


**<http://www.cs.utexas.edu/~bajaj/cs384R08/>**

The figure consists of four panels illustrating different ways to visualize a scalar field:

- Top Left:** A 3D perspective view of a surface. The surface has a green upper portion and a red lower portion. Two prominent peaks are highlighted with red dots and labeled with their values: 20 and 25.
- Top Right:** A 2D grid representation of the field. Black dots represent the grid nodes. Colored paths highlight specific contours or trajectories: a red path, a green path, and a yellow path. Some nodes are also labeled with numerical values like 18, 10, 7, 9, 16, 20, 13, 15, 22, 25, 12, 10, 5, 17, 10, 15, 10, 10.
- Bottom Center:** A 2D contour plot showing the field's distribution. It features a Y-shaped structure. Three points are labeled: 20 at the top left branch, 25 at the top right branch, and 0 at the base of the Y.
- Bottom Right:** A 2D visualization showing a set of black nodes connected by several colored paths (black, brown, green). The paths represent different trajectories or level sets through the field. Labeled values include 20, 25, 15, 12, 10, 9, and 0.



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# Surface Splines from Volumes III: Contouring Scalar Functions



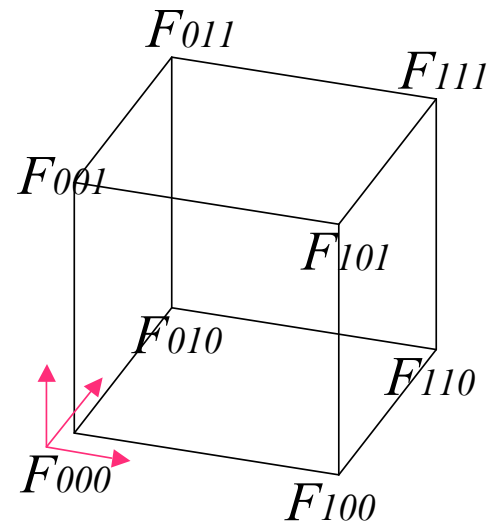
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September 2008

# Isosurface of Trilinear Function

- Trilinear Function

$$\begin{aligned} F(x,y,z) = & F_{000}(1-x)(1-y)(1-z) \\ & + F_{001}(1-x)(1-y)z \\ & + F_{010}(1-x)y(1-z) \\ & + F_{011}(1-x)yz \\ & + F_{100}x(1-y)(1-z) \\ & + F_{101}x(1-y)z \\ & + F_{110}xy(1-z) \\ & + F_{111}xyz \end{aligned}$$



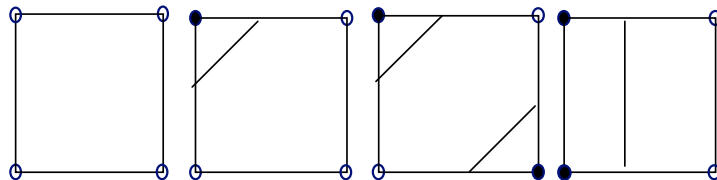
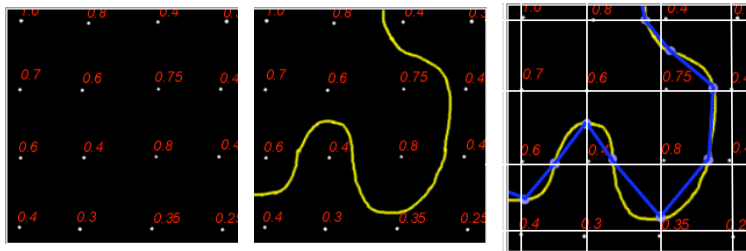
- Bilinear Function

$$\begin{aligned} F^f(x,y) = & F_{00}(1-x)(1-y) + F_{01}(1-x)y \\ & + F_{10}x(1-y) + F_{11}xy \end{aligned}$$



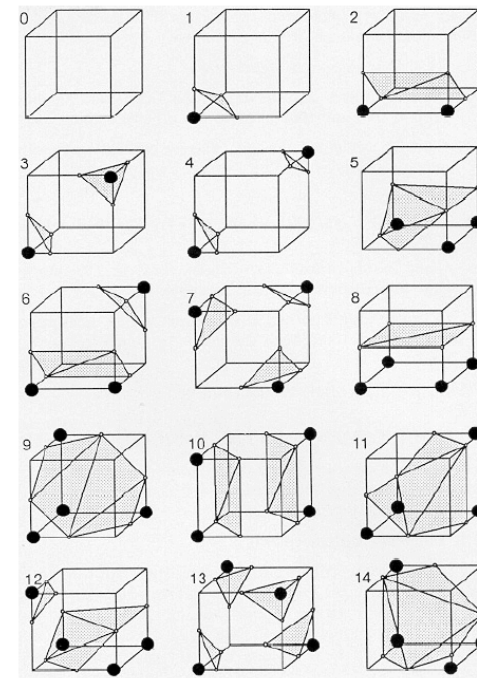
# Marching Cubes (MC) : Triangular Approximation

- 2D rectangle



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- 3D cube :  
15 Cases for Triangulation



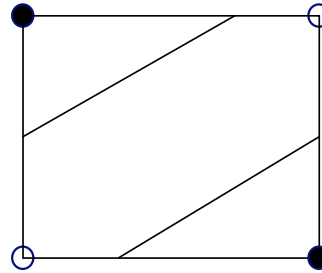
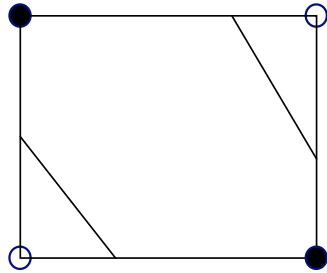
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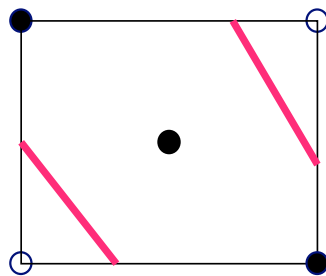
# Decision on Contour Topology ( Nielson 92 : Asymptotic Decider )

- Resolving Face Ambiguity

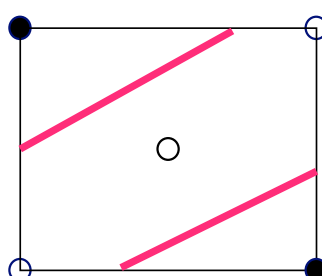
- Ambiguity ( face saddle )



- Decision based on the value  $s$  of saddle point



*$s$  is positive*



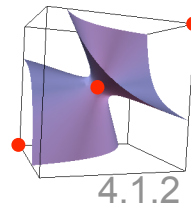
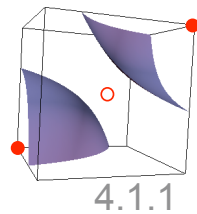
*$s$  is negative*





# Decision on Contour Topology ( Natarajan 94 )

- Resolving Internal Ambiguity
  - Ambiguity ( body saddle )



–Decision based on the value  $s$  of saddle point

- (i)  $s$  is positive  $\rightarrow$  tunnel
- (ii)  $s$  is negative  $\rightarrow$  two pieces



# Saddle Points Computation

- Face Saddle Point

$$F(x,y) = ax + by + cxy + d \quad (\text{bilinear interpolant})$$

$$\text{First derivatives : } F_x = a + cy = 0, F_y = b + cx = 0$$

$$\text{Saddle point } S = (-b/c, -a/c)$$

- Body Saddle Point

$$F(x,y,z) = a + ex + cy + bz + gxy + fxz + dyz + hxyz$$

First derivatives = 0 :

$$F_x = e + gy + fz + hyz = 0$$

$$F_y = c + gx + dz + hxz = 0$$

$$F_z = b + fx + dy + hxy = 0$$



# Face and Body Saddle Points

- We obtain saddle points :

$$x = -\frac{c+dz}{g+hz}$$

$$y = \frac{k_0+k_1z}{k_2}$$

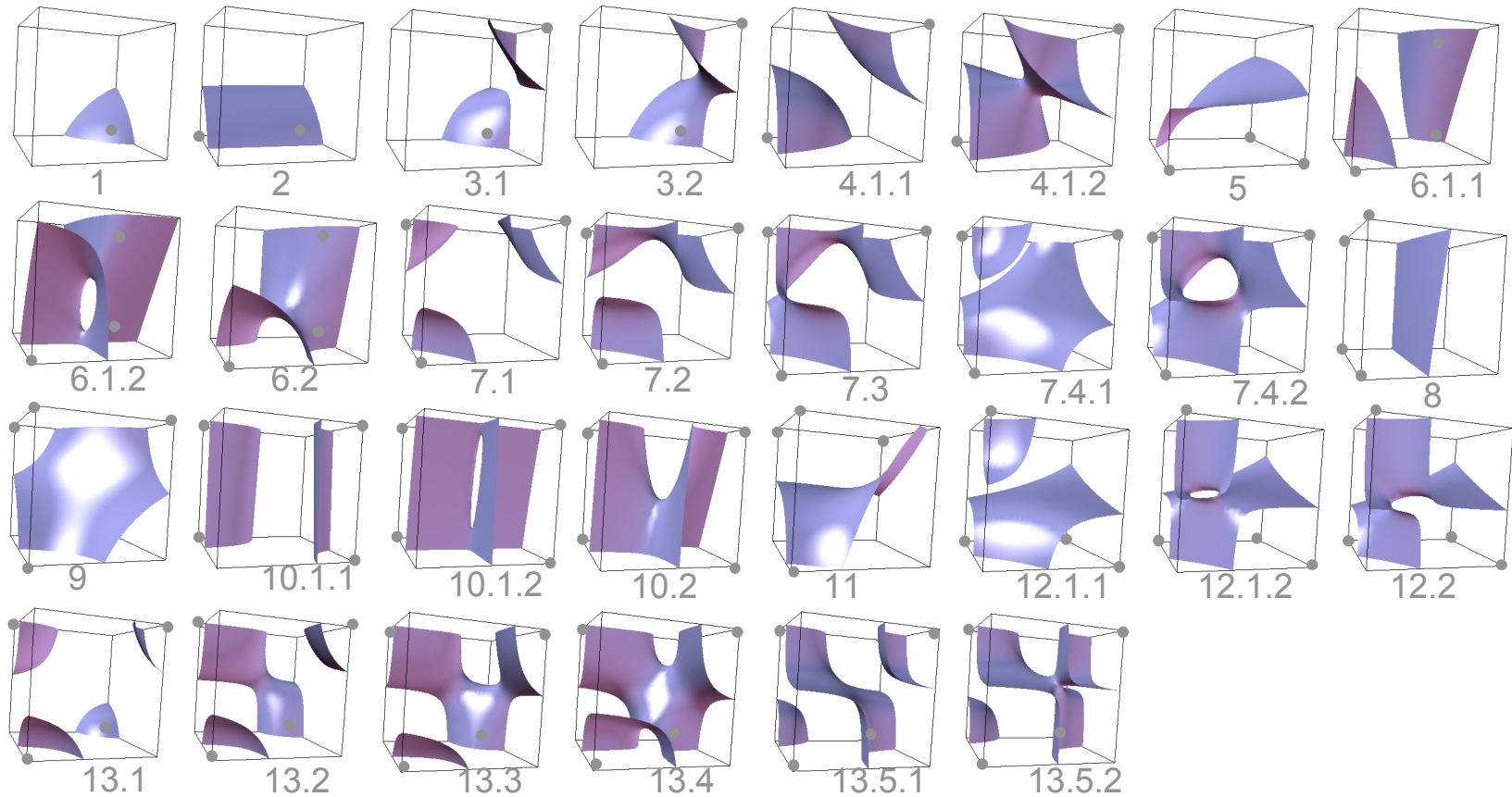
$$z = -\frac{g}{h} \pm \frac{\sqrt{g^2k_1^2 - hk_1^{1/2}(ek_2 + gk_0)}}{h}$$

$$k_0 = cf - bg, k_1 = df - bh, k_2 = dg - ch$$

- saddle point outside the cube → discard  
( only case 13.5 has more than one valid body saddle point. )



