# Vaa3D: an extendible and versatile open-source tool for 3D visualization-assisted analysis of large-scale bioimages

#### 8<sup>th</sup> Advanced School on Scientific Visualization @CINECA 14 - 18 October 2013 CINECA - Bologna

#### speaker Ing. Alessandro Bria

PhD student at University of Cassino and L.M., Cassino, Italy Collaborator at University Campus Bio-Medico of Rome, Rome, Italy Collaborator at Allen Institute for Brain Science, Seattle, WA, USA Collaborator at Radboud University Nijmegen, The Netherlands Member of the Projectome project at ICON foundation, Florence, Italy





#### The context: bioimage informatics (1/2)

- using computational techniques to analyze (= extract useful information from) multidimensional **bioimages** at molecular, cellular or systemic scale, e.g. :
  - high-content screening (or *visual screening*) for drug discovery
  - cells segmentation
  - mapping brain circuits





#### The context: bioimage informatics (2/2)

- automated microscopes and the increase in resolution has led to bioimage data explosion
  - terascale has become a reality
- need of automatic processing
  - fully- or semi-automatic?
  - human intervention might be needed
    - post-processing proofreading
    - semi-automatic analysis

2-5D visualizationassisted analysis





# The goal: visualization-assisted analysis of large bioimages (1/2)





# The goal: visualization-assisted analysis of large bioimages (2/2)





- free and/or open-source visualization tools
  - Voxx, OME, ImageJ, Icy, ilastik, CellProfiler, CellOrganizer, CellExplorer, FARSIGHT, Bisque, BrainExplorer, BrainAligner, **3D Slicer**, ParaView
- commercial tools
  - Amira (VSG), Imaris (Bitplane), ImagePro (MediaCybernetics), Neurolucida (MBF) Bioscience)
- standalone 3-5D visualization-assisted analysis of large images not feasible with any of these tools at present
  - - large  $\neq$  terascale
    - missing 3-5D visualization
    - low versatility: supported image formats, cross-platform, etc.
    - low extendibility: how many available plugins? Are they easy-to-write?
    - high memory requirements: both system RAM and GPU RAM

## Vaa3D<sup>(1)</sup>: enjoying working with 3D image data!

"Vaa3D is designed expressly for working with 3D volumetric data and is built on an efficient 3D renderer that allows real-time visualization and manipulation of multigigabyte—sized data on a standard computer. [...] it may not be as **fun** as 3D gaming but Vaa3D promises to make working with 3D image data in the lab much more **enjoyable**"

Daniel Evanko, "Connecting the dots in 3D", Nature Methods highlights, 2010

developed and under development at Peng Lab



<sup>(1)</sup> Peng, H. et al, "V3D enables real-time 3D visualization and quantitative analysis of large-scale biological image data sets", Nature Biotechnology 28, 348-353, 2010







- 3D color image stacks
  - Tiff stack (.tif, .tiff), Zeiss LSM (.lsm), MRC (used for electron microscopy images) (.mrc), Vaa3D's raw file (.v3draw, .raw)
  - any bioimage format supported by LOCI Bioinformats Java library (using the Vaa3D-bioformats plugin)
- 5D time series of color image stacks
  - each time point saved as a separate file (end with suffix like 000.tif, 001.tif, ...)
  - each time point saved as a single slice of a 3D image stack of whatever formats Vaa3D supports (e.g. tiff, or Vaa3D's raw)
- *3D irregular shaped surfaces*: Wavefront .OBJ files, Vaa3D's surface format (.v3ds)
- *3D point cloud*: .apo file (a simple CSV format with fixed number of columns)
- 3D landmarks: .marker (indeed a simple CSV format), .csv







