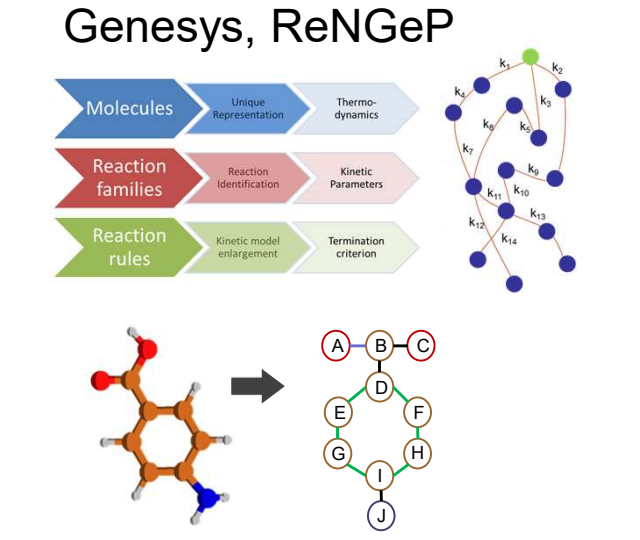
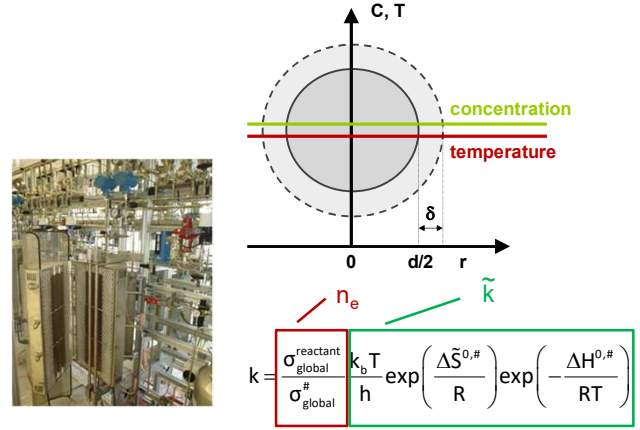
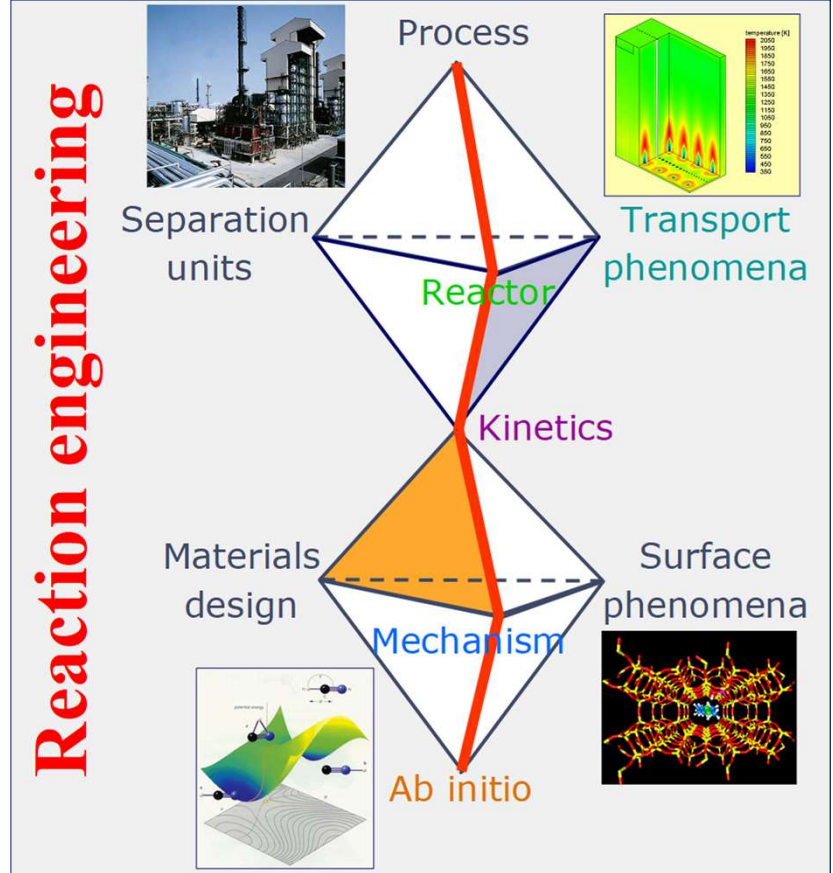
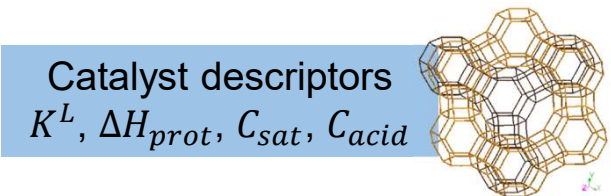
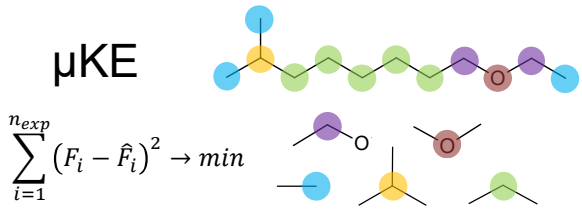
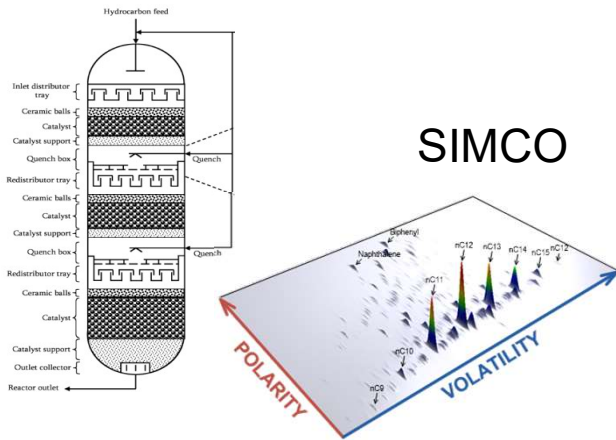


Large-scale production of fuels and chemicals:  
coping with complex feedstocks and chemistry

Guy B. Marin, Joris W. Thybaut and Kevin M. Van Geem  
Laboratory for Chemical Technology



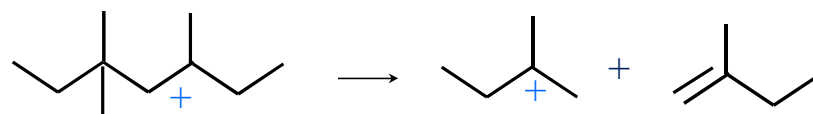
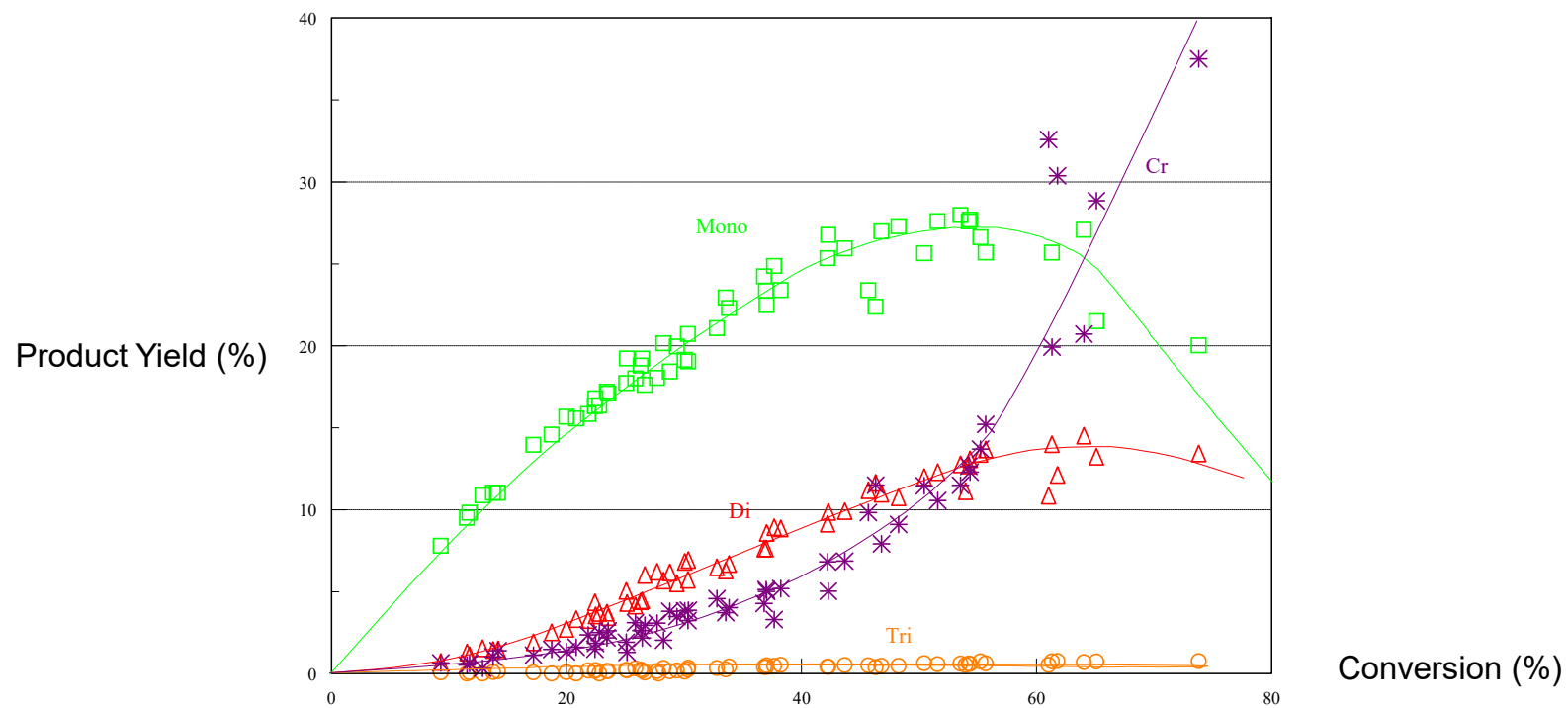
# Kinetics: methods and applications



# Outline

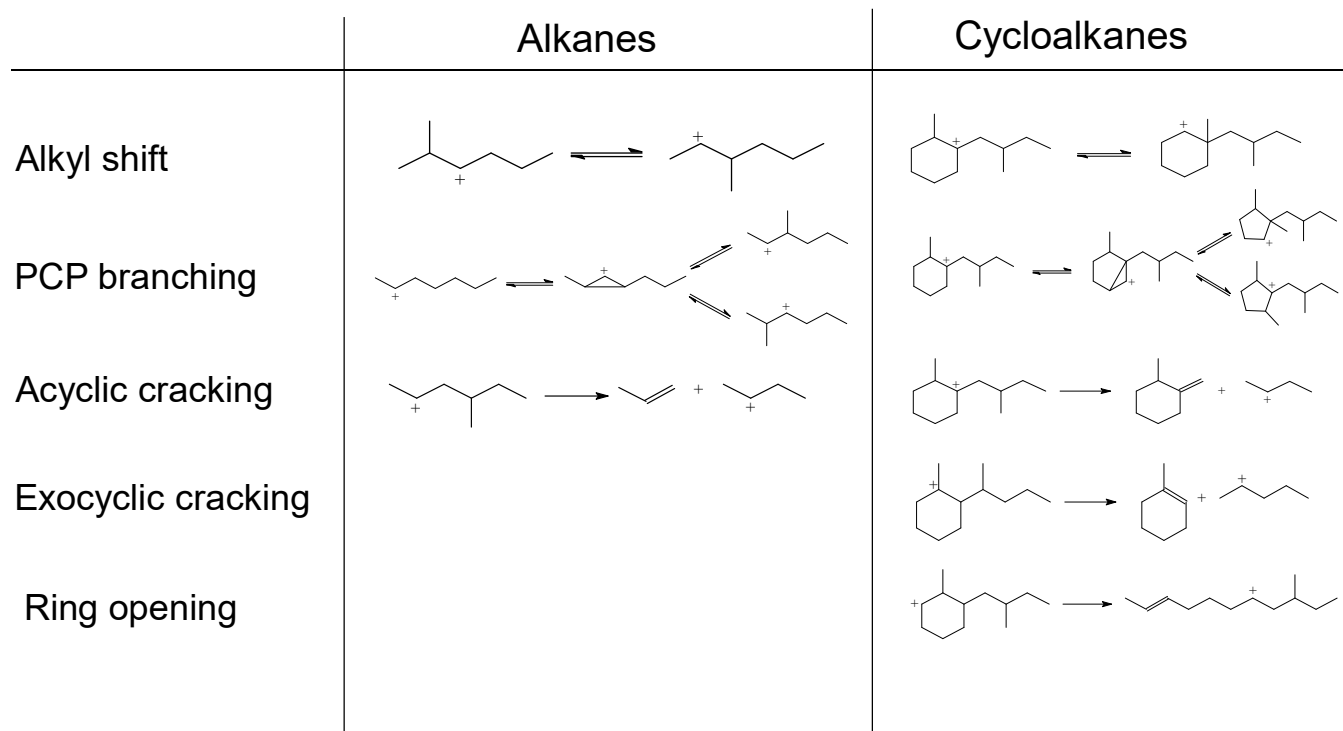
- Introduction
- **Complex feedstocks: ideal hydroconversion**
- Feedstock characterization and reconstruction
- Reaction-network generation
- Complex chemistry: parameter determination
- Reactor design
- Conclusions

# Hydrocracking and isomerization: reaction families

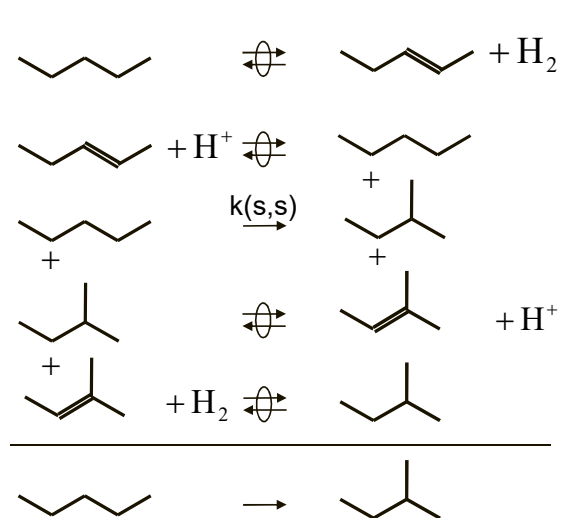


(t,t) β-scission

# Free carbenium ion reaction families: kinetic descriptors



# Kinetic versus Catalyst descriptors



Strength of acid sites

$$\Delta H_{\text{prot}}$$

Concentration of acid sites

$$r = \frac{C_{\text{sat}} C_t k K_{\text{prot}} K_{\text{deh}} K_L p_P p_{H_2}^{-1}}{1 + K_L p_P}$$

$$k = \frac{\sigma_{gl,r}}{\sigma_{gl,\ddagger}} \frac{k_B T}{h} \exp\left(\frac{\Delta \tilde{S}^{0,\ddagger}}{R}\right) \exp\left(\frac{\Delta H^{0,\ddagger}}{RT}\right)$$

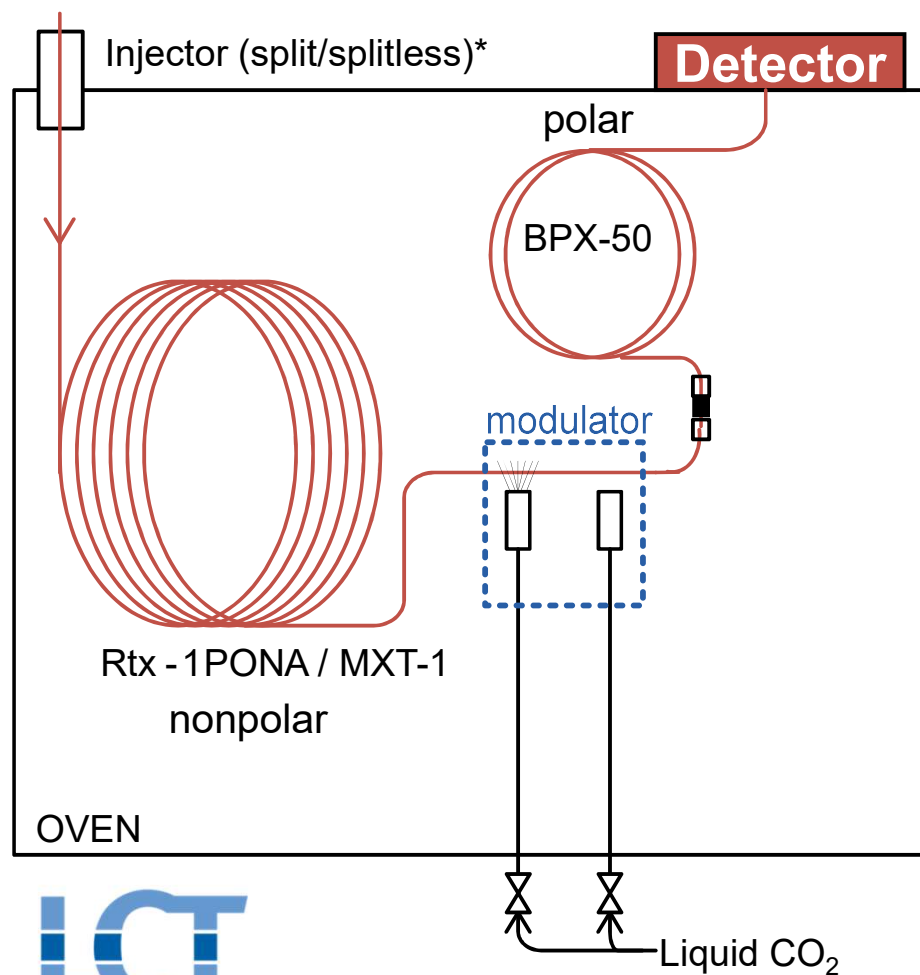
determined by the reaction network generation code

constant for a reaction family (chain length independent)

