
IOCB SERVICE DAYS

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Research-Service Groups / Service Groups / Core Facilities

20
23

Electromigration Methods

Václav Kašička



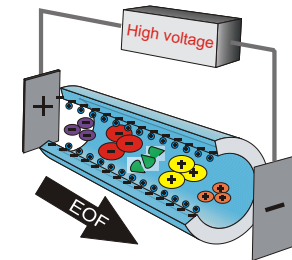
ÚOCHB AV
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IOCB PRAGUE



Electromigration Methods

Research-Service Group

Analytical Chemistry & Separation Sciences, cluster PHYS



People

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Scientists: Dušan Koval

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Ishak Kovač (0.2)



Location

Building A, S-E wing, 2nd floor, A 2.91



ÚOCHB AV ČR, v.v.i.

Capillary Electromigration (CE) methods

Electrophoretic:

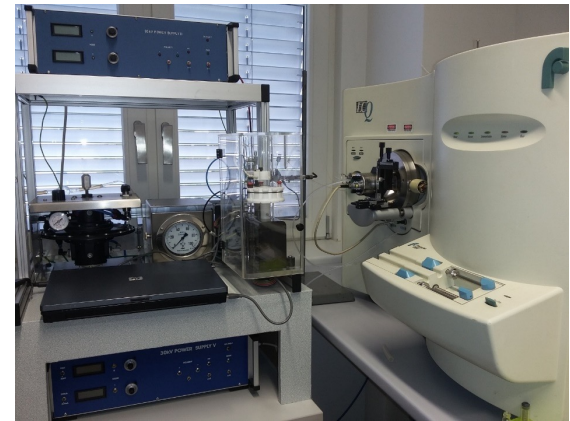
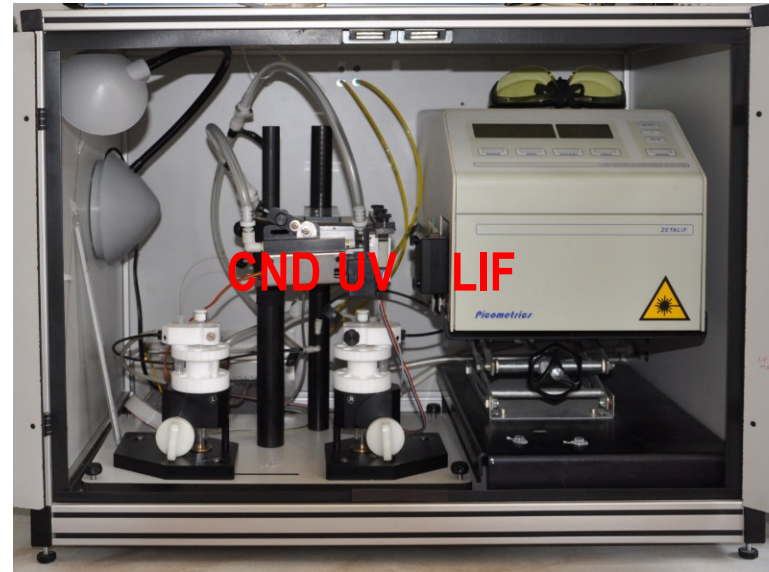
- Zone Electrophoresis (CZE (FSCE, CGE))
- Isotachopheresis (CITP)
- Isoelectric focusing (CIEF)
- Affinity electrophoresis (ACE)

Electro-chromatographic:

- Electrokinetic chromatography (CEKC)
- Electrochromatography (CEC)

Instrumentation

- **Two CE analyzers 7100 Agilent**, fully automated, UV-vis & CND detection
- **Two CE analyzers MDQ Beckman Coulter**, fully automated, UV-vis or LIF detection
- **Two home-made CE devices**, semi-automated, CND, UV, LIF or MS detection



CE methods = high performance techniques (HPCE)

