

## Vectorization of a statistical segmentation

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# Paper: Vectorization of a statistical segmentation

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## Abstract

We propose an efficient vectorial implementation of a region merging segmentation algorithm. In this algorithm the merging order is based on edge value, and the merging predicate exploits recent statistical investigations. A notable acceleration is obtained by exploiting two forms of parallelism, firstly the Data Level Parallelism by processing edges of the same weight in parallel, secondly the Instruction Level Parallelism. Moreover, the classical UNION-FIND data structure is improved by using local registers to reduce the access time of FIND operations. Finally the implementation could be easily tuned to extract textures (object analysis) or all edges (image enhancement).

## INTRODUCTION

Researchers have been working on image segmentation for more than 30 years. Image segmentation is an ill-defined problem, so the optimal solution could not exist, and until now no standards were defined for this field. Nevertheless, many applications could benefit widely from a good segmentation algorithm, for example object oriented compression, pattern recognition, 2D/3D conversion and many others. The image segmentation algorithms could be classified into two categories, namely contour-based and region-based methods. In the first category we find out the significant object boundaries and extract connected components [1]. The main difficulty in this category is to find boundaries closed over objects especially in noisy images. Moreover this approach doesn't benefit from statistical properties of the image. Because of these limitations, the second category, i.e. region-based, is more often used. In these methods, we merge neighbours regions that verify a certain similarity criterion. Two important points define completely a region-based algorithm; the first one is the similarity criterion used to indicate whether two regions should merge or not, the second one is the order in which the similarity test should be done. There is an important gap about the way these two points interact. Many similarity criterions have been used in the segmentation literature, In [2] the most used criterion are reviewed, In [3, 4, 5] some robust criterion are proposed. These similarity criterions are combined with a data structure that establishes an order in similarity test. In [6], they use a tree structure and propose two merging order, "mergesquare", which is claimed as a parallel algorithm, and "scanline", which is sequential. The main drawback of these orders of merging is that they don't depend on the image content, which influence the segmentation result. The region adjacency graph approach (RAG) avoids this drawback. In this approach, we can achieve the best local merge, i.e. every region will merge with the most similar of its neighbours. In [7, 8], a Valued region adjacency graph is computed and decomposed in a set of partial complete graph. The RAGs are also used in pyramidal structure [9, 10, 11]. But RAGs approaches still don't exploit global information of the image. In the implementation point of

view, segmentation algorithms are very computing-intensive. Many works proposed parallel algorithms of segmentation to solve the implementation issue. The irregular pyramids were particularly designed to fit a massively parallel architecture. We can also cite [12] in the scope of parallelizing segmentation algorithms. But all these works don't conciliate the exploitation of global information with the parallelization issue.

In this paper we propose an implementation tending toward this conciliation. We use the algorithm proposed in [13] which combine an order of merging that depends on the content of the image with an adaptive threshold for fusion. We propose an original implementation where the main parts of the algorithm are simplified or vectorized. In the following sections, firstly the algorithm is described, then we propose some implementation solutions where the main steps of the algorithm are vectorized or simplified, and finally we propose a method to tune the algorithm in order to extract textured regions or to extract edges.

## THE SEGMENTATION ALGORITHM

In [13], Nock et al proposed a region-based merging. In this algorithm, they combine a specific order of merge with an original similarity criterion. As far as notations are concerned, let consider an image  $I$ . The notations  $h$  and  $w$  denotes respectively the horizontal and vertical size of the image,  $|I| = h * w$  is the total size of the image,  $a(p)$  is the pixel colour level at position  $p$  and  $g$  denotes the maximum colour level. In the two following sections we explain the order of merging and we present the similarity criterion.

## ORDER OF MERGING

The order of merging is built based on the edges values as in [13, 3]. The idea behind this order of merging is to merge first what is similar before merging what is different.