# Trilinear Isosurface Topology 31 cases



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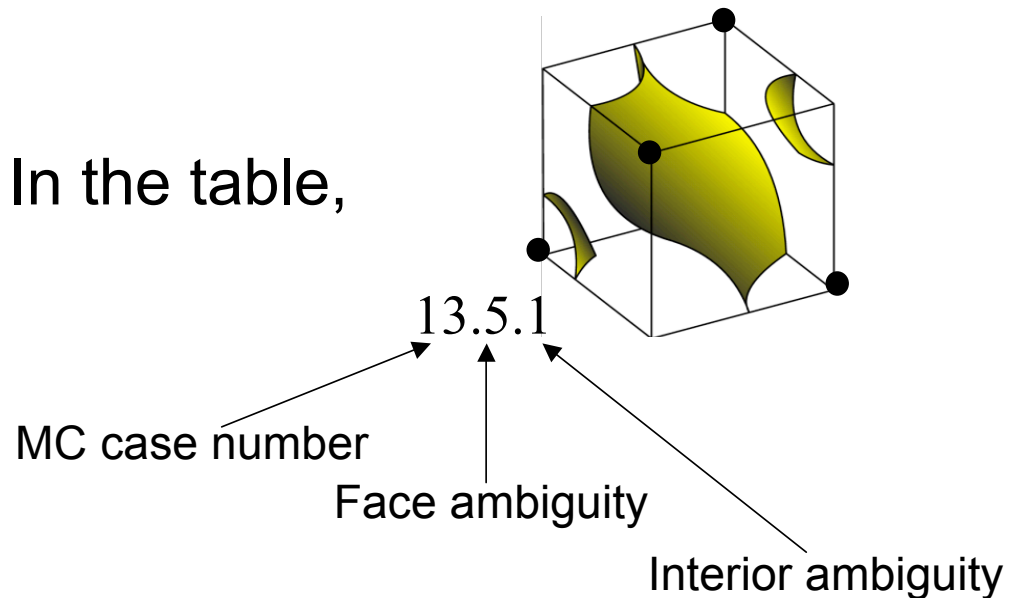
Lopes, Brodlie 2003

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## 31 Cases

- In the table,



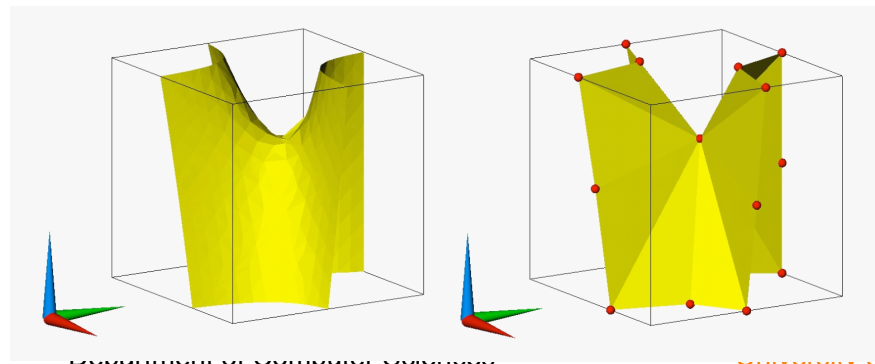
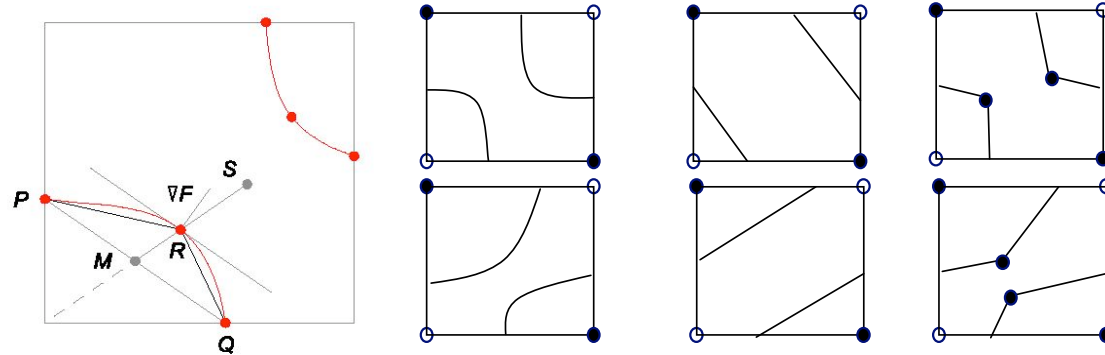
- MC : 15(further reduced to 14) cases based on vertex coloring (symm).
- 31 cases ← (vertex coloring , face ambiguity , internal ambiguity)

Symmetry of different configurations are used to reduce the cases.



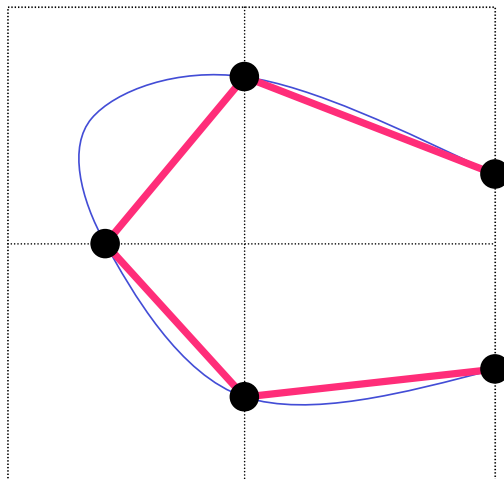
# Geometric Approximations

- Better approximation of trilinear interpolant
  - Adding a shoulder and inflection points

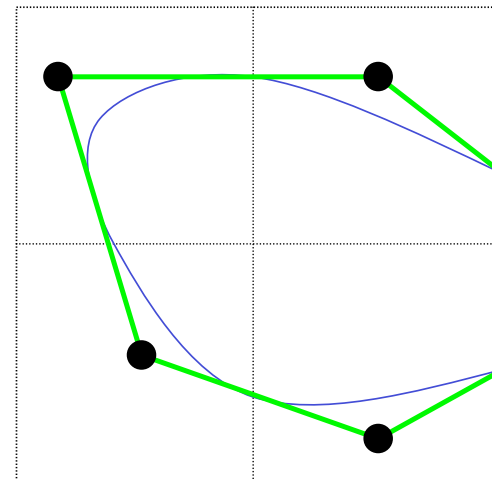


# Dual Contouring

- Primal Contouring vs Dual Contouring



Primal contour

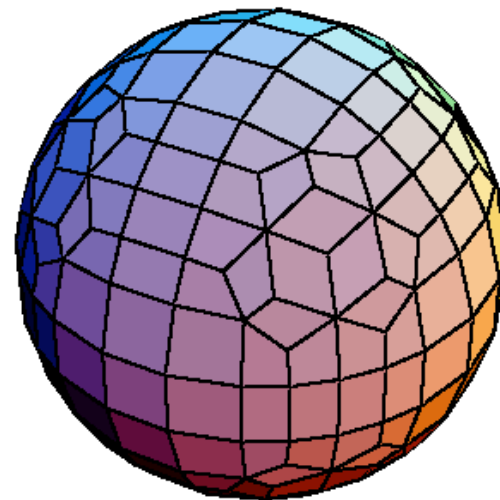
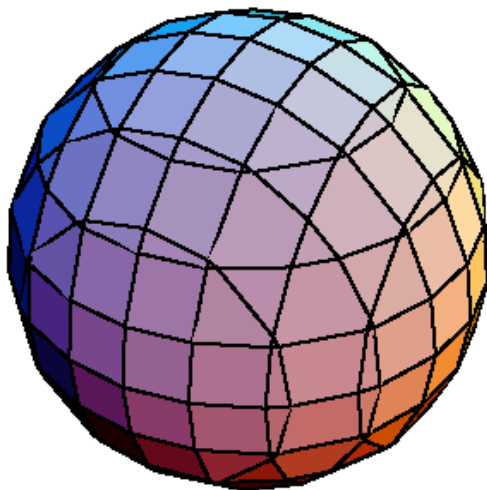


Dual Contour



# Dual Contouring

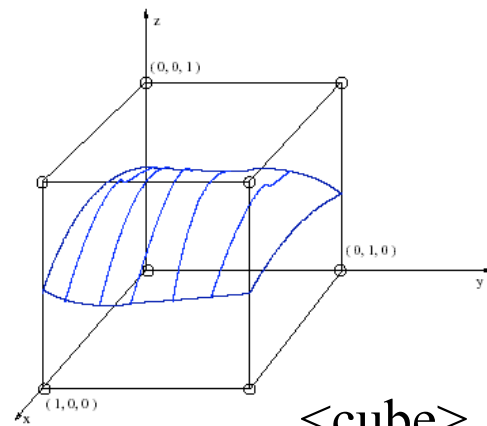
- Polygons with better aspect ratio



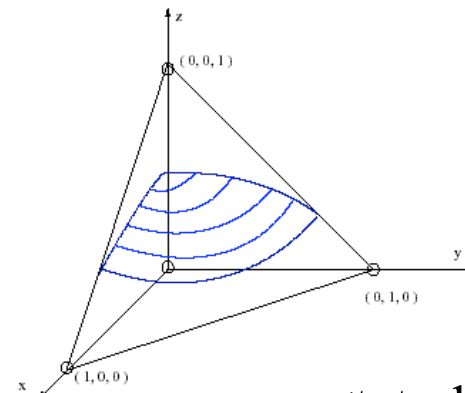


## Algebraic Patches: Smooth Boundary Elements

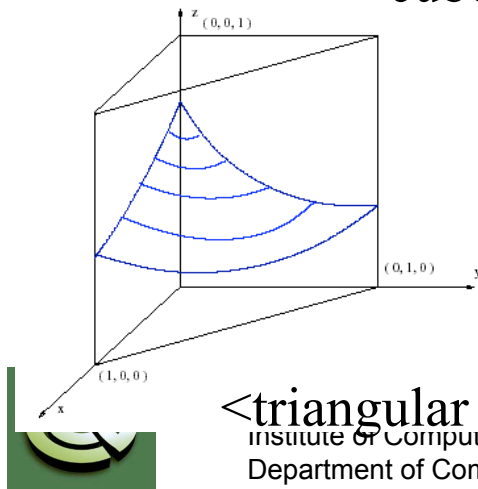
- Implicit form of Isocontour :  $f(x,y,z) = w$



<cube>

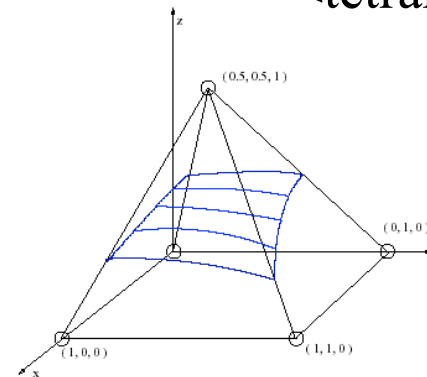


<tetrahedron>



<triangular prism>

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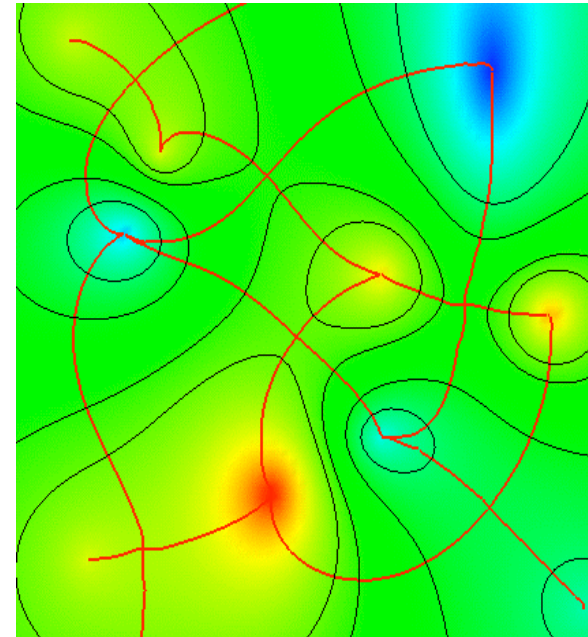
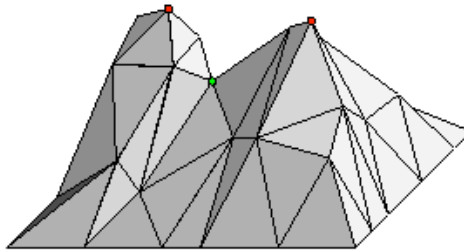
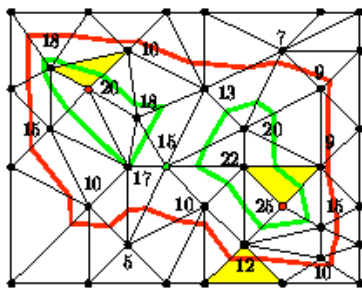
<square pyramid>

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# Interactive Isocontour Queries

- Input:
  - Scalar Field  $F$  defined on a mesh
  - Multiple Isovalues  $w$  in unpredictable order
- Output (for each isovalue  $w$ ):  
Contour  $C(w) = \{x \mid F(x) = w\}$



