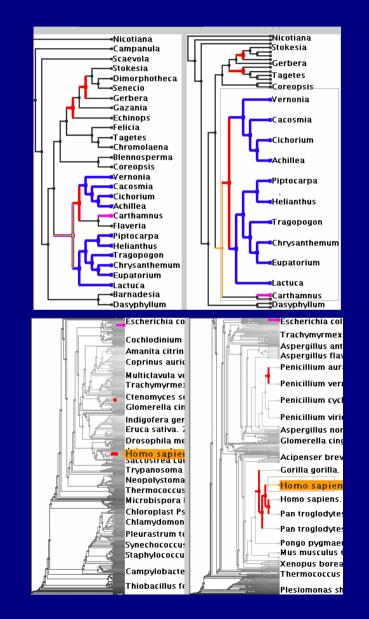
Information Visualization with Accordion Drawing

> Tamara Munzner University of British Columbia

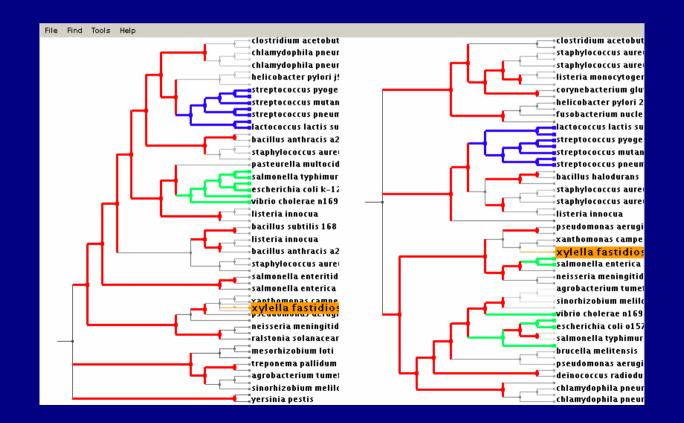
# **Accordion Drawing**

- 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



## **Guaranteed Visibility Challenges**

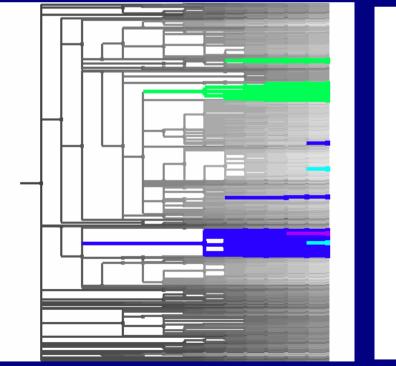
- hard with larger datasets
- reasons a mark could be invisible
  - outside the window
    - AD solution: constrained navigation
  - underneath other marks
    - AD solution: avoid 3D
  - smaller than a pixel
    - AD solution: smart culling

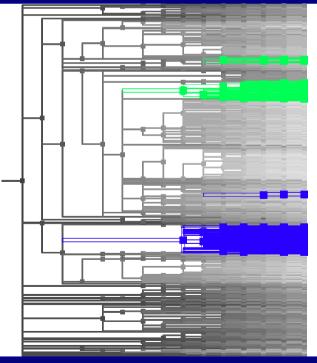
## **Guaranteed Visibility: Small Items**

