

StratomeX

Visual Analysis of Large-Scale Heterogeneous Genomics Data for Cancer Subtype Characterization

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Cancer Subtypes

Cancer Subtypes have

- different histology

- different molecular alterations

Subtypes have serious implications

- different treatment for subtypes

- prognosis varies between subtypes

Cancer Subtype Analysis

Modern cancer subtype analysis based on
biomolecular data

Our Goal:

Support cancer subtype characterization through
integrative visual analysis of multiple relevant datasets.

Gene expression patterns of breast carcinomas distinguish tumor subclasses with **clinical implications**

Theresa Sorlie et al.

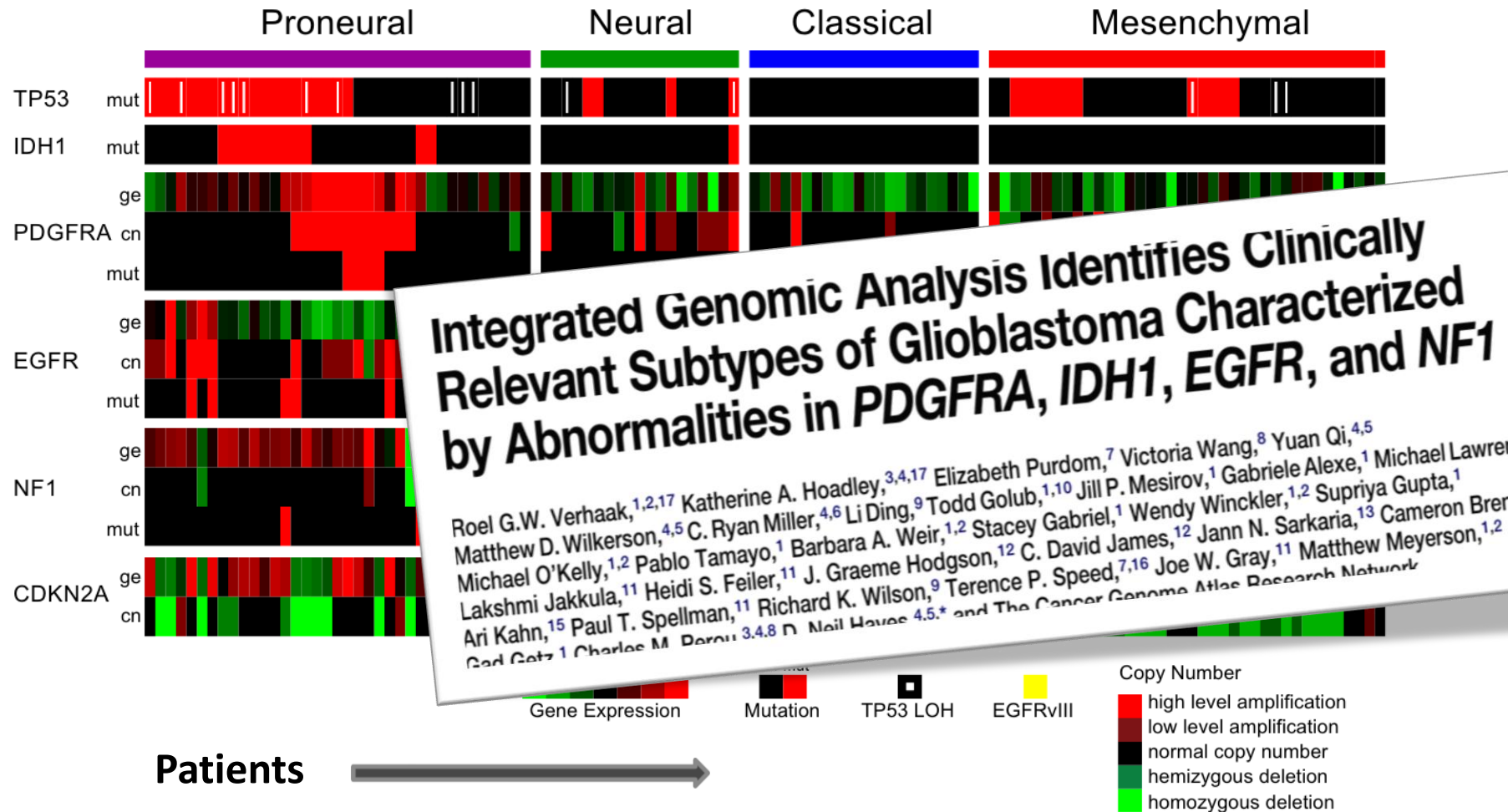
Molecular subclasses of high-grade glioma **predict prognosis**, delineate a pattern of disease progression, and resemble stages in neurogenesis

Gene expression profiling identifies **clinically relevant** subtypes of prostate cancer

Integrated Genomic Analysis Identifies Clinically Relevant Subtypes of Glioblastoma Characterized by Abnormalities in *PDGFRA*, *IDH1*, *EGFR*, and *NF1*

Roel G.W. Verhaak,^{1,2,17} Katherine A. Hoadley,^{3,4,17} Elizabeth Purdom,⁷ Victoria Wang,⁸ Yuan Qi,^{4,5} Matthew D. Wilkerson,^{4,5} C. Ryan Miller,^{4,6} Li Ding,⁹ Todd Golub,^{1,10} Jill P. Mesirov,¹ Gabriele Alexe,¹ Michael Lawrence,^{1,2} Michael O'Kelly,^{1,2} Pablo Tamayo,¹ Barbara A. Weir,^{1,2} Stacey Gabriel,¹ Wendy Winckler,^{1,2} Supriya Gupta,¹ Lakshmi Jakkula,¹¹ Heidi S. Feiler,¹¹ J. Graeme Hodgson,¹² C. David James,¹² Jann N. Sarkaria,¹³ Cameron Brennan,¹⁴ Ari Kahn,¹⁵ Paul T. Spellman,¹¹ Richard K. Wilson,⁹ Terence P. Speed,^{7,16} Joe W. Gray,¹¹ Matthew Meyerson,^{1,2} Gad Getz,¹ Charles M. Perou,^{3,4,8} D. Neil Hayes,^{4,5,*} and The Cancer Genome Atlas Research Network

