

SYSTEMS BIOLOGY IN BIOTECH

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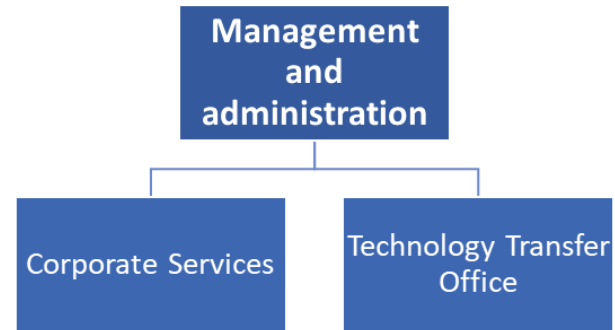
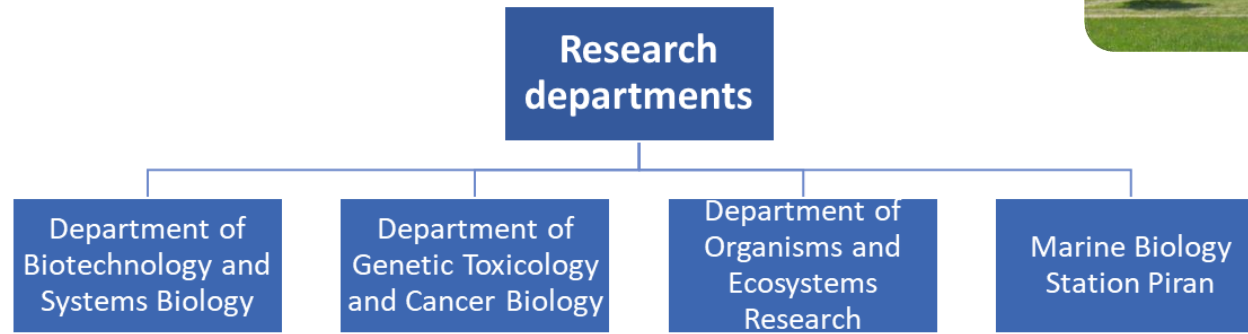
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NACIONALNI INŠTITUT ZA **BIOLOGIJO**
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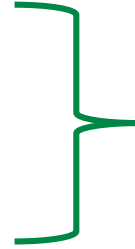
National Institute of Biology



Department of Biotechnology and Systems Biology

Fields of research and applications:

- Microbiology
- GMO
- Systems biology
- Bacteriology and metrology



Basic and applicative research
Contract research for industry

Advanced molecular biology techniques

Quantitative real-time PCR, digital PCR, LAMP, NGS, barcoding...

Statistical modeling and data integration

Workshops and trainings

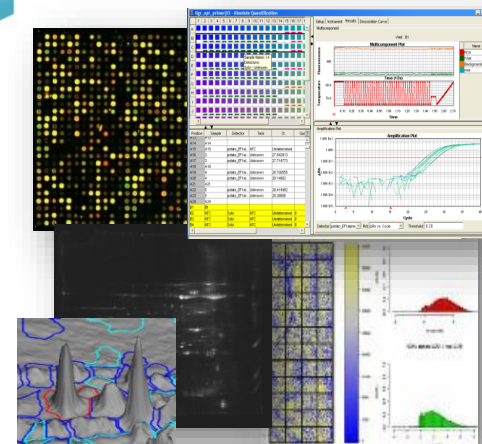
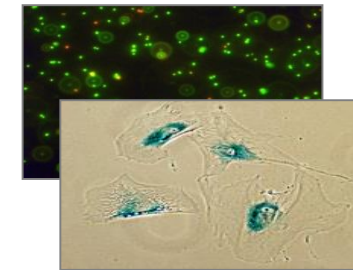
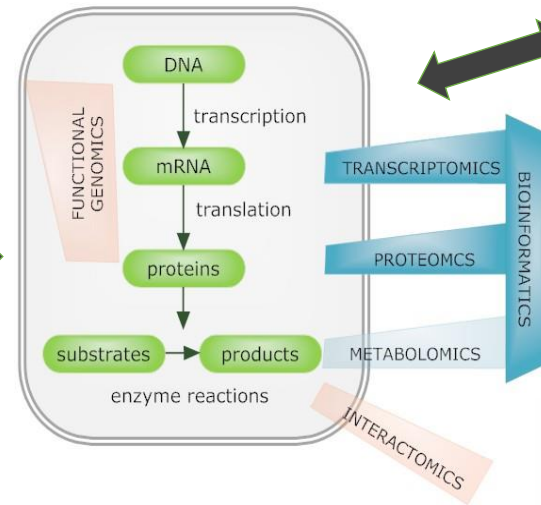
Organization of proficiency tests

