



Polymeric polypeptides



How structure influence functional properties of branched chain polymeric polypeptides ?

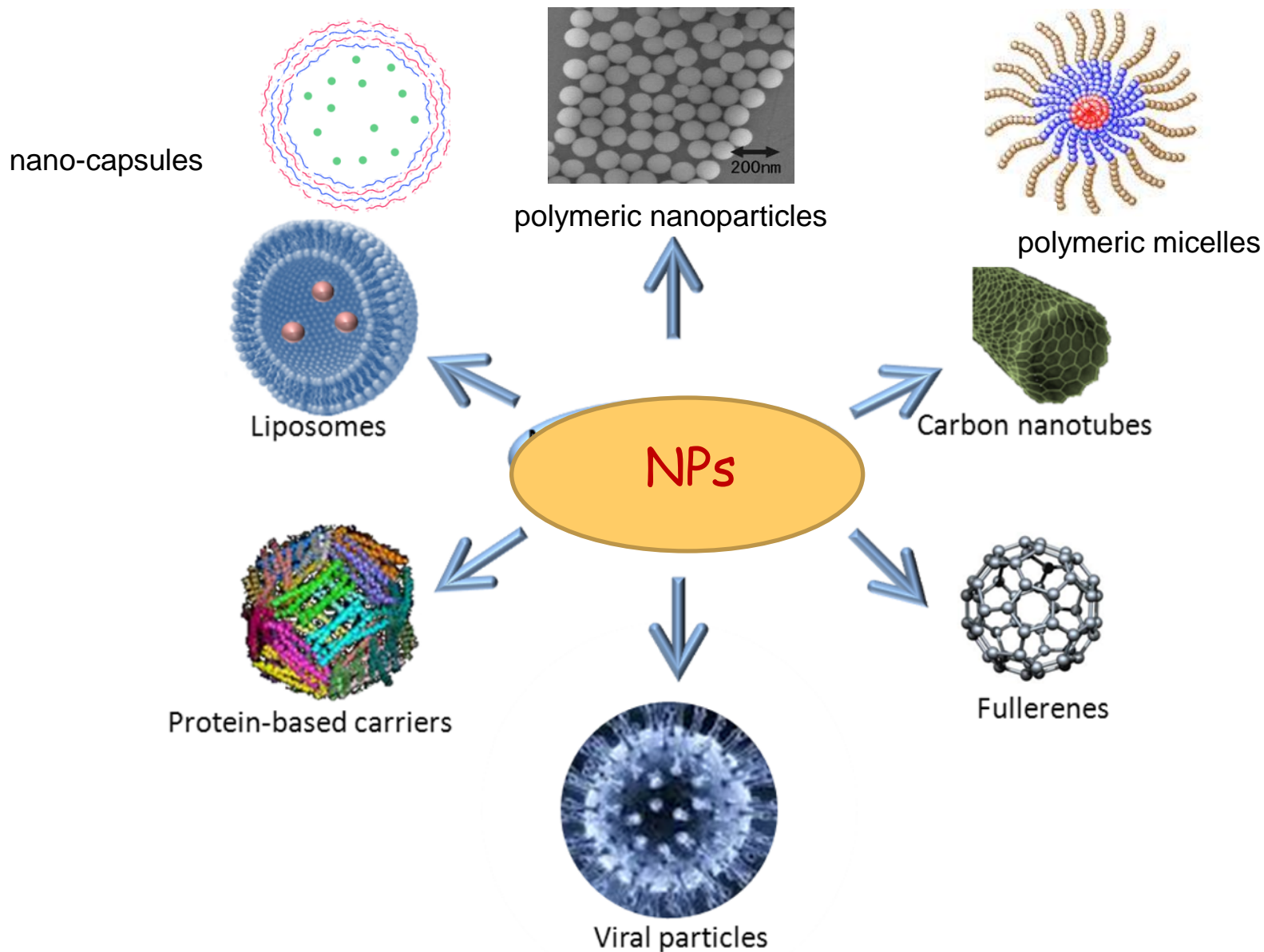
Ferenc Hudecz^{1,2}

¹Department of Organic Chemistry, Eötvös Loránd University, Budapest

²Research Group of Peptide Chemistry, Hungarian Academy of Sciences

Nanoparticle based carriers for biomedical applications

Ryvolova, M. et al. *Sensors* 12: 14792-14820 (2012)



