

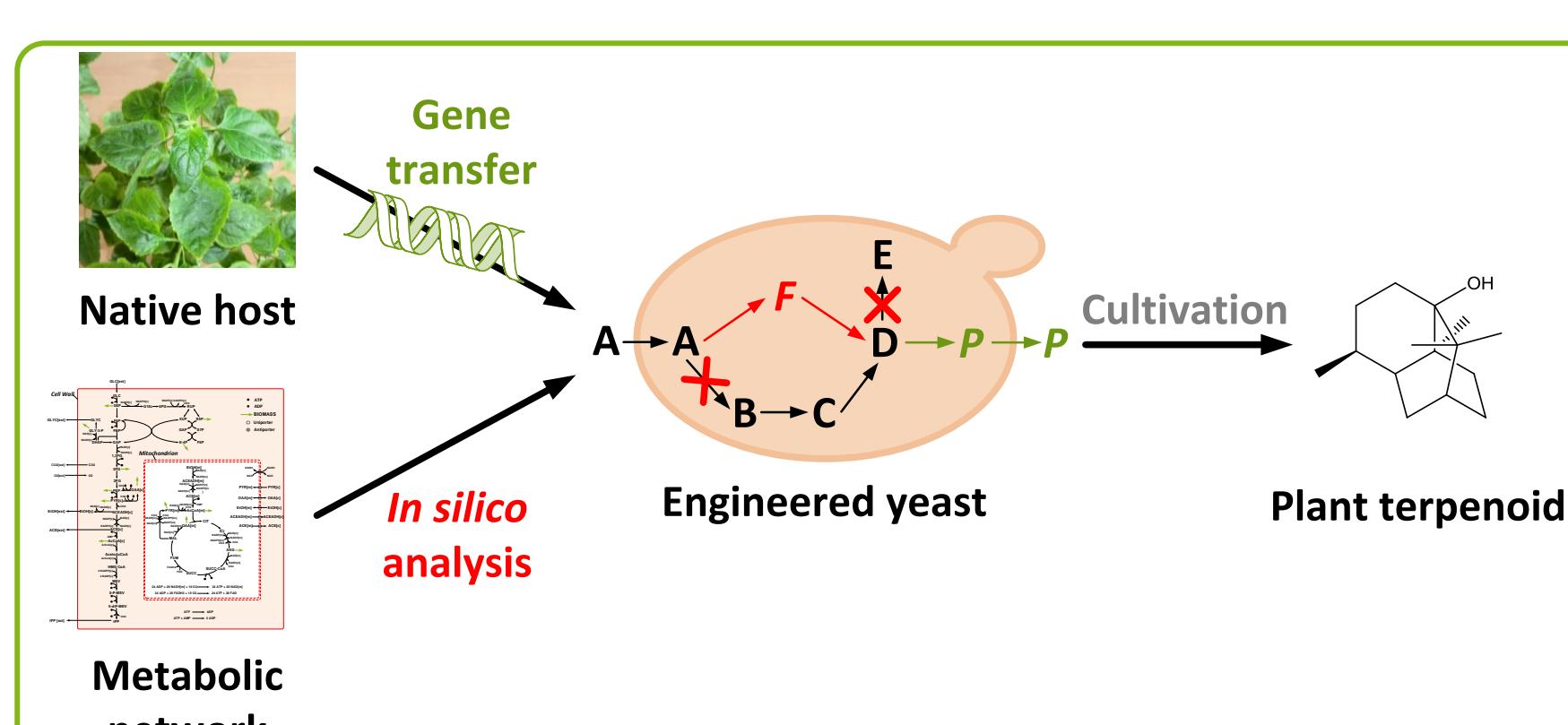
# Towards a platform organism for terpenoid production – *in silico* analysis of *Saccharomyces cerevisiae* as potential host

Evamaria Gruchattka, Verena Schütz, Oliver Kayser

Technische Universität Dortmund, Technische Biochemie  
Evamaria.Gruchattka@bci.tu-dortmund.de; Oliver.Kayser@bci.tu-dortmund.de