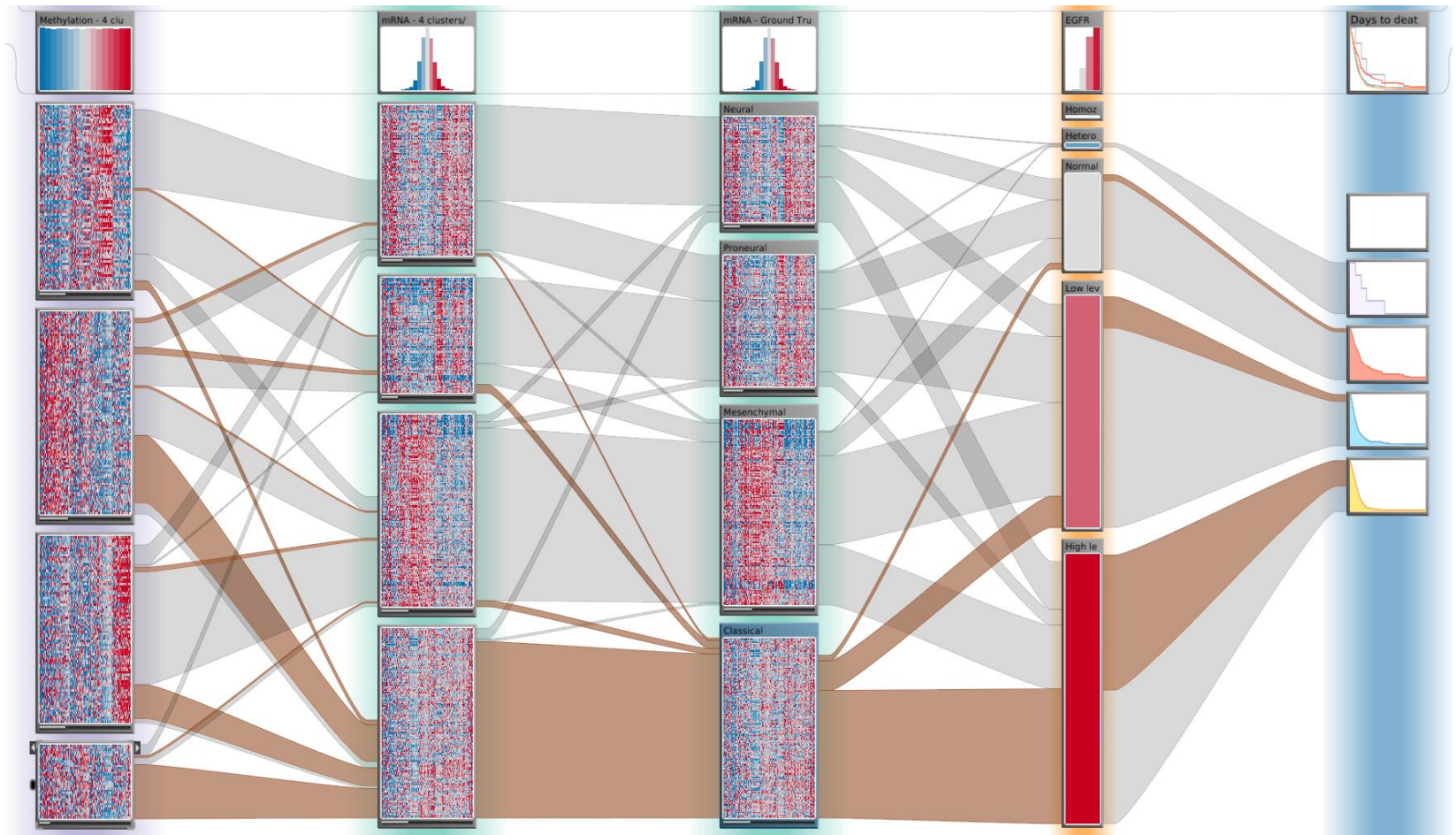
How Subtypes are Visualized



Challenges

Challenge 1: Visualize complex interdependencies between multiple datasets

StratomeX: Interdependencies between Stratifications of Datasets

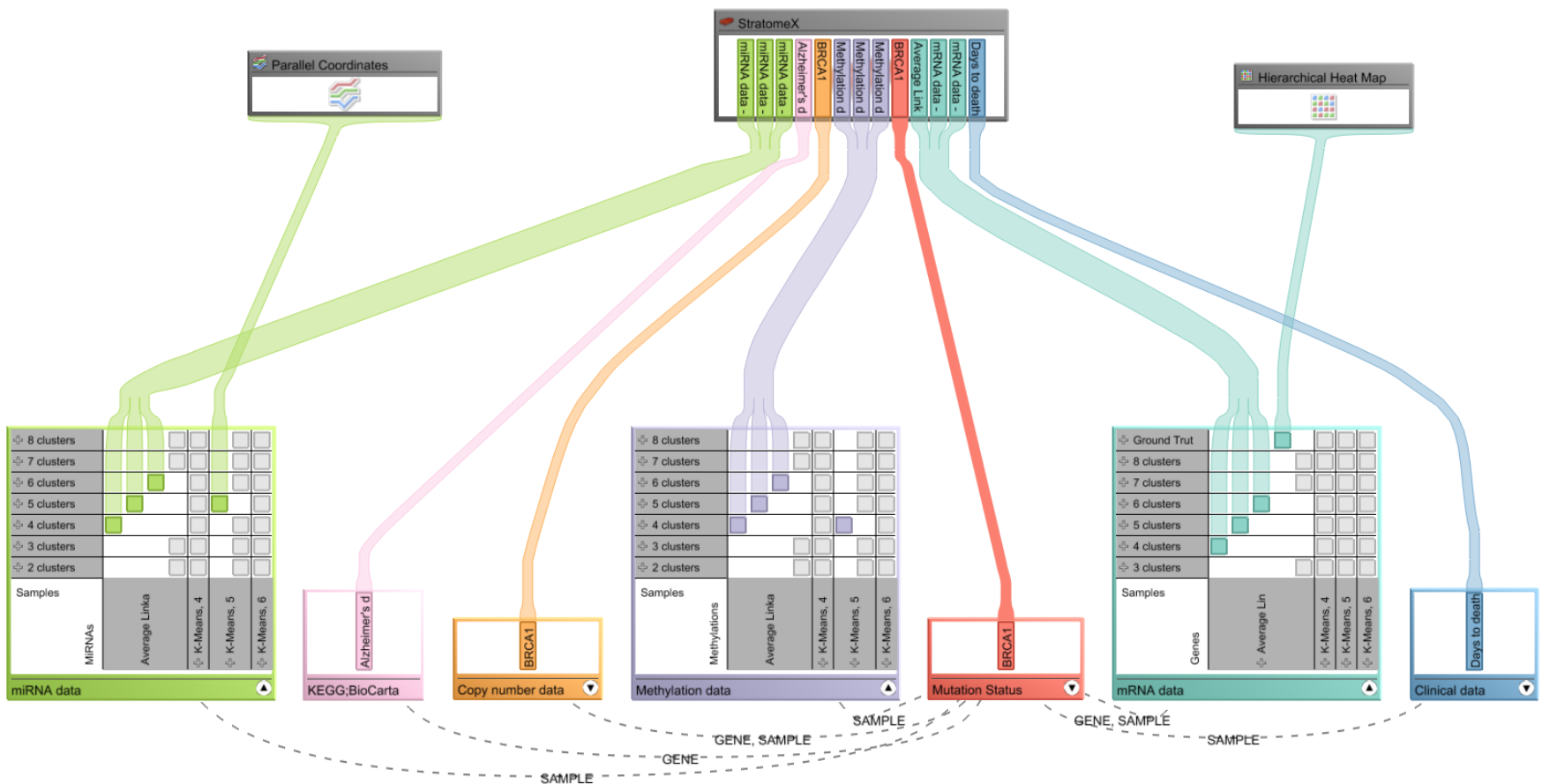


Challenges

Challenge 1: Visualize complex interdependencies between multiple datasets

Challenge 2: Manage complex setup of multiple datasets, multiple stratifications and multiple views