# Outline

- Introduction
- Complex feedstocks: ideal hydroconversion
- **Feedstock characterization** and reconstruction
- Reaction-network generation
- Complex chemistry: parameter determination
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# GC×GC-FID/TOF-MS/NCD/SCD setups



Normal Phase: nonpolar × polar  
Reversed Phase: polar × nonpolar

## Detectors:

250 kEuro

## TOF-MS:

➤ Qualitative analysis

## NCD:

25 kEuro

➤ Quantitative analysis

➤ Selective to N- compounds

## SCD:

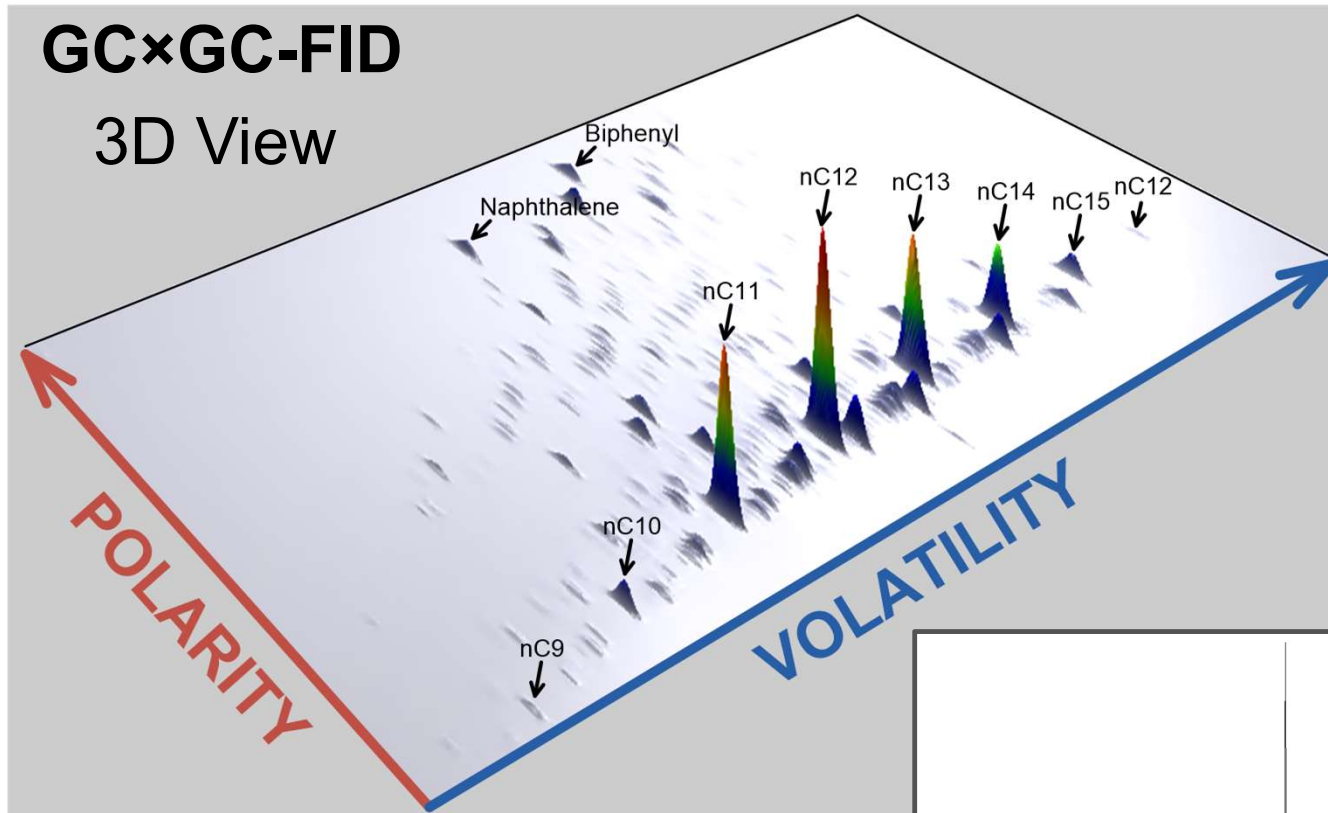
25 kEuro

• Quantitative analysis

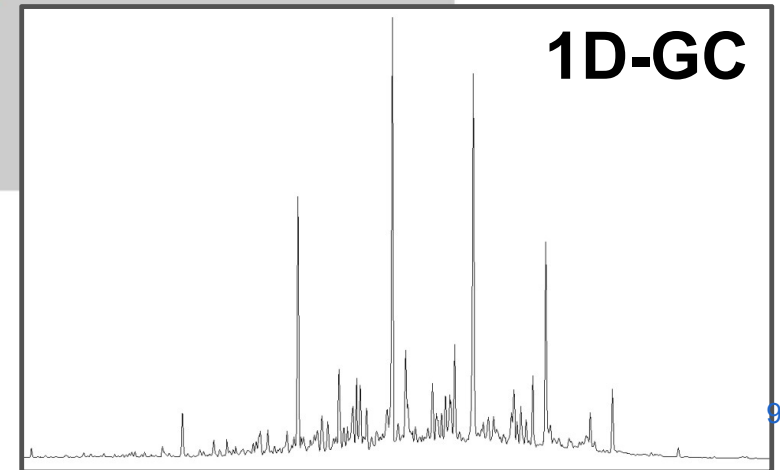
• Selective to S- compounds



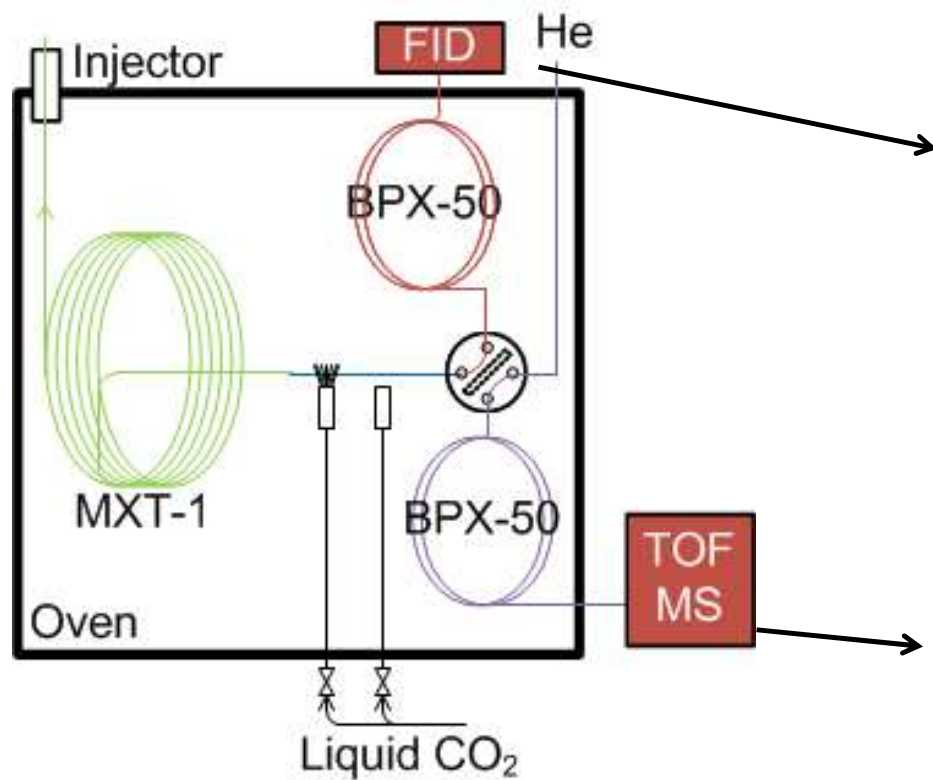
# Example: Middle Distillate Analysis



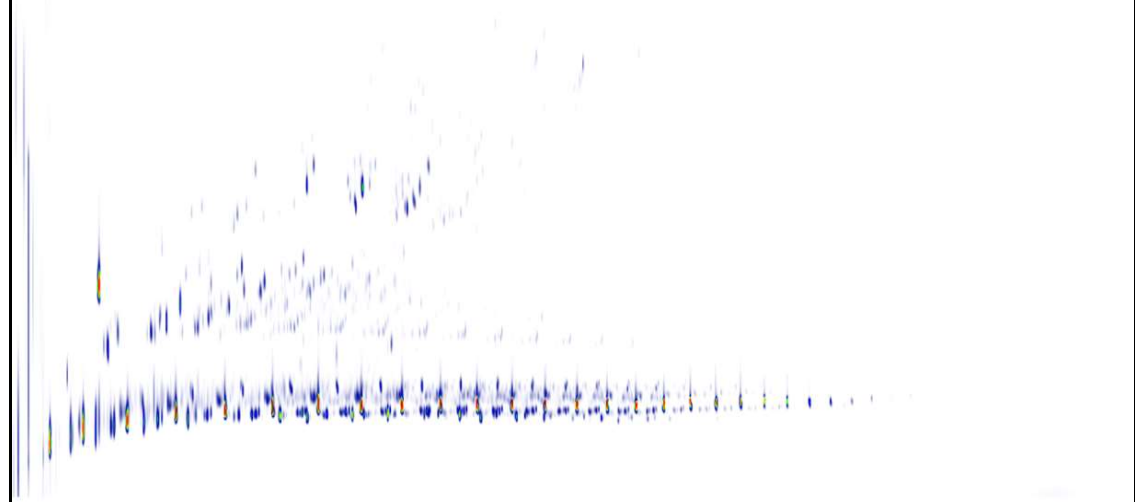
2 independent separation mechanisms  
→ Enhanced resolution compared to 1D-GC



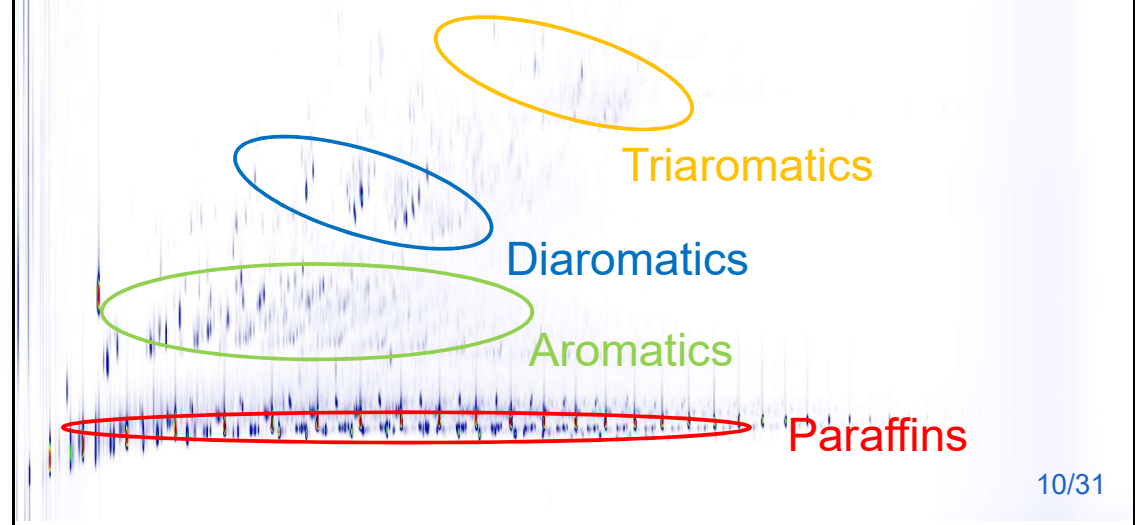
# GC × GC-FID/TOF-MS



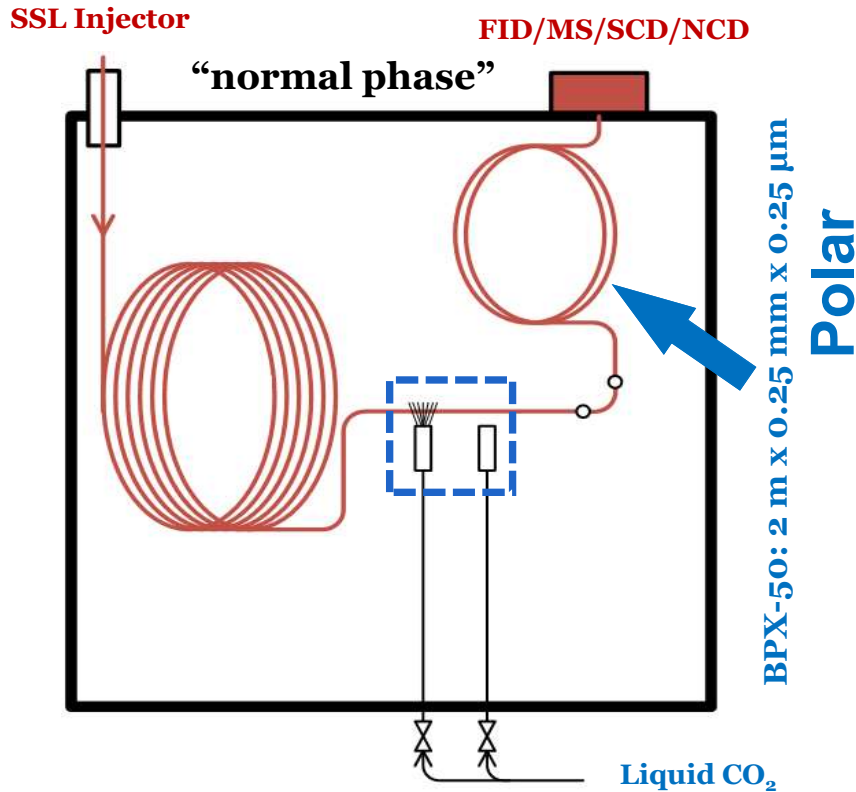
Quantitative but no direct identification possible



Qualitative. difficult to quantify

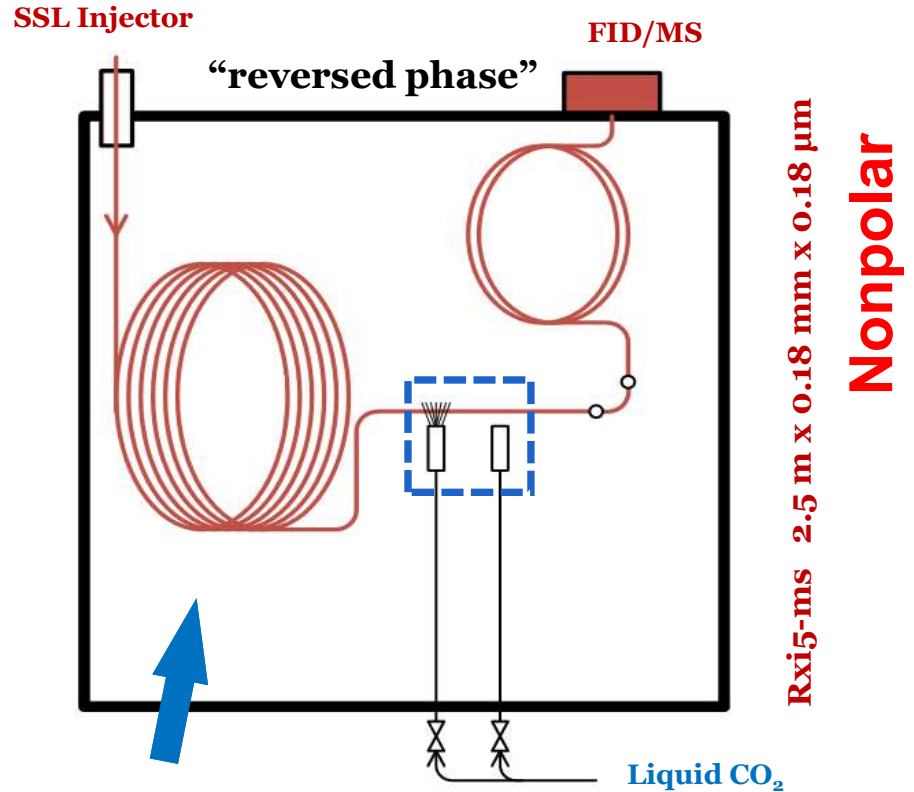


# GC × GC column configuration for Shale Oil Analysis



PONA: 30m x 0.25 mm x 0.25  $\mu$ m

**Nonpolar**



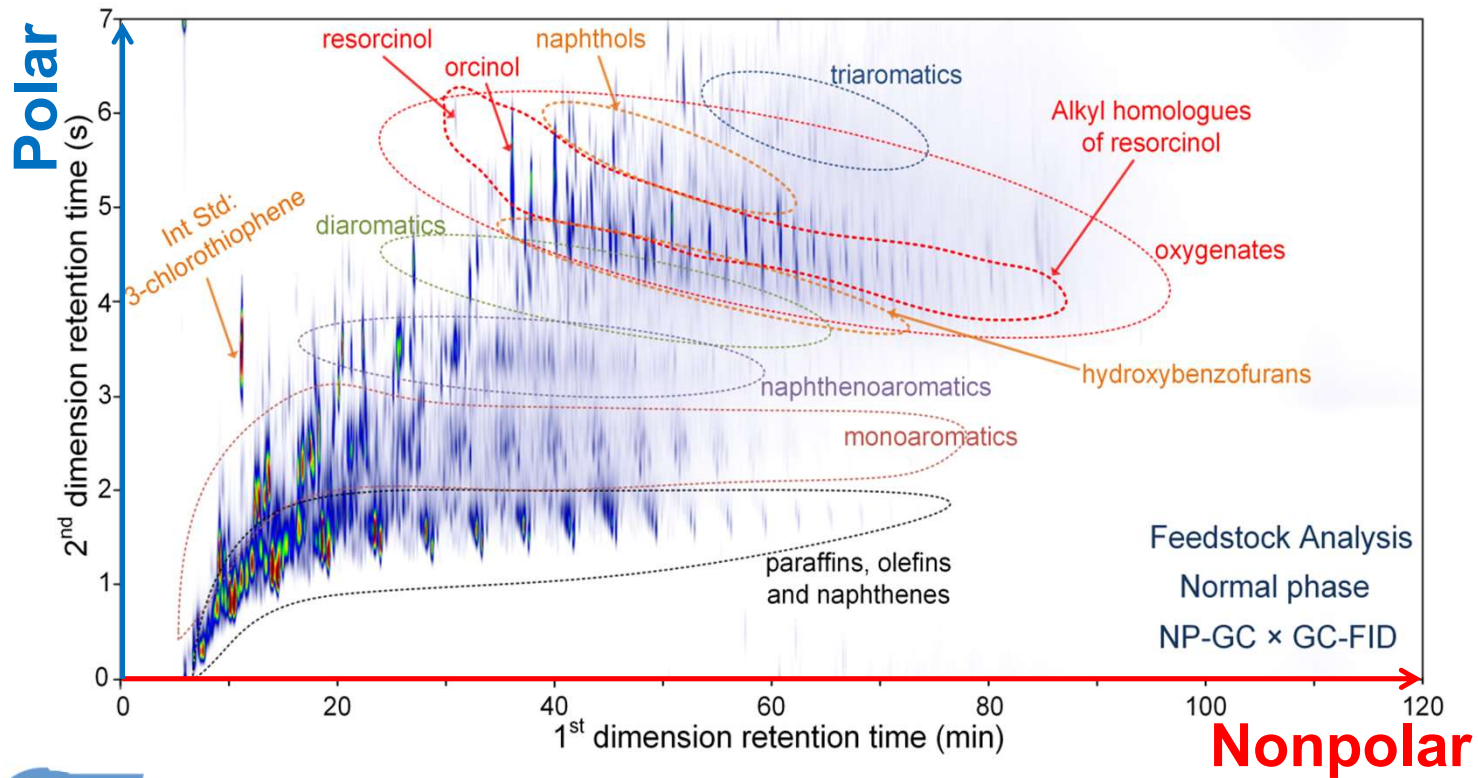
1. Stabilwax; 60 m x 0.25 mm x 0.25  $\mu$ m (240 °C)

