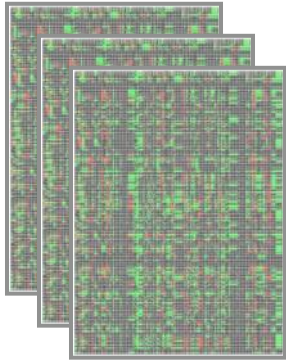


Genome Model Construction

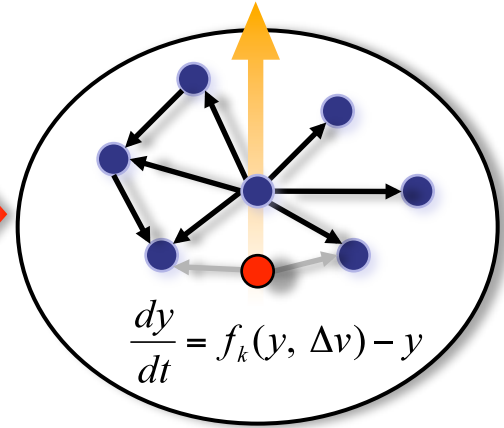
“-Omics” data



- Transcriptomics (microarrays)
- Metabolomics (metabolic profiles)
- Phenomics (agronomical properties)

Reverse Engineering

Agronomic properties



$$\frac{dy}{dt} = f_k(y, \Delta v) - y$$

Effective regulatory model

Genome Redesign

Compute predicted agronomical properties

Solve gene expression and metabolic profile in steady-state

Suggested mutation

Multiple-objective optimization

Loop N-times

Experimental Validation

