



Mechanisms of retention in HPLC

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Part 4



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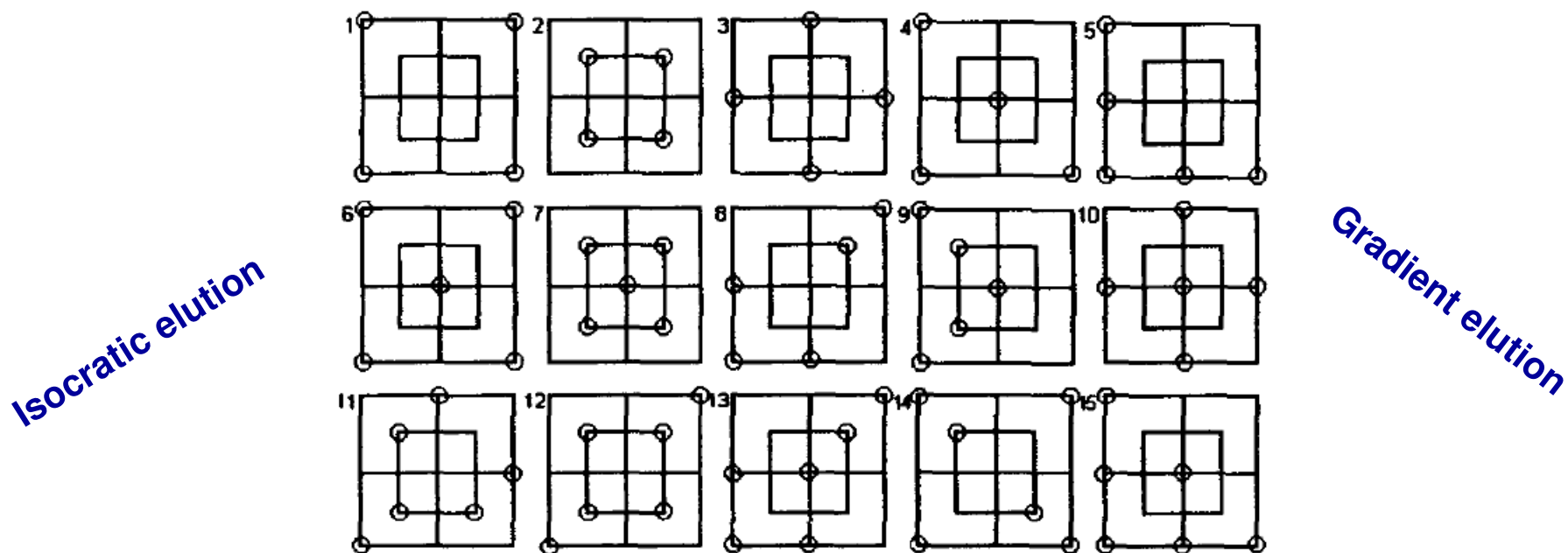
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4.1. Introduction

Interpretive optimisation strategies are based on the **accurate description (modelling) of chromatographic behaviour**. The first step in these strategies consists of **gathering information** about the chromatographic behaviour of the compounds in the sample, covering a reasonably wide factor region usually attending only to retention. For this purpose, either **isocratic or gradient experiments** can be used.



Experimental design : distribution of the experiments within the factor domain



For each solute, a **relationship that describes the retention** is obtained as a function of **the experimental factors**. This function will allow the prediction of **isocratic** or **gradient** retention times at **different conditions**.

Alternatives

- convenient **algebraic expression** (empirical or mechanistic)
parameters searched on a **least-squares basis**
- **black-box algorithm** (as neural networks)

Quality (accuracy and precision) of predictions is **decisive for the reliability** of optimisations or any other applications, and **depends on:**

- **information richness** provided by the training data in the experimental design
- **distribution of the experimental points**
- **equation selected** to fit the training data
- **fitting procedure**
- **ranges of experimental factors**
- **elution mode** (isocratic or gradient)



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4.2. Isocratic elution in RPLC at fixed pH

4.2.1. Use of polynomial models with hydro-organic mobile phases

The **concentration of organic solvent** in the mobile phase is the factor most frequently optimised in RPLC.

Advantages

- **large impact** on both elution strength and selectivity
- a very convenient property:
flexibility and accuracy in the implementation of changes in this factor



Hence, the **importance of obtaining reliable retention models involving the concentration of organic solvent as factor !!!**



RPLC retention (solute i) in terms of solubility parameters (Schoenmakers et al.):

$$\ln k_i = \frac{t_{Ri} - t_0}{t_0} = \frac{v_i}{RT} [(\delta_M - \delta_i)^2 - (\delta_S - \delta_i)^2] + \ln \frac{n_S}{n_M} \quad (4.1)$$

k_i : solute retention factor

t_{Ri} : solute retention time

t_0 : dead time

R : gas constant (1.9865 cal K⁻¹ mole⁻¹)

T : absolute temperature (K)

v_i : solute molar volume (cm³ mole⁻¹)

δ : polarity parameter (cal^{0.5} cm^{-1.5})

n_M : moles of mobile phase

n_S : moles of stationary phase

Water (w) / **organic solvent** (o) binary mixtures:

$$\delta_M = \sum_j \delta_j \varphi_j \quad (1.2)$$



$$\delta_M = (1 - \varphi) \delta_w + \varphi \delta_o \quad (4.2)$$

φ : organic solvent volume fraction



$$\ln k_i = \frac{t_{Ri} - t_0}{t_0} = \frac{v_i}{RT} [(\delta_M - \delta_i)^2 - (\delta_S - \delta_i)^2] + \ln \frac{n_S}{n_M}$$

$$\delta_M = (1 - \varphi) \delta_w + \varphi \delta_o$$

$$\ln k = c_0 + c_1 \varphi + c_2 \varphi^2 \quad (4.3)$$

Quadratic model

Parameters c_0 , c_1 and c_2 gather all the constants and solute/mobile phase/stationary phase properties, and have **particular values** for a **given solute** and **column/solvent system**.