Omics approaches

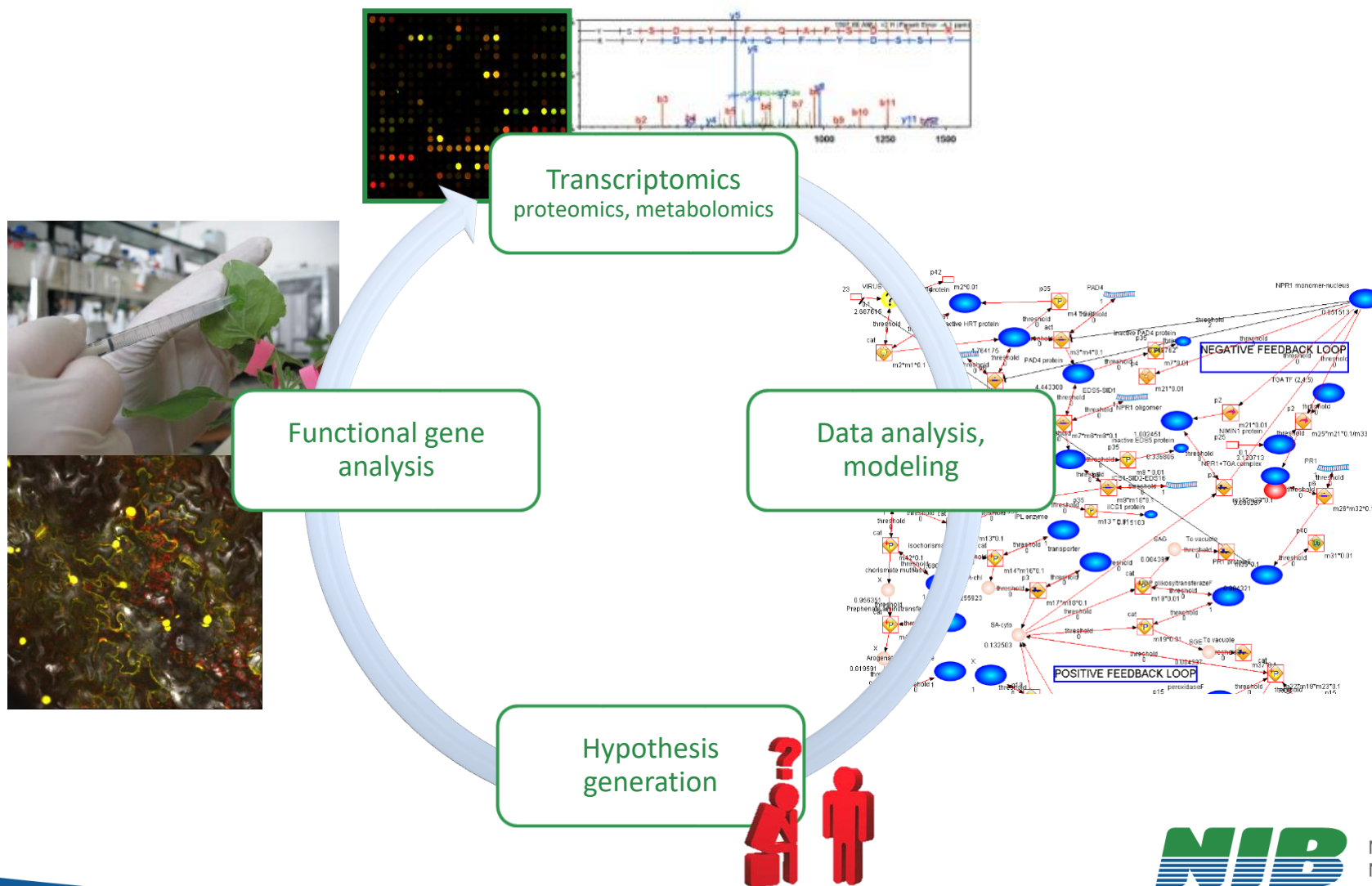
Plant- pathogen/pest interactions



Projects with pharmaceutical industry:
Improved drug production
Development of cancer therapies