In our algorithm, an edge corresponds to a couple of pixels  $(p, p')$  in 4-connectivity. The edge values  $v$  correspond to the maximum of the three differences over the three colour components  $\{r, g, b\}$  :

$$v(p, p') = \max_{a \in \{r, g, b\}} (|a(p) - a(p')|). \quad (1)$$

The edges are then sorted in an increasing order of their values and corresponding pixels are treated in this order for fusion.

## THE CRITERION OF MERGING

We use the criterion of merging proposed in [13]. Let's explain briefly how this criterion works. Given two neighbours regions  $s_1$  and  $s_2$ , the average of the three colour components within these regions are denoted by  $\mu_{a_1}, \mu_{a_2}$  with  $a \in \{r, g, b\}$ . The region cardinal of  $s_i$  is denoted  $|s_i|$ . The criterion for merging the two regions is the

following:

$$Pr(s_1, s_2) = \begin{cases} true & \text{if } \Delta\mu(s_1, s_2) \leq g * \sqrt{f(s_1) + f(s_2)} \\ false & \text{otherwise} \end{cases} \quad (2)$$

$$\Delta\mu(s_1, s_2) = \max_{a \in [r, g, b]} (\mu_{a_1} - \mu_{a_2}).$$

The adaptive threshold  $f(s_i)$  takes into account the region size  $|s_i|$  as follows:

$$f(s_i) = \min(g, |s_i|) * \frac{\ln(|s_i| + 1) + \ln(\gamma)}{2 * Q * |s_i|}. \quad \gamma = 6 * |I|^2.$$

This threshold is based on a statistical model of the image and obtained using McDiarmid's inequality, see[13] for more details.  $Q$  is a parameter set by the user that could tune the coarseness of the segmentation.

## IMPLEMENTATION

As shown in Fig.1 the algorithm can be decomposed in three main steps. The first step corresponds to histogramming where the histogram of edges values is computed. This histogram is then used to order edges. In the third step we do the merging following this order of edges. The parallelism in the three steps is not obvious to extract. Indeed the three operations are irregular both in data access from the memory and in computations. In this paper we focus on the vectorization of computation. In this vectorization we process a vector of  $n$  data  $D = [d_1, d_2 \dots d_n]$  in parallel way. In the following we detail the vectorization of the main steps of the algorithm, i.e histogramming, sorting of edges and merging.

### HISTOGRAMMING VECTORIZATION

Let us consider that  $H$  denotes the histogram of edge values that is computed in this step. To compute  $H$ , firstly we compute edges values  $v$  as detailed in equation (1), secondly we compute the distribution  $H$  of these values.

There is no data dependency in the computation of edges values, so we can achieve this operation in vectorial way over a vector of edges  $E = [(p_1, p_2) \dots (p_{2n-1}, p_{2n})]$  which result on a vector of values  $V = [v_1 \dots v_n]$ . However, computing the distribution  $H$  of the edges values in vectorial way, is not straightforward. Indeed two edges values could be equal, and incrementing the histogram's bin corresponding to this value in parallel way will give incorrect result. To solve this data dependency, we propose the following method:

We consider an array  $T$  of  $g$  cells, each cell is  $n$  bits width. Each  $v_i$  in  $V$  set the  $i^{th}$  bit of the  $v_i^{th}$  cell of  $T$ . Then we add the bits of each cell of  $T$  in one instruction. The result in  $T$  is used to update the histogram  $H$ . The algorithm is described in details in Algorithm.1.

From the hardware point of view, this vectorization requires binary adders with  $n$  input, which is very simple.

Let us explain how the histogram  $H$  is used for the sorting step. We consider an array  $M_v$  of size  $h * (w - 1) + (h - 1) * w$  which is the number of edges in the 4-connectivity in the whole image. This array  $M_v$  will be used to store the order of edges. We compute the accumulated histogram  $H_a$  as detailed in equation (3). This  $H_a$  is used to partition  $M_v$  in  $g + 1$  parts, the  $i^{th}$  part is limited between  $H_a[i]$  and  $H_a[i + 1]$  addresses. In this  $i^{th}$  part of  $M_v$  we will store edges with values equal to  $i$ .