2. MegaWax; 30 m x 0.25 mm x 0.15  $\mu$ m (300 °C)

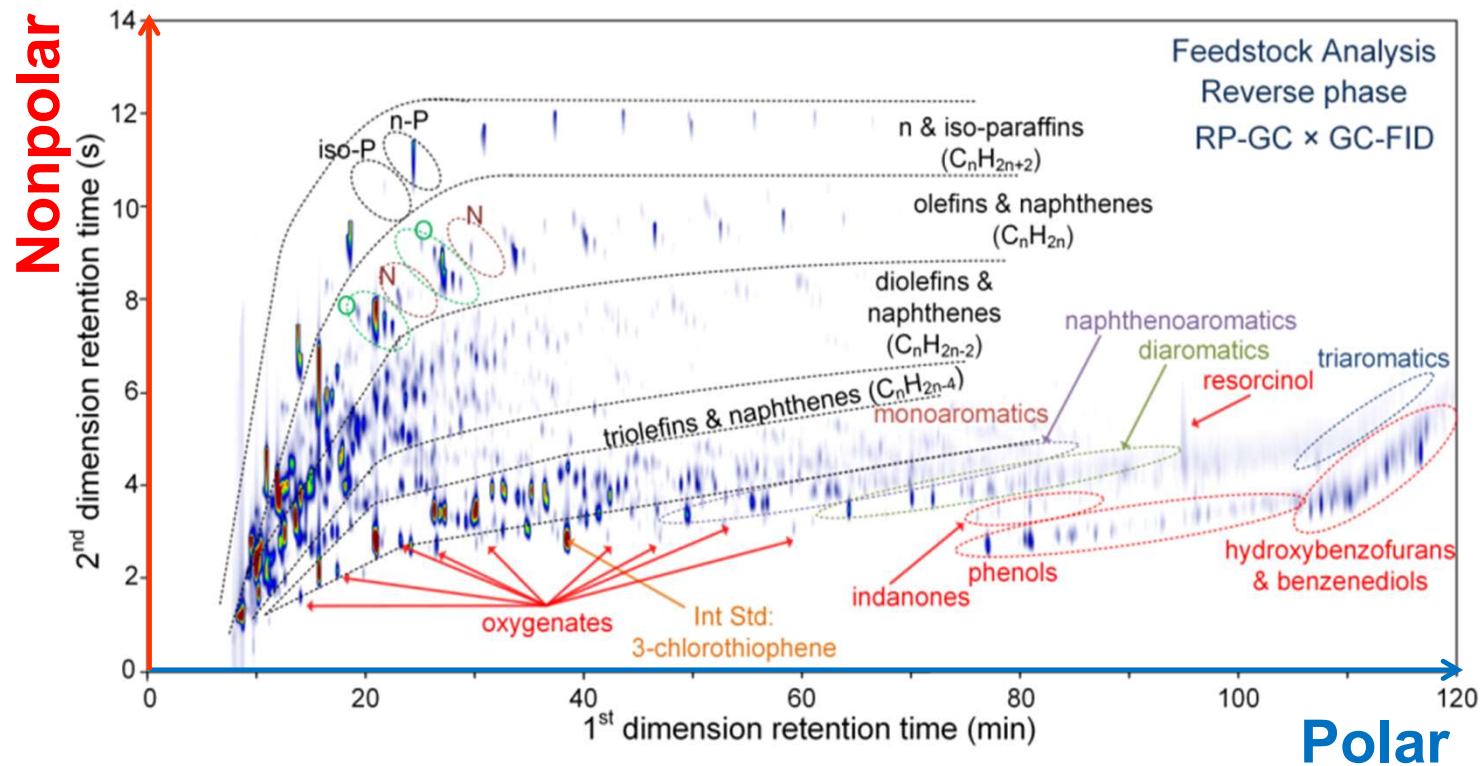
**Polar**

# Shale Oil: NP-GC × GC –FID/TOF-MS

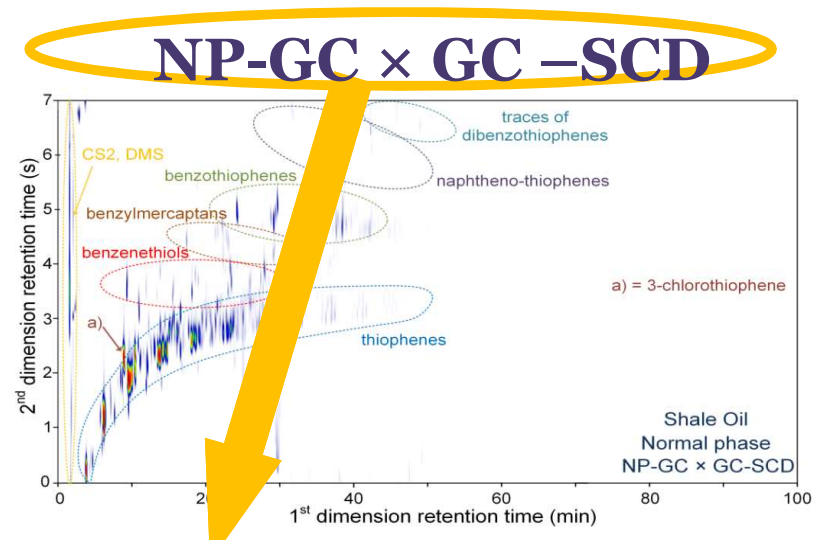
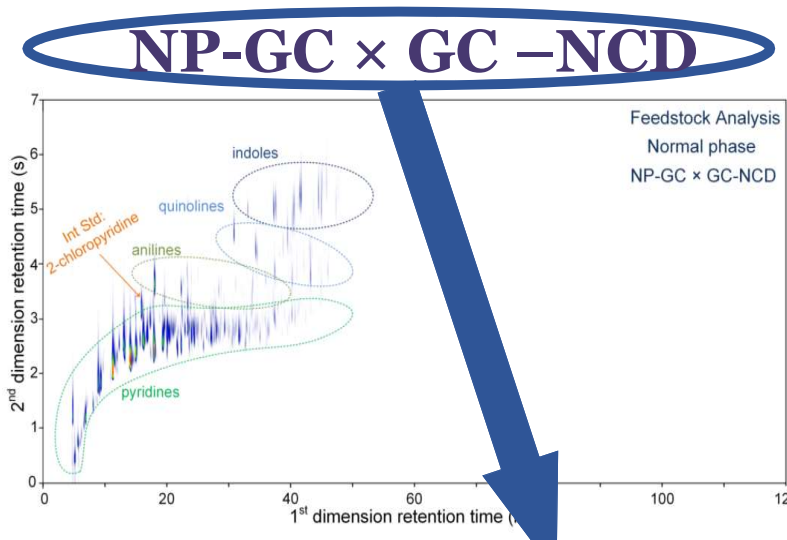
Estonian Pyrolysis Shale Oil (~ 6 wt% of Oxygen)



# Shale Oil: RP-GC × GC –FID



# Shale Oil: NP-GC × GC –NCD/SCD



Measured **elemental nitrogen** using:  
NP-GC × GC –NCD  
is **0.03 wt.%**  
EA is **0.03 wt.%**

Nitrogen containing chemical classes	Sulfur containing chemical classes
<ul style="list-style-type: none"> <li>Pyridines</li> <li>Anilines</li> <li>Quinolines</li> <li>Indoles</li> </ul>	<ul style="list-style-type: none"> <li>Thiols/Sulfides</li> <li>Thiophenes</li> <li>Benzenethiols</li> <li>Benzylmercaptans</li> <li>Benzothiophenes</li> <li>Naphthenobenzothiophenes</li> <li>Dibenzothiophenes</li> </ul>

Measured **elemental sulfur** using:  
NP-GC × GC –SCD  
is **0.68 wt.%**  
EA is **0.7 wt.%**

# Outline

- Introduction
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# Macroscopic Properties

HPLC  
• PIONA  
• SARA



Distillation



- ✓ ASTM D86
- ✓ ASTM D1160

Sim Dist



- ✓ ASTM D2887
- ✓ ASTM D7169



CHNS-O  
Elemental Analysis



- ✓ ASTM D5291
- ✓ ASTM D5622
- ✓ Total sulfur using FPD

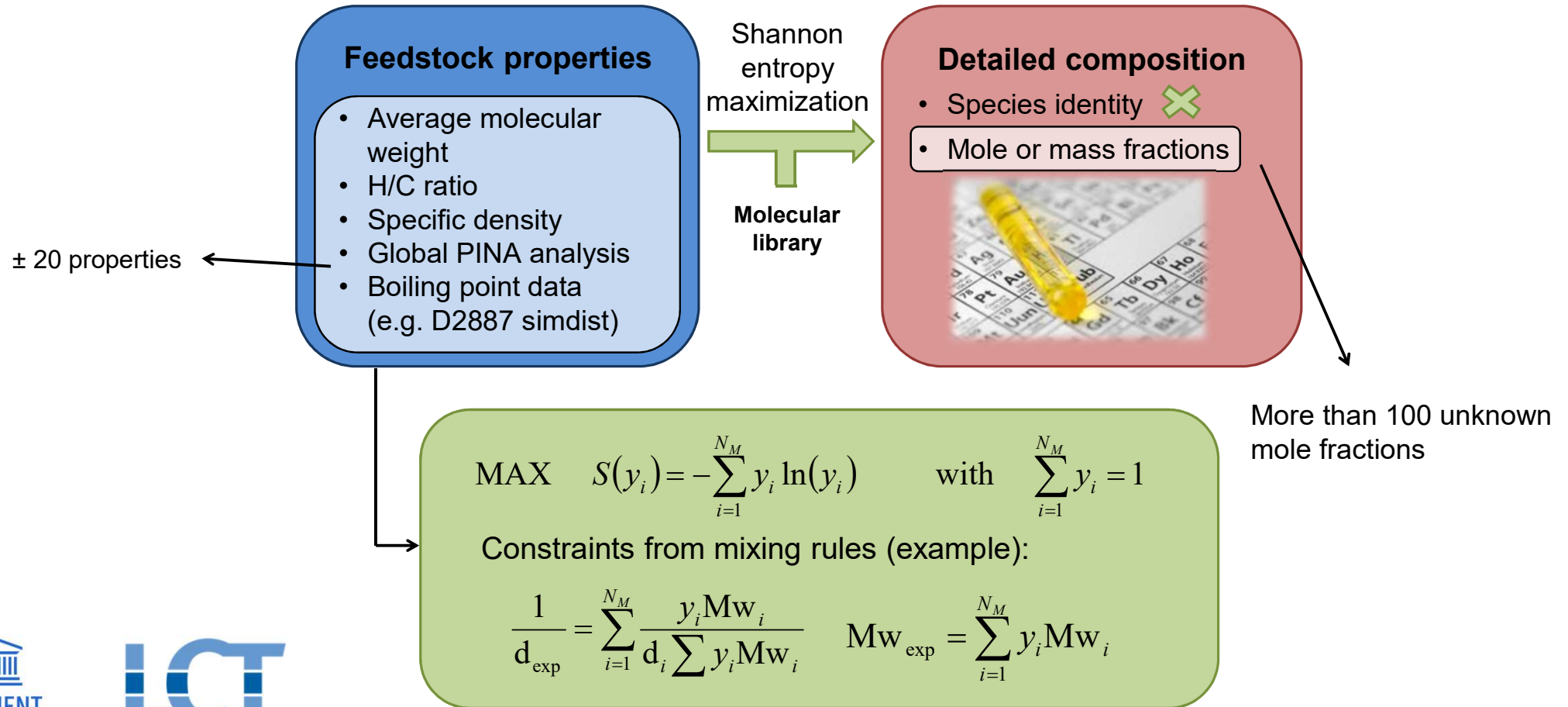
Density



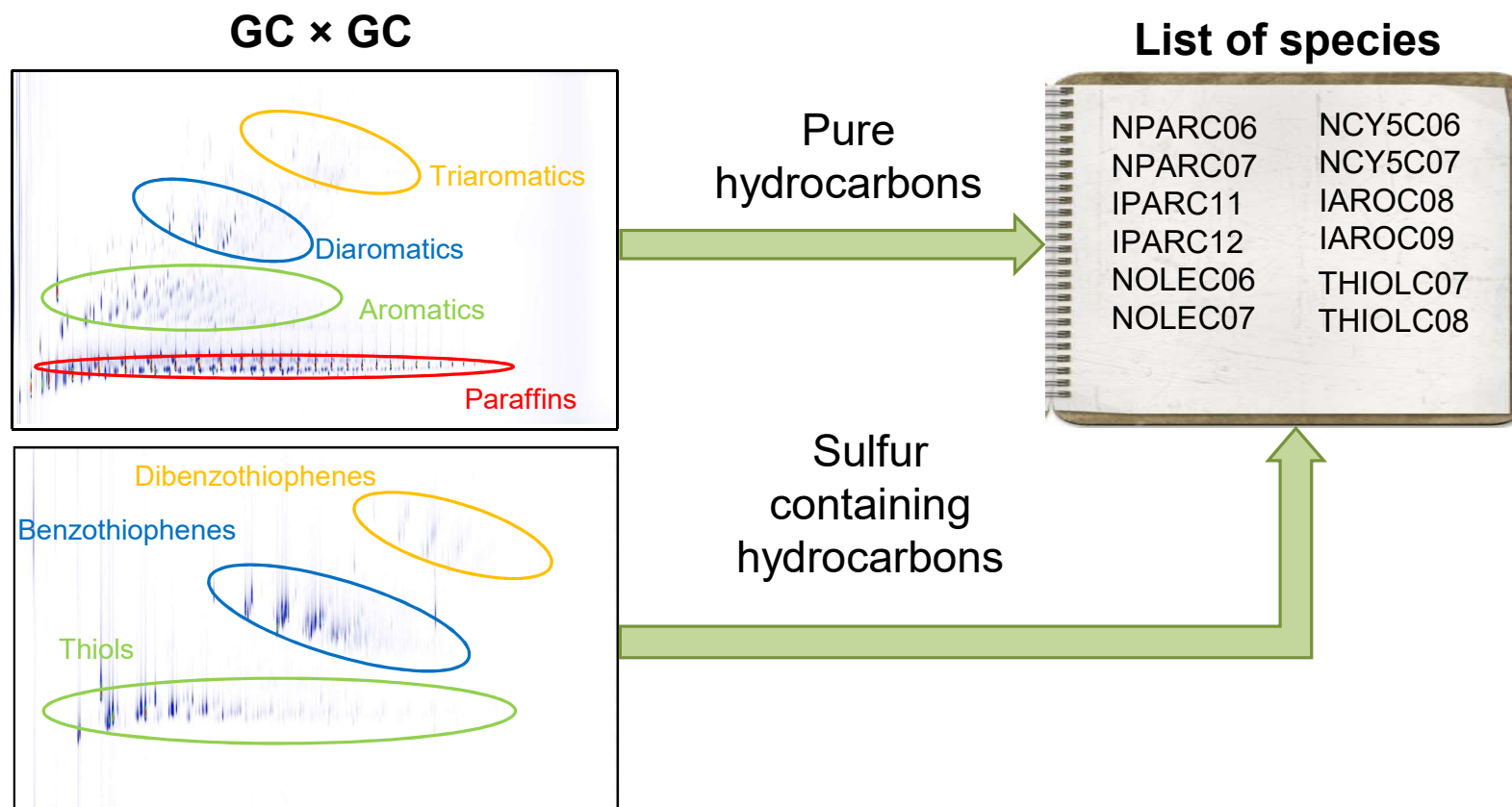
Access to: Average MW  
 $^1\text{H}$  and  $^{13}\text{C}$  NMR data ...



