

Comparison of polysaccharide stationary phases in supercritical fluid chromatography

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Why supercritical fluid chromatography ?

Packed column SFC uses a **carbon dioxide - solvent** mobile phase

Viscosity is lessened : higher flow rates are possible and column efficiency is improved (peaks are thinner)

Solvent consumption is usually reduced as compared to HPLC

Collected fractions are more concentrated in prep

Overall productivity is usually improved



Current state of chiral SFC

Similarly to HPLC:

Limited understanding of the factors favouring successful separation

Screening strategies are the only ones used everywhere

- time-optimized strategies: SFC can be very fast with short columns, gradient elution and parallel screening

- usually optimized in terms of complementarity of the systems screened

Is it good enough?

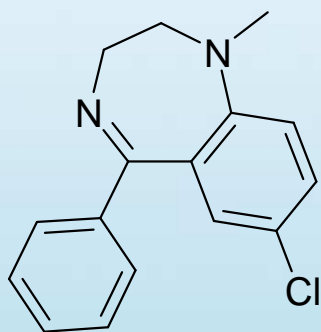
Can we do anything better?

Objectives of our study

To propose a classification of chiral stationary phases for SFC use

The knowledge acquired in HPLC is not sufficient

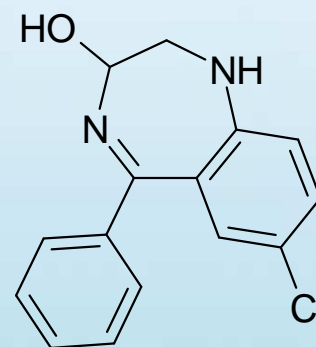
HPLC-SFC transfer is not always immediate



α (LC) = 1

α (SFC) = 8.09

diazepam



α (LC) = 1.39

α (SFC) = 1

oxazepam

Lux Cellulose-1, CO₂-iPrOH or HPT-iPrOH 90:10 (v/v), 25°C, SFC : 150 bar 3 mL/min, HPLC : 1 mL/min



Objectives of our study

To propose a classification of chiral stationary phases for SFC use

To propose compound-tailored screening strategies rather than generic strategies

To possibly predict the outcome of a separation

To improve our understanding of enantioseparation mechanisms

How to proceed?

Need for consistent data to compare chromatographic systems

Litterature is a mess !!!

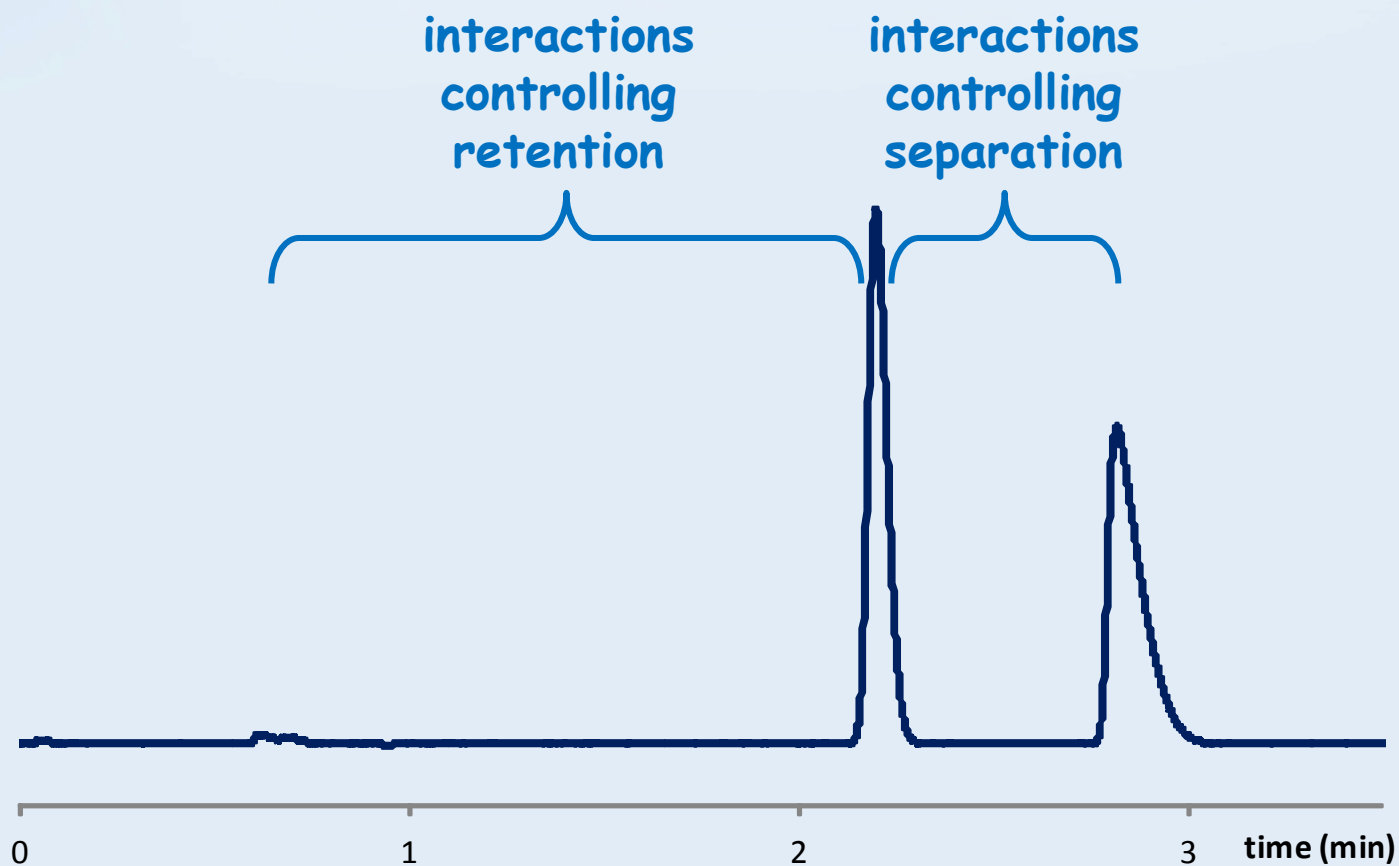
Large set of compounds with structural diversity

Identical operating conditions for all columns

CO₂-MeOH 90:10 (v/v), 25°C, 150 bar, 3 mL/min

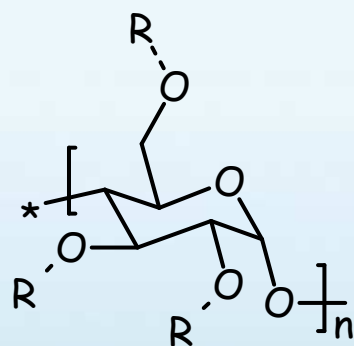
How to proceed?

Need to distinguish between

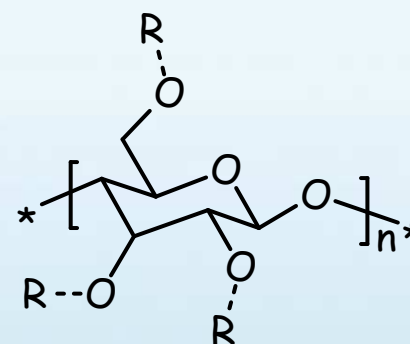


Modified polysaccharide chiral stationary phases

Amylose

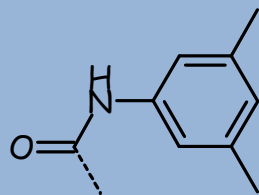


Cellulose

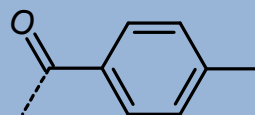


Phenomenex chiral stationary phases (CSP)

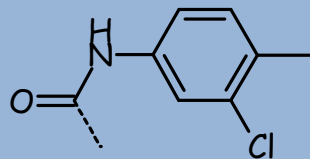
4 CSPs based on Cellulose



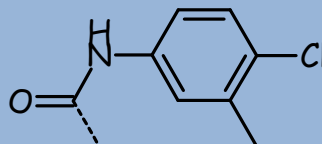
Lux Cellulose-1 (LC1)



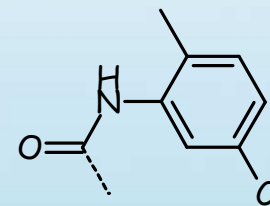
Lux Cellulose-3 (LC3)



Lux Cellulose-2 (LC2)

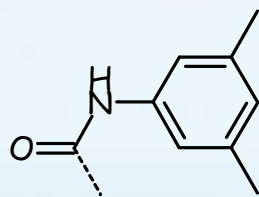


Lux Cellulose-4 (LC4)

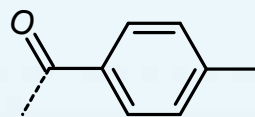


Lux Amylose-2 (LA2)

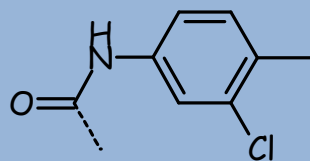
Phenomenex chiral stationary phases (CSP)



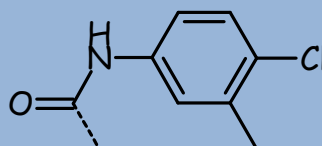
Lux Cellulose-1 (LC1)



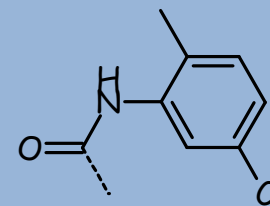
Lux Cellulose-3 (LC3)



Lux Cellulose-2 (LC2)



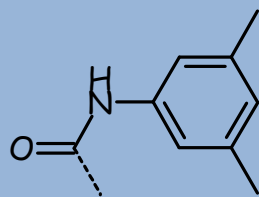
Lux Cellulose-4 (LC4)



Lux Amylose-2 (LA2)

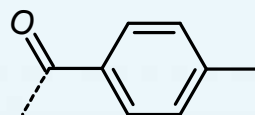
3 chlorinated CSPs

Phenomenex chiral stationary phases (CSP)

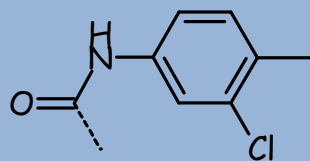


Lux Cellulose-1 (LC1)

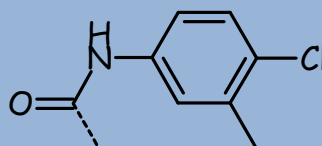
1 ester ligand



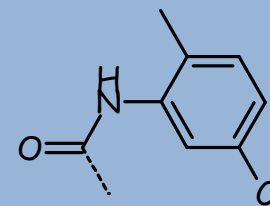
Lux Cellulose-3 (LC3)



Lux Cellulose-2 (LC2)



Lux Cellulose-4 (LC4)



Lux Amylose-2 (LA2)