### Algorithm 1 Vectorization of histogramming

---

```

for  $i \in [0 : n - 1]$  do
   $T[i] = 0$ 
end for
for  $i \in [0 : n - 1]$  do
   $T[v_i][i] = 1$ 
end for
for  $i \in [0 : n - 1]$  do
   $H[v_i] = H[v_i] + \sum_{j=0}^{i-1} T[v_i][j]$ 
end for

```

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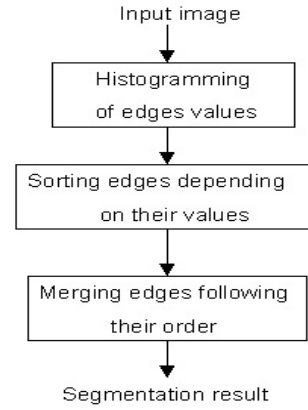


Figure 1. General diagram of image segmentation.

$$\begin{aligned} H_a[0] &= 0; \\ H_a[i] &= H[i - 1] + H_a[i - 1]; \end{aligned} \quad (3)$$

### SORTING VECTORIZATION

In this step, we want to assign to a vector  $E$  of edges a vector of addresses  $A$  where they will be stored in  $M_v$ . There is a data dependency in this step; if two edges have the same value, the assignment of two different addresses to these two edges in parallel way is not obvious. To solve this dependency, we use the same idea detailed in the previous section. Each edge  $E[i]$  with value equal to  $v_i$  set the  $i^{th}$  bit of the  $v_i^{th}$  cell in  $T$ . Then we assign to  $E[i]$  an address  $A[i]$  as detailed in Algorithm.2

### MERGING VECTORIZATION

The algorithm of merging is described in Algorithm.3. This algorithm uses the UNION-FIND data structure. For an edge that corresponds to a couple of pixels  $(p_1, p_2)$ , we use the "FIND" operation to find the couple of segments  $(s_1, s_2)$  containing these two pixels, then the predicate is evaluated for  $(s_1, s_2)$  as described in equation(2). If the predicate is true, we make the "UNION" of  $s_1$  and  $s_2$ . After the "UNION" operation, we compute the new segment properties  $(|s_i|, \mu_{r_i}, \mu_{g_i}, \mu_{b_i})$  and update the main memory with these information. In this paper we focus on the vectorization of the predicate evaluation and the "UNION" operation for a vector  $S$  of couples  $(s_1, s_2)$ . The FIND operation still be hard to parallelize.

Firstly we propose a simplification for the thresholds computation detailed in equation (2). We used a linearization by di-

**Algorithm 2** Vectorization of sorting

```

for  $i \in [0 : n - 1]$  do
   $T[i] = 0$ 
end for
for  $i \in [0 : n - 1]$  do
   $T[v_i][i] = 1$ 
end for
for  $i \in [0 : n - 1]$  do
   $A[i] = H_a[v_i] + H[v_i] + \sum_{j=i}^{j=n-1} T[v_i][j]$ 
end for
for  $i \in [0 : n - 1]$  do
   $H[v_i] = H[v_i] + \sum_{j=0}^{j=n-1} T[v_i][j]$ 
end for

```

**Algorithm 3** Vectorization of sorting

```

for all the edges in the sorted list do
  p1 and p2 are the pixels connected by the edge
   $s1 = FIND(p1)$ 
   $s2 = FIND(p2)$ 
  if  $(Pr(s1, s2) = True)$  then
     $UNION(s1, s2)$ 
  end if
end for

```

chotomy. The linearization provides a Look Up Table. To compute a threshold value  $f$  corresponding to one region cardinal  $|s_i|$ , we read the coefficients  $\alpha$  and  $\beta$  from the LUT, and the computing is  $f(s_i) = \alpha * |s_i| + \beta$ . This trick simplifies the computation a lot without any loss of quality.