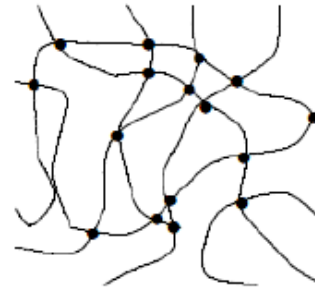
Branched polymers



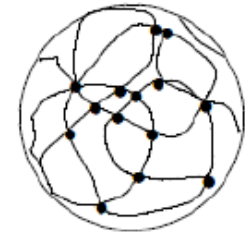
short chain branching



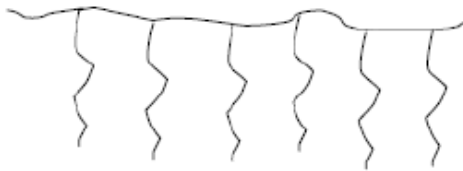
highly branched



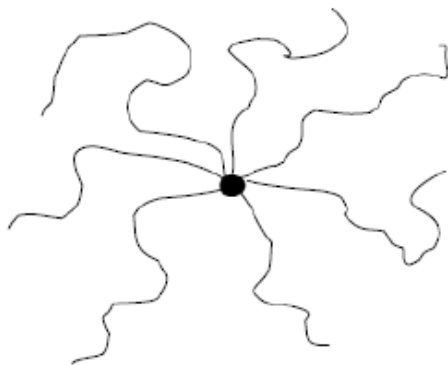
networks



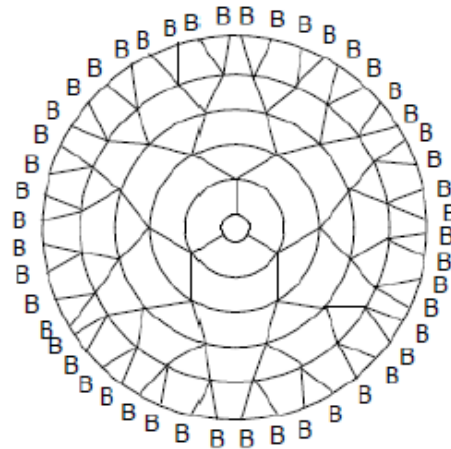
microgels



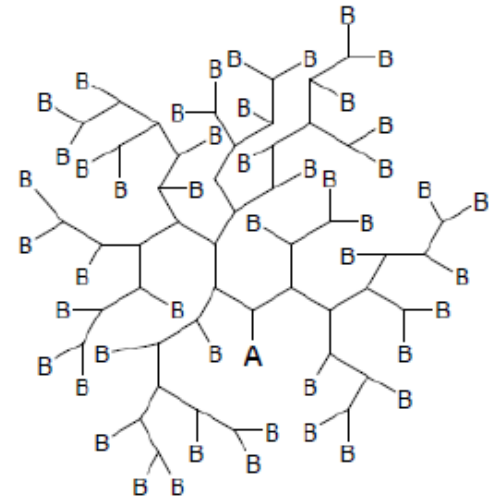
comb



stars



dendrimers



hyperbranched polymers

Conjugates with polymeric polypeptides



Synthetic antigens

- peptide vaccine
- immunodiagnostics

HSV gD
glycoprotein

mucin
glycoprotein(s)

tuberculin
proteins

Drug/reporter targeting

- increased specificity
- prolonged effect

antitumor drugs

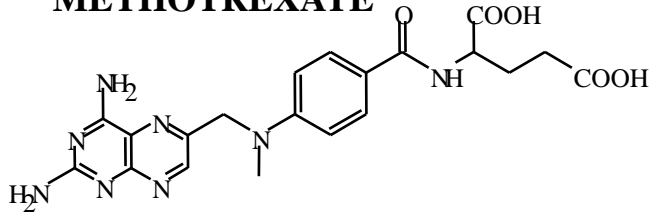
- daunomycin
- vinblastin

antimicrobials

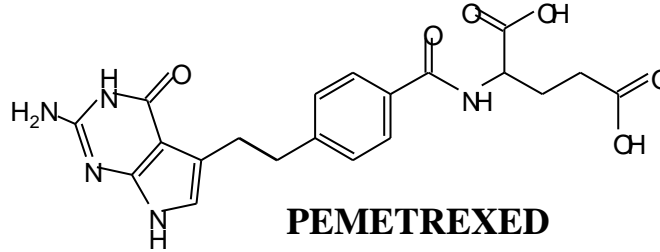
- *Leishmania don.* (e.g. methotrexate)
- *M. tuberculosis* (e.g. isoniazid)

Drug, epitope, reporter molecule

METHOTREXATE

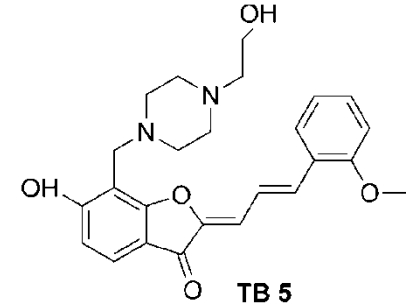


Hudecz F. et al. *Bioconjugate Chem.* **4**: 25 (1993)
Kóczán Gy. et al. *Bioconjugate Chem.* **13**: (2002)