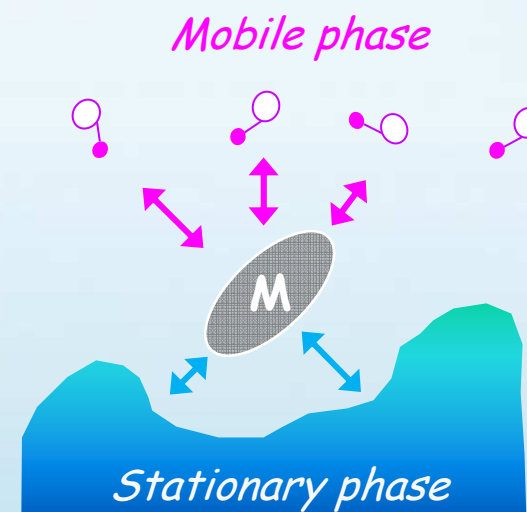
4 carbamate ligands

A method to compare stationary phases

The analyte interacts with the **stationary phase** and the **mobile phase**

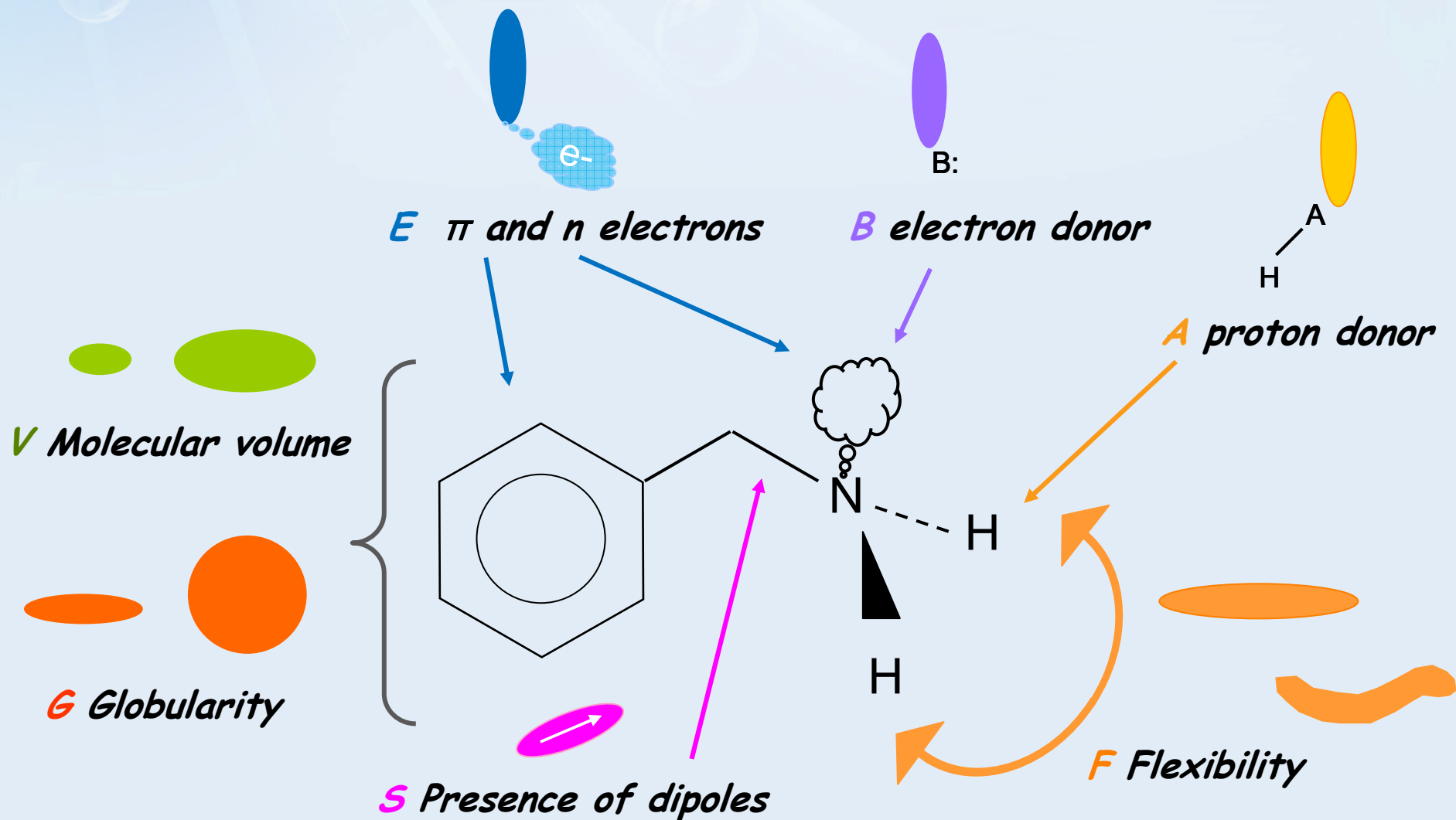
depending on molecular characteristics inducing different capabilities for interaction

Example: an acidic compound can interact via hydrogen-bonding



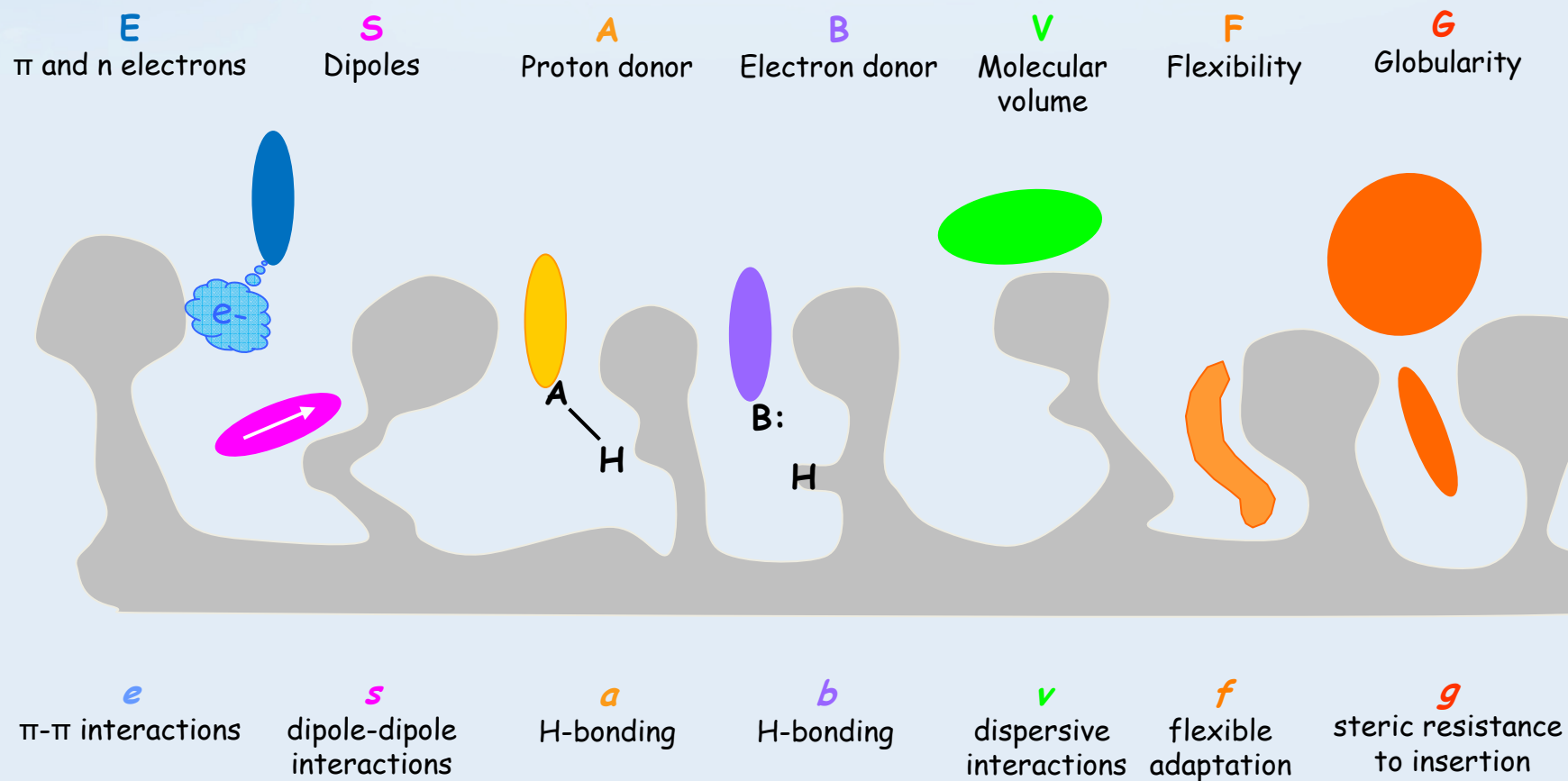
A method to compare stationary phases

Analyte capabilities for interaction



A method to compare retention properties

Interactions in the chromatographic system



A method to compare retention properties

Modified solvation parameter model

$$\log k = c + eE + sS + aA + bB + vV + fF + gG$$

E
π and n electrons

S
Dipoles

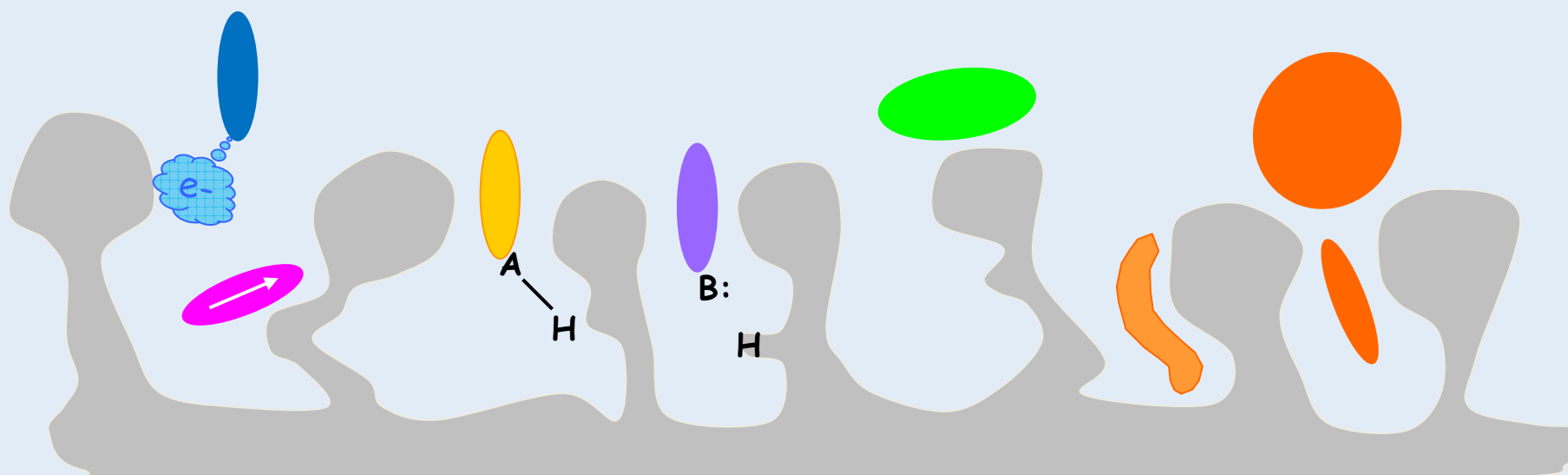
A
Proton donor

B
Electron donor

V
Molecular
volume

F
Flexibility

G
Globularity



e
π-π interactions

s
dipole-dipole
interactions

a
H-bonding

b
H-bonding

v
dispersive
interactions

f
flexible
adaptation

g
steric resistance
to insertion

A method to compare retention properties

Modified solvation parameter model

$$\log k = c + eE + sS + aA + bB + vV + fF + gG$$

Determine the properties of a group of probe analytes (>200):
E, S, A, B, V, F and G with a dedicated software
(or literature data: 4000 compounds)

Measure retention *k* for each of them in the chromatographic system you wish to characterize

Do a multiple linear regression to determine the system coefficients: *e, s, a, b, v, f and g*.