Systems biology circle

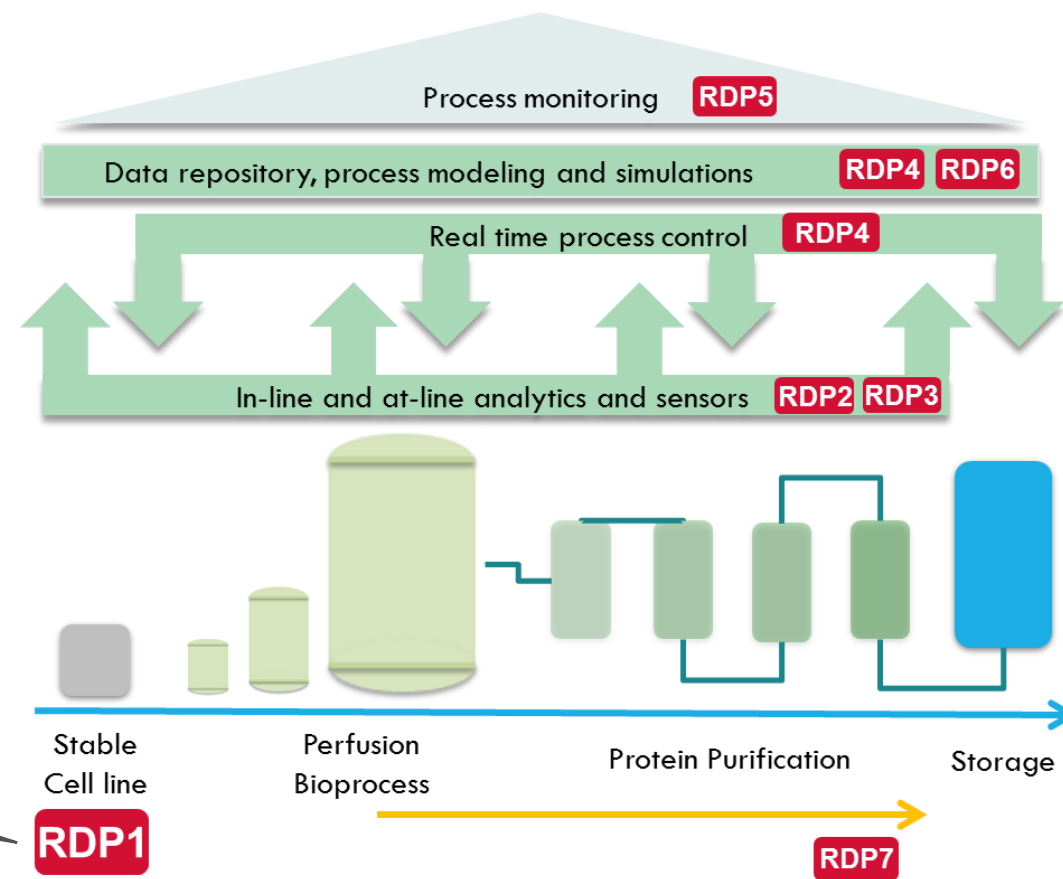


Use case 1: Improving productivity of CHO cells



- 70% of all biologics produced globally is produced in CHO cells
- BioPharmaceuticals Mengeš, part of Lek Sandoz, Slovenia - largest biosimilars development center in Europe
- Biopharm.si project aims to develop continuous manufacturing processes and supporting innovative devices and services

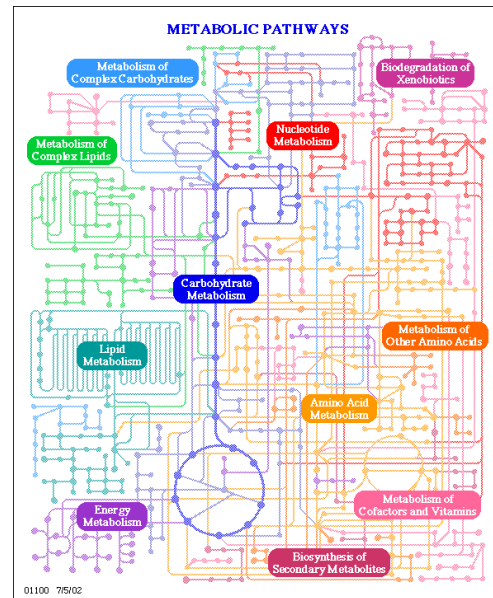
Cell line development
Process optimization



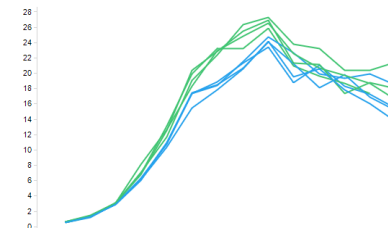
Systems biology approaches to increase the production

- Better designed parental cell lines
- Decrease cell line development time
- Monitor cell line well-being in the bioreactor

From Genomes & Genes



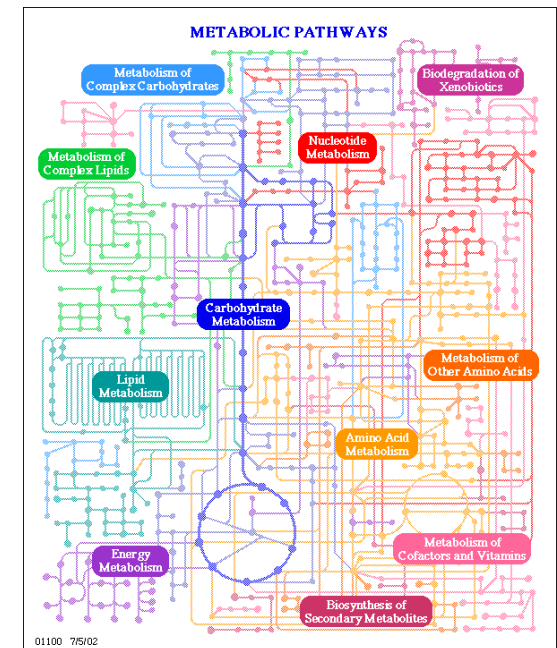
To the Phenotype



Better parental cell lines

Approach: Find the genes with higher expression in low producers
→ knock-out can lead to improved parental lines with increased potential for high productivity

