Visual Analytics for Plant Pangenomes

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About me

Dept. Of Mathematics & Computer Science

Visualization Cluster



PhD Candidate @ Eindhoven University of Technology

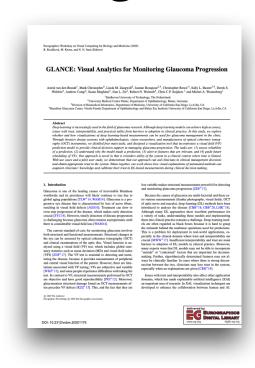
Msc Data Science and Entrepreneurship

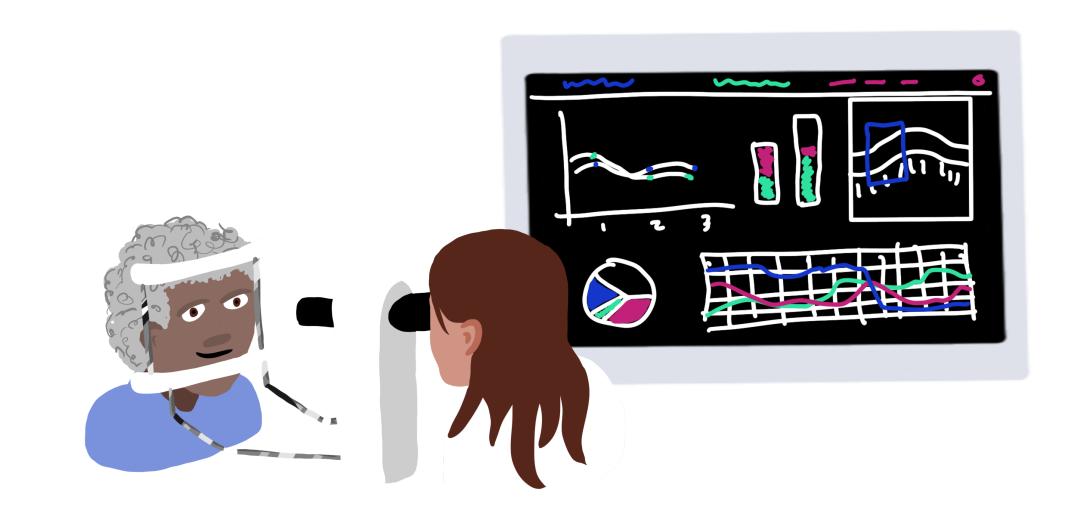
Bsc Industrial Design

