





#### **GPU-**ACCELERATED SNAKE

GPU IMPLEMENTATION OF A REGION-BASED SEGMENTATION ALGORITHM (SNAKE) FOR LARGE IMAGES

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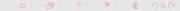




#### Image segmentation

#### Definition, goal

- Dividing an image in two homogeneous regions.
- Reducing the amount of data needed to code information.
- Helping the human perception in certain cases.



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#### Images of our interest

#### Origins

- Synthetic Aperture RADAR (S.A.R.),
- Ultrasonic (medical imaging),
- Photographic (IR, nightshots).

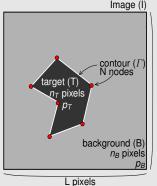
#### Characteristics

- 16 bit-coded gray levels,
- From 10 Mpixels to more than 100 Mpixels,
- Very noisy.





#### Algorithm basics: criterion



H pixels

- The goal is to find the most likely contour  $\Gamma$  (number and positions of nodes).
- The criterion used is a Generalized Likelihood one. In the Gaussian case, it is given by

$$GL = \frac{1}{2} \left[ n_B.log\left(\widehat{\sigma_B}^2\right) + n_T.log\left(\widehat{\sigma_T}^2\right) \right]$$

where  $\widehat{\sigma}_{\Omega}$  is the estimation of the deviation  $\sigma$  for the region  $\Omega$ .

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### Algorithm basics: parameters estimation

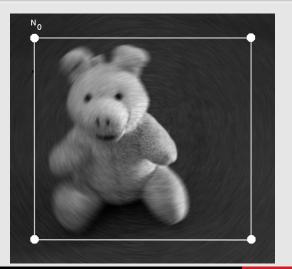
- Based on the Green-Ostogradsky theorem, Chesnaud has shown how to replace those 2-dimensions sums inside the contour by 1-dimension sums along the contour.
- This optimization implies:
  - the precomputation of a few matrices (called cumulated images) containing the potential contributions of each pixel of the image,
  - the use of constant lookup tables of weighting coefficients to determine the *contributions* of each segment of pixels.



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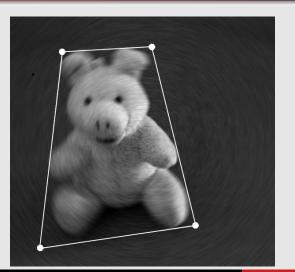




- 15 Mpixels image (SSE implementation limit).
- Initial contour: 4 nodes.





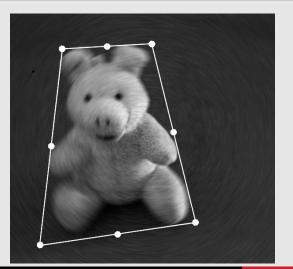


 End of first iteration: no more move can be of interest.

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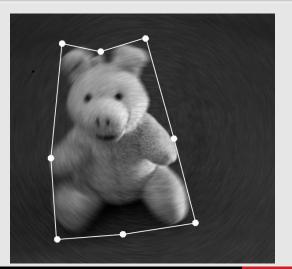


 Nodes added in the middle of segments.

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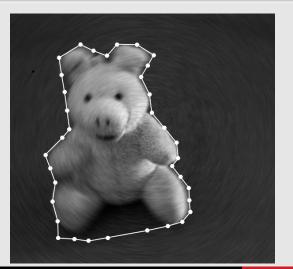


• End of second iteration.

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End of fifth iteration (36 nodes).



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# GPU implementation: prior knowledge

- The parallelism of a modern GPU lays on a SIMT paradigm (Single Instruction Multiple Threads): the same instruction is processed by a great number of threads at a time (up to 2<sup>16</sup>).
- Threads are compounded in independants blocks with no possible synchronization between blocks.
- Threads in a block share a small amount of shared memory (16-48 KBytes).
- There are restrictive conditions to be fullfilled in order to make efficent accesses to global and shared memory.
- Data transfers between CPU and GPU are slow.

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# GPU implementation: precomputations

- One of the cumulated images is not to be computed anymore: values are evaluated on the fly.
- An inclusive parallel prefixsum is performed on each row of the image for each matrix to be processed  $(z,z^2)$ .
- Speedup is around x7 for images larger than 100 MPixels. Comparison is done with the SSE/CPU implementation of the PhyTI group.
- ➡ Higher speedups (x15) are obtained with specific versions for constant image sizes.