High: Separation efficiency: 10^5 – 10^6 theor. plates/m

Sensitivity: fmol–amol in nL–pL inj. volume; μ M–nM

Typical samples: 1–100 μ M, 5–20 μ L, 100 μ g

Speed: 5–20 min

Selectivity: charge, size, shape, mobility | affinity, hydrophobicity

Flexibility: CZE, CTP, CIEF, CGE | ACE, CEKC, CEC

Applicability: ionogenic, electroneutral, LMW, HMW, (bio)particles,

Biocompatibility: free aqueous buffered solution

Medium: Concentration sensitivity: μ mol/L

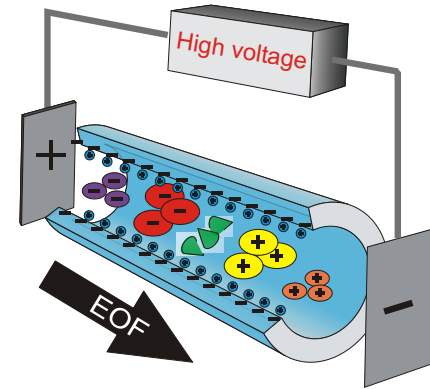
Repeatability: 1–5 %

Robustness

Low: Pressure: mbar–10 bar

Running costs: 5–10 EUR / 1 m capillary

Environmental pollution: mostly in aqueous media – green methods



Provided services/collaborations (1)

Qualitative and quantitative analysis

- **Quality control:** determination of chemical purity degree
- **Monitoring of chemical & enzymatic reactions, isolation, purification**
- **Determination of LMW ionic admixtures (salts)**
- **Determination in (bio)matrices:** biofluids, tissue extracts, reaction mixtures
- **Separation of complex mixtures:** peptide mapping, bottom-up proteomics
- **Separation of stereoisomers:** chiral analysis, monitoring of racemization

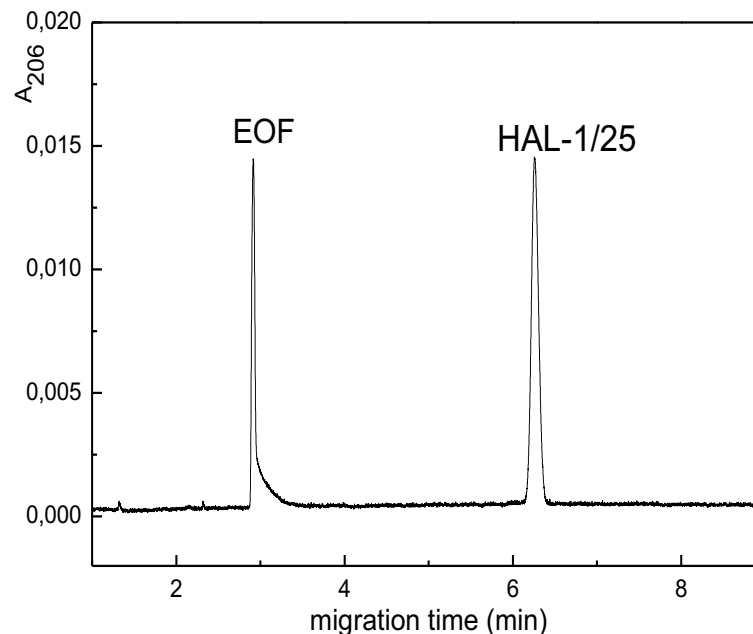
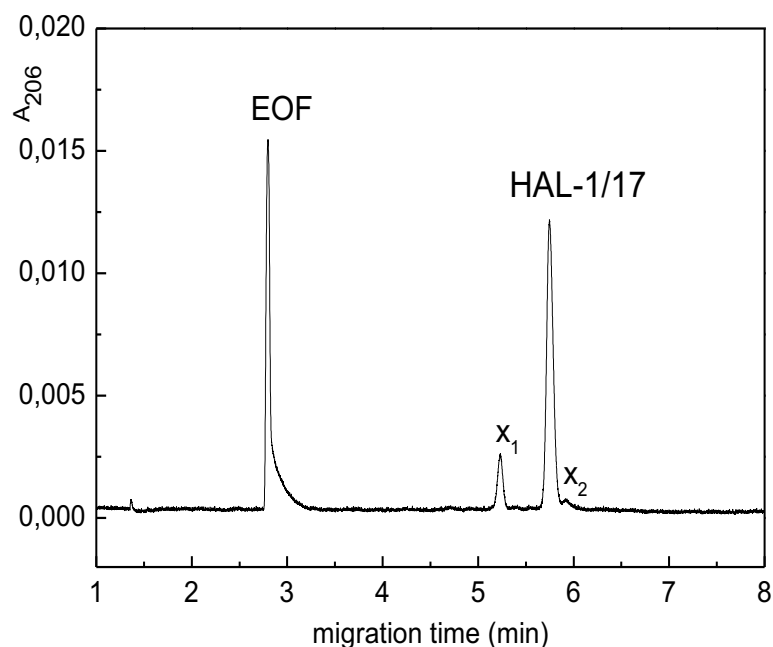
Analyzed (bio)molecules and (bio)particles:

