



Linux Clustering Software

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Surface Reconstruction from Point Clouds



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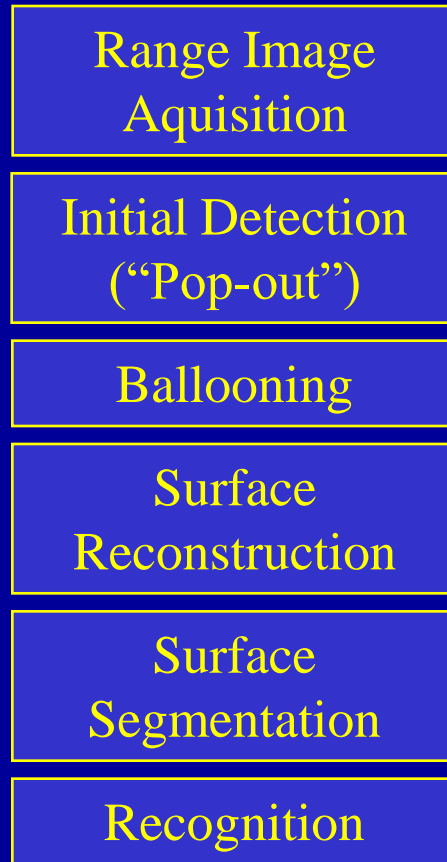
The Ohio State University

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Topic Motivation

- Computationally intensive work
- Many “small” execution jobs at each phase



Multiple balloons must be merged:

- Segmentation requires dense meshes
- Non-sparse balloons are “leaky”
- Merging overlapping balloons is non-trivial

Outline

- Part 1: Linux Clustering Software
 - ClusterIt
 - Portable Batch System
- Part 2: Surface Reconstruction
 - Preliminaries
 - Curve Reconstruction
 - Cocone Algorithm
 - Undersampling
 - Some Results
- Current Status
- Further Reading





Cluster Software: ClusterIt

- What is it?
 - Simultaneous execution on a set of Linux boxes
 - Execution on “any” Linux box
(but PBS is much better)
- What do I have to do to use it?
 - See <http://sampl.eng.ohio-state.edu/~dalleyg/faq/index.html> for instructions on configuring SSH



Cluster Software:

SAMPL Cluster Groups

Group	Comments
a	All processors (use with <i>jsd</i> , etc.)
a1	All boxes (use with <i>dsh</i>, etc.)
b	All Beowulf processors
b1	All Beowulf boxes
m	All of the Micron boxes
dc	All processors on the dual Celerons
dc1	All dual Celeron boxes

See `/etc/dsh.cluster` on a Linux box for the full set of groups



Cluster Software: dsh

- When would I use it?
 - Same command, many machines
 - Batch job preparation
 - Administrative work
- Usage
 - `dsh -g om 'hostname -i'`
Group (required!) Remote command (quoted)
 - Results
 - `samp101: 164.107.162.9`
 - `samp102: 164.107.162.10`
 - `samp110: 164.107.162.18`
 - `samp112: 164.107.162.20`



Cluster Software: dsh Examples

- Good examples
 - `dsh -g om 'hostname -i'`
 - `dsh -g om 'hostname > /tmp/tmpfile'`
 - `dsh -g al 'smbmount //samplf04/dalleyg /u/dalleyg -o umask=077,dmask=077, password=PASSWORD'`
- What to be careful about
 - Using quotes (single or double)
 - Redirecting input or output



Cluster Software:

Other ClusterIt Examples

- **pcp -g a1 datafile /tmp/datafile**
 - Copies file **datafile** from the current directory to **/tmp/datafile** on all machines in the **a1** group
 - Useful for copying files to local temporary storage
- **prm -g a1 /tmp/datafile**
 - Deletes **/tmp/datafile** on all machines in the **a1** group
 - Useful for cleanup of local temporary storage, etc.



Cluster Software:

Portable Batch System

- What is it?
 - Manages submission and execution of large batch jobs
 - Allows balancing between users
 - Used for single multi-processor machines all the way up to compute farms with many Crays (e.g. OSC)



Cluster Software:

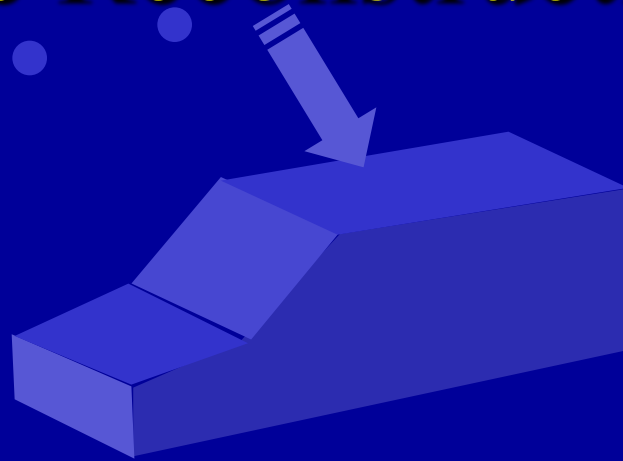
Portable Batch System (cont'd.)