naive culling may not draw all marked items





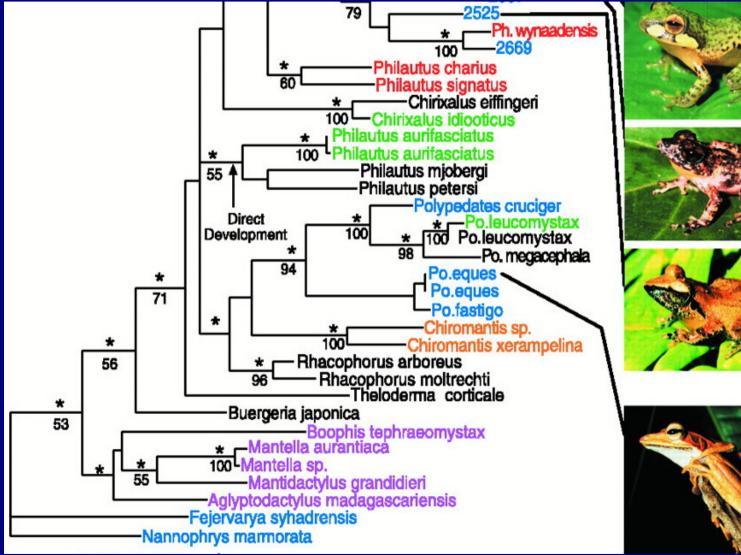




# Outline

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

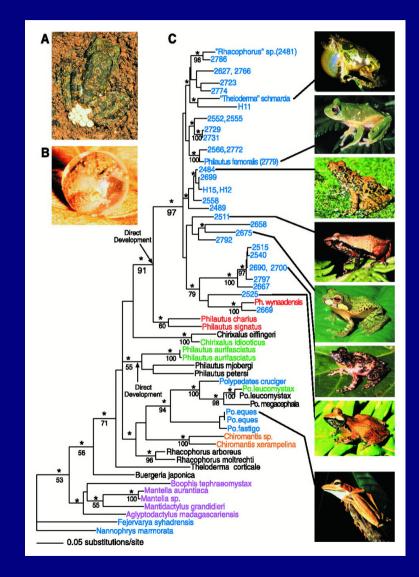
## **Phylogenetic/Evolutionary Tree**



7

M Meegaskumbura et al., Science 298:379 (2002)

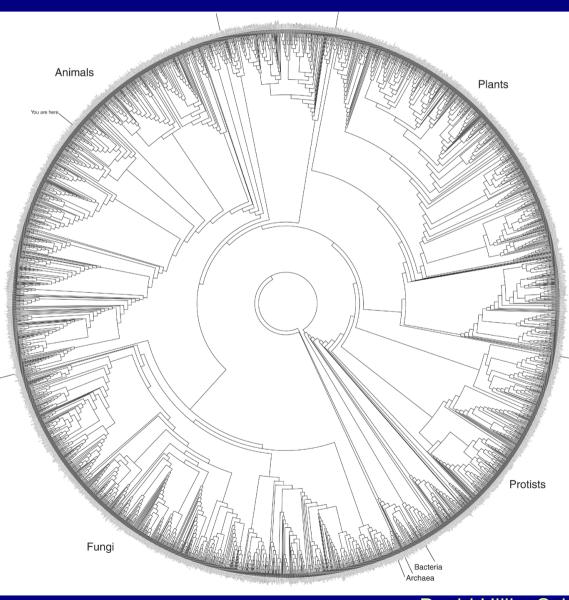
#### **Common Dataset Size Today**

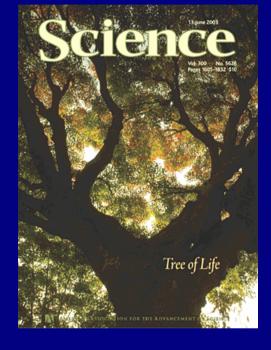


M Meegaskumbura et al., Science 298:379 (2002)

8

#### Future Goal: 10M Node Tree of Life



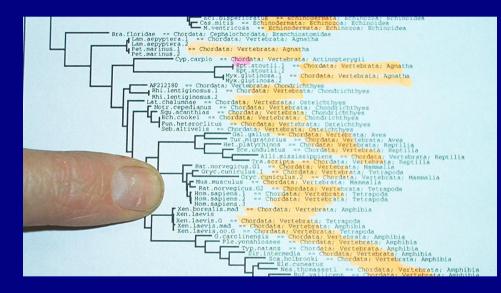


9

David Hillis, Science 300:1687 (2003)