(e, s, a, b, v, f, g)
represent retention in the chromatographic system

A method to compare retention properties

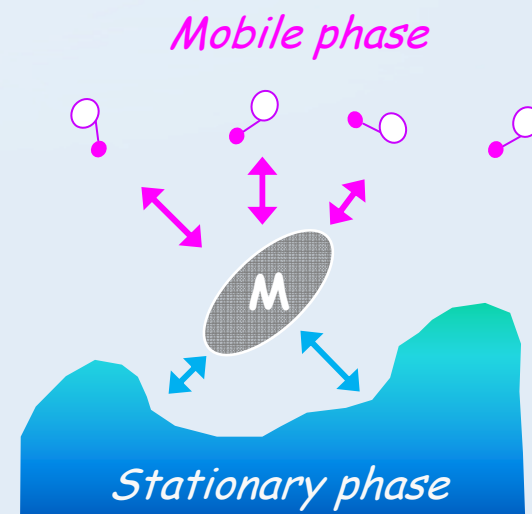
Modified solvation parameter model

$$\log k = c + eE + sS + aA + bB + vV + fF + gG$$



For each coefficient x :

$$x = x_{\text{stationary}} - x_{\text{mobile}}$$

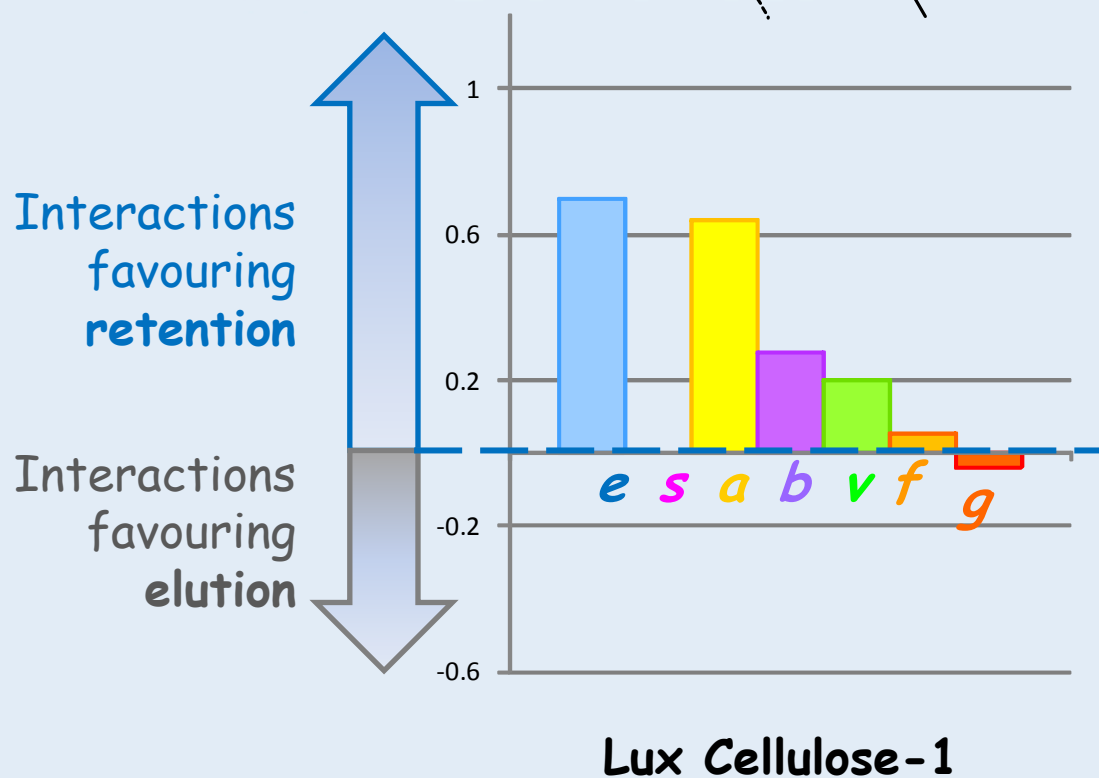
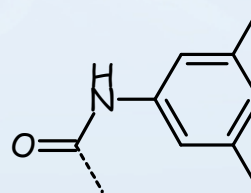


If $x_{\text{stationary}}$ is constant, *mobile phases can be compared*

If x_{mobile} is constant, *stationary phases can be compared*

A method to compare stationary phases

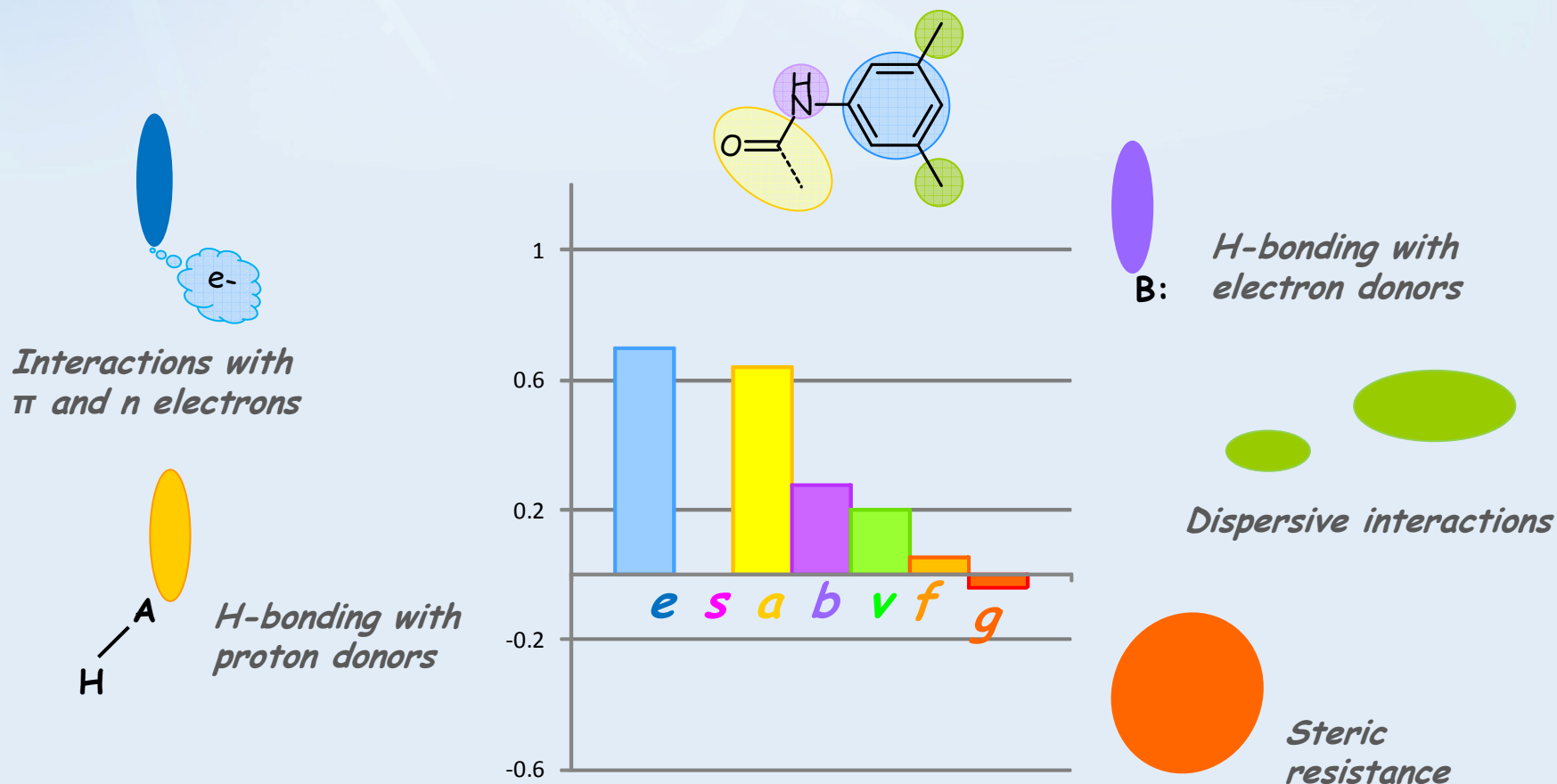
The solvation parameter model



Operating conditions:
CO₂-MeOH 90:10 (v/v)
25°C, 150 bar 3 mL/min

A method to compare stationary phases

The solvation parameter model



Lux Cellulose-1

Operating conditions:
 CO_2 -MeOH 90:10 (v/v)
 25°C, 150 bar 3 mL/min

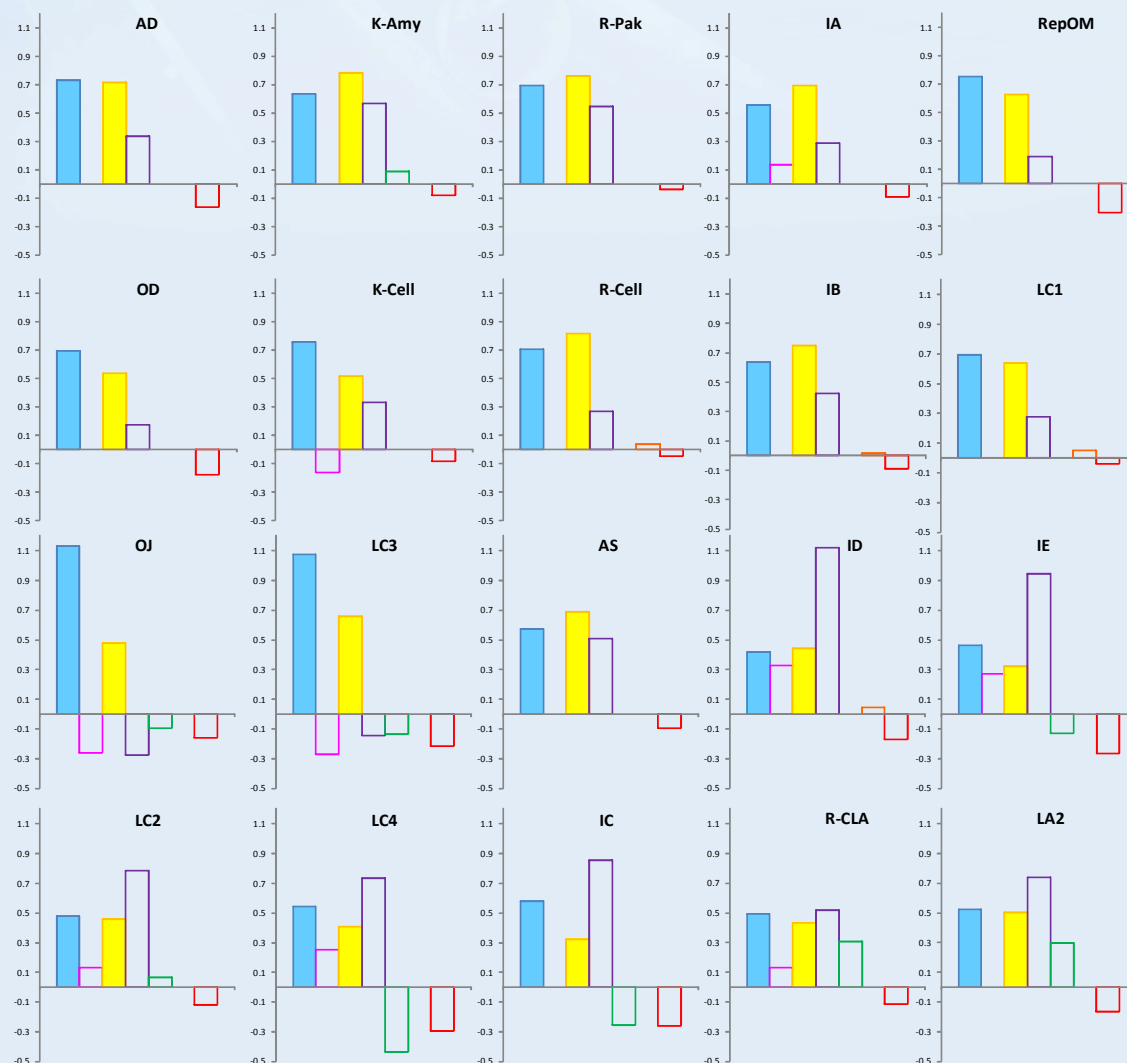
A method to compare retention properties

Modified solvation parameter model



A method to compare retention properties

Modified solvation parameter model

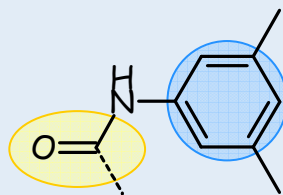


A method to compare retention properties

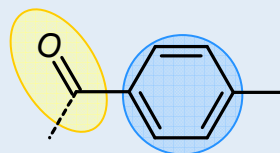
Modified solvation parameter model



*Interactions with
 π and n electrons*



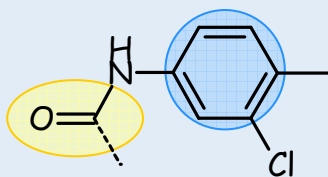
Lux Cellulose-1 (LC1)



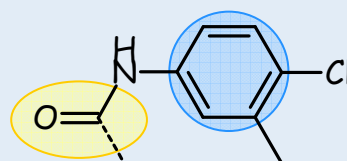
Lux Cellulose-3 (LC3)



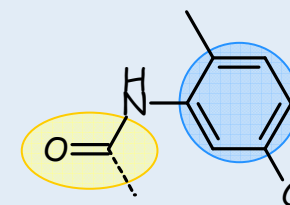
*H-bonding with
proton donors*



Lux Cellulose-2 (LC2)