decimal logarithms

$$\log k = c_0 + c_1 \varphi + c_2 \varphi^2 \quad (4.4)$$

$$\log k = c_0 + c_1 \varphi \quad (4.5)$$

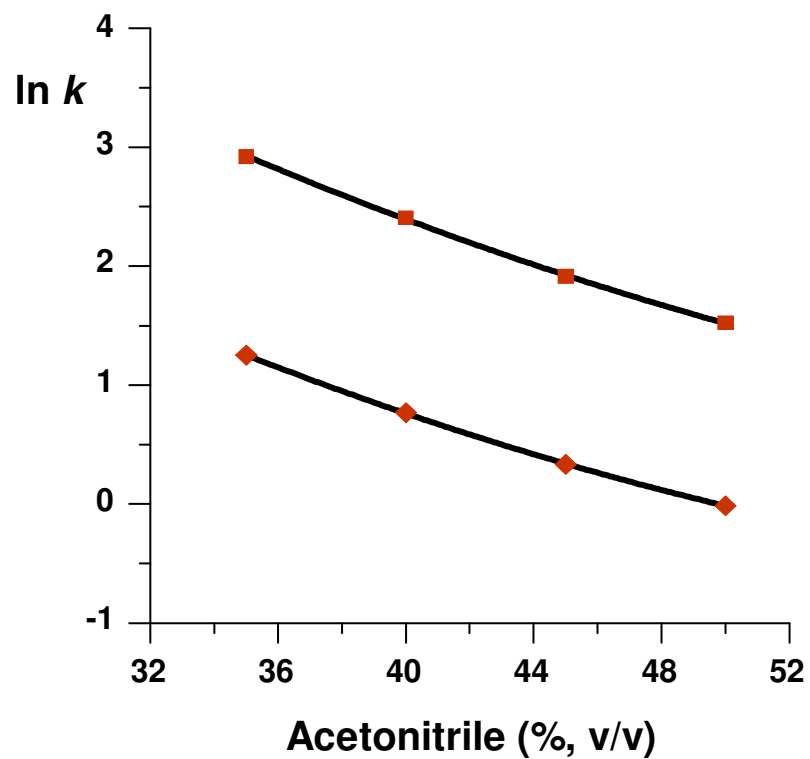
linear model

narrow concentration ranges

These equations are also valid in the presence of specific interactions in the **presence of additives**.



Two flavonoids eluted with acetonitrile/water mixtures



$$\log k = c_0 + c_1 \varphi + c_2 \varphi^2$$

$$\log k = c_0 + c_1 \varphi$$



Linear model

$$\log k = c_0 + c_1 \varphi = \log k_w - S \varphi \quad (4.5)$$

c_0 (intercept): extrapolated $\log k$ value if **water** were the mobile phase

S (slope): sensitivity of retention to changes in the organic solvent content (**elution strength**)

- **Simplifies** the retention model to a **straight-line**.
- Valid for **narrow concentrations of organic solvent**, which is usual in RPLC.
- In wide concentration ranges, **deviations from linearity** are observed.
- The deviations are especially **significant** at the **highest and lowest concentrations** of organic solvent.



Ternary systems (two organic solvents and water)

$$\log k = c_0 + c_1 \varphi_1 + c_2 \varphi_2 + c_{12} \varphi_1 \varphi_2 + c_{11} \varphi_1^2 + c_{22} \varphi_2^2 \quad (4.6)$$

φ_1, φ_2 : concentrations of organic solvents

The search for an appropriate model has not finished here !!!

- Review Nikitas and Papa-Louisi (J. Chromatogr. 1216 (2009) 1737–1755)
- Most models are logarithmic, but there are other proposals.
- Model proposed by Jandera for NPLC, applied to RPLC:

excellent accuracy in RPLC $\longrightarrow$ $\frac{1}{k} = [a + b\varphi]^n \quad (4.7)$



More retention models !!!

$$k = e^{\ln k_w - a\varphi + b\varphi^2}$$

$$k = e^{\ln k_0 (1 - a\varphi(1 - \varphi))}$$

$$k = e^{\ln k_w - a \frac{\varphi}{1 + b\varphi}}$$

$$k = e^{\ln k_0 - \frac{a\varphi}{1 + 2\varphi}}$$

$$k = e^{\ln k_w - \ln(1 + b\varphi) - c \frac{\varphi}{1 + b\varphi}}$$

$$k = e^{\ln k_w - \ln(1 + 2\varphi) - c \frac{\varphi}{1 + 2\varphi}}$$

$$k = \frac{e^{\ln k_0}}{(1 + b\varphi)^4}$$

$$k = \frac{e^{\ln k_0}}{1 + a\varphi + b\varphi^2}$$

$$k = \frac{e^{\ln k_0}}{(1 + b\varphi)^3}$$

$$k = e^{\ln k_0} \frac{1 - a\varphi}{1 + b\varphi}$$

$$k = \frac{e^{\ln k_0}}{1 + b\varphi(1 + b\varphi)}$$

$$k = e^{\ln k_0 - a\varphi}$$

$$k = e^{\ln k_0 - a \ln \varphi / \varphi_0}$$



4.2.2. Polarity models

Linear relationships between **log k** and **mobile phase polarity parameters** that depend on **solvent content** are interesting models.

- Dimroth-Reichardt parameter $E_T(30)$
- Normalised $E_T^N(30)$

limited range of linearity : 0 – 80% acetonitrile and 20 – 100% methanol

- Normalised polarity parameter proposed by Bosch and Rosés P_M^N

extends the linearity to 0 – 100% acetonitrile and methanol

For **acetonitrile-water**:

$$P_M^N = 1.00 - \frac{2.068\varphi}{1 + 1.341\varphi} \quad (4.8)$$

For **methanol-water**:

$$P_M^N = 1.00 - \frac{1.33\varphi}{1 + 0.47\varphi} \quad (4.9)$$



A model was developed, which isolates the **polarity contributions** of the three agents involved in the separation system (**solute, stationary phase and mobile phase**).

$$\log k = (\log k)_0 + p_S (P_M^N - P_S^N) \quad (4.10)$$

Descriptors

P_M^N : mobile phase polarity

P_S^N : stationary phase polarity

$(\log k)_0$: retention in a hypothetical mobile phase
with the same polarity as the stationary phase

p_S : solute polarity

The **separation of the polarities of solute, mobile phase and stationary phase in the polarity model is not perfect**: p_S mainly gathers the solute contributions. However, it is not an absolute value, but a relative measurement that **depends on the column environment** (stationary and mobile phase nature).



$$\log k = (\log k)_0 + p_s (P_M^N - P_S^N) \quad (4.10)$$

fitted with the data
for a **set of compounds**

Advantages of the polarity model

- **A simple model**: only needs knowledge of

one parameter per solute: p_s

two descriptors for the column ($(\log k)_0$ and P_S^N)

P_M^N is obtained from the mobile phase composition

- This facilitates **transference of k values to new columns and solvent systems** by establishing simple correlations between p_s -values for **small sets of selected reference compounds**.



