# WIEN Informatics

## Visualization and Interaction Techniques for the **Exploration of the Fruit Fly's Neural Structure**

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Synapses are located at a neuron's arborisation. Where two or more neurons' arborisations overlap, they display a potential connectivity.

Neurobiologists use the information on potential connectivity to formulate new hypotheses and experiments. Visualising and computing overlaps between more than two arborisations (higher order overlaps) is difficult.



#### **Data Visualisation**

With the goal to closely imitate the design, we chose to combine a variety of non-photorealistic rendering techniques.



The 3D visualisation is a composite of five separate renderings. Among those are a silhouette rendering, a watercolour shader for arborisations, and a transparent rendering for optional

developed a beautiful novel design in cooperation with the neurobiologists at the IMP in Vienna.

### **Goals of the Thesis**

- Realise the novel design
- Compute volumes of higher order overlaps on demand
- Provide interaction techniques to communicate the data

#### **Computing Intersections**

A major contribution of the thesis is the algorithm to compute mesh volumes and mesh intersection volumes on the GPU. We use A-Buffers on the GPU—which are typically used to achieve order-independent transparency—to compute volumes. A framebuffer typically stores a single colour value per screen pixel. The A-Buffer, however, can record multiple values in a single screen pixel by allocating linked lists.

The graph below illustrates a simplified version of our algorithm, three passes on the GPU are sufficient. The first pass renders meshes and stores depth



The silhouette of the brain

neural structures. All these serve as context for the point of interest, the overlaps. These intersections of arborisations are rendered in bright colours (yellow pair-wise overlaps, orange shades for higher order overlaps). The visualisation shows overlaps on demand. They are abstracted as interactive glyphs. The glyphs (small circles) can be used to select/show overlaps or blend in the computed volumetric data. The shaders for overlaps and their glyph abstractions use A-Buffers on the GPU, the same data structure The final 3D visualisation shows a pair-wise overlap used for their volume computation (compare the box on the right).



values (pictured as red dots on the right).

screen frustum

render first A-Buffer meshes

sorting

The rendering pipeline does not guarantee the order of fragments in each pixel. Thus, a compute shader sorts the depth values from the first A-Buffer and writes to a second A-Buffer.

The second A-Buffer contains depth values sorted by depth. In the third and final pass, a compute shader iterates the linked list in each pixel to find depth values relevant for a specific volume.



The examples on the left illustrate the depth differences in two dimension. In the simplest case we compute depth differences for a single mesh For a pair-wise overlap the algorithm finds the depth values of the intersection volume.

The depth differences stored in the final A-Buffer can be summed up on the CPU to compute a volume.

by depth

A-Buffer

Visual design by Judith Moosburner

second A-Buffer

compute depth differences



menus that display the computed volumetric data. Tooltips for glyphs are directly embedded into this 3D view. A separate menu in the viewport provides, in addition to the volumetric data, settings for the visibility of neuronal structures.





Arborisations and their colour palette

Detail: a tooltip of a 4-overlap with enough volumetric data to judge the importance of the overlap

#### Conclusion

• Features described by the design successfully implemented

• Very fast on demand computation of intersection volumes on the GPU

 Visual and quantitative investigation of higher order overlaps of arbitrary order

 Integrated with the BrainGazer framework used by neurobiologists • Interactive links with other tools in Brain-Gazer

final A-Buffer