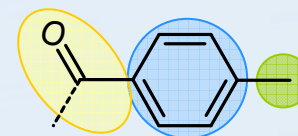
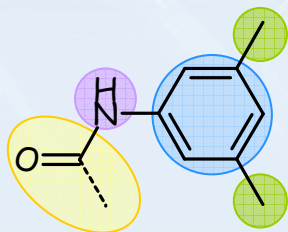
Lux Cellulose-4 (LC4)



Lux Amylose-2 (LA2)

A method to compare stationary phases

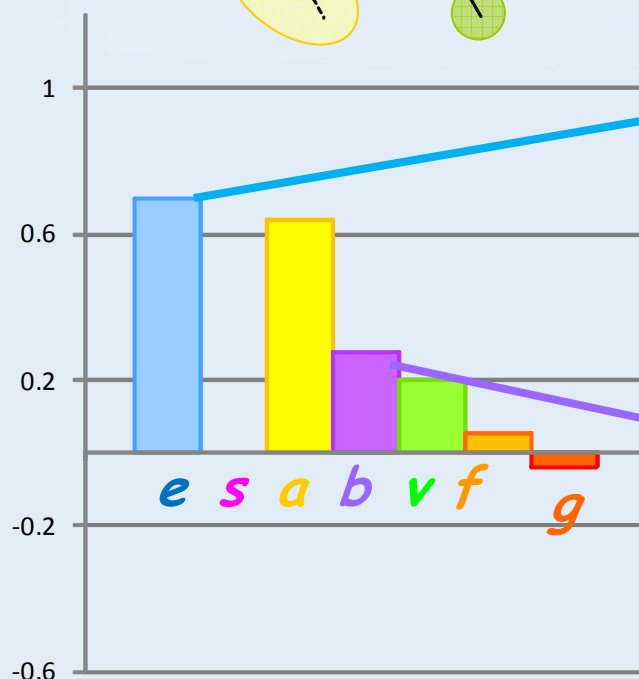
The solvation parameter model



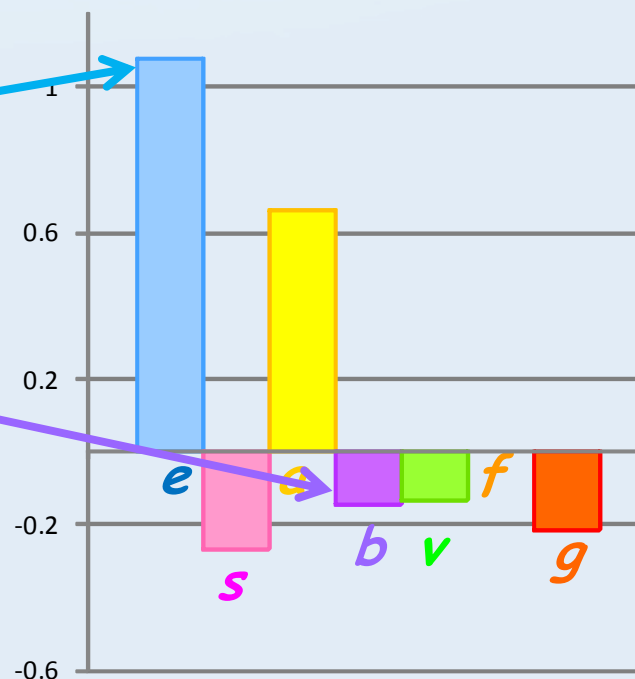
Interactions with π and n electrons



H-bonding with electron donors



Lux Cellulose-1

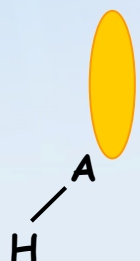


Lux Cellulose-3

Operating conditions:
 CO_2 -MeOH 90:10 (v/v)
 25°C, 150 bar 3 mL/min

A method to compare stationary phases

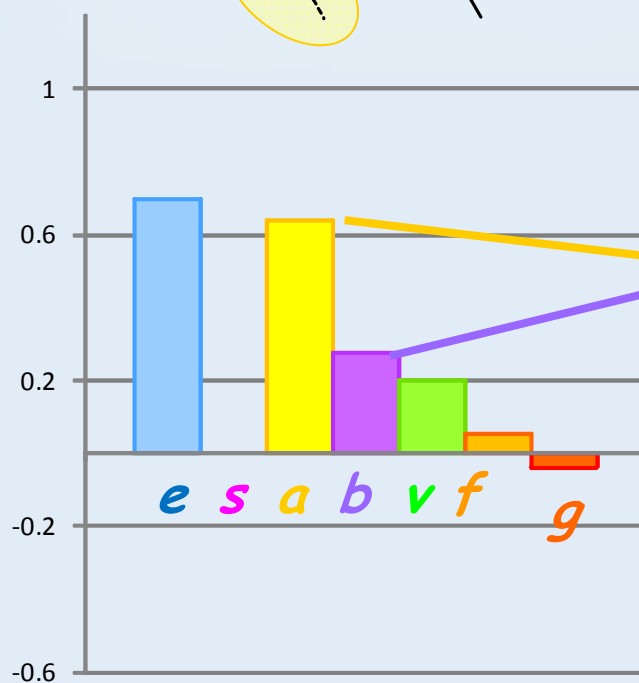
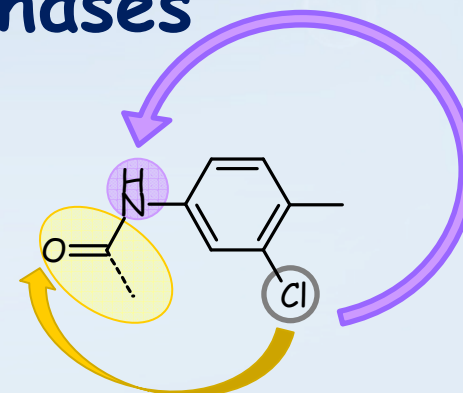
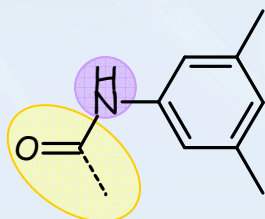
The solvation parameter model



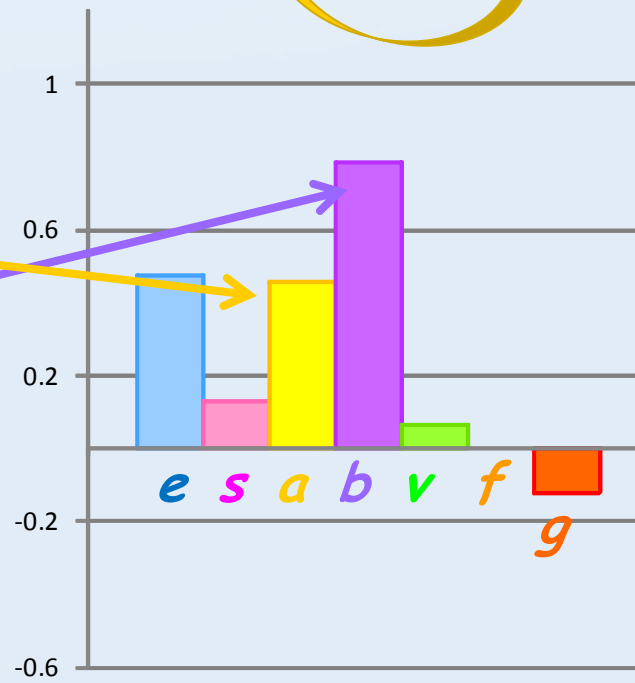
H-bonding with proton donors



H-bonding with electron donors



Lux Cellulose-1



Lux Cellulose-2

Operating conditions:
 CO_2 -MeOH 90:10 (v/v)
 25°C, 150 bar 3 mL/min

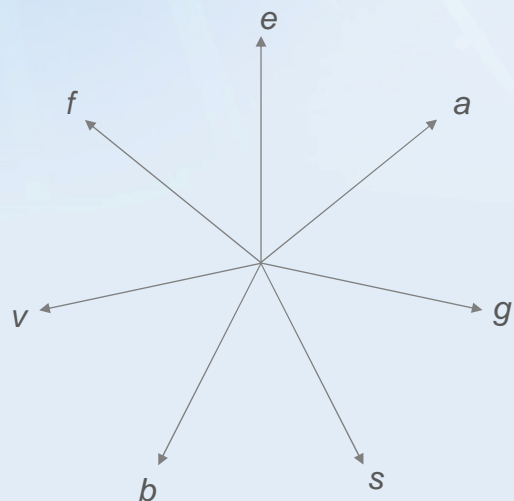
A method to compare retention properties

Modified solvation parameter model




















A method to compare stationary phases

A retention map



OJ   LC3 Ester CSPs

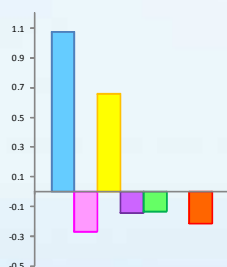
OD  RepOM  LC1  R-Cell 
K-Cell  AD  IB  IA 
K-Amy  AS  R-Pak 

R-CLA  LA2 
IC  LC2  LC4 
Chlorinated CSPs
IC 

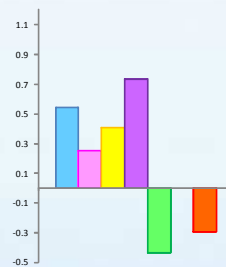
The solvation parameter model As a tool to predict retention