- When $(\log k)_0$ and P_S^N are fitted **individually for each solute**:

$$\log k = q + p_s P_m^N = c_0 + c_1 P_m^N \quad (4.11)$$

$$\log k = (\log k)_0 + p_s (P_M^N - P_S^N) \quad (4.10)$$

Predictions performed by **imposing common column parameters** are necessarily poorer than those when all parameters in the equation are fitted for each individual solute, which are preferable for optimisation purposes !!!



$$\log k = c_0 + c_1 \varphi$$

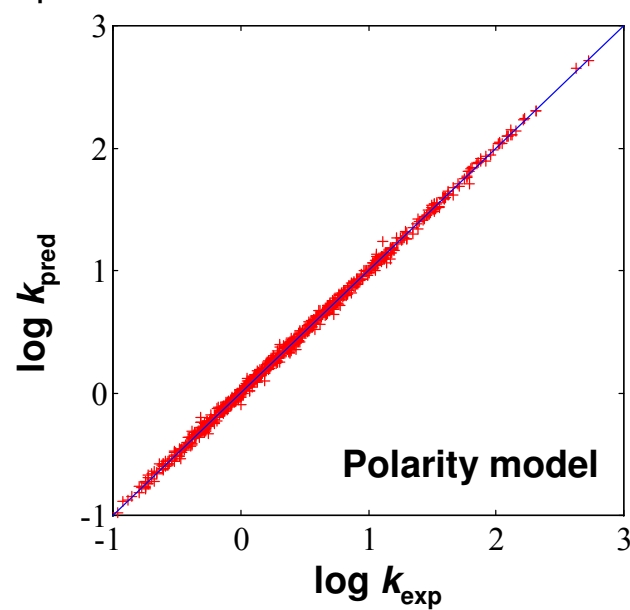
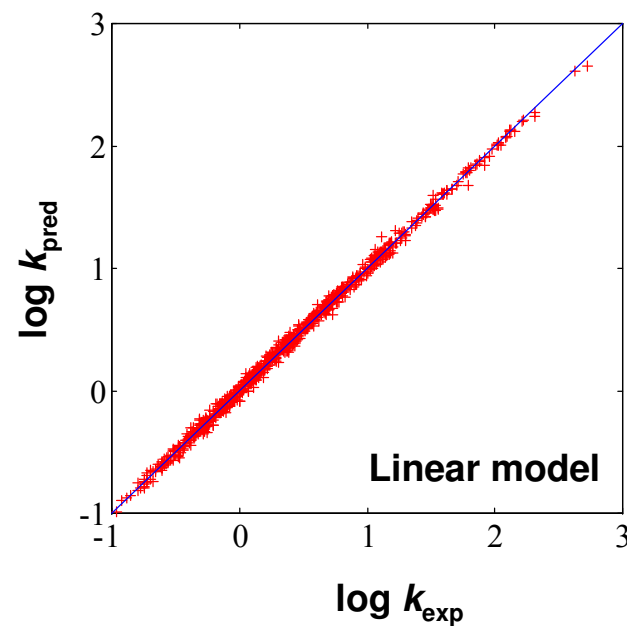
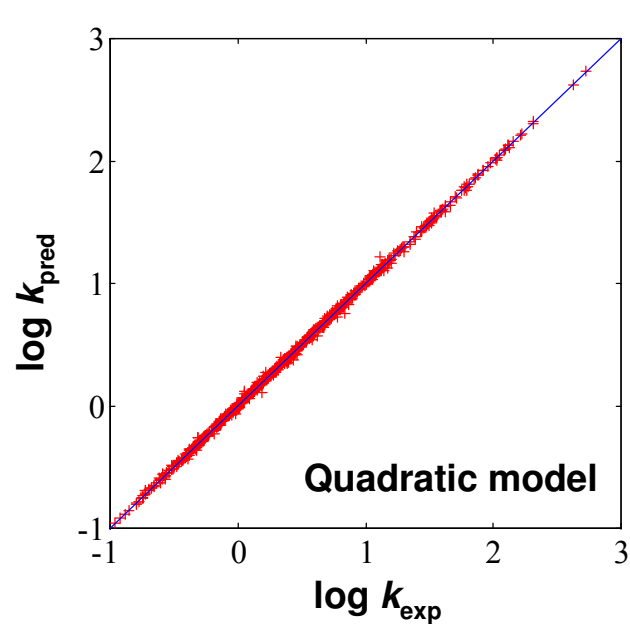
$$\log k = c_0 + c_1 P_M^N$$

two-parameter models

$$\log k = c_0 + c_1 \varphi + c_2 \varphi^2$$

three-parameter model

The two-parameter polarity model is valid for **wider concentration ranges** and **comparable in accuracy to the quadratic model !!!**

**152 compounds eluted with methanol / water**



4.2.3. Micelles and ionic liquids in the mobile phase

Among the **chromatographic modes using mobile phases with additives**, **micellar liquid chromatography** deserves a special mention.

Modelling

- **Polynomial** equations
- **Mechanistic** models based on equilibrium considerations
 - ➡ gave a meaning to the coefficients in the polynomial equations
 - ➡ improved understanding of the retention mechanism

Pure MLC (mobile phases containing only surfactant):

$$k = \frac{K_{AS}}{1 + K_{AM} [M]} \quad (4.12)$$

$$\frac{1}{k} = \frac{1}{K_{AS}} + \frac{K_{AM}}{K_{AS}} [M] = c_0 + c_1 [M] \quad (4.13)$$

