

# GPU OPTIMIZER: A 3D RECONSTRUCTION ON THE GPU USING MONTE CARLO SIMULATIONS

## *How to Get Real Time without Sacrificing Precision*

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Abstract: The reconstruction of a 3D map is the key point of any SLAM algorithm. Traditionally these maps are built using non-linear minimization techniques, which need a lot of computational resources. In this paper we present a highly parallelizable stochastic approach that fits very well on the graphics hardware. It can achieve the same precision as non-linear optimization methods without losing the real time performance. Results are compared against the well known Levenberg-Marquardt algorithm using real video sequences.

## 1 INTRODUCTION

Real time simultaneous localisation and mapping (SLAM) consists of calculating both the camera motion and the 3D reconstruction of the observed scene at the same time. This work addresses the 3D reconstruction problem, i.e., obtaining a set of 3D points that represents the observed scene using only the information provided by a single camera.

If the required precision is high, existing reconstruction algorithms are usually very slow and not suitable for real time operation. This work proposes an implementation that can achieve a high level of accuracy in real time, taking advantage of the graphics hardware available in any desktop computer.

This work develops a new 3D reconstruction algorithm based on Monte Carlo simulations that can be directly executed on a modern GPU. The algorithm consists of approximating the maximum likelihood estimator, random sampling from the space of possible locations of the 3D points. Since each sample is independent from others, this method exploits well the data level parallelism required by this programming model.

For validating it, we have compared both precision and performance with the implementation of the Levenberg-Marquardt non-linear minimization algorithm given in (Lourakis, 2004).

## 2 PROBLEM DESCRIPTION

It is assumed that there is an image source that feeds the algorithm with a constant flow of images. Let  $I_k$  be the image of the frame  $k$ . Each image has a set of features associated to it given by a 2D feature tracker  $Y_k = \{\bar{y}_1^k, \dots, \bar{y}_n^k\}$ . Feature  $\bar{y}_i^k$  has  $(u_i^k, v_i^k)$  coordinates. The 3D motion tracker calculates the camera motion for each frame as a rotation matrix and a translation vector  $\mathbf{x}_k = [\mathbf{R}_k | \vec{t}_k]$ , where the set of all computed cameras up to frame  $t$  is  $X_t = \{\mathbf{x}_1, \dots, \mathbf{x}_t\}$ . The problem consists of estimating a set of 3D points  $Z_t = \{\vec{z}_1, \dots, \vec{z}_n\}$  that satisfies the following equation:

$$\bar{y}_i^k = \Pi(\mathbf{R}_k \vec{z}_i + \vec{t}_k) \quad \forall i \leq n, \forall k \leq t \quad (1)$$

where  $\Pi$  is the pinhole projection function. For simplicity, the calibration matrix can be obviated in Equation 1 if 2D feature points are represented in normalized coordinates instead of pixel coordinates.

### 2.1 Proposed Algorithm

A 3D structure optimization method is proposed, that performs a global minimization using a probabilistical approach based on the Monte Carlo simulation paradigm (Metropolis and Ulam, 1949). This approach consists of generating inputs randomly from the domain of the problem. These possible solutions are then weighted using some type of function depending on the measurement obtained from the system. Monte Carlo simulations are suitable for

problems were it is not possible to calculate the exact solution from a deterministic algorithm, i.e., the case of 3D reconstruction from 2D image features, since the direct method is ill-conditioned.

However, this strategy leads to very computationally intensive implementations that makes it unusable for real time operation. One of the key features of these simulations is that each possible solution is computed independently from others, making it optimal for data-streaming architectures, like GPUs.

This system complements the 3D camera tracker presented in (Eskudero et al., 2009) that also runs on the GPU using a similar paradigm.

Every new frame, at time  $t + 1$ , the set  $Z_t$  is enlarged with new points and refined with the new observations  $Y_{t+1}$  provided by the feature tracker, getting a new set  $Z_{t+1}$ . Unlike probabilistic batch methods, the proposed optimizer uses all the available frames for doing this optimization, since the GPU can handle them comfortably. Of course, there is a limit in the amount of frames that the GPU can process in real time. The overall view of the proposed method has the following steps:

1. **Initialize New 3D points.** The algorithm tries to triangulate new 3D points using the feature points provided by the tracker.
2. **Generate Samples from Nnoisy 3D Points.** The system generates new hypotheses about the location of the 3D points using the available 3D structure as initial guess.
3. **Evaluate the Hypotheses.** Hypotheses are evaluated using an objective function that computes the projection residual of all the samples against all the available measurements. The best one is used as new location for the 3D point.

