

EPSO Biofuels Workshop

London, 28-29 May 2008

Metabolic engineering for production of fuels and chemicals from biomass



Teknologiasta liiketoimintaa

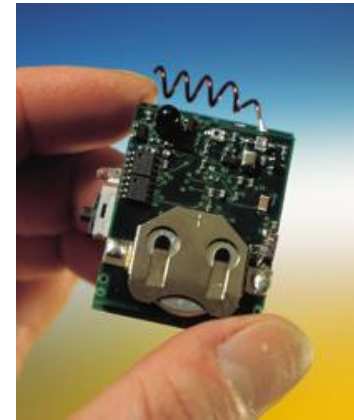
Merja Penttilä

VTT

VTT

(Valtion Teknillinen Tutkimuskeskus)

Through creating and applying technology, we actively enhance the competitiveness of industry and other business sectors, and thus increase the welfare of society.

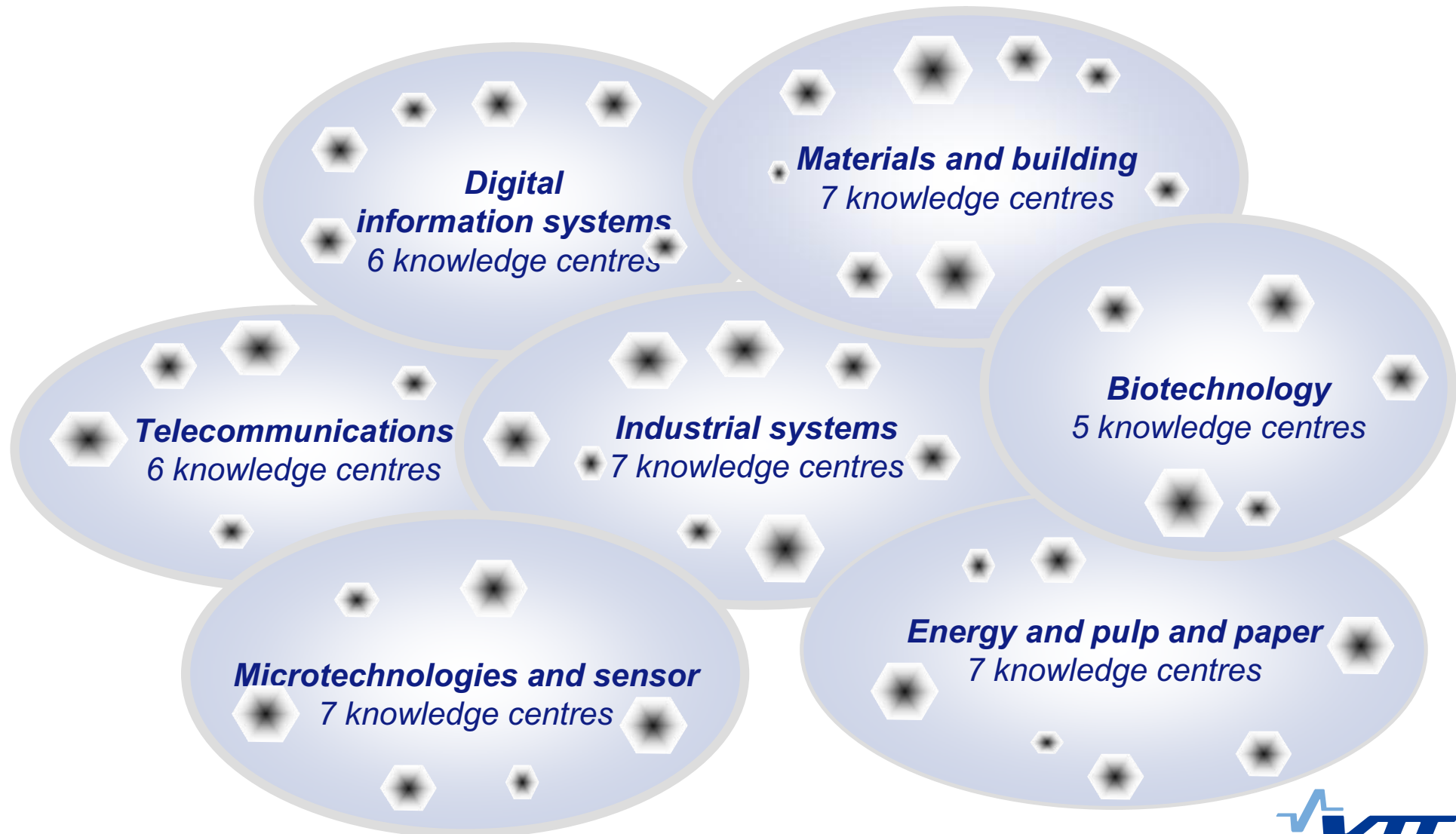


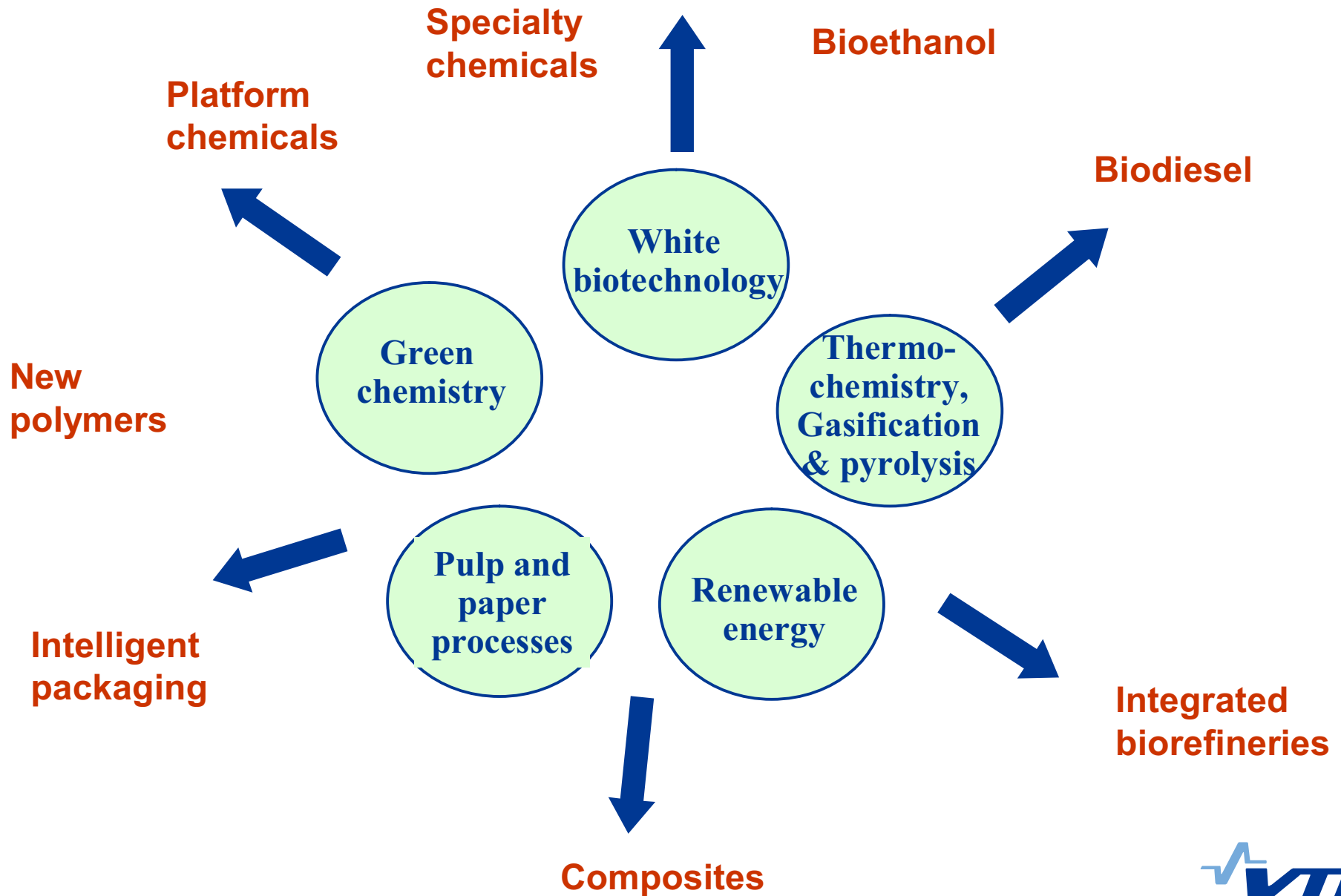
- Governmental, not-for profit research organisation
- Biggest in Nordic countries
- Staff: ~ 2800
- R&D activities range from fundamental public research to confidential industrial projects