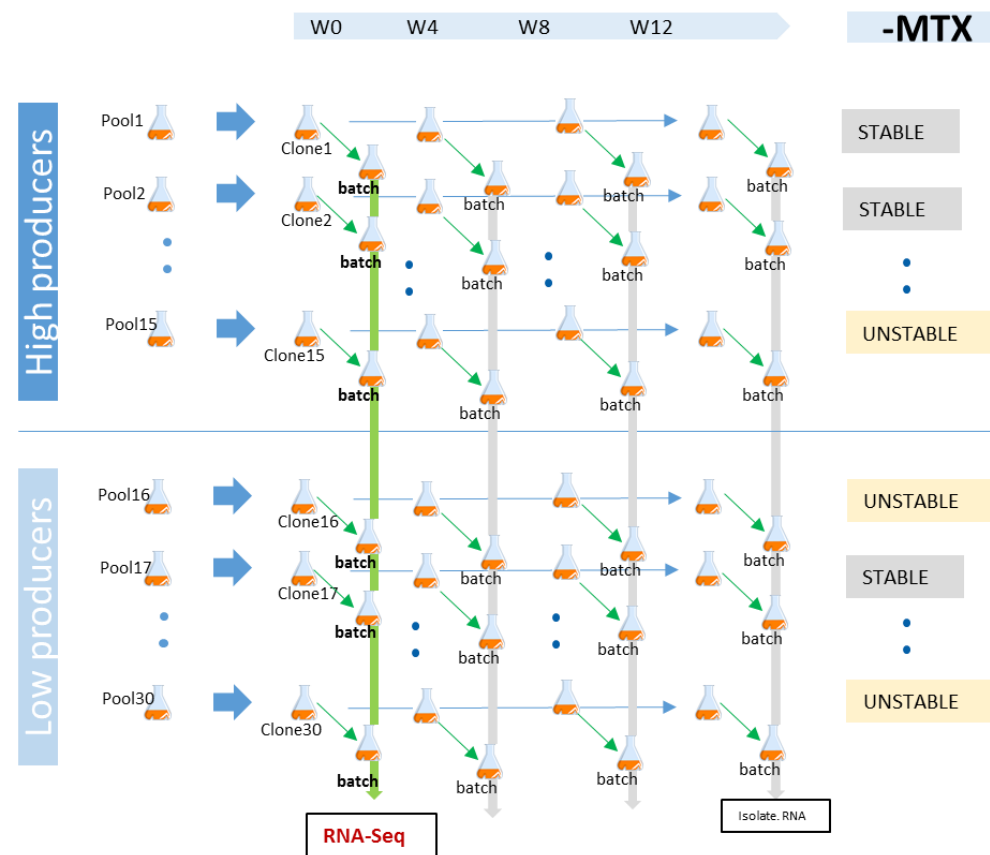
- Available transcriptomics dataset: 5 different parental lines, each from 2 – 5 different production clones, +/-MTX
- Statistical pipeline to identify genes with differential expression
 - no overlap between different cell lines
- Annotation of genes in background knowledge – involvement in relevant processes



Decrease the development time

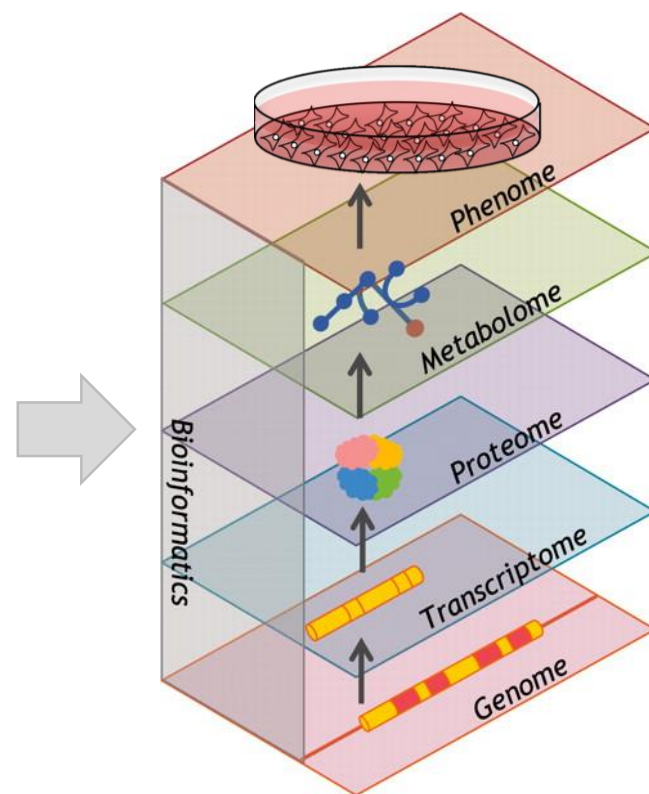
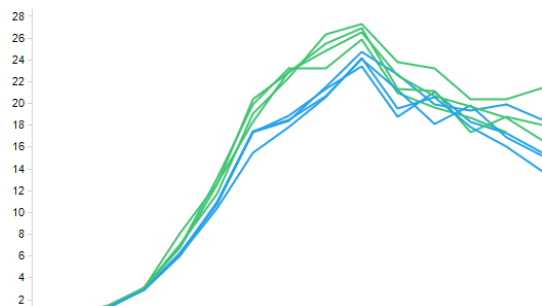
- Checking the stability of cell lines takes 12 weeks: can we predict which cell line is stable already in first weeks?
- Using relevant transcriptomic datasets, we can find markers of stability that we can use in the early stages of cell line development

Jamnikar et al., BMC Biotechnology 2015

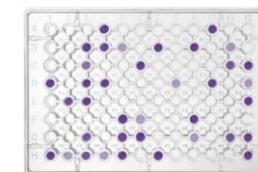


Monitoring the well-being of cells within bioreactor

- Important for the continuous bioprocess
- Metabolic or transcriptomic markers (implemented as at-line sensors)



Metabolic phenotype
(BioLog)