Data View Integrator



THE DATA

Large-scale project to catalogue genetic mutations responsible for cancer

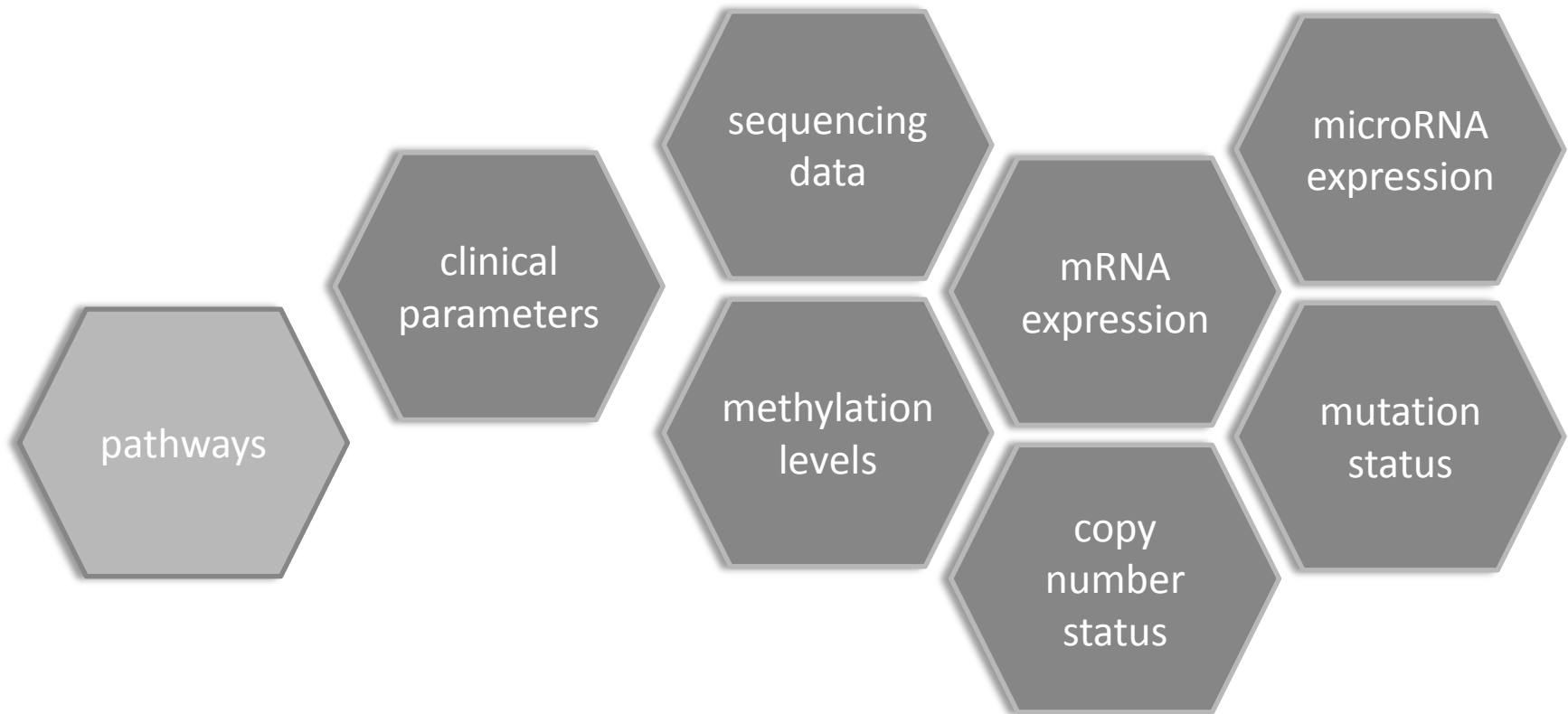
20 tumor types

500 patient samples each

Extensive molecular profiling for each patient



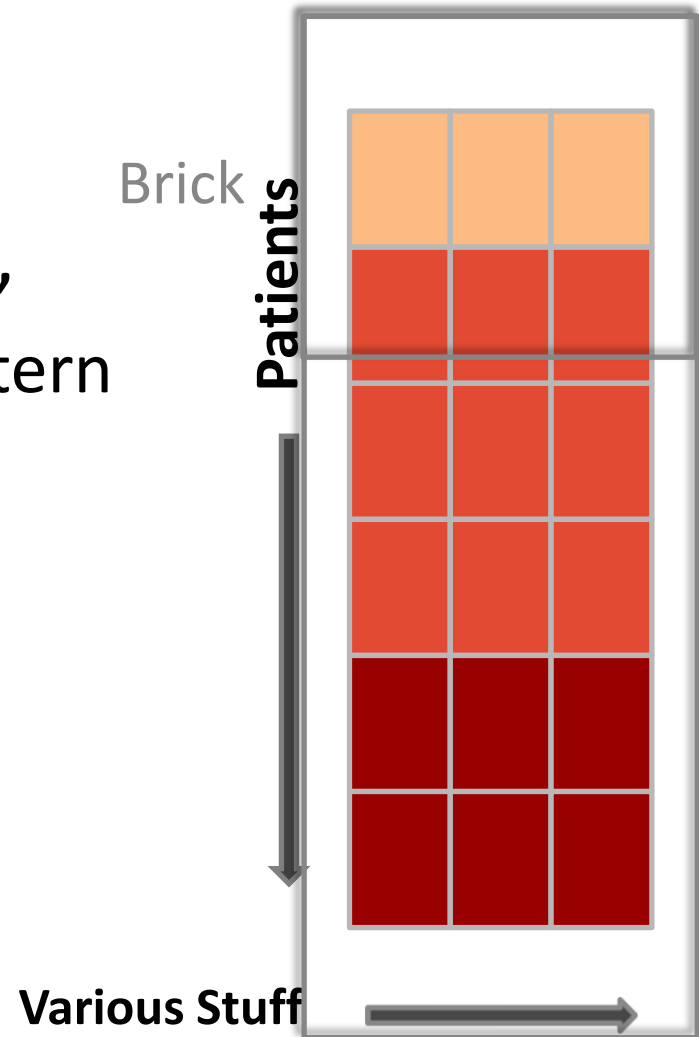
TCGA Data



THE TECHNIQUE

Tabular Data

Subtypes are identified
by stratifying datasets, e.g.,
based on an expression pattern
a mutation status
a copy number alteration
a combination of these