- ❖ **Small organic & inorganic ions and molecules:** acids, amino acids, metal ions, (nucleo)bases, nucleosides, nucleotides, monosaccharides, steroids, etc.
- ❖ **Medium size (bio)molecules:** peptides, oligonucleotides, oligosaccharides, helquats
- ❖ **Large (bio)molecules:** proteins, polynucleotides, DNA/RNA fragments, polymers
- ❖ **(Bio)particles:** viruses, bacteria, cells, cell organelles, nanoparticles

Soluble in: buffered solutions in water or organic solvents (MeOH, EtOH, IPA, ACN)
or mixed hydro-organic solvents

CZE analysis of antimicrobial peptides

Derived from: H-Gly-Met-Trp-Ser-Lys-Ile-Leu-Gly-His-Leu-Ile-Arg-NH₂ (+4e)



Halictine analog: KMWSKILGHLIR-NH₂, +5e

Halictine fragment: WSKILGHLIR-NH₂, +4e

CE: home-made IOCB, capillary: FS dynamically coated with DDAB, 50 μ m x 31/20 cm

Detection: UV-abs @ 206 nm, BGE: 25/30 mM Tris/H₃PO₄, 0.1 mM DDAB, pH 2.8, -12 kV, 35 μ A

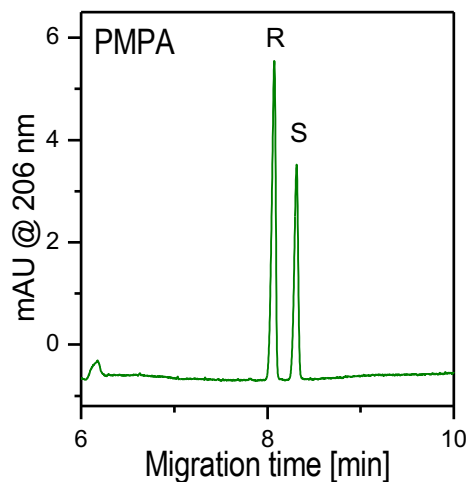
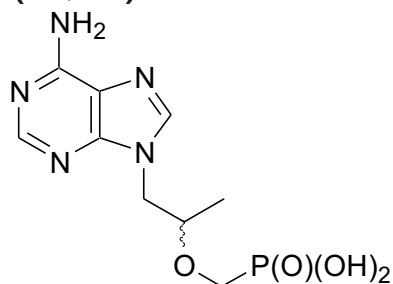
Sample: 100 μ g peptide/100 μ L BGE, hydrodynamic injection, 500 Pa, 20 s

CE of enantiomers of acyclic nucleotide phosphonates

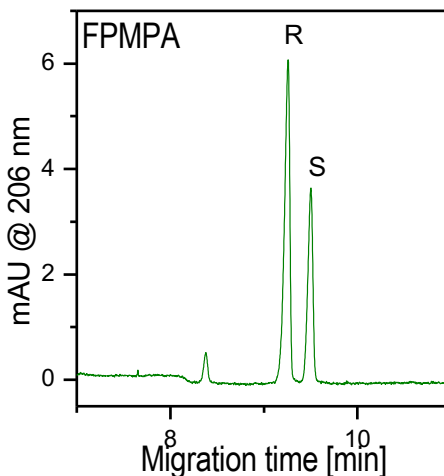
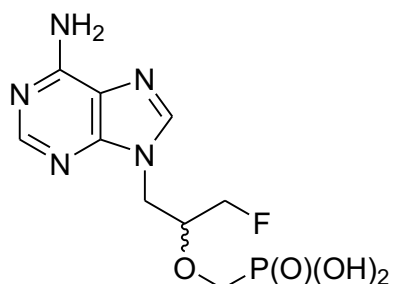
BGE: 30 mM $\text{Na}_2\text{B}_4\text{O}_7$ + NaOH, pH 10.0

