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SIMULATING DEPTH PERCEPTION IN VIRTUAL MICROSCOPY

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ABSTRACT

3D datasets production capacity in bioimaging has widely evolved in recent years. This trend also results in a growing demand of more suitable display procedures. In this paper, we propose a new virtual microscopy approach aiming at recovering the third dimension, in order to get closer to analogue microscopes. For this purpose, we rely on multi-view autostereoscopic display with off-centered parallel virtual cameras to ensure 3D perception for a more realistic user experience. Also, this approach handles very large volume data size thanks to an out-of-core data management structure which offers interactive navigation by using complete GPU algorithms.

Index Terms— Virtual microscopy, 3D display, GPU rendering.

1. INTRODUCTION

Nowadays, the volume of 3D datasets tends to increase in nearly all application fields. Such an observation is especially true with the current image acquisition technologies used in biology [1]. Despite the large amount of generated data, efficient visualization tools are needed to improve three-dimensional structures understanding. Among the possible methods available to overcome this problem, virtual microscope is a tool of choice in the biological field.

A virtual microscope could be defined as a system simulating the observation of microscopic samples on a computer. It aims to mimic a conventional microscope, enabling observation, navigation [2, 3], and annotation of virtual slides [4]. The latest virtual microscopy developments [5, 6, 7] have addressed the issue of huge data size. However, as of today, another crucial issue has not been considered: users were frustrated by the availability of only one single plain focus, and the induced loss of three dimensional perception [8]. To improve the user experience quality and to recover the depth perception, we propose to use autostereoscopic display [9] for the 3D visualization [10, 11]. To complete the system, we allow the users to freely navigate and zoom in or zoom out in the whole volume rather than in a single slide.

To address this problem, we rely on an out-of-core data management architecture to handle the large volume of data. In most cases, a resolution level pyramid is created where each level is subdivided into data blocks (bricks). Conversely to tree based structures (like Gigavoxels [12]) that can lead to deep tree traversal to access data, we focus our work on the method proposed by Hadwiger *et al.* [13]. They provide a virtual memory approach with a multi-level, multi-resolution page table mechanism. Validated with a concrete application with interactive exploration of petascale volumes by Beyer *et al.*

[14], this structure offers constant access time between all resolution levels.

Inspired by this out-of-core method, we propose a visualization-driven approach which generates stereoscopic multi-view frames on GPU in interactive time. The aim is to recover the depth perception in a most realistic way with the use of off-centered parallel virtual cameras focused on a shared point of interest.

A brief overview of the system and data access is provided in section 2. After a detailed description of our virtual microscope approach in section 3), experimental results are presented and discussed in section 4. Finally, section 5 provides concluding remarks and perspective works.

2. SYSTEM OVERVIEW

The proposed system fully runs on GPUs [15] and is composed of the virtual microscope and the out-of-core data management. The whole pipeline is called *visualization-driven*. According to the camera position given by the virtual microscope, data is accessed using the out-of-core data manager. We base our approach on the one proposed by Hadwiger *et al.* [13]. The data is represented with a multi-resolution bricked pyramid and addressed using a virtualized page table hierarchy. When a chunk of data is missing in the hierarchy, a brick request is raised and handled by the CPU. Bricks are then loaded from its own cache or a mass storage to the GPU. This method allows to get the data on-demand and to address very large volume of data. The whole process is triggered by the virtual microscope depending on the required data.

Data addressing. The voxel access is performed using a virtual volume representation. A pair $[l, p]$ is composed to get through the virtualization hierarchy. The level of detail is determined by l while $p \in [0, 1]^3$ is the 3D normalized coordinate vector of the requested voxel position. Using normalized coordinate vector allows the use of the hardware trilinear interpolation. Finally, the pair $[l, p]$ is sent to the GPU out-of-core memory manager which sends back three possible answers: the requested voxel itself, an *empty* status if the brick contains no values, or *unmapped*, warning the system that the brick is not in cache and needs to be fetched.

3. VIRTUAL MICROSCOPY

The notion of depth perception is usually lost in conventional 2D virtual microscopy. In this work, we provide an approach to recover this perception somehow available with genuine microscopy of thick slices. To achieve this goal, we use multi-view autostereoscopic displays. Such displays rely on N multi-view images shot according to a specific geometry implying off-centered parallel cameras. The number of views N is directly dependent on the display. In our application, the multi-view image generation follows three steps: *i*)

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The compositing is the final step of the pipeline, providing a final multi-view frame \mathcal{F} . This frame is generated using the N pre-build views \mathcal{V}^i , and is composed by using filtering masks \mathcal{M} on the views (Fig. 4). These masks are binary filters ($\mathcal{M}^i = \{0, 1\}^3 [\mathbb{Z}^2]$), and determine, for all views, which color components will contribute to the final frame composition. This step can be expressed as $\mathcal{F}_c = \sum_i \mathcal{V}_c^i \mathcal{M}_c^i$ where $c \in \{R, G, B\}$. The filtering masks mechanism is dependent of the display. This process may differ, and take more or less computing time, depending on the used hardware.