### Vaa3D: surface rendering and creation







#### Vaa3D: neuron editing





### Vaa3D: visualizing 5D data

• On-air demostration





### Vaa3D: 2-mouse click 3D pinpointing (1/2)





#### Vaa3D: 2-mouse click 3D pinpointing (2/2)



- be A and B the two non-parallel rays generated at two viewing angles, corresponding to 2-mouse clicks
  - each click defines a ray through the current cursor location orthogonal to the screen
- a marker is created at the point in space for which the sum of its Euclidean distance to A and B is minimal
  - robust to inaccuracy in the user's 2D clicks



### Vaa3D: 1-mouse click 3D pinpointing (1/2)







#### Vaa3D: 1-mouse click 3D pinpointing (2/2)



- the most probable location on A is estimated by applying the mean-shift algorithm on the intensity distribution
  - the initial center of mass (CoM) p<sub>1</sub> is computed along the whole ray [0, R<sub>1</sub>] intersecting the volume
  - the CoM is repeatedly reestimated by using progressively smaller intervals around the proceeding CoM until convergence
- can be used for quick manual cellcounting or for quantitatively profiling the voxel intensity along the straight line segment connecting two markers

#### Vaa3D: semi-automatic neuron tracing (1/2)





#### Vaa3D: semi-automatic neuron tracing (2/2)



- searching the "optimal path" connecting a set of markers
  - voxels are considered as graph vertexes
  - edges connect each pair of adjacent voxels
  - edge weight is the inverse of the average intensity of the two voxels
  - Dijkstra's algorithm is used to find the least-cost path between pairs of markers
- individual segments are defined as paths between markers and branching points

#### Vaa3D: 1-stroke 3D curving

🕷 v3d64 File Image/Data Process Visualize Plug-In Window







#### Vaa3D: built-in plugins

Vaa3D plugin creator

Vaa3D plugin creator celegans atlasquided seganno celegans straighten data IO bioimagelO Bioformat data type 5D stack Converter datatype converter Convert Image to AtlasViewMode image blending blend multiscanstcks image edge detection edge of maskimg image filters Distance Transform Gaussian Filter Grevscale Distance Transform minMaxfilter image geometry montage image sections recenterimage Rotate Image image registration ssd registration ⊟ image resolution XYZ Resolution image ROI

ROI Editor

image stitching ifusion istitch Map View image thresholding Simple adaptive thresholding linker file Linker File Generator movies Simple Movie Maker neuron utilities Enhanced SWC Format Converter Global Neuron Feature Resample SWC Sort SWC SWC to Maskimage pixel intensity Canvas Eraser Change Single Pixel Value principal skeleton detection Principal Skeleton Detection reference extract refextract Vaa3D PluginInterface Demos 3D viewer data push and display Plugin Call Each Other Mouse Event Monitor Multi Image Interface Single Image interface

#### a plugin to create plugins!

# ON A

#### importing ANY bioimage format

image filters



image registration

image stitching

movie maker

### Vaa3D: a tutorial for developing a plugin (1/8)

- Setting up the development environment under Linux / MacOS
  - get and build (or install, if available for your platform) Qt 4.7.2 (or .3, .4)
  - get and build Vaa3D source code by following instructions at <a href="https://code.google.com/p/vaa3d/">https://code.google.com/p/vaa3d/</a>
  - get and install Qt Creator (optional)
- Setting up the development environment under Windows
  - download the precompiled binaries Qt 4.7.2 for Visual Studio 2008
  - check the prerequisites at <a href="https://code.google.com/p/vaa3d/">https://code.google.com/p/vaa3d/</a>
  - get and build Vaa3D source code by following the build instructions for CMake at the Vaa3D's google code webpage. <u>Use Visual Studio 2008</u> as both project generator and compiler.
  - get and install Qt Creator (optional)