Let us explain how the vectorization of the merging step is achieved. We consider a vector of couples of segments  $S$  which is the result of the "FIND" operation applied on a vector of edges  $E$ . We load the vector  $S$  and the data corresponding to each segments  $s_i$  ( $|s_i|, \mu_{r_i}, \mu_{g_i}, \mu_{b_i}$ ) in local registers. Processing the vector  $S$  in parallel way is not straightforward because one segment label  $s_i$  could be equal to another  $s_j$ , and the result of merging will be incorrect as described in the example Fig.3. So the level of parallelism depends highly on the image content.

The vectorization of the merging of  $S$  is done in the following way: Firstly the elements of the vector  $S$  are classified into two parts, the first one contains independent couples  $(s_i, s_{i+1})$  where one label  $s_i$  figures once only, the second part contains dependent couples where one label  $s_i$  figures in another couple in  $S$ . Firstly we process the first part of  $S$  in parallel way. Secondly we process in sequential way the second part. We then investigated the optimal width  $n$  of the vector  $S$  that gives the best data level parallelism (DLP) exploitation. In this investigation, we computed the ratio between the number of operations in the vectorized of merging, over the number of operations in the sequential merging described in Algorithm.3. In Fig.2 we show this ratio for many vector width and for many real sequences. The acceleration is maximal for a vector length around 25.

In addition to the DLP exploited by processing the first part of  $S$  in parallel, the processing of the second part of  $S$  benefits from the locality of data. Indeed, if two couples  $(s_i, s_{i+1}), (s_j, s_{j+1})$  share one label, they are processed sequentially. The latest one will use the result of merge of the first one, which still be available in local

	$n=20$	$n=25$	$n=50$	$n=100$	$n=200$	$n=300$	$n=400$	$n=500$	$n=600$
Bicycle	1.310	1.312	1.308	1.296	1.281	1.273	1.269	1.266	1.264
Boat	1.504	1.510	1.501	1.475	1.454	1.446	1.442	1.439	1.437
Soccer	1.438	1.446	1.443	1.423	1.405	1.398	1.393	1.391	1.389
Sport	1.297	1.305	1.315	1.304	1.286	1.279	1.275	1.273	1.271

Figure 2. ratio between the number of operations of the sequential merging and the merging when exploiting data level parallelism.

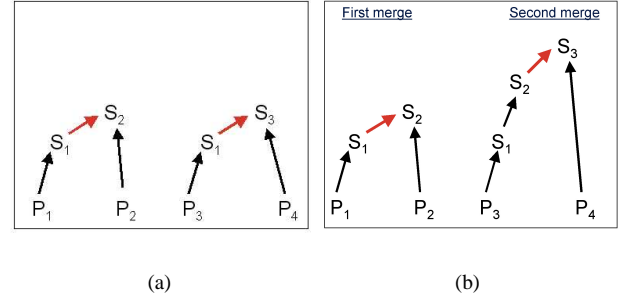


Figure 3. Merging two couple of segments  $(S_1, S_2)$  and  $(S_1, S_3)$ : a.Parallel merge, after merging,  $S_1$  belongs to  $S_2$  and  $S_3$  which are two different segments, so the result is incorrect. b.Sequential merge : after the first merge  $S_1$  belongs to  $S_2$ , after the second merge both  $S_1$  and  $S_2$  belong to  $S_3$ , the result is correct

registers, instead of getting it from the main memory as in the simple sequential merging.

In the other hand when updating segment's properties some of the operations are independent and could be parallelized. Let us consider that  $s_1$  and  $s_2$  are merged, and  $s_2$  becomes the representative of the two segments. The properties of  $s_2$  should be updated. The operations used for this updating are :

$$\begin{aligned}
 sum &= |s_1| + |s_2|. \\
 \mu_{r_2} &= (\mu_{r_1} * |s_1| + \mu_{r_2} * |s_2|) / sum. \\
 \mu_{g_2} &= (\mu_{g_1} * |s_1| + \mu_{g_2} * |s_2|) / sum. \\
 \mu_{b_2} &= (\mu_{b_1} * |s_1| + \mu_{b_2} * |s_2|) / sum.
 \end{aligned}$$