Tasks

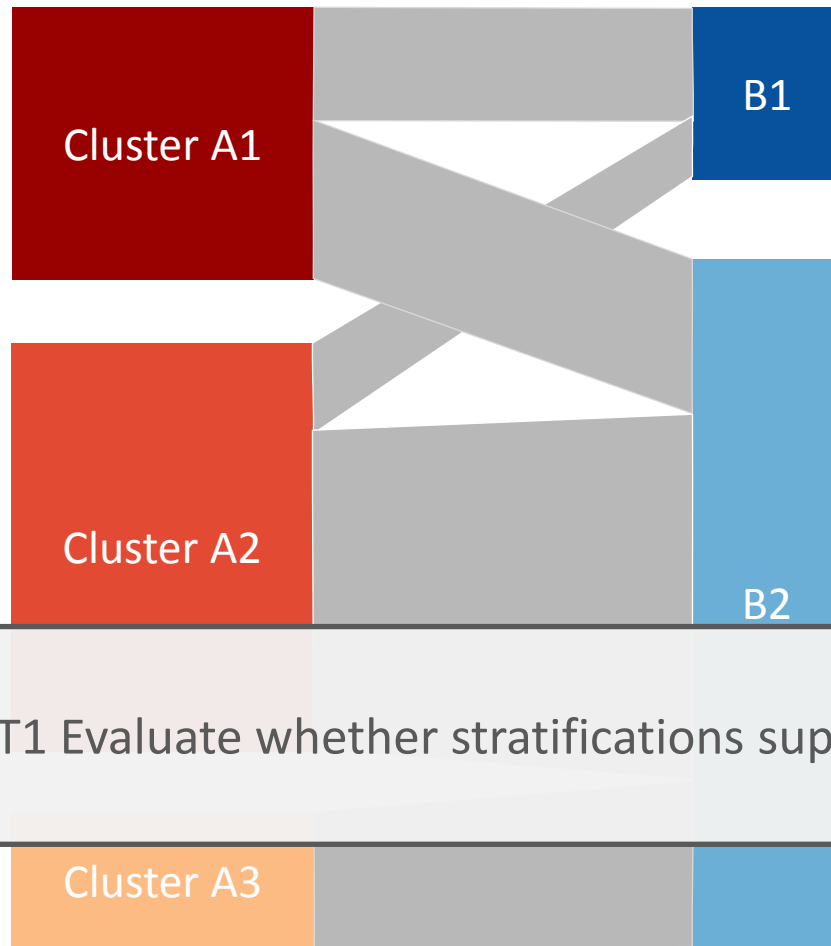
- T1 Evaluate whether stratifications **support each** other
- T2 **Refine** stratifications
- T3 Review **effect** of stratifications
 - on clinical outcomes
 - on pathways
- T4 Show expression **patterns** in subtypes

Elicited in expert interviews and literature review

Stratification of a Single Dataset



Stratification of Multiple Datasets

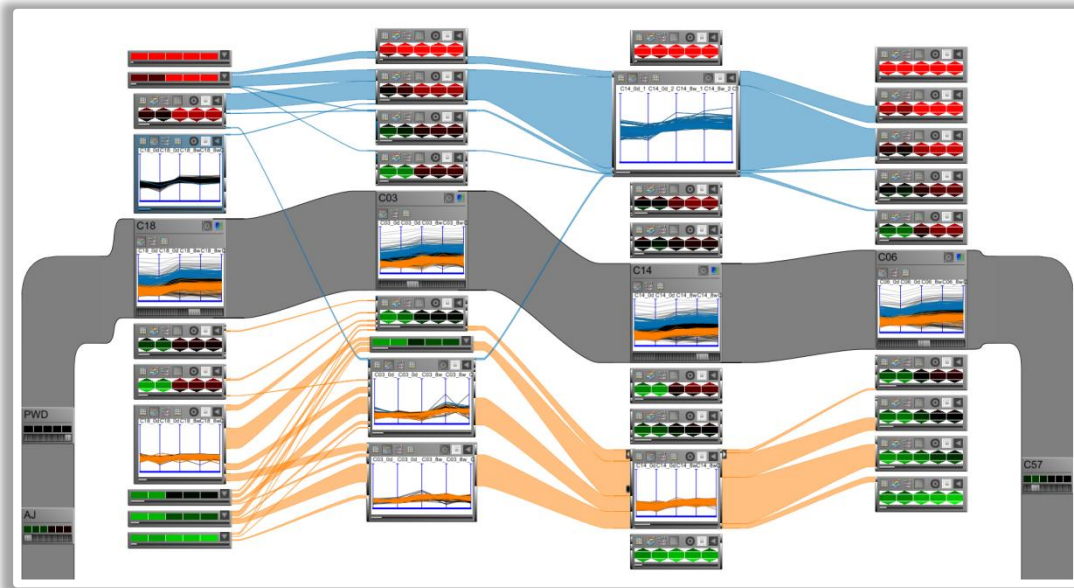


T1 Evaluate whether stratifications support each other

Tabular Data

Categorical Data

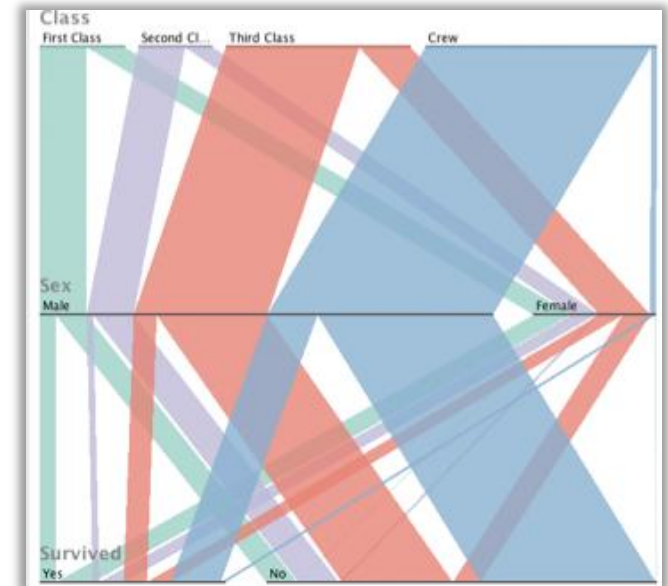
VisBricks



[Lex, InfoVis 2011]