GLANCE: Visual Analytics for Monitoring Glaucoma Progression



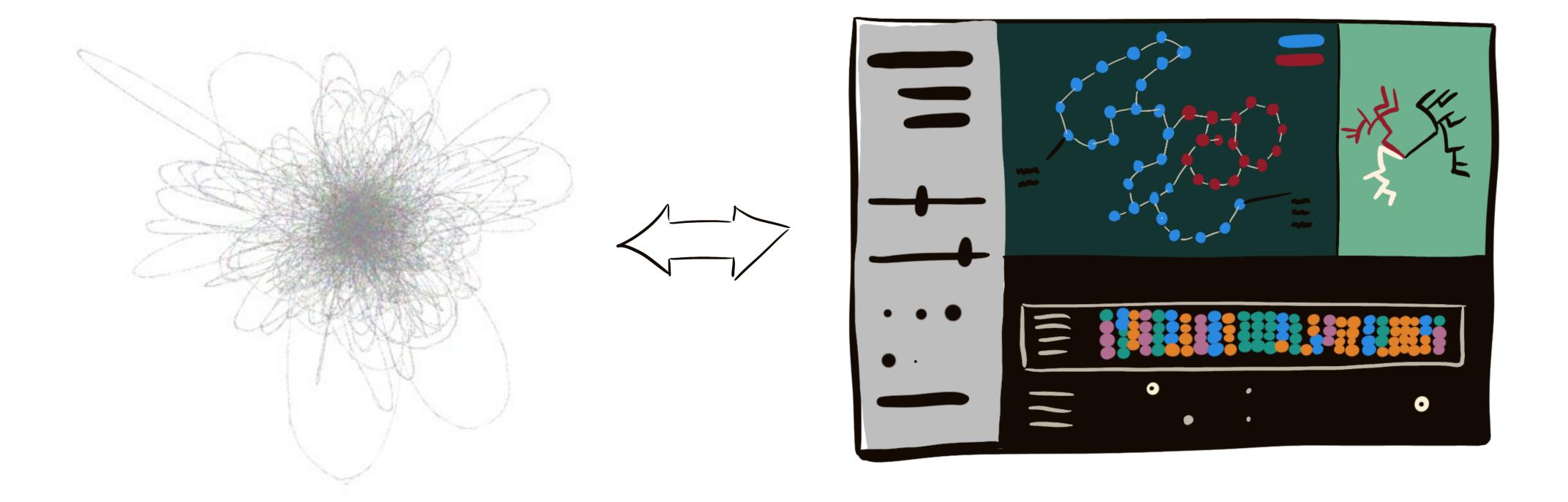
Design Study





VA for Plant Pangenomes

Translating pan genome information into meaningful views for genome scientists



What do we need?



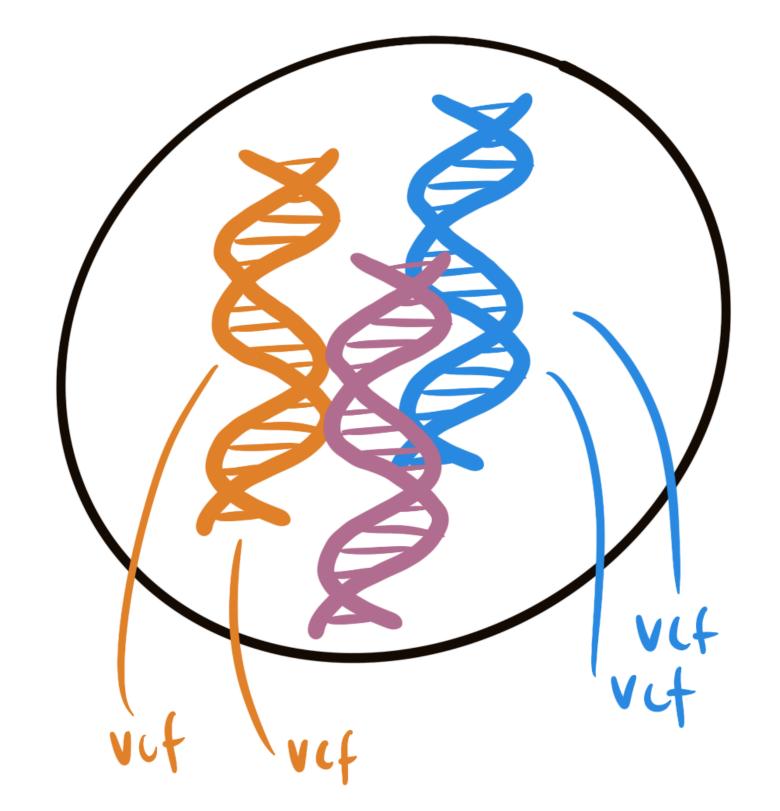
Use Cases: genomic diversity

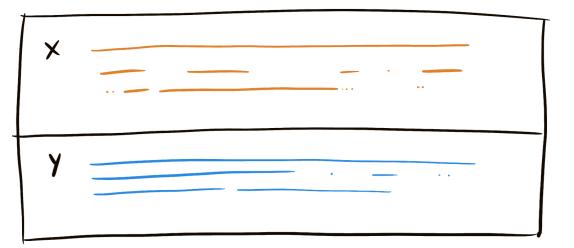
1. Sequence Variation: allelic variation, genetic variation