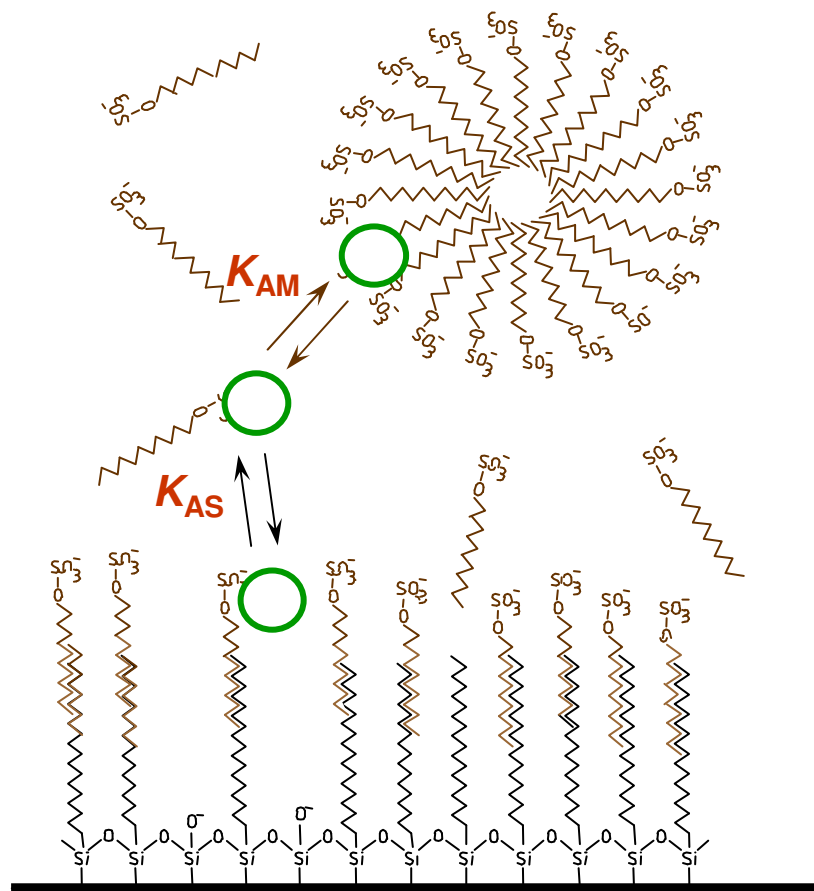
[M] : concentration of surfactant molecules involved in micelle formation

K_{AS} : partition of the solute between bulk water and stationary phase

K_{AM} : partition of the solute between bulk water and micelle



Pure MLC





Hybrid MLC

Hybrid micellar medium (mobile phases containing both surfactant and organic solvent):

$$k = \frac{K_{AS} \frac{1 + K_{SD} \varphi}{1 + K_{AD} \varphi}}{1 + K_{AM} \frac{1 + K_{MD} \varphi}{1 + K_{AD} \varphi}} [M] \quad (4.14)$$

K_{SD} , K_{AD} and K_{MD} : account for the displacement of the partition equilibria by the organic solvent

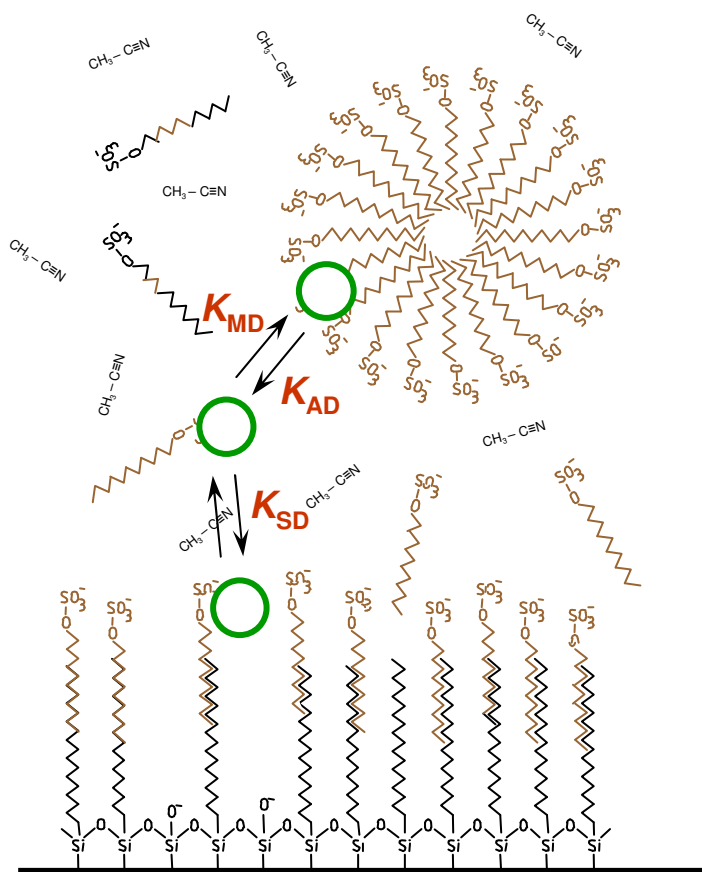
K_{SD} only for highly hydrophobic solutes. In other cases:

$$k = \frac{K_{AS} \frac{1}{1 + K_{AD} \varphi}}{1 + K_{AM} \frac{1 + K_{MD} \varphi}{1 + K_{AD} \varphi}} [M] \quad \longrightarrow \quad \log k = c_0 + c_1 \varphi + c_2 [M] + c_{12} \varphi [M] \quad (4.15)$$

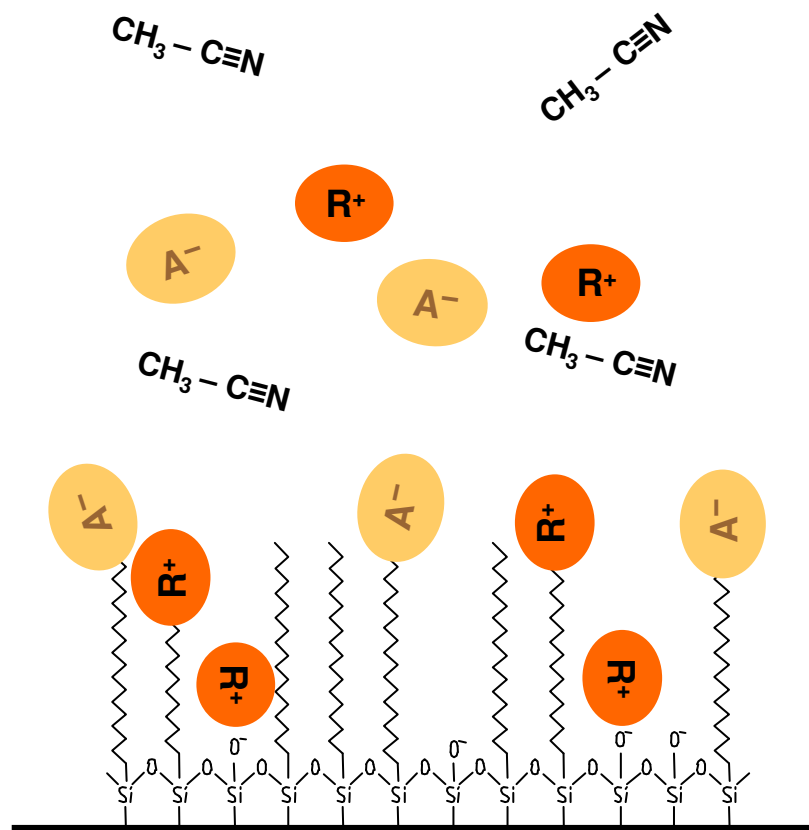
Eq. (4.15) is also valid for **mobile phases containing ionic liquids**, where the molar concentration of ionic liquid $[IL]$ is used instead of $[M]$.