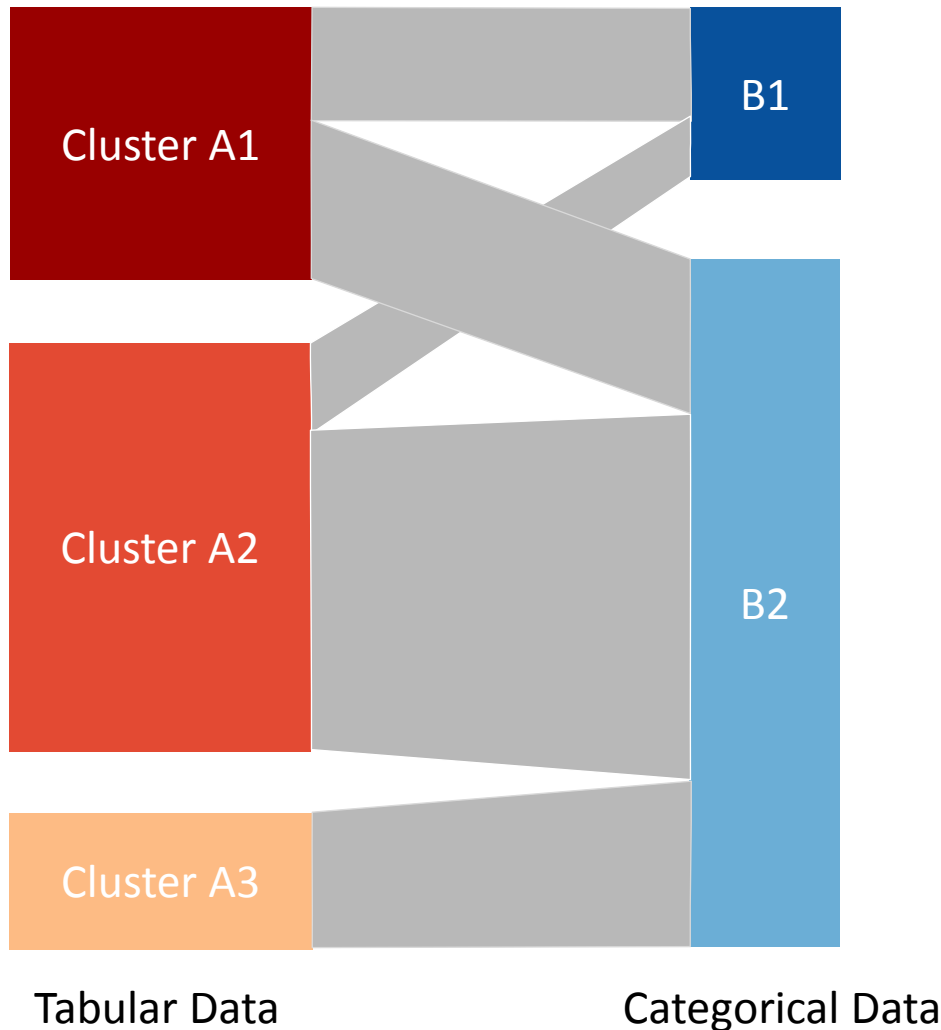
Alexander Lex, Graz University of Technology

Parallel Sets

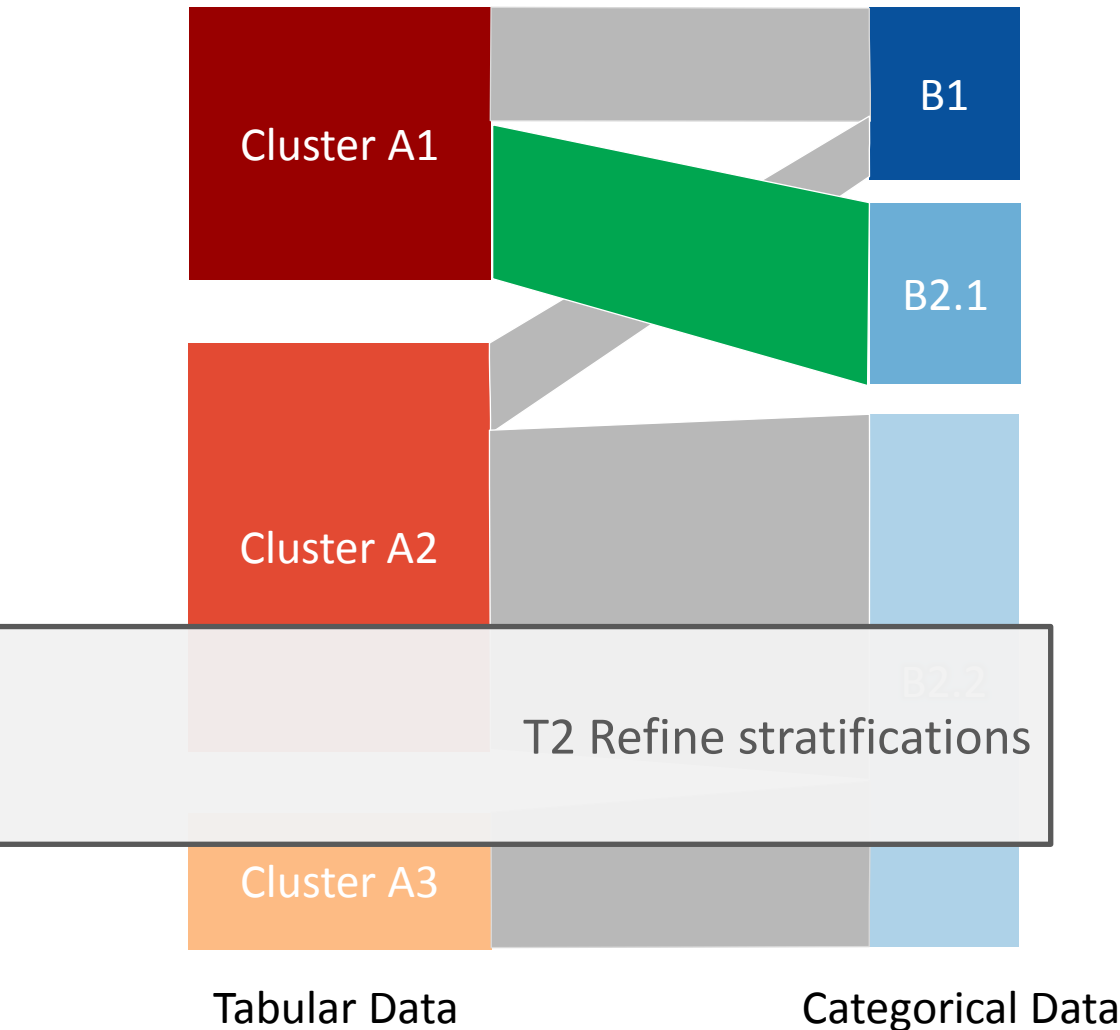


[Kosara, InfoVis 2006]