Some of these operations could be executed in parallel way. The whole computing could be done in three steps, in each step we execute the independent operations as shown below :

$$\begin{aligned}
 \text{1- First step:} \\
 o_1 &= |s_1| + |s_2|; o_2 = \mu_{r_1} * |s_1|; o_3 = \mu_{r_2} * |s_2|; \\
 o_4 &= \mu_{g_1} * |s_1|; o_5 = \mu_{g_2} * |s_2|; \\
 o_6 &= \mu_{b_1} * |s_1|; o_7 = \mu_{b_2} * |s_2|.
 \end{aligned}$$

$$\begin{aligned}
 \text{2- Second step:} \\
 o_8 &= o_2 + o_3; o_9 = o_4 + o_5; o_{10} = o_6 + o_7
 \end{aligned}$$

$$\begin{aligned}
 \text{3- Third step:} \\
 o_{11} &= \frac{o_8}{o_1}; o_{12} = \frac{o_9}{o_1}; o_{13} = \frac{o_{10}}{o_1}
 \end{aligned}$$

Therefore, if we have enough resources (6 Multiplier, 3 adder, 3 divider) we can do the updating in 3 operations instead of 12 operations.

## TUNING THE SEGMENTATION

The image segmentation requirements are different depending on the application. In Image enhancement, the main properties to find are edges in order to process pixels belonging to homogeneous regions in the same manner, while in many image analysis applications like pattern recognition, texture extraction is fundamental. We propose a very simple method to switch the segmentation from a texture-oriented to an edge-oriented segmentation. When using the predicate of equation (2), we find out textures. But if we replace this predicate by the one described in equation (4), we will find out all the edges higher than a fixed threshold.

$$P(s_1, s_2) = \begin{cases} true & \text{if } \Delta v(p_1, p_2) \leq tr \\ false & \text{otherwise} \end{cases} \quad (4)$$

$$\Delta v(p_1, p_2) = \max_{a \in [r, g, b]} (a_1 - a_2).$$

Where  $(p_1, p_2)$  is the edge being processed,  $s_1, s_2$  are segments containing  $p_1$  and  $p_2$ ,  $tr$  is a threshold fixed experimentally to 10 for  $g = 255$ . In Fig.4 we show the result of segmentation of one image for the two predicate. In Fig.4(a) we show the result of a textured-oriented segmentation with the first predicate of one textured image. Notice that the textures are well segmented. In Fig.4(b) we show the result of an edge-oriented segmentation by using the second predicate. We can see all the edges in white colour.

## CONCLUSION

Actually we are investigating to build a memory system where data is accessed by content instead of address. With such system we will implement the "FIND" operation efficiently. The solutions proposed in this paper were tested in c language and we are looking for a real hardware implementation.

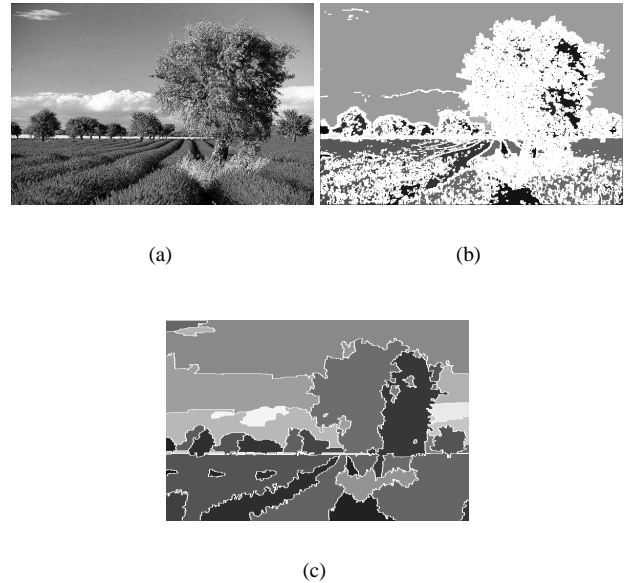


Figure 4. Tuning segmentation: a. The original image. b. An edge-oriented segmentation. c. A texture-oriented segmentation

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