PEMETREXED

Miklán Zs. et al.
J. Peptide Sci. **17**: 805 (2011)



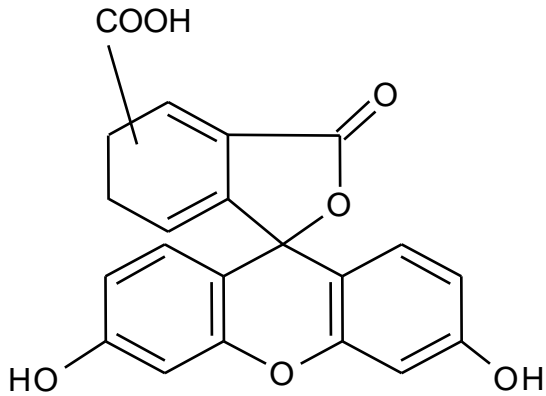
TB 5

Horvati, K.. et al.
Bioconjugate Chem. **22**:981 (2012)

D-Trp-*D*-Cpa-*D*-Trp-Ser-Tyr-*D*-Lys-Leu-Arg-Pro-*D*-Ala

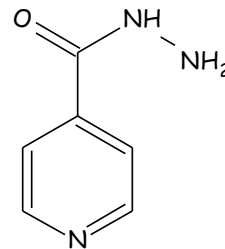
GN-RH ANTAGONIST, MI-1544

Mező, G. et al. *Bioconjugate Chem.* **7**: 642 (1996)
Vincze, B. et al. *J. Cancer Res. Clin. Onc.* **120**: 578 (1994)



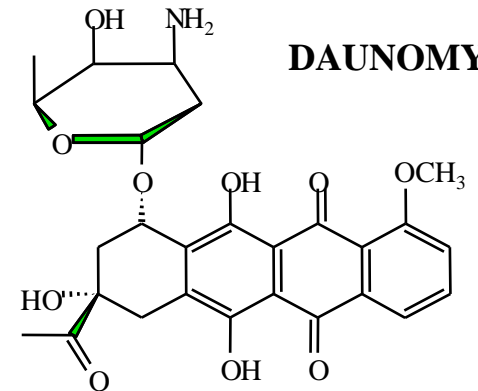
5(6)-CARBOXYFLUORESC EIN

Szabó R. et al. *Bioconjugate Chem.* **19**: 1078 (2008)
Bánóczy Z. et al. *Bioconjugate Chem.* **19**: 1375 (2008)



ISONIAZID

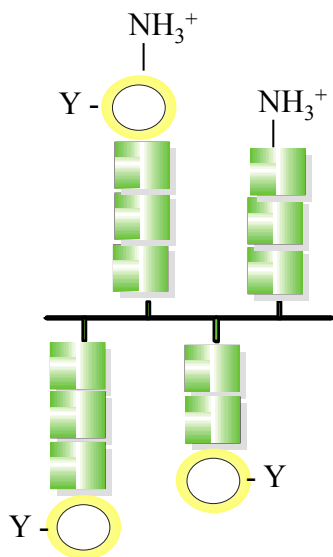
Horvati, K.. et al.
J. Peptide Sci. **15**:385 (2009)



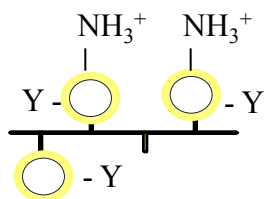
DAUNOMYCIN

Bánóczy Z. et al. *Archivoc* **143** (2008)
Miklán Zs. et al. *Biopolymers* **92**:489 (2009)
Szabo R. et al. *BBA* **1798**: 2209 (2010)

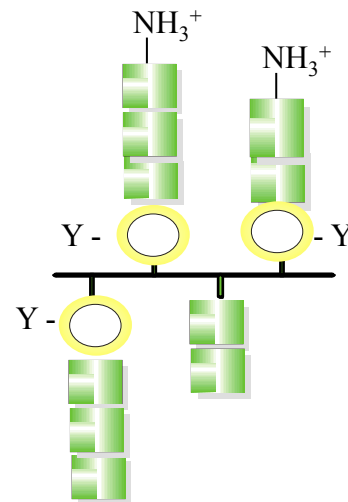
Branched polypeptides



XAK, poly[Lys(X_i -DL-Ala $_m$)]



X_iK , poly[Lys(X_i)]



AXK, poly[Lys(DL-Ala $_m$ - X_i)]

Hudecz, F.: (Eds.: Agelli, A., Boden, N., Zhang, S.) Kluwer Academic Publisher (2001), pp. 139-160