#### Vaa3D: a tutorial for developing a plugin (2/8)

• Using *Vaa3D plugin creator* for creating a Qt project of a new plugin





#### Vaa3D: a tutorial for developing a plugin (3/8)

• myplugin.pro





#### Vaa3D: a tutorial for developing a plugin (4/8)

myplugin\_plugin.h





### Vaa3D: a tutorial for developing a plugin (5/8)







#### Vaa3D: a tutorial for developing a plugin (6/8)

#### • V3DPluginCallback (1/3)

#### ////invoke a Vaa3D plugin function

#### ////get opened images

//obtain a list of all currently opened images
virtual v3dhandleList getImageWindowList() const = 0;

//obtain the \*current\* selected image window, defined as the tri-view window currently selected in Vaa3D main window
virtual v3dhandle currentImageWindow() = 0;

//obtain the \*current\* selected image window, defined as the currently being operated 3D viewer
//curHiddenSelectedWindow may not be the \*currentImageWindow\* if the selection is done from a 3d viewer
virtual v3dhandle curHiddenSelectedWindow() = 0;

#### ////set computed image content/result

//create a new image window for returning some computed image content virtual v3dhandle newImageWindow(QString name="new\_image") = 0;

//directly output computed image content to an existing image window. The size of the window will be changed
//automatically.
virtual void updateImageWindow(v3dhandle image\_window) = 0;



#### Vaa3D: a tutorial for developing a plugin (7/8)

#### • V3DPluginCallback (2/3)

#### ////manipulate image names

```
virtual QString getImageName(v3dhandle image_window) const = 0;
virtual void setImageName(v3dhandle image_window, QString name) = 0;
```

#### ////access the actual 4D image data structure

```
virtual Image4DSimple * getImage(v3dhandle image_window) = 0;
virtual bool setImage(v3dhandle image_window, Image4DSimple * image) = 0;
```

#### ////access the 3D landmark list defined for an image

```
virtual LandmarkList getLandmark(v3dhandle image_window) = 0;
virtual bool setLandmark(v3dhandle image_window, LandmarkList & landmark_list) = 0;
```

#### ////access the 3D region of interest (ROI) defined for an image

```
virtual ROIList getROI(v3dhandle image_window) = 0;
virtual bool setROI(v3dhandle image_window, ROIList & roi_list) = 0;
```

////access the 3D curve, 3D curve set, and 3D reconstructed neuron structure for an image

```
virtual NeuronTree getSWC(v3dhandle image_window) = 0;
virtual bool setSWC(v3dhandle image_window, NeuronTree & nt) = 0;
```



#### Vaa3D: a tutorial for developing a plugin (8/8)

#### • V3DPluginCallback (3/3)

#### ////operating (open/close) the rendering windows

```
//open and close a global 3D viewer
virtual void open3DWindow(v3dhandle image_window) = 0;
virtual void close3DWindow(v3dhandle image_window) = 0;
```

//open and close a local 3D viewer virtual void openROI3DWindow(v3dhandle image\_window) = 0; virtual void closeROI3DWindow(v3dhandle image\_window) = 0;

#### ////Data pushing functions

```
//update the surface objects currently being displayed in a 3D viewer
virtual void pushObjectIn3DWindow(v3dhandle image_window) = 0;
```

//update the content in a 3D viewer directly virtual void pushImageIn3DWindow(v3dhandle image\_window) = 0;

```
//update the time point of a 3D viewer if it is displaying 5D (temporal) data
virtual int pushTimepointIn3DWindow(v3dhandle image_window, int timepoint) = 0;
```

#### ////direct pointers to Vaa3D internal data structure

virtual View3DControl \* getView3DControl(v3dhandle image\_window) = 0; virtual View3DControl \* getLocalView3DControl(v3dhandle image\_window) = 0; virtual TriviewControl \* getTriviewControl(v3dhandle image\_window) = 0; Direct access to Vaa3D core! To be used with care.