### 3 GPU IMPLEMENTATION

The algorithm is composed by three shader programs. These programs will run sequentially for each point to be optimized. The first shader program will generate all the hypotheses for a single point location, the second shader program will compute the weight of each hypothesis and the third shader program will choose the best candidate among the hypotheses. Algorithm 1 shows a general overview of the proposed method. The parts executed on the GPU have the GPU\_ prefix.

#### 3.1 Data Structures

Since the GPU is a hardware designed to work with graphics, the way to load data on it, is using image

textures. The output is obtained using the render-to-texture capabilities of the graphics card. It is very important to choose good memory structures since the transfers between the main memory and the GPU memory are very slow.

In our case, the hypotheses for the location of a 3D point are stored in a RGBA texture. Each hypothesis has its coordinates stored in the RGB triplet and the result of evaluating the objective function in the alpha channel. Another similar texture is used as framebuffer. Each texel of these textures will be a single hypothesis, so the total number of hypotheses for each point will be the size of the texture squared.

Another RGB texture is used for storing random numbers generated in the CPU. This is because graphics hardware lacks random number generating functions. This texture is computed in preprocessing stage and remains constant, converting this method in a pseudo-stochastic algorithm. Interested readers can refer to (Eskudero et al., 2009) for more details.

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**Algorithm 1:** Overview of the GPU minimization.

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for all  $\bar{z}_i$  in  $Z_t$  do
  SendToGPU( $\bar{z}_i$ )
  GPU_SampleHypotheses()

  for all  $\bar{y}_i^k$  in  $\{Y_1, \dots, Y_t\}$  do
    SendToGPU( $\bar{y}_i^k, \mathbf{x}_k$ )
    GPU_EvaluateHypotheses()
  end for

   $\hat{z}_i = \text{GPU\_GetBestHypothesis}()$ 
   $Z_{t+1} \leftarrow \text{ReadFromGPU}(\hat{z}_i)$ 
end for

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#### 3.2 Initialization

New points are initialized via linear triangulation. This is a very ill-conditioned procedure and its results are unusable, but it is a computationally cheap starting point for the minimization algorithm. This stage is implemented in the CPU since it runs very fast, even when triangulating many points.

#### 3.3 Sampling Points

In this step all the 3D points in the map are subject to be optimized. This stage runs when new points are triangulated and when new frames are tracked. Triangulated points have large error due to ill-conditioned systems of equations, and existing points can be improved with the new measures provided by the feature tracker.

For each point  $\vec{z}_i$ , a set of random samples  $S_i = \{\vec{z}_i^{(1)}, \dots, \vec{z}_i^{(m)}\}$  is generated around its neighborhood. The stochastic sampling function used is a uniform random walk around the initial point:

$$\vec{z}_i^{(n)} = f(\vec{z}_i, \vec{n}_i) = \vec{z}_i + \vec{n}_i, \quad \vec{n}_i \sim U_3(-s, s) \quad (2)$$

where  $\vec{n}_i$  is a 3-dimensional uniform distribution having minimum in  $-s$  and maximum in  $s$ . The parameter  $s$  is chosen to be directly proportional to the prior reprojection error of the point being sampled. In this way, the optimization behaves adaptively avoiding falling into local minimums and handling well points far from the optimum. The GPU implementation is performed using a fragment shader. The data needed are the 3D point to be optimized and the texture with the random numbers. The output is a texture containing the coordinates for all hypotheses. The only datum transferred is the 3D point coordinates, because the the random numbers are transferred in preprocessing stage. It is not necessary to download the generated hypotheses to main memory, because they are only going to be used by the shader that evaluates the samples.

### 3.4 Evaluating Samples

All the set  $S_i$  for every point  $\vec{z}_i$  is evaluated in this stage. The objective function is the residual of Equation 1 applied to every 3D point for every available frame:

$$\operatorname{argmin}_j \sum_{k=1}^t \sqrt{\Pi(\mathbf{R}_k \vec{z}_i^{(j)} + \vec{t}_k) - \vec{y}_i^k} \quad (3)$$

Equation 3 satisfies the independence needed in stream processing, since each hypothesis is independent from others.

Hypotheses are evaluated using a different shader program. This shader runs once for each projection  $\vec{y}_i^k$  using texture ping-pong (Pharr, 2005), avoiding to use loops inside the shader. The only data needed to be transferred are the camera pose and the projection of the 3D point for each frame. This shader program must be executed  $t$  times for each 3D point.

When all the passes are rendered, the output texture will contain the matrix with all the hypotheses weighted. Now there are two ways to proceed. The first one is to download the entire texture to main memory and then search the best candidate using the CPU. The second one is to search directly in the GPU. Experimentally, we concluded that the second one is the best way if the size of the texture is big enough. This search is performed in a parallel fashion using reduction techniques (Pharr, 2005).