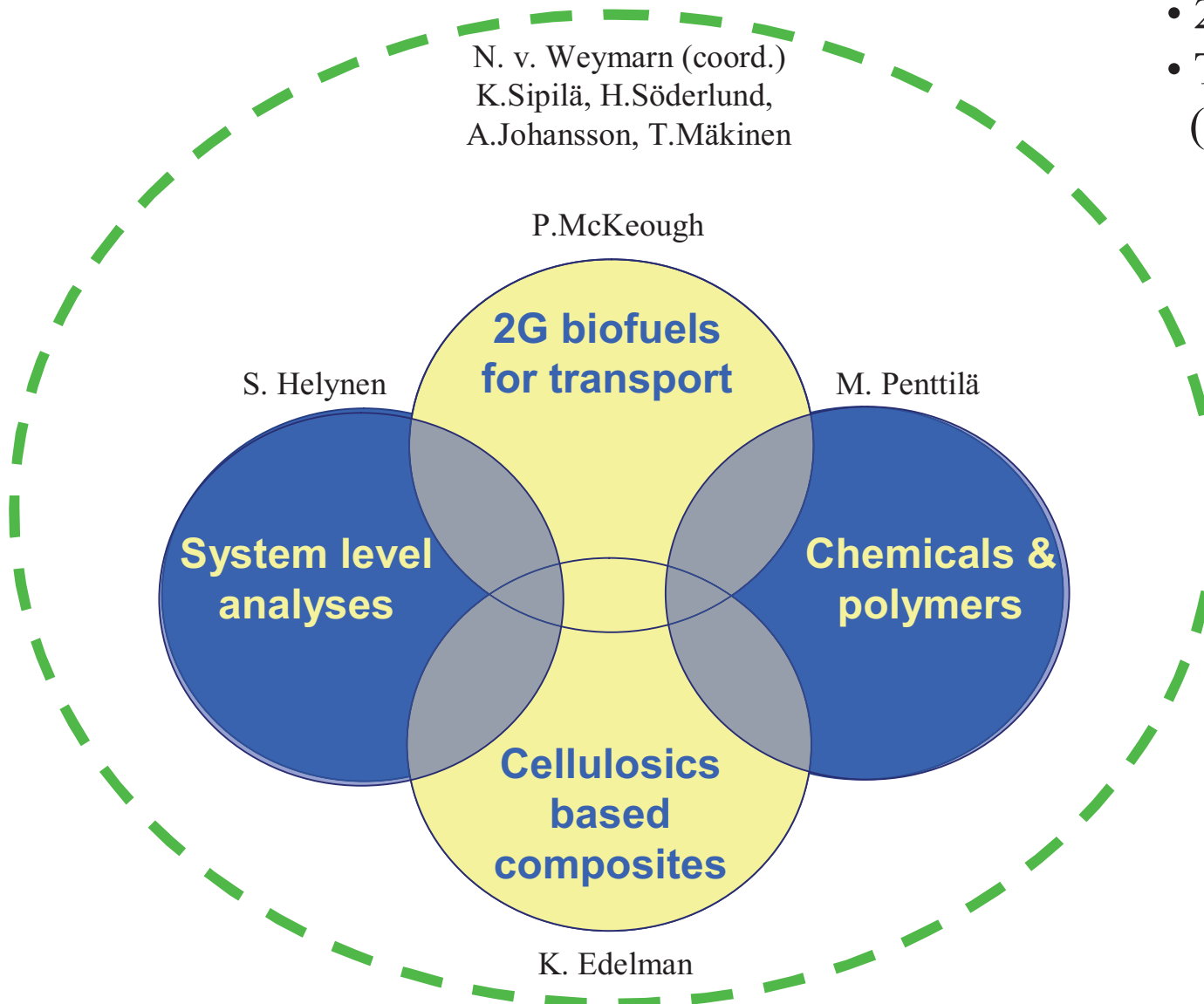
RESEARCH AND DEVELOPMENT

7 knowledge clusters and 45 knowledge centres

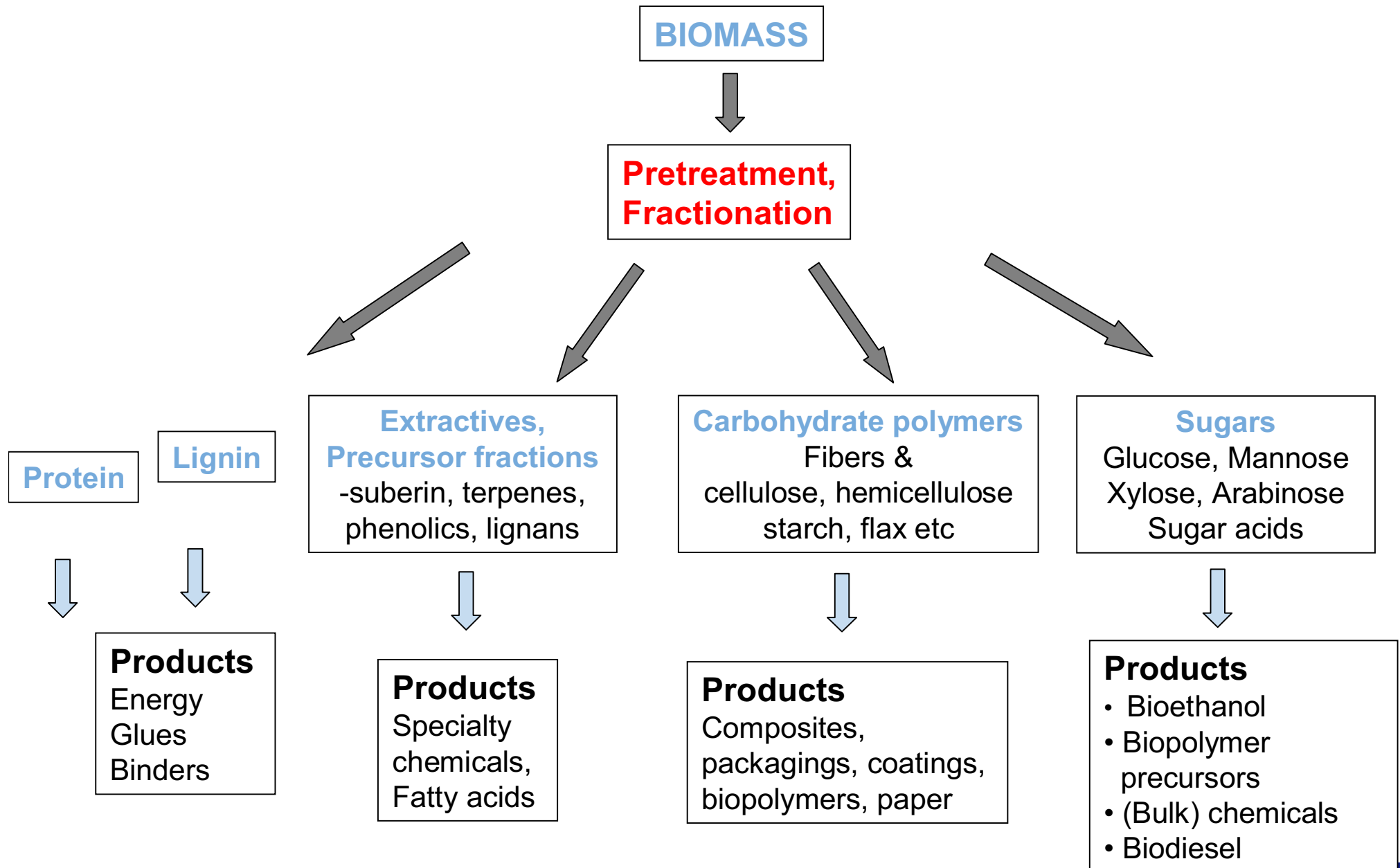




- 2007-2011
- Total budget ~ 40 mill.€
(from VTT ~ 20 mill.€)

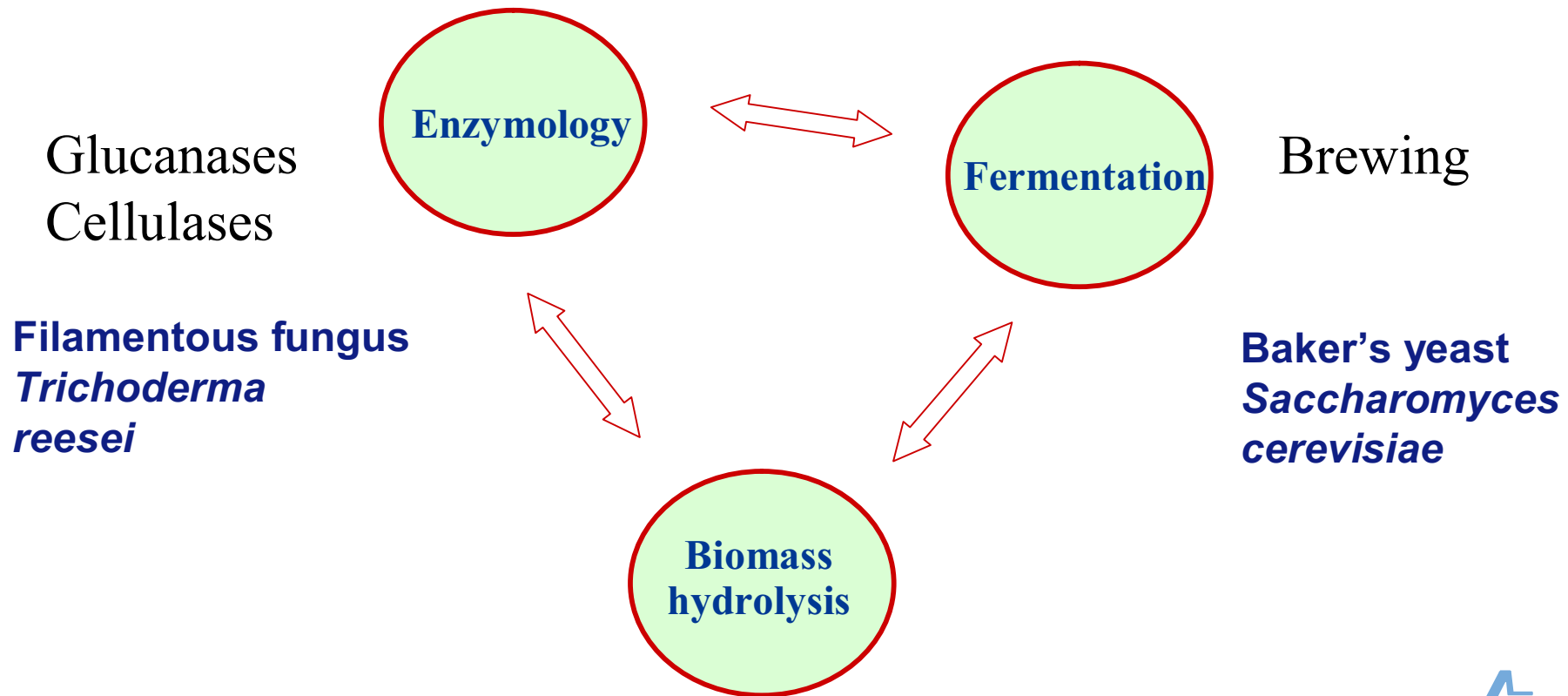


Good fibers, easy sugars, interesting extractives ?



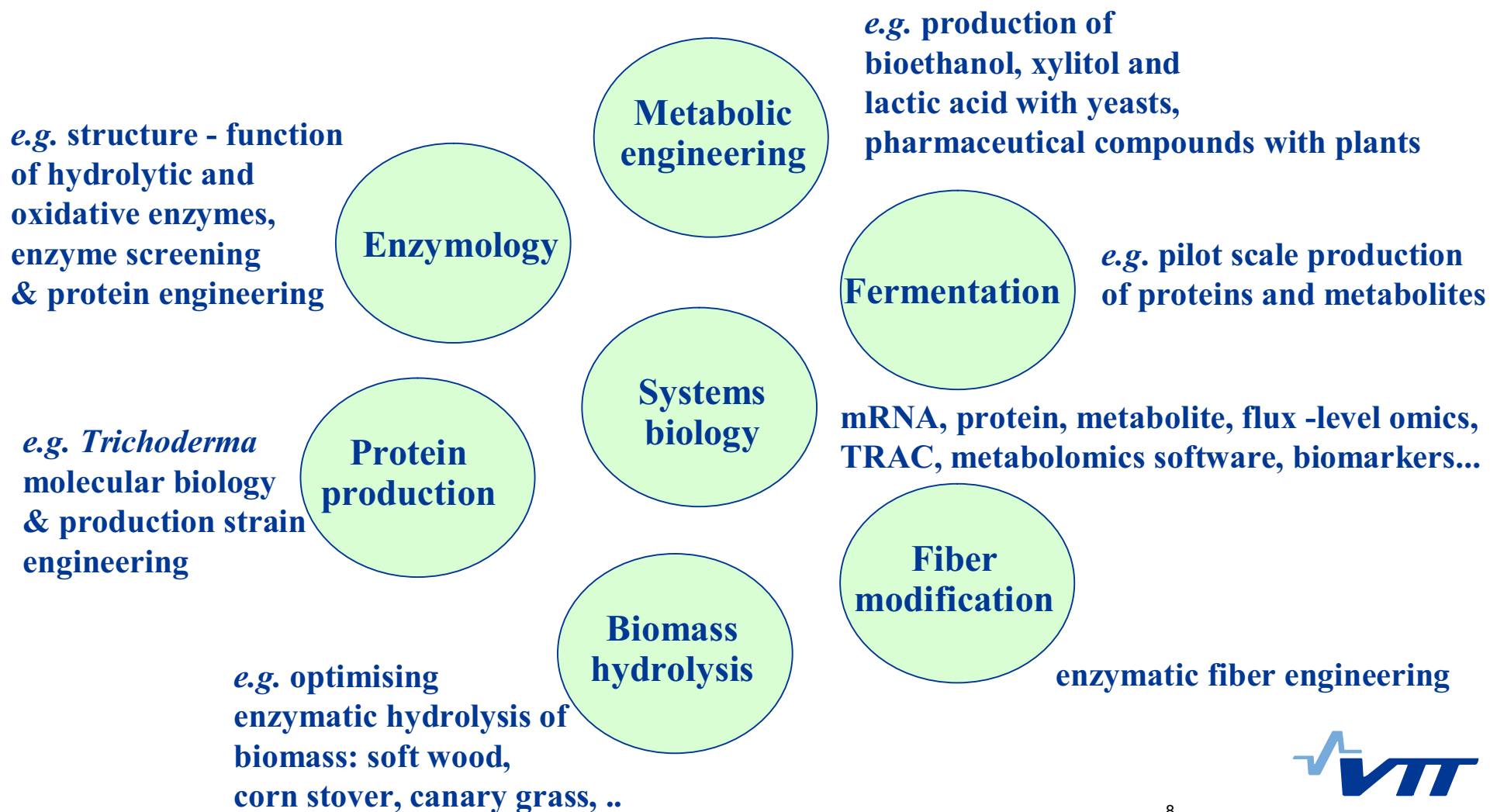
VTT is an expert in the essential disciplines of Industrial Biotechnology