# Vaa3D: implementing an example plugin

• On-air demostration





#### Moving towards terascale bioimages: The Projectome Project

- High Performance Computational Infrastructure for processing and visualizing neuro-anatomical information obtained using CLSM
- Whole mouse brain 3D imaging with Confocal Light-Sheet Microscopy:
  - micrometer resolution
  - cm-sized specimen
  - TeraVoxel-sized dataset



International Center of Computational Neurophotonics





#### Moving towards terascale bioimages: CLSM microscopy



- neurons and other structures are selectively labeled with a fluorescent protein
- the specimen is optically cleared and fixed
- the specimen is illuminated by a thin sheet of light and the fluorescence emission is observed from the scanning axis perpendicular to the illumination plane
- translations of the light sheet along the scanning axis produce a stack of 2D slices
- the field of view of the confocal microscope is limited, so translations of the system along V, H axes are needed to produce different overlapping stacks



#### Moving towards terascale bioimages: TeraStitcher (1/3)

- fast 2D approach to align adjacent stacks
- efficient use of memory resources (< 2GB of memory peak)</li>
- the stitched volume is saved into a multiresolution representation suited for further processing



## Moving towards terascale bioimages: TeraStitcher (2/3)

#### results

	Tool	Displ. comp.	Merging	Total	Memory occ.
	Fiji	5.4	8	13.4	800 <i>MB</i>
	XuvStitch	3.5	11	14.5	320 мв
┣	iStitch	9	2	11	600 мв
	Our tool	2.2	1.7	3.9	200 мв

#### the Vaa3D's built-in stitching plugin

Gvoxel	Displ. comp.	Merging	I/O	Total	Memory occ.
63,25	23	37	157	217	821 <i>MB</i>
126,50	68	77	318	433	1132 мв
198,78	109	123	539	771	1132 мв
1315,04	n.a.	n. a.	n. a.	6050	2450 мв







#### Moving towards terascale bioimages: TeraStitcher (3/3)

C TeraStitcher plugin File Options Help							
Importing Aligning Merging	000 Sel	lect volume's directory		-			
Import form	Look in:	/home/utente	e] (	0 0	<b>6</b>		
Re-import (check this to re-scan all acquisition file)	Comp	Name 🔓	* Size	Type Folder	Date Modif 31/05/ 16		
Volume's informations	-	Filiapp         Fol           globusconnect-1.6         Fol           immagini         Fol           ikDDS         Fol           JoVE-Captures         Fol           Musica         Fol           Projects         Fol	Folder Folder	07/06/ 09 06/06/ 17			
Absolute path:			Folder Folder	31/05/ 16 06/06/ 10			
Stacks dimensions (voxels);			Folder	12/06/ 14 31/05/ 16			
Voxel's dims (µm):			Folder	31/05/ 16 05/06/ 10			
Origin (mm):		Scaricati		Folder	31/05/ 16 05/06/ 16		
Stitch test: Slice 0/0		Scrivania Fr	Folder	31/05/ 16 32/02/ 00 -			
	Directory:	100					
	Files of type	Directories			Cancel		
What's this? Move the mouse over an object and its descrip	ion will be displaye	d here.		1			
Ready				-			
1			Start Wiscop	1			



#### TeraFly: overview (1/3)

- Vaa3D's 3D rendering cannot handle very large 3D images
  - e.g. 1 GigaVoxels images require at least a video card with 1 GB of dedicated memory
- **TeraFly** extends the Vaa3D software to cope with (potentially) **unlimited** sized bioimages even on laptops with a limited amount of system memory ( $\leq$  4 GB) and video card memory ( $\leq 1$  GB)
  - easy zoom-in/out with mouse-scroll
  - 4D supported (5D support work in progress 🚸 )
  - automatic scaling of 3D markers and 3D curves throughout 3D navigation
  - annotation of 3D objects
  - basic caching (advanced caching work in progress **(**)
  - separate translations along X, Y, Z
  - separate threads for GPU and I/O
  - fast zoom-in by interpolation + subsequent refinement by image slicing (web-like)





#### TeraFly: overview (2/3)

- the underlying idea is to mimic the behavior of **Google Earth** 
  - what you see is what you need (WYSIWYN)
  - multiresolution representation
  - mouse scroll for zoom-in/out