# Shannon entropy maximization



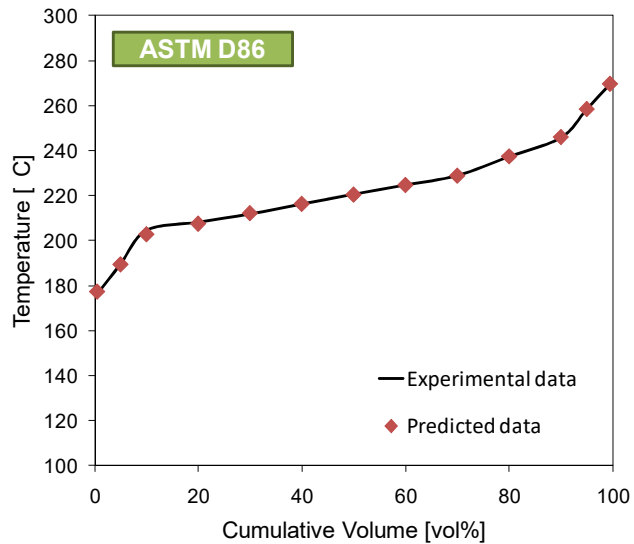
# Molecular library: considered species



# Example: Middle distillates

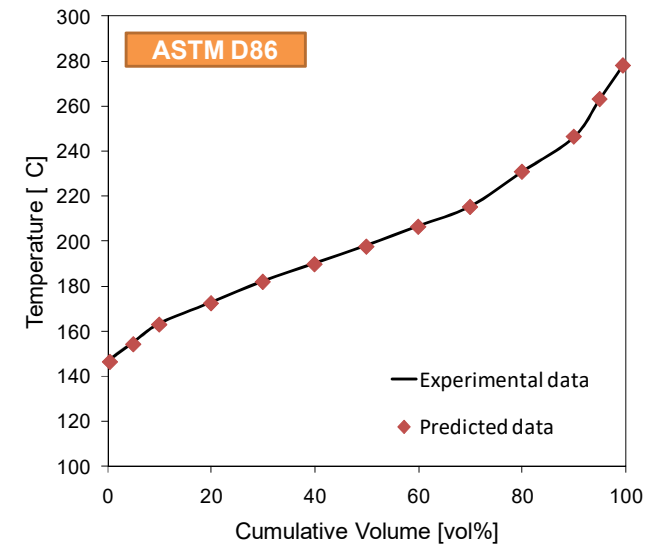
## SAMPLE 1: Middle East

		INPUT	CALC.
<b>d</b>	(g/ml)	0.7983	0.7983
<b>P</b>	(wt%)	23.459	23.458
<b>I</b>	(wt%)	25.370	25.368
<b>O</b>	(wt%)	0.000	0.000
<b>N</b>	(wt%)	34.074	34.078
<b>A</b>	(wt%)	17.097	17.097



## SAMPLE 2: Brazil

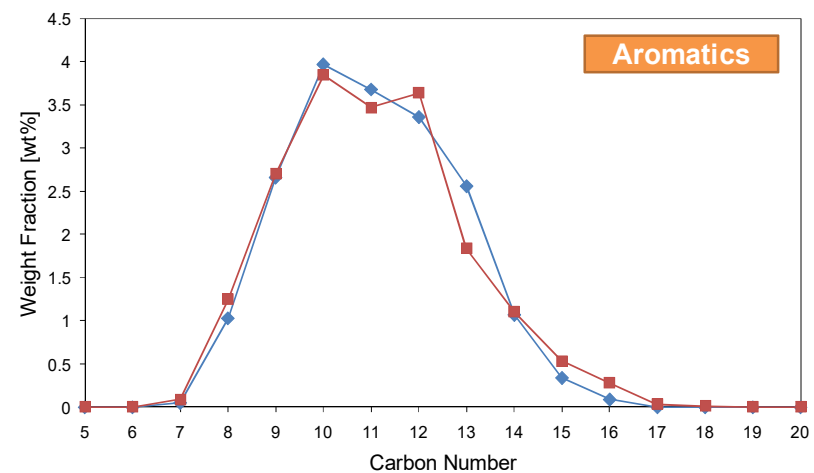
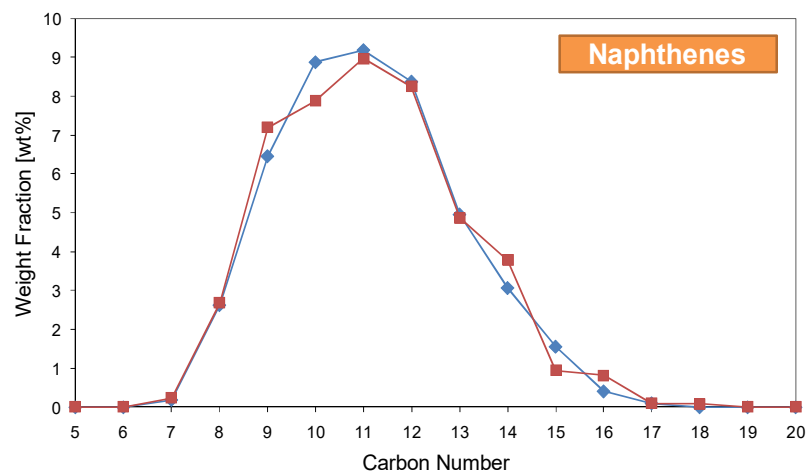
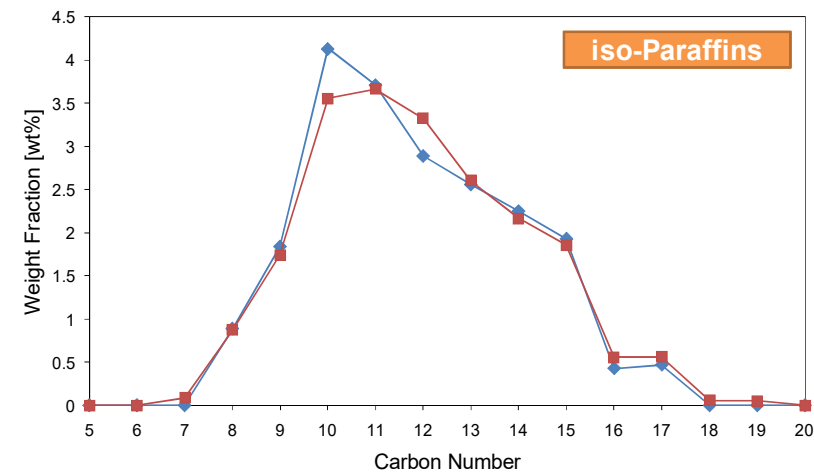
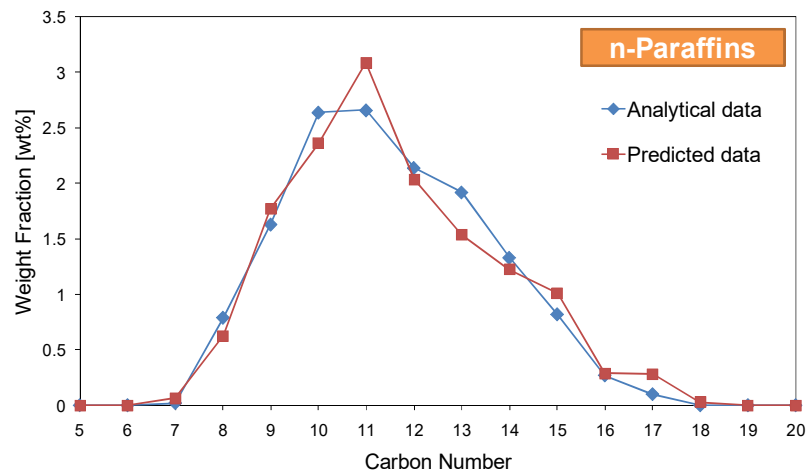
		INPUT	CALC.
<b>d</b>	(g/ml)	0.7987	0.7986
<b>P</b>	(wt%)	14.311	14.305
<b>I</b>	(wt%)	21.087	21.087
<b>O</b>	(wt%)	0.000	0.000
<b>N</b>	(wt%)	45.803	45.812
<b>A</b>	(wt%)	18.799	18.797



# Example: Middle distillates

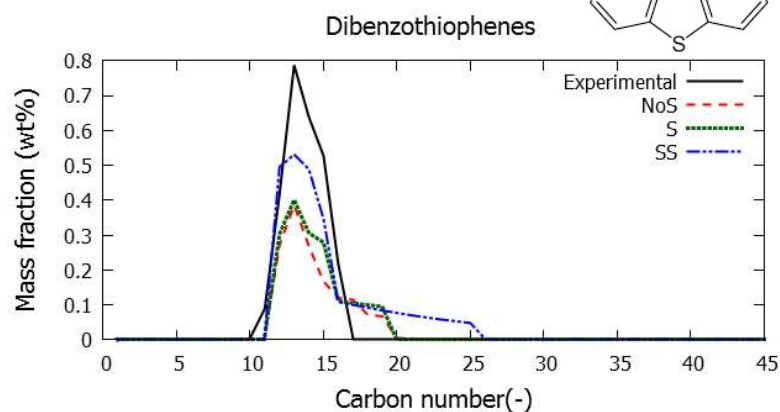
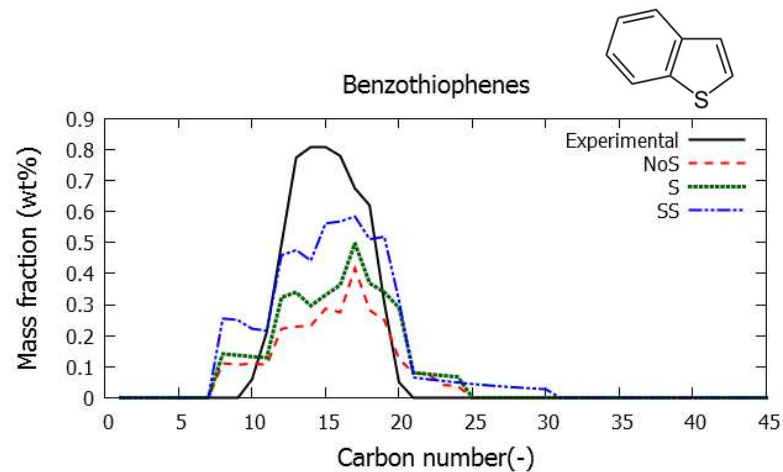
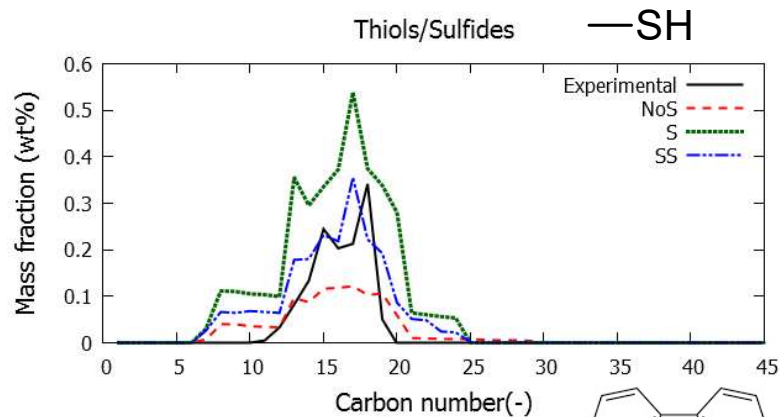
SAMPLE 2: *Brazil*

## Carbon Number Distribution



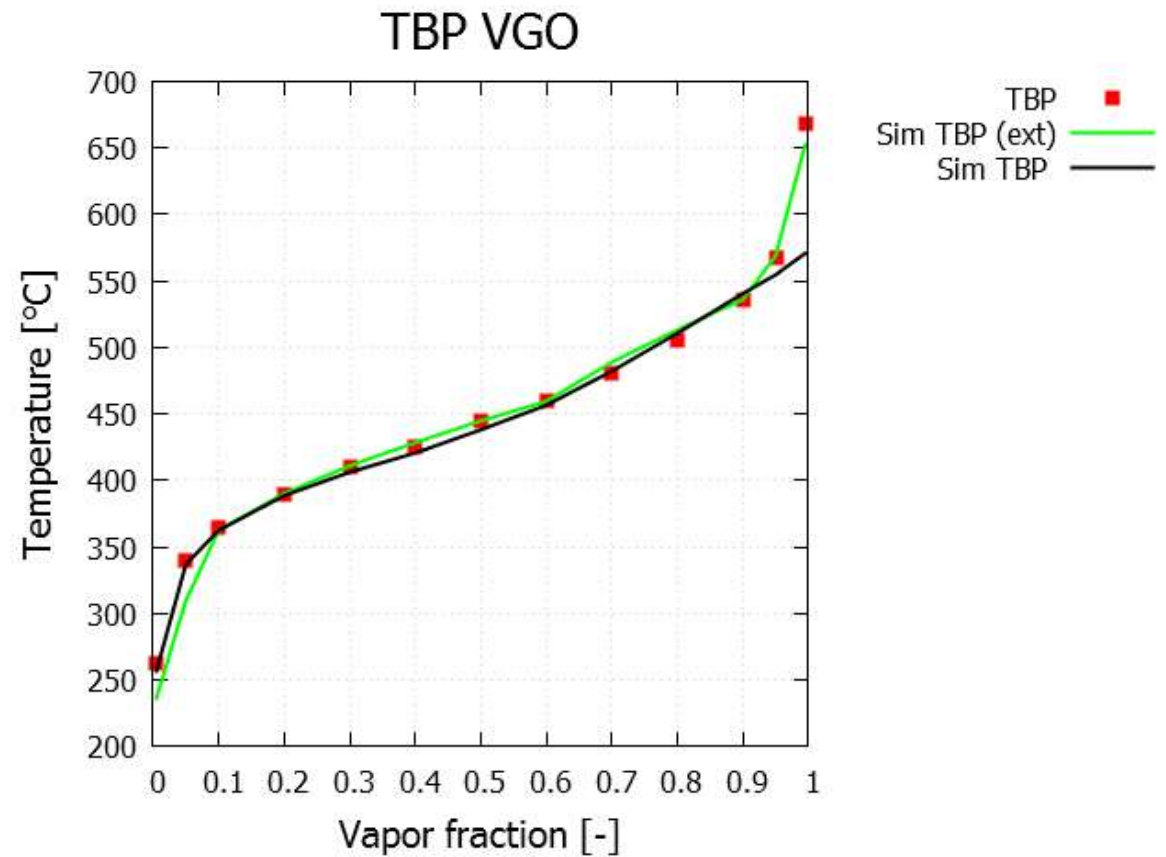
# Example: sulfur containing components

	Set 1	Set 2	Set 3
Density	Yes	Yes	Yes
Total Sulfur	No	Yes	Yes
Aromatic Sulfur	No	No	Yes
Group Type analysis	PINA		
Boiling point curve	ASTM-D2887		



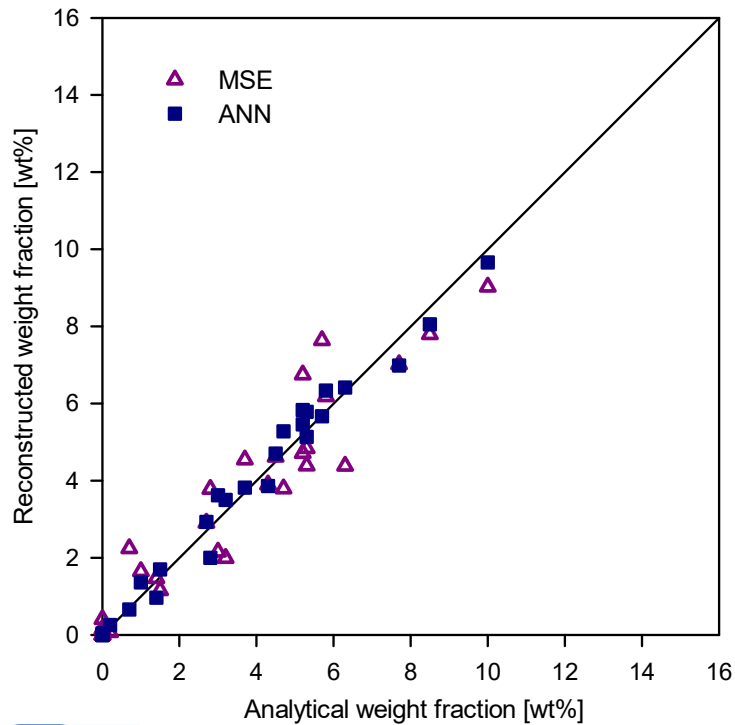
# Example: Vacuum Gas Oil (VGO)