MLC



RPLC with ionic liquids





4.2.4. The silanol effect

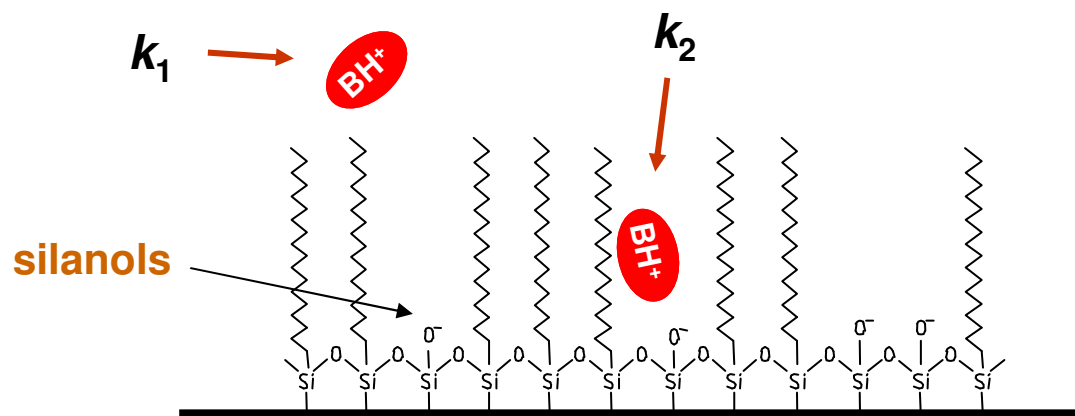
Basic compounds experience a combined retention mechanism when separated with bonded-silica stationary phases and organic solvent-water mixtures that includes **hydrophobic** (partition into the bonded phase), and **silanophilic interactions** (attraction of the cationic solutes to the anionic silanols). In the absence of any additional interaction, masking of silanols will decrease the retention.

$$k_0 = k_1 + k_2 \quad (4.16)$$

k_0 : retention factor with silanols totally exposed for interaction

k_1 : hydrophobic contribution

k_2 : silanophilic contributions





Horváth model

Evaluation of the silanol suppressing potency by addition of amines

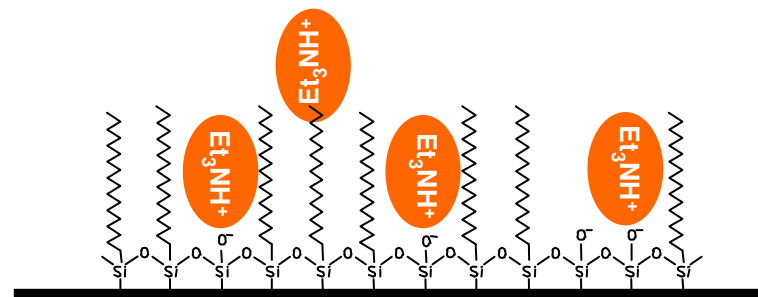
without additive $\longrightarrow k_0 = k_1 + k_2$ (4.16)



with additive $\longrightarrow k = k_1 + \frac{k_2}{1 + K_A [A]}$ (4.18)

K_A : silanol / protonated amine binding constant

$[A]$: amine molar concentration



Silanophilic contribution to retention

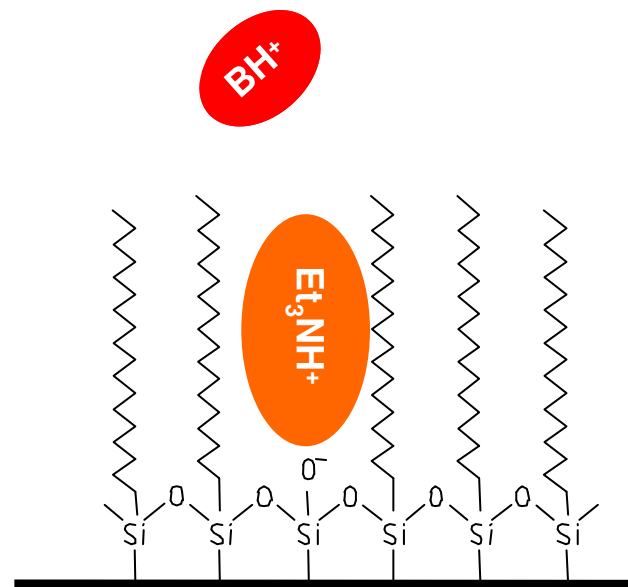
$$k_0 - k = k_2 - \frac{k_2}{1 + K_A [A]} = \frac{k_2 K_A [A]}{1 + K_A [A]} \quad (4.19)$$



$$\frac{[A]}{k_0 - k} = \frac{1}{k_2 K_A} + \frac{[A]}{k_2} \quad (4.20)$$



$$\frac{[A]}{k_0 - k} = \frac{1}{k_2 K_A} + \frac{[A]}{k_2} \quad (4.20)$$



The binding constant K_A :

- evaluates the **ability** of the amine to **block silanol sites**
- can be obtained by **regressing the left term versus the concentration of amine**



