

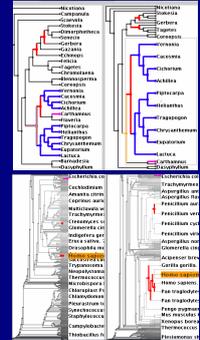
Information Visualization with Accordion Drawing

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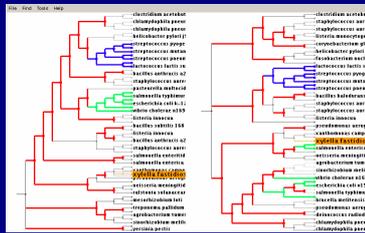
Accordion Drawing

- rubber-sheet navigation
 - stretch out part of surface, the rest squishes
 - borders nailed down
 - Focus+Context technique
 - integrated overview, details
 - old idea
 - [Sarkar et al 93], ...
- guaranteed visibility
 - marks always visible
 - important for scalability
 - new idea
 - [Munzner et al 03]



Guaranteed Visibility

- marks are always visible
- easy with small datasets



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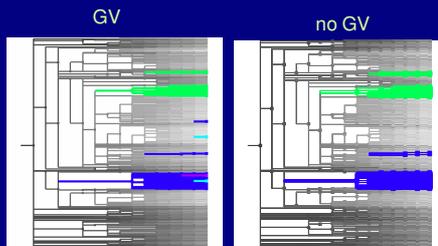
Guaranteed Visibility Challenges

- hard with larger datasets
- reasons a mark could be invisible
 - outside the window
 - AD solution: constrained navigation
 - underneath other marks
 - AD solution: avoid 3D
 - smaller than a pixel
 - AD solution: smart culling

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Guaranteed Visibility: Small Items

- naive culling may not draw all marked items

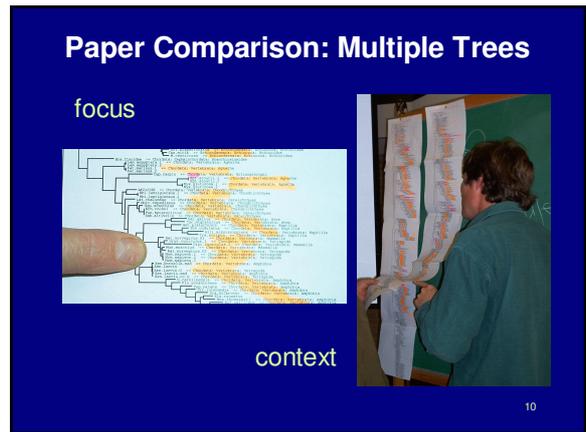
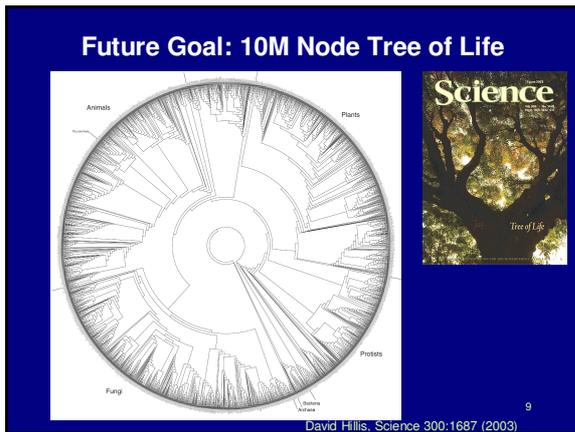
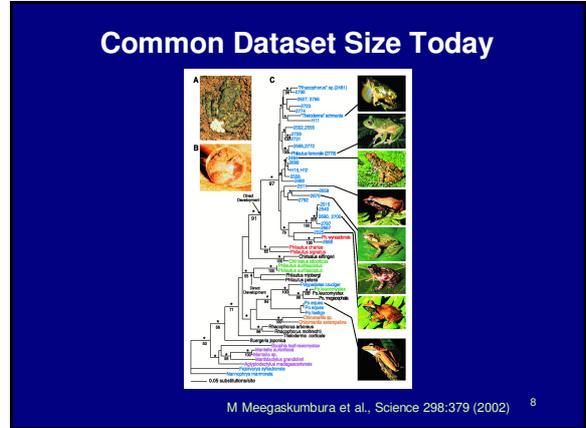
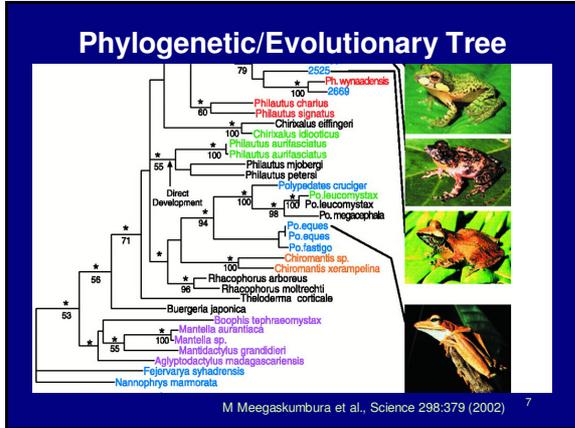