## **Paper Comparison: Multiple Trees**

#### focus

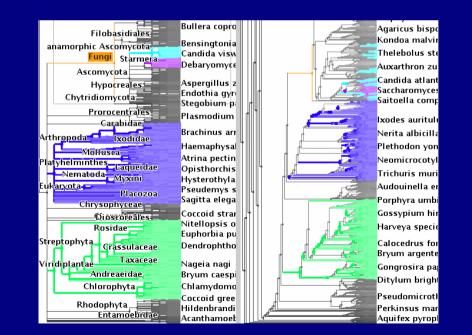


#### context



#### TreeJuxtaposer

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



# **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)

#### 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

#### SequenceJuxtaposer

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

Whale	Т	С	Ν	Т	Т	А	G	G	G	А	Т	A (	C C	G	G	С	Т	С	С	СТ	G	А	С	С	СС	G	т	G A	AA	۹ G	Сu	, A
Ruminant	т	С	С	т	Т	А	G	G	G	G	Т	A		I N	N	N	Т	С	G	с т	G	А	С	С	СС	G	T C	C A	AA	A G	сc	; A
Rhino	Т	С	С	т	Т	А	G	G	G	G	Т	A	C C	T	G	G	Ν	А	А	тт	G	А	С	С	СC	G	то	G A	AA	A N	NC	, A
Cat	т	С	С	т	T	A	G	A	G	А	Т	A	5 0	T	G	G	т	А	G	A T	G	A	С	С	СС	G	A (	C A	A A	۹ G	сc	; A
Caniform	т	С	С	т	т	A	G	G	G	A	т	A (	с с	т	G	G	т	A	G	ΑT	G	A	С	С	сс	G	тс	G A	AA	۹ C	сc	A
Pangolin	т	с	С	т	т	A	G	G	G	G	т	A	c c	т	G	G	N	А	A	ΑT	т	A	С	С	СС	G	A C	G A	AA	A G	cc	Â
Flying_Fox	т	С	С	т	т	A	G	G	G	A	т	A (	C A	т	G	С	т	А	A	A T	т	A	С	С	CN	I G	A (	G A	AA	A G	cc	A
Rousette_Fruitbat	т	С	С	т	т	A	G	G	G	А	т	A	c c	T	G	G	т	А	А	ΑT	т	A	С	С	СС	G	A C	C A	AA	A N	NC	; A
False_vampire_bat	т	С	С	Т	т	А	G	G	G	А	т	A	c c	т	G	G	N	А	А	A T	G	A	С	С	сс	G	ТС	G A	AA	A C	СС	, A
Shrew	т	С	N	т	т	A	G	G	G	A	т	A	с с	G	G	G	т	С	G	тт	G	С	С	С	сс	с	тс	G A	AA	۹ C	СС	; A
																										Ē						
Armadillo	т	С	С	т	т	A	G	G	G	A	G	A (	c 0	G	G	G	т	A	A	тт	G	A	Т	С	сс	G	тс	C A	AA	A N	NC	; A
Tenrecid	N	N	N	т	Т	A	G	G	G	A	т	A (	C N	I N	N	N	т	С	A	СТ	G	С	С	С	сс	G	тс	C A	AA	A N	N C	; A
Golden_Mole	т	с	С	т	т	т	G	A	G	A	G	Α (	c c	т	G	G	N	A	A	A T	G	A	С	С	тт	С	тс	G A	. A <i>A</i>	A N	N A	A
Sh_Ear_Ele_Shrew	N	С	С	т	т	A	G	A	G	A	A	A (	C A	Т	G	G	т	С	A	A T	G	С	С	С	тс	G	тс	G N	N	NN	сc	; A
Lo_Ear_Ele_shrew	т	С	С	т	т	A	G	A	G	A	т	A	r A	т	G	С	N	т	G	ΤТ	G	С	С	С	тс	G	тс	C A	AA	A G	NN	A
Aardvark	т	С	Т	т	Т	A	G	G	G	A	Т	A (		I N	N	N	Т	А	А	тт	Т	A	С	С	с т	G	тс	G A	AA	A G	C A	A