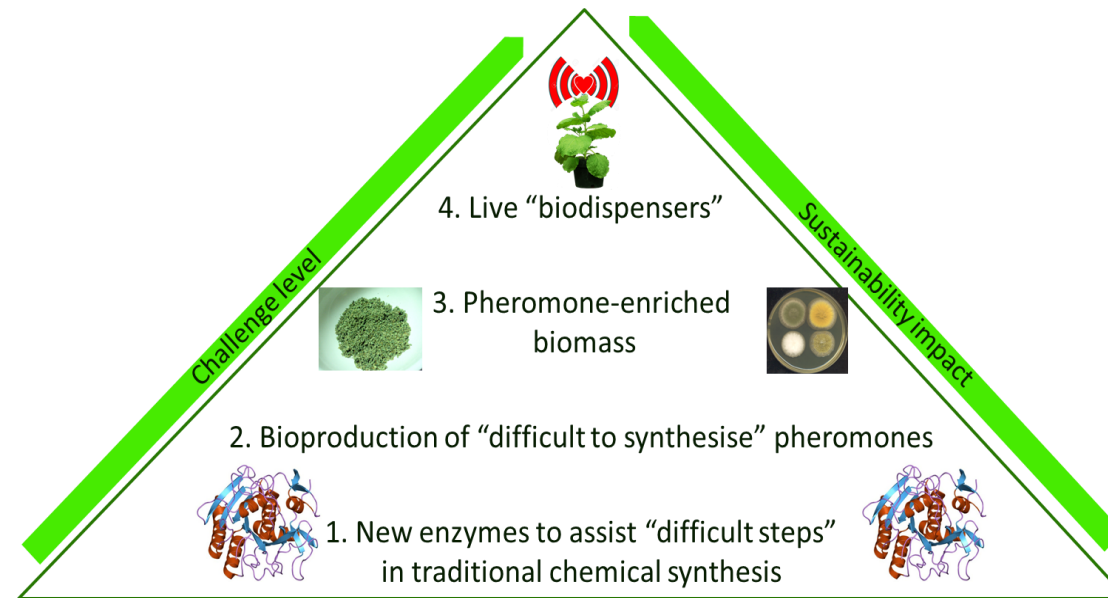
RNA markers
(NanoString, qPCR)



Use case 2: Sustainable Bioproduction of Pheromones for Insect Pest Control in Agriculture

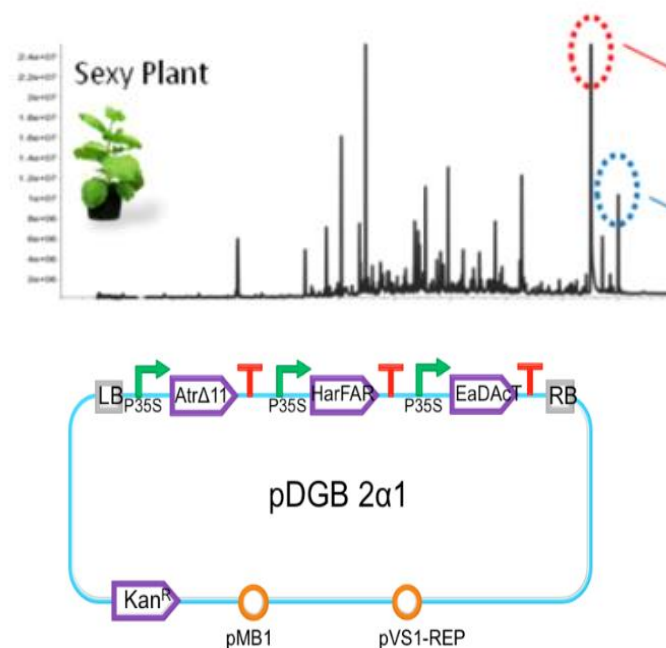
SUSPHIRE project aims to provide sustainable, low-cost biomanufacturing (plant & fungal) platforms for insect pheromones

- Expand the use of sex pheromones to improve agricultural sustainability
- Environmentally-friendly production



Aim 1: Demonstrate high yields of lepidopteran pheromones

Sexy plants 1.0: expressing 3 lepidoptera pheromone biosynthesis genes



Low-producing line

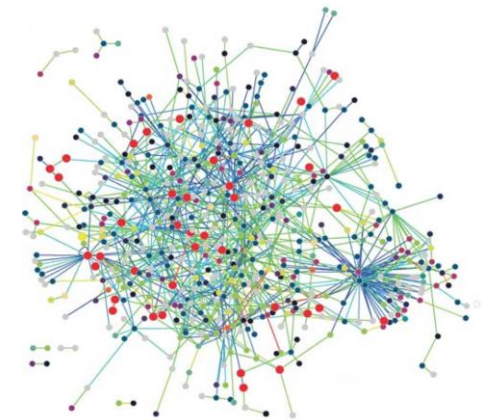


High-producing line

→ Use of systems biology tools to identify regulatory bottlenecks causing growth penalty

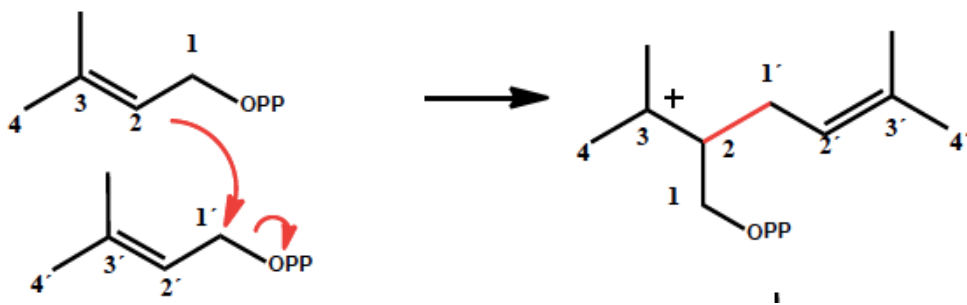
Aim 1: Demonstrate high yields of lepidopteran pheromones