#### TeraFly: overview (3/3)





#### TeraFly: multiresolution representation



- the volume is saved in tiled format at different resolutions i = 0, ..., k
  - *i*-th resolution is obtained by dividing *i*-1-th resolution by 2, that is equivalent to divide the original resolution by  $2^i$
  - k is chosen so as the k-th resolution has size
     <100 MegaVoxels, thus it can be easily handled</li>
     by the Vaa3D renderer
  - e.g. for a ~1 TeraVoxels volume whose size is  $10.000 \times 10.000 \times 10.000$
  - k = 5 and the 5-th resolution has size ~29 MegaVoxels
- tiles dimensions is typically in [256, 512]



#### TeraFly: architecture (1/2)





### TeraFly: architecture (2/2)





#### TeraFly: flying through the brain on a laptop

On-air demostration

easy zoom-in/out with mouse scroll

fast zoom-in by interpolation + 1-step refinement

separate threads for GPU and I/O

basic caching





# TeraFly: zoom-in method (1/3)



- 3D exploration starts with a precomputed 3D image of the whole volume at low resolution
  - the first time a multiresolution volume is imported into TeraFly, the resolution that is best suited for the computer hardware capabilities is chosen and saved in a fast-to-load format (vmap.bin)
  - the low-res volume map so obtained will be used for starting the 3D exploration every time the user will open the volume
  - usually has size < 100 MegaVoxels</li>



- zoom-in is triggered when mouse scroll exceeds a fixed threshold
- since the Vaa3D renderer zoom-in is center-based, we look at the center of the viewport
  - random 1-click pinpointing actions are triggered around the center of the viewport
  - the majority of markers is created on the foreground tissue/cells
  - we take the centroid as the center of the next higher-res view
  - the VOI is defined using the view size (can be set by the GUI)

# TeraFly: zoom-in method (3/3)



- data I/O starts asap in a different thread
- meanwhile, VOI is extracted from the current view by interpolation and passed to the next view
- the interpolated VOI is shown in Vaa3D (GPU thread)
- meanwhile, data I/O ends and a signal is emitted
- the GPU thread catches the signal and triggers an update in the current view with the high res data just loaded

#### TeraFly: zoom-out method and basic caching



- zoom-out is triggered when the mouse scroll down exceeds a fixed threshold (can be tuned in the GUI)
- the current view is hidden and the previous view is restored
- the higher res view just hidden is maintained in memory for basic caching
  - when zooming-in again, the cached view is simply restored if the overlap between its VOI and the requested VOI is above a certain percentage (can be tuned in the GUI)

#### TeraFly: image slicing (1/2)

#### On-air demostration

fast zoom-in by interpolation + n-step refinement (by *image slicing*)





# TeraFly: image slicing (2/2)



- depending on hardware speed, it might be convenient to use *image slicing* so as to load the first chunk of high res data and display it asap
- convenient when image updates are very fast and I/O is quite slow (tradeoff)
- the optimal number of steps should be automatically detected given the hardware specs

#### TeraFly: annotating 3D objects (1/4)

• On-air demostration

automatic scaling of 3D markers and 3D curves throughout 3D navigation

annotation of 3D objects

separate translations along X, Y, Z





# TeraFly: annotating 3D objects (2/4)

Octrees











### TeraFly: annotating 3D objects (3/4)



- we use a point region (PR) octree for storing 3D markers and 3D curves
  - the node stores an explicit 3dimensional point, which is the "center" of the subdivision for that node
- compact representation
- fast search
- for 3D curves, additional linking between nodes is introduced for loading whole segments