Chiral selector: 20 mg/mL β -cyclodextrin

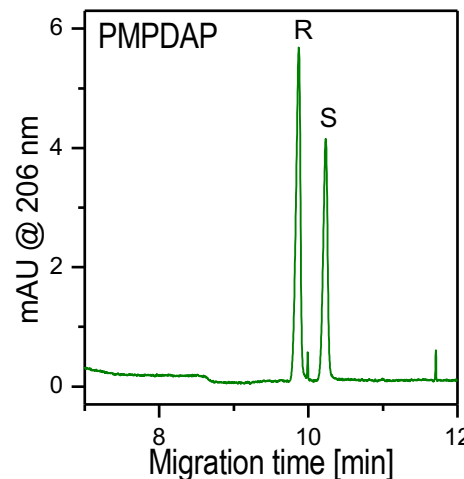
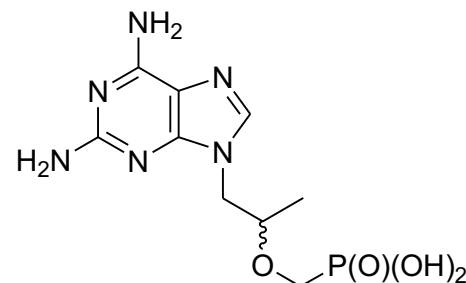
(*R,S*)-PMPA