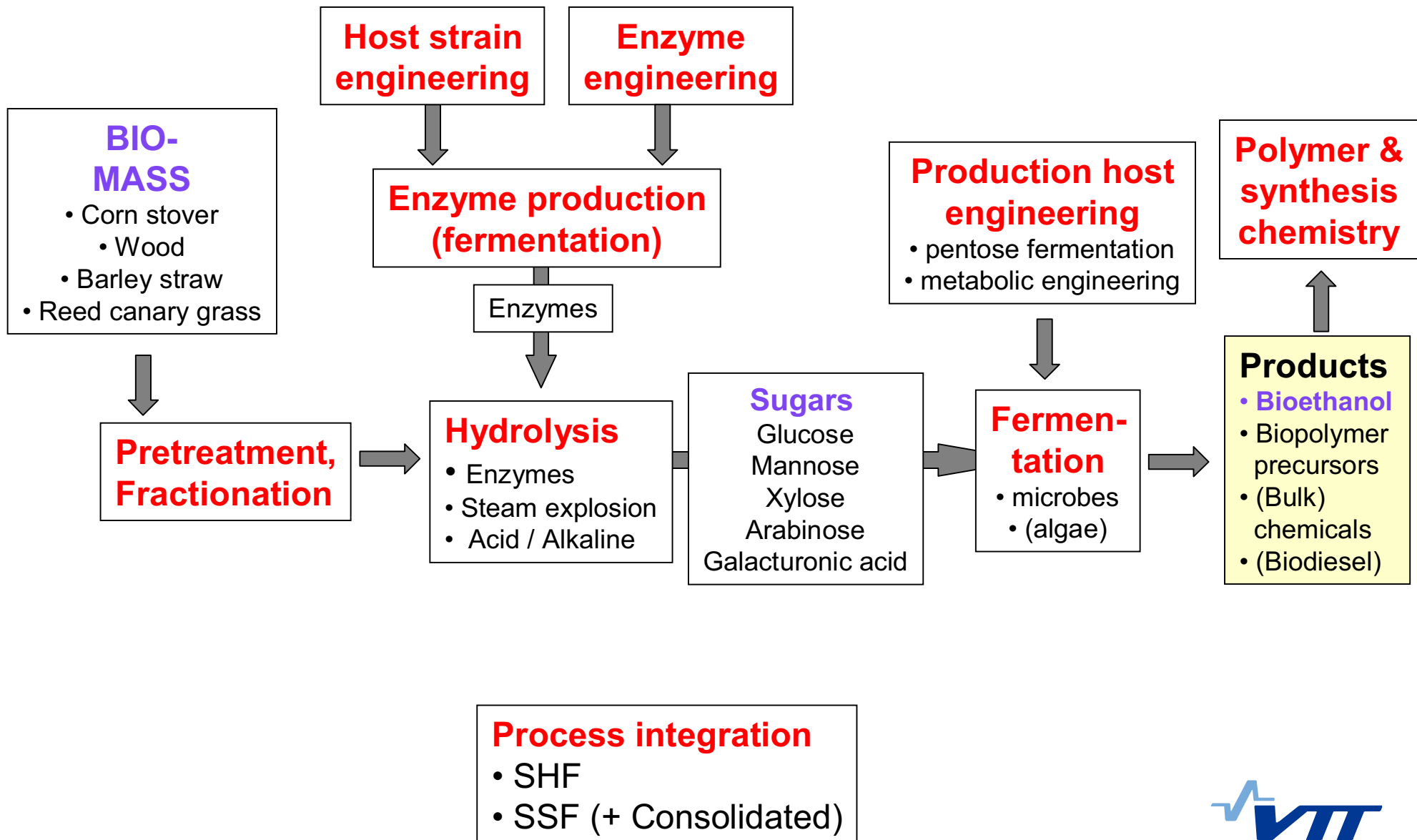
Research started in the 70's



VTT is an expert in the essential disciplines of Industrial Biotechnology / White Biotechnology

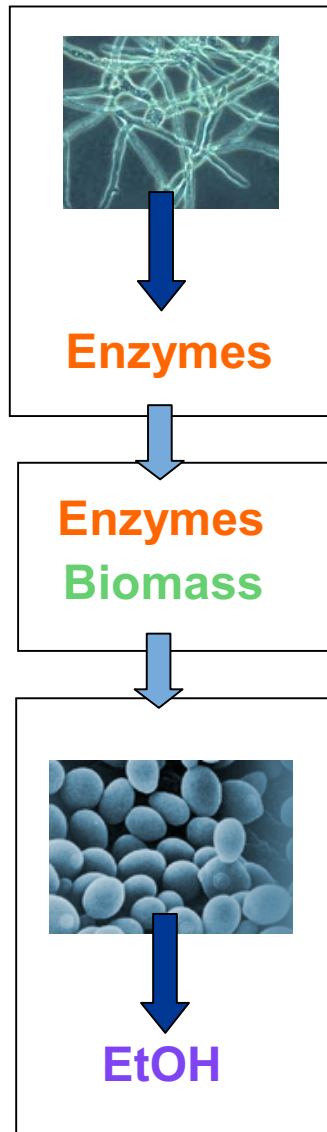
Last 25 years: > 1000 publications, > 100 patents





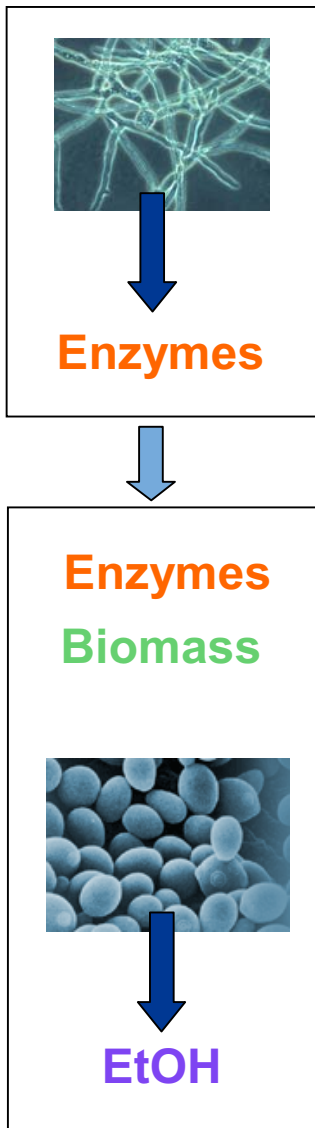
SHF

Separate hydrolysis
and fermentation



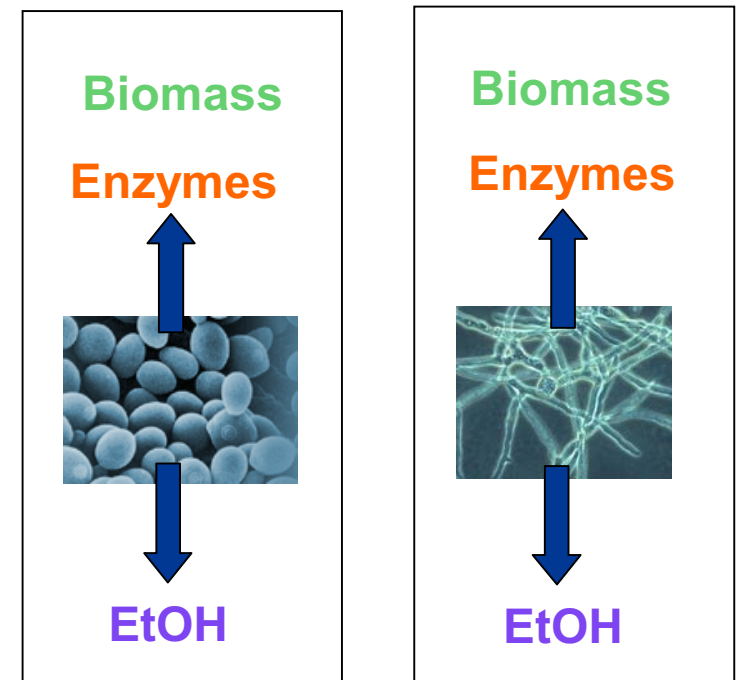
SSF

Simultaneous saccharification
and fermentation



CBP

Consolidated bioprocessing



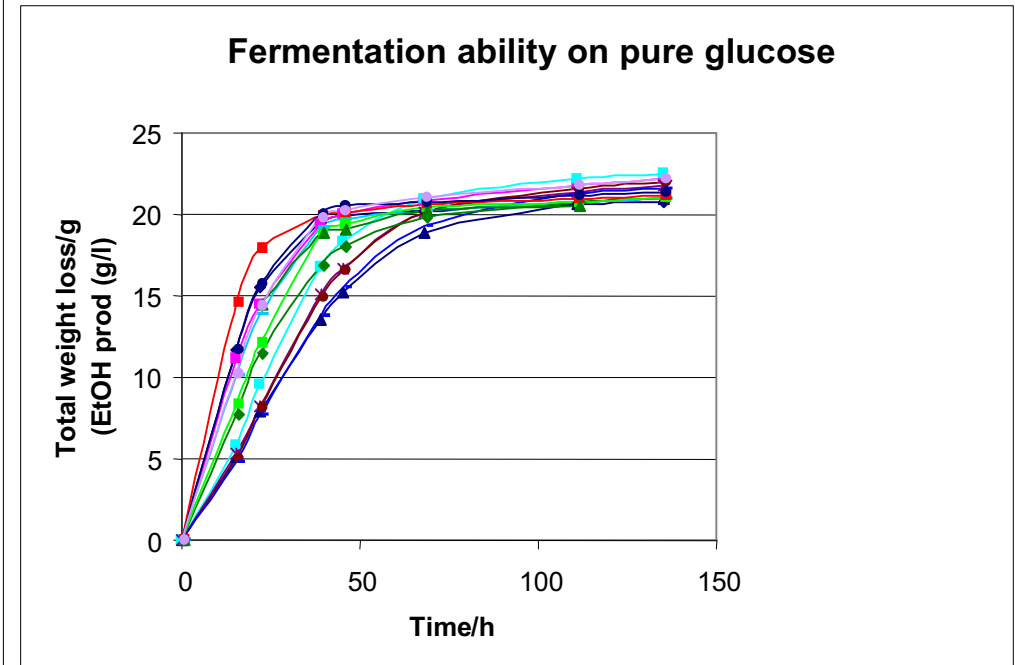
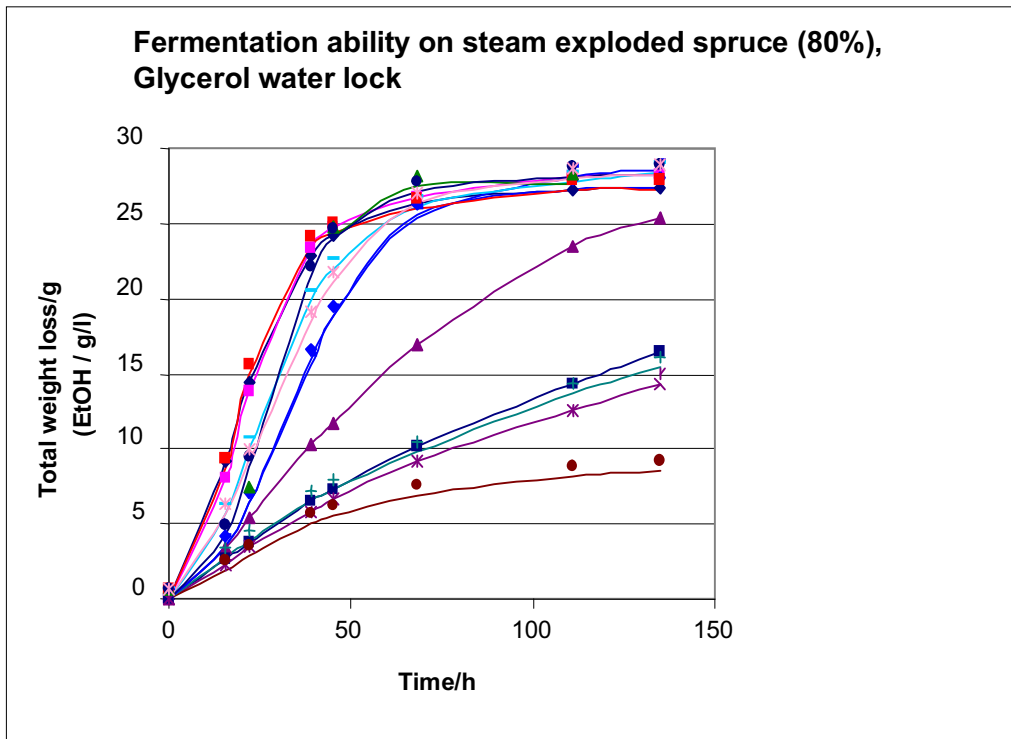
In SSF and CBP temperature for
hydrolysis and fermentation needs to
be compatible

Requirements of the production organism for 2nd generation bioethanol

- Generally robust in large scale production (~ 1 000 m³)
- Ethanol tolerant (> 8%)
- Good fermentation rate (>1g/L/h)
- Compatible with processes not prone to contaminations (low pH, high T)
- Require minimal nutrients
- Flexible with air (facultative anaerobes)