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Outline

- trees
 - TreeJuxtaposer
- sequences
 - SequenceJuxtaposer
- scaling up trees
 - TJC
- general AD framework
 - PRISAD
- power sets
 - PowerSetViewer
- evaluation

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TreeJuxtaposer

- comparison of evolutionary trees
 - side by side
- demo
 - olduvai.sf.net/tj

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TJ Contributions

- first interactive tree comparison system
 - automatic structural difference computation
 - guaranteed visibility of marked areas
- scalable to large datasets
 - 250,000 to 500,000 total nodes
 - all preprocessing subquadratic
 - all realtime rendering sublinear
- introduced accordion drawing (AD)
- introduced guaranteed visibility (GV)

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Joint Work: TJ Credits

Tamara Munzner, Francois Guimbretiere, Serdar Tasiran, Li Zhang, and Yunhong Zhou.

TreeJuxtaposer: Scalable Tree Comparison using Focus+Context with Guaranteed Visibility.
SIGGRAPH 2003

www.cs.ubc.ca/~tmm/papers/tj

James Slack, Tamara Munzner, and Francois Guimbretiere.

TreeJuxtaposer: InfoVis03 Contest Entry. (Overall Winner)
InfoVis 2003 Contest

www.cs.ubc.ca/~tmm/papers/contest03

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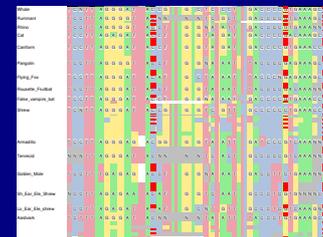
Genomic Sequences

- multiple aligned sequences of DNA
- now commonly browsed with web apps
 - zoom and pan with abrupt jumps
- check benefits of accordion drawing
 - smooth transitions between states
 - guaranteed visibility for globally visible landmarks

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SequenceJuxtaposer

- dense grid, following conventions
 - rows of sequences partially correlated
 - columns of aligned nucleotides
 - videos



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SJ Contributions

- accordion drawing for gene sequences
- paper results: 1.7M nucleotides
 - current with PRISAD: 40M nucleotides
- joint work: SJ credits

James Slack, Kristian Hildebrand, Tamara Munzner, and Katherine St. John.

SequenceJuxtaposer: Fluid Navigation For Large-Scale Sequence Comparison In Context.

Proc. German Conference on Bioinformatics 2004

www.cs.ubc.ca/~tmm/papers/sj

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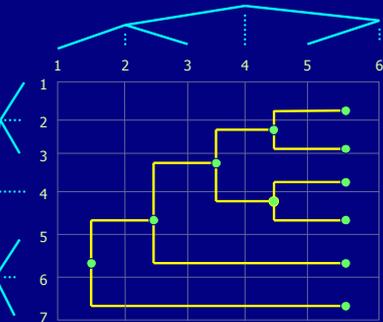
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Scaling Up Trees

- TJ limits
 - large memory footprint
 - CPU-bound, far from achieving peak rendering performance of graphics card
- quadtree data structure used for
 - placing nodes during layout
 - drawing edges given navigation
 - culling edges with GV
 - selecting edges during interaction

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Navigation Without Quadtrees



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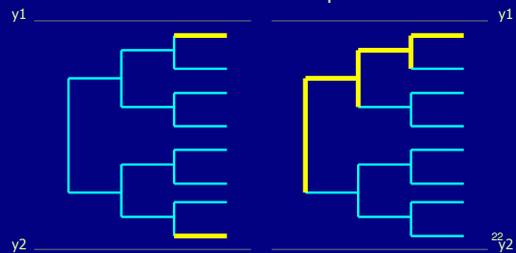
Eliminating the Quadtree