Small-scale variation

Pangenome

- Genomic regions
- VCF files

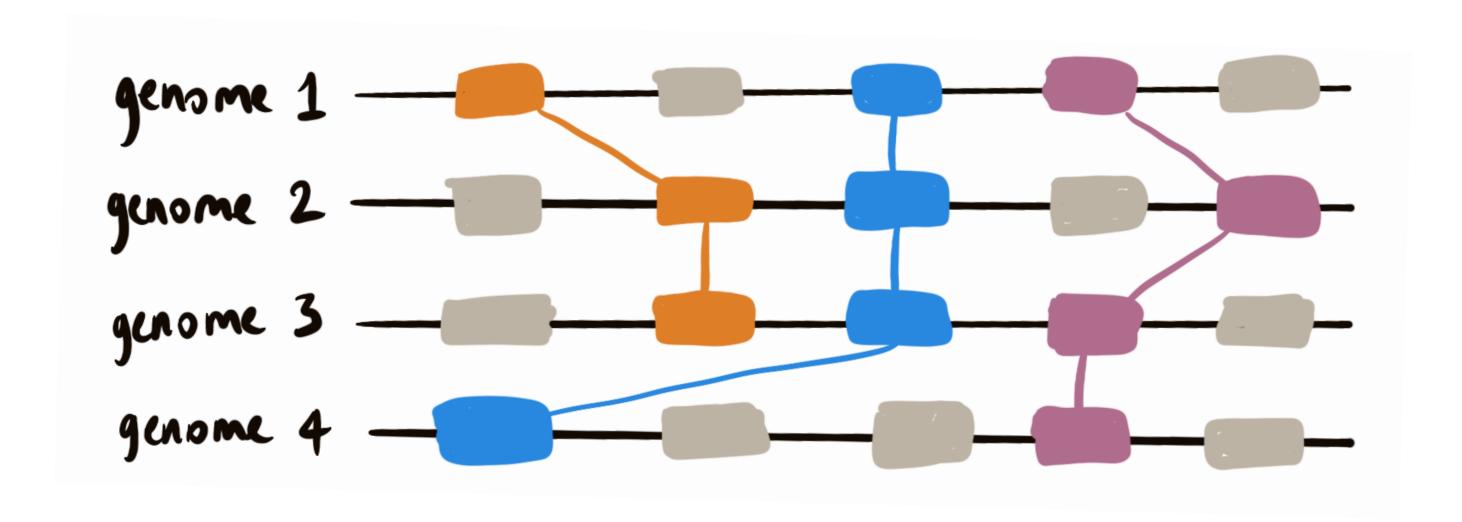




Use Cases: genomic diversity

2. Structural variation: collinearity/synteny, gene order

Large-scale variation



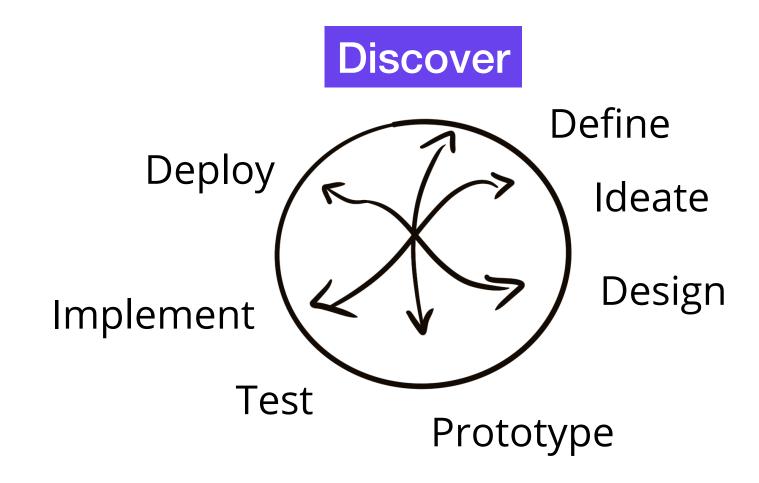
Visualization challenges

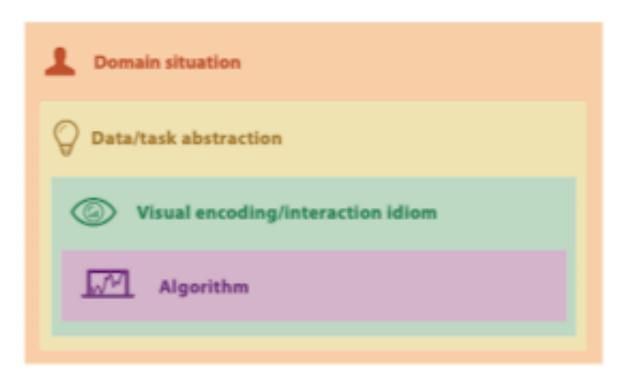
- Complex data structure (large and heterogenous)
- Intuitive exploration between different levels of organisation