Retention models + molecular descriptors

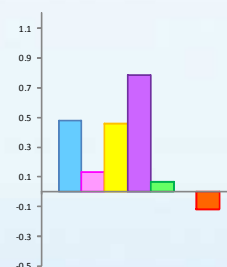
E, S, A, B, V, F, G



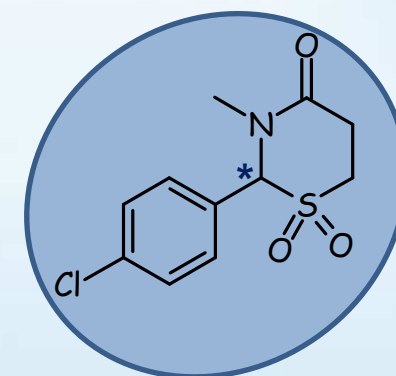
LC3



LC4



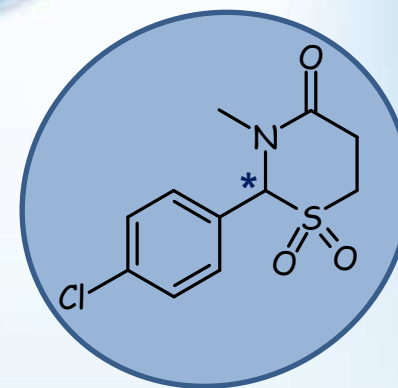
LC2



Chlormezanone

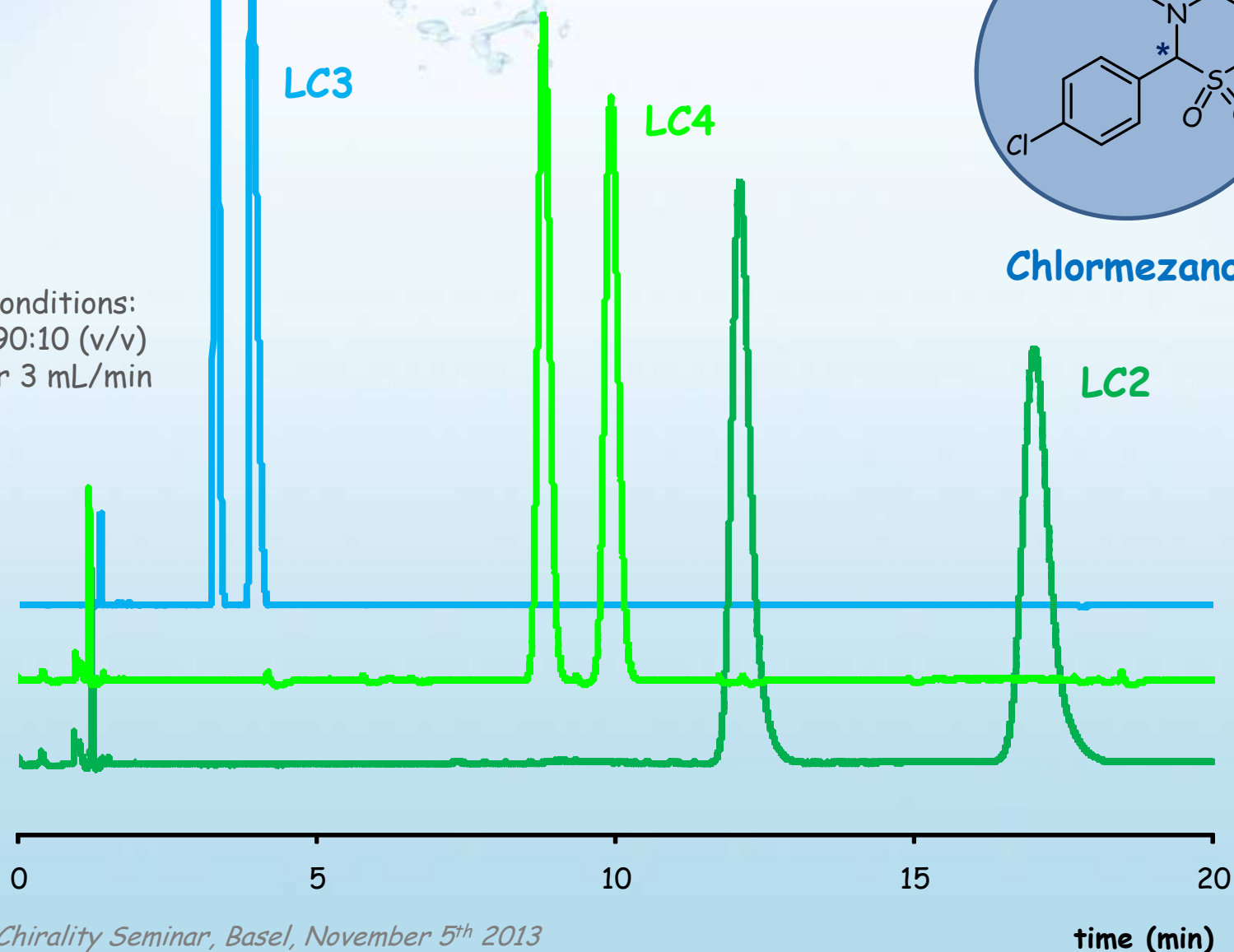
Predicted retention k

The solvation parameter model As a tool to predict retention



Chlormezanone

Operating conditions:
CO₂-MeOH 90:10 (v/v)
25°C, 150 bar 3 mL/min





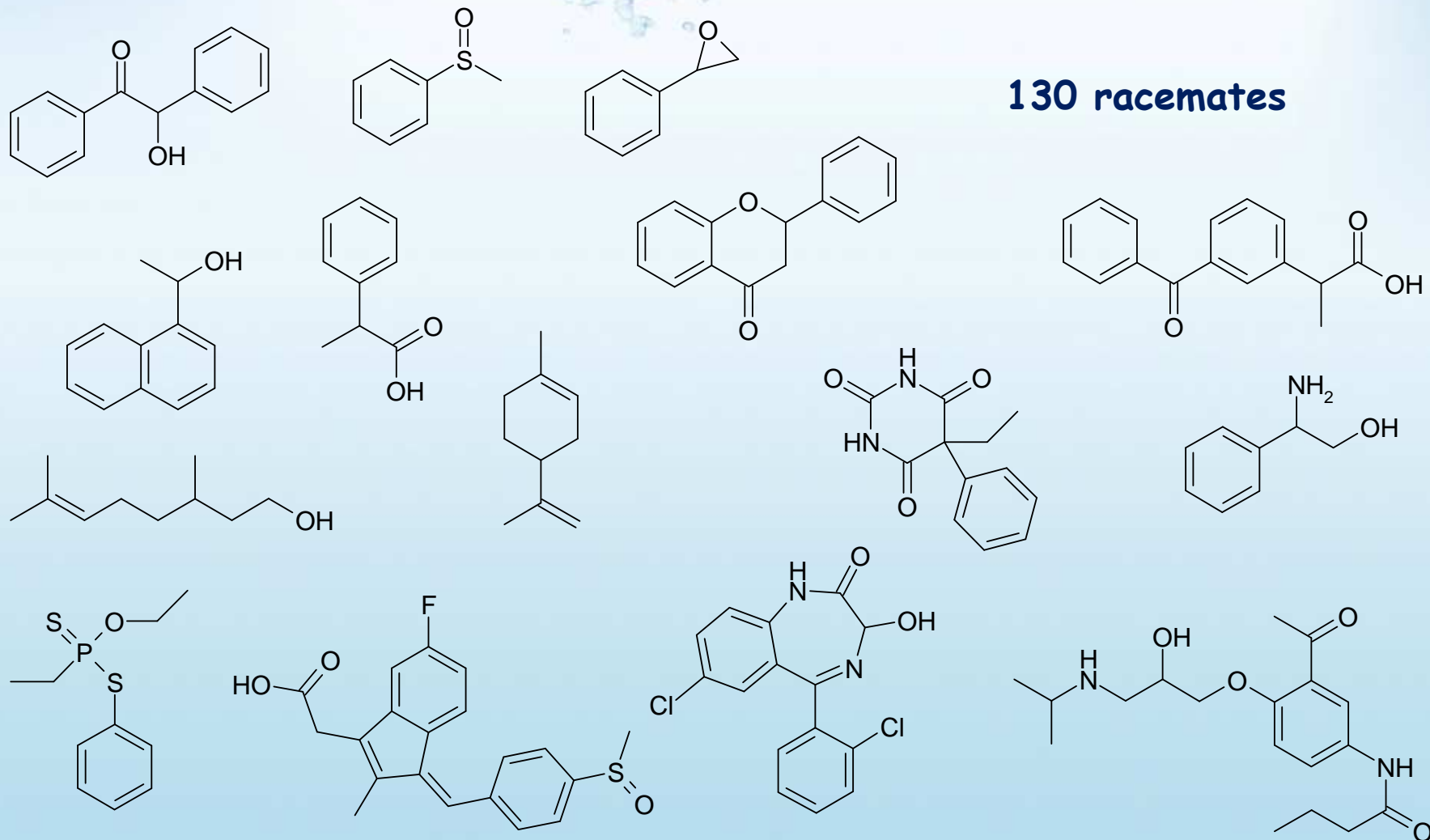
Now what about separation ?

How can we compare the separation capabilities of different CSPs ?

Most frequent method encountered in the literature :
analyze a large group of chiral probe compounds

Compare success rate

A selection of chiral probe analytes



Comparison based on success rate

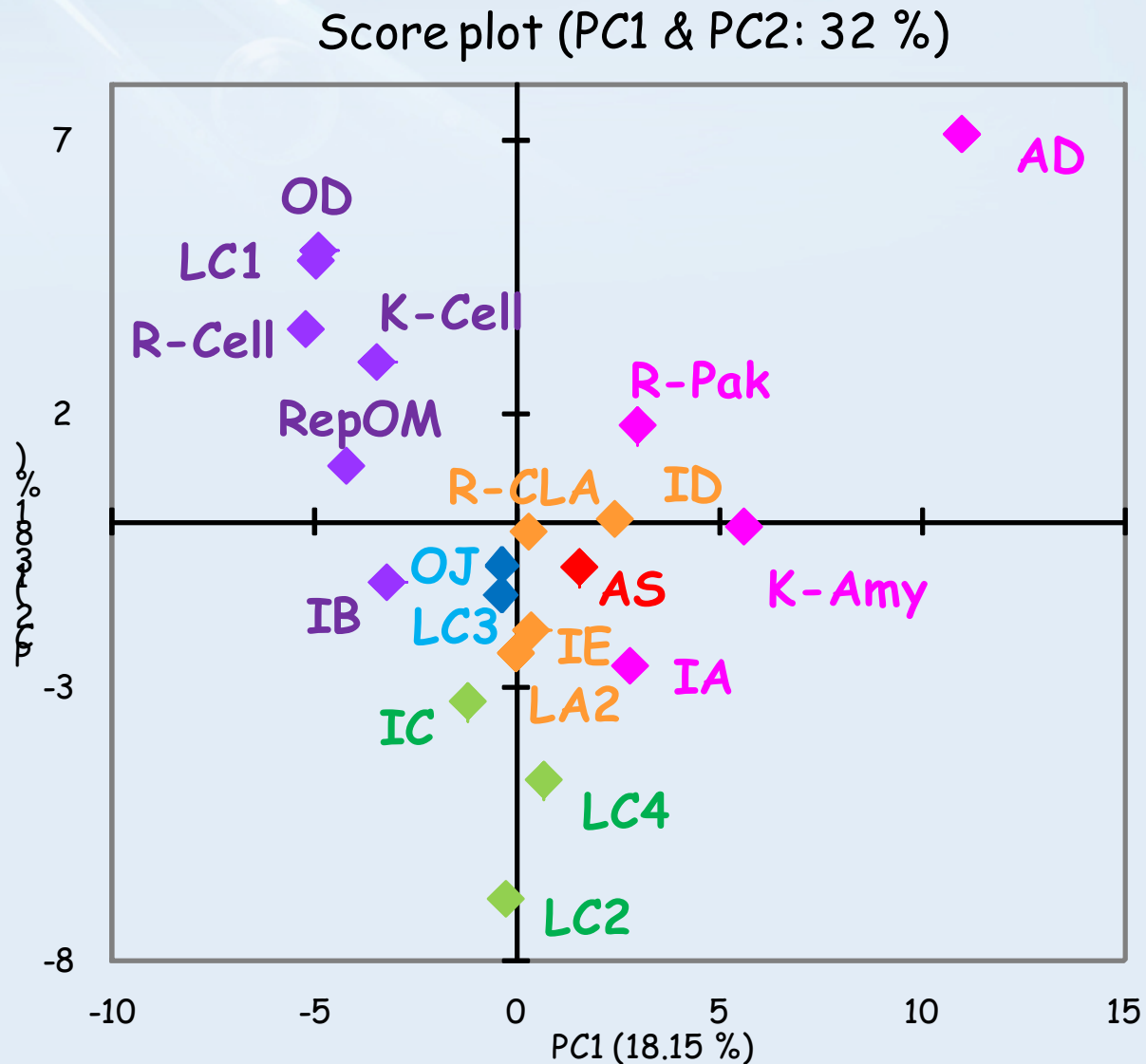
So what ??