## 4 EXPERIMENTAL RESULTS

Both precision and performance of the proposed method have been measured in order to validating it. All tests are executed on a real video recorded in  $320 \times 240$  using a standard webcam. Results are compared with the implementation of the Levenberg-Marquardt algorithm given by (Lourakis, 2004). In our setup, the GPU optimizer runs with a viewport of  $256 \times 256$ , reaching a total of  $2^{16}$  hypotheses per point. The maximum number of iterations allowed to the Levenberg-Marquardt algorithm is 200.

### 4.1 Precision

Various optimizations on triangulated 3D points have been executed to measure the precision of the GPU optimizer. In each run, 25 different points are reconstructed using 15 consecutive frames tracked by the algorithm described in (Eskudero et al., 2009). Figure 1 shows the mean reprojection residual. The figure is in logarithmic scale. This test shows that the GPU optimizer gets on average 1.4 times better results than Levenberg-Marquardt, demonstrating that both Levenberg-Marquardt and GPU optimizer get equivalent results.

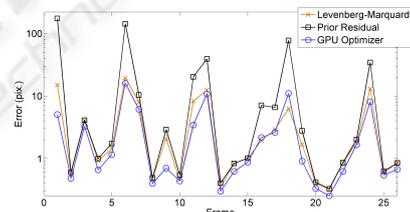


Figure 1: Residual error on real images.

### 4.2 Performance

The PC used for performance tests is an Intel C2D E8400 @ 3GHz with 4GB of RAM and a nVidia GeForce GTX 260 with 896MB of RAM memory. Following tests show the performance comparison between the GPU optimizer and Levenberg-Marquardt. In Figure 2, 15 points are used, incrementing in each time step the number of frames and Figure 3 shows a test running with 10 frames incrementing the number of points in each time step.

Note that both figures are in logarithmic scales. Figure 2 shows that the GPU optimizer runs approximately 30 times faster than Levenberg-Marquardt when the number of frames is increased, being capable to run at 30fps. even when optimizing 15 points over 60 frames.

Next tests analyze deeper the time needed by the GPU optimizer in its different phases. Figure 4 shows

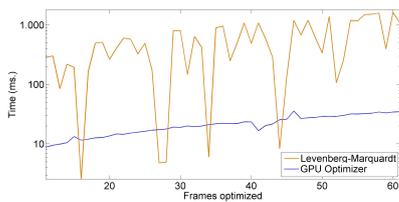


Figure 2: Performance with constant number of points.

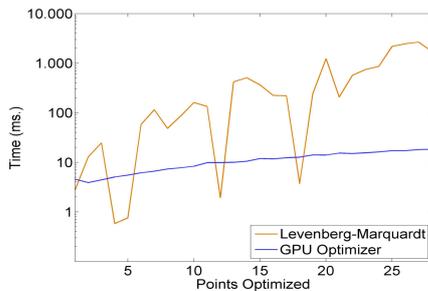


Figure 3: Performance with constant number of frames.

the time needed to run the optimizer with 15 points incrementing the number of frames in each time step, and Figure 5 shows the the time needed when the number of points to optimize is increased in each time step, using always 10 frames.

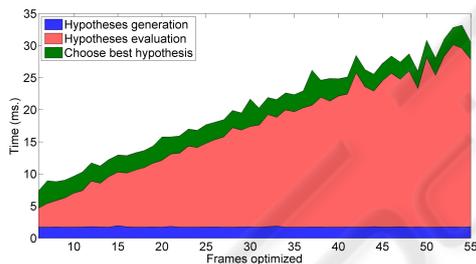


Figure 4: Performance with constant number of points.

From Figure 4 follows that the point evaluation is the only stage that depends on the number of optimized frames. The total time depends linearly on both number of points and number of frames optimized as seen in Figure 5.

## 5 CONCLUSIONS

The proposed GPU optimizer runs a Monte Carlo simulation locally on each point to be optimized, making it very robust to outliers and highly adaptable to different level of errors on the input data.

For validating it, a GPU implementation is proposed and compared against the Levenberg-Marquardt algorithm. Tests on real data show that GPU optimizer can achieve better results than

Levenberg-Marquardt in much less time. This gain of performance allows to use more data on the optimization, obtaining better precision without losing the real time operation. Moreover, the GPU implementation leaves the CPU free of computational charge so it can dedicate its time to do other tasks. In addition, the tests have been done in a standard PC configuration using a standard webcam, making the method suitable for middle-end hardware.

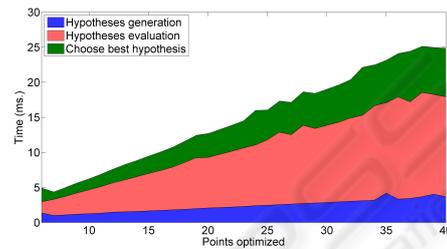


Figure 5: Performance with constant number of frames.

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