FEMTO-ST





GPU-accelerated snake.

Implementation of a region-based segmentation algorithm (snake).

Gilles Perrot









Image segmentation – Definition, goals

- 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 – Characteristics

- 16 bit-coded gray levels,
- From 1 Mpixels to more than 100 Mpixels,
- Corrupted by additive white Gaussian noise.









- The goal is to find the most likely contour Γ (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 σ for the region Ω .





Algorithm basics – Parameters estimation

• In the Gaussian case, Probability Density Function (PDF) p_{Ω} has two parameters, average μ_{Ω} and standard deviation σ_{Ω} , which are estimated by maximum likelihood.

If z is the gray level of the pixel of coordinates (i, j):

$$\left(\begin{array}{c} \widehat{\mu_{\Omega}} = \frac{1}{n_{\Omega}} \sum_{\substack{(i,j) \in \Omega \\ \sigma_{\Omega}^2}} z(i,j) \\ \widehat{\sigma_{\Omega}^2} = \frac{1}{n_{\Omega}} \sum_{\substack{(i,j) \in \Omega \\ (i,j) \in \Omega}} (z(i,j) - \widehat{\mu_{\Omega}})^2 \end{array}\right)$$

• These estimations have to be computed for each test state of the contour Γ : time-consuming.





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- These estimations have to be computed for each test state of the contour Γ : time-consuming.
- 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 three cumulated images, each one containing a single parameter needed to compute the corresponding pixel's *contribution* to the above sums.







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







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







Nodes added in the middle of segments.







• End of second iteration.



Institut FRESNEL

Algorithm basics – Simplified flowchart





Snake Algorithm



Algorithm basics - Simplified flowchart





Algorithm basics - Simplified flowchart





Snake Algorithm













Algorithm basics - GL criterion computation





Design facts

- Parallelism needs reside essentially in two clearly identified code blocks : GL criterion (60%) and cumulated images (20%)
- Keeping data in GPU memory avoids costly transfers. Thus, the whole computation is being performed by the GPU.
- Parallelism level is set to one thread per pixel of the contour. Alternate choices lead to lower performance (1 thread/segment, 1 thread/contour).
- The innermost loop, among contour nodes, is being parallelized.
- The two other loops are CPU driven. Each loop only requires a single byte of data to be transferred from GPU to CPU at each step.
- The precomputations of two cumulated images are being parallelized. The element values of the third cumulated image are being computed on the fly.





GL criterion – Parallelism level



- 8 test positions around each node *P_i*, denoted *T_{i,0}* to *T_{i,7}*.
- Each test position defines a pair of segments.
- Each combination of test positions defines a contour.
- GPU implementation evaluates in parallel every possible combination of test positions
- Even nodes and odd nodes are moved independently.





GL criterion - Data structure







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GL criterion - Data structure







GL criterion – Parallel computations

- Each active thread corresponds to a pixel of a segment. There are inactive threads.
- The main kernel (kernel1) computes coordinates of every pixels, fetches parameter values from cumulated images and achieves the first reduction stage at block level.
- Another kernel (kernel2) achieves a second reduction stage, providing individual segments contributions.
- A third kernel (kernel3) computes the GL criterion value of each evaluated contour and returns the best.





Focus on kernel1 – Coordinates of pixels

```
global void kernel1(int nb nodes, uint32 npix max, int 1, uint2 *liste pix, bool pairs)
      // indexes of elements
      int blockSize = blockDim.x :
      int tib = threadIdx.x :
      int nblocs noeud = gridDim.x / (nb nodes/2 + (nb nodes%2)*pairs) ;
      int nblocs seg = nblocs noeud / 16 ;
      int id interval = blockldx.x / nblocs noeud ;
      int id segment = ( blockldx.x - id interval*nblocs noeud )/nblocs seg ;
      int tis = idx - ( id interval*nblocs noeud + id segment*nblocs seg )*blockDim.x :
      //in registers
      uint2 p ;
      int dx=x2-x1:
      int dy=y2-y1;
      int abs dx = ABS(dx);
      int abs dv = ABS(dv):
      int nb pix = abs dy>abs dx?(abs dy+1):(abs dx+1):
      int incx, incy ;
      if (tis < nb pix){</pre>
         if (abs dy > abs dx)
          //1 thread per row
          double k = (double) dx/dy;
          p = make uint2(y1 + incy*tis, x1 + floor((double)incy*k*tis+0.5));
        } else {
          //1 thread per column
          double k = (double) dv/dx :
          p = make uint2(y1 + floor((double)(incx*k*tis)+0.5), x1 + incx*tis);
      syncthreads();
```





Focus on kernel1 – Coordinates of pixels

- A Bresenham algorithm is not efficient here to discretize segments into a set of individual pixels. It would generate too much branches in the code.
- As contour is always processed counterclockwise, we use a simpler but more efficient method.
- There are only few branches in the code but they do not lead to any overhead compared to sequential Bresenham.
- Not enough computation to perform. Unable to hide arithmetic operations latency.
- Use of thread registers for better throughout.