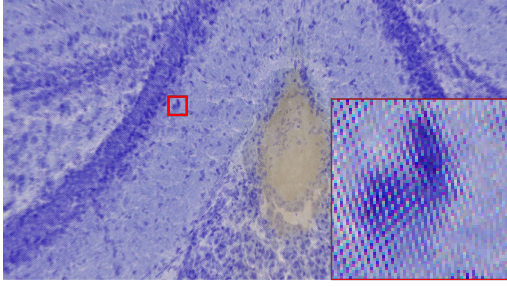


Fig. 5. Multi-view frame generated. Example resulting from the histological volume.

4. RESULTS AND DISCUSSION

The experiments were made using three different datasets:

- (a) A 114 histological slices stack of a mouse brain with a resolution of 64000×50000 RGBA pixels (1.459 TB);
- (b) A $2160 \times 2560 \times 1072$ volume with grayscale 16 bits voxels (11 GB) of a primate hippocampus from a light sheet microscope;
- (c) A 645 blockface slices stack of the mouse brain (a) with a resolution of 823×202 RGBA pixels (428 MB).

The dataset (a) is the proof of concept of our system (Fig. 5). The dataset (b) demonstrates that our system can be applied to different modalities. Finally, the dataset (c) is used to compare the result with the dataset (a), acknowledging the former has a strongest anisotropy than the latter. The display used was a HD (16:10) autostereoscopic 8-views display with a 1920×1200 viewport and requiring a distance – display of 2 m. The same test scenario was applied on the three datasets, and consists of a zoom-in, a pan navigation, then a navigation in the stacks. Table 1 shows the average time recorded to generate a complete multi-view frame.

	(a)	(b)	(c)
Generate frame	11	12	7
Set alpha	8	5	8
Alpha blending	5	5	6
Compositing	5	5	5
Sum	29	27	26

Table 1. Average rendering time in milliseconds of a frame for three different datasets.

Multi-view frame generation. The first noticeable fact is the independence between the frame rendering time and the volume data size or the acquisition modality. Considering the time allowed to other required processes (out-of-core data management) the method offers a real-time navigation with an average of 30 fps. In the presented test, the rendering pipeline was performed at each frame to simulate the worst case scenario in which this needs to be done every time. Nevertheless, in concrete situation, the whole pipeline should be triggered only when a change is detected in the camera position. The first step recorded is the volume projection on the views only. The average time to generate the views was around 11-12 ms. The dataset (c) took less time to project because of its size, as the volume did not fill entirely the views. In fact, on pixels in the outside

borders of the views, we try to project information that is outside the volume. In that case, we are outside the bound of normalized volume sizes and there is no need to continue to process the projection; therefore the frame are generated faster.

The *set alpha* step consists in computing the Euclidean distance between the voxels and different clusters. In the tests, we used five clusters computed beforehand using a k-means algorithm. The choice in the classification algorithm is of importance: the computation runtime can affect the time to render a view, therefore the navigation fluidity. The difference noticed in the results is related to the data type differences between the volumes. While the dataset (b) uses grayscale pixels, the others use RGBA pixels. Therefore, the pixels assignment is performed faster on (b) than (a) and (c).

Depth perception. As stated in section 3.1, the value Δ_z is of importance in the proposed method, as a non optimal value may induce an uncomfortable visualization. During the tests, we noticed that the differences between two slices were small in the dataset (b). The depth perception was recovered by using $\Delta_z = 3$. However, this also underlines that visualizing a grayscale works better with high-contrast or color data. An extra step may be required to colorize the data via LUTs. The extreme case happened with the dataset (a). As expected, the volume anisotropy was significant and required to use a $\Delta_z < 1$. Depending on Δ_z value, the number of slices the user can see in a multi-view frame may be drastically reduced, however the final result is softened. Having $\Delta < 1$ is possible because we are addressing the data using normalized coordinates. The used out-of-core data structure relies on the GPU texture memory and allows trilinear interpolations. However, misusing this value may reduce the depth effect and in turn make the approach irrelevant. Yet, the data anisotropy is an important factor to consider when using this approach.

Camera positioning. In the given approach, the views are positioned on the same plane as the slices. This could be improved using an orbital camera. One could be able to analyse the volumes in a different view angle. However, the data anisotropy would be a major concern and would require a better alternative than simply using a Δ_z parameter.

To test our method, an autostereoscopic display was used, where the user has to move in front of the display to move inside the multi-view frame. With current technologies, it could be interesting to use the same approach on tablet devices, as it may be more convenient for the user to move the device left and right than to move in front of a screen. Finally, all tests were performed and assessed by the authors. The method is ready for statistical studies with the end-users, using lightsheet or histological datasets and using different parameters.

5. CONCLUSION

We introduced a proof of concept to virtualize a microscope using off-centered parallel cameras and an autostereoscopic display. With the help of current GPUs, this approach allows the user to interactively navigate through multi-resolution images. This allows data visualization similar to an analogue microscope visualization by recovering the depth perception. Taking advantage of the out-of-core data management, the proposed approach works independently of the data size, from MB to several TB. In addition, the concept offers a similar user experience across different volume modalities, without significant differences in rendering time. In the future, the development of an approach offering the ability to use an orbital cameras would be of great benefits to this method.

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