# Isocontour Query Problem

Lower Bound

Input size  $n$   
Output size  $m$  }  $m + \log(n)$

The search for  $a_h$   
takes at least  $\log(n)$

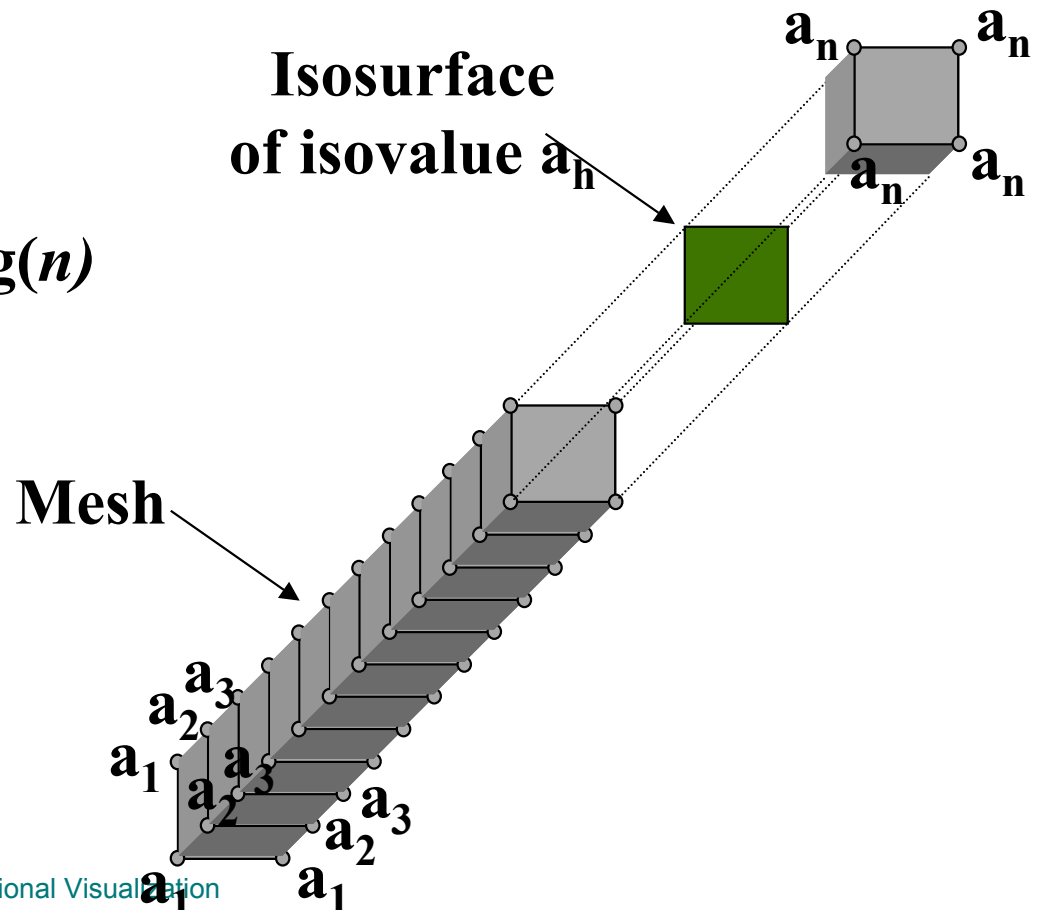
$a_1, a_2, a_3, a_4, \dots, a_n$



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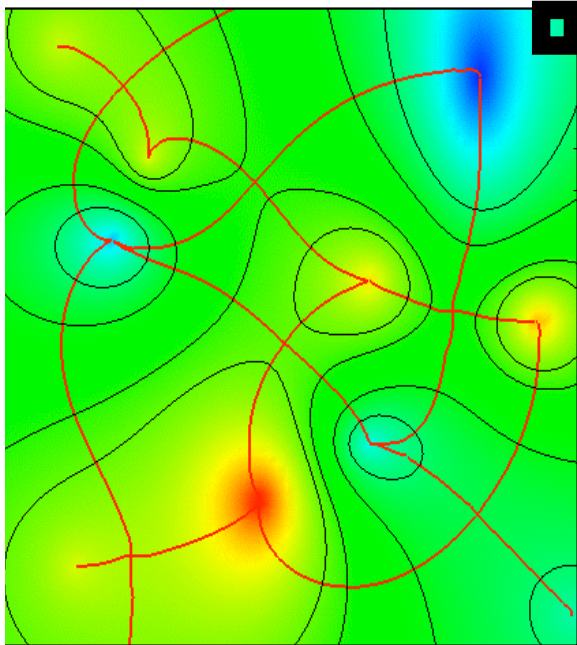
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# Optimal Single-Resolution Isocontouring

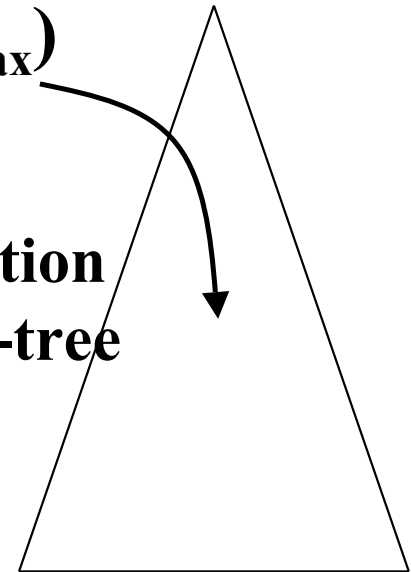
## The basic scheme



Preprocessing:  $(f_{\min}, f_{\max})$

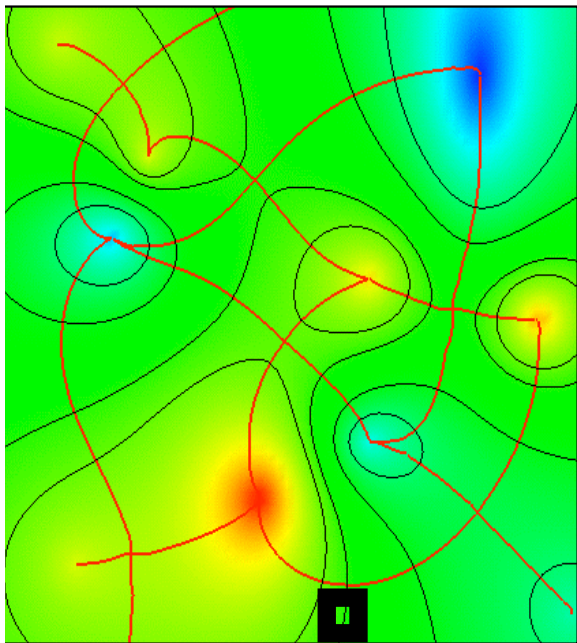
For each cell  $c$  in  $M$

Enter its range of function values into an interval-tree



# Optimal Single-Resolution Isocontouring

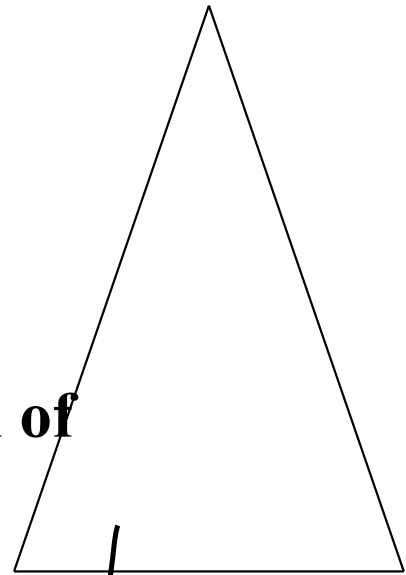
## The basic scheme



**Isocontour query  $W$**

**For each interval  
containing  $W$**

**Compute the portion of  
isocontour in the  
corresponding cell**



$(f_{\min}, f_{\max})$



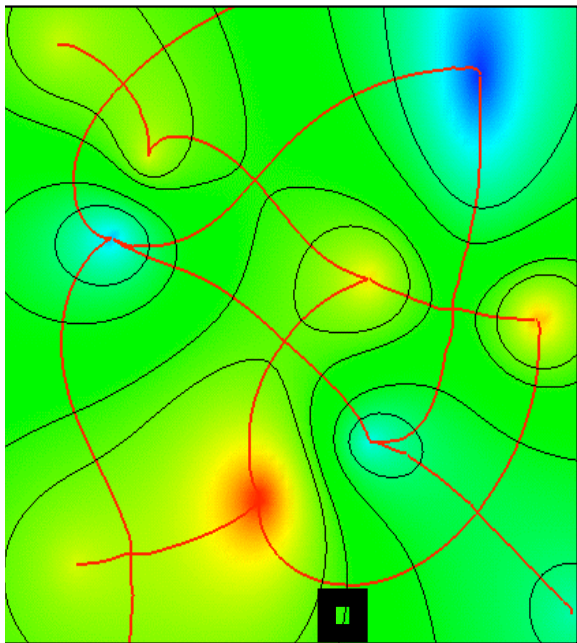
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# Optimal Single-Resolution Isocontouring

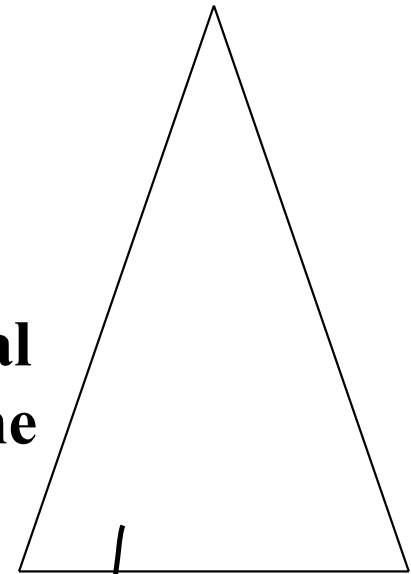
## The basic scheme



Isocontour query  $W$

Complexity:  $m + \log(n)$

Optimal but impractical  
because of the size of the  
interval-tree



$(f_{\min}, f_{\max})$



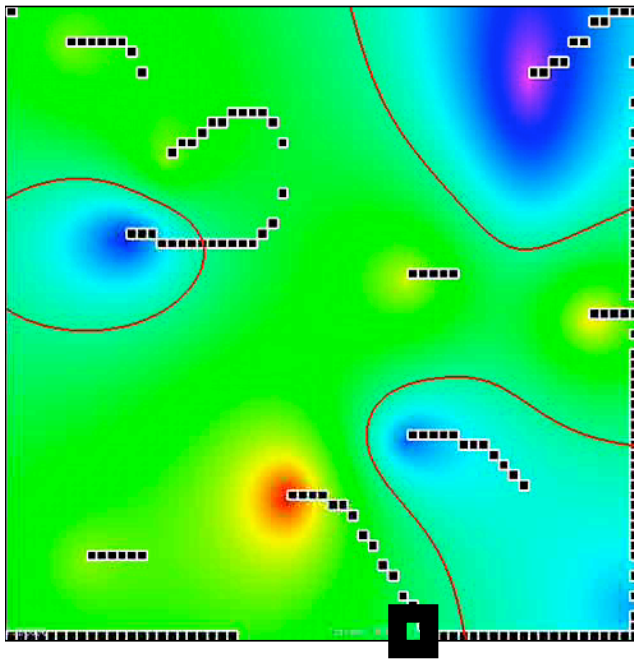
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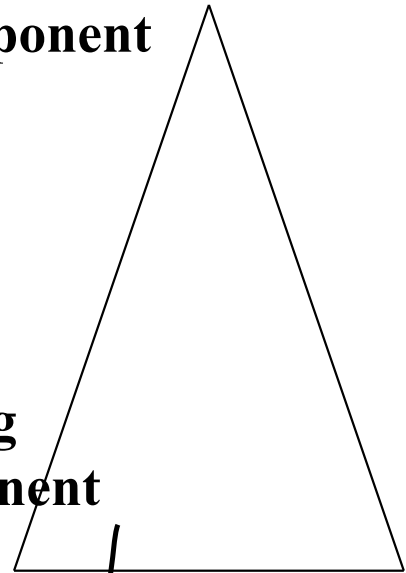
# Optimal Single-Resolution Isocontouring