# **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

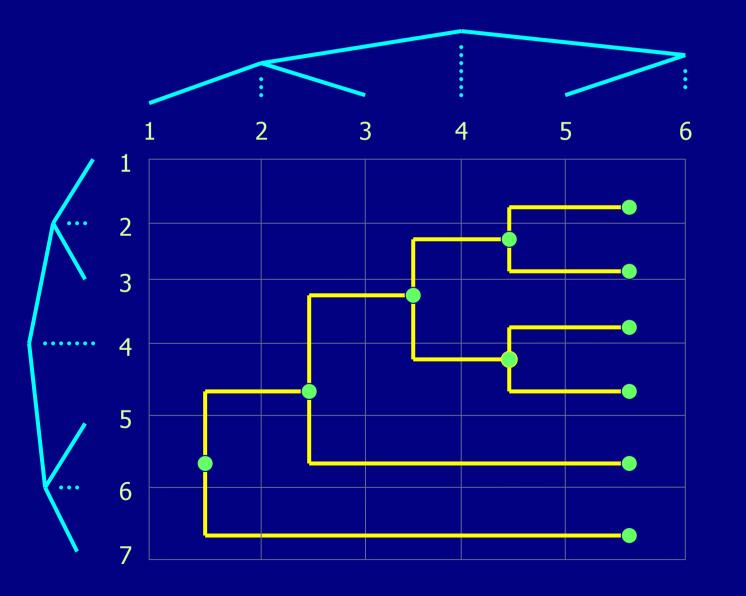
# Outline

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

# **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

# **Navigation Without Quadtrees**



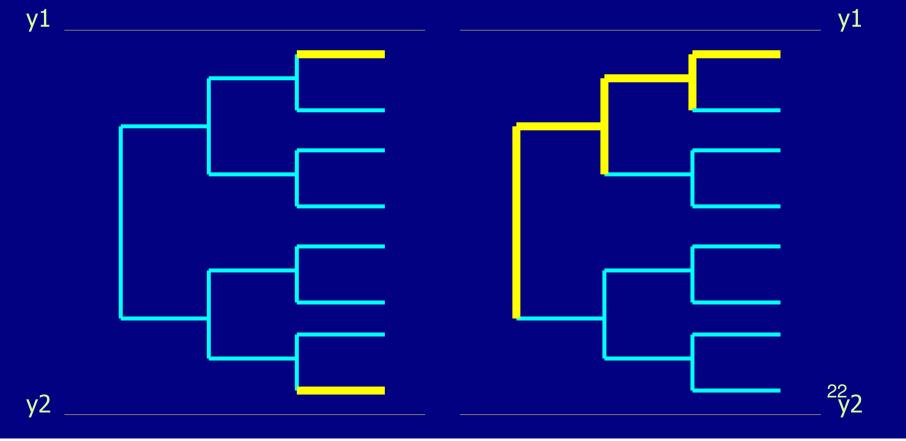
20

# **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

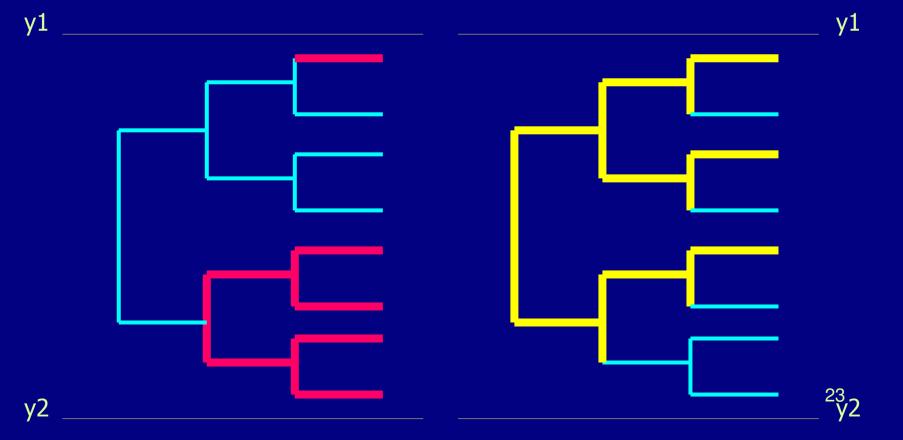
## **Drawing the Tree**

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



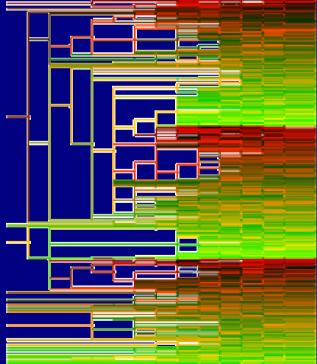
# **Guaranteed Visibility**

 continue recursion only if subtree contains both marked and unmarked nodes



# **Picking Edges**

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



# **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

#### 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

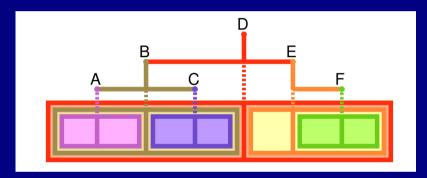
- generic accordion drawing infrastructure
  - handles many application types
- efficient
  - guarantees of correctness: no overculling
  - tight bounds on overdrawing
    - handles dense regions efficiently
  - new algorithms for rendering, culling, picking
    - exploit application dataset characteristics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

