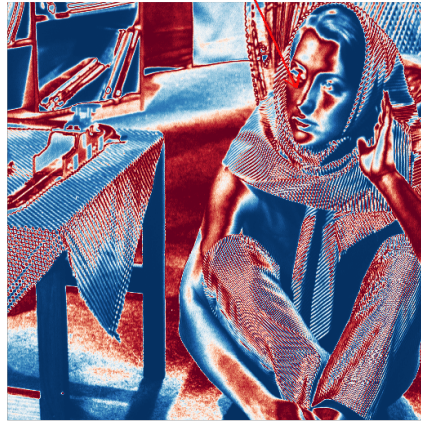
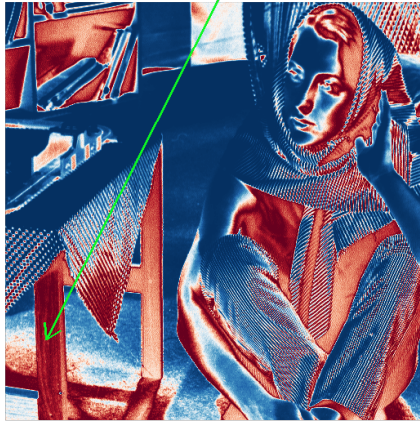
**Photometric gaussian Kernel** The formula of this kernel is

$$K(z_i, z_j) = \exp\left(-\frac{\|z_i - z_j\|^2}{h_z^2}\right).$$

Affinity matrices with  $h_z = 10$ :

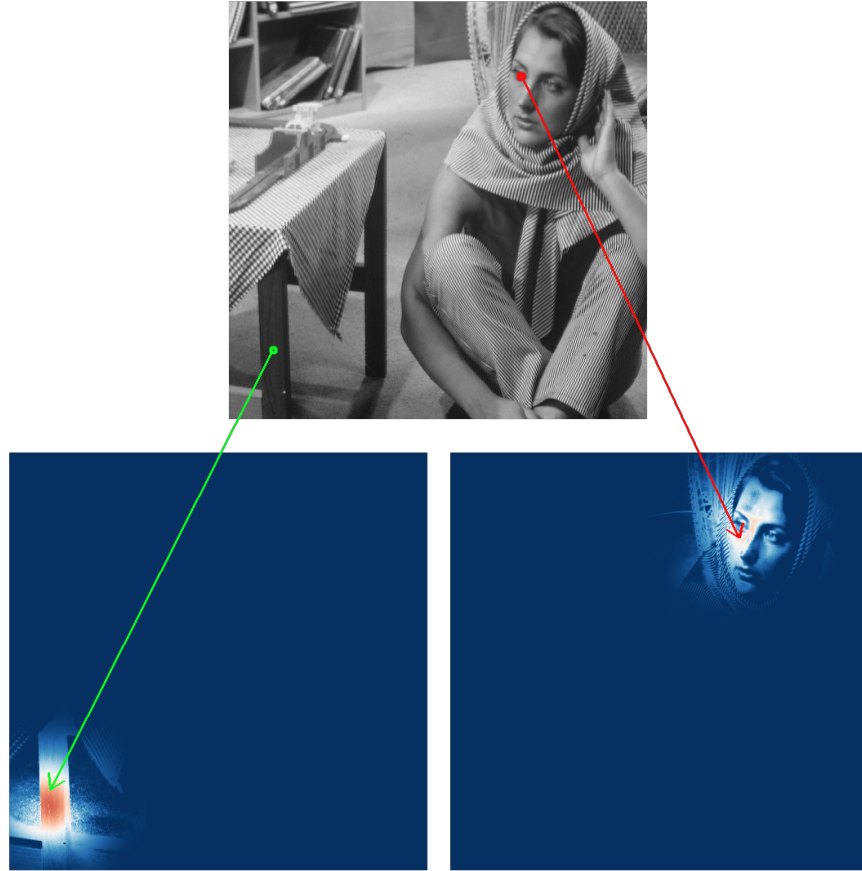


Affinity matrices with  $h_z = 50$ :



Generally, the  $h$  parameters in both kernel functions here are smoothing parameters. If  $h$  is small, it is more discriminating between the affinity of different pixels.

**Bilateral Kernel** Affinity matrices with  $h_x = 50$  and  $h_z = 35$ :



Remember that each matrix here is only the affinity matrix of one pixel. In a very heterogeneous image, the bilateral kernel will be useful to keep the spatial similarity, but with excluding very dissimilar neighbour pixels.

### 2.4.1 Pixel degree

An interesting matrix that can be observed is the degree of the pixels. A pixel, in our case, is a node of the graph, so we sum the weights of outgoing edges from each node. On the picture below, red pixels indicate that the pixel has a lot of similarity with all other pixels in the image, which can be roughly interpreted as the pixel is a common one:



(a) Original image

(b) Pixel degrees

We see that the most uncommon pixels are very light ones and very dark ones in this image.

## 2.5 Adaptive Sharpening

Sharpening is usually using a highpass filter and therefore leads to amplifying high frequency noise components even more. We will have an effect of over-sharpening.

$\beta$  controls the amount of sharpening of the image.

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