## Seed Set Optimization



**For each connected component  
we need only one cell  
(and then propagate by  
adjacency in the mesh)**

**Seed Set:  
a set of cells intersecting  
every connected component  
of every isocontour**



$(f_{\min}, f_{\max})$



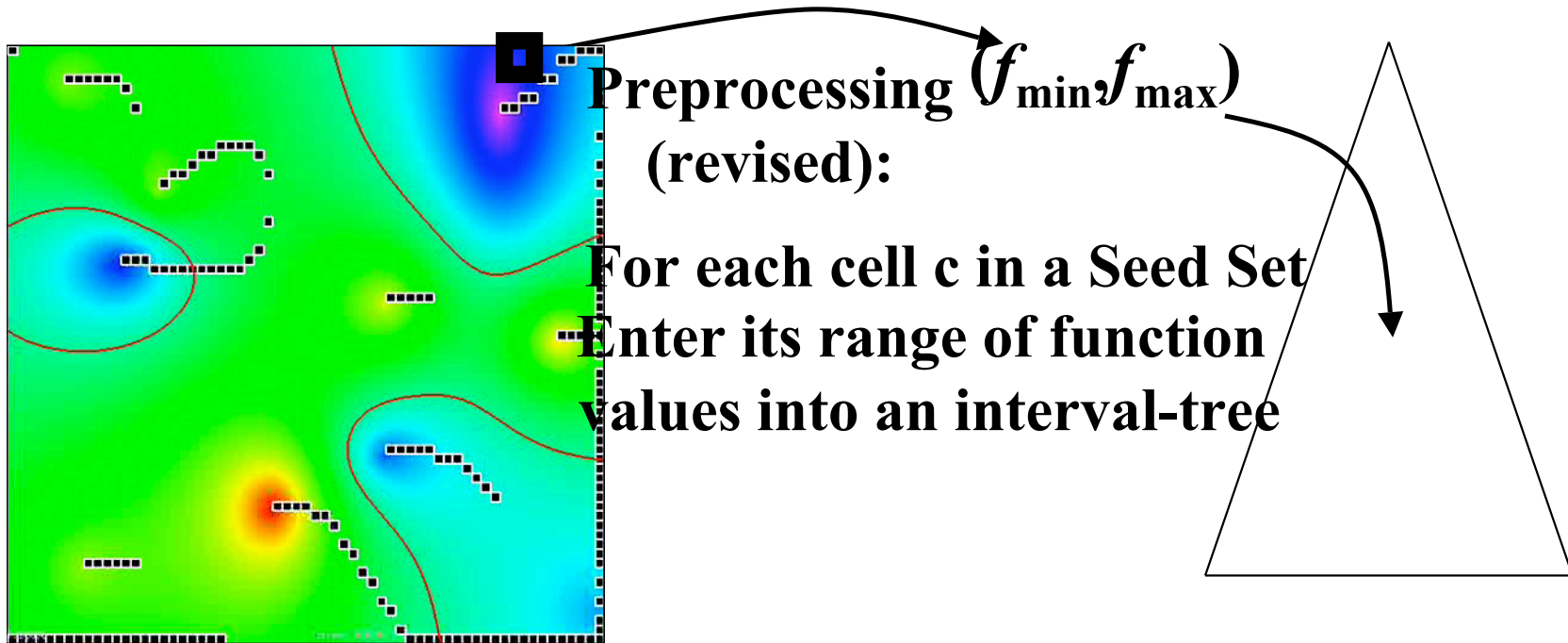
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# Optimal Single-Resolution Isocontouring

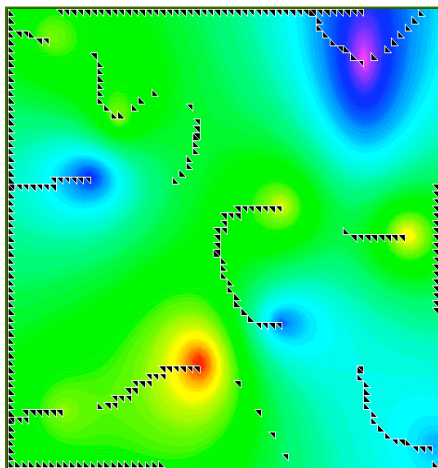
## The basic scheme



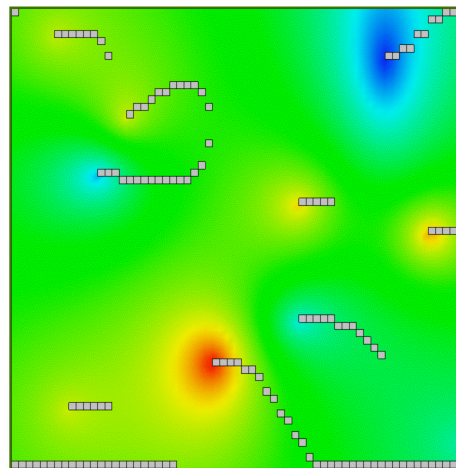


# Optimal Single-Resolution Isocontouring

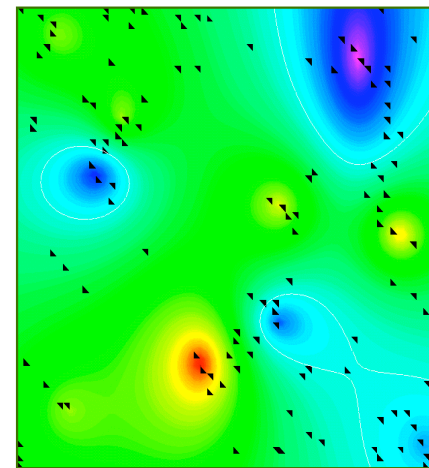
Seed Set Generation ( $k$  seeds from  $n$  cells)




Domain Sweep



Responsibility Propagation



Range Sweep

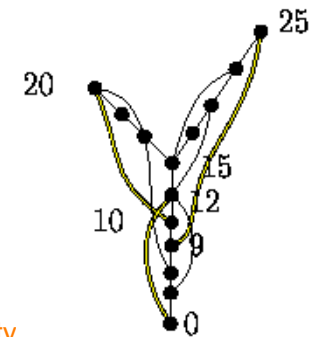
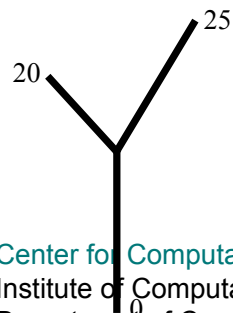
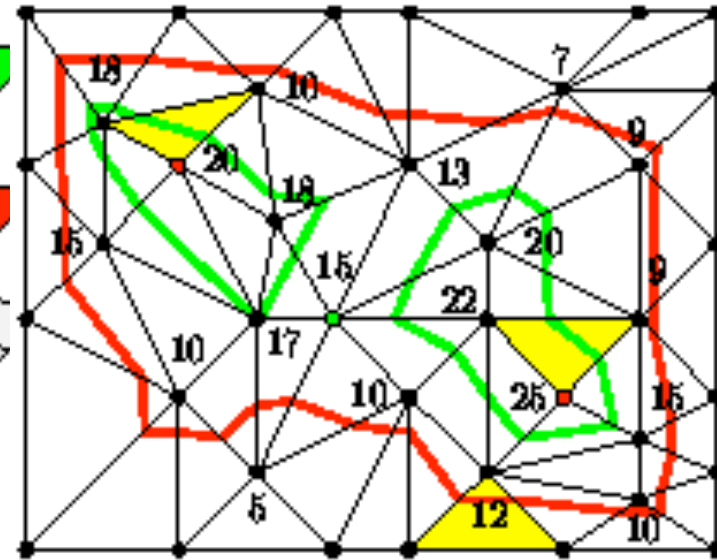
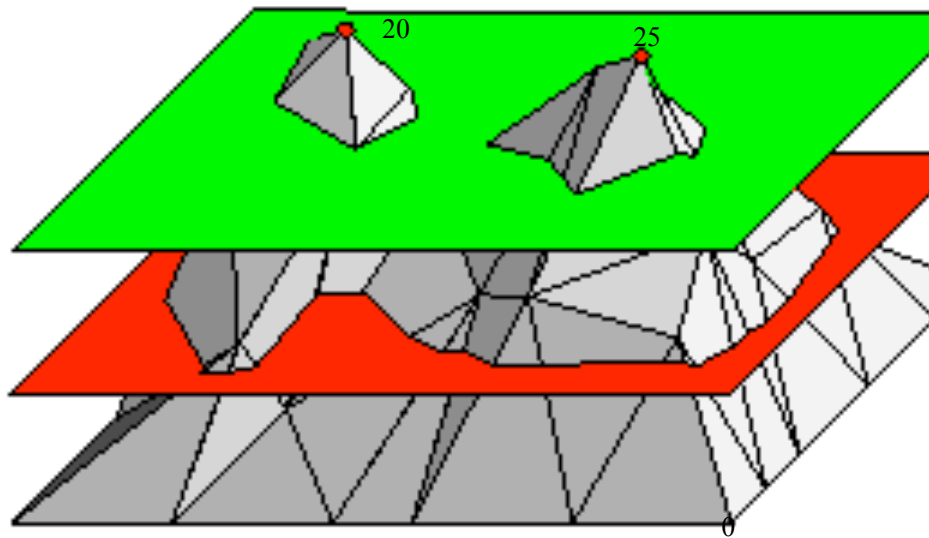
Time	$O(n)$	$O(n)$	$O(n \log n)$
Space	$O(k)$	$O(k)$	$O(n)$
$k =$	?	?	$2 k_{\min}$
Test	 <p>238 seed cells 0.01 seconds</p>	<p>177 seed cells 0.05 seconds</p>	<p>59 seed cells 1.02 seconds</p>

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# Optimal Single-Resolution Isocontouring