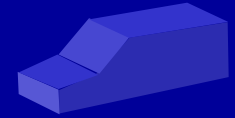
- Submitting a job (from *eepc359* **only**)*
 - `qsub -N jobname jobprogram`
 - `qsub -N Test /home/dalleyg/nfs/test2.pl`
- Cluster status
 - `qstat -f -B`
- **xpbs**
 - Helps you create job scripts, configure advanced options, etc.
- **xpbsmon**
 - Helps you monitor the state of the machines on the cluster



Part 2:

Surface Reconstruction...

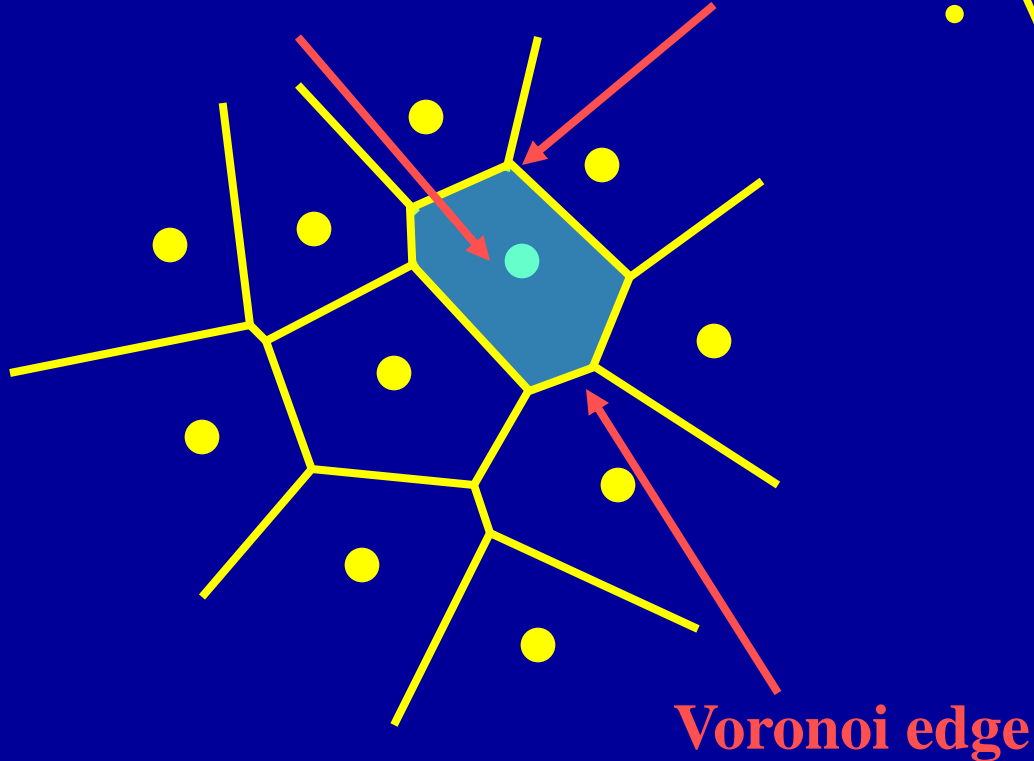




Voronoi Diagrams

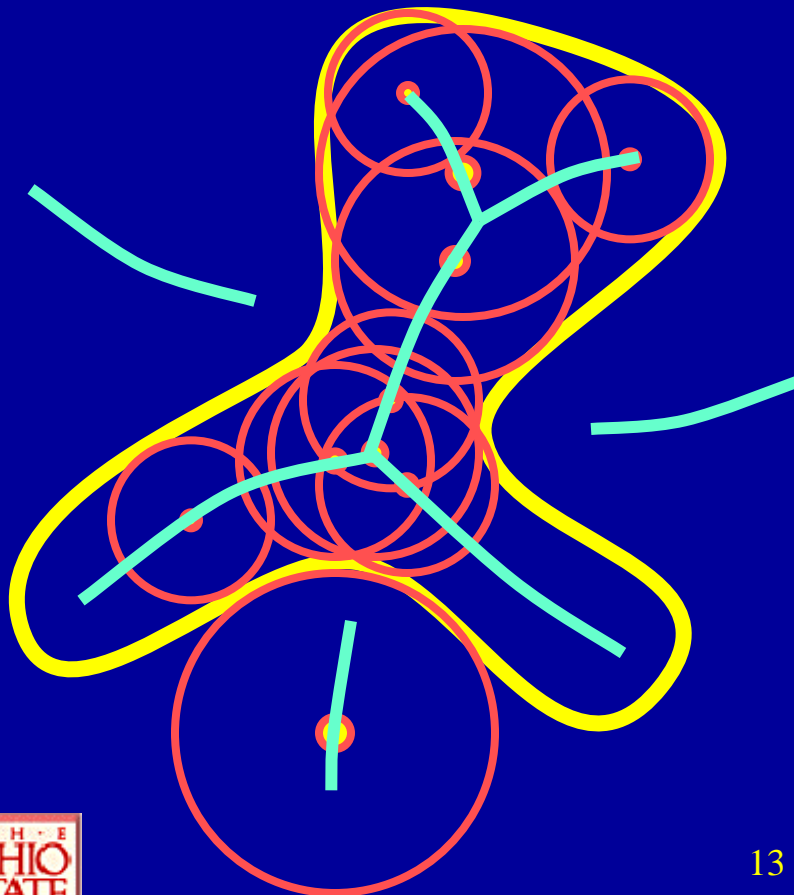
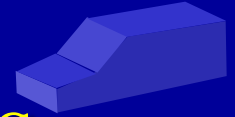
Voronoi cell