- RNAseq analysis: 2 low producing, 2 high producing lines, wt
- The majority of gene expression changes are linked to pheromone production, but not to transgene insertion
- Stress response in high producing plants
- Further network analyses to identify regulatory bottlenecks



Aim 2: Discover & engineer mealybug pheromone biosynthesis pathways

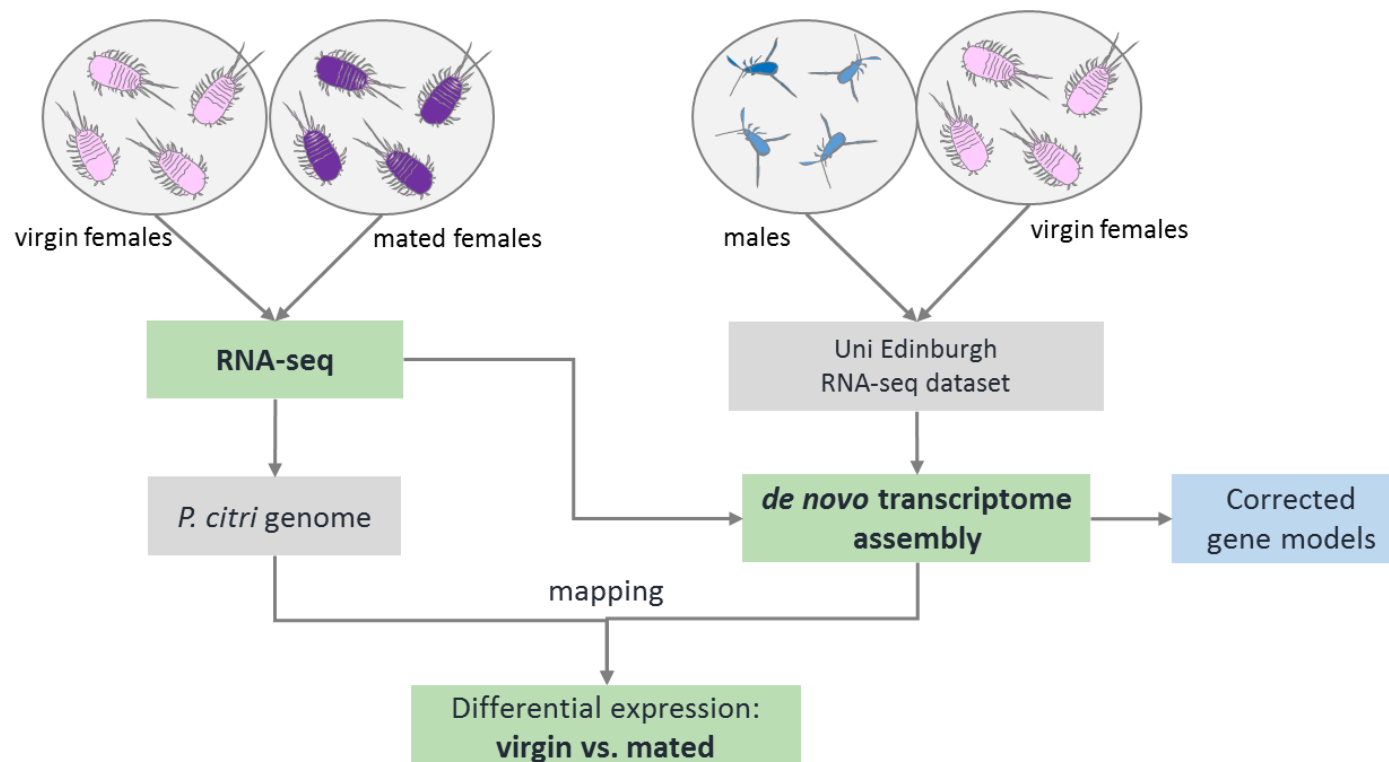
- Mealybugs – important plant pests
- Mealybug pheromones = irregular monoterpeneoids (unusual NON-head-to-tail linkage of two isoprene units)
- Chemical synthesis difficult & biosynthesis unknown