## Contour tree

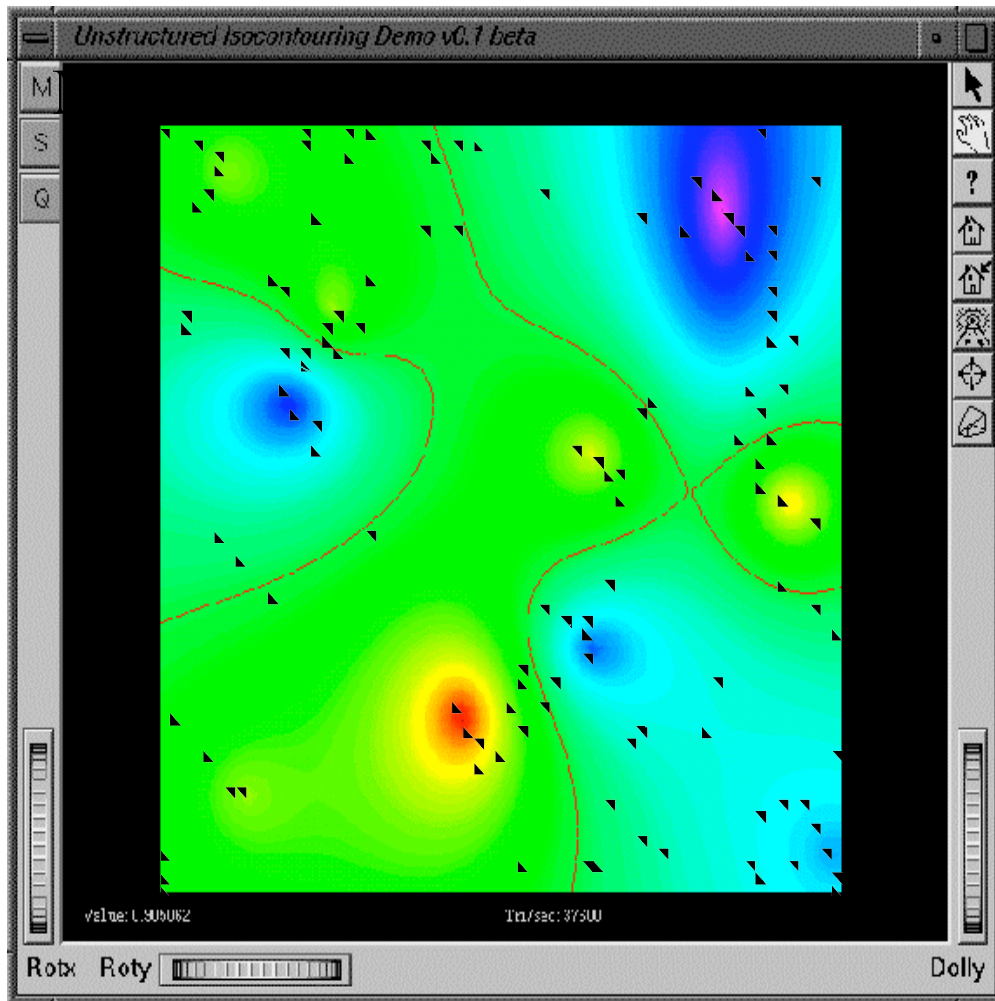


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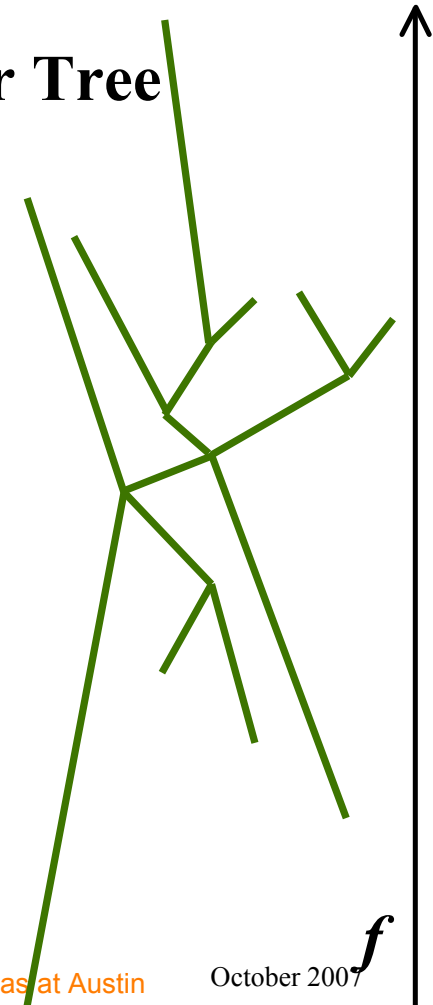
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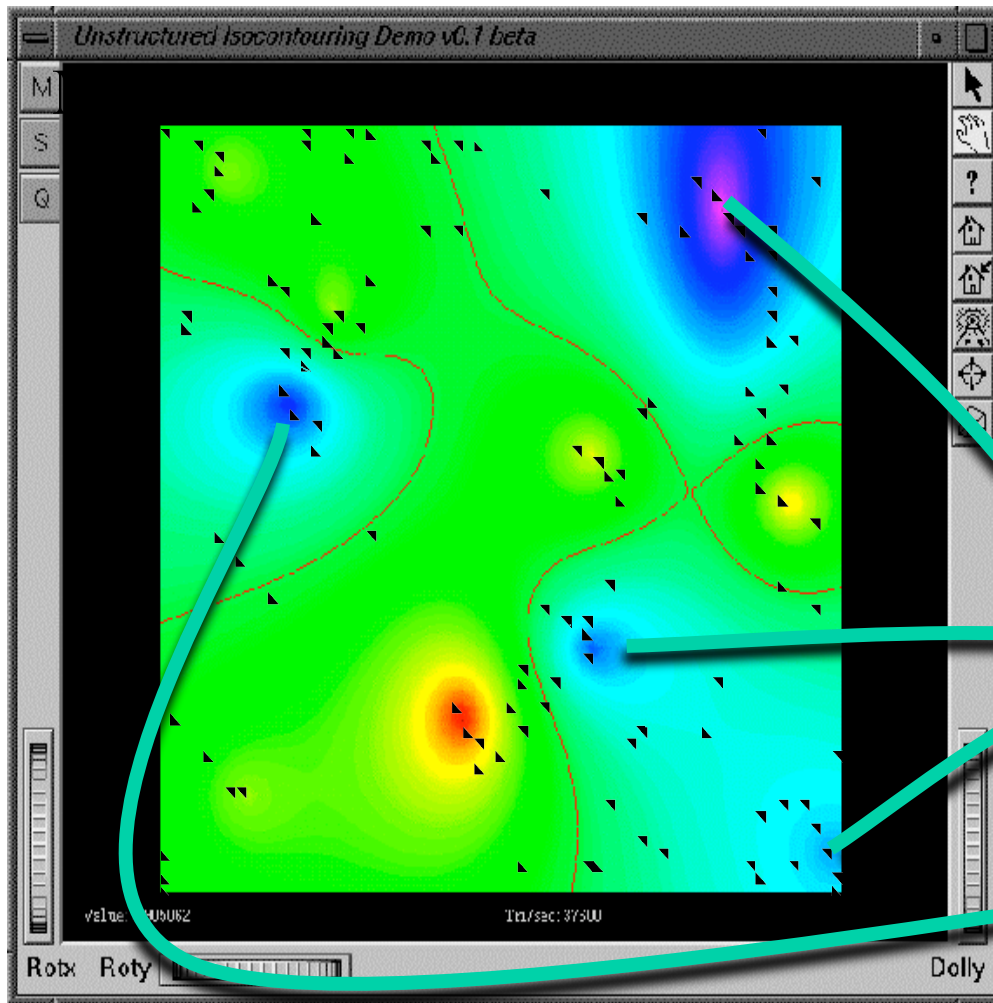
## Optimal Single-Resolution Isocontouring



## Contour Tree



## Optimal Single-Resolution Isocontouring



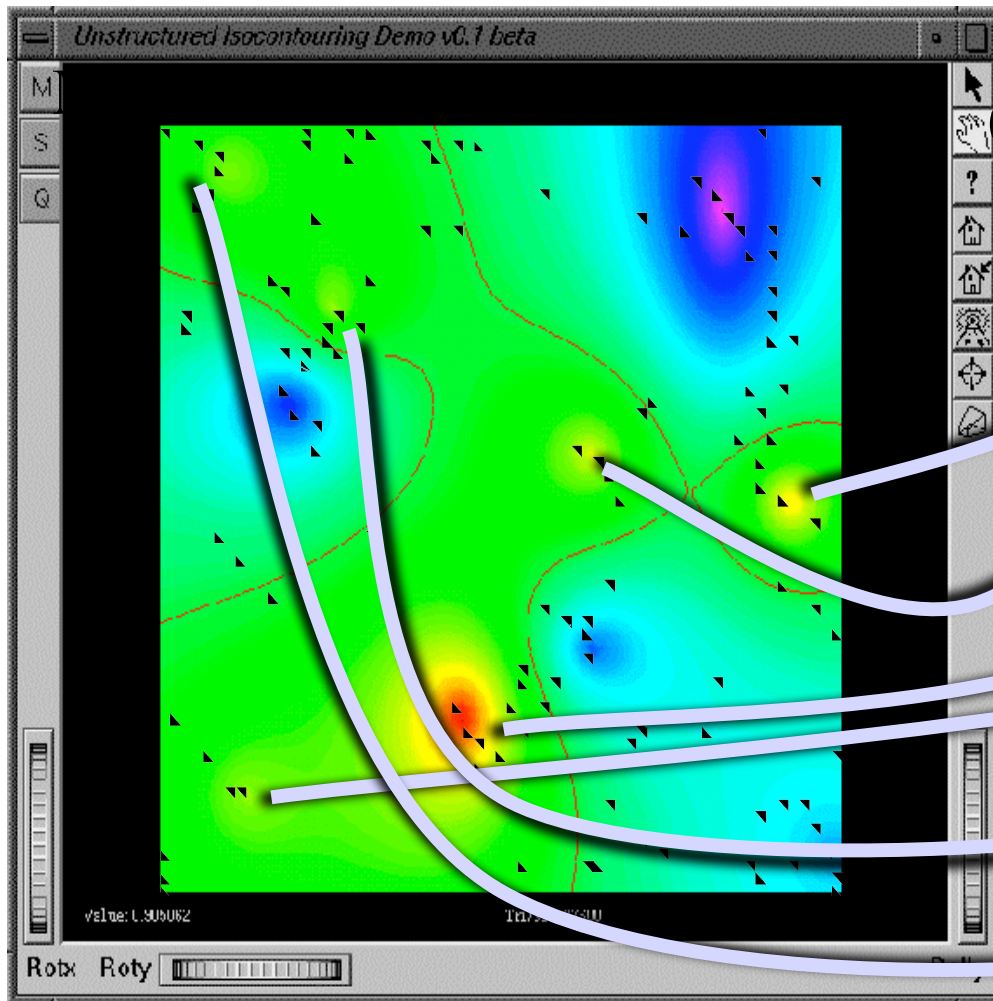
**Contour Tree  
(local minima)**

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$f$

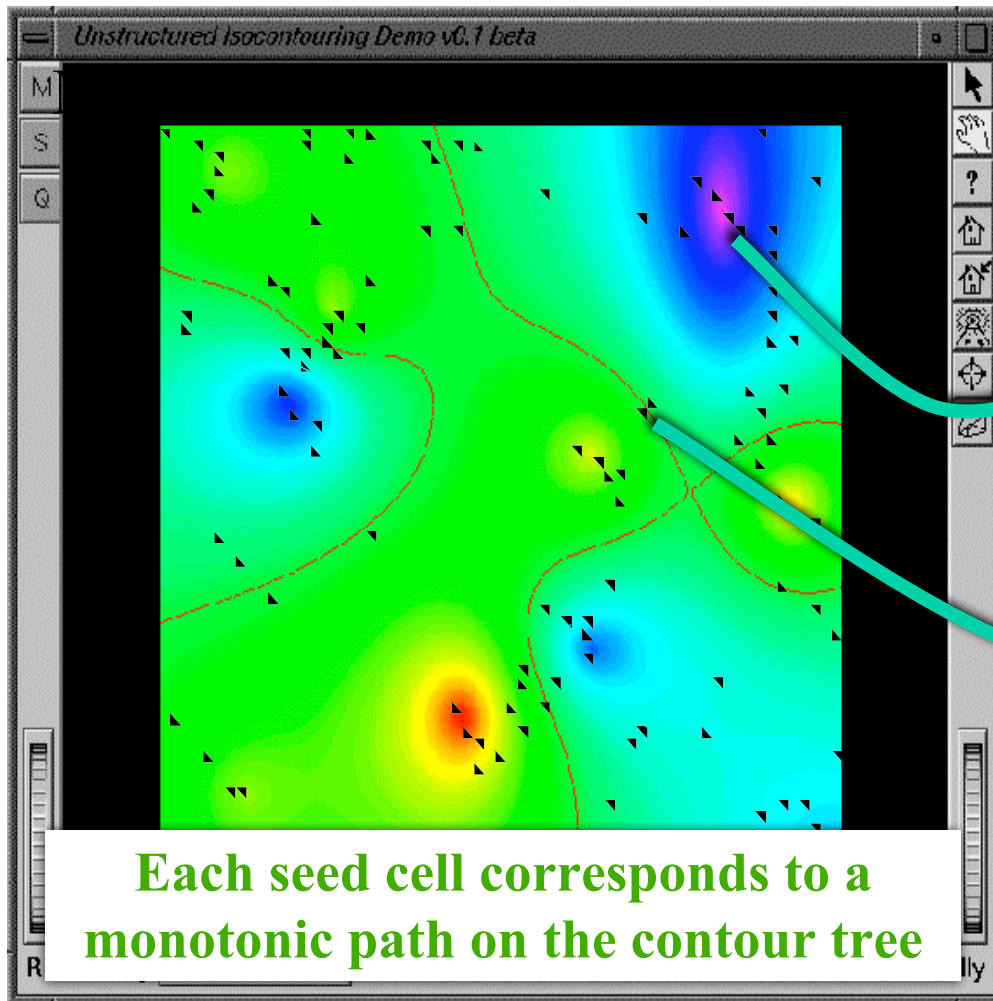
## Optimal Single-Resolution Isocontouring



Contour Tree  
(local maxima)