Voronoi vertex



- Voronoi Cell of x
 - The set of points that are closer to x than to any other sample point

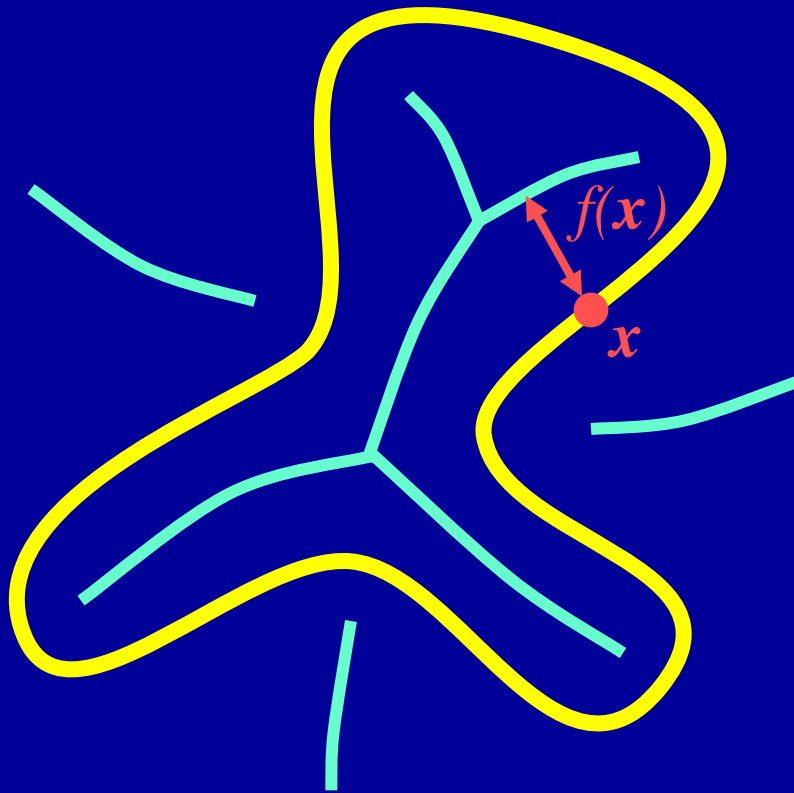
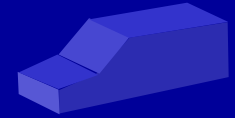
Preliminaries: Medial Axis



- Medial Axis:
 - Find all circles that tangentially touch the curve in at least 2 points
 - Medial axis = centers of all those circles

Surface Reconstruction Preliminaries:

ε -sampling



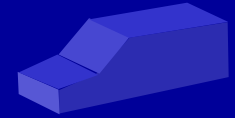
$f(x) \equiv$ feature size at point x
= distance to the medial axis
at point x

Sampling criterion:

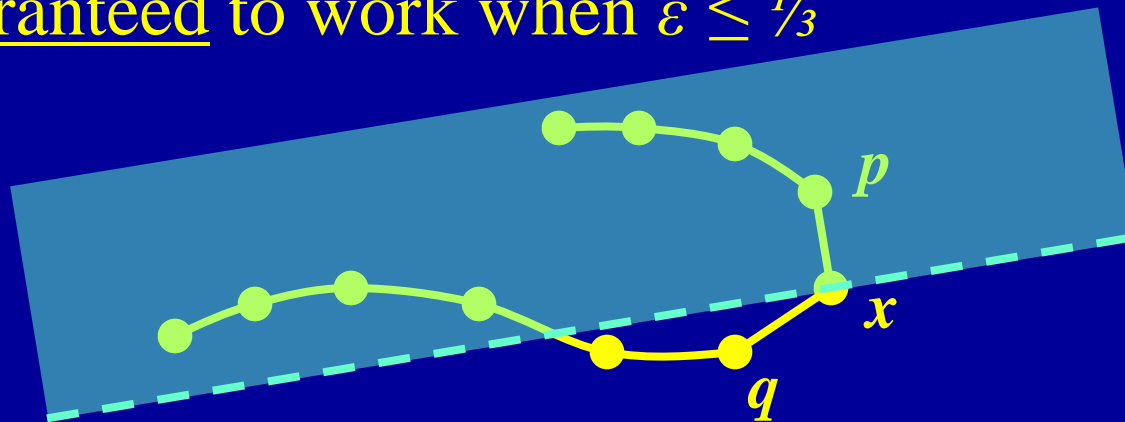
each sample point x is at
most $\varepsilon f(x)$ from the next
closest sample ($0 < \varepsilon < 1$,
typically).

Important note: When ε is
small, the curve locally
looks flat

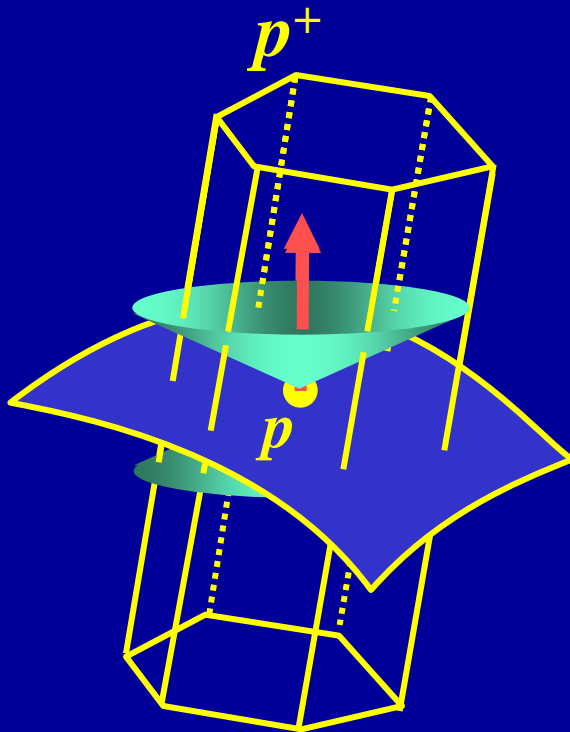
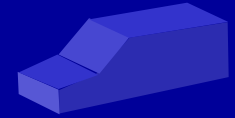
Surface Reconstruction: Curve Reconstruction



- Algorithm:
 - Find the closest point, p , to x and connect them
 - Find the closest point, q , to x such that the angle pxq is at least 90° .
- Guaranteed to work when $\varepsilon \leq 1/3$