Focus on kernel1 – Segments contributions, fetching pixel's parameters

```
//shared memory vectors
extern shared t sum 1 scumuls 1[]
t sum x * scumuls x = (t sum x*) & scumuls 1[CFI(blockDim.x)] ;
t sum x2 * scumuls x2= (t sum x2*) & scumuls x[CFI(blockDim.x)] ;
if ((tis > 0) \&\& (tis < nb pix - 1))
  && ( ((abs dy <= abs dx) && ( ( xprec > p.x) || ( xsuiv > p.x)))
   || (abs dy > abs dx) ))
  //fetch two parameters in the cumulated images, the third is computed on the fly.
  int pos = p.x * l + p.y;
  scumuls 1[ CFI(tib)] = 1 + p.y ;
  scumuls x[ CFI(tib)] = cumul x[ pos ] ;
  scumuls x2[CFI(tib)] = cumul x2[ pos ];
} else {
  //pixel with null contribution and padding in the last block of each segment
  scumuls 1[ CFI(tib)] = 0;
  scumuls x[ CFI(tib)] = 0;
  scumuls x2[CFI(tib)] = 0:
  syncthreads();
```





Focus on kernel1 – Segments contributions, fetching pixel's parameters

- No possible coalescence for global memory accesses as segment's geometry always vary.
- Two ways shared memory bank conflicts exists as shared data is 64 bits-coded.
- But shared memory is still the best choice because of the reduction to be done.





Focus on kernel1 – Segments contributions, first reduction stage

```
uint offset:
#pragma UNROLL
for (offset=1024: offset>32: offset/=2) {
  if (blockSize >= 2*offset) {
    if (tib < offset) {
      scumuls 1[ CFI(tib)] += scumuls 1[ CFI(tib + offset) ];
      scumuls x[ CFI(tib)] += scumuls x[ CFI(tib + offset) ]:
      scumuls x2[CFI(tib)] += scumuls x2[CFI(tib + offset) ];
    }
      syncthreads();
if (tib < 32) {
  #pragma UNROLL
  for (offset=32; offset>0; offset/=2)
    scumuls 1[ CFI(tib)] += scumuls 1[ CFI(tib + offset) ];
    scumuls x[ CFI(tib)] += scumuls x[ CFI(tib + offset) ];
    scumuls x2[CFI(tib)] += scumuls x2[CFI(tib + offset)];
  }}
if (tib == 0) \{
  asombloc[ block[dx,x] = scumuls 1[0] :
  gsombloc[ blockldx.x + gridDim.x ] = scumuls x[0] ;
  gsombloc[blockldx.x + 2*gridDim.x] = scumuls x2[0];
```





Focus on kernel2 – Segments contributions, second reduction stage

```
global void somsom full(uint64 * somblocs, int nb nodes, unsigned int nb bl seg,
                           uint64 * somsom, bool pairs){
//registers
uint64 sdata[3];
unsigned n = nb nodes/2 + pairs * (nb nodes%2);
unsigned int seg = blockldx.x ;
unsigned int nb seg = 16*n :
//1 thread per segment
sdata[0] = 0:
sdata[1] = 0:
sdata[2] = 0;
for (int b=0; b < nb bl seq ; b++){
  sdata[0] += somblocs[seg*nb bl seg + b];
  sdata[1] += somblocs[(seg + nb seg)*nb bl seg + b];
  sdata[2] += somblocs[(seg + 2*nb seg)*nb bl seg + b];
//sums ~ seament contribution ---> alobal memory
  somsom[3 * seg] = sdata[0];
  somsom[3 * seg + 1] = sdata[1]:
  somsom[3 * seg + 2] = sdata[2];
```





Results and analyse

- On images of size between 10 and 150 Mpixels : speedup around x7 compared with SSE2 implementation.
- The first iteration is fast, while the followings are sometimes lower than the reference CPU implementation. It is due to larger segments and thus less inactive threads in the grid.
- Algorithm that do not fit very well GPU architecture.
- Numerous lines of code. For example 200 lines vs 20 lines to compute cumulated images.





Conclusion

- Speedups are not so impressives.
- Another data structure may be more suited.
- A 2D process would be far easier to code, but would not bring such a speedup that the 2D→1D transform brought.
- When designing an algorithm, the targetted host properties should be taken into account.
- Some processes may actually not be suited to GPUs.