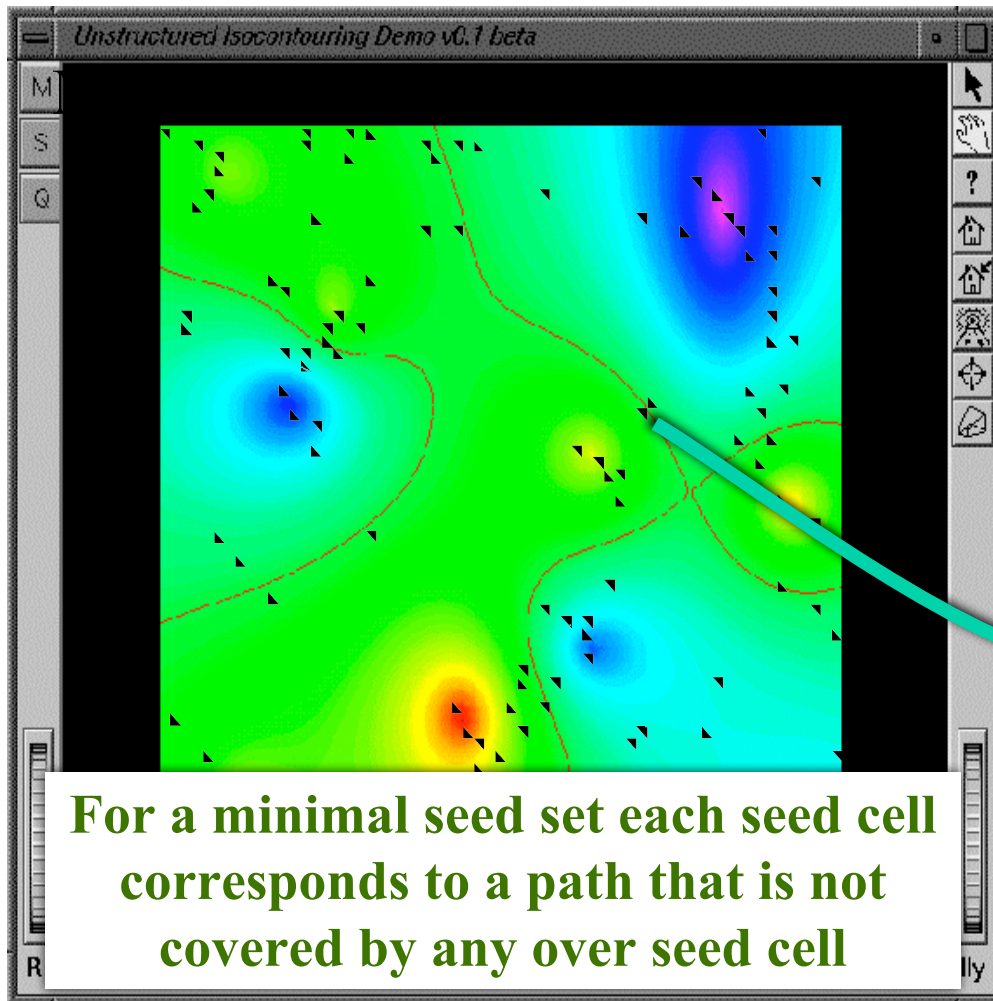


## Optimal Single-Resolution Isocontouring



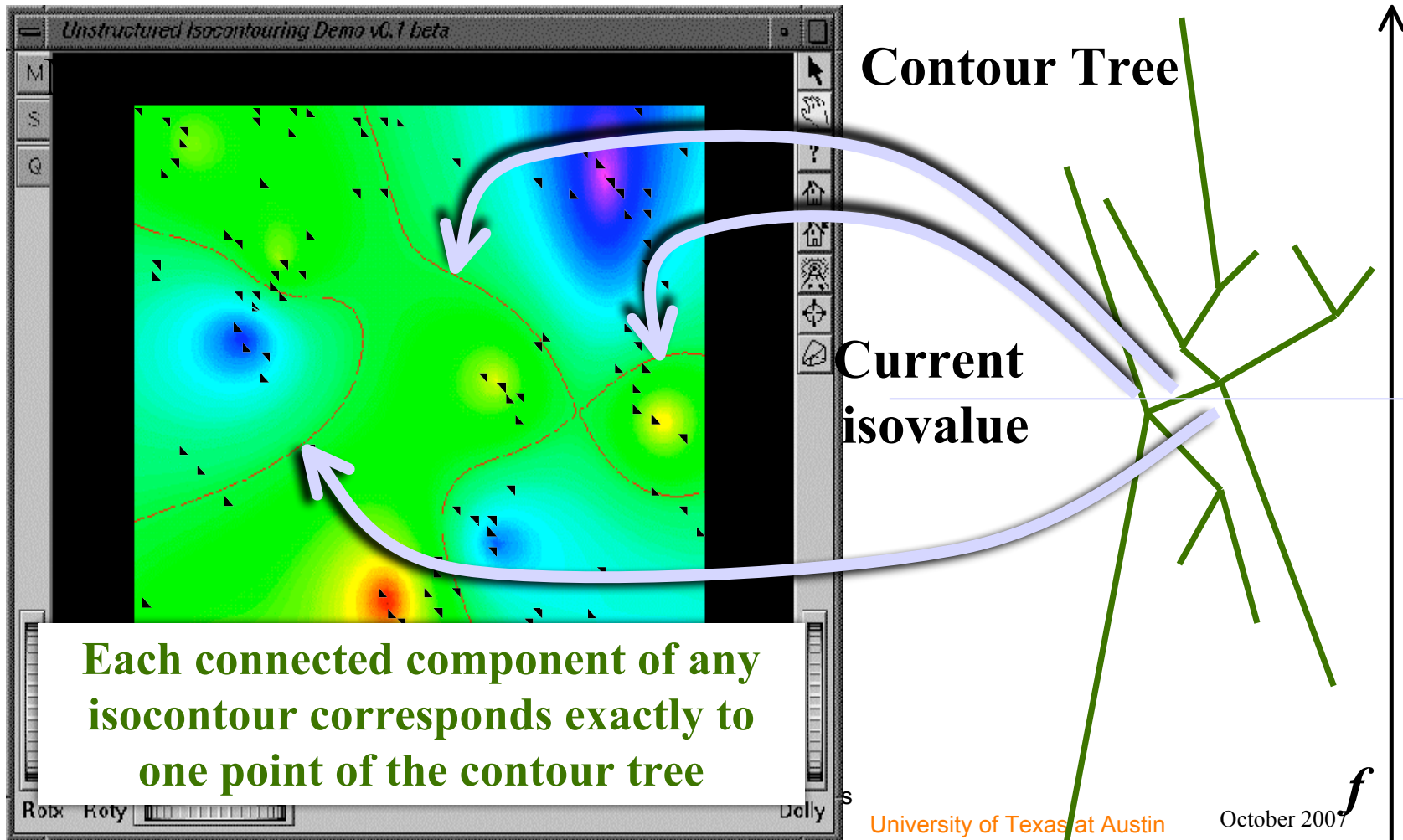
Contour Tree

## Optimal Single-Resolution Isocontouring



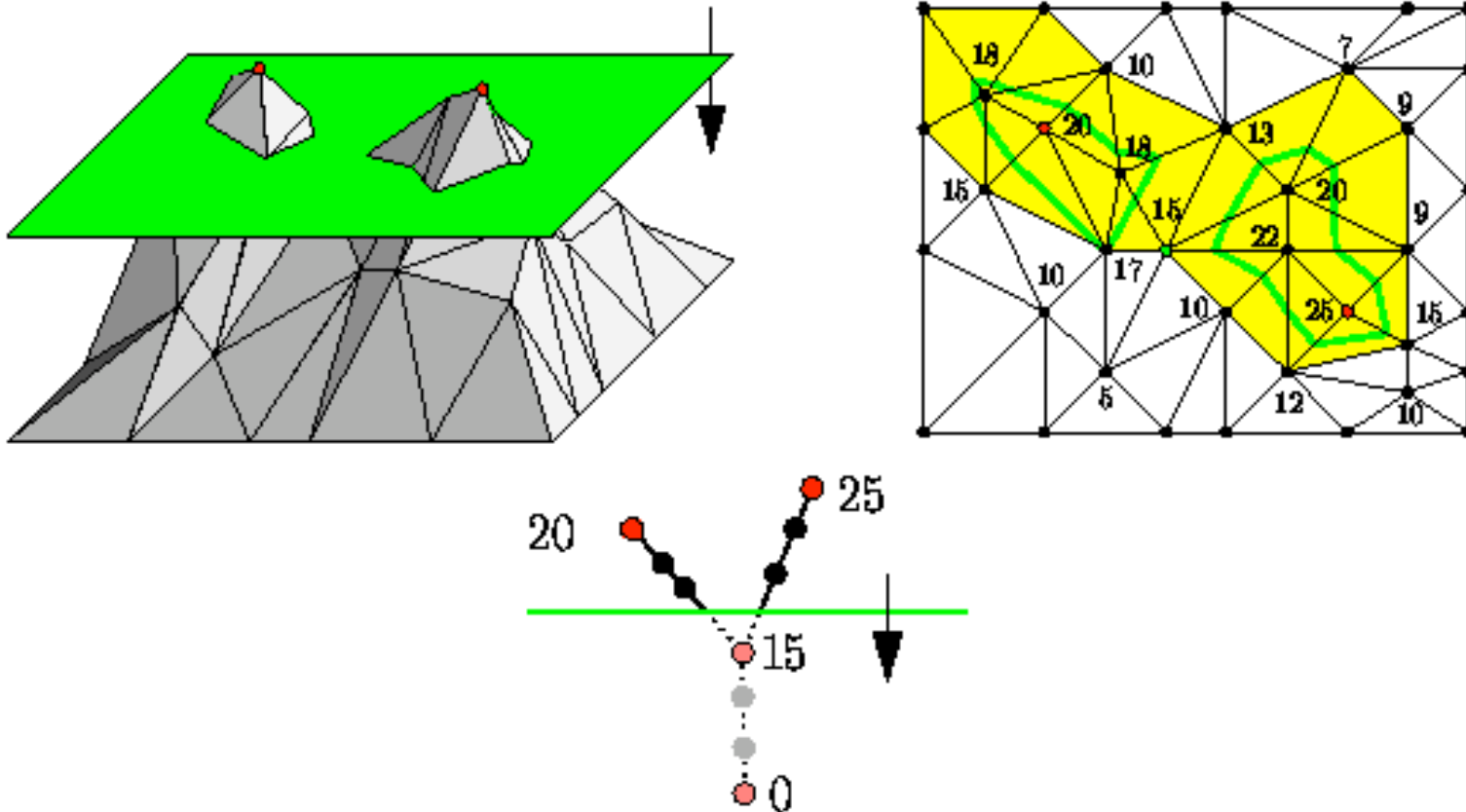
Contour Tree

## Optimal Single-Resolution Isocontouring



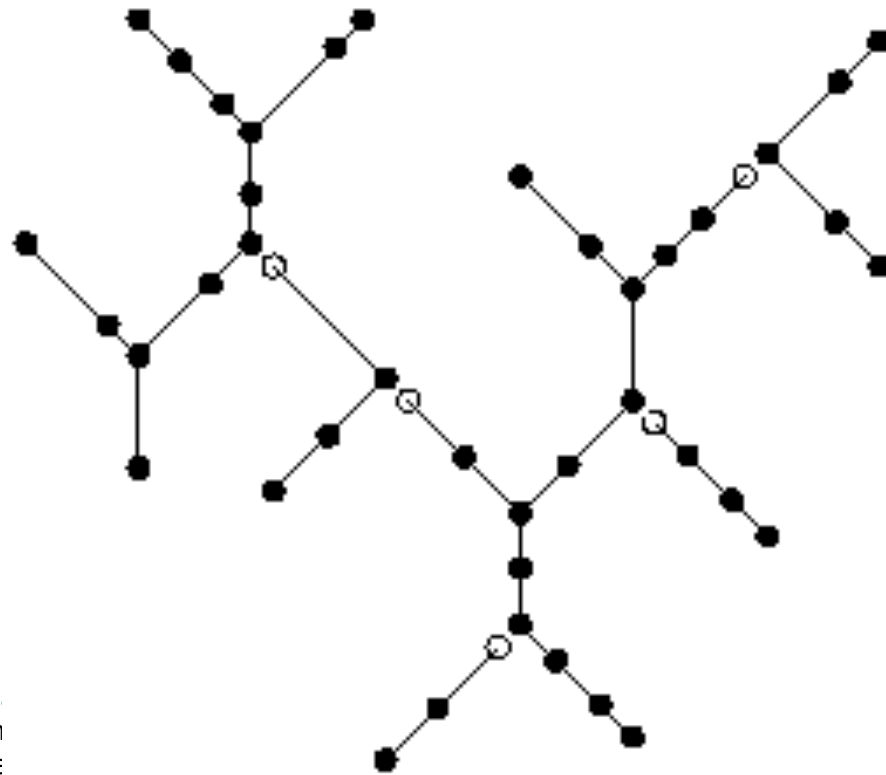


## Optimal Single-Resolution Isocontouring



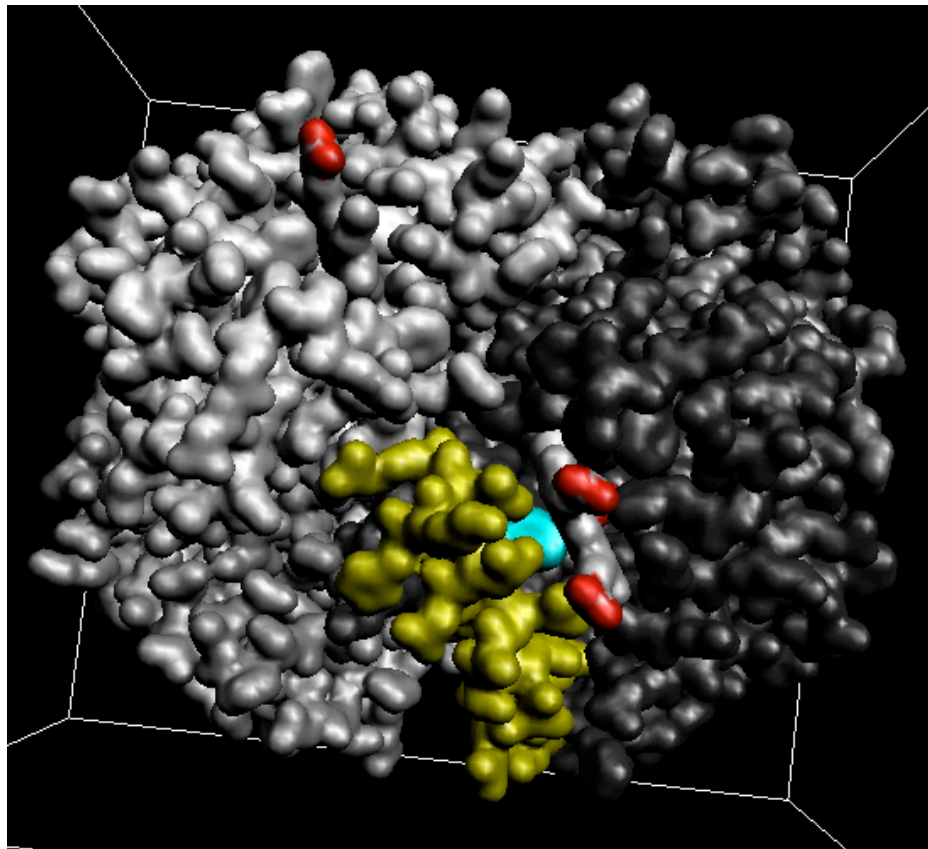
## Optimal Single-Resolution Isocontouring

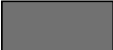


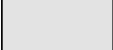
- The number of seeds selected is the minimum plus the number of local minima.






# Structural Analysis

## Contour Spectrum and Contour Tree on Hemoglobin Dynamics



	Subunit A
	Subunit B
	Subunit C
	Subunit D

Within Subunit A

	F helix
	Histidine Ligand(HIS87)
	O <sub>2</sub>

- Oxy process : O<sub>2</sub> binds to the Fe<sup>2+</sup> ion on the opposite side of the histidine ligand. F helix shifts position through the oxy-deoxy cycle.

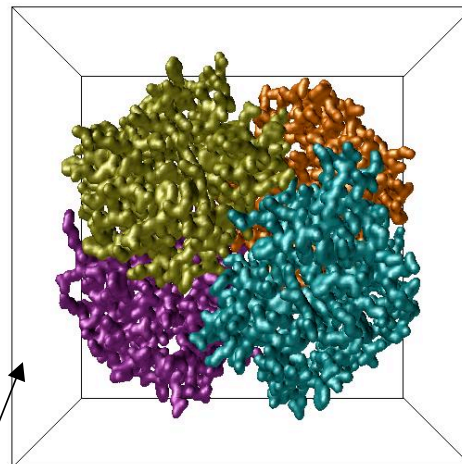
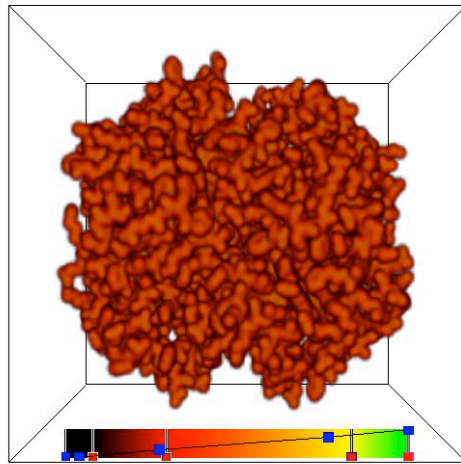


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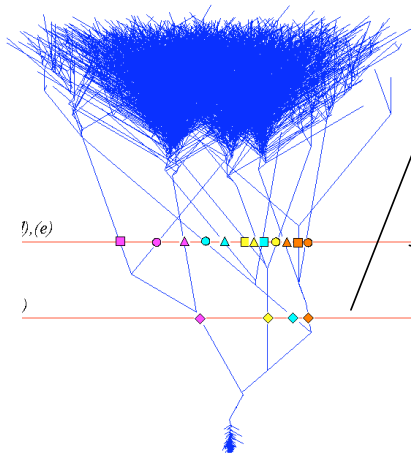
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# Topological Analysis & Visualization



Four Polypeptide chains