Molecular library extended  
from  $C_{33}$  to  $C_{60}$

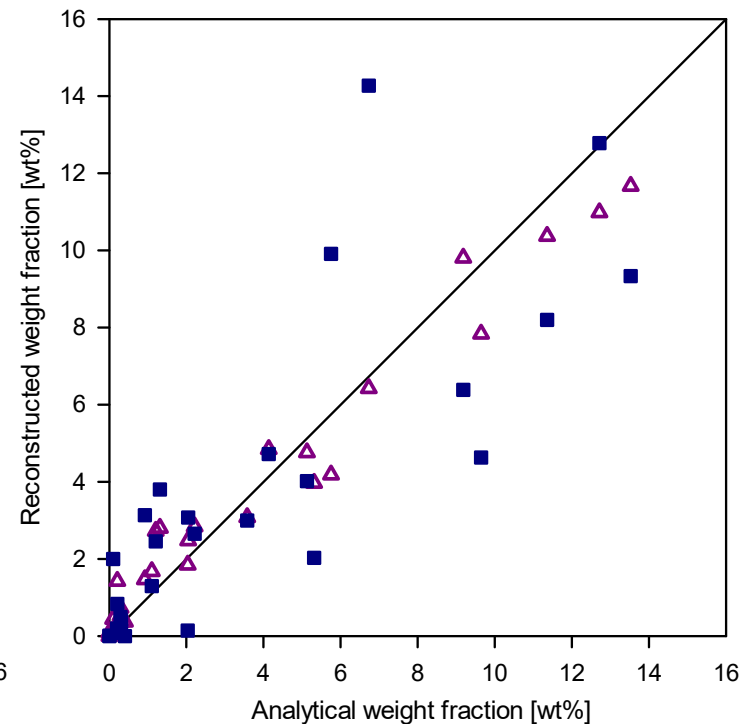


# Example: naphtha

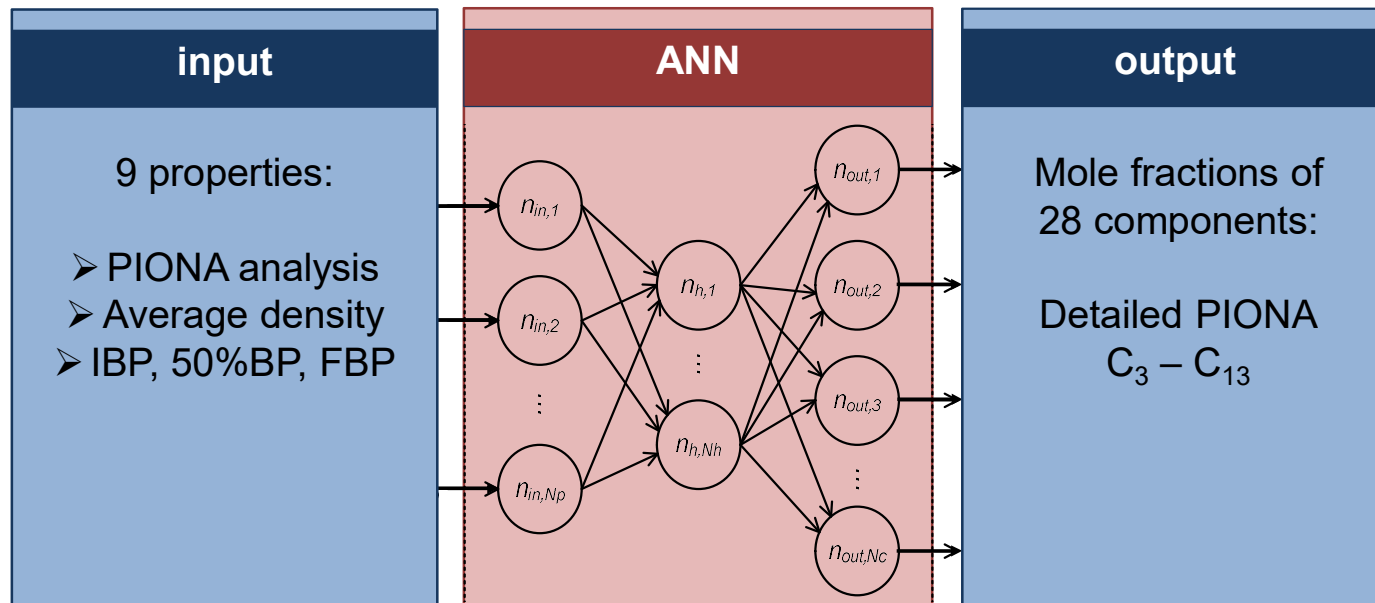
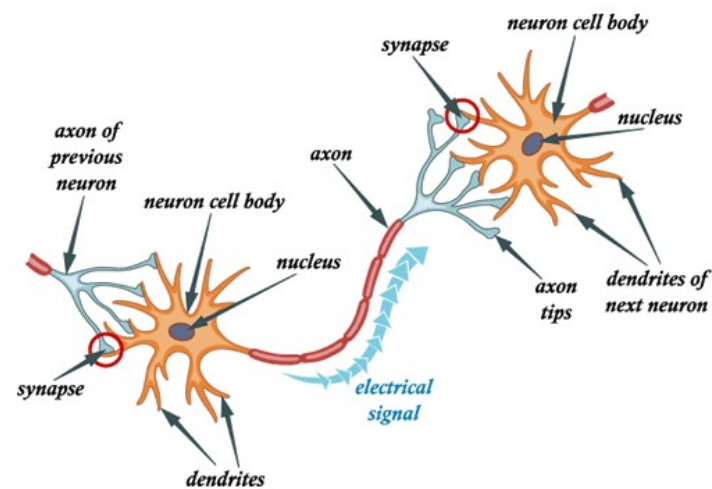
**Naphtha A – in range of ANN training set**



**Naphtha B – out of range of ANN training set**



# Artificial neural network (ANN): e.g. naphtha



ANN = nodes + weighted links

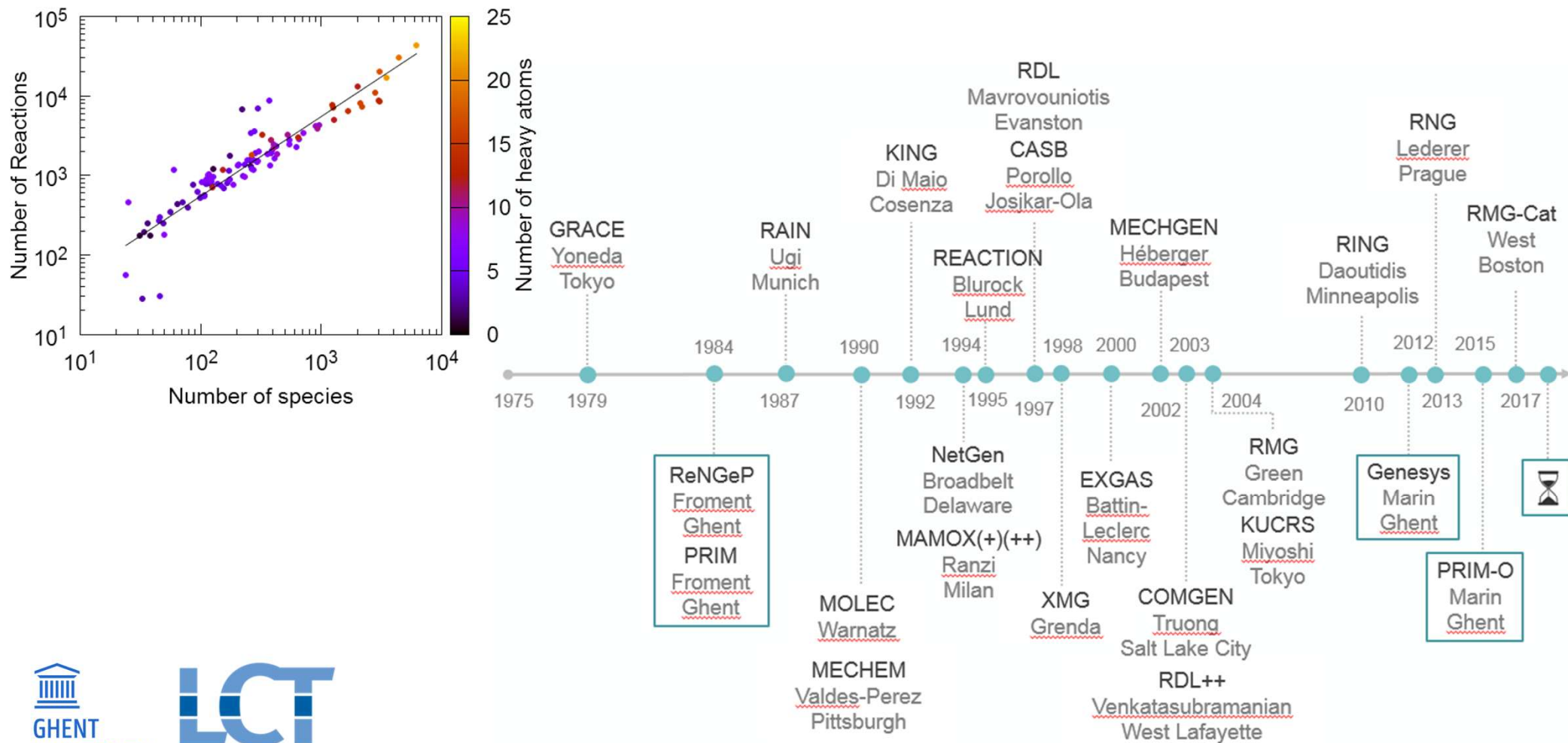
Weights and activation functions from training data set



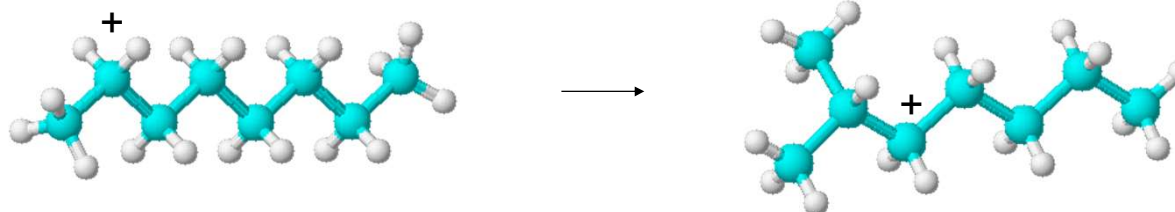
# Outline

- Introduction
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- Complex chemistry: parameter determination
- Reactor design
- Conclusions

# Network generation



# RENGEP: Matrix operations



	1	2	3	4	5	6	7	8	
1		1							
2	1		1						
3		1		1					
4			1		1				
5				1		1			
6					1		1		
7						1		1	
8							1		
+2									
2									
1 2 2 2 2 2 2 1									
26 26 26 26 26 26 26 26									

	1	2	3	4	5	6	7	8	
1		1							
2	1		1					1	
3		1		1					
4			1		1				
5				1		1			
6					1		1		
7						1			
8		1							
+2									
4									
1 3 2 2 2 2 1 1									
26 26 26 26 26 26 26 26									

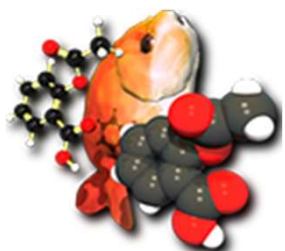
# GENESYS : GENERation of reacting SYStems



Open-Source Cheminformatics  
and Machine Learning



CDK



OpenBabel



- Computer sciences, chemistry and mathematics
- Open source projects
- Advanced species representation
- Advanced algorithms
  - Substructure matching
  - Transition state identification
  - 3D representation
    - Chirality

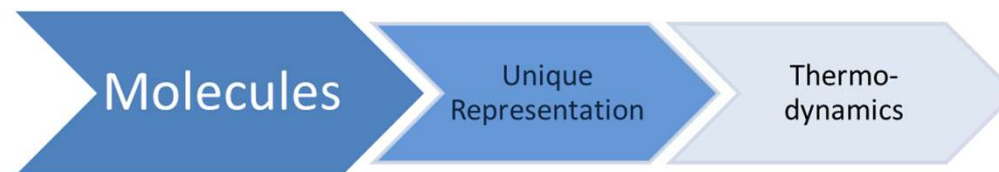
Vandewiele, N.M. et al., Chem. Eng. J., 207-208, 526-538, 2012

Van de Vijver, R. et al. Int. J. Chem. Kinet., 47 (4), 199-231, 2015

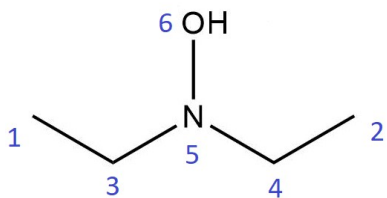
Vandewiele, N.M. et al., J. Comput. Chem., 36 (3), 181-192, 2015

Van de Vijver, R. et al., "Automatic kinetic model generation for heterogeneous catalysis", ISCRE 2018

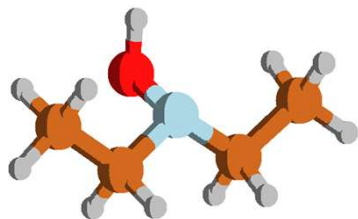
# Molecules



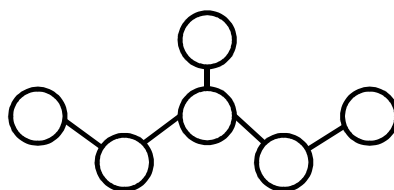
**Structural formula**



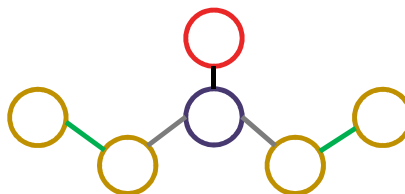
**3D structure**



**Graph structure**



**Weighted graph structure**



## - Molecular identifiers

### - SMILES:

CCN(O)CC

### - InChI:

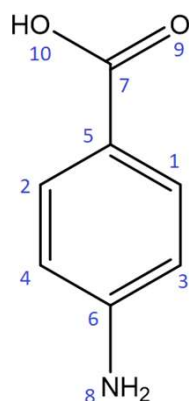
InChI=1S/C4H11NO/c1-3-5(6)4-2/h6H,3-4H2,1-2H3

## - Graph Theory

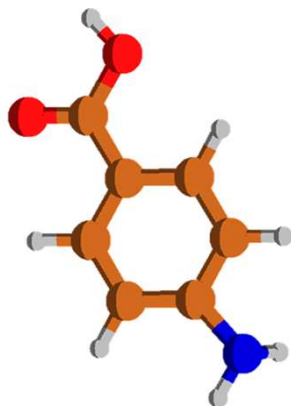
"Kinetics of Chemical Reactions : Decoding Complexity"  
G.B. Marin, G.S. Yablonsky, D. Constales Wiley-VCH Verlag,  
2<sup>nd</sup> edition, 2018

# Molecules

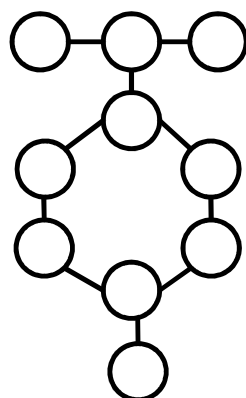
## Structural formula



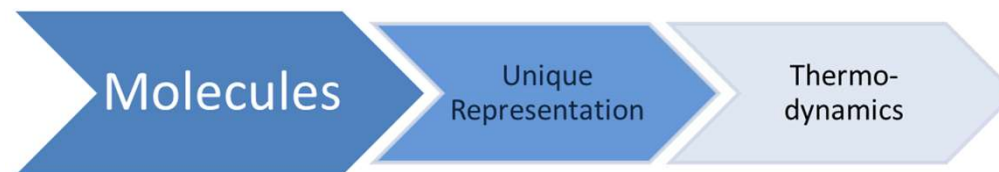
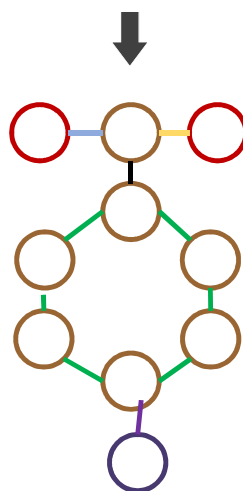
## 3D structure



## Graph structure



## Weighted graph structure



## - Molecular identifiers

### - SMILES:

NC1=CC=C(C(O)=O)C=C1

### - InChI:

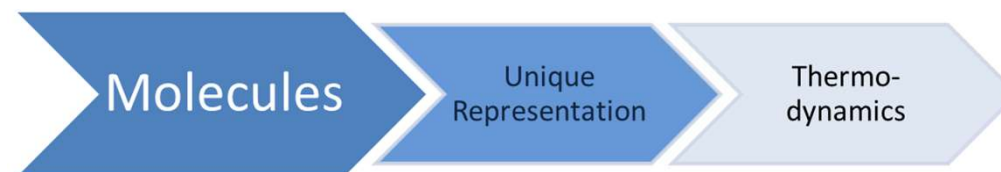
InChI=1S/C7H7NO2/c8-6-3-1-5(2-4-6)7(9)10/h1-4H,8H2,(H,9,10)

## - Graph Theory

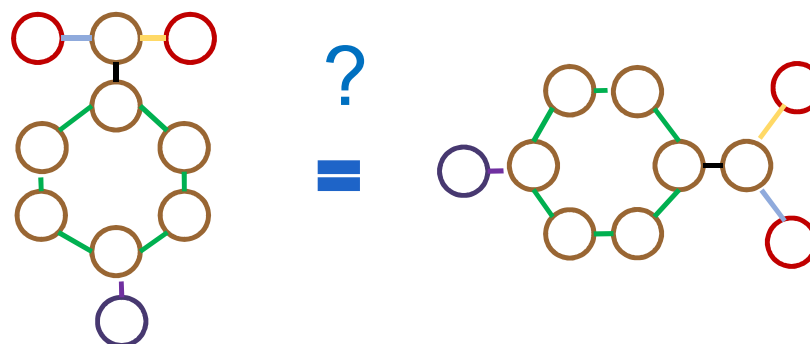
“Kinetics of Chemical Reactions : Decoding Complexity”  
G.B. Marin, G.S. Yablonsky, D. Constales Wiley-VCH Verlag, 2<sup>nd</sup> edition, 2018

Vandewiele, N.M. et al., Chem. Eng. J., 207-208, 526-538, 2012  
Van de Vijver, R. et al. Int. J. Chem. Kinet., 47 (4), 199-231, 2015  
Vandewiele, N.M. et al., J. Comput. Chem., 36 (3), 181-192, 2015

# Graph theory

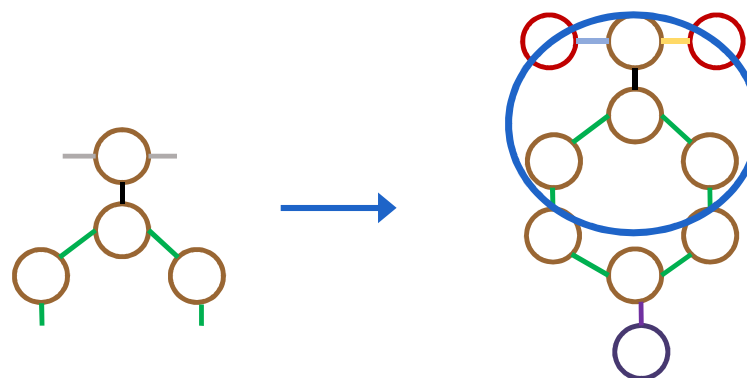


- Graph isomorphism

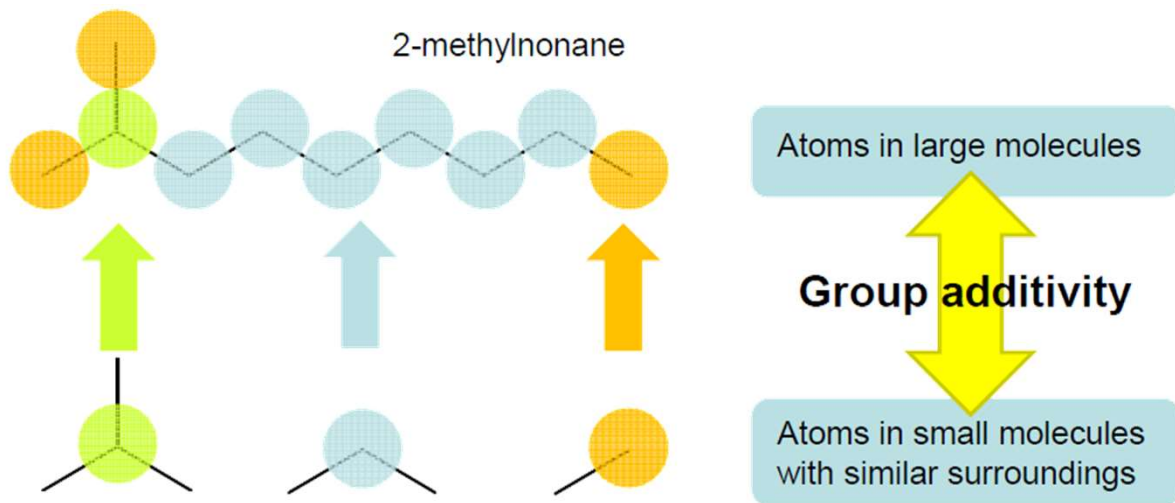
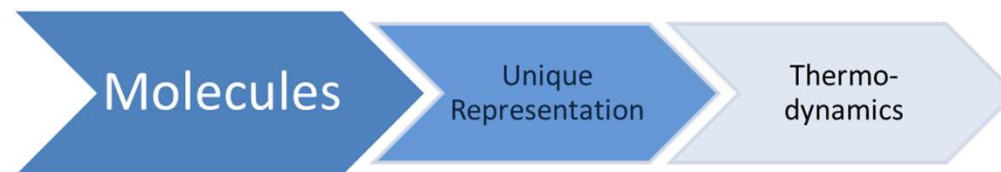