- Tolerant towards inhibitors in lignocellulosic hydrolysates
(acetic acid, furfural, phenolic compounds)
- Able to ferment all major biomass C6 and C5 sugars (glucose, xylose, arabinose)
- Able to ferment certain sidestream sugars (arabinose, galacturonic acid)

Comparison of different *S.cerevisiae* yeast strains



Examples of microbes studied for bioethanol production

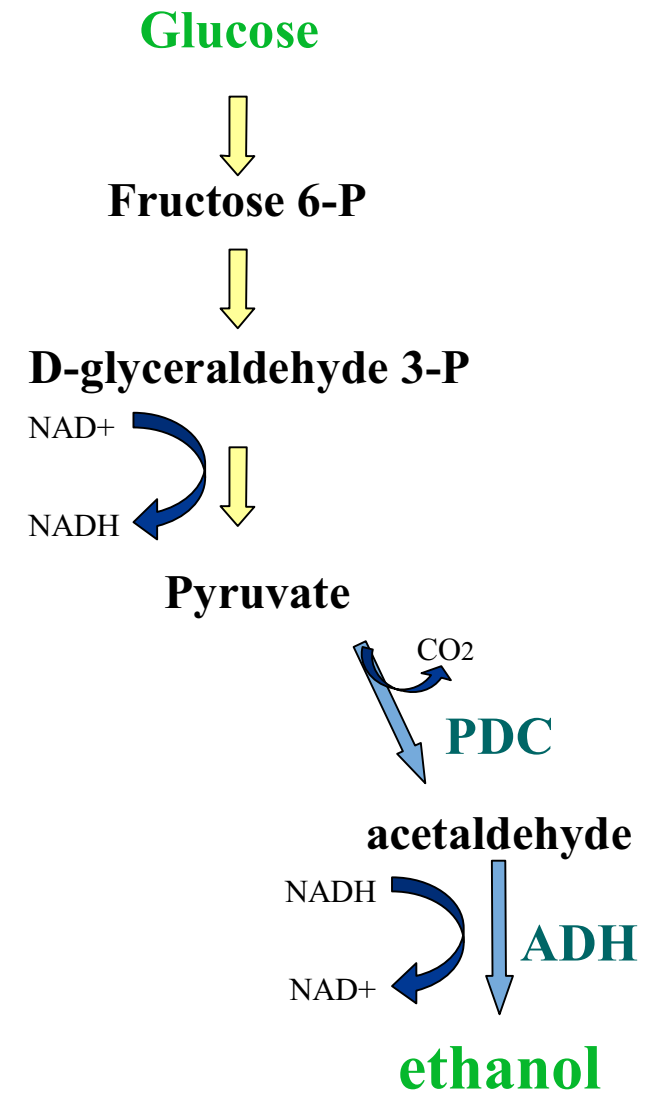
- *Zymomonas mobilis*; naturally ethanol producing
engineered to ferment xylose and arabinose
- *E.coli*; utilises naturally xylose, engineered to produce ethanol
- *Klebsiella*; engineered to produce ethanol
- *Clostridia* (gas fermentation)

General drawbacks with bacteria: low inhibitor, low pH, ethanol tolerance and process robustness, more complex nutrient requirements

- *Pichia stipitis*; naturally xylose utilising
- *Saccharomyces cerevisiae*; current production host for ethanol, GRAS
 - engineered to ferment xylose and arabinose
 - engineered for reduced byproduct formation, e.g. xylitol and glycerol

Drawbacks with engineered *S.cerevisiae*: insufficient C5 fermentation rate, C5 uptake