(*R,S*)-FPMPA



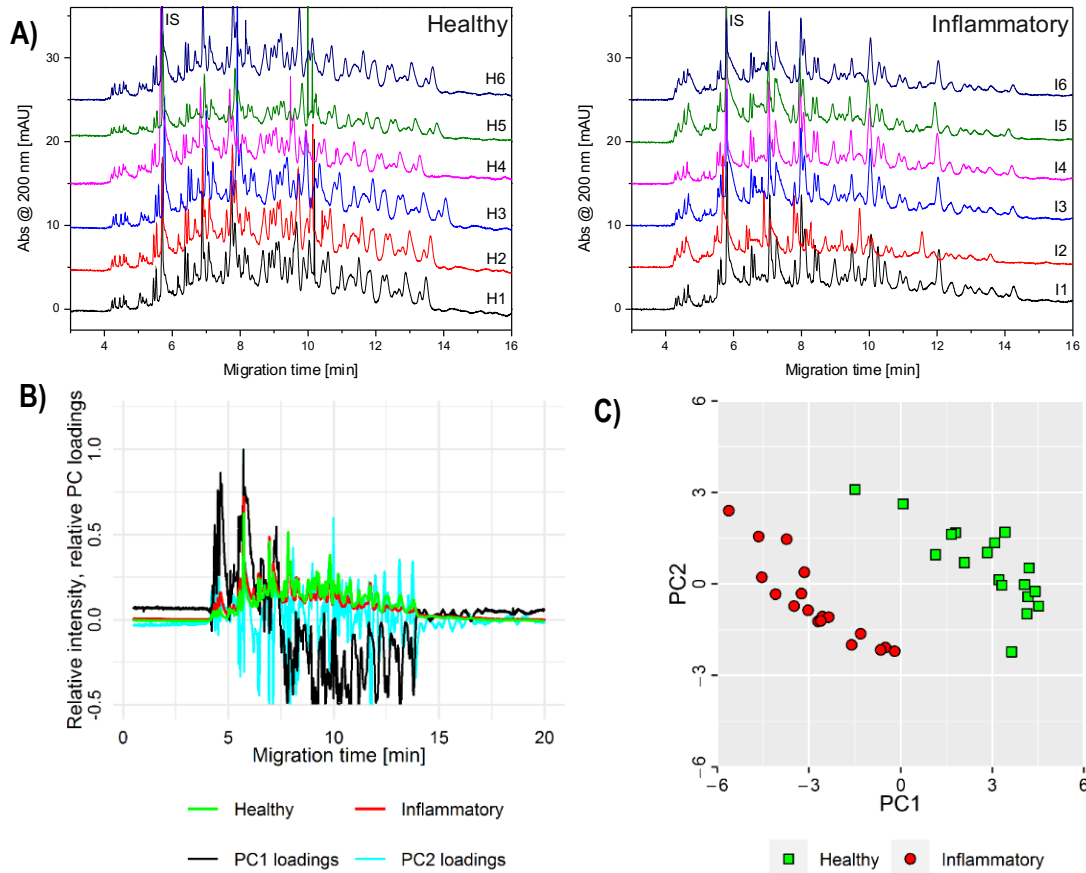
(*R,S*)-PMPDAP



Fused silica capillary 50/375 μm , 39.2/29.0 cm, 15 kV, 66.5 μA , 20°C, inj. 6.9 mbar, 5 s

CE separations of complex peptide mixtures

- A) **CE-UV profiles** of tryptic peptides of proteins of **healthy (left) & inflammatory (right) bone tissues**
- B) Averaged normalized CE-UV profiles and relative loadings of the first two principal components
- C) PCA scores plot of healthy and inflammatory tissue samples



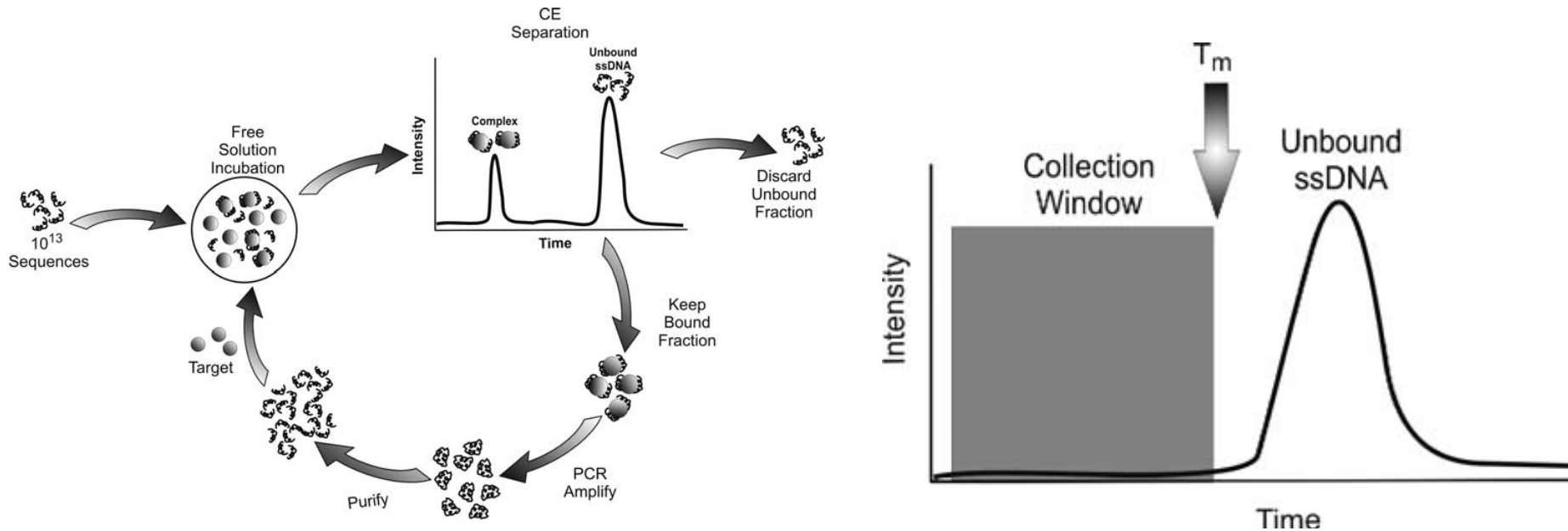
FS capillary: 50/375 μm id/od, 30/40 cm effect./total length, UV det. @ 200 nm

BGE 5: 55 mM H_3PO_4 , 14 mM Tris, pH 2.01; 12 kV, 20°C, IS, internal standard (0.2 mM tyramine)

Provided services/collaboration (2)

Microscale isolation: ng – μ g scale isolation (selection of binding aptamers)

CE-SELEX (CE – Systematic Evolution of Ligands by Exponential Enrichment)



Scheme of the CE-SELEX process. A random sequence DNA library is incubated with the target. Sequences bound to the target are separated and isolated by CE, PCR amplified and made single stranded, generating a new pool suitable for further rounds of enrichment.