# TeraFly: annotating 3D objects (4/4)

3D View [E:/TOMO_250611/V_000_MULTI_RES/adv2.ano]			
		Controls Volume Surf/Object Others V Markers Sync Tri-view Lobel Size x15 Surfaces Load/Save Sur Stretch with Volume	Dbjs \$
	Object Manager		8 2
3	ne Structure (1) Point Cloud (97) I on/off color x 1 V #a05dae 1016.36 5	Point Cloud Set (1)	elect All
	2 2 #9e08c7 3176.33 6	643.77 1373.85 0 Sere	a inverse
	Object no 5 Name  a neuron Comments		
4th	Other properties		
	Cancel		ЭК

- markers are saved into 3D point cloud files (.apo)
- curves are saved into SWC files (.swc)
- a link file (.ano) is automatically generated for grouping heterogeneous 3D objects and annotations
- .ano files can be simply drag-anddropped into Vaa3D or loaded by

TeraFly on the image



# TeraFly: I/O and the importance of the file format (1/3)



- tiled format pros:
  - small files
  - can copy a subvolume (volume slicing) by simply copying directories
  - when a VOI is requested, only the tiles intersecting the VOI are involved

## TeraFly: I/O and the importance of the file format (2/3)



- in the beginning, TeraFly could handle only the "Image series (tiled)" format
- for very large volumes along X and Y, this led to a huge number of slice files
  - 10.000 slices × 50 × 50 stacks= 25 millions of files!
  - almost impossible to move
  - slow to access data: need to open thousands of files



### TeraFly: I/O and the importance of the file format (3/3)



- TeraFly now uses the "Vaa3D raw (tiled, 4D)" format
- blocks along Z instead of slices!
  - 10.000 slices × 50 × 50 stacks
     = 50.000 files with blocks containing each 500 slices
  - each block is a 3D singlechannel Vaa3D raw (random access, "almost 10 times faster than TIFF")
  - one volume per channel (i.e. channels is the 4<sup>th</sup> dimension)



TeraConverter						
-Step 1: Import volume from:						
Image series (tiled)	E:\TOMO_250611\V_000_MULTI_RES\R	ES(14261x6814x7828) Browse for dir				
Step 2: Convert volume to:						
Vaa3D raw (tiled, 4D)		Browse for dir				
Image series (tiled) Image series (nontiled) 3D TIFF (tiled) 3D TIFF (nontiled) Vaa3D raw Vaa3D raw (tiled, RGB) Vaa3D raw (tiled, 4D) Vaa3D raw (series)	Select       Resolution (X ×         ♥       14261 × 6814         ♥       7130 × 3407 ×         ♥       3565 × 1703 ×         ♥       1782 × 851 ×         ♥       891 × 425 ×         ♥       445 × 212 ×	Y × Z) Size (GVoxels)     × 7828 708.440     × 3914 88.549     × 1957 11.065     × 978 1.381     < 489 0.172     × 244 0.021				
Stacks dims:	256 (X) 🚔	256 (Y) 💂 256 (Z)				
Estimated RAM usage:	2.896 GB					
Three-leveled folder structure (first level for channels, other two levels for tiling X vs Y) with each tile composed by a series of 3D blocks stored into Vaa3D raw files containing a single channel.						
Ready to convert volume.						

- for the conversion of terascale volumes from one format to another
  - Image series (tiled / nontiled): any image format is supported (tiff, png, jpeg, bmp, etc.)
  - 3D tiff 🚸
  - Vaa3D raw (single file / tiled with blocks / series), RGB or 4D
- RAM usage estimation
- selection of the resolutions to be produced

- Vaa3D is a free, open-source, cross-platform, extendible and versatile tool for visualizing and analyzing 3-5D bioimages on workstations and even on laptops
- the existing standalone tools, both free and commercial, still cannot deal with terascale images and/or do not embed such a powerful and user-friendly 3Dvisualization-assisted analysis of bioimages
- TeraFly enables Vaa3D to handle terascale 4D images, thus making it possible to *fly through* terabytes of images almost instantly and even on laptops
- thanks to TeraConverter, TeraFly is also independent from the file format