- new drawing algorithm
 - addresses both ordering and culling
- new way to pick edges
 - uses advances in recent graphics hardware
- find a different way to place nodes
 - modification of O-buffer for interaction

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Drawing the Tree

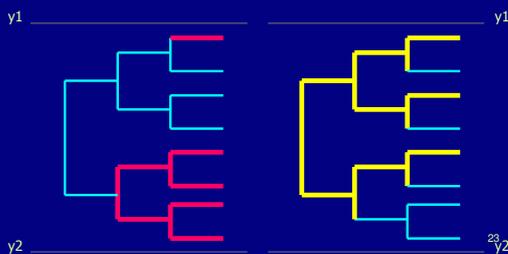
- continue recursion only if sub-tree vertical extent larger than apixel
 - otherwise draw flattened path



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Guaranteed Visibility

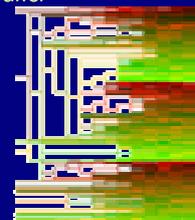
- continue recursion only if subtree contains both marked and unmarked nodes



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Picking Edges

- Multiple Render Targets
 - draw edges to displayed buffer
 - encoding edge identifier information in auxiliary buffer



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TJC/TJC-Q Results

- TJC
 - no quadtree
 - requires HW multiple render target support
 - 15M nodes
- TJC-Q
 - lightweight quadtree
 - 5M nodes
- both support tree browsing only
 - no comparison data structures

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Joint Work: TJC, TJC-Q Credits

Dale Beermann, Tamara Munzner, and Greg Humphreys.
 Scalable, Robust Visualization of Large Trees.
 Proc. EuroVis 2005
www.cs.virginia.edu/~gfx/pubs/TJC

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PRISAD

- generic accordion drawing infrastructure
 - handles many application types
- efficient
 - guarantees of correctness: no overculling
 - tight bounds on overdrawing
 - handles dense regions efficiently
 - new algorithms for rendering, culling, picking
 - exploit application dataset characteristics instead of requiring expensive additional data structures

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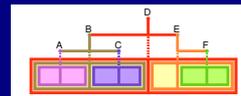
PRISAD vs Application Interplay

| | Application | PRISAD | |
|----------------------------|-------------|------------------------------------|-----------------------|
| World-space discretization | Layout | (x, y) size Initialize | |
| | Gridding | {S _x , S _y } | |
| | | S, node Mapping | |
| Screen-space rendering | Render | S, τ Partition | |
| | Seed | S ranges | |
| | | Queue | |
| | Draw | Object | Progressive Rendering |
| | | | |

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PRISAD Responsibilities

- initializing a generic 2D grid structure
 - split lines: both linear ordering and recursive hierarchy



- mapping geometric objects to world-space structures
- partitioning a binary tree data structure into adjacent ranges
- controlling drawing performance for progressive rendering

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Application Responsibilities

- calculating the size of underlying PRISAD structures
- assigning dataset components to PRISAD structures
- initiating a rendering action with two partitioning parameters
- ordering the drawing of geometric objects through seeding
- drawing individual geometric objects

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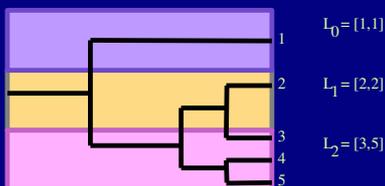
Example: PRITree

- rendering with generic infrastructure
 - partitioning
 - rendering requires sub-pixel segments
 - partition split lines into leaf ranges
 - seeding
 - 1st: roots of marked sub-trees, marked nodes
 - 2nd: interaction box, remainder of leaf ranges
 - drawing
 - ascent rendering from leaves to root

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Tree Partitioning

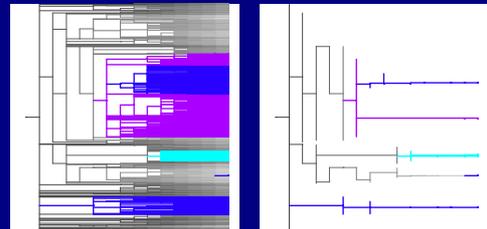
- divide leaf nodes by screen location
 - partitioning follows split line hierarchy
 - tree application provides stopping size criterion
 - ranges [1,1]; [2,2]; [3,5] are partitions