Glucose fermentation is efficient

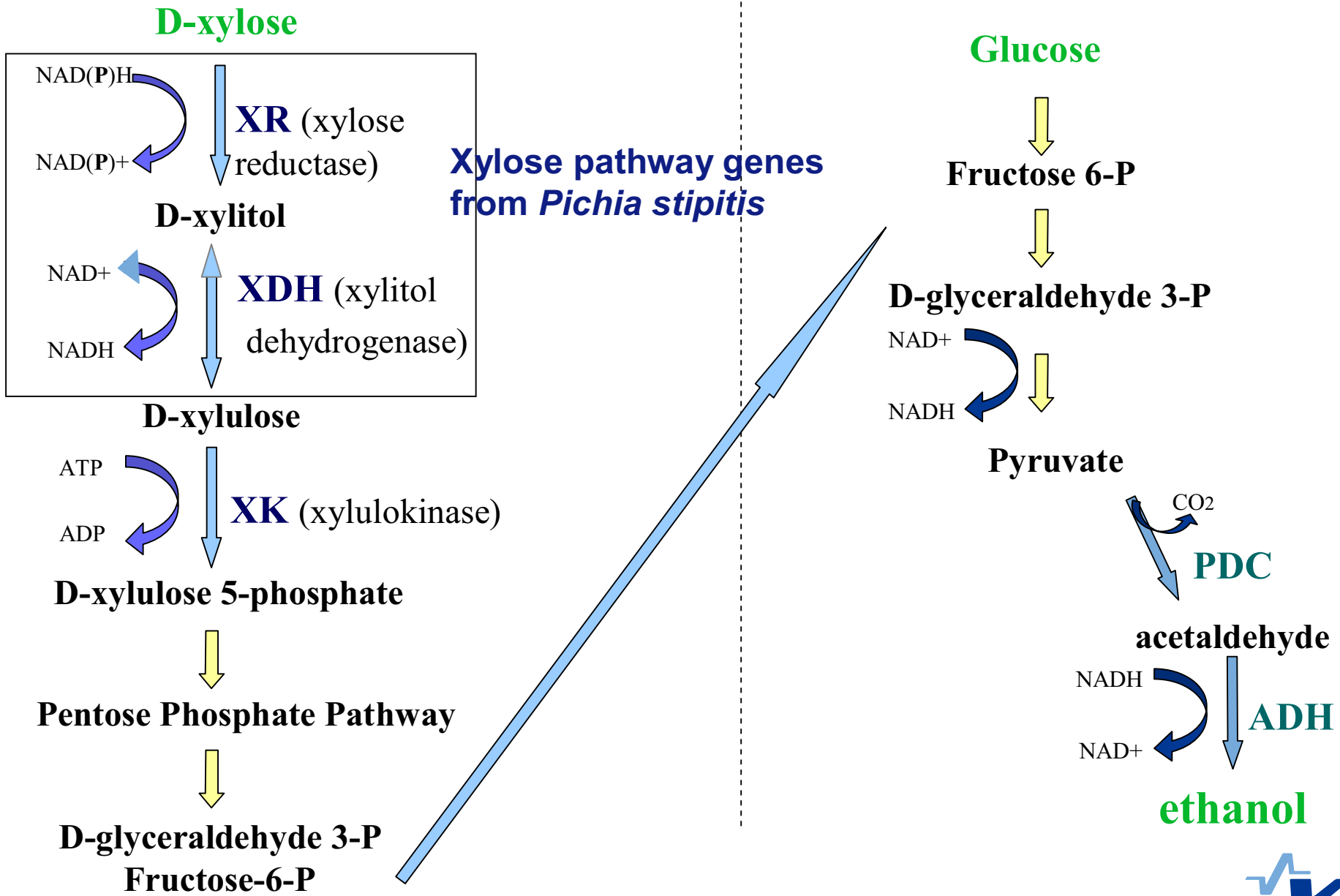


Saccharomyces cerevisiae yeast



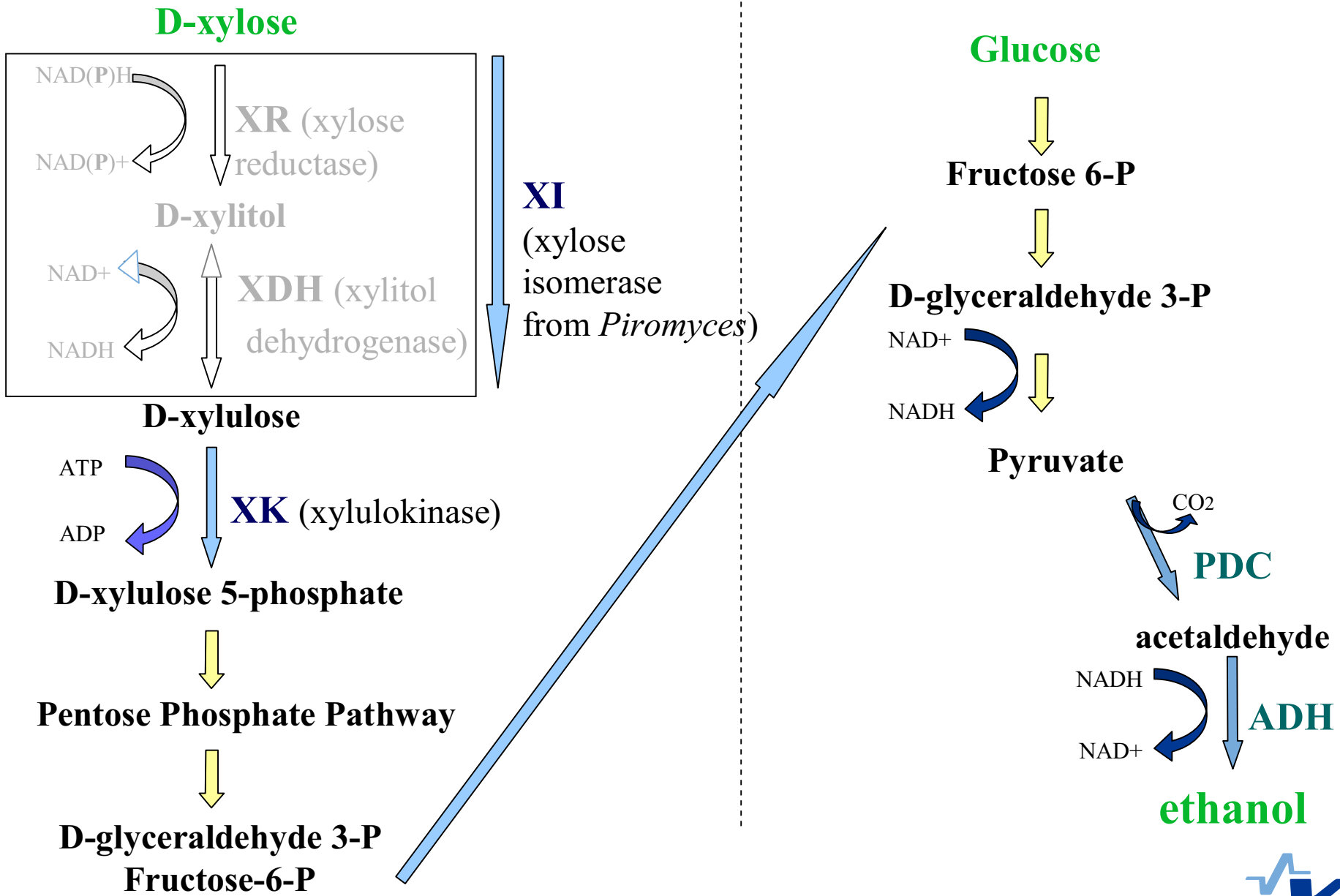
Pentose utilisation

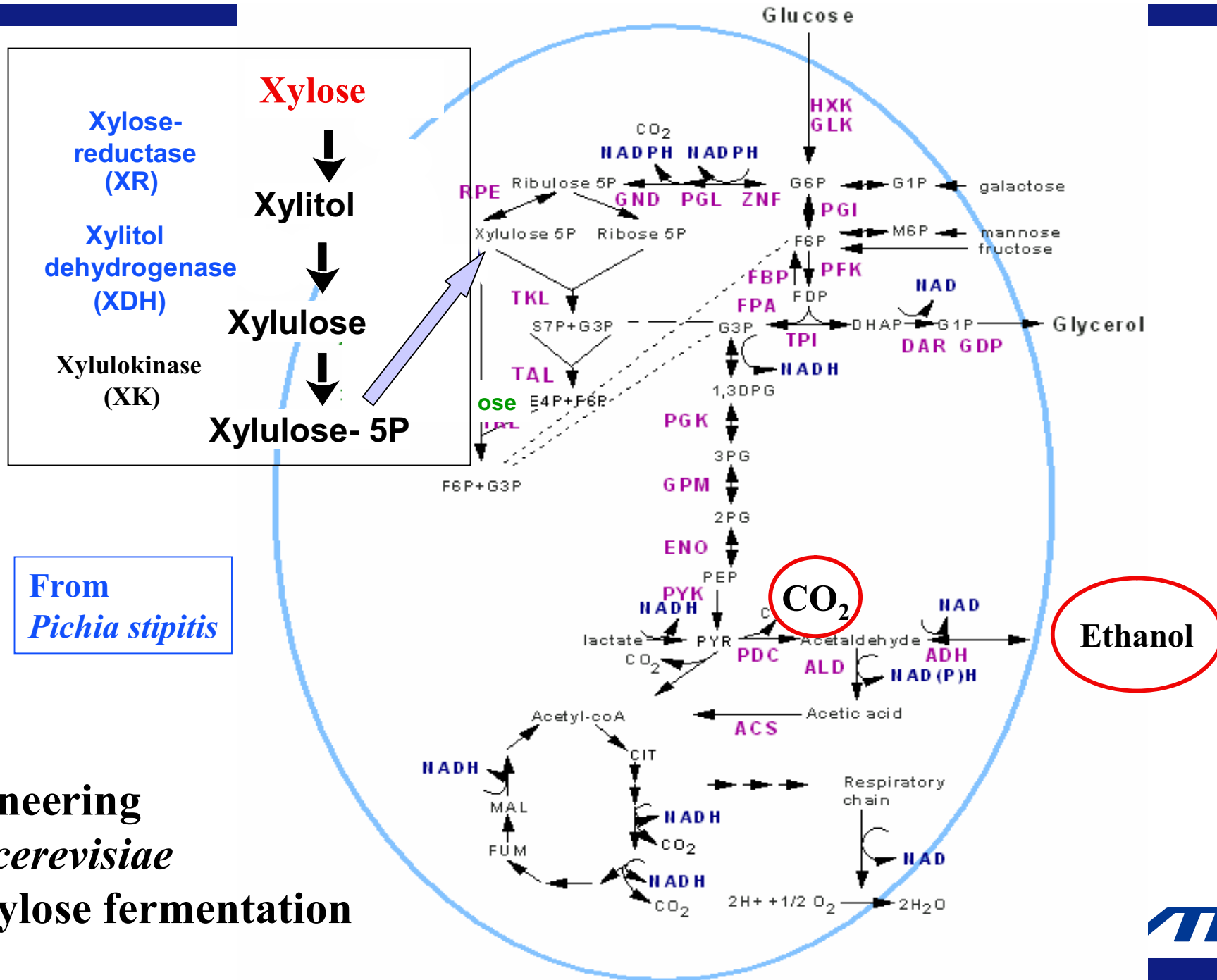
Glycolysis



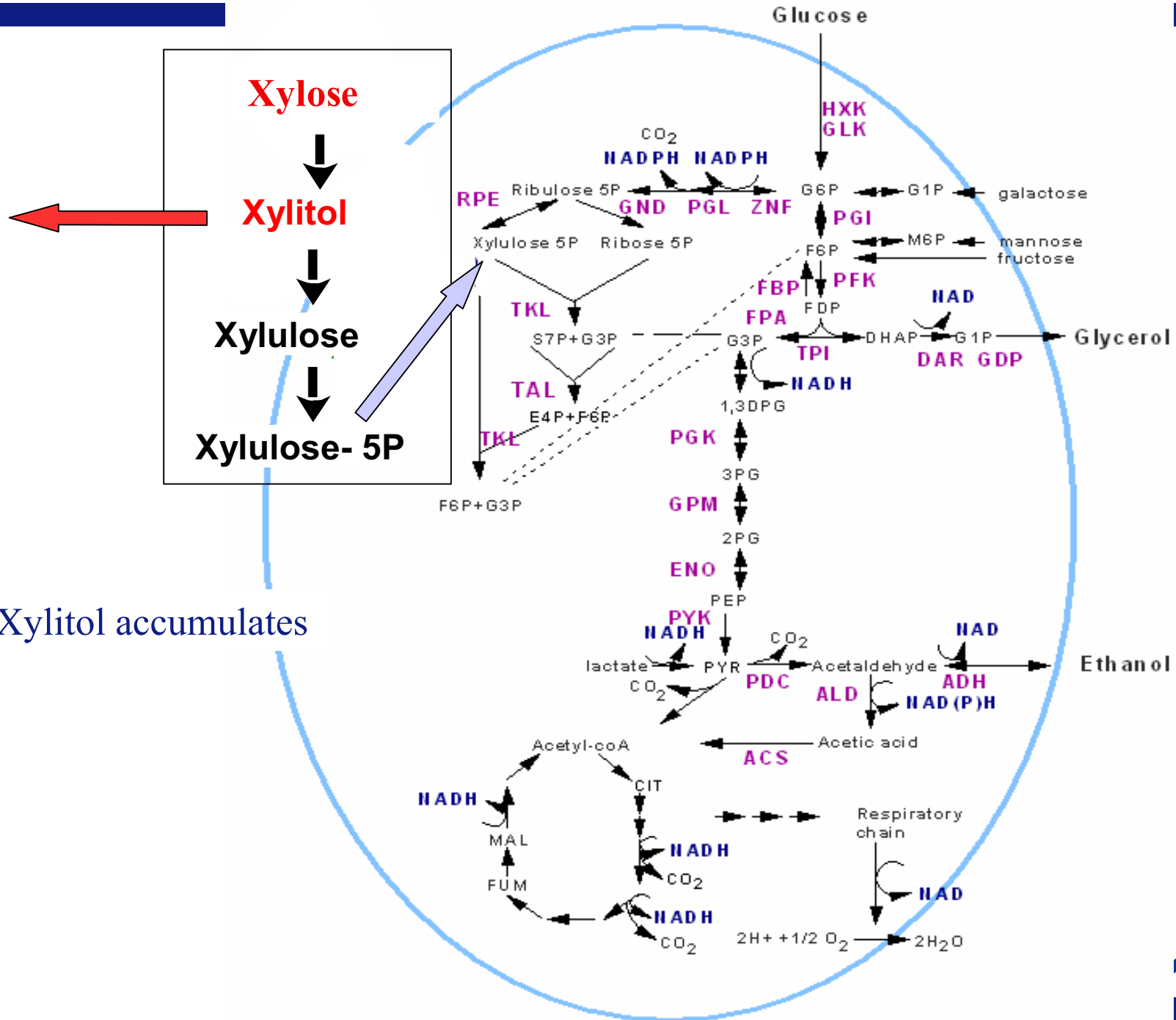
Pentose utilisation

Glycolysis





Engineering of *S.cerevisiae* for xylose fermentation

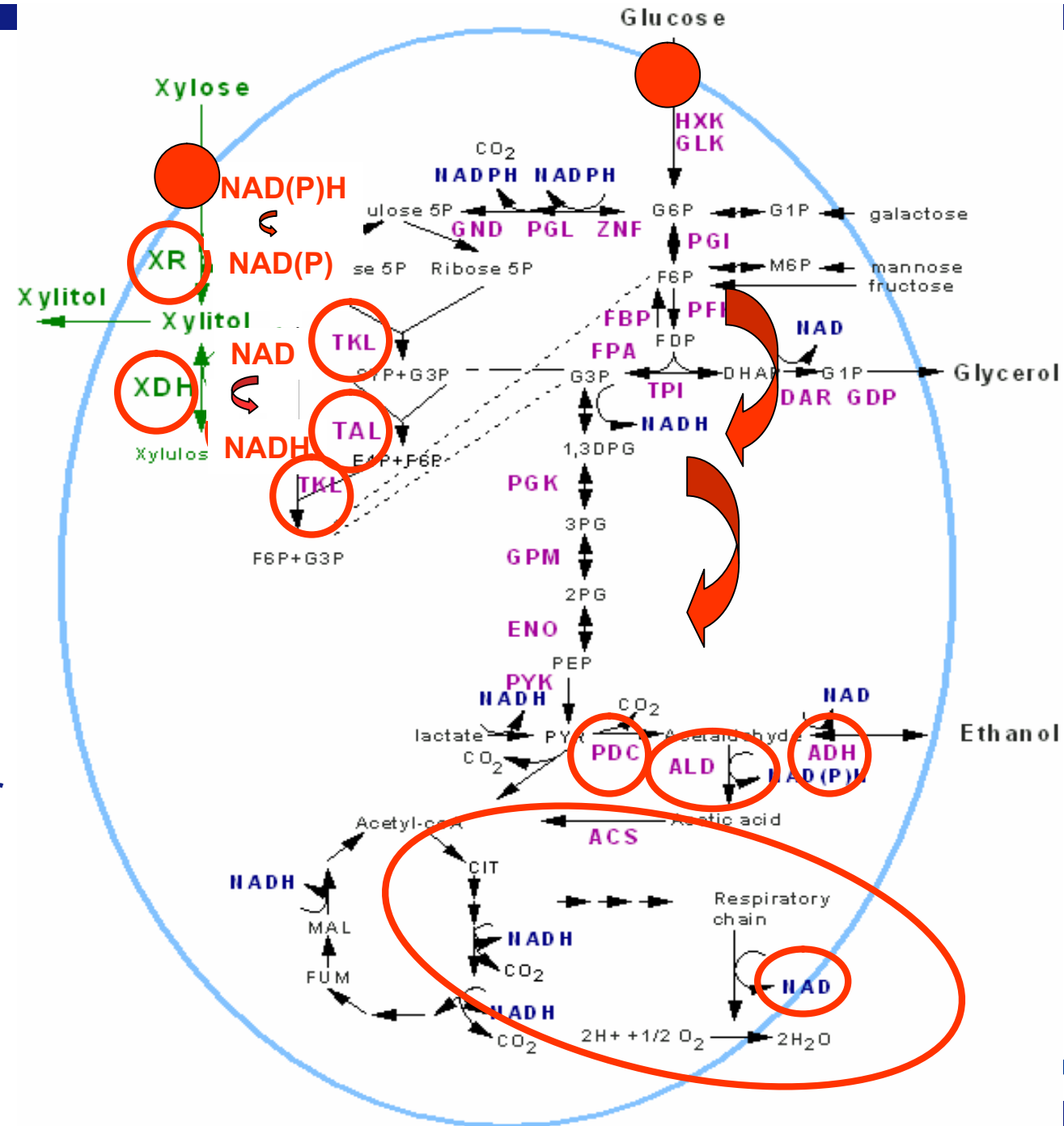


• Xylitol accumulates

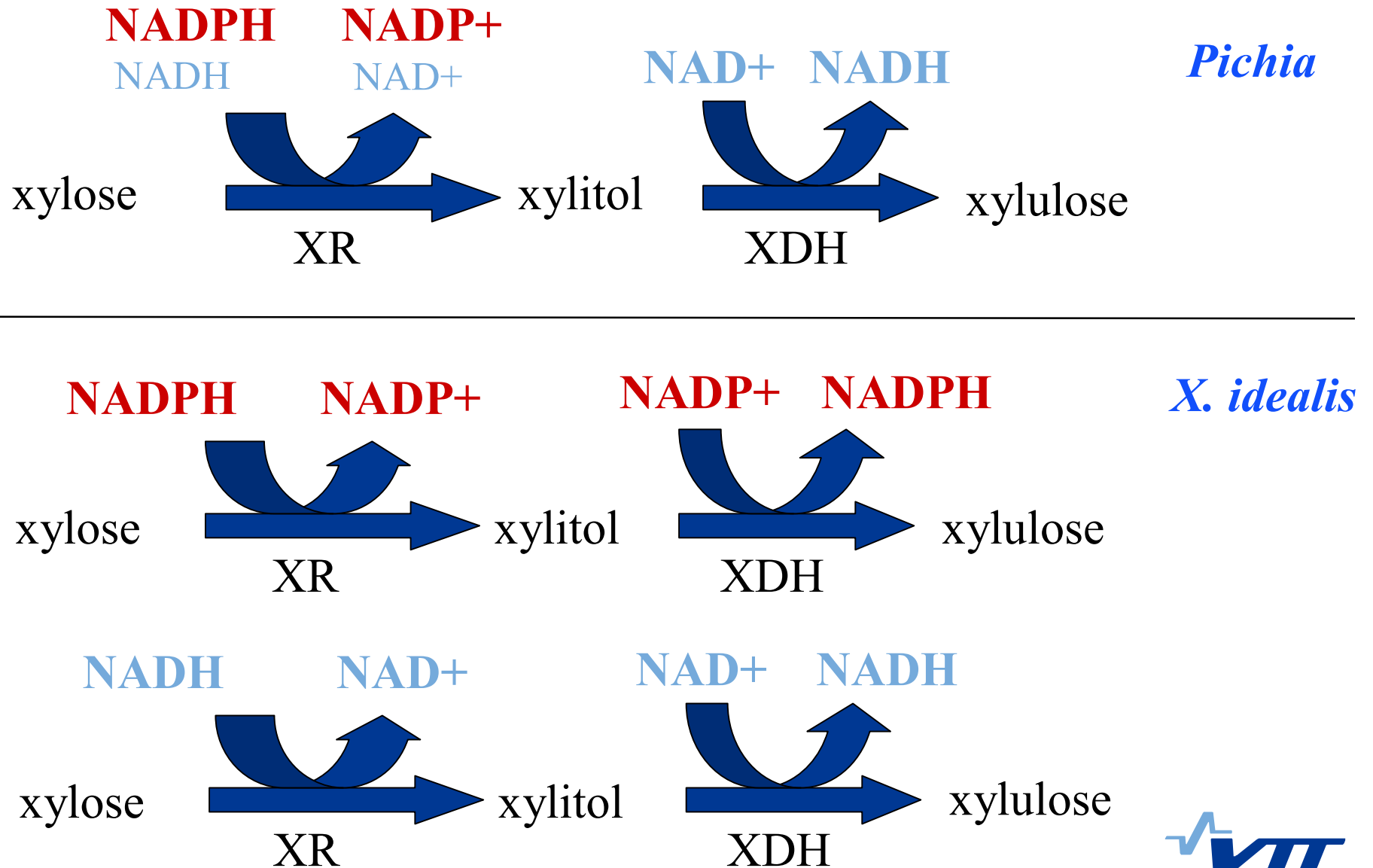
Bottle neck suspects in xylose fermentation in recombinant *S.cerevisiae*