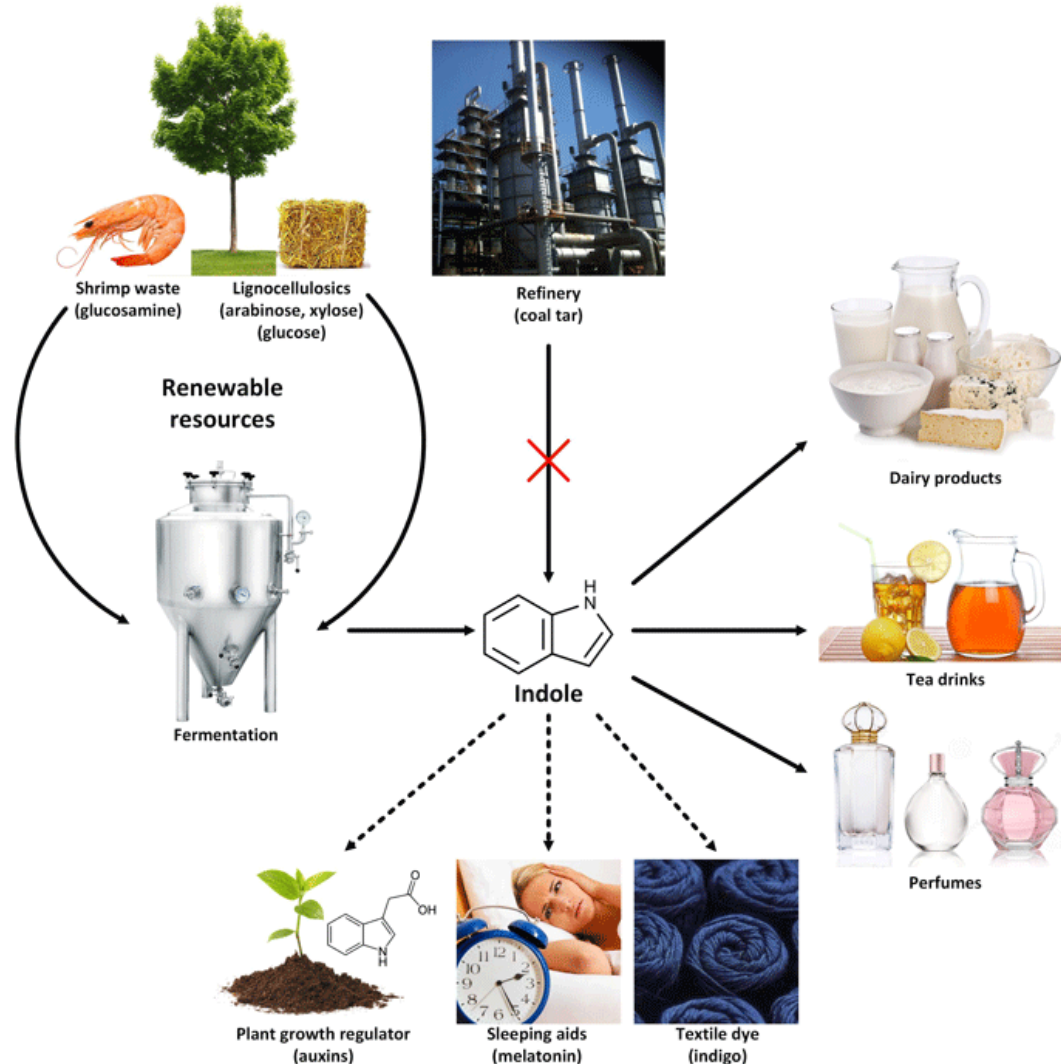
ECONOMIC IMPORTANCE

Aim 2: Discover & engineer mealybug pheromone biosynthesis pathways

Identification of isoprenyl diphosphate synthase (IDS), first step in *Planococcus citri* pheromone biosynthesis) from transcriptome data



Use case 3: Biotechnological production of indole



Establish sustainable production of indole

- *Corynebacterium glutamicum* (industrial amino acid producer)
- Identification of biosynthetic enzymes in bacteria in plants
- Pathway design and optimization
- Set up industrial scale production, using renewable resources

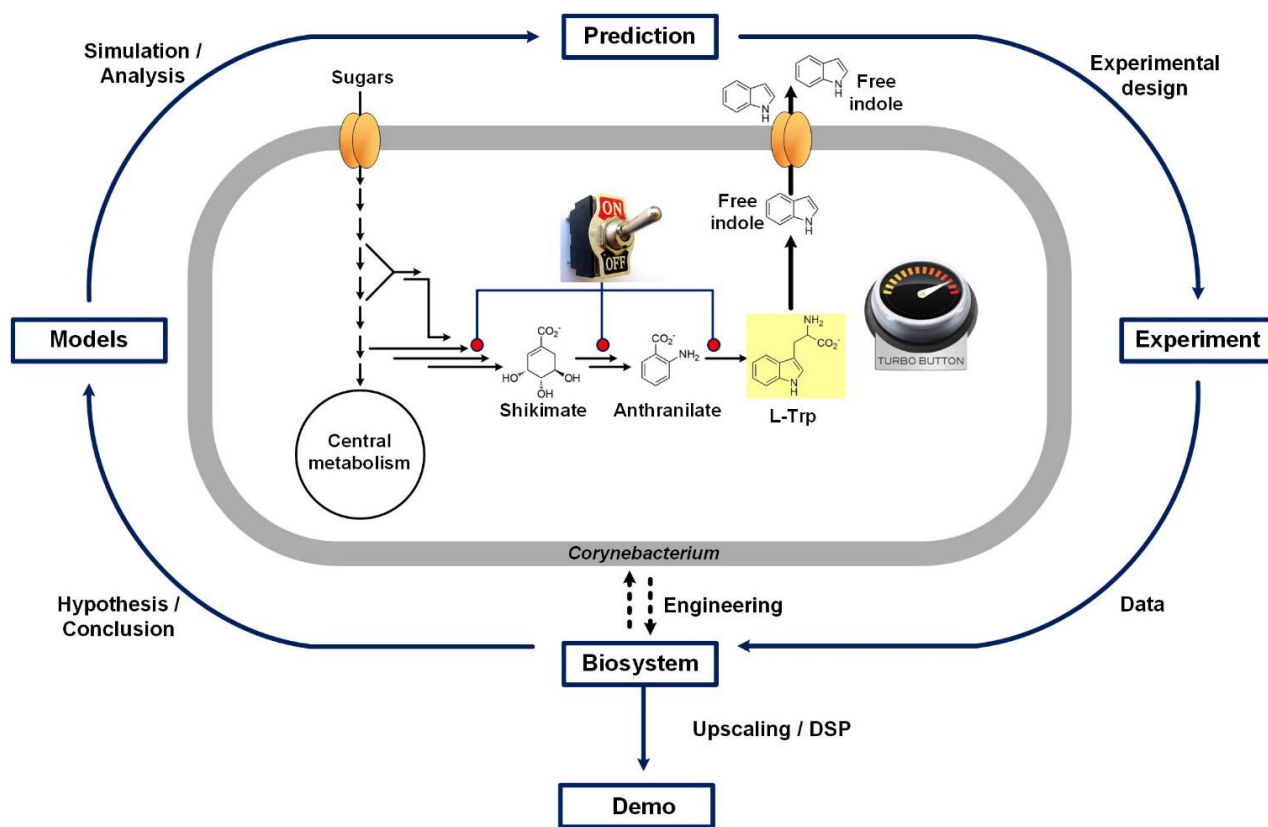
Identification of indole synthases in plants