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Tree Seeding

- marked subtrees not drawn completely in first frame
 - draw “skeleton” of marks for each subtree for landmarks
 - solves guaranteed visibility of small subtree in big dataset



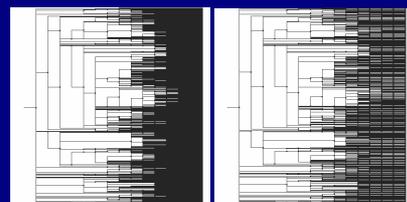
Tree Drawing Traversal

- ascent-based drawing
 - partition into leaf ranges before drawing
 - TreeJuxtaposer partitions during drawing
 - start from 1 leaf per range, draw path to root
 - carefully choose starting leaf
 - 3 categories of misleading gaps eliminated
 - leaf-range gaps
 - horizontal tree edge gaps
 - ascent path gaps

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Leaf-range Gaps

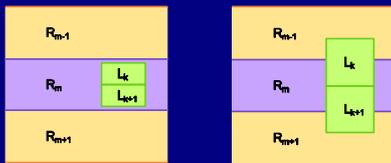
- number of nodes rendered depends on number of partitioned leaf ranges
 - maximize leaf range size to reduce rendering
 - too much reduction results in gaps



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Eliminating Leaf-range Gaps

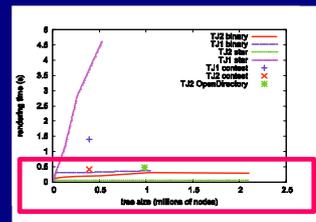
- eliminate by rendering more leaves
 - partition into smaller leaf ranges



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Rendering Time Performance

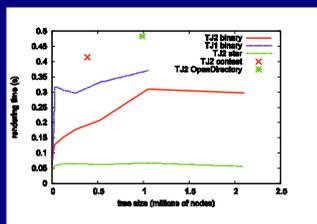
- TreeJuxtaposer renders **all** nodes for star trees
 - branching factor k leads to $O(k)$ performance
- we achieve 5x rendering improvement with contest comparison dataset
- constant time, after threshold, for large binary trees



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Rendering Time Performance

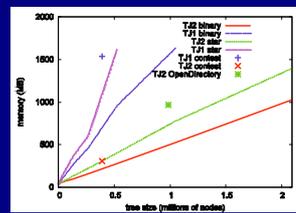
- constant time, after threshold, for large binary trees
 - we approach rendering limit of screen-space
- contest and OpenDirectory comparison render 2 trees
 - comparable to rendering two binary trees



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Memory Performance

- linear memory usage for both
 - generic AD approach 5x better
- marked range storage changes improve scalability
 - 1GB difference for contest comparison



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PRISAD Results

- video
- joint work: PRISAD credits

James Slack, Kristian Hildebrand, and Tamara Munzner.
 PRISAD: A Partitioned Rendering Infrastructure for Scalable Accordion Drawing.
 Proc. InfoVis 2005, to appear

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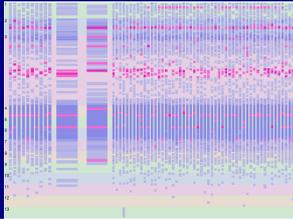
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PowerSetViewer

- data mining market-basket transactions
 - items bought together make a set
 - space of all possible sets is power set
 - place logged sets within enumeration of power set



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PSV Results

- dynamic data
 - show progress of steerable data mining system with constraints
 - all other AD applications had static data
- handles alphabets of up to 40,000
- handles log files of 1.5 to 7 million items
- joint work in progress with
 - Qiang Kong, Raymond Ng

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Evaluation

- how focus and context are used with
 - rubber sheet navigation vs. pan and zoom
 - integrated scene vs. separate overview
- user studies of TJ
 - tasks based on biologist interviews
- joint work in progress, with
 - Adam Bodnar, Dmitry Nekrasovski, Joanna McGrenere

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Conclusion

- accordion drawing effective for variety of application datasets
 - trees, sequences, sets
- guaranteed visibility is powerful technique
 - computational expense can be handled by generic algorithms

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More Information

- papers, videos, images
 - www.cs.ubc.ca/~tmm
- free software
 - olduvai.sourceforge.net/tj
 - olduvai.sourceforge.net/sj

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