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Contour Tree  
of  
Electron Density Map

3D chemical bonding  
structures  
with different levels

## Functional groups

Atoms belonging to the same  
contour have stronger linkage

Each chain consists of  
heme, iron, and globin

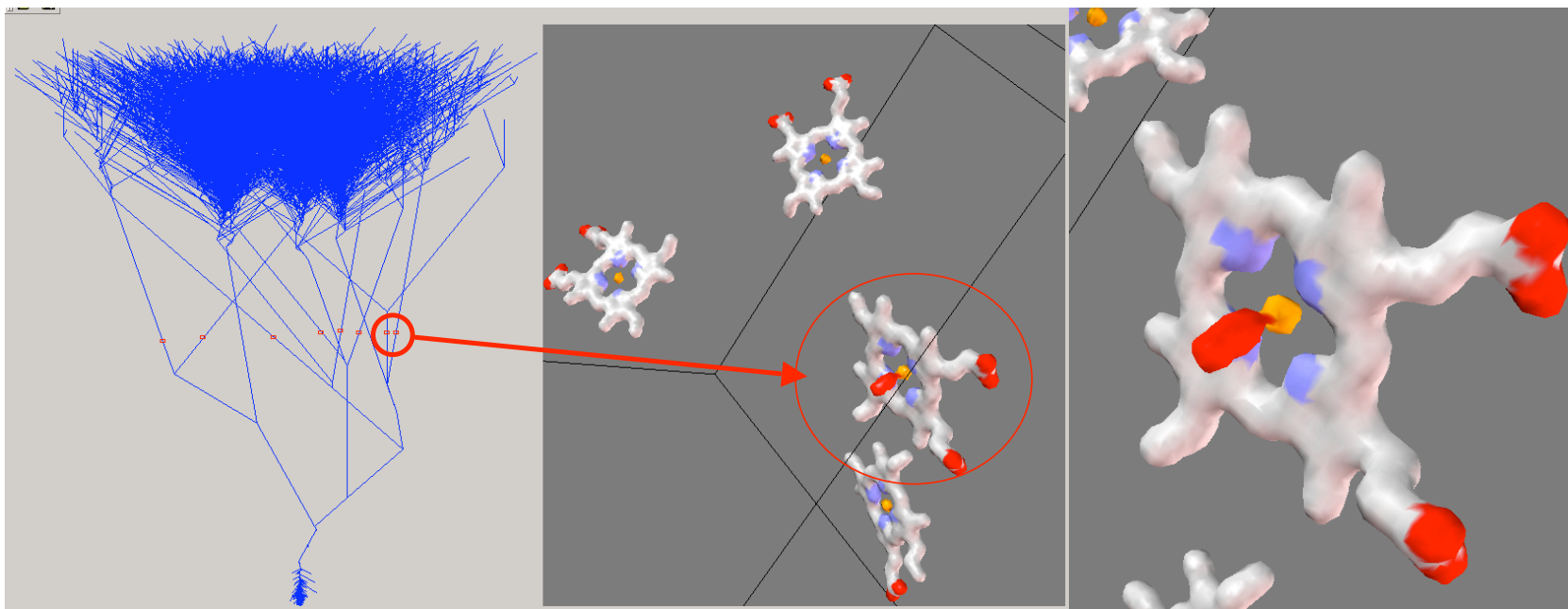
- M. van Kreveld, R. van Oostrum, C. Bajaj, V. Pascucci, and D. Schikore, *Chap5, pg 71 - 86, 2004 ed. by S. Rana, John Wiley & Sons, Ltd, 2004*
- C. Bajaj, V.Pascucci, and D.Schikore, *Proceedings of the 1997 IEEE Visualization Conference, 167-173, October 1997 Phoenix, Arizona*

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# Topological Analysis using the CONTOUR TREE

- Oxygenated Hemoglobin ( T=1 )



<isovalue = 31>



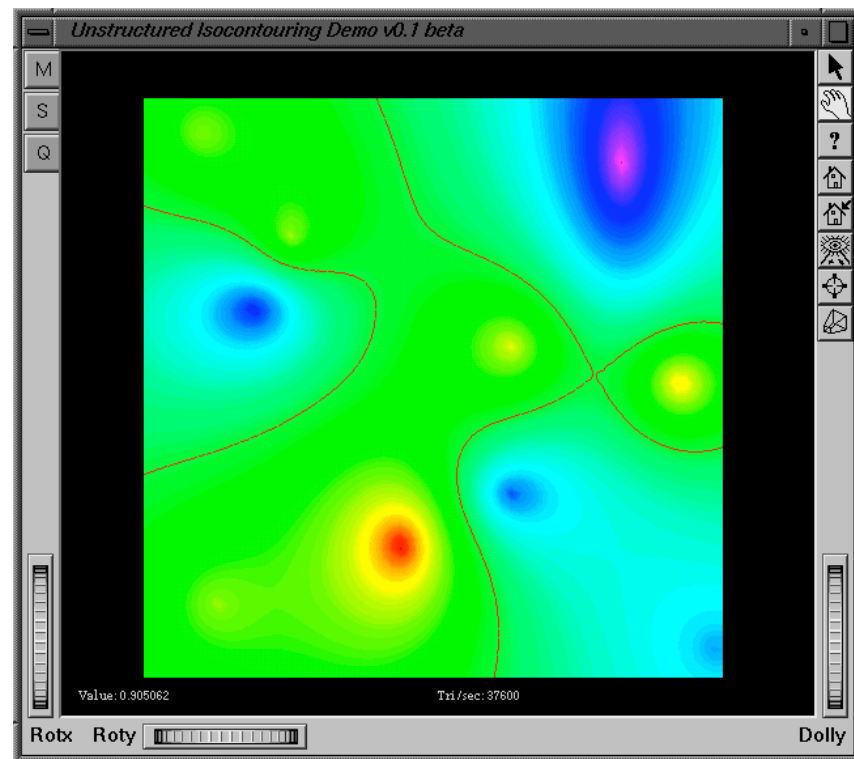
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# Spectral Analysis

- Consider a terrain of which you want to compute the length of each isocontour and the area contained inside each isocontour.



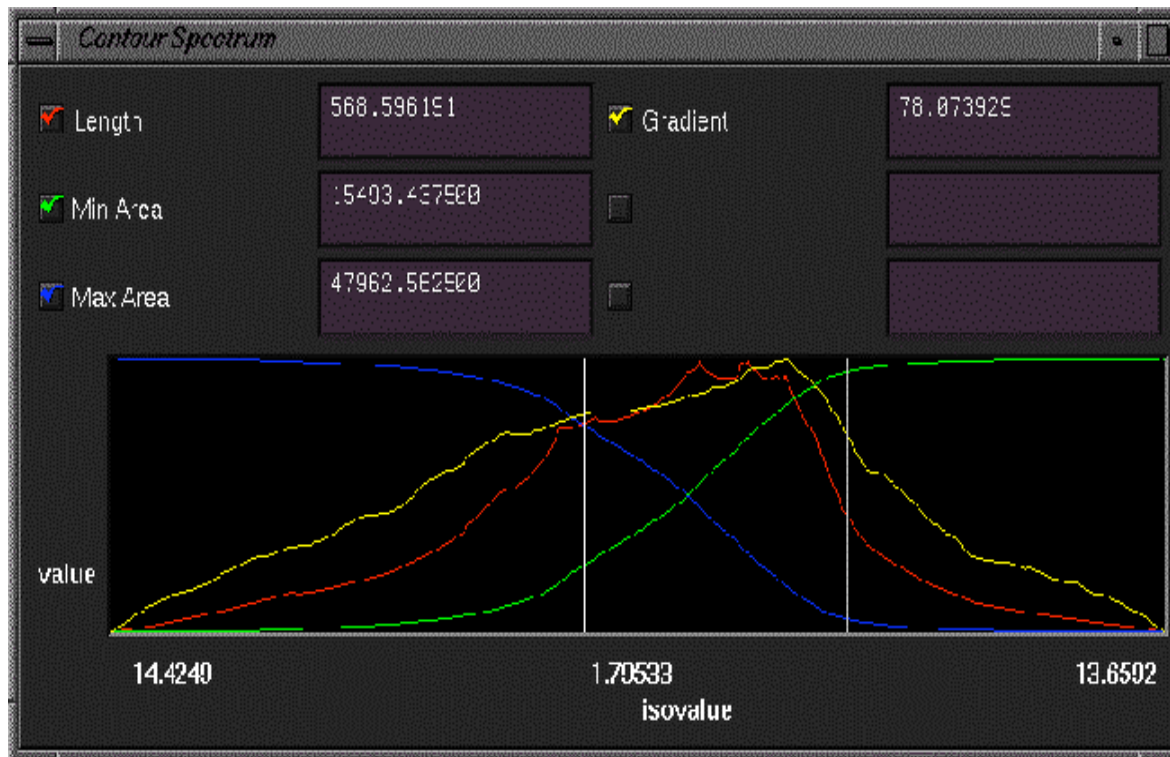
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# Spectral Analysis

## Graphical User Interface for Static Data



- The horizontal axis spans the scalar values  $\alpha$ .
- Plot of a set of signatures (length, area, gradient ...) as functions of the scalar value  $\alpha$ .