Comparison based on similar separations (α)

Principal component analysis furnishes a view of columns based on separation factors

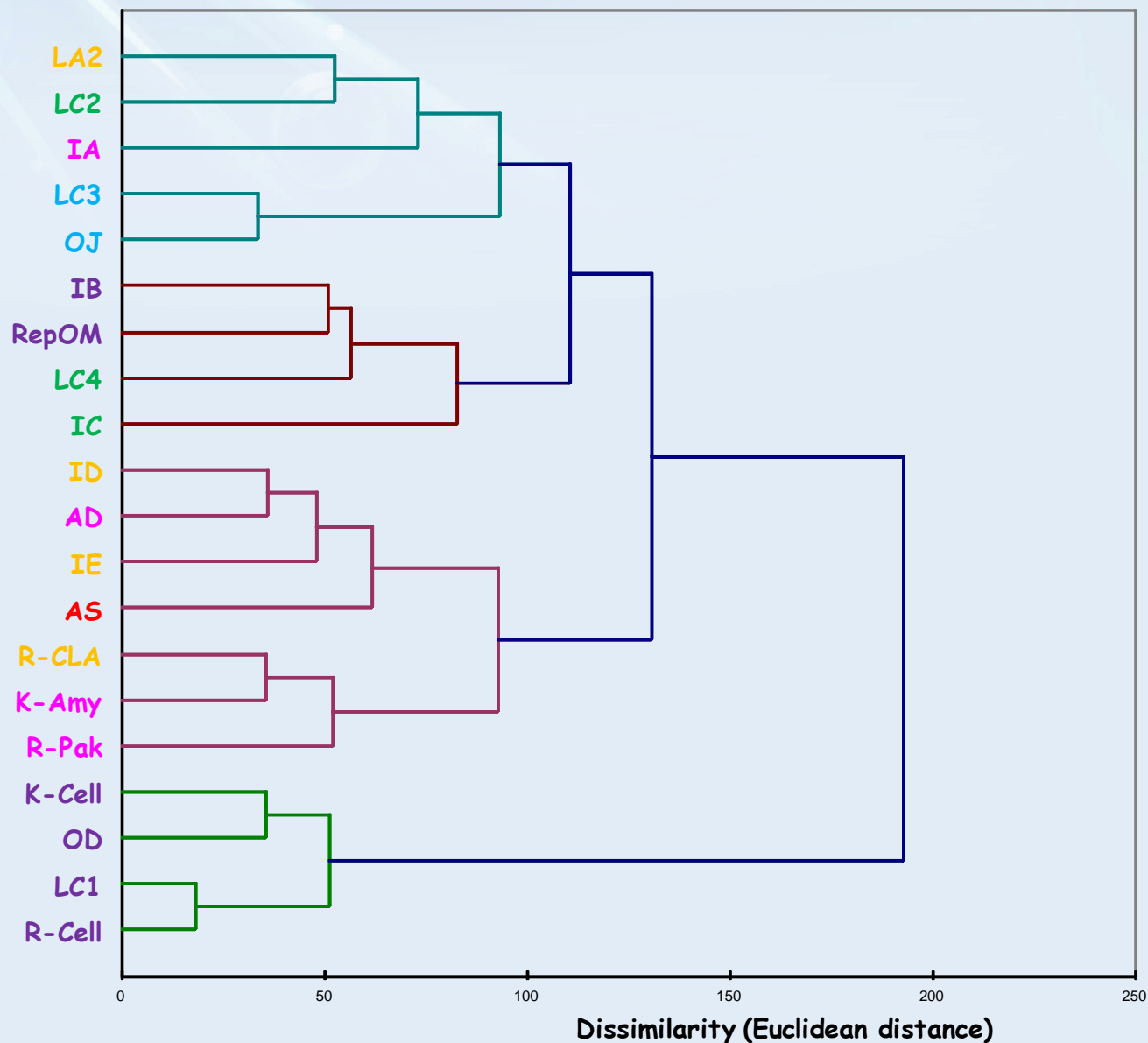
So what ??



Comparison based on similar separations (α)

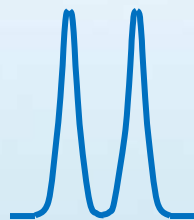
Cluster analysis
furnishes a
classification of
columns based on
separation factors

So what ??



Looking for another (more informative) method

For each chromatographic system, divide the racemates into two classes



Separated



Not separated

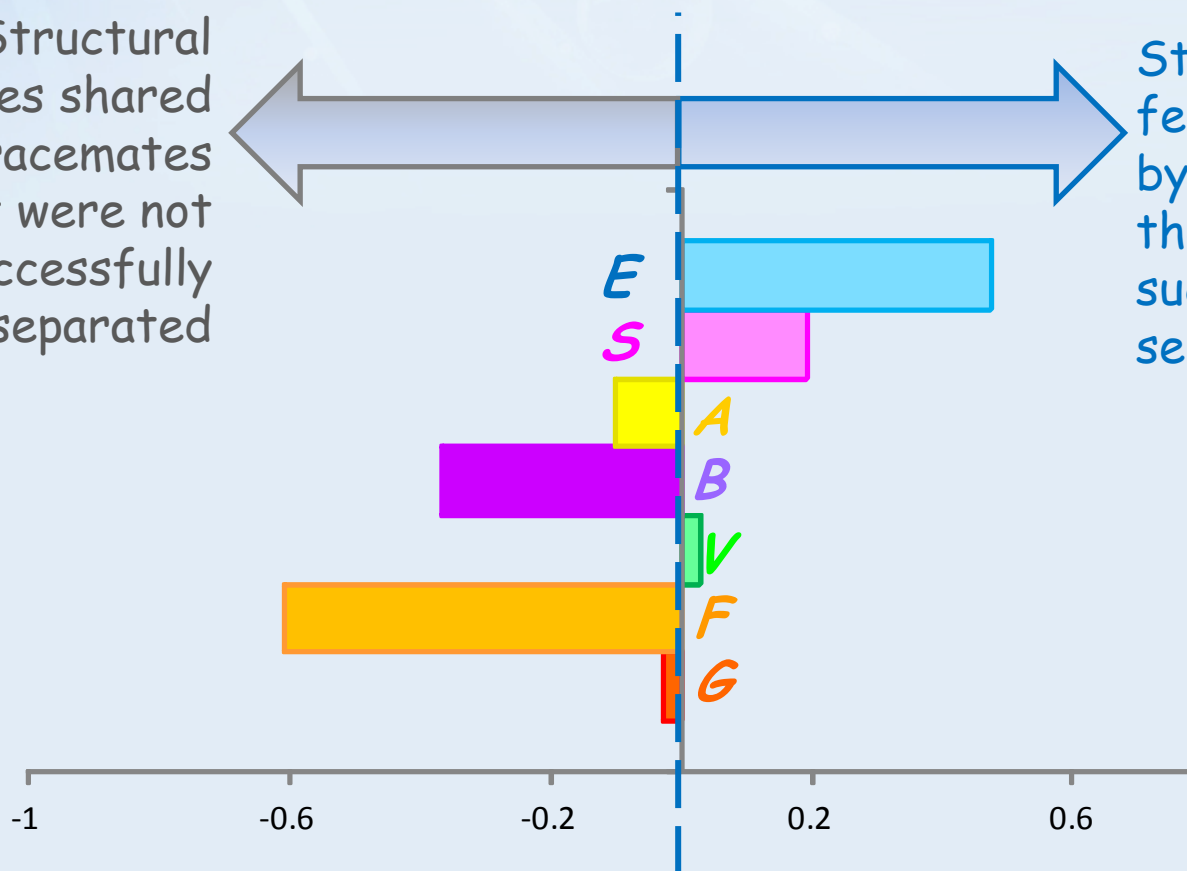
Calculate factorial discriminant analysis to determine the features that are common to the racemates in each class