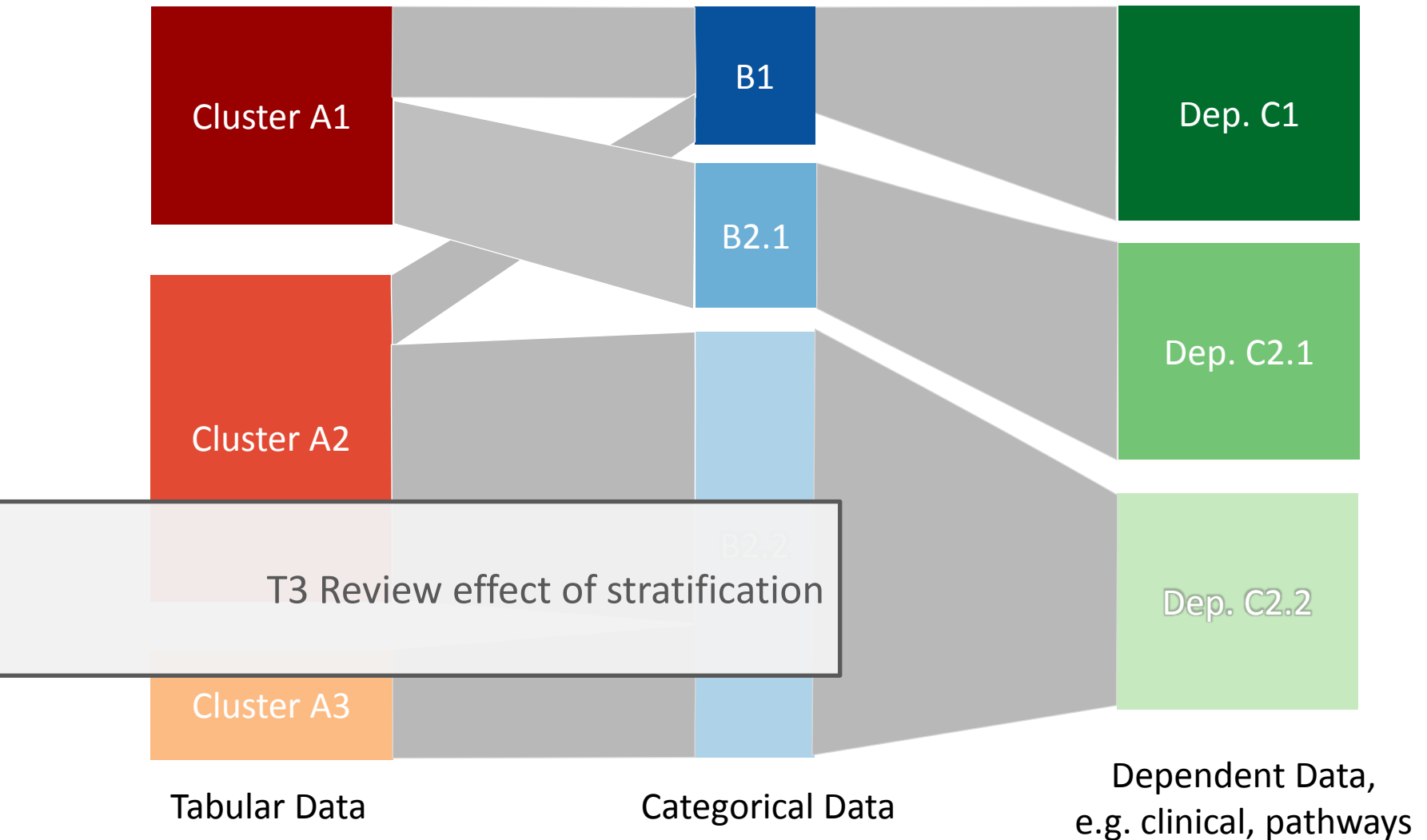
Stratification of Multiple Datasets



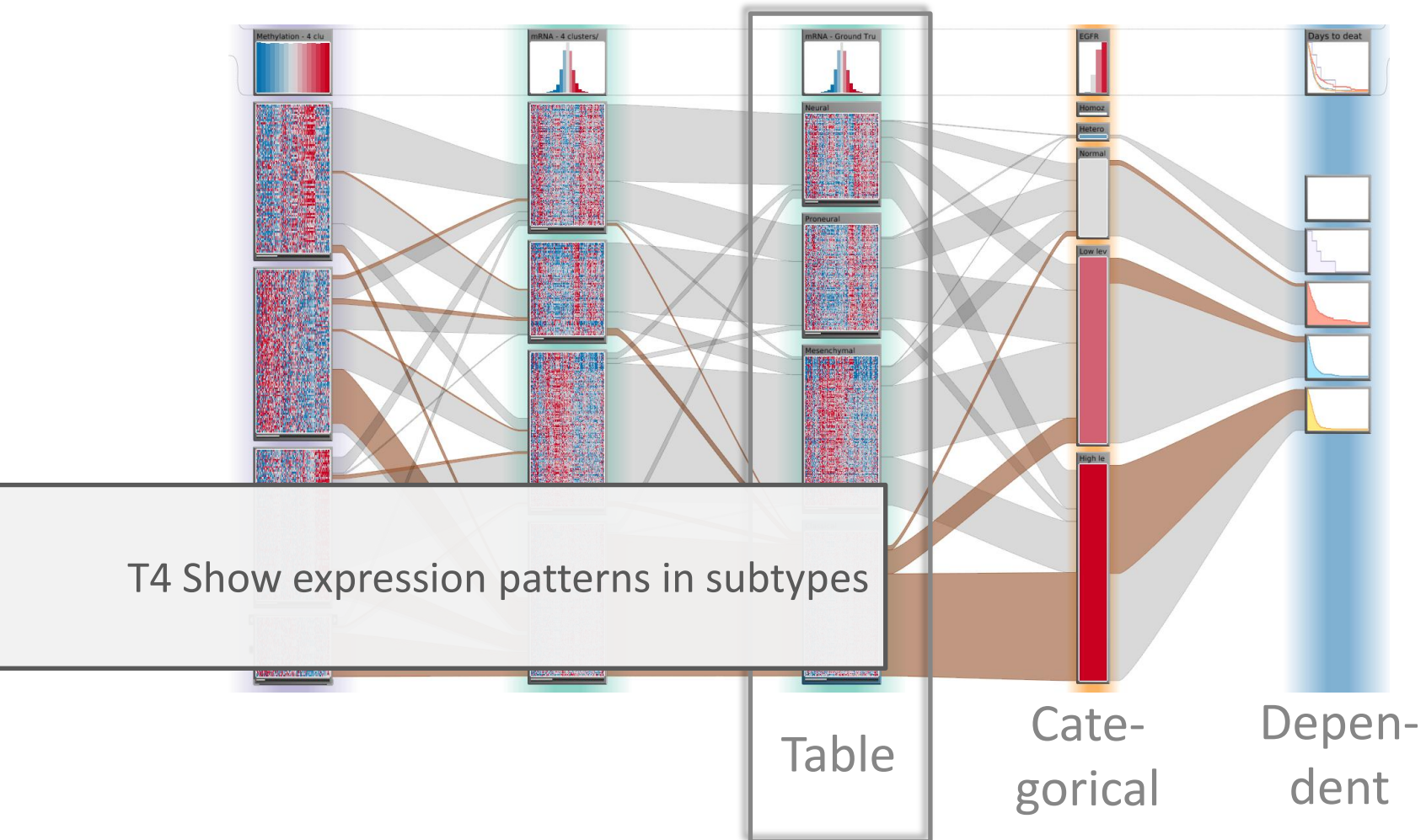
Stratification of Multiple Datasets



Stratification of Multiple Datasets



Column Classes



Live Demo

Glioblastoma Multiforme

Case Studies

Conducted with two domain experts
from Broad Institute of MIT and Harvard

Datasets

Glioblastoma Multiforme

Breast Invasive Carcinoma

Report on findings

Methylation subtypes