Looking for INS motifs in proteomes of plant clades that produce indole derivatives using iGLOSS tool



→ Candidates for functional analysis identified

Optimization of designed production strains



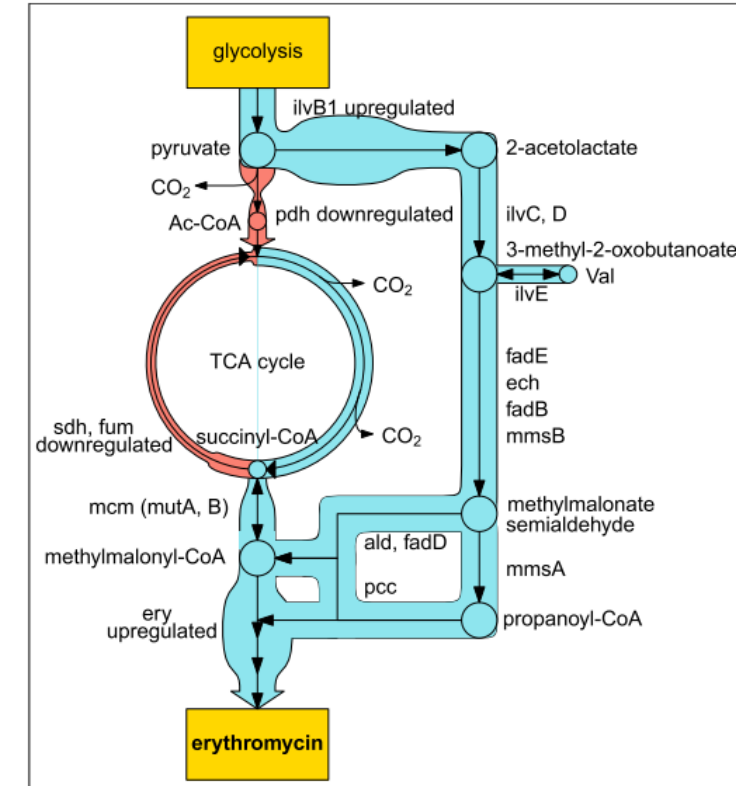
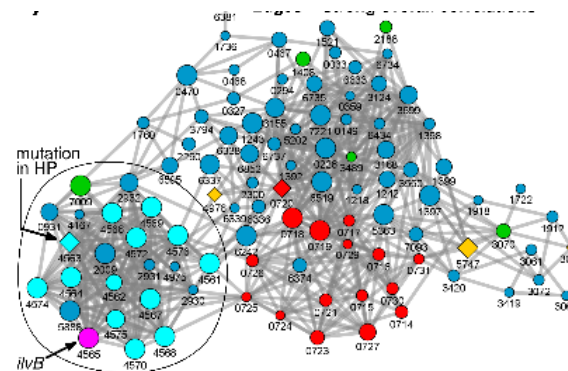
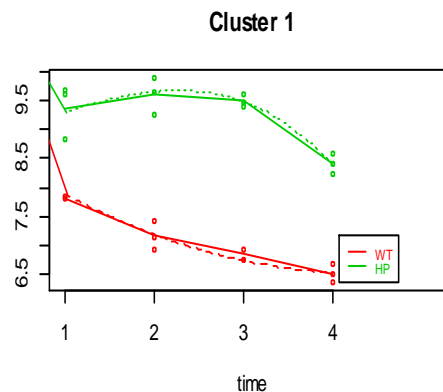
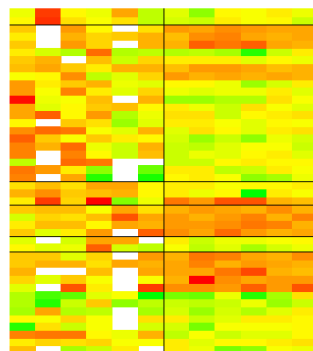
Integration of transcriptomic and metabolomic data with the *C. glutamicum* metabolic model

→ Identification for additional targets for optimization

Use case 4: Improving antibiotic production

Improvement of *Saccharopolyspora erythraea* production strain

- Transcriptomics, proteomics in high-producing and wt strain
 - Time-course statistical analysis
 - Data integration, visualization & network analysis
- Identification of key regulatory elements and targets for pathway reengineering



Karničar et al., *Microbial Cell Factories*, 2016