A method to compare separation capabilities

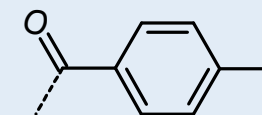
Discriminant analysis

Structural features shared by the racemates that were not successfully separated

Structural features shared by the racemates that were successfully separated

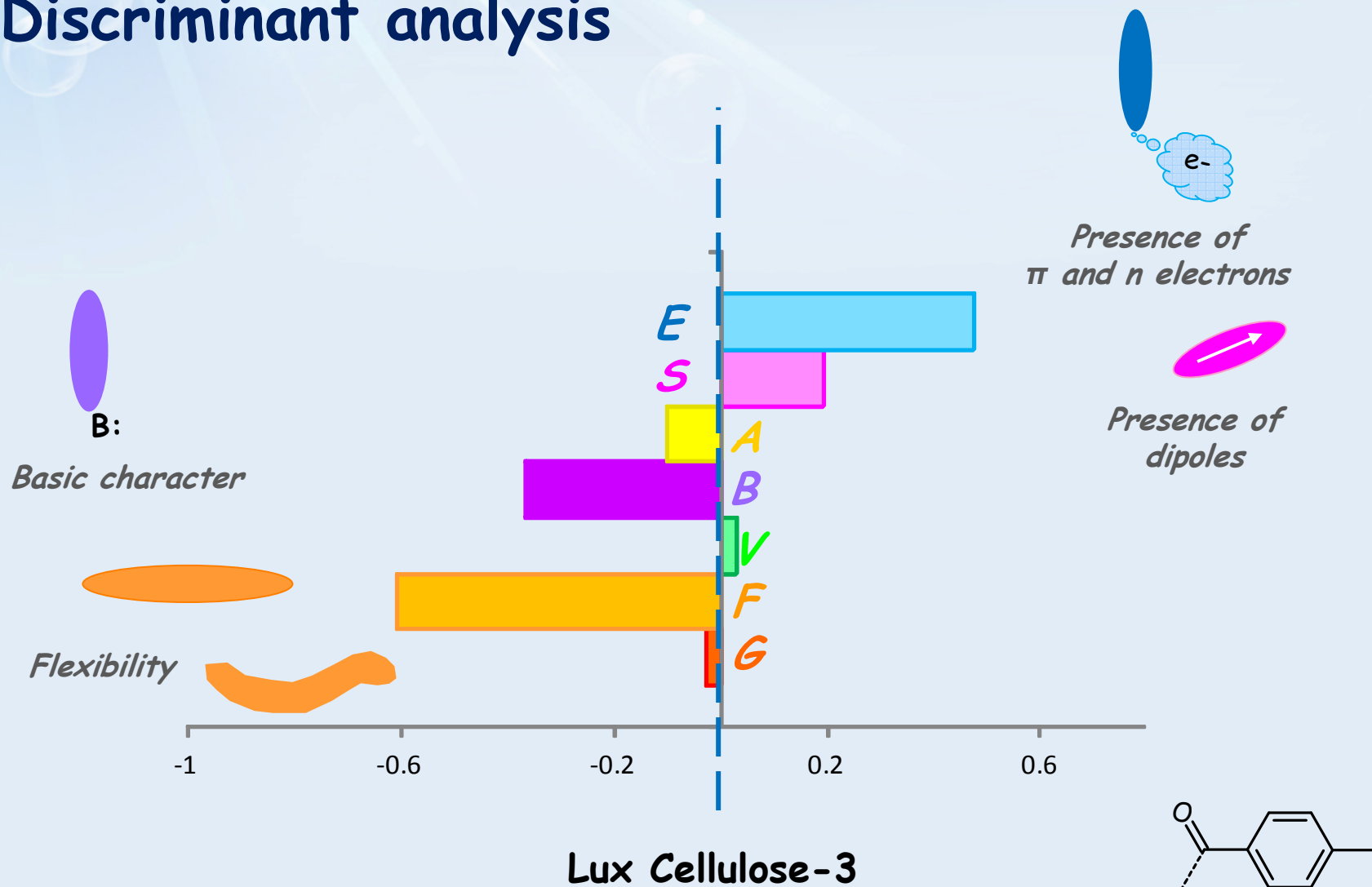


Lux Cellulose-3



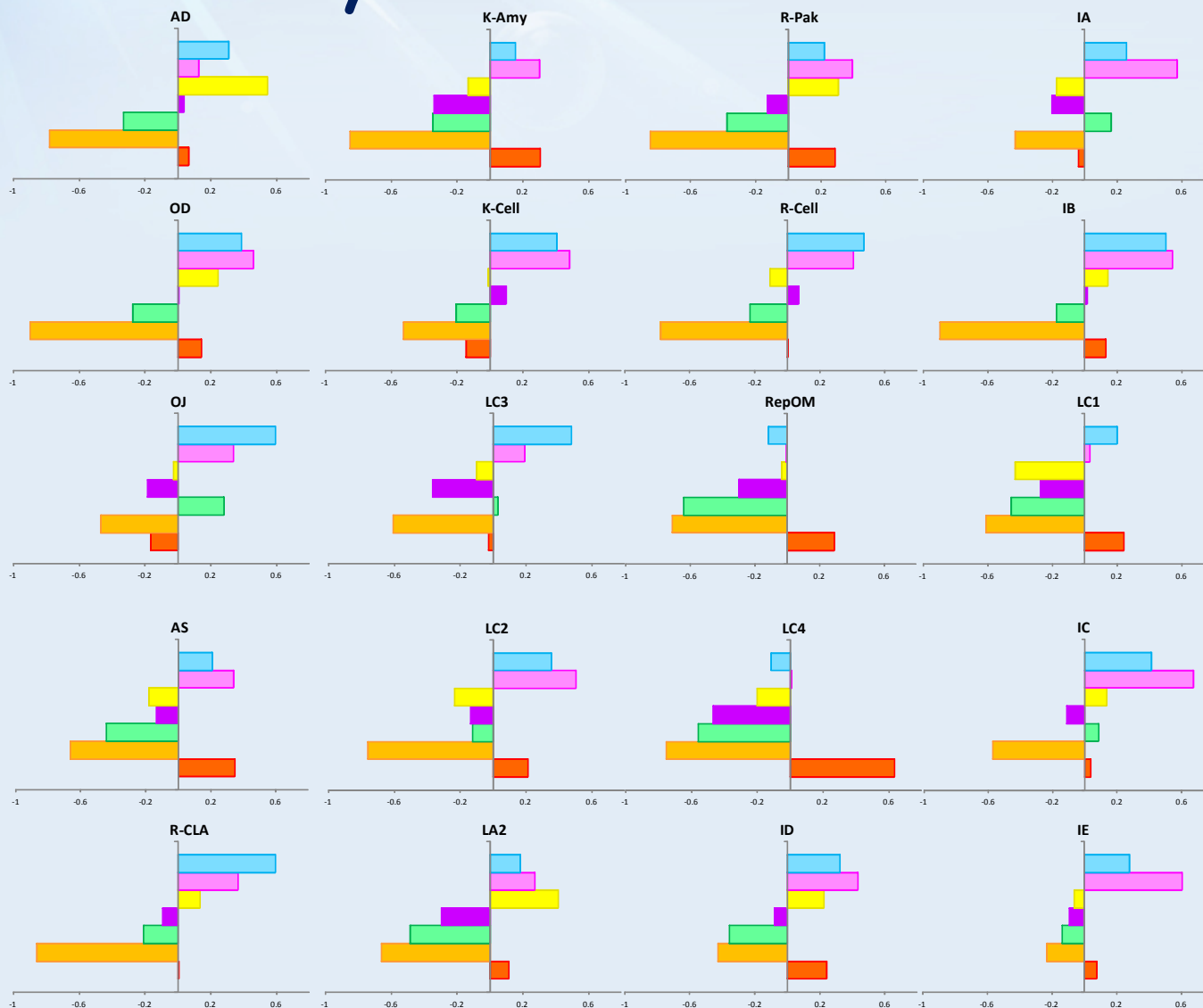
A method to compare separation capabilities

Discriminant analysis



A method to compare separation capabilities

Discriminant analysis



A method to compare separation capabilities

Discriminant analysis

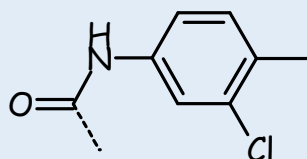
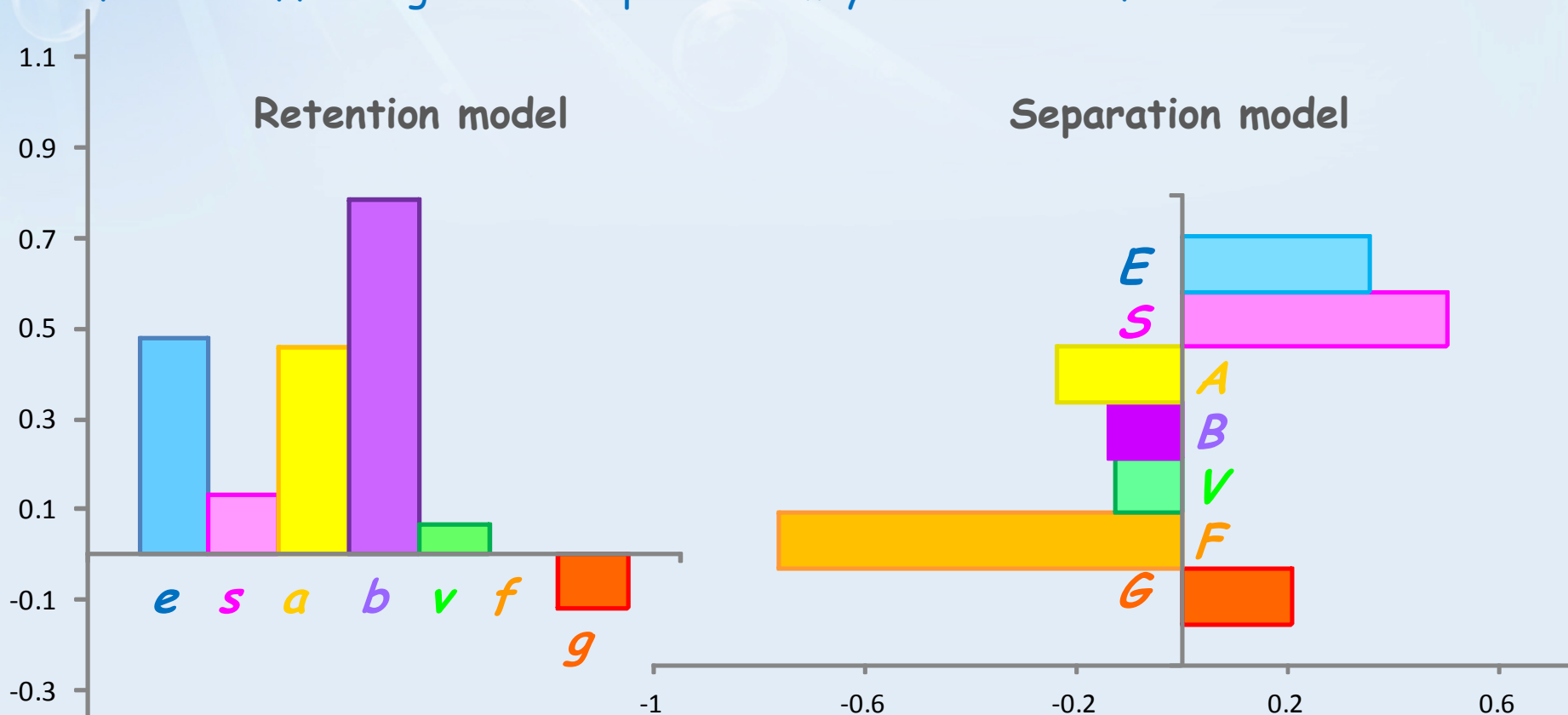


Flexibility



A method to compare separation capabilities

The factors affecting enantioseparation may have little influence on retention

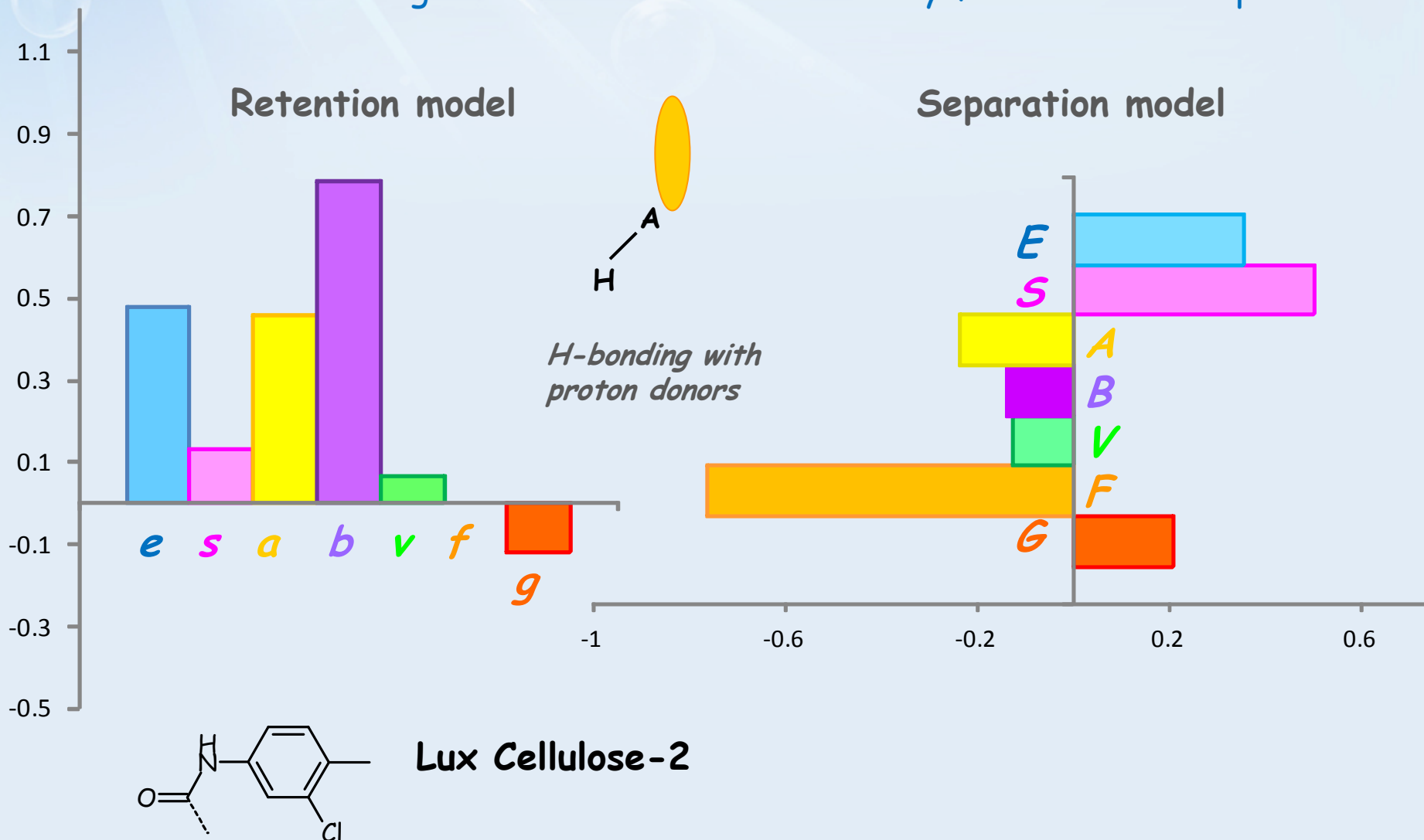


Lux Cellulose-2



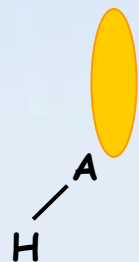
A method to compare stationary phases

The interactions causing retention are not necessarily favourable to separation

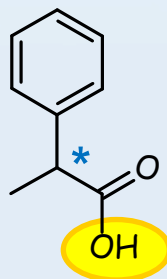


A method to compare stationary phases

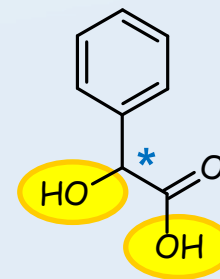
The interactions causing retention are not necessarily favourable to separation