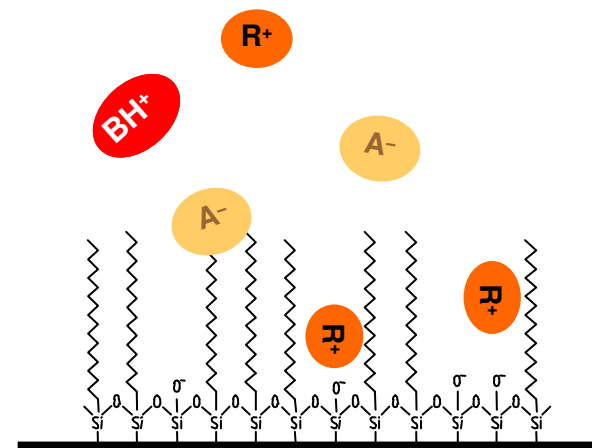
Horváth model applied to ionic liquids

$$\frac{[A]}{k_0 - k} = \frac{1}{k_2 K_A} + \frac{[A]}{k_2} \quad (4.20)$$

- K_A was also applied to measure the **ability of the cation of an ionic liquid to block silanol sites**.
- The retention is **affected by** the attraction of the cationic solute to the **adsorbed anion** on the stationary phase.
- **A third contribution** should be considered corresponding to the extra interaction:

$$k_0 = k_1 + k_2 + k_3$$

The **Horváth equation** can only be applied to **ionic liquids**, where the **absorption of the anion is negligible**.





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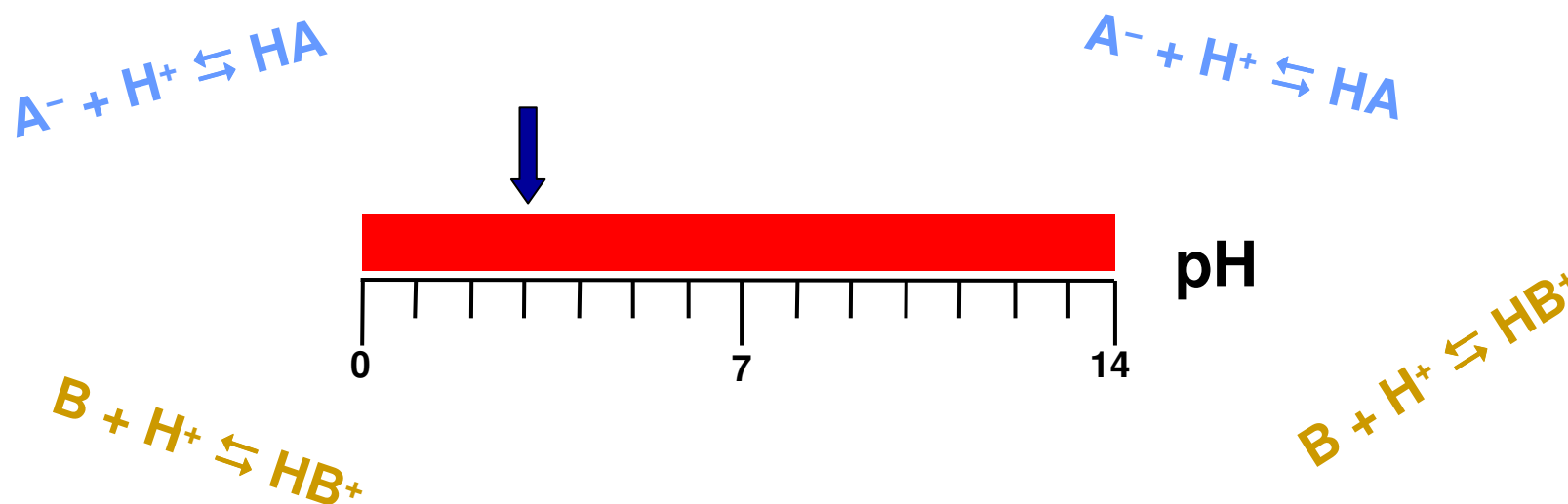




4.3. pH as experimental factor

4.3.1. Hydro-organic mobile phases

For **ionisable solutes**, **pH** is an additional experimental factor with a **large influence on retention and selectivity**. When the analysed mixture contains one or more compounds with acid-base behaviour, pH tuning can offer unique opportunities to reach resolution. However, **changes in retention with pH are particularly hard to model**. Not surprisingly, a widely extended practice consists of fixing the pH at a convenient value. With this practice, the **benefits of this experimental factor on selectivity are lost**.





RPLC retention of ionisable compounds

Weighted mean of the retention of the basic and acidic species

$$k = k_A \delta_A + k_{HA} \delta_{HA} = k_A \frac{1}{1 + K h} + k_{HA} \frac{K h}{1 + K h} = \frac{k_A + k_{HA} K h}{1 + K h} \quad (4.21)$$

k : the measured retention factor

k_A : retention factor for **basic species**

k_{HA} : retention factor for **acidic species**

δ_A : molar fraction for **basic species**

δ_{HA} : molar fraction for **acidic species**

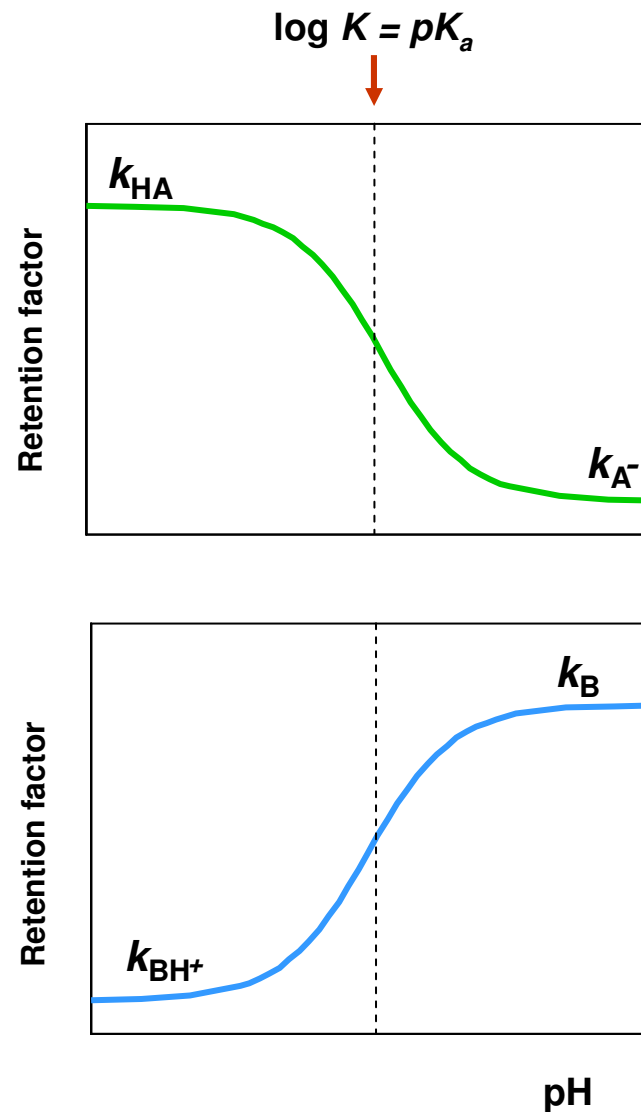
K ($= K_a^{-1}$) : apparent protonation constant

(takes into account all interactions of both acid-base species with the stationary phase and mobile phase components)



Remember that ...