# **PRISAD vs Application Interplay**

	Application		PRISAD				
World-space discretization	Layout	(x, y) size	la iti a lin a				
	Gridding	$\{S_X,S_Y\}$	Initialize				
	andding	S, node					
			Mapping				
Screen-space	Render	S, τ					
rendering		S, 1					
		S ranges	Partition				
	Seed						
		Queue					
		Object	Progressive Rendering				
		Object	richidening				
	Draw						

# **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

# **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

#### **Example: PRITree**

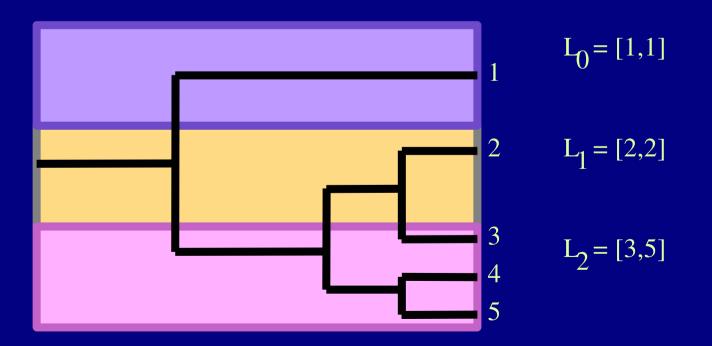
- rendering with generic infrastructure
  - partitioning
    - rendering requires sub-pixel segments
    - partition split lines into leaf ranges
  - seeding
    - 1<sup>st</sup>: roots of marked sub-trees, marked nodes
    - 2<sup>nd</sup>: interaction box, remainder of leaf ranges