Provided services/collaboration (3)

Phys-chem. & biochem. characterization

$$m_{\text{eff}} = \frac{v}{E} \doteq \frac{q}{6\pi\eta r}$$

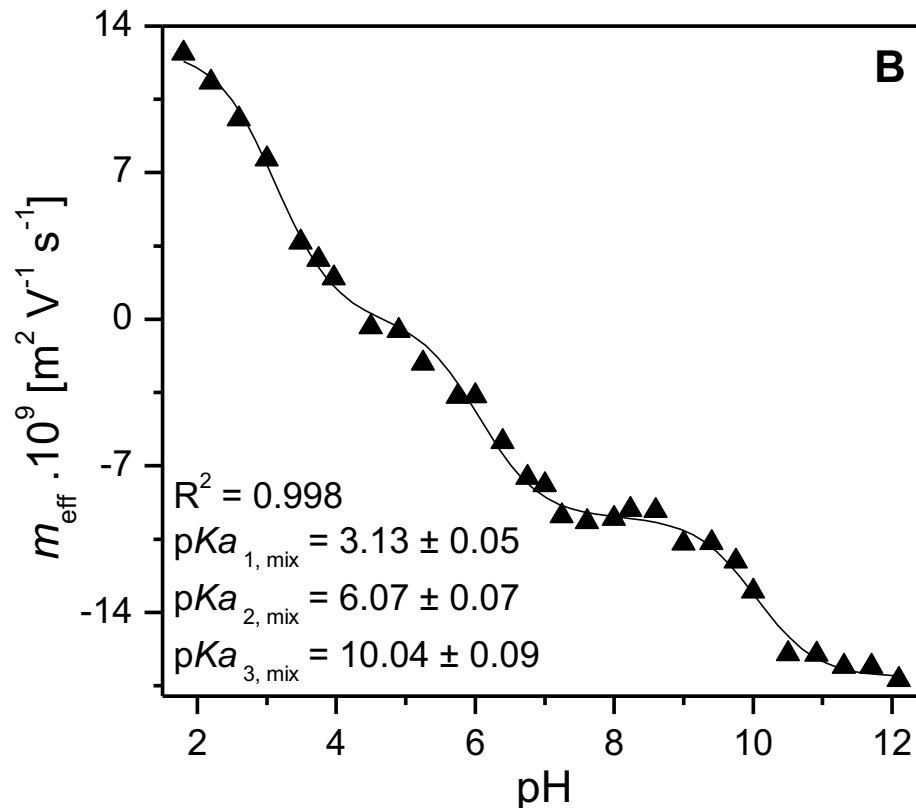
$$m_{\text{eff}} = \frac{L_{\text{tot}} L_{\text{eff}}}{U_{\text{sep}}} \left(\frac{1}{t_{\text{mig}}} - \frac{1}{t_{\text{eof}}} \right)$$

- Effective, actual & limiting (absolute) electrophoretic mobilities
- Effective charges
- Stokes radii
- Relative molecular masses (SDS-CGE)
- Diffusion coefficients
- Isoelectric points
- Acidity (ionization) constants
- Association (binding) constants of complexes
- Interconversion barriers of enantiomers
- Rate constants of chem. & enzymatic reactions
- Gibbs energy, enthalpy, entropy

Determination of acidity constants (pKa) of trivalent peptide by CE



$$m_{eff} = \frac{m_{P^+} 10^{pK_{a,1}^{mix} - pH} + m_{P^-} 10^{pH - pK_{a,2}^{mix}} + m_{P^{-2}} 10^{2pH - pK_{a,2}^{mix} - pK_{a,3}^{mix}}}{10^{pK_{a,1}^{mix} - pH} + 10^{pH - pK_{a,2}^{mix}} + 10^{2pH - pK_{a,2}^{mix} - pK_{a,3}^{mix}} + 1}$$



CONCLUSIONS

Do you need:

- High-sensitive analysis
- High-efficient separation
- Microscale isolation
- Phys-chem. & biochem. characterization
of your (bio)molecules?

Do not hesitate to contact us:

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Since ca. April 2023:

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