- Since the intrinsic retention for each acid-base species **is different**, a **sudden change** in retention will happen at **pH** values **close to the logarithm of the apparent protonation constant** (or pK_a).
- Depending on the **charge of the acidic and basic species**, the retention **may decrease, increase, or remain constant with pH**.

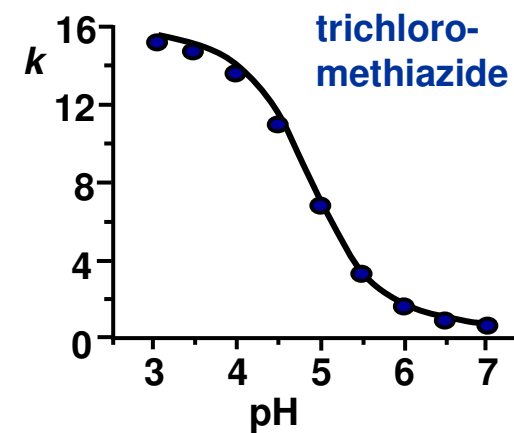
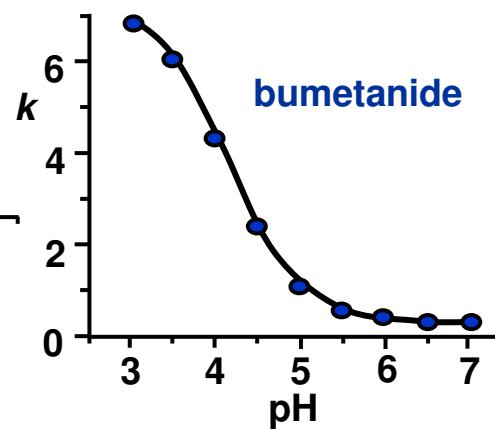
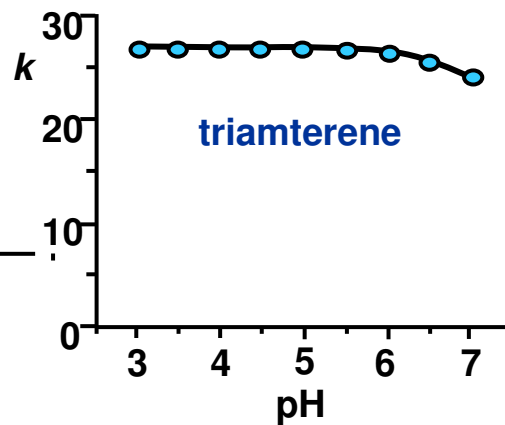
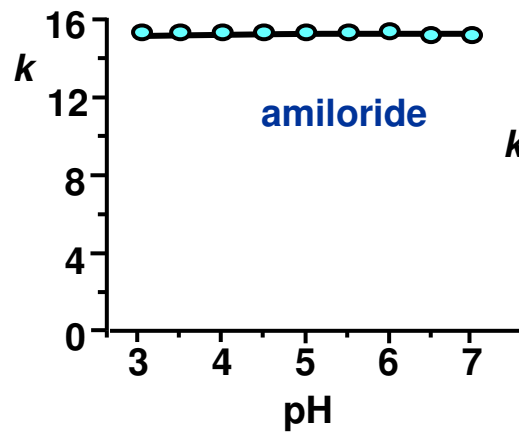




Incomplete equilibria of ionisable compounds

- The protonation process covers **several pH units**.
- Therefore, when the **working pH range** of the stationary phase is **narrow** (e.g. 3–7 for conventional RPLC columns), the change in **retention will be only fully sampled for some compounds**.
- For this reason, the retention plots of compounds with acid-base behaviour often show **partial equilibria**, giving rise to different patterns.

As a result of the incomplete information, **fitting** of the experimental data can be **hard**, since the **retention** of the acidic and basic species should be **extrapolated**.



Diuretics

0.05 M sodium dodecyl sulphate

4% propanol



Fixed organic solvent content

$$k = k_A \delta_A + k_{HA} \delta_{HA} = k_A \frac{1}{1 + Kh} + k_{HA} \frac{Kh}{1 + Kh} = \frac{k_A + k_{HA} Kh}{1 + Kh} \quad (4.21)$$

Dividing by k_{HA} ($f = k_A/k_{HA}$) yields a more convenient fitting behaviour:

$$k = \frac{\frac{k_A}{k_{HA}} + Kh}{1 + Kh} k_{HA} = \frac{f + Kh + fKh - fKh}{1 + Kh} k_{HA} = \left(f + \frac{Kh}{1 + Kh} (1 - f) \right) k_{HA} \quad (4.22)$$

$$\log k = \log k_{HA} + \log \left[f + \frac{Kh}{1 + Kh} (1 - f) \right] = \log k_{HA} + \log \left(f + \frac{10^{\log K} h}{1 + 10^{\log K} h} (1 - f) \right) \quad (4.23)$$



Simultaneous effect of organic solvent and pH

To account for the simultaneous effect of organic solvent and pH, the **equations that describe the retention with regard to both factors are combined.**

$$k = \frac{k_A + k_{HA} K h}{1 + K h}$$

$$\log k = \log k_w + S\phi + T\phi^2 \quad (4.25)$$

- The **change in the protonation constant with changes in the concentration of organic solvent** should also be considered.

$$\log K = \log K_w + Q_1\phi + Q_2\phi^2 \quad (4.26)$$

$$\log K = \log K_w + Q_1\phi \quad \leftarrow \text{for simplicity}$$

k_w : retention in pure water

K_w : protonation constant in pure water

- As a consequence of this relationship, **the retention surfaces are twisted and shifted.**