- Subgraph recognition

SMARTS c(:c)(:c)-[C;X3] →



# Group additivity

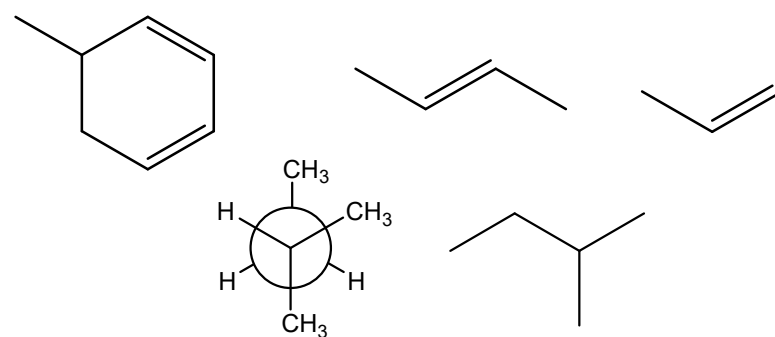


Group definition based on surroundings (ligands)

Additivity:



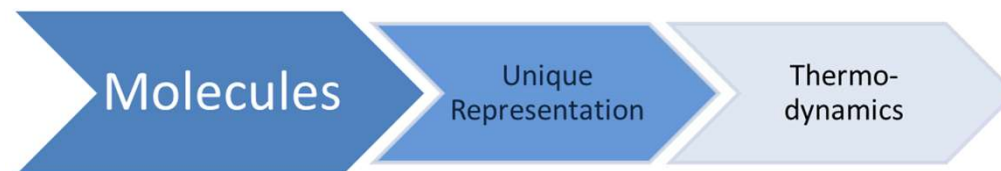
- Ring strain corrections
- Non-nearest neighbor interactions
  - Gauche interactions
  - Trans interactions
- Symmetry contribution



$$S = S_{int} - R \ln \sigma$$



# Thermodynamics

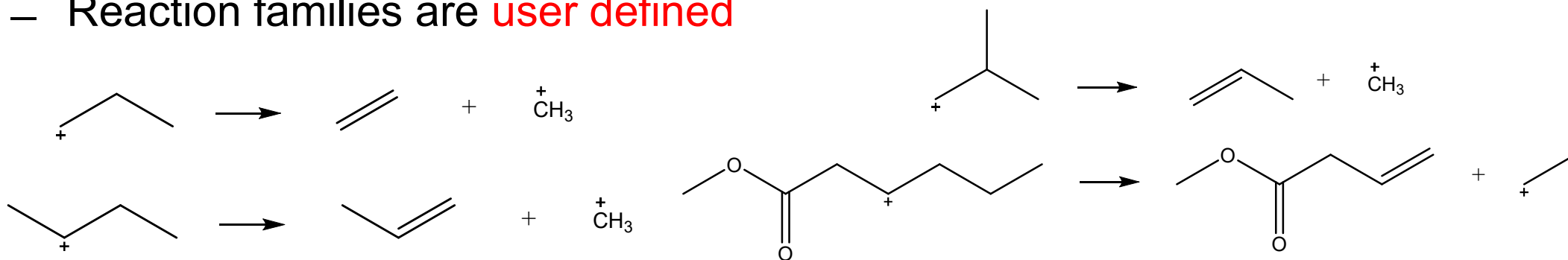


- “SpeciesThermo” database
  - Single entry per species
  - Originates from experimental work or high level quantum chemistry calculations
  - Limited in size
  - Database search via isomorphisms or InChI identifiers
- “BensonThermo” database
  - Group additive values
  - Originate from high level quantum chemical calculations (CBS-QB3)
  - Database search via SMARTS

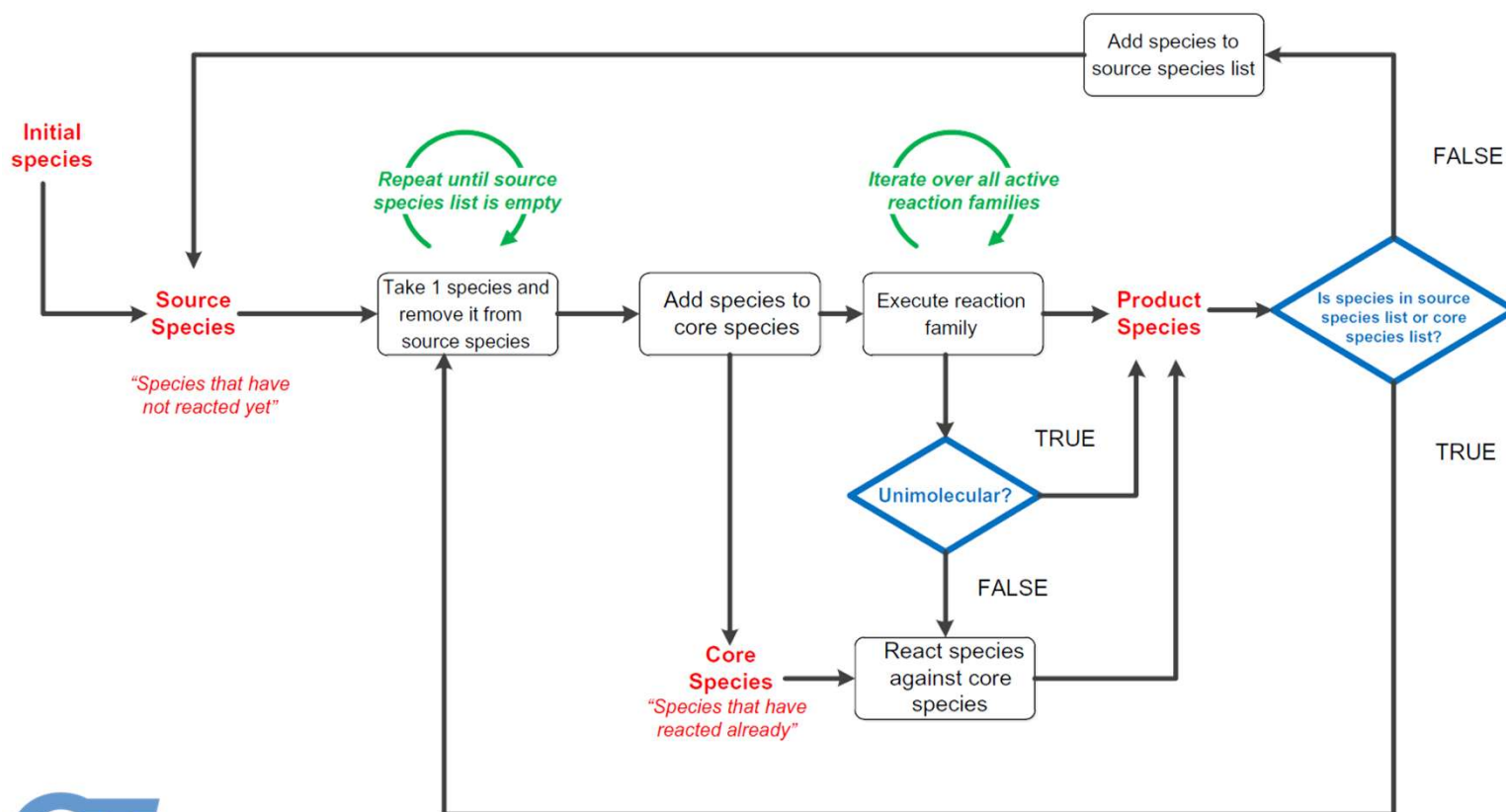
# Reactions



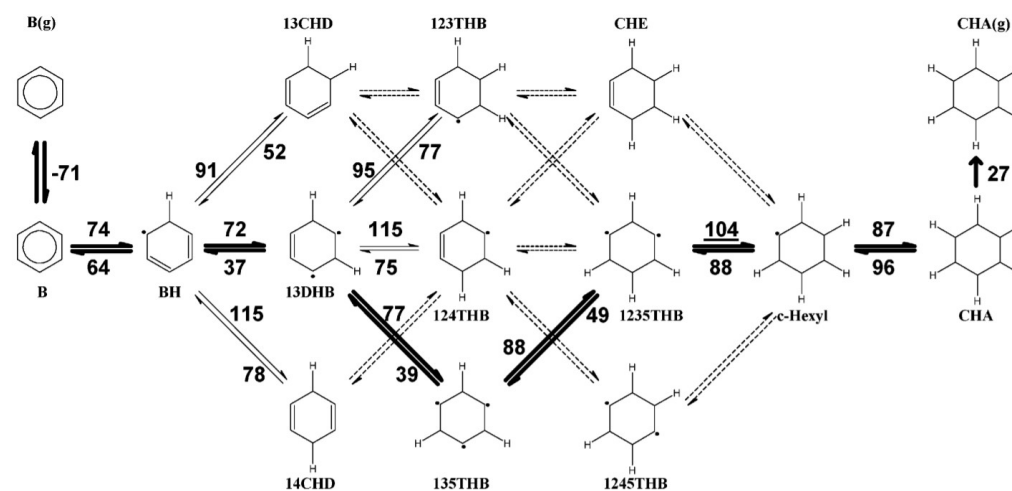
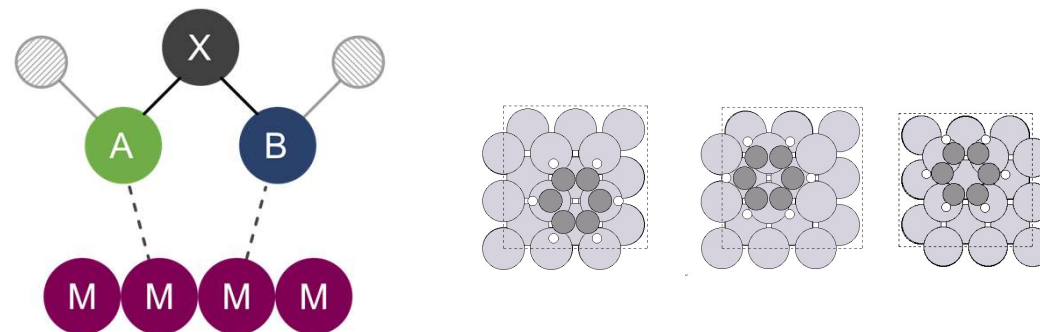
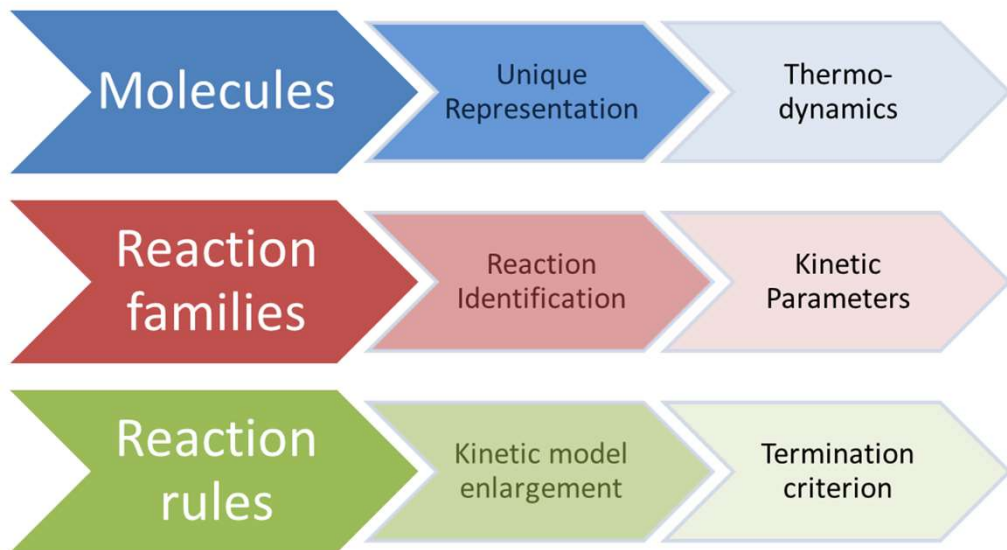
- Thousands of reactions can be classified in **only a few reaction families**
- All reactions within one reaction family have a very similar change in electronic structures of the atoms
- The reaction is mainly governed by a few atoms rather than the reactants in their entirety
- Reaction families are **user defined**



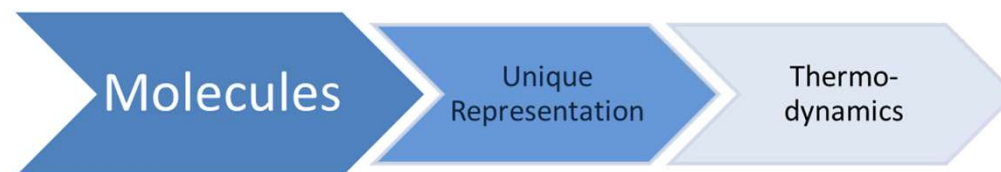
# Network Generation



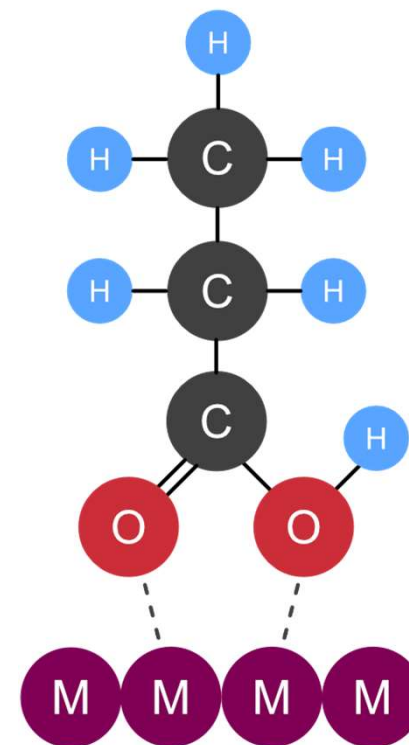
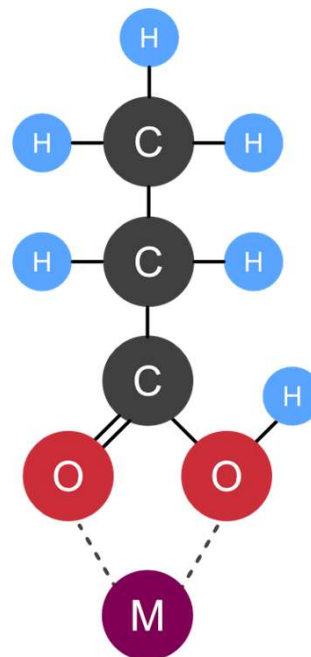
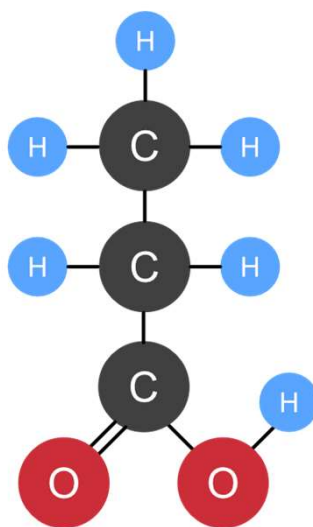
# Genesys: Catalytic Systems



# Molecules: Catalysis



- Dummy atoms for representation of surface
- No specific metal atoms: generation of general mechanism
- Properties of surface in separate input file
- Active site = multiple dummy atoms