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#### GPU implementation: nodes move

To select the possible next position of a node:

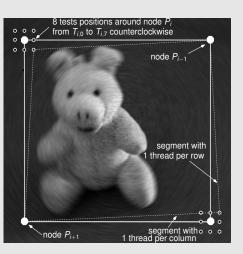
- Parameters of the corresponding contour have to be estimated.
- Then the value of the criterion can be obtained and compared with the previous one.
- The parallelization needs reside essentially in the parameters estimation.
  Two possible parallelism levels:
  - One contour per thread.
  - One pixel per thread.
  - ➤ The one pixel per thread rule is far more efficient, due to memory access constraints.







#### GPU implementation: parallelization



- Every 16 segments for every even/odd nodes are processed in parallel.
- Fits GPU specific parallelism: each pixel is processed by a thread.





The main idea is to organize, in a single array, every pixels of every segments to be processed.

Thus, for a given state of the contour (*N* nodes), we:

- Find the largest segment to be processed. It gives:
  - the block size bs of the computing grid,
  - the number of blocks needed for each segment  $(N_{TB})$ .
- Compute in parallel, the coordinates of every pixels of the 16.N segments to be considered,
- Make some parallel reductions to finally obtain parameters estimation.



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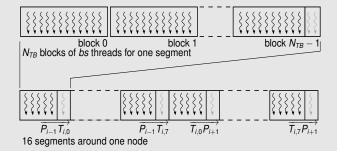
N<sub>TB</sub> blocks of bs threads for one segment

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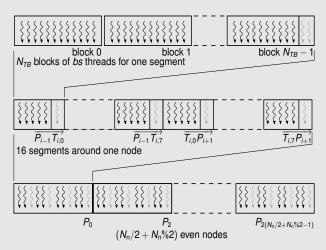
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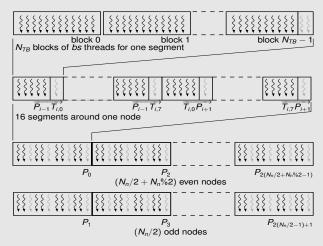


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#### GPU implementation: data structure



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### GPU implementation: first results

- Global speedup around x7-x8 for image sizes from 15 to 150 Mpixels.
- First iterations have higher speedups:
  - several large segments,
  - few inactive threads in the grid.
- Last iterations are sometimes slower than on CPU:
  - a lot of small segments,
  - more inactive threads in the grid.



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#### GPU implementation: smart init (reasons)

- The target shape is often far from initial contour,
- It causes the very first iteration to be much more time-consuming than the other ones.
- Horizontal segments contributions are null.
- Vertical segments contributions computations can be fast, through a specific process.
- >> It's fast to find a rectangle near the target.

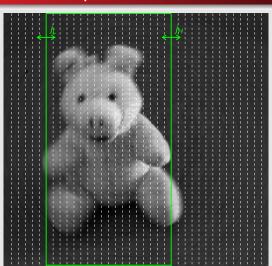


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#### GPU implementation: smart init (process)

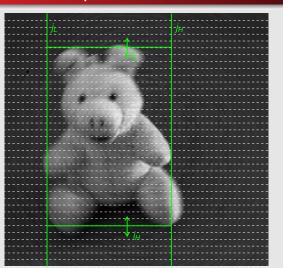


- Realize a periodic sampling of a few hundreds of J-coordinates.
- Evaluate in parallel every possible rectangle of diagonal (0, j<sub>L</sub>) – (H, j<sub>H</sub>).
- Select the one with the best GL criterion.
- j<sub>L</sub> and j<sub>H</sub> are now considered as constants.





#### GPU implementation: smart init (process)



- Given j<sub>I</sub> and j<sub>H</sub>.
- Realize a periodic sampling of a few hundreds of I-coordinates.
- Evaluate in parallel every possible rectangle of diagonal (i<sub>L</sub>, j<sub>L</sub>) – (i<sub>H</sub>, j<sub>H</sub>).
- Select the one with the best GL criterion.

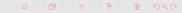
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### GPU implementation: enhancement

- Global speedup around x10 for image sizes from 15 to 150 Mpixels and a small enough target (as in the example)
- Less than 0.6 second for the 150 Mpixels image of this example.



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#### Conclusion, future works

- Interesting speedups
- Original algorithm is not GPU-friendly
- Future works:
  - Finding a more suited structure to describe the contour.
  - Switching to a statistical model independent from a PDF: the potts model.
  - Benefit from recent features of CUDA v4 (overlapping, multiple kernels)
  - Extend to a multiple targets algorithm, based on this single target elementary piece of code.



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