## Terpenoids

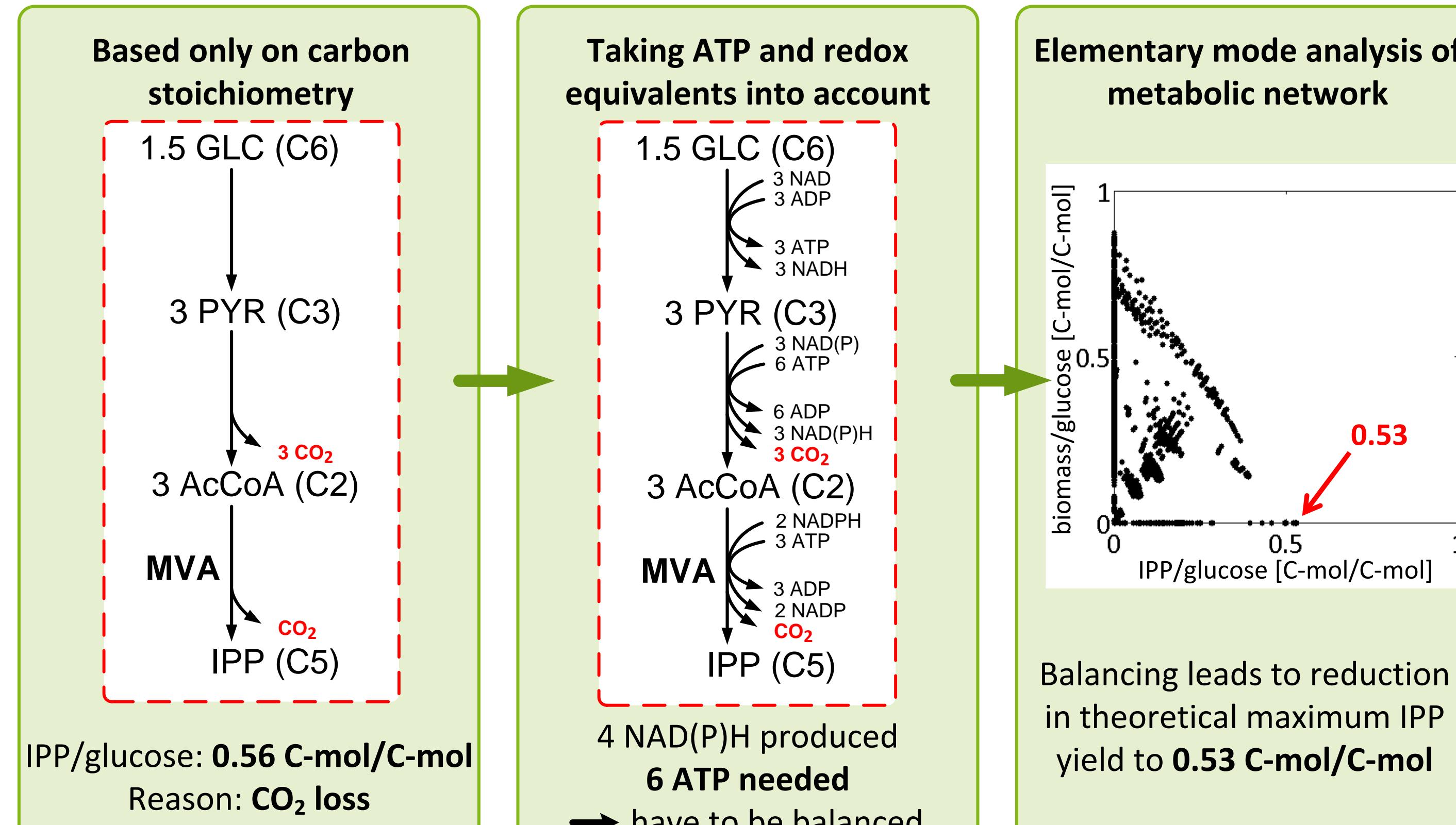
- One of the largest classes of natural products
- Possess important medicinal and industrial properties
- Some are rare and produced in low amounts in plants
- Heterologous microbial production may help to overcome supply problems and high purification costs
- Necessity of optimization of yield and productivity in yeast e.g. via metabolic engineering [1]



## Objectives

- Development of a platform organism for the efficient supply of isopentenyl diphosphate (IPP), the biosynthetic precursor of all terpenoids using *in silico* modelling
- S. cerevisiae* is analyzed by means of elementary mode analysis [2] regarding the metabolic potential to supply IPP and to identify overexpression candidates
- Knockout-strategies for an enhanced terpenoid yield are identified using constrained minimal cut sets [3]

## Analysis of Terpenoid Pathway and Central Metabolic Network: Identification of Limitations and Solutions



## Constrained Minimal Cut Sets (cMCS): Enhanced Terpenoid Production

C-source	Theoret. max.	Experimental *
Glucose	0.53	0.003-0.01 [9-11]
Galactose	0.53	0.004 [12]
Fructose	0.53	-
Xylose	0.53	-
Glycerol	0.56	-
Ethanol	0.68	0.19 [11]

\* yeast strains optimized within terpenoid pathway

Experimental IPP yields are very low

→ Change flux distribution to force the cell to produce high yields

### Constrained minimal cut sets: theory

- Minimal set of structural interventions (gene knockouts)
- Repressing a certain functionality (deletion task: low product yield)
- Preserving a certain functionality (desired modes: high product yield)



### cMCS for enhanced terpenoid yield on glucose

#### Feasible set of interventions

- Prevention of acetate secretion
- Prevention of ethanol secretion or production (alcohol dehydrogenases)
- Partial disruption of citric acid cycle, e.g.:
  - mitochondrial  $\alpha$ -ketoglutarate dehydrogenase
  - mitochondrial succinyl-CoA ligase

#### Consequence for flux distribution

- Carbon cannot be oxidized completely to  $\text{CO}_2$
- Carbon flux is redirected towards AcCoA

#### Predicted results

Feature	Wild type	cMCS knockout mutant
Elementary modes	9,844	19-142
Minimal IPP carbon yield	0	0.27-0.33
Maximal IPP carbon yield	0.53	0.53
Maximal biomass yield	0.87	0.32-0.4

[13]

## Summary/Conclusion

- Heterologous enzymes/pathways with decreased energy consumption or less  $\text{CO}_2$  loss are identified leading to a higher theoretical maximal IPP yield
- Knockout strategies are identified for a coupling of biomass production to a minimal IPP yield which is higher than published yields to date

## Future Prospects

- In vivo* validation of predicted results using patchoulool as a reporter (a sesquiterpenoid and fragrance compound used in perfume industry)

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