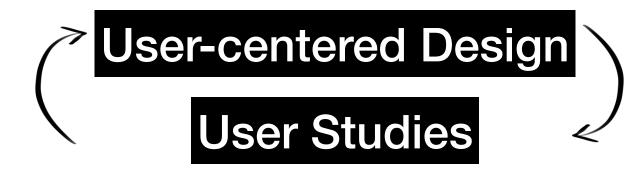
Beyond pairwise comparisons

Approach

Visualization Design Process







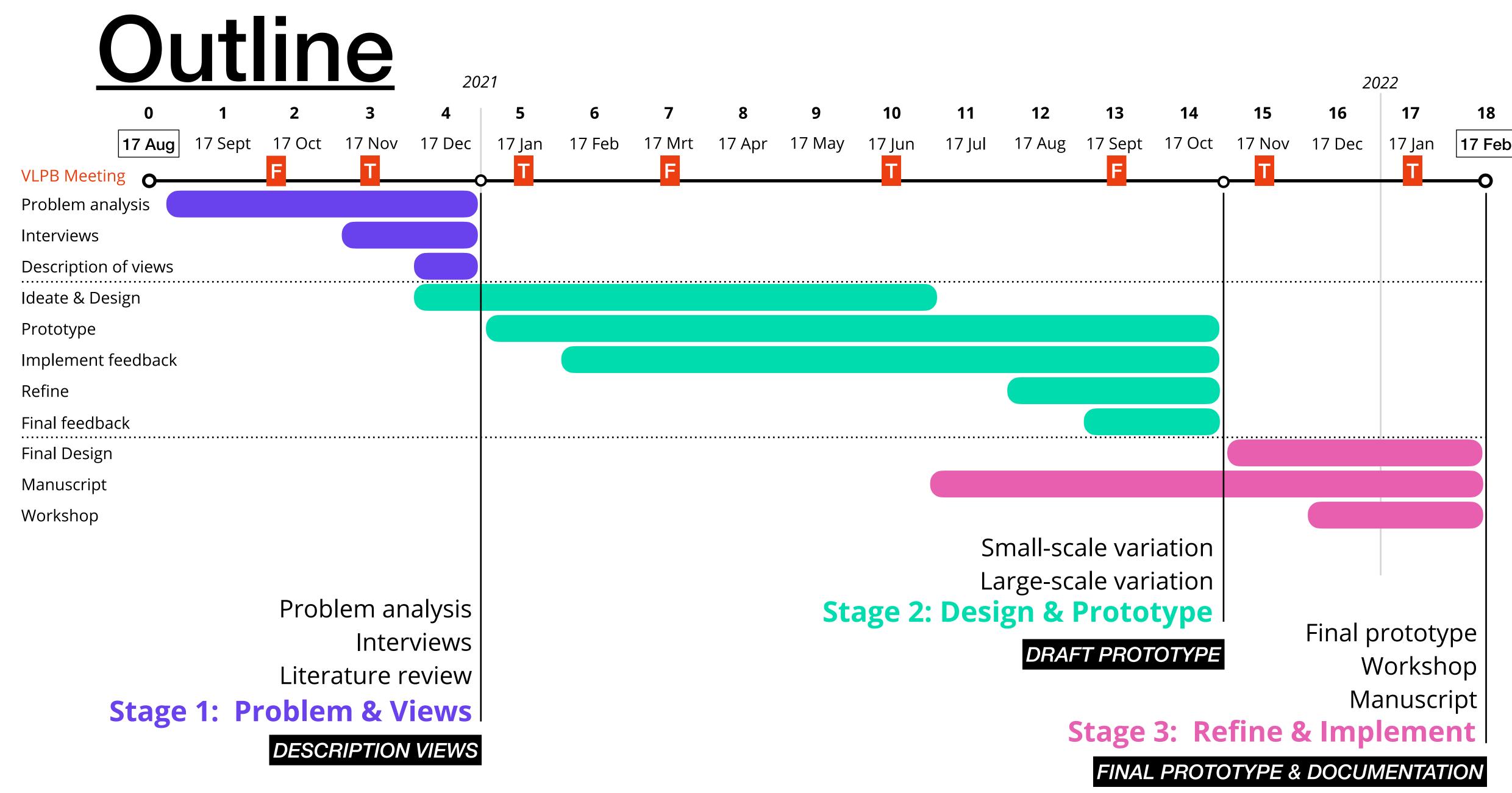
Munzner (2014). Four nested levels of vis design.

Task Analysis

Domain situation

Questions & Needs — > Qualitative assessment





Communication

Communicating progress VLPB meetings, telecons, Shortcase

Interviews & visits Face-to-face, telecons

T1: 17/11/'20 between 9:00 and 12:00 (?)

T2: 19/01/'21 between 9:00 and 12:00 (?)

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