Effects of clustering

Effects on Pathways



Implementation

Part of Caleydo – <http://caleydo.org>

Caleydo is now **open source!**

Release 2.0 - 3 weeks ago

Includes Glioblastoma dataset



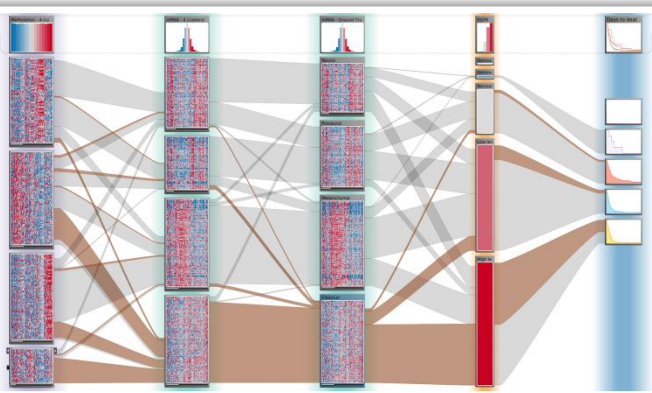


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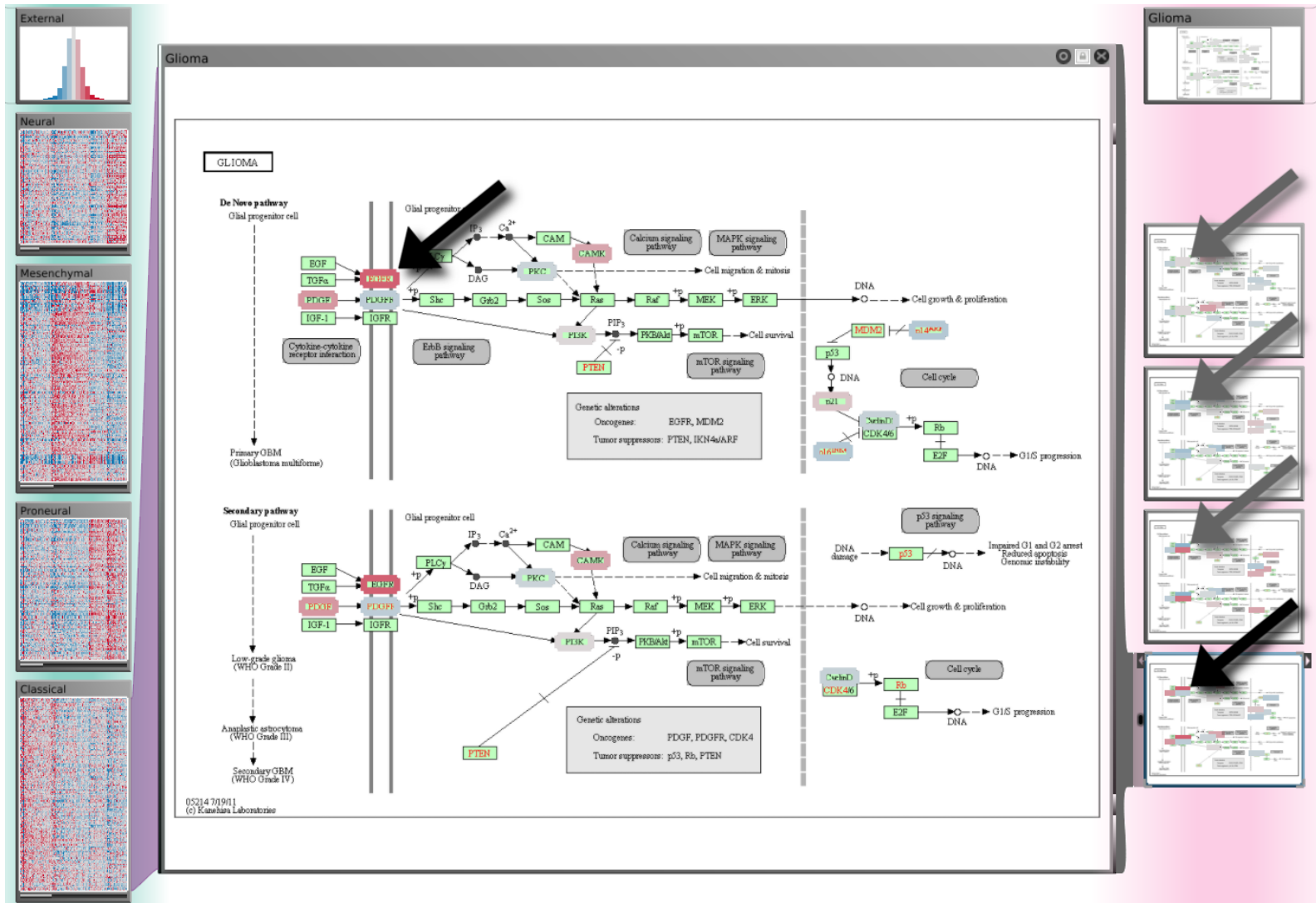
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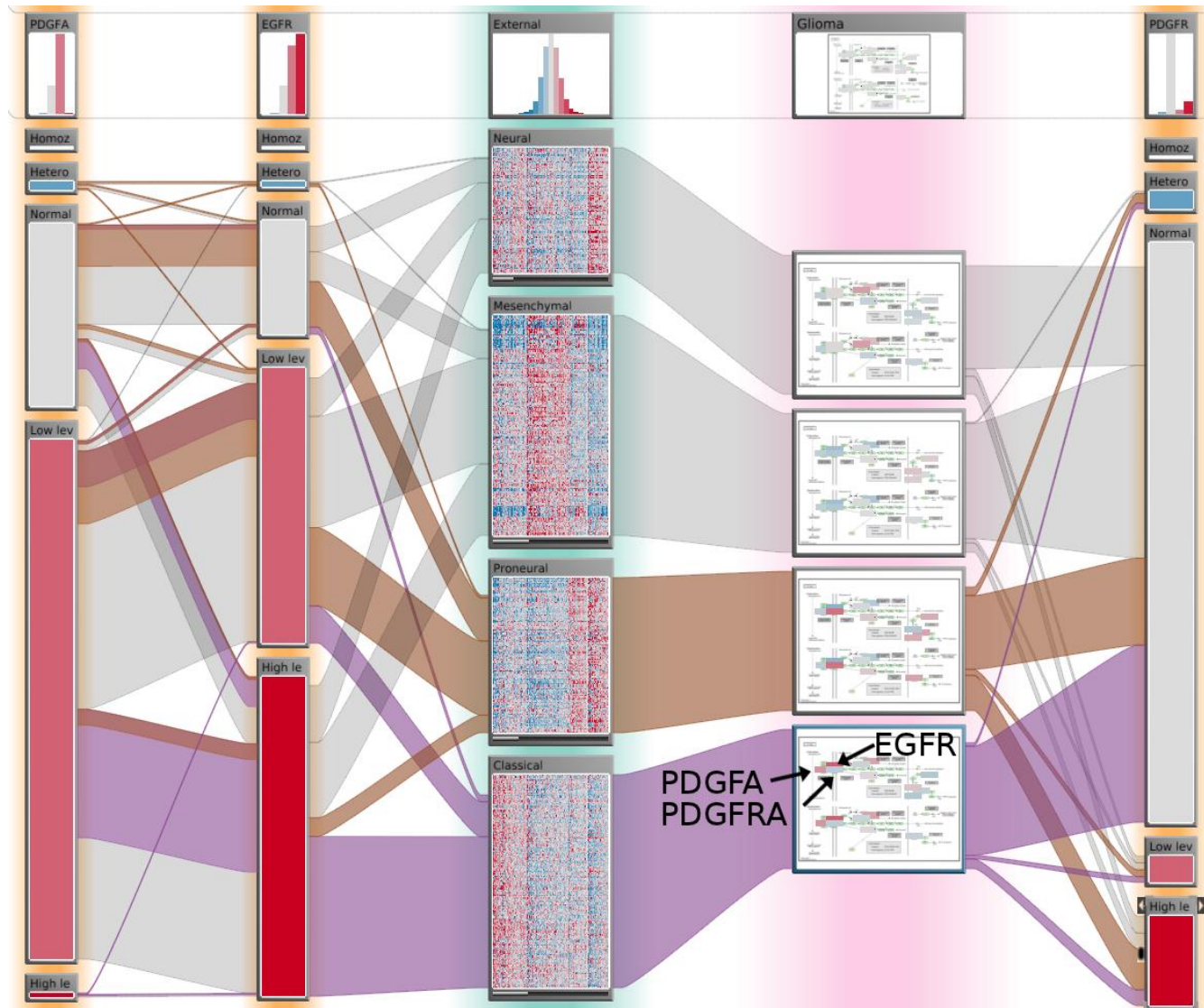
Case Studies