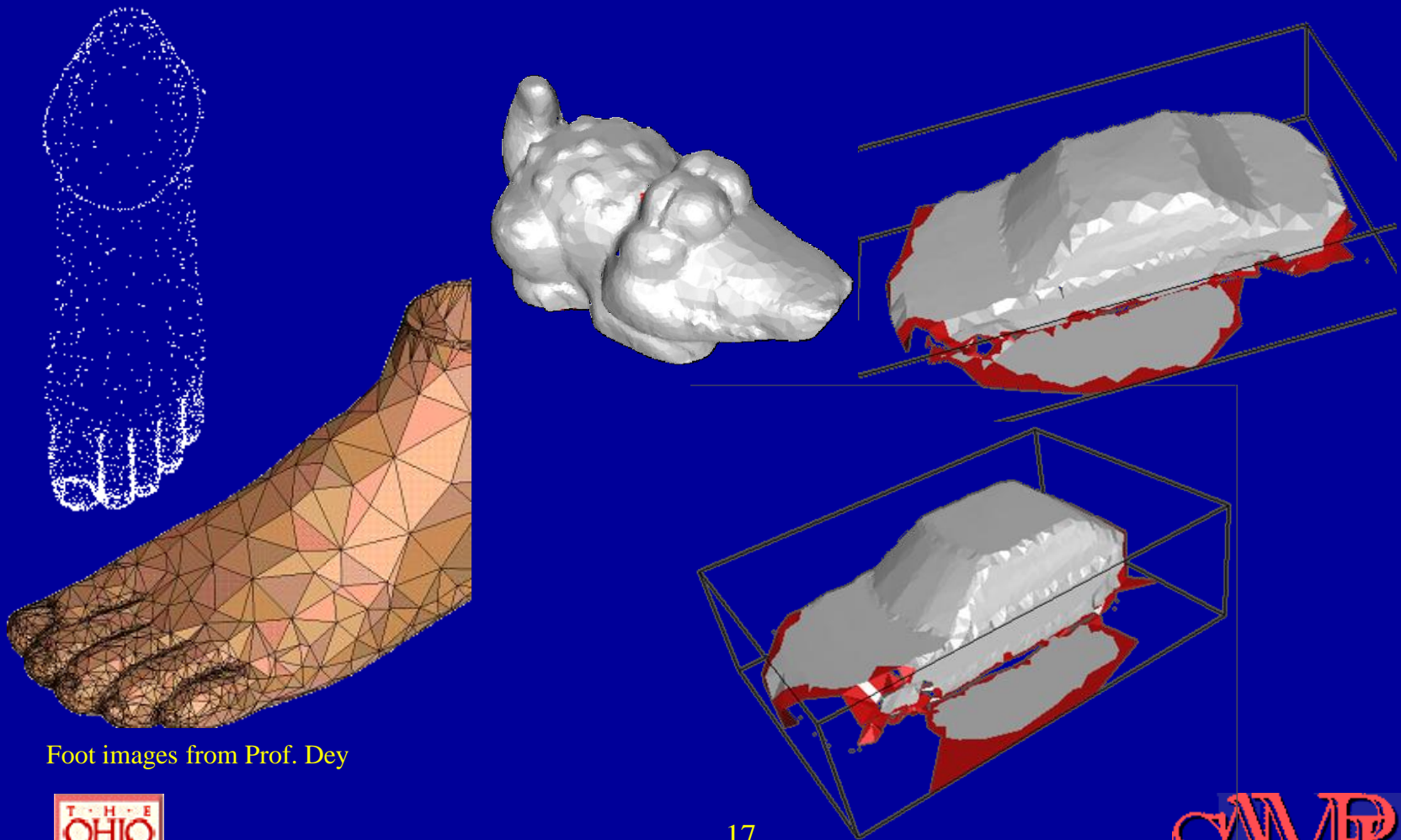
Surface Reconstruction: Cocone Algorithm



Voronoi cell of p

- $p^+ \equiv$ pole of p = point in the Voronoi cell farthest from p
- $\varepsilon < 0.1 \rightarrow$
 - the vector from p to p^+ is within $\pi/8$ of the true surface normal
 - The surface is nearly flat within the cell

Sample Reconstructed Surfaces



Foot images from Prof. Dey

TuT Status

Range Image
Aquisition

Software written

Initial Detection
("Pop-out")

See Kanu

Ballooning

Software written (by Kanu), generating data

Surface
Reconstruction

Software written (by Prof. Dey)

Surface
Segmentation

Evaluating several variants

Recognition

To be done

Further Reading

– Clusterit

- <http://www.garbled.net/clusterit.html>
- <http://sampl.eng.ohio-state.edu/~dalleyg/faq/index.html>

– Portable Batch System

- <http://www.openpbs.org>
 - Official web site (ask if you need the password)
- http://www-itg.lbl.gov/Grid/public/pbs/pbs.v2.3_admin.pdf
 - Detailed administration/usage guide
- <http://www-math.cudenver.edu/~jmandel/mri/Schedulers-overview.pdf>
 - Slideshow introduction to PBS by Doug Johnson from the Ohio Supercomputer Center

– Surface Reconstruction

- <http://www.cis.ohio-state.edu/~tamaldey/>
- N. Amenta, S. Choi, T. K. Dey and N. Leekha. [A simple algorithm for homeomorphic surface reconstruction](#). Proc. 16th ACM Symposium on Computational Geometry, 2000, 213--222.