- drawing

ascent rendering from leaves to root

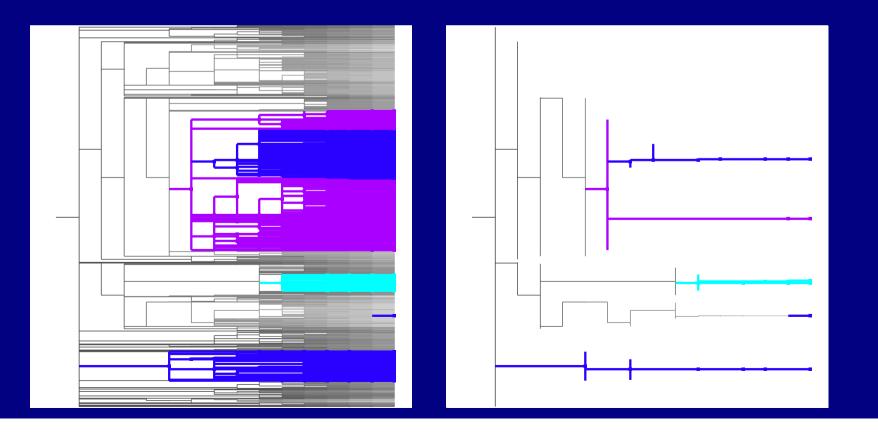
#### **Tree Partitioning**

- 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



#### **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



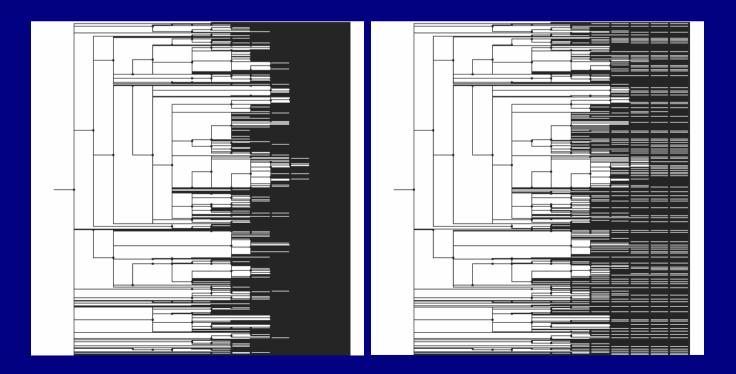
#### **Tree Drawing Traversal**

# 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

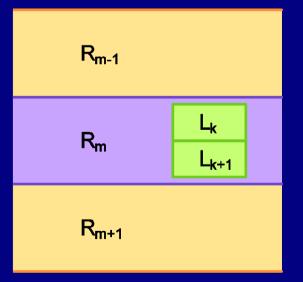
#### Leaf-range Gaps

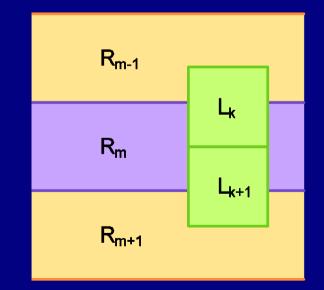
- number of nodes rendered depends on number of partitioned leaf ranges
  - maximize leaf range size to reduce rendering
  - too much reduction results in gaps



### **Eliminating Leaf-range Gaps**

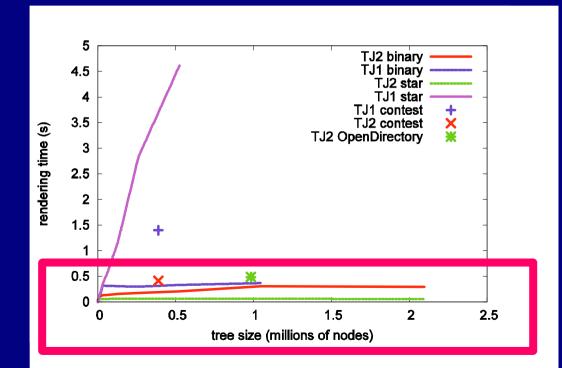
eliminate by rendering more leaves
 – partition into smaller leaf ranges





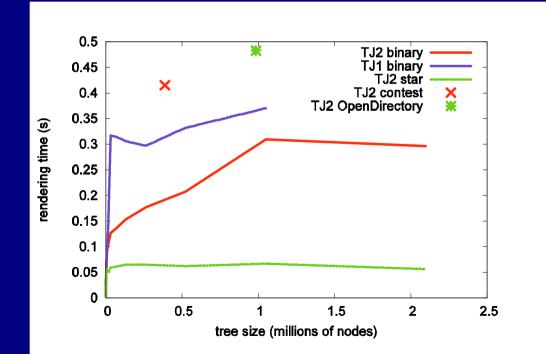
### **Rendering Time Performance**

- 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



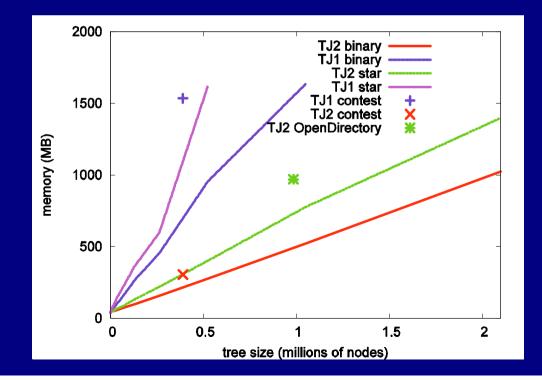
### **Rendering Time Performance**

- 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



#### **Memory Performance**

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



### **PRISAD Results**

#### video

joint work: PRISAD credits

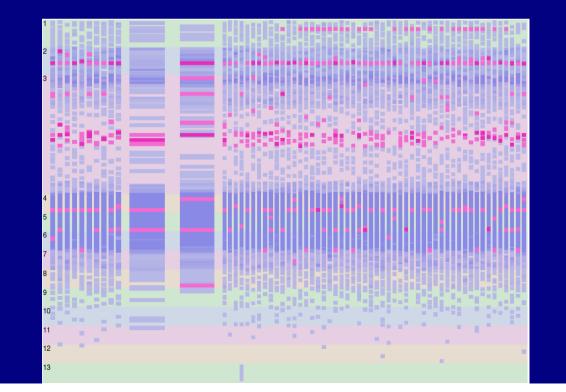
James Slack, Kristian Hildebrand, and Tamara Munzner. PRISAD: A Partitioned Rendering Infrastructure for Scalable Accordion Drawing. Proc. InfoVis 2005, to appear

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#### **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



### **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

## 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

### **More Information**

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