- *sugar uptake*
- *inefficient PPP*
- *general regulatory issues?*
- *redox imbalance;*

problematic in particular in anaerobic conditions, where respiration is not functioning in NAD⁺ regeneration



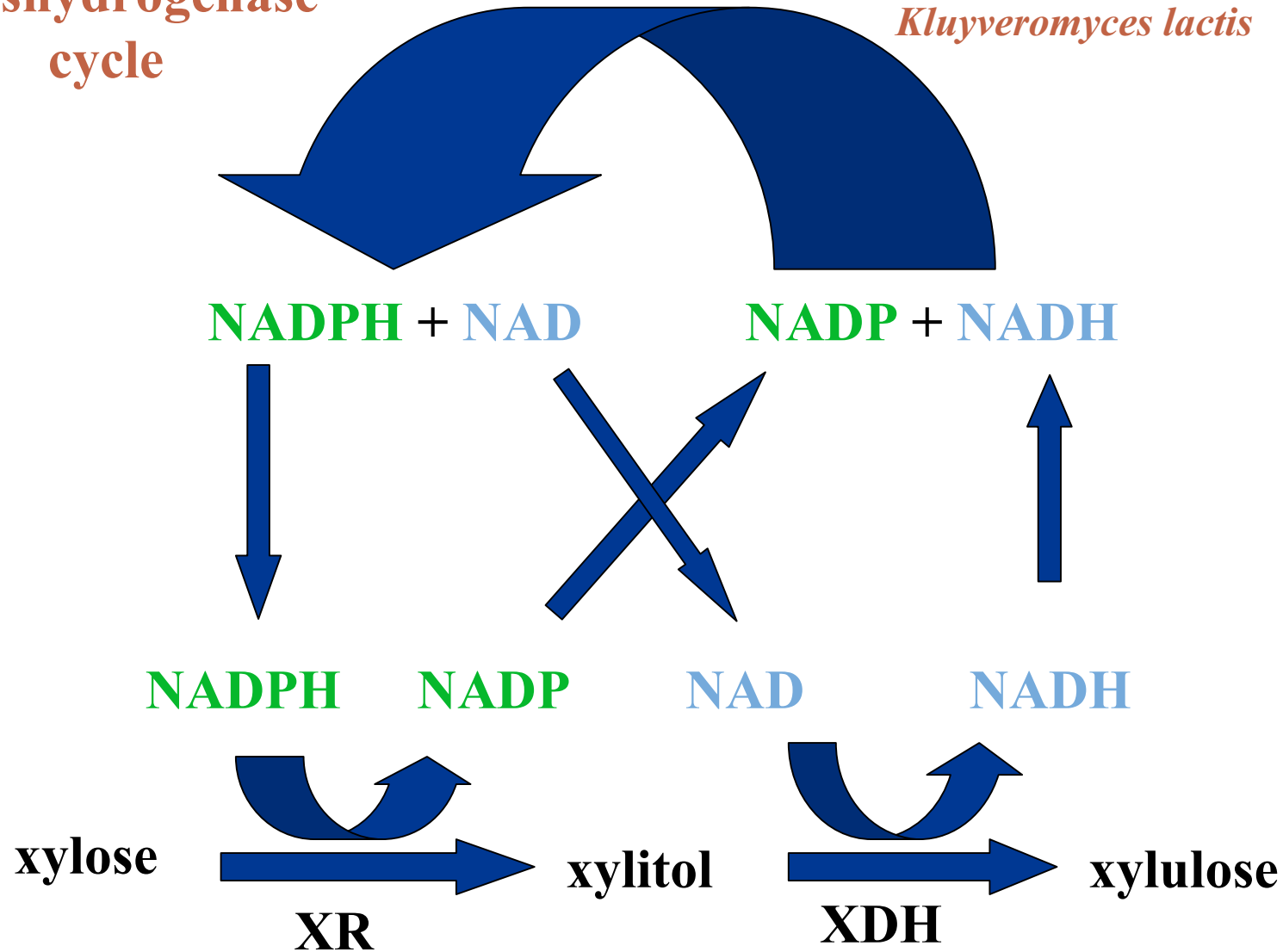
Cofactor imbalance in the xylose pathway



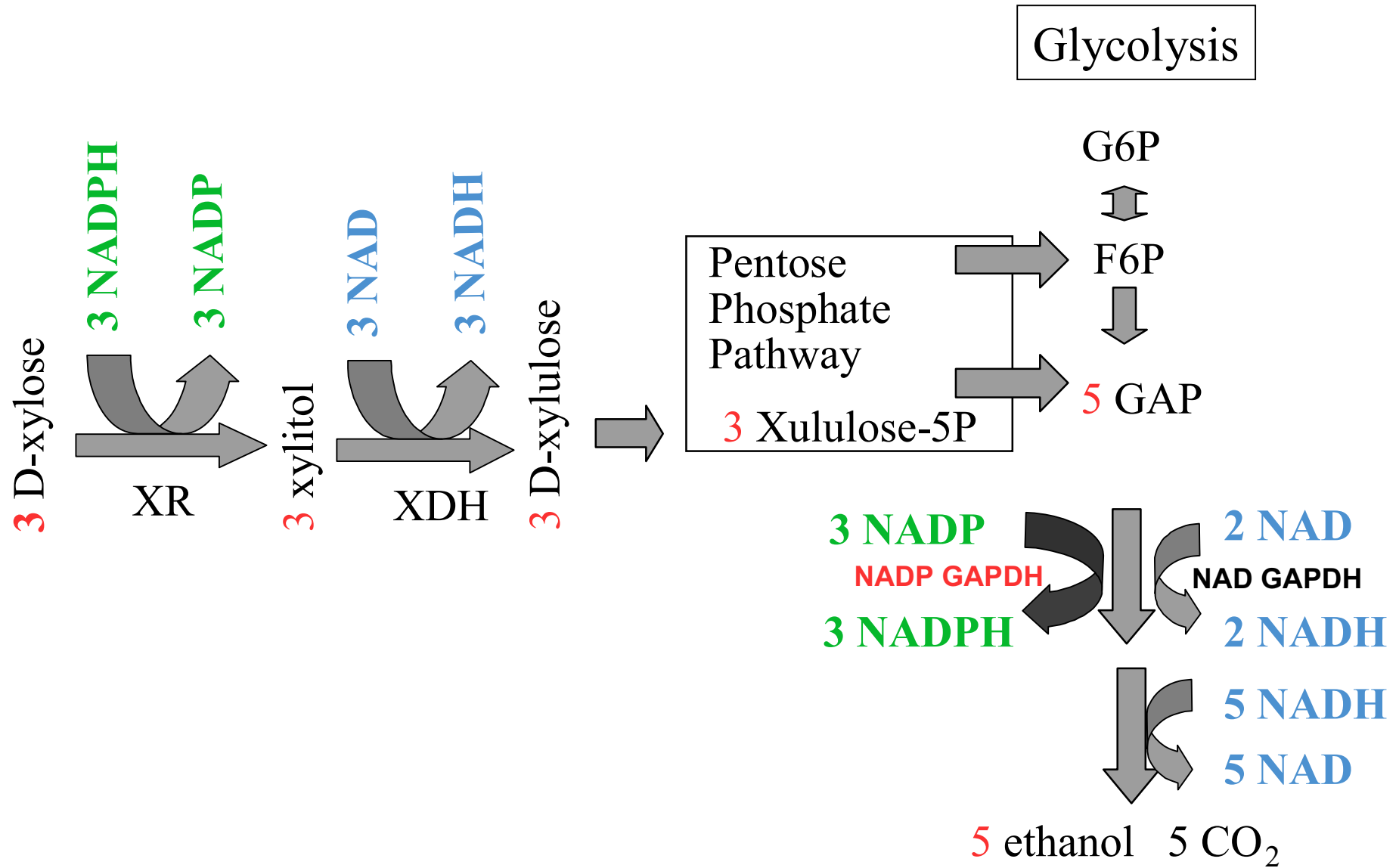
Engineering of cellular redox balances

- coexpression NAD & NADP-dep. GDH
- constitutive expression of malic enzyme
- expression of NADP-dep GAPDH of *Kluyveromyces lactis*

Transhydrogenase cycle



Cofactor balancing with the NADP dependent glyceraldehyde-P dehydrogenase

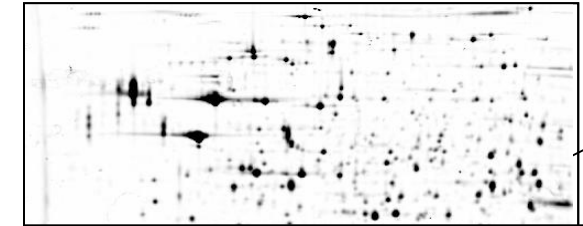
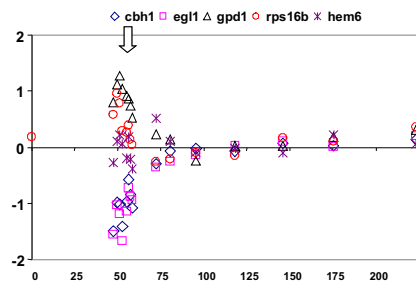


Significant reduction in xylitol/EtOH ratio

VTT Systems Biology Experimental platform

Transcriptomics

- commercial oligo-nucleotide arrays
- rapid at line analysis of selected transcripts by TRAC



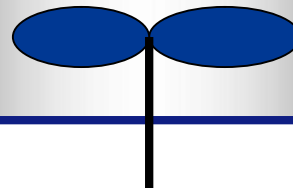
Proteomics

- 2D electrophoresis
- P-proteomics 2D + MS
- Mitochondrial and nuclear proteomics

