Methods for Visualizing Biodiversity & **Building Rewarding Collaborations**

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UBC Biodiversity Challenge Retreat, Hakai Institute 11 lune 2019

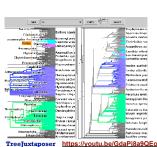
www.cs.ubc.ca/~tmm/talks.html#hakai19-methods

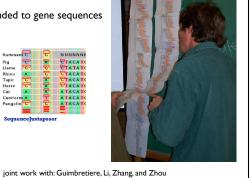


@tamaramunzner

• driving problem from UT Austin Hillis Lab in 2001 phylogenetic trees algorithm focus on scale, later extended to gene sequences

Tree uxtaposer: Visual tree comparison





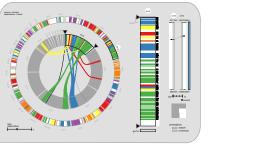
 Cytoscape plugin, funded by Agilent Cerebral https://youtu.be/76HhG1FQngl

multiple views with linked highlighting and navigation

automatic network layout by subcellular location, like hand-drawn diagrams Robert Kincaid

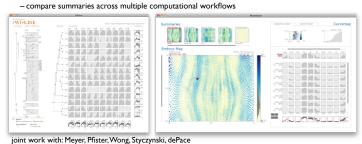
MizBee: Comparing genomes between species

- driving problems: Broad Inst. biologists studying fungus (Ma) and stickleback/pufferfish (Grabherr)
- two use phases: first fully validate computational pipeline, then can analyze biological questions
- investigated whole-genome duplication events, refined syntenic block construction algorithm



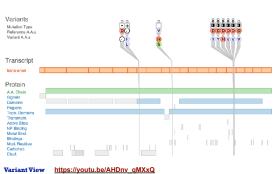
Comparative functional genomics

- · Pathline: multiple pathways, multiple genes, multiple species over time - Broad Institute, Regev Lab
- curvemap as alternative to heatmap MulteeSum: all that + spatial location (cells within fruitfly embryo)
- Harvard Med School, dePace Lab



Variant View: Visualizing sequence variants in genetic context

• concise overview supports reasoning about variant type & location - across several levels of biological context (vs extensive navigation w/ genome browsers)







Aggregated Dendrograms: Visual comparison between many phylogenetic trees

· concisely summarize trees interactively wrt bio meaningful criteria

Cerebral: Integrating gene expression w/ interaction network

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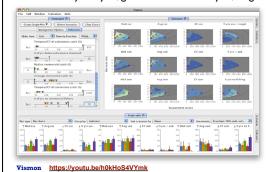
- one use case: compare gene trees to species trees





Vismon: Fisheries simulation

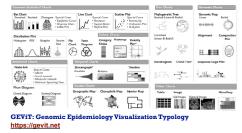
• supporting decision-makers not expert in simulation & stats - sensitivity analysis, global trade-offs analysis, staged uncertainty





Integrating visualization & biostats methods

- · Human-centered design & qualitative coding
- Epidemiology/health expectations & constraints
- · Mixed initiative: automation and manual analysis
- Mixed methods: when to use qual & when to use quant







Anamaria



₩ Algorithm

A Nested Model

for Visualization Design and Validation

http://www.cs.ubc.ca/labs/imager/tr/2009/NestedMode

A Nested Model for Visualization Design and Validation.

Munzner. IEEE Trans. Visualization and Computer Graphics (Proc. InfoVis 09), 15(6):921-928, 2009.

benchmarks can't

confirm design

lab studies can't

confirm task

Nested model: Four levels of visualization design

- · domain situation
- -who are the target users?

Aggregated Dendrograms https://youtu.be/2SLcz7KNLJw

- abstraction

 - what is shown? data abstraction
- -how is it shown?
- · visual encoding idiom: how to draw
- interaction idiom: how to manipulate
- algorithm
- efficient computation

[A Nested Model of Visualization Design and Validation Munzner. IEEE TVCG 15(6):921-928, 2009

- translate from specifics of domain to vocabulary of visualization
- why is the user looking at it? task abstraction

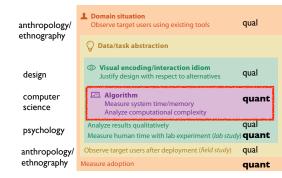
abstraction ____

[A Multi-Level Typology of Abstract Visualization Tasks Brehmer and Munzner. IEEE TVCG 19(12):2376-2385, Different threats to validity at each level



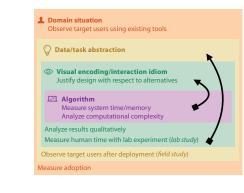
Interdisciplinary: need methods from different fields at each level

• mix of qual and quant approaches (typically)



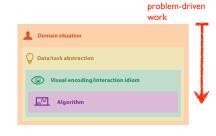


Mismatches: Common problem



Problem-driven collaborations • working with domain scientists

- translating from domain-specific language
- -how to pull this off?