Glioblastoma Multiforme

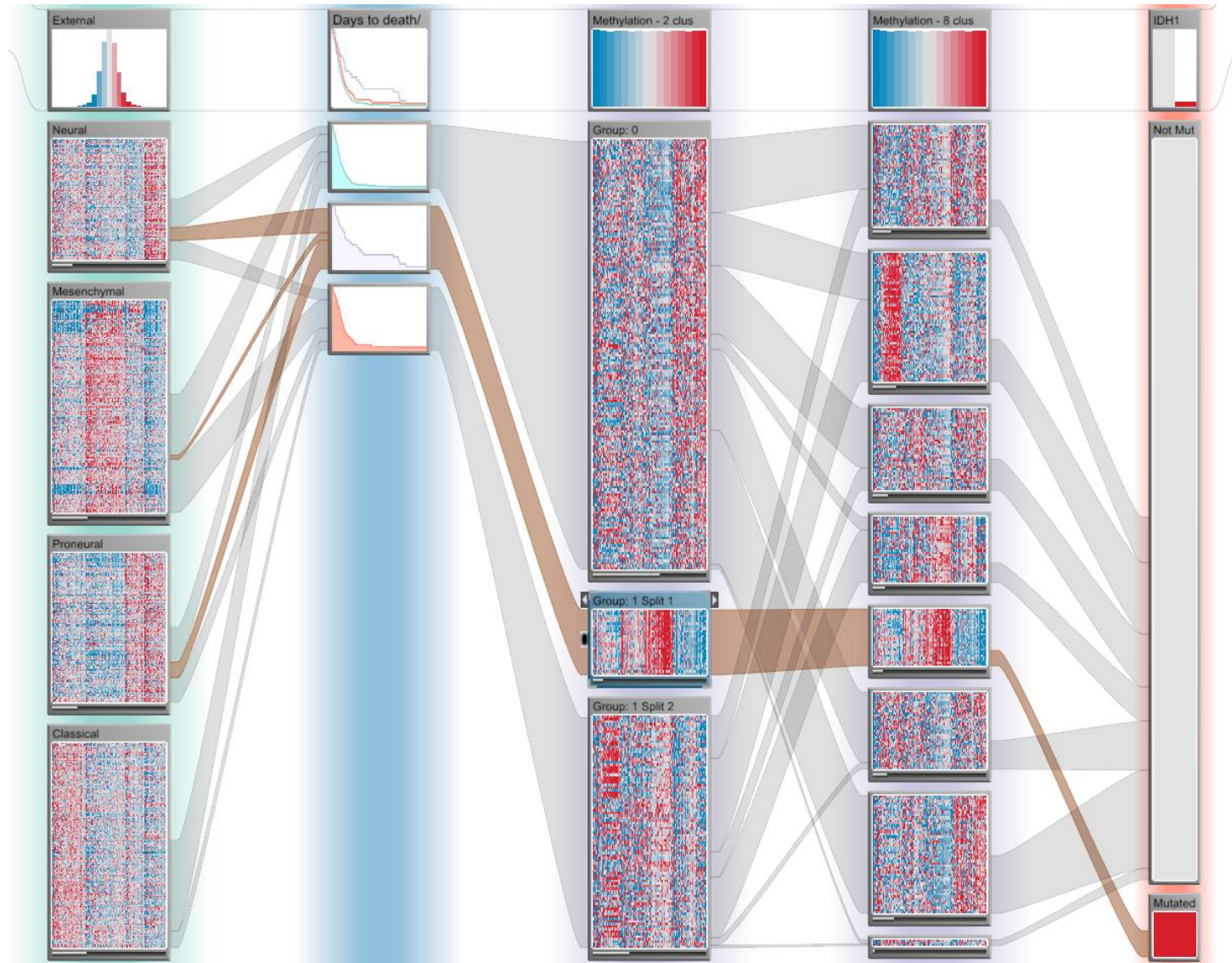
Effects on Pathways



Effects on Pathways



Methylation Subtypes



Clustering Effects