Fermentation facilities

for batch, steady-state and transient experiments

- engineered strains
- specific culture conditions



Metabolic flux analysis

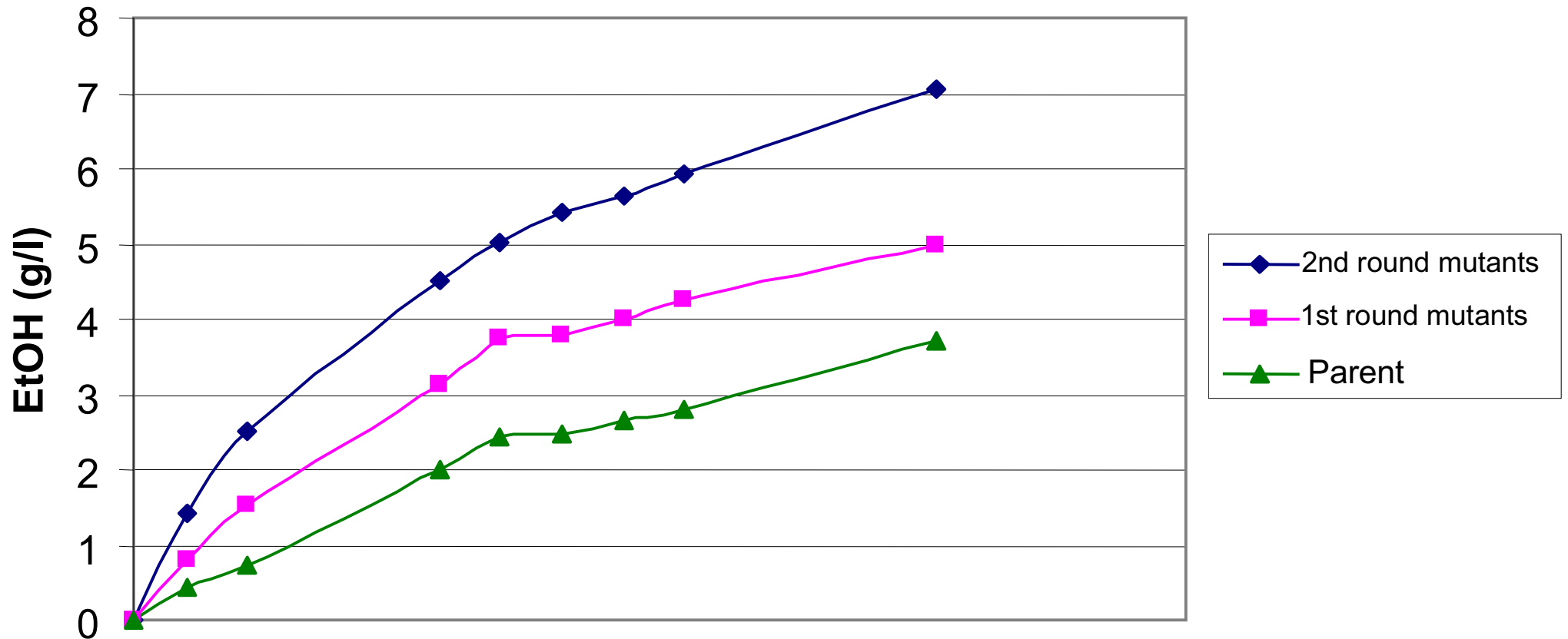
- metabolic flux analysis (MFA)
- ^{13}C tracer analysis (NMR, MS)

Metabolite analysis

- LC-MS: central C-metabolites
- GC-MS: organic & amino acids
- NMR with cryogenic probes

Data integration and modelling, identification of biomarkers

Classical mutagenesis is a useful tool in strain improvement



Ethanol production of mutants in anaerobic (glycerol lock flasks), cultures on xylose, mean values of several mutants

Xylose fermentation to bioethanol by recombinant *Saccharomyces cerevisiae*

XR-XDH pathway (e.g. VTT)

- ~ 70% yield of EtOH from xylose of the theoretical in anaerobic conditions (strain with various genetic modifications)
- Hydrolysate tolerant yeasts ferment biomass hydrolysates

XI pathway (e.g. Delft Univ.)

- XI from an anaerobic fungus *Piromyces*
- Close to theoretical yields of ethanol (strain with various genetic modifications)

Rates of xylose fermentation still need improvement

“2nd generation” raw materials under study

Whole crop approach

- Straw, hulls
- Corn stover, cobs
- Wood chips, bark

Industrial process sidestreams

- Wheat bran, spent grain
- Bagasse
- Beet pulp
- Food industry waste (peels etc, oil industry waste)
- P&P waste streams
- Recycled paper

Dedicated energy plants

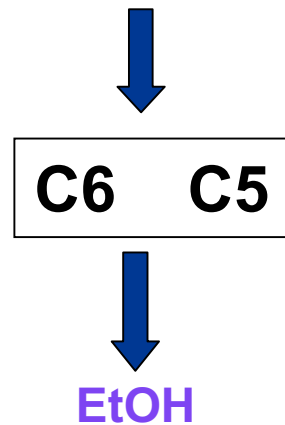
- *Miscanthus*
- *Sorghum*
- *Reed canary grass*

Etc



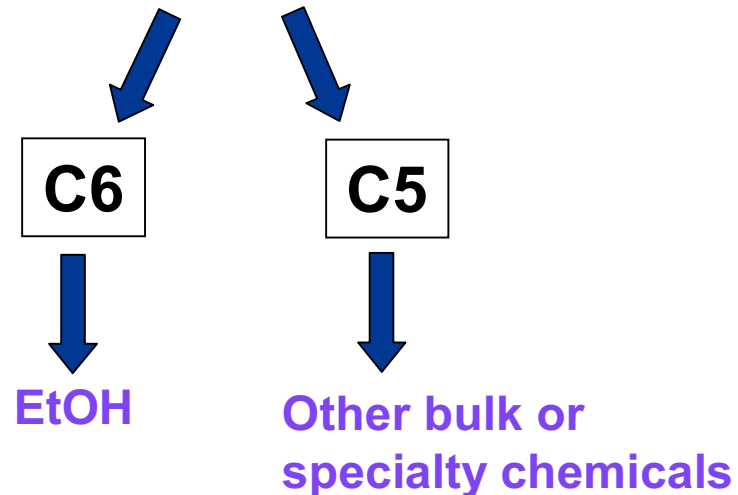
Process options

Biomass



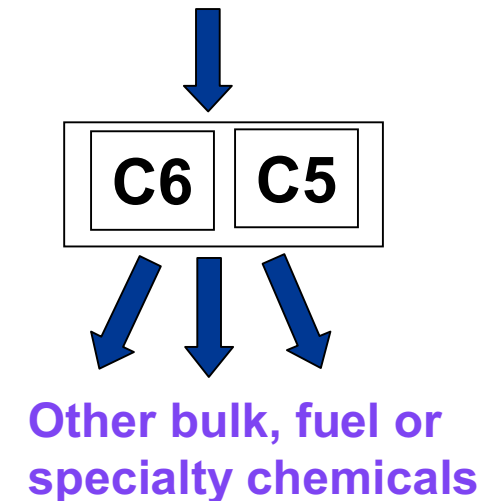
- efficient co-fermentation needed
- high yield of bioethanol per biomass
- in principle a simple process

Biomass



- separation of sugar fractions needed
- economics of bioethanol biorefinery can be improved
- novel product options

Biomass

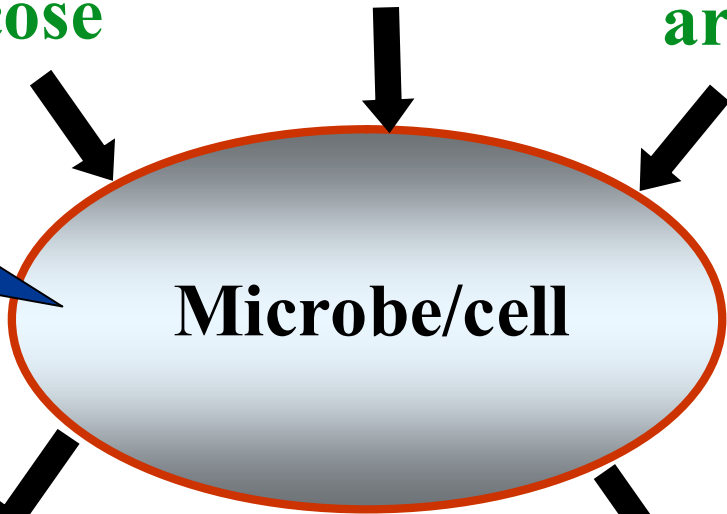


- infrastructure and biomass treatment methods available
- bioethanol process easily convertible to other products
- flexible product options

Corn **Municipal waste** **Lignocellulosics**
 Plant waste

glucose **xylose** **arabinose**

A cheap raw material can be converted to a valuable product through cellular metabolism