- Vertical axis spans normalized ranges of each



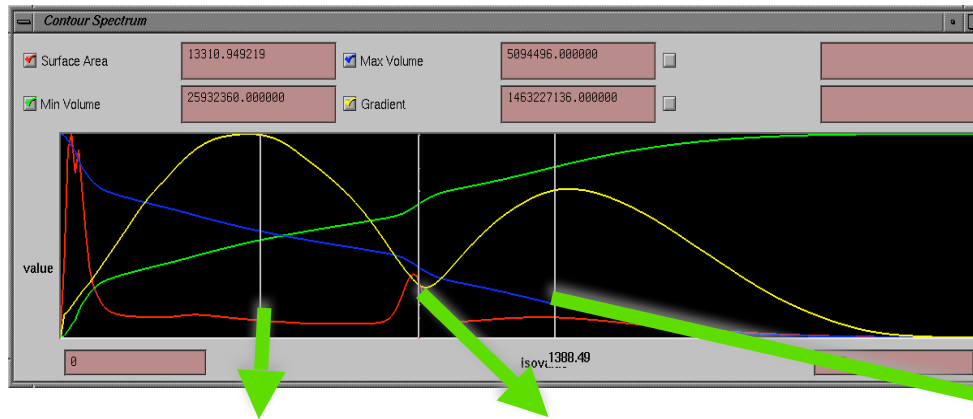
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Department of Computer Sciences

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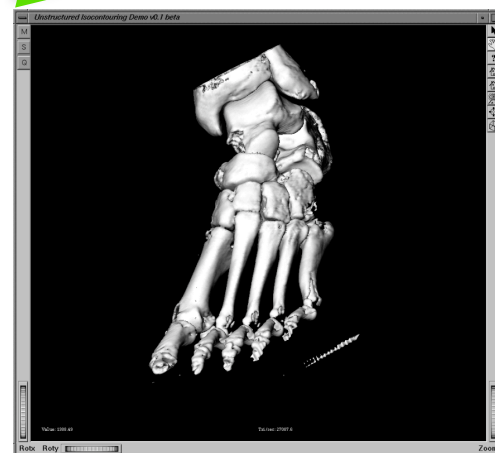
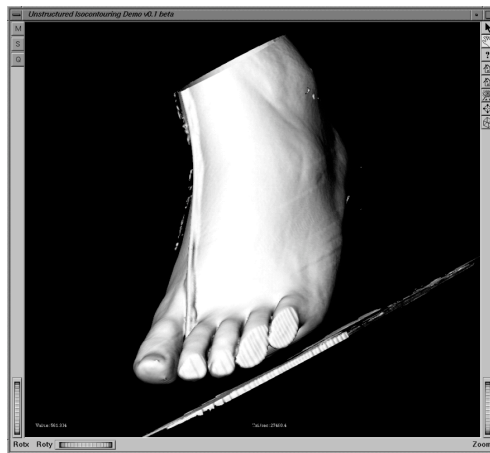
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## Contouring based Selection

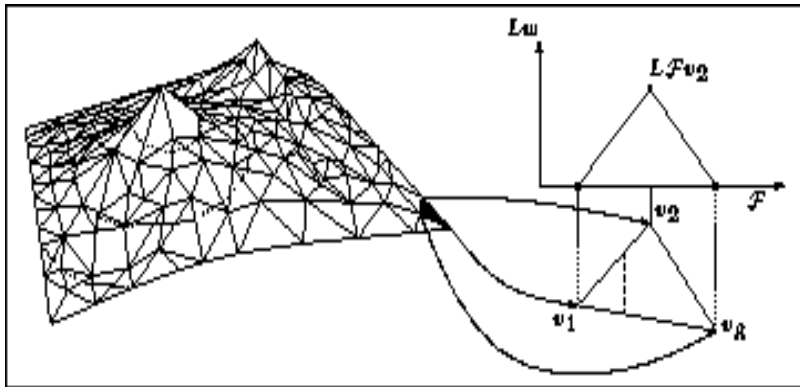


*The contour spectrum allows the development of an adaptive ability to separate interesting isovalues from the others.*





# Spectral Analysis (signature computation)

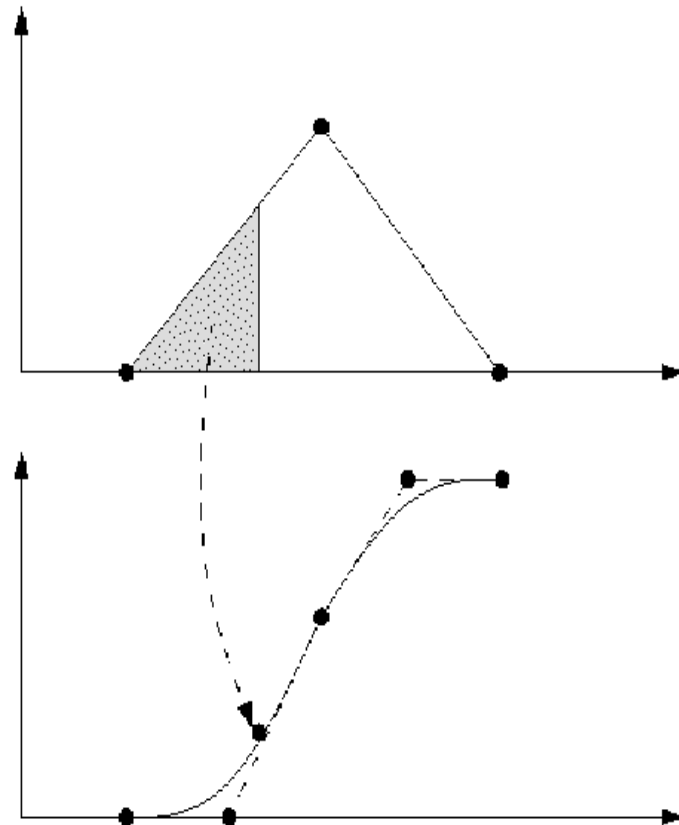


- The length of each contour is a  $C^0$  spline function.

The area inside/outside each isocontour is a  $C^1$  spline function.



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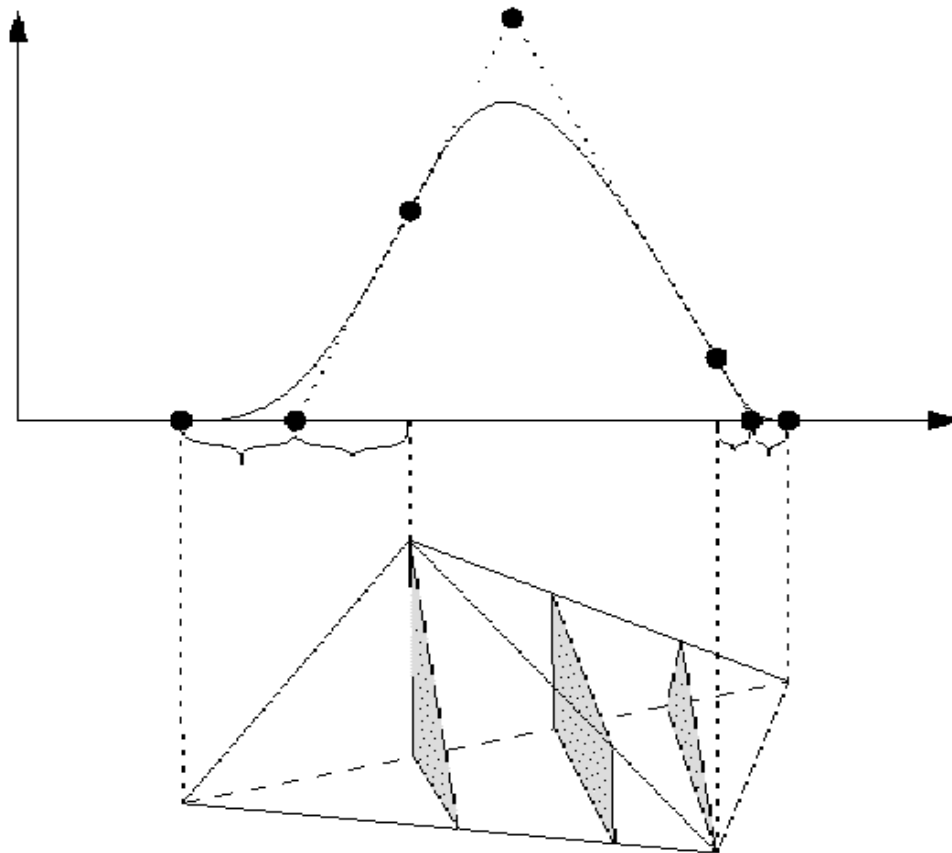


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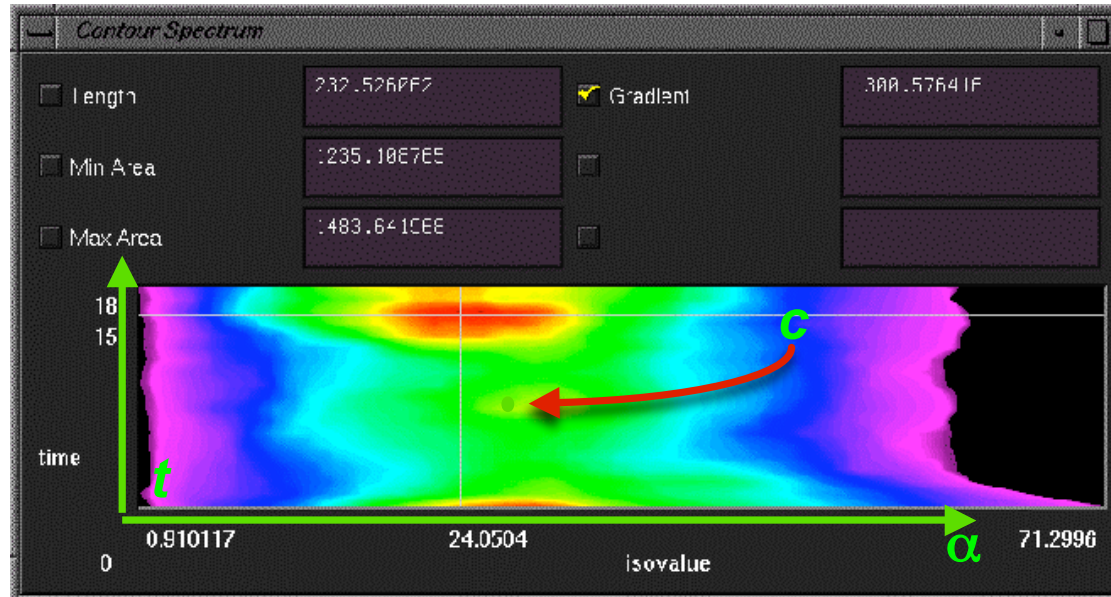
# Spectral Analysis (signature computation)

- In general the size of each isocontour of a scalar field of dimension  $d$  is a spline function of  $d-2$  continuity.
- The size of the region inside/outside is given by a spline function of  $d-1$  continuity



# Spectral Analysis

## Graphical User Interface for time varying data



The horizontal axis spans the scalar value dimension  $\alpha$

The vertical axis spans the time dimension  $t$

high  
 $(\alpha, t) \rightarrow c$   
The color  $c$   
is mapped  
to the  
magnitude  
of a  
signature  
function of  
time  $t$  and  
isovalue  $\alpha$   
low

magnitude



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## Further Reading

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