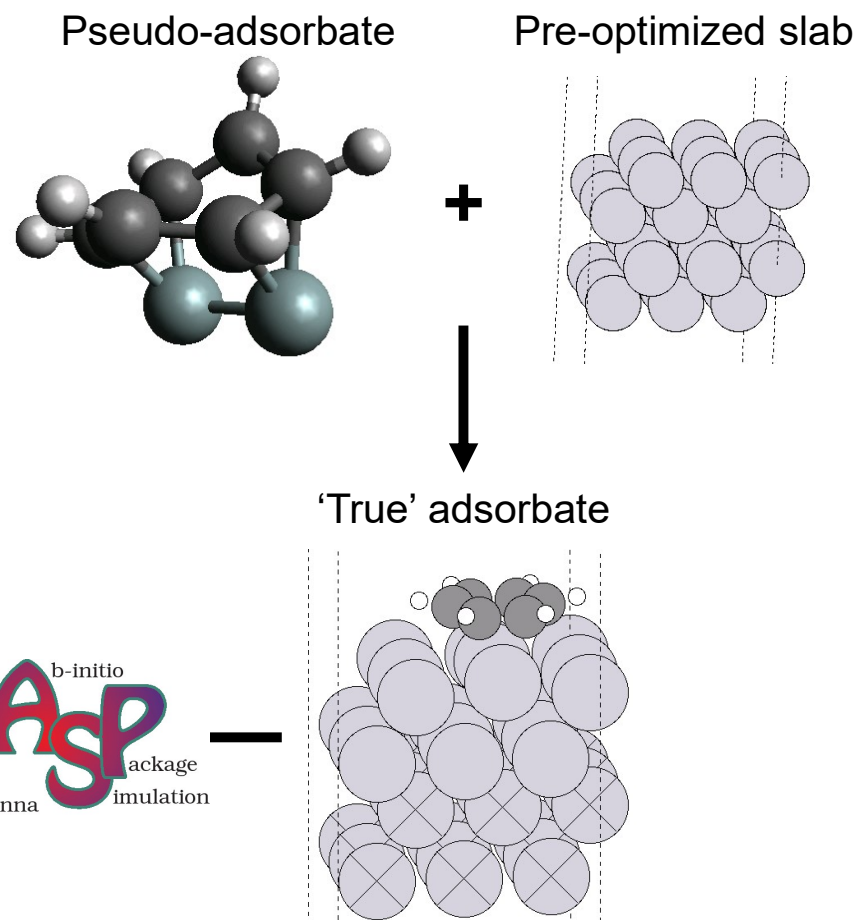
# Reaction Kinetics: catalysis

Reaction families

Reaction Identification

Kinetic Parameters

- Automated Ab-Initio calculations
  - Pre-optimized catalyst geometries
  - Pseudo-adsorbate
  - Combine + re-optimize
- Use obtained reaction (activation) energies to develop fast estimation methods

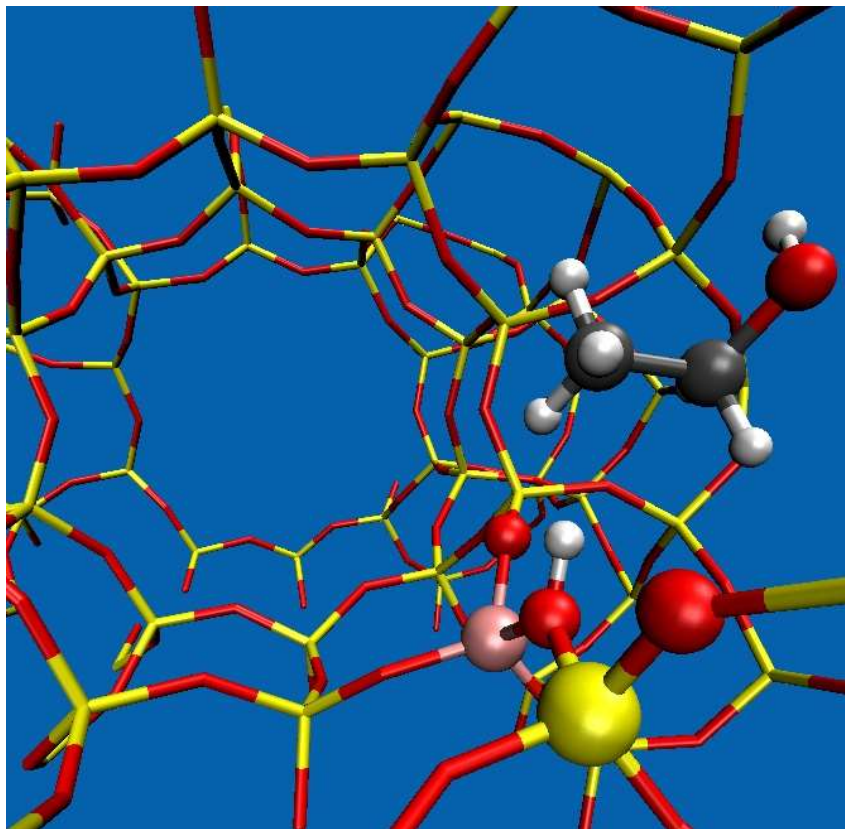


# Outline

- Introduction
- Complex feedstocks: ideal hydroconversion
- Feedstock characterization and reconstruction
- Reaction-network generation
- **Complex chemistry: parameter determination**
- Reactor design
- Conclusions

# Alcohol conversion in zeolites: e.g. $C_2H_5OH \rightarrow C_2H_4 + H_2O$

## Ethanol physisorption in H-ZSM-5

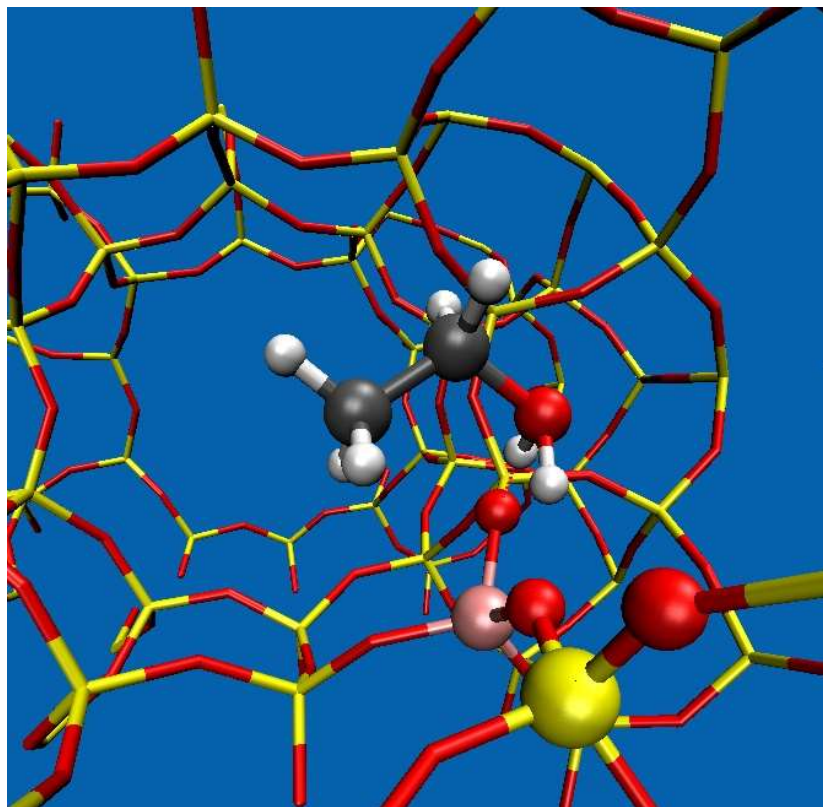


- van der Waals:
  - dipole-dipole
  - dipole-induced dipole
  - dispersive
- H-bonding
- electrostatic interactions

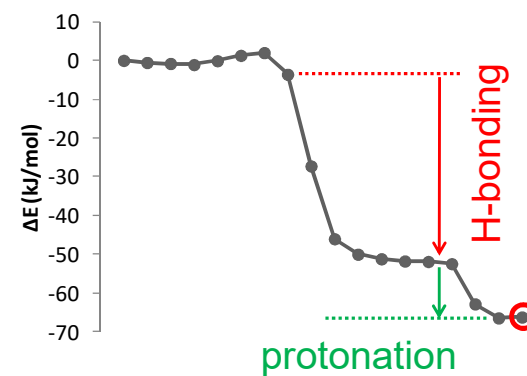


# H-bonding and protonation: chemisorption

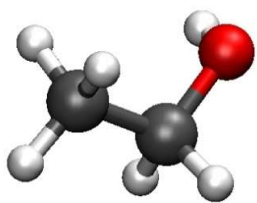
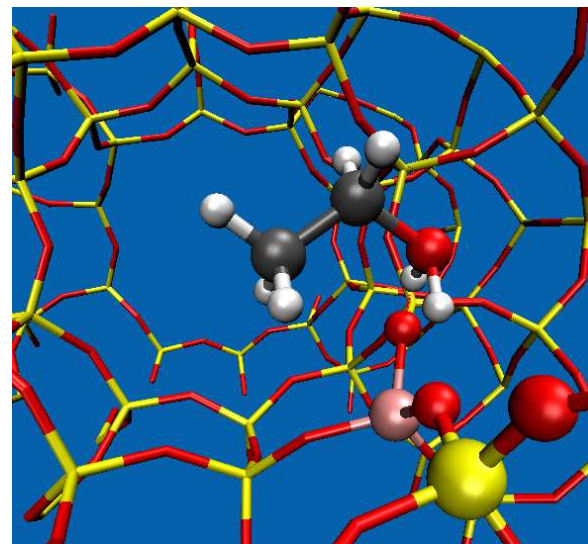
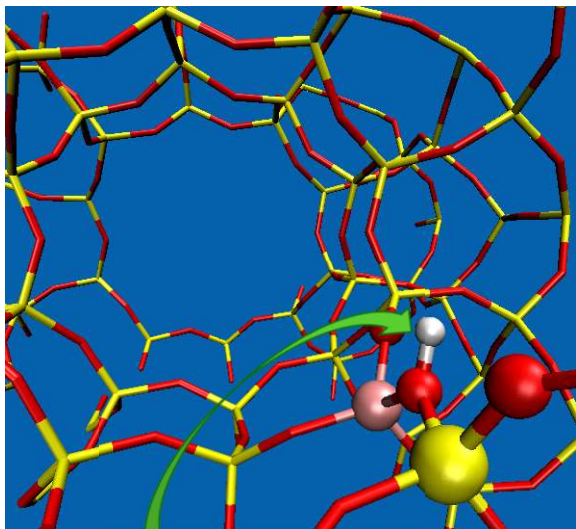
Ethanol adsorption and protonation in H-ZSM-5



Nudged Elastic Band calculation with PBE-D2 functional



# Alcohol adsorption & reaction in zeolites



$$r = k \theta_{ads,alcohol} = k K_{ads} P_{alcohol}$$

Ab initio calculation of:

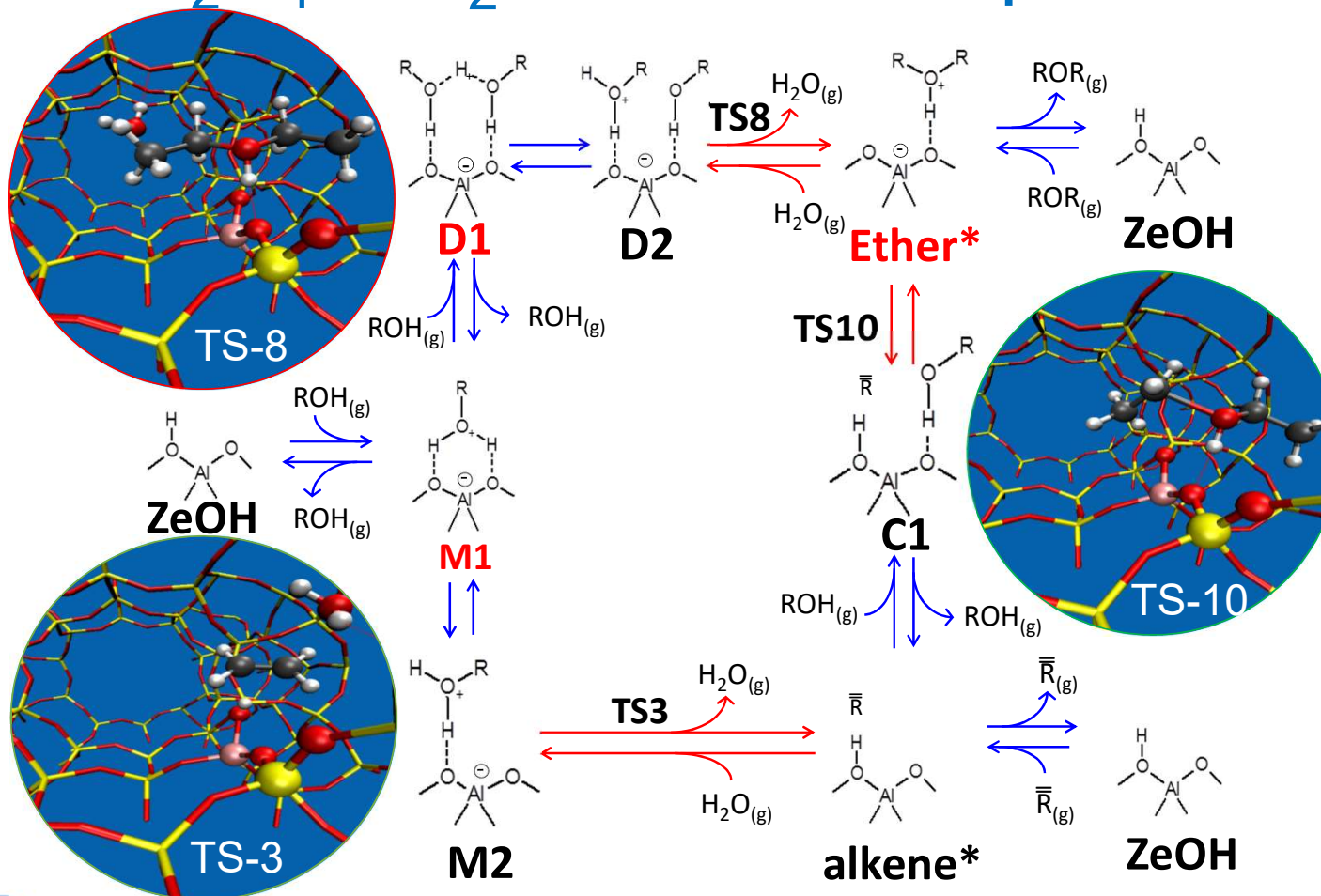
$$K_{ads} = \exp\left(-\frac{\Delta G_{ads}^0}{RT}\right) = \exp\left(-\frac{\Delta H_{ads}^0 - T\Delta S_{ads}^0}{RT}\right)$$

thermodynamics

$$k = \frac{k_b T}{h} \exp\left(\frac{\Delta S^{0,\#}}{R}\right) \exp\left(-\frac{\Delta H^{0,\#}}{RT}\right) = \frac{k_b T}{h} \exp\left(-\frac{\Delta G^{0,\#}}{RT}\right)$$

kinetics

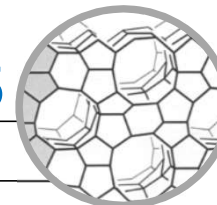
# $C_2H_5OH \rightarrow C_2H_4 + H_2O$ : dominant paths



# Experimental procedures and conditions



HZSM-5



## Properties

Si/Al	15	40
$c_{H^+}$ (mol kg <sup>-1</sup> )	0.77	0.36
BET (10 <sup>3</sup> m <sup>2</sup> kg <sup>-1</sup> )	430	436
$V_{\text{micro}}$ (10 <sup>-5</sup> m <sup>3</sup> kg <sup>-1</sup> )	1.1	1.1

## Experimental conditions

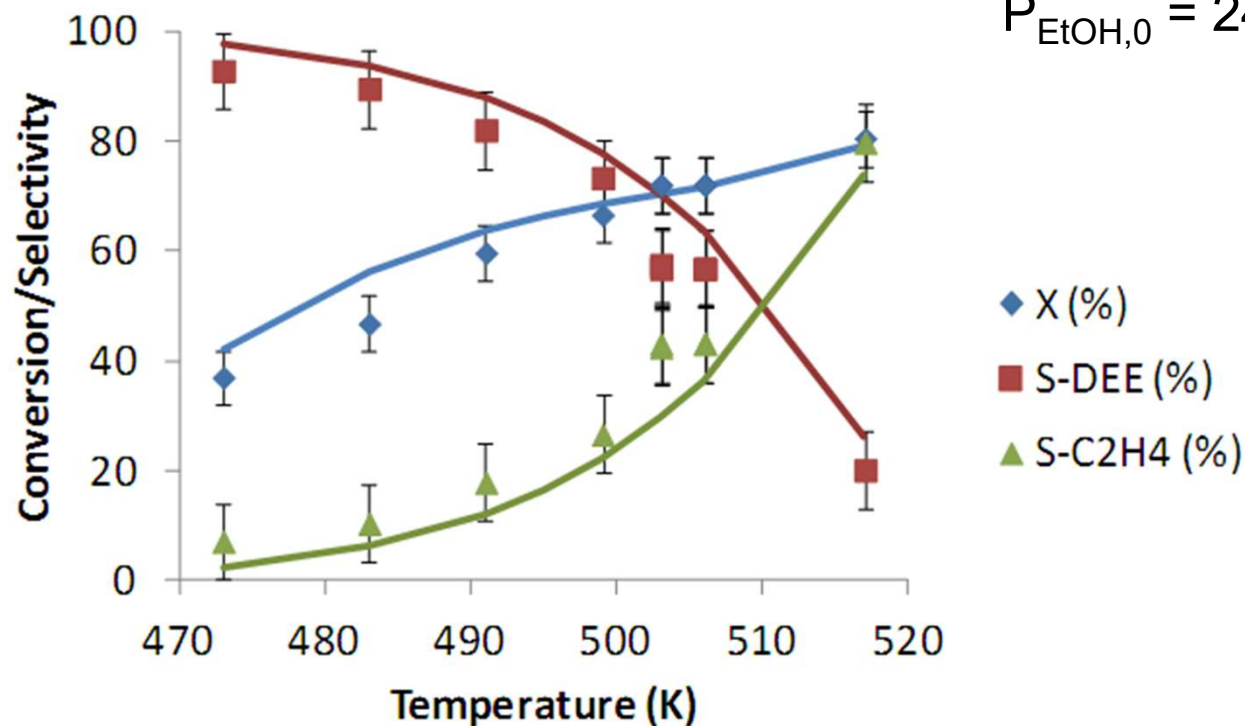
Temperature (K)	453 – 523
$p_{\text{EtOH,in}}$ (kPa)	8 – 50
$W/F_{\text{EtOH,in}}$ (kg s mol <sup>-1</sup> )	1.5 – 17.0