Microbe/cell

Fine chemicals

vitamins
 xylitol
 amino acids

Bioactive compounds

antibiotics

Bioethanol
Butanol
Biodiesel

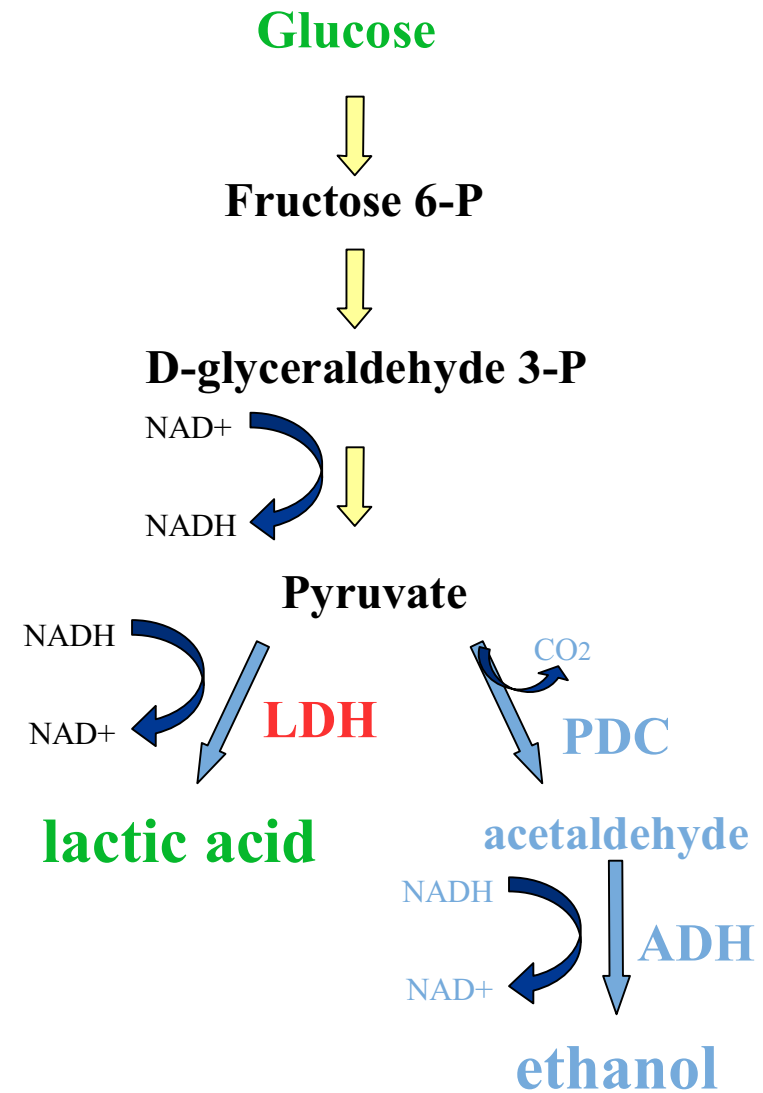
Polymer-precursors

butyric acid comp. -> PHB
 lactic acid -> PLA
 1,3-propanediol -> polyesters

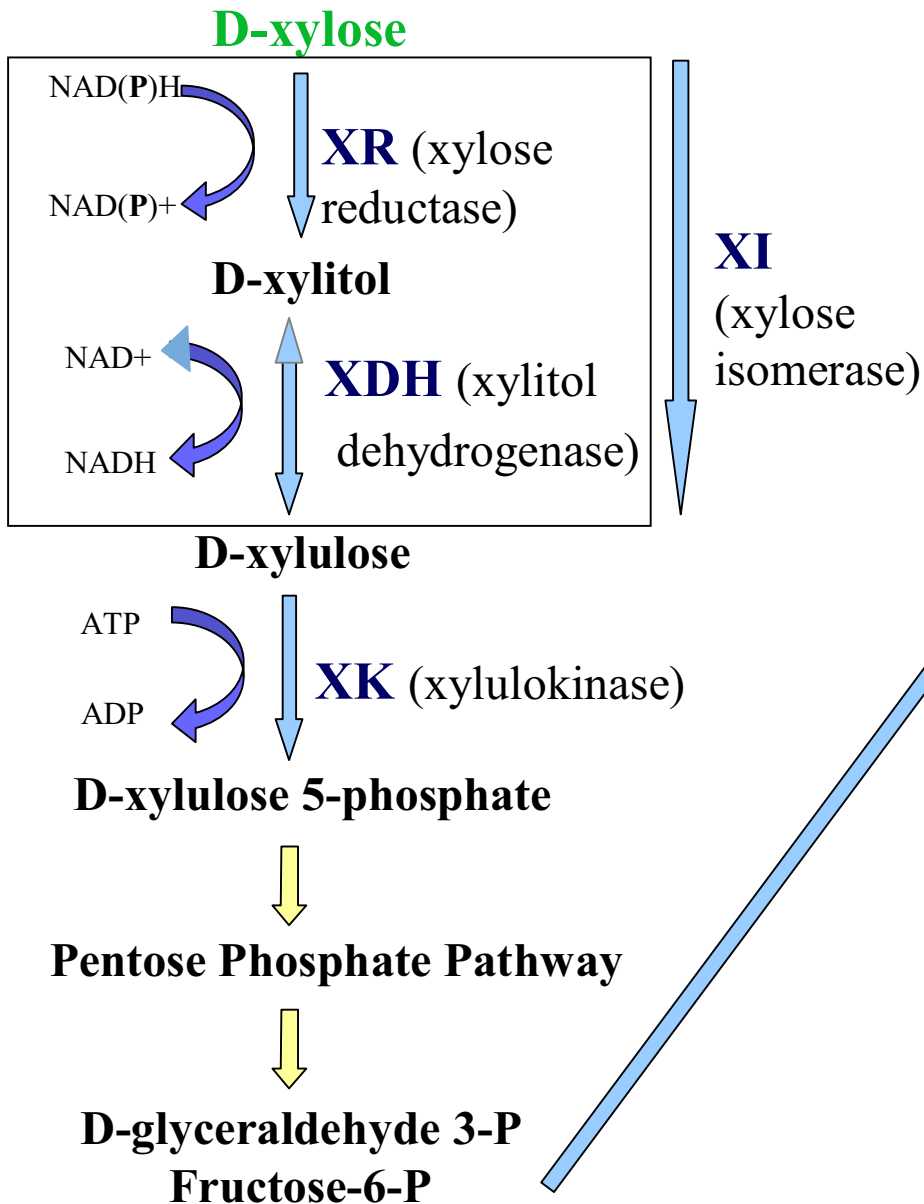
Production of lactic acid, PLA monomer

Lactic acid production from glucose

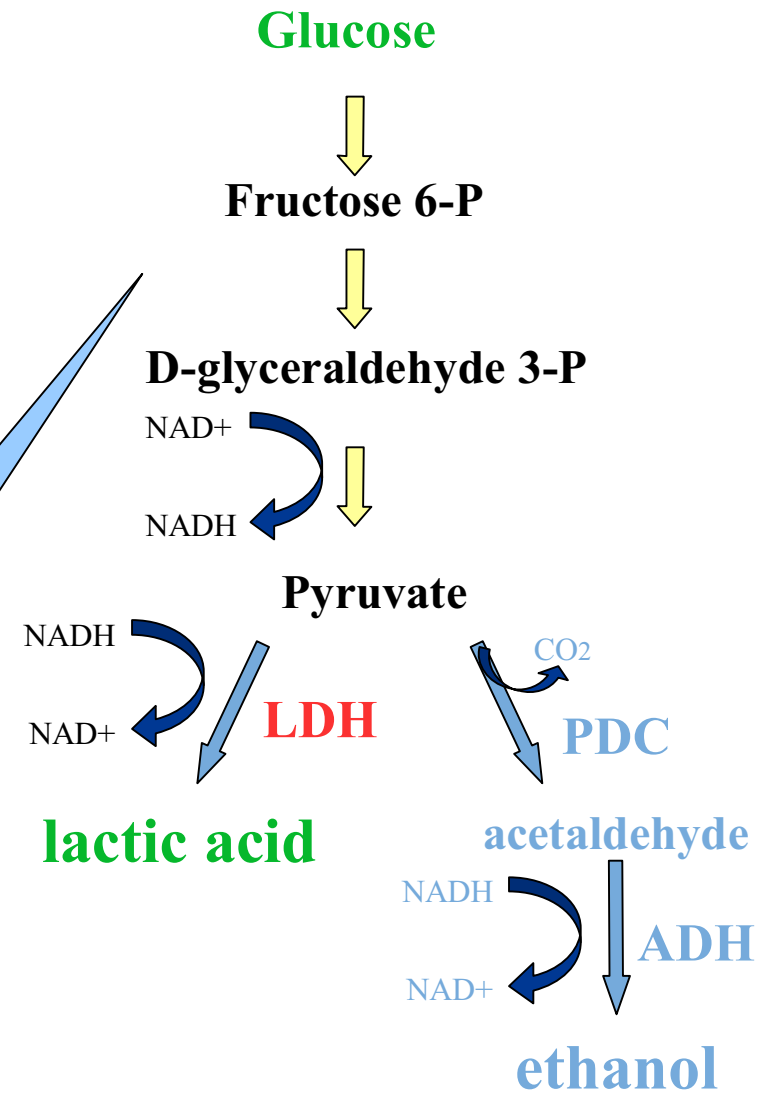
- introduction of bacterial lactate dehydrogenase (LDH)



Pentose utilisation



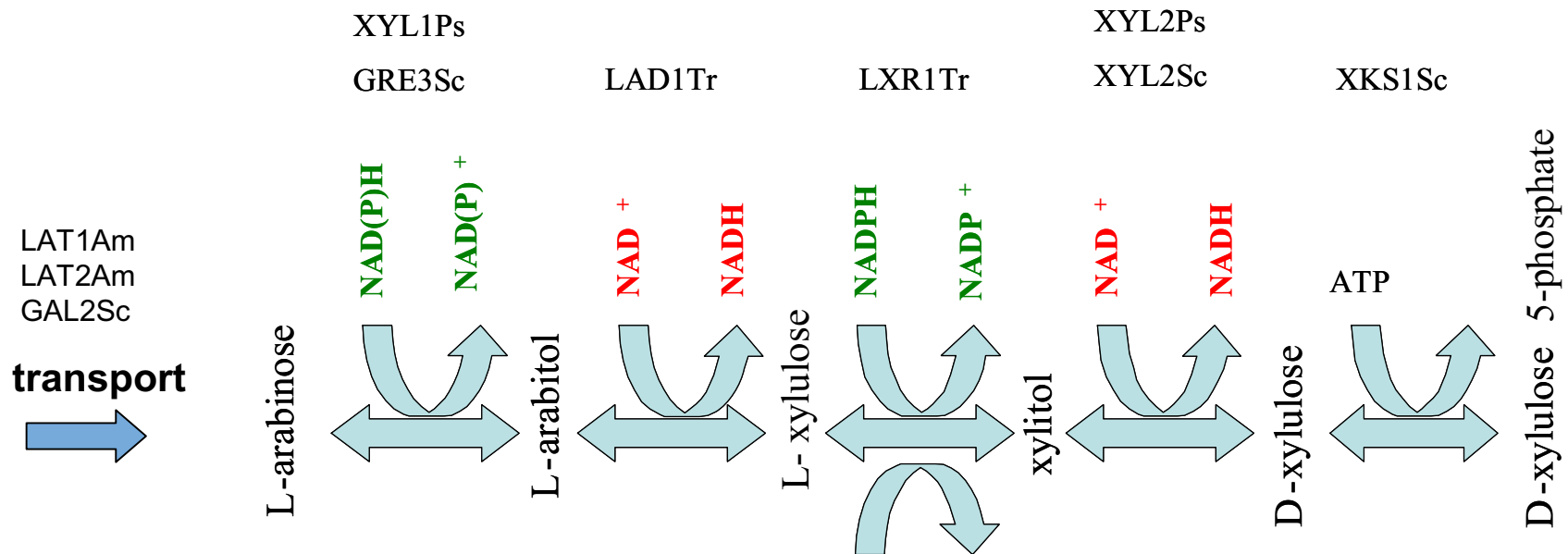
Glycolysis



VTT has also characterised the fungal pathways for

- L-arabinose utilisation
- D-galacturonic acid utilisation

and expressed these pathways in *S.cerevisiae*

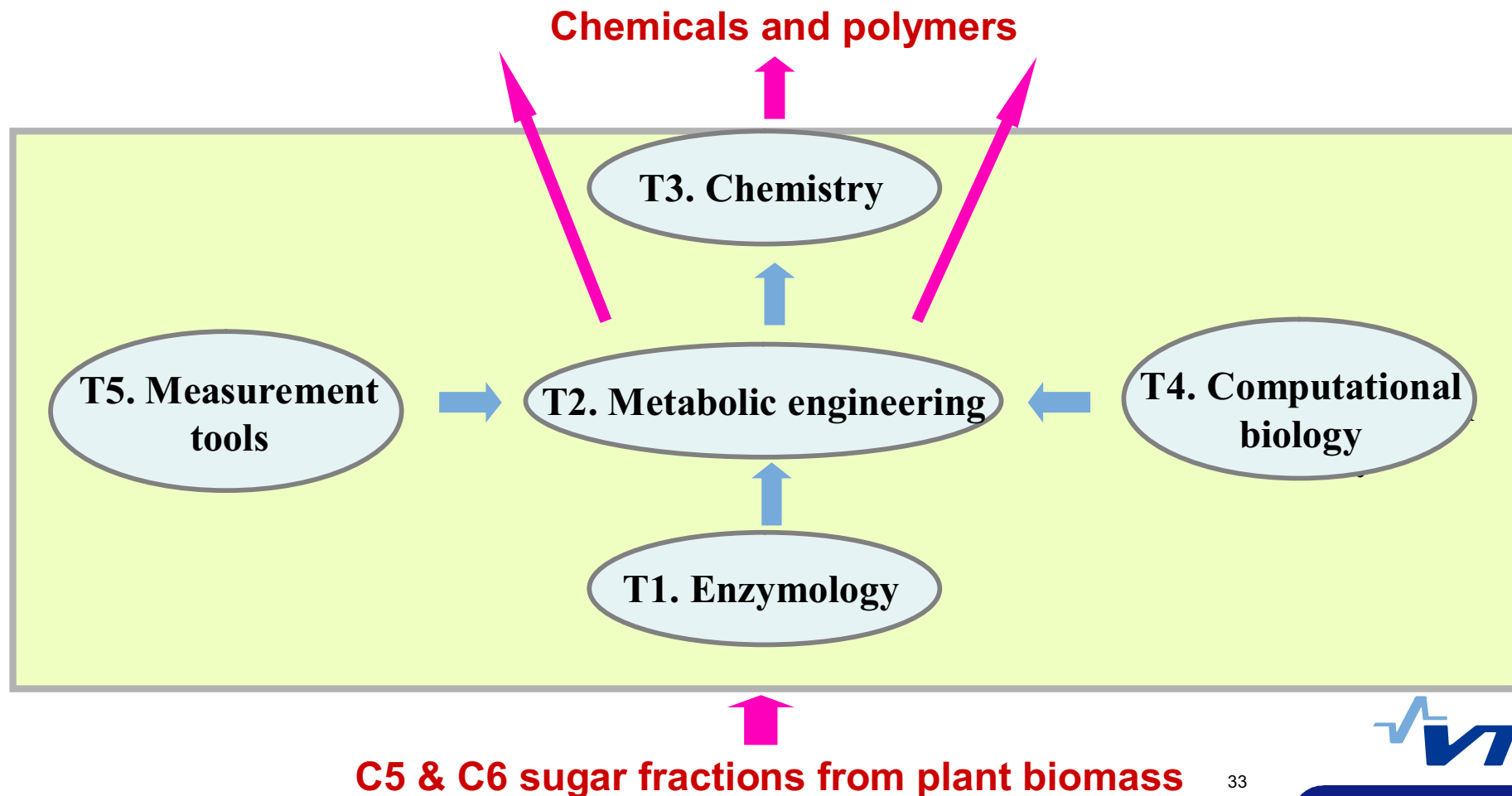


The fungal pathway for L-arabinose utilisation with the cofactors involved. The genes studied are shown above and below the pathway with their origin indicated (Ps= *P. stipitis*, Sc=*S. cerevisiae*, Tr=*T. reesei*, Am=*Ambrosiozyma monospora*).

Centre of Excellence status

by the basic science funding body Academy of Finland

2008-2013 in White Biotechnology – Green Chemistry



Issues

- Pretreatment methods under rapid development
 - Reduced enzyme requirements, consolidated processes
 - Reduced toxicity
 - Separation of sugar streams
 - Feedstock development
 - Reduced lignin, “self-hydrolysis”
 - Altered sugar composition
 - New product options
 - Essential for biorefinery concepts and their economy
 - Market need & volumes need to be considered
 - New challenges for metabolic engineering
- Organisms robustness & performance remain always key issues
 - Process and end product tolerance
 - Rate and yield of production