
Isosurface Computation and Rendering Tool

Vishal Peter, Srishty Suman
M. Tech, CSA

1 Problem Formulation

For 2D or 3D meshes that represent the domain of a continuous function of real values, the contours or iso-surfaces of a specified scalar value are an important way to visualize the function. To find such contours, a seed set can be used as the starting point from which the traversal of the contours can begin. In this project, our aim is to extract iso-surfaces of 3D meshes containing scalar field visualization data and render them for different scalar values using contour propagation using seed sets and display them in an interactive, user friendly manner.

2 Related Work

Previous works which we have studied are listed here. [4] [3][1]. We have implemented marching cube algorithm using [2] as reference.

3 Work Completed

3.1 Real Time Contour Propagation

We have implemented a pre-processing based algorithm for calculating contours in real time for different scalar values. Display is supported for a predefined list of scalar values(say S). By default, the display tool supports all scalar values between 0 to 1 with a step-size of 0.01(ie $\{0.01, 0.02 \dots 1.00\}$). Additionally, the user can also specify more specific scalar values in a config file for a finer control over isosurfaces to be displayed. We have made our code publicly available at <https://github.com/viz27/visualize3D>. The algorithm is described below.

- Let L_i denote the list of grid cells that contain scalar value i .
- First, do a sweep of all the cells in data grid and add the index of each grid cell to the list L_i if scalar value i passes through the grid cell. For eg. if scalar values 0.01 and 0.02 passes through grid cell of index 1000, then 1000 will be added to $L_{0.01}$ and $L_{0.02}$.
- Once we have the list L_i , we can directly look at the grid cells in the list and extract contours from those cells as required. We call this mode of operation **FAST Mode**. In this mode, index of all the intersecting cells for all the scalar values in S need to be kept in memory. Hence, it is memory inefficient, but contour propagation is fast.
- We also describe another mode called **SEED Mode**, where seed sets are calculated for every scalar values in S .
- Given cell IDs $a, b \in L_i$, we can combine a and b to a single set if a and b are neighbors in the data grid. Checking all entries in L_i in this manner and combining neighbors to single set, items in L_i will be divided into disjoint sets and elements in each set will belong to the same connected component in the isosurface of scalar value i . This combination can be done efficiently using union-find algorithm. Taking a single representative item from each set in L_i , we will get a small number of seed cells from which all the other cells can be identified via propagation.

- Instead of calculating seed sets for each scalar value one after another, we describe a method to calculate it in one sweep of the entire grid cells. For a grid cell with ID j , consider all its neighbors (up to 28 neighbors for a cubic cell) and whenever cell j and neighbor k belong to the list L_i (ie. scalar value i passes through both cells j and k), merge them in L_i (using union find algorithm). After sweeping over the entire grid cells in this manner, each L_i will be split into disjoint sets from which seed cells can be extracted.
- Once we have the seed set for each scalar value $i \in S$, this can be saved and for subsequent runs, this can be retrieved and contours can be calculated based on the seed set, eliminating the pre-processing phase in subsequent runs.

3.2 Interactivity

1. “Space key input” to stop and start rotating the iso-surface.
2. Trackball to rotate the object along x and y axes.
3. Scrollbar for changing the camera position and re-rendering simultaneously.
4. Scrollbar for changing the iso-values and re-rendering.
5. Keyboard input to specify new scalar values.

4 Experiments and Results

Here we report the processing time and number of vertices generated per iso-surface for 4 different scientific visualization datasets. The numbers are shown for the program when run in FAST Mode.

Dataset	Preprocessing Time			Contour Extraction Time				Number of vertices in isosurface			
	data read	generate cells	total	0.01	0.1	0.2	0.5	0.01	0.1	0.2	0.5
teapot_256x256x178_uint8	1s	1s	2s	0.32s	0.24s	0.23s	0.007s	2.9M	2.2M	2.1M	72K
bonsai_256x256x256_uint8	2s	1s	3s	1.1s	0.58s	0.16s	0.17s	11.5M	5.8M	1.7M	1.9M
csafeheptane_302x302x302_uint8	3s	2s	5s	0.35s	0.39s	0.24s	0.15s	3.0M	3.4M	2.2M	1.5M
stent_512x512x174_uint16	8s	2s	10s	0.69s	0.38s	0.31s	0.02s	7.2M	4.0M	3.3M	0.3M

Table 1: Performance of visualization tool in FAST Mode

5 Discussion

As seen from Table 1, our tool is quite fast for both pre-processing as well as real time contour extraction for Medium sized datasets when run in FAST Mode. When run in SEED Mode, the pre-processing and contour extraction times are greater. This is likely due to union-find algorithm overhead for seed cell extraction and Breadth First Search overhead for contour propagation.

6 Contributions

- Vishal Peter: Algorithm and some interactivity parts.
- Srishty Suman: Interactivity part.

References

- [1] A. Bock, H. Doraiswamy, A. Summers, and C. Silva. Topoangler: Interactive topology-based extraction of fishes. *IEEE transactions on visualization and computer graphics*, 24(1):812–821, 2017.
- [2] P. Bourke. Polygonising a scalar field, <http://www.paulbourke.net/geometry/polygonise/>, 1994.
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- [4] M. Van Kreveld, R. van Oostrum, C. Bajaj, V. Pascucci, and D. Schikore. Contour trees and small seed sets for isosurface traversal. In *Proceedings of the thirteenth annual symposium on Computational geometry*, pages 212–220, 1997.