# Conversion and selectivities

HZSM-5

$W_{\text{cat}}/F_{\text{EtOH},0} = 6.5 \text{ kg s} / \text{mol}$

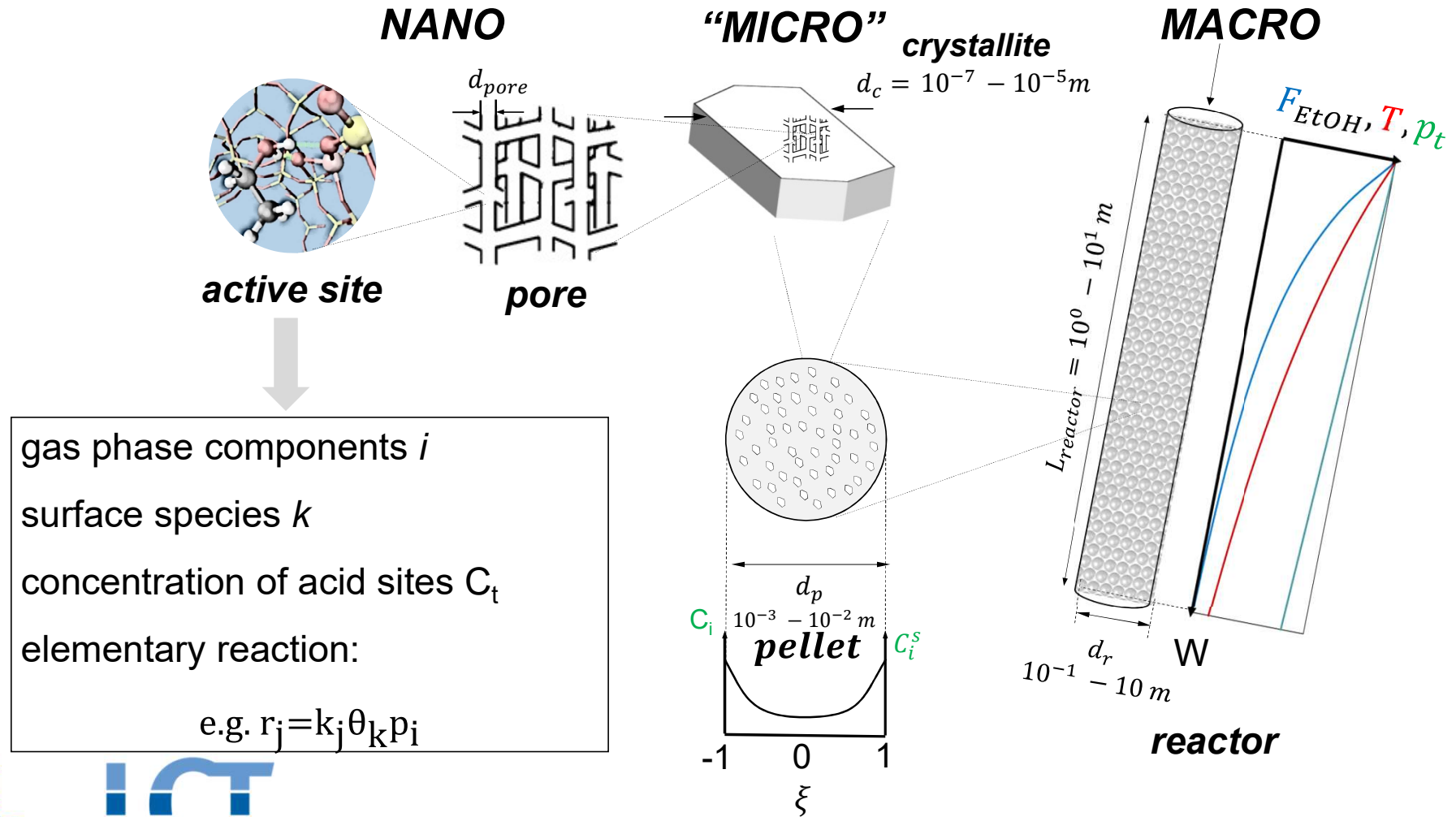
$P_{\text{EtOH},0} = 24 \text{ kPa}$



# Outline

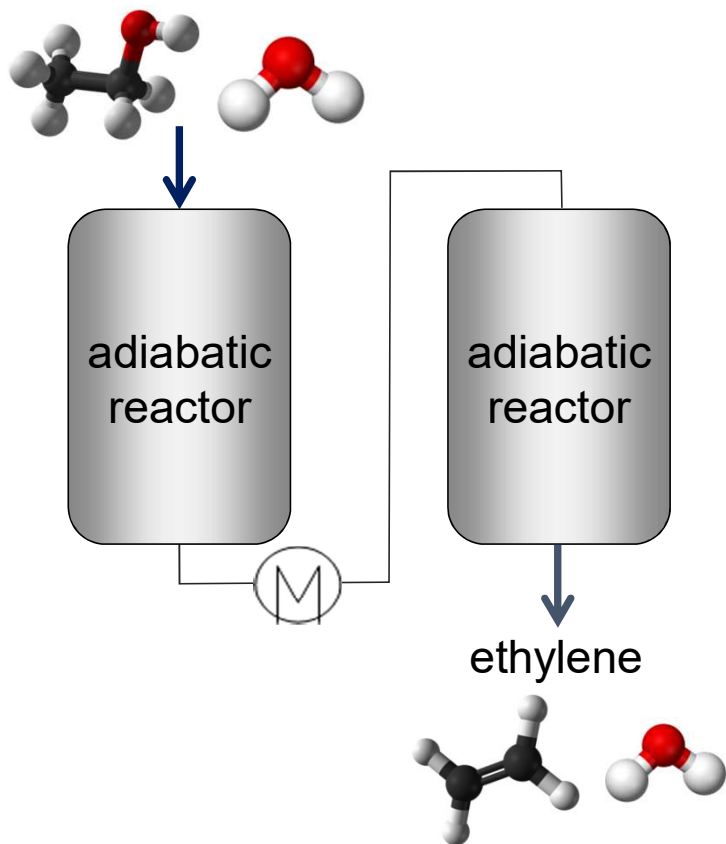
- Introduction
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- Conclusions

# Multiscale modeling of an industrial reactor



# Industrial dehydration reactor

bio-ethanol (aqueous ethanol solution)



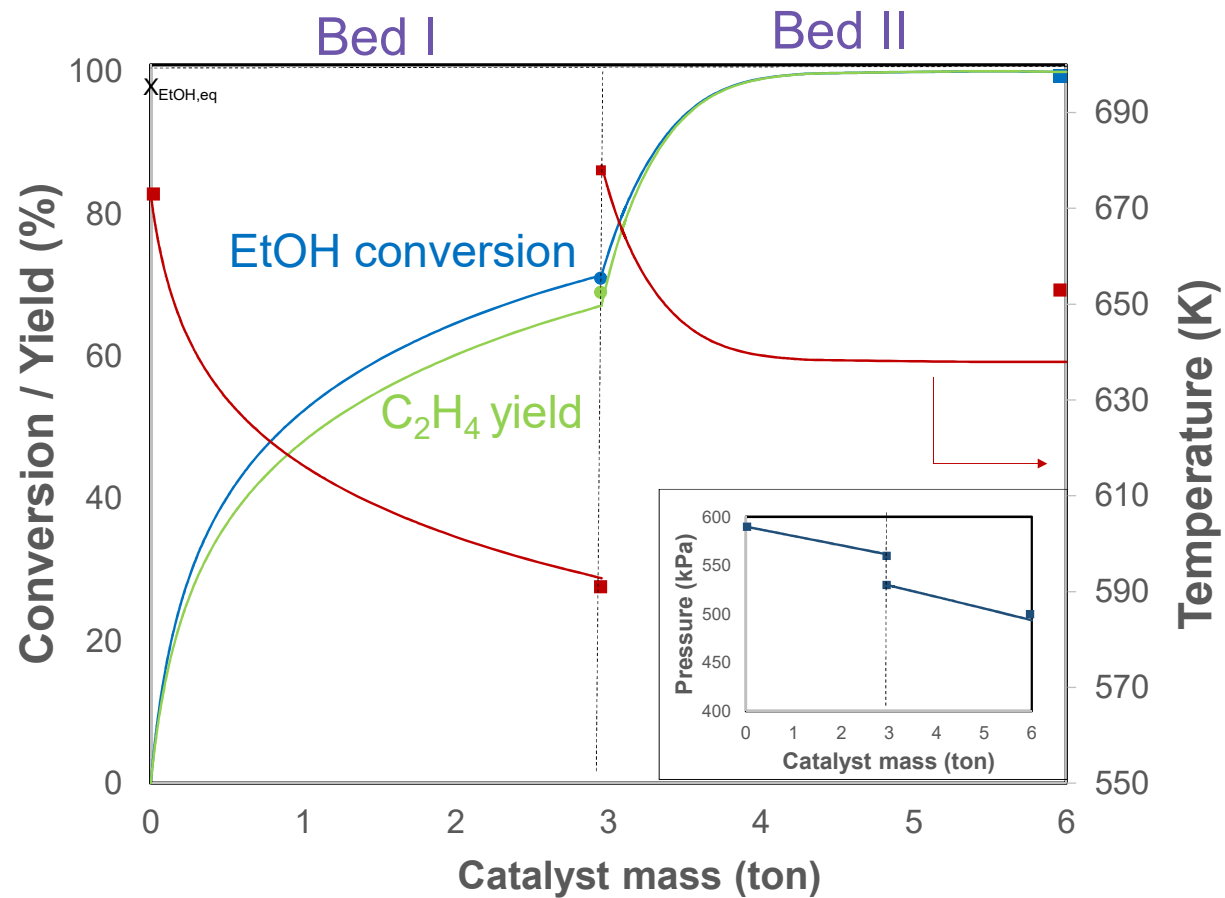
## Design specifications<sup>1</sup>

$T^0$ (K)	673
$P^0$ (kPa)	590
Ethylene production (kT y <sup>-1</sup> )	220
Ethanol content (wt.%)	26
HZSM-5 mass (ton)	6





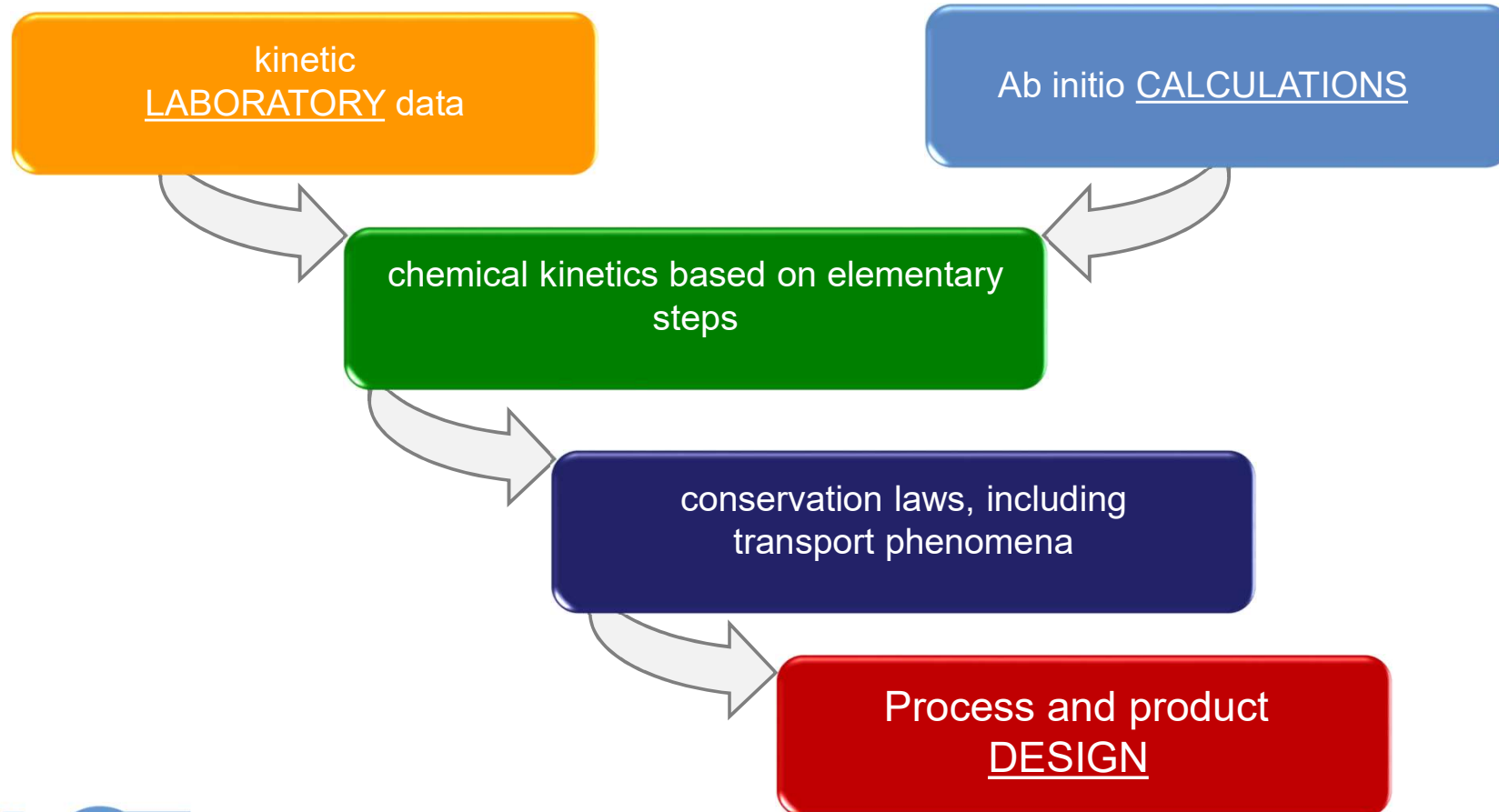
# Industrial multibed adiabatic operation



# Outline

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- Reactor design
- **Conclusions**

# Kinetics: From Molecules to Process/Products



# Acknowledgments

Profs. Marie-Françoise Reyniers and Maarten Sabbe

Dr. Kostas Alexopoulos

Dr. Manh Cuong Nguyen

Dr. Ruben Van de Vijver

Dr. Jonas Van Belleghem

Dr. Steven Pyl

Dr. Chetan Raghuvier

Dr. Thomas Dijkmans

Dr. Nick Vandewiele

Dr. Bart Vandegehuchte

Dr. Gisela Lozano-Blanco

Dr. Indranil Choudhary

Dr. Laxmi Narasimhan

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# Glossary

GPC	Gel permeation chromatography
FBP	Final boiling point
FT-NIR	Transform-near infrared
FT-NMR	FT-nuclear magnetic resonance
GC	Gas chromatography
GC × GC	Comprehensive two-dimensional gas chromatography
HPLC	High-performance-liquid-chromatography
IBP	Initial boiling point
CE	Capillary electrophoresis
MS	Mass spectrometry
GFC	Gel Filtration Chromatography
PIONA	Paraffins Isoparaffins Olefins Naphthenes and Aromatics
SFC	Supercritical fluid chromatography
IC	Ion Chromatography

# Glossary

Roof-tilt	In GCxGC: ascending band of components with similar features
TOF-MS	Time-of-Flight Spectrometry
SSL	Split/Splitless
PTV	Programmable Temperature Vaporizing
COC	Cold on Column
NP	Normal Phase
RP	Reversed Phase
SCD	Sulfur Chemiluminescence Detector
NCD	Nitrogen Chemiluminescence Detector
DAD	Diode-Array Detector
ELSD	Evaporative Light Scattering Detector
FPD	Flame Photometric Detector
SARA	Saturates Aromatics Resins Asphaltenes
FID	Flame Ionization Detector

# Glossary

- **SMILES** (Simplified Molecular Input Line Entry System): line notation for representing chemical structures. By making use of a canonicalization algorithm, a “unique SMILES” can be generated. SMILES are also quite compact compared to other methods for representing chemical structures (making them more human readable) and they allow isotopic and chiral specifications.
- **SMARTS** (SMiles ARbitrary Target Specification): line notation that allows to specify substructures using rules that are straightforward extensions of SMILES. In addition to the atomic and bond symbols, wildcard atoms and bonds can be specified and logical operators can be used, allowing an extremely precise sub structural specification. Using SMARTs flexible and efficient substructure-search specifications can be made (database searching, reactive center identification ...)



# Glossary

- **InChI** (IUPAC International Chemical Identifier): InChI encodes a chemical structure into “layers”, with each layer holding a distinct class of structural information (main layer, charge layer, stereochemical layer, isotopic layer, fixed-H layer and reconnected layer). Only the main layer specifying the chemical formula, atoms, and bonds between them, is required for all InChIs. The InChI is open-source and the software for generating InChI strings is freely available (in contrast to SMILES). However, they are less human readable compared to SMILES strings, because InChI was developed as a “machine-readable” chemical identifier.
- **Fixed H-layer**: When potentially mobile hydrogen atoms are present and the user specifies that they should be immobile (e.g. tautomerism not allowed), this layer binds these hydrogen atoms to the atoms specified in the input structure.
- **Reconnected layer**: when e.g. all metal atoms are disconnected in the main layer. The user can request to add a “reconnected” layer that generates an InChI containing all bonds given in the input structures.

# Glossary

- **CDK** (Chemistry Development Kit): collection of modular Java libraries for processing chemical information (Cheminformatics). The modules are free and open-source and are easy to integrate with other open-source or in-house projects. Functionalities include: molecule and reaction valence bond representation, reading and writing file formats such as SMILES and InChI, canonical identifiers for fast exact searching, substructure and SMARTS pattern searching...
- **RDKit**: cheminformatics toolkit similar to CDK, with the same core functionalities, but written in another language (C++ versus Java).
- **OpenBabel**: software mainly used to interconvert chemical file formats or representations. Other functionalities include generation of 3D molecular structure, generation of molecular fingerprints, calculation of partial charges ...

# Glossary

- **Benson group:** polyvalent atom, i.e. a central atom with ligancy  $\geq 2$ , together with all of its ligands. The corresponding notation is  $X-(A)_i(B)_j(C)_m(D)_l$ , in which X represents the central atom surrounded by  $i$  ligands of atom A,  $j$  ligands of atom B and so on.
- **(Un)weighted graph:** An unweighted graph object consists of a set of nodes or vertices representing the atoms, connected by edges representing the chemical bonds. A weighted graph introduces additional information. The vertices can bear information about the atom type, its electronic configuration etc. The edges can refer to single, double, triple, transition-state etc. bonds.