Both factors (organic modifier and pH) **interact with each other.**



This complicates the treatment notably.



$$\log k = \log k_{\text{HA}} + \log \left[f + \frac{10^{(\log K_w + m\varphi)} h}{1 + 10^{(\log K_w + m\varphi)} h} (1 - f) \right] =$$

$$(\log k_{w,\text{HA}} - S\varphi) + \log \left[f + \frac{10^{(\log K_w + m\varphi)} h}{1 + 10^{(\log K_w + m\varphi)} h} (1 - f) \right] \quad (4.24)$$

$$k = \frac{k_{w,A} 10^{(S_A \varphi + T_A \varphi^2)}}{1 + K_w h 10^{(Q_1 \varphi + Q_2 \varphi^2)}} + \frac{k_{w,HA} K_w h 10^{[(S_{HA} + Q_1) \varphi + (T_{HA} + Q_2) \varphi^2]}}{1 + K_w h 10^{(Q_1 \varphi + Q_2 \varphi^2)}} \quad (4.27)$$

quadratic relationships 

$$k = \frac{10^{(R_A + S_A \varphi + T_A \varphi^2)} + 10^{(R_{HA} + S_{HA} \varphi + T_{HA} \varphi^2)} h}{1 + 10^{(Q_0 + Q_1 \varphi + Q_2 \varphi^2)} h} \quad (4.28)$$

easier fitting 

S , T , Q_1 , Q_2 : coefficients that describe the influence of the organic modifier on the retention factor and protonation constant



Polarity model

$$k = \frac{k_A + k_{HA} K h}{1 + K h}$$

$$\log k = c_0 + c_1 P_M^N$$

$$\log k = c_0 + c_1 P_M^N + \log \left(f + \frac{10^{(\log K_w + m\phi)} h}{1 + 10^{(\log K_w + m\phi)} h} (1 - f) \right) \quad (4.29)$$

c_0 and c_1 correspond to the acidic species



4.3.2. Micellar mobile phases

$$k = \frac{k_A + k_{HA} K h}{1 + K h}$$

$$k = \frac{K_{AS} \frac{1}{1 + K_{AD} \varphi}}{1 + K_{AM} \frac{1 + K_{MD} \varphi}{1 + K_{AD} \varphi} [M]}$$

$$k = \frac{\left(K_{AS} \frac{1}{1 + K_{AD} \varphi} \right) + \left(K_{HAS} \frac{1}{1 + K_{HAD} \varphi} \right) K h}{\left(1 + K_{AM} \frac{1 + K_{MD} \varphi}{1 + K_{AD} \varphi} [M] \right) + \left(1 + K_{HAM} \frac{1 + K_{HMD} \varphi}{1 + K_{HAD} \varphi} [M] \right) K h} \quad (4.30)$$

Polynomial models: acceptable accuracy in **narrow pH ranges**

$$\frac{1}{k} = a_0 + a_1 [M] + a_2 \varphi + a_3 \text{pH} + a_{12} [M] \varphi + a_{13} [M] \text{pH} + a_{23} \varphi \text{pH} + a_{123} [M] \varphi \text{pH} + a_{33} \text{pH}^2 \quad (4.31)$$



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4.4. Quantitative structure-retention relationships (QSRR)

The ideal optimisation is that one requiring **no experiment**. Accordingly, considerable effort has been made to estimate retention **without any other extra knowledge that** several **experimental solute descriptors** or **solute chemical structures**.

General solvation model (Abraham model)

A property for a **series of solutes at fixed solvent conditions** can be predicted from several **solute molecular descriptors**:

$$SP = eE + sS + aA + bB + vV + c \quad (4.32)$$

SP : solute property (**log k** or polarity parameter p_s)

E, S, A, B, V : solute descriptors

e, s, a, b, v : characterise **differences between the mobile phase and stationary phase in HPLC**

E : excess molar refraction

S : dipolarity/polarizability

A : hydrogen-bond acidity

B : hydrogen-bond basicity

V : McGowan volume

e : capability of the environment to interact
with solute π and n -electron pairs

s : dipolarity/polarizability

a : hydrogen-bond basicity

b : hydrogen-bond acidity

v : hydrophobic contribution

} polar
contributions



Correlations between retention properties and octanol-water partition coefficients

$$SP = c_0 + c_1 \log P_{o/w}$$

$$SP = \log k \text{ or } p_s$$

- However, conventional RPLC columns still have **silanol groups accessible to solutes and solvent**, even when heavily coated and end-capped.
- Silanols are **weakly acidic** and may affect the retention: **ionised silanols** can retain **protonated bases** by cation-exchange, and **neutral silanols** can **hydrogen bond proton-acceptor solutes**.

➔ Therefore, the conversion of $\log P_{o/w}$ into $\log k$ or p_s values **needs a hydrogen-bond acidity term** or any other term accounting the hydrogen-bond donor interactions:

$$SP = c_1 + c_2 \log P_{o/w} + c_3 A \quad (4.33)$$

- Comprehensive **databases** listing **$P_{o/w}$ values** are available. Also, **$P_{o/w}$ values** and **hydrogen-bond acidity** can be **predicted from molecular structures**.



Use of molecular descriptors predicted from molecular structures

- Software such as **CODESSA** or **Dragon** are able to **calculate multiple molecular descriptors**.
- **Correlations** between retention data and the molecular descriptors can be obtained.
- **A step is required where the number of descriptors** is reduced so that only a small number of uncorrelated descriptors is kept.

Example: in a study including a set of 233 compounds showing a wide variation of chemical characteristics, the solute polarity parameter p_s was best described by a model containing, besides $\log P_{o/w}$, three more descriptors:

$$p_s = c + p_{o/w} \log P_{o/w} + p_{o/d}(\text{pol}/d^2) + hl(\text{HOMO-LUMO}) + hd(\text{HDCA2}) \quad (4.34)$$

pol/d² : related to the electrostatic nature

HOMO-LUMO : energy-related descriptor

HDCA2 : accounted for hydrogen-bond interactions



Unfortunately, current QSRR predictions are **not yet sufficiently accurate for optimisation purposes**. Reaching the high accuracy level needed in optimisation still requires the assistance of experimental data. However, this approach is interesting for **prospective purposes**, such as the selection of the separation system.