*H-bonding with
proton donors*



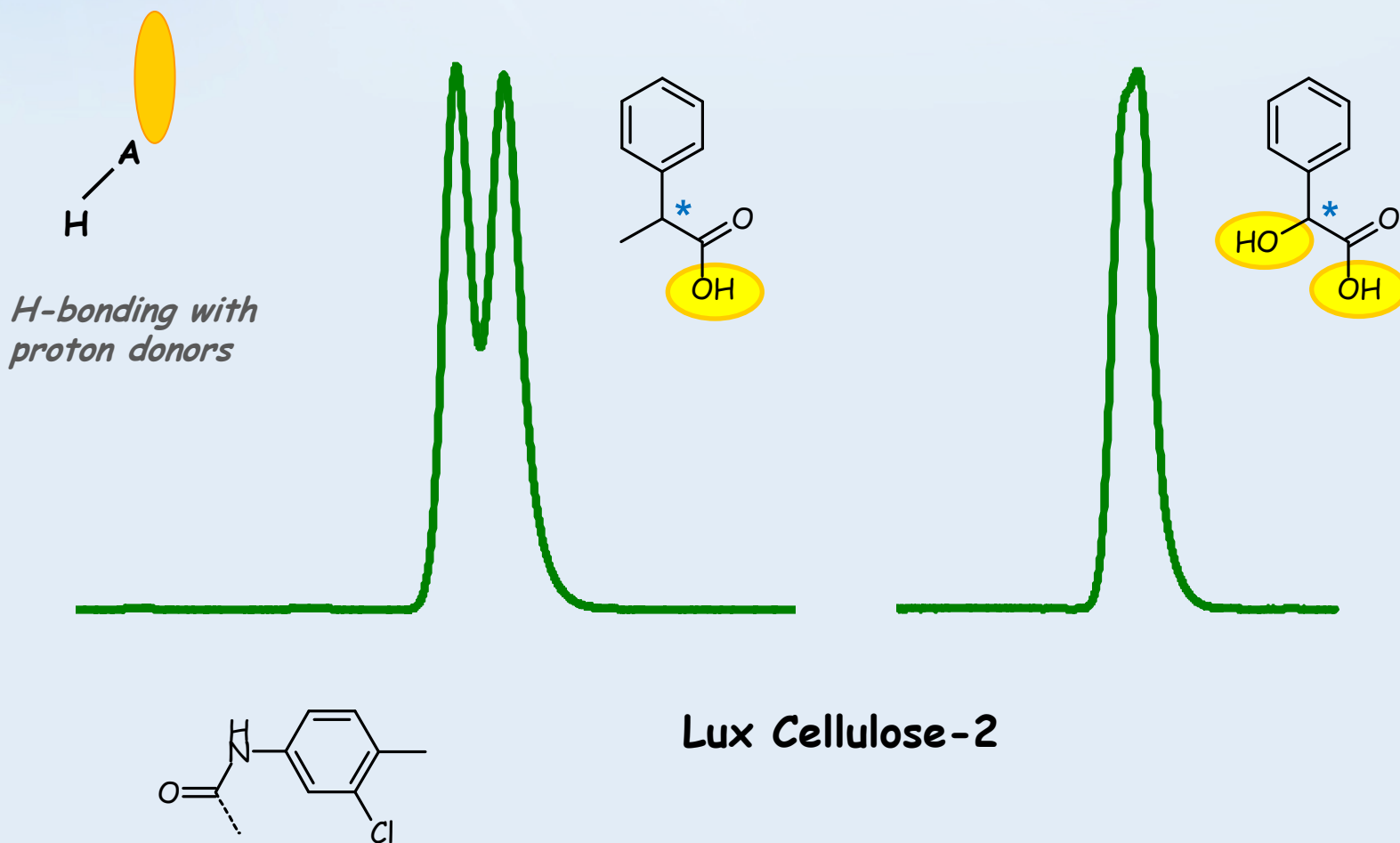
Phenylpropionic acid



Mandelic acid

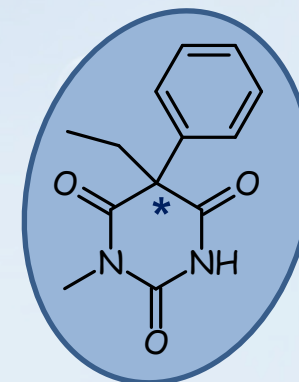
A method to compare stationary phases

The interactions causing retention are not necessarily favourable to separation

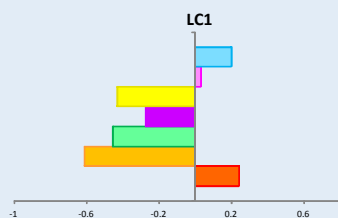


A method to compare stationary phases

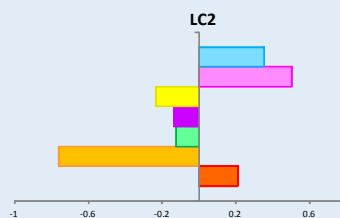
Mephobarbital has little **flexibility** :
rigidity is favourable to separation on all columns



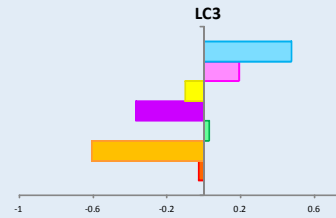
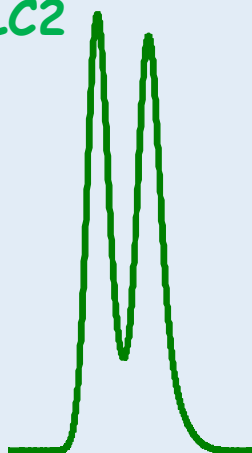
But it is **globular** (not a planar molecule) : unfavourable on LC4



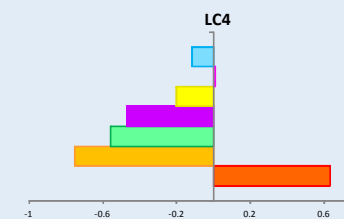
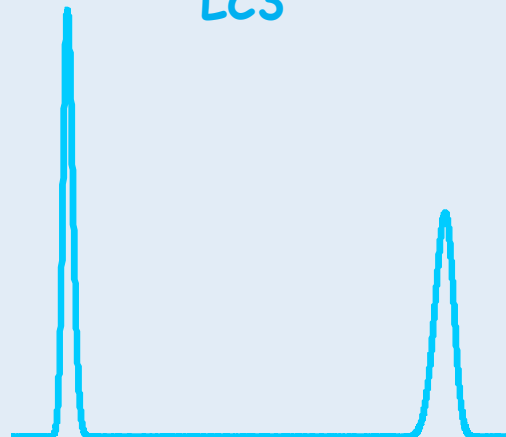
LC1



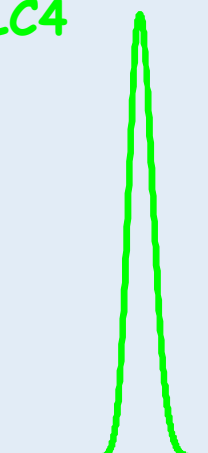
LC2



LC3

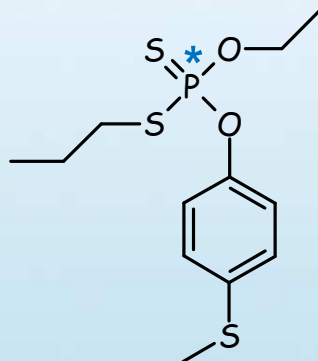


LC4

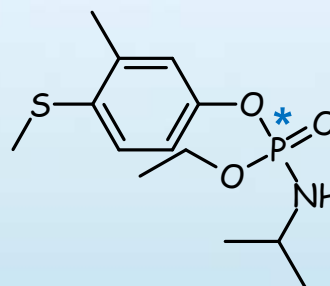


What else can we do ?

Is it possible to use the separation models
to predict the outcome of a separation ?



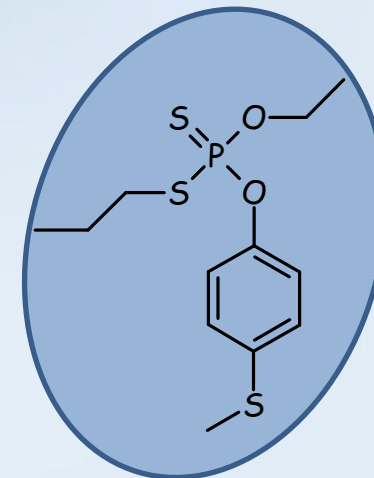
Sulprofos



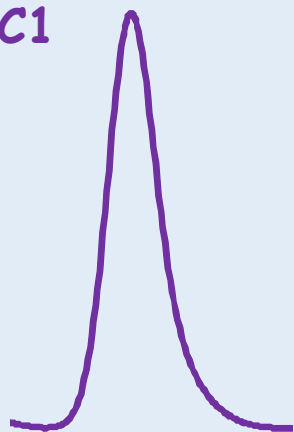
Fenamiphos

Can we predict the outcome of a separation ?

Sulprofos

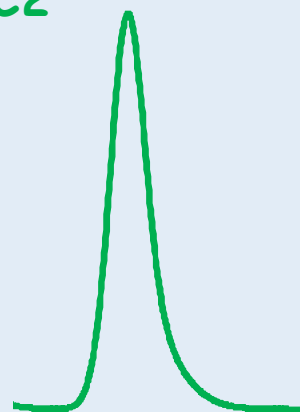


LC1



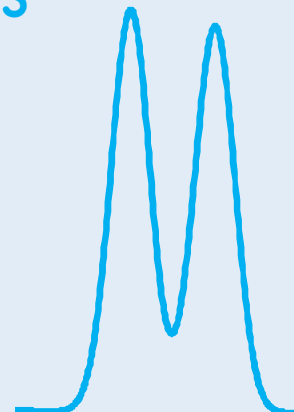
57%

LC2



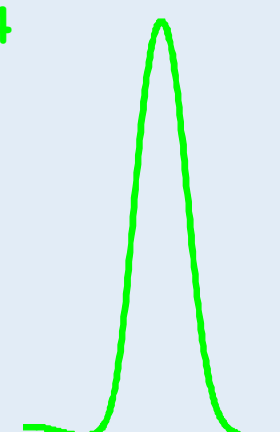
7%

LC3



90%

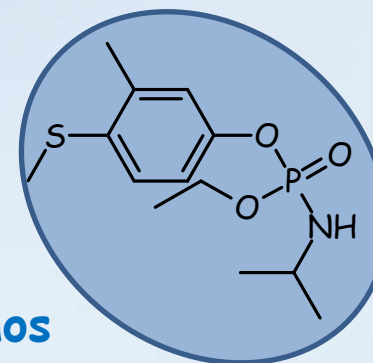
LC4



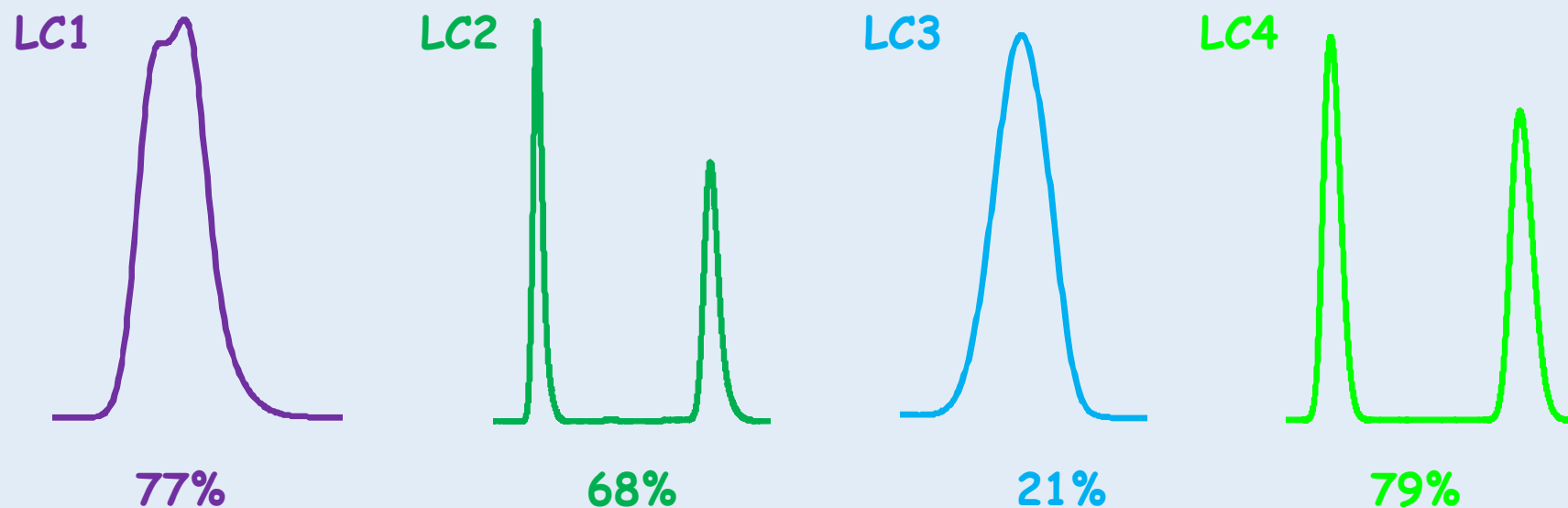
11%

Predicted probability of success on each stationary phase

Can we predict the outcome of a separation ?



Fenamiphos



Predicted probability of success on each stationary phase



What else can we do ?

Is it possible to use the separation models
to predict the outcome of a separation ?

The method is not perfect ($\approx 70\%$ success)
needs some refinement

Molecular descriptors are the weakest link



Conclusions

Multivariate data analyses provide some insight into

Retention mechanisms

Separation mechanisms

Prediction is

Quite good for retention

Possible but not always successful for separation



A bit of reading...

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