Building Rewarding Collaborations



Design Study Methodology

Reflections from the Trenches and from the Stacks

Tamara Munzner

@tamaramunzner http://www.cs.ubc.ca/labs/imager/tr/2012/dsm/

Design Study Methodology: Reflections from the Trenches and from the Stacks. Sedlmair, Meyer, Munzner. IEEE Trans. Visualization and Computer Graphics 18(12): 2431-2440, 2012 (Proc. InfoVis 2012).



Lessons learned from the trenches: 21 between us

















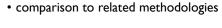






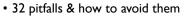


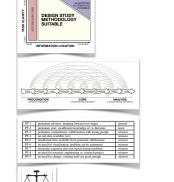






• 9-stage framework



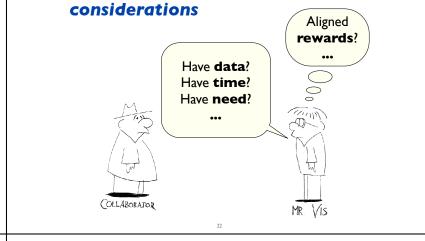


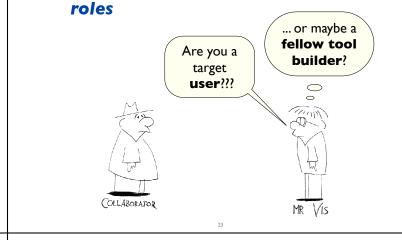
Design study methodology: 32 pitfalls

• and how to avoid them

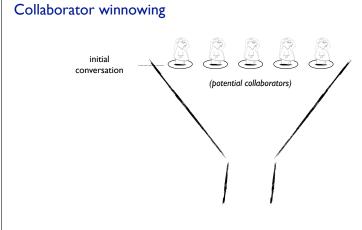
PF-1	premature advance: jumping forward over stages	general
PF-2	premature start: insufficient knowledge of vis literature	learn
PF-3	premature commitment: collaboration with wrong people	winnow
PF-4	no real data available (yet)	winnow
PF-5	insufficient time available from potential collaborators	winnow
PF-6	no need for visualization: problem can be automated	winnow
PF-7	researcher expertise does not match domain problem	winnow
PF-8	no need for research: engineering vs. research project	winnow
PF-9	no need for change: existing tools are good enough	winnow

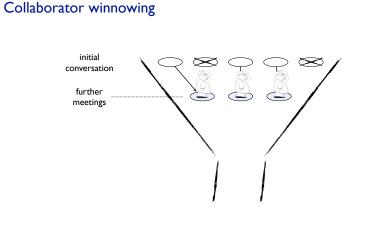


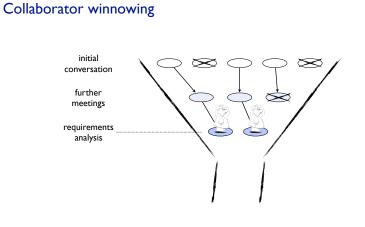


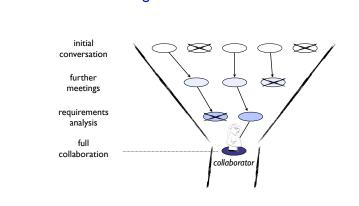














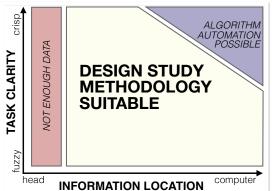
Talk with many, stay with few!

Design study methodology: 32 pitfalls

• and how to avoid them

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More Information

https://www.cs.ubc.ca/~tmm/talks.html#hakai19-methods

 papers, videos, software, talks, courses http://www.cs.ubc.ca/group/infovis http://www.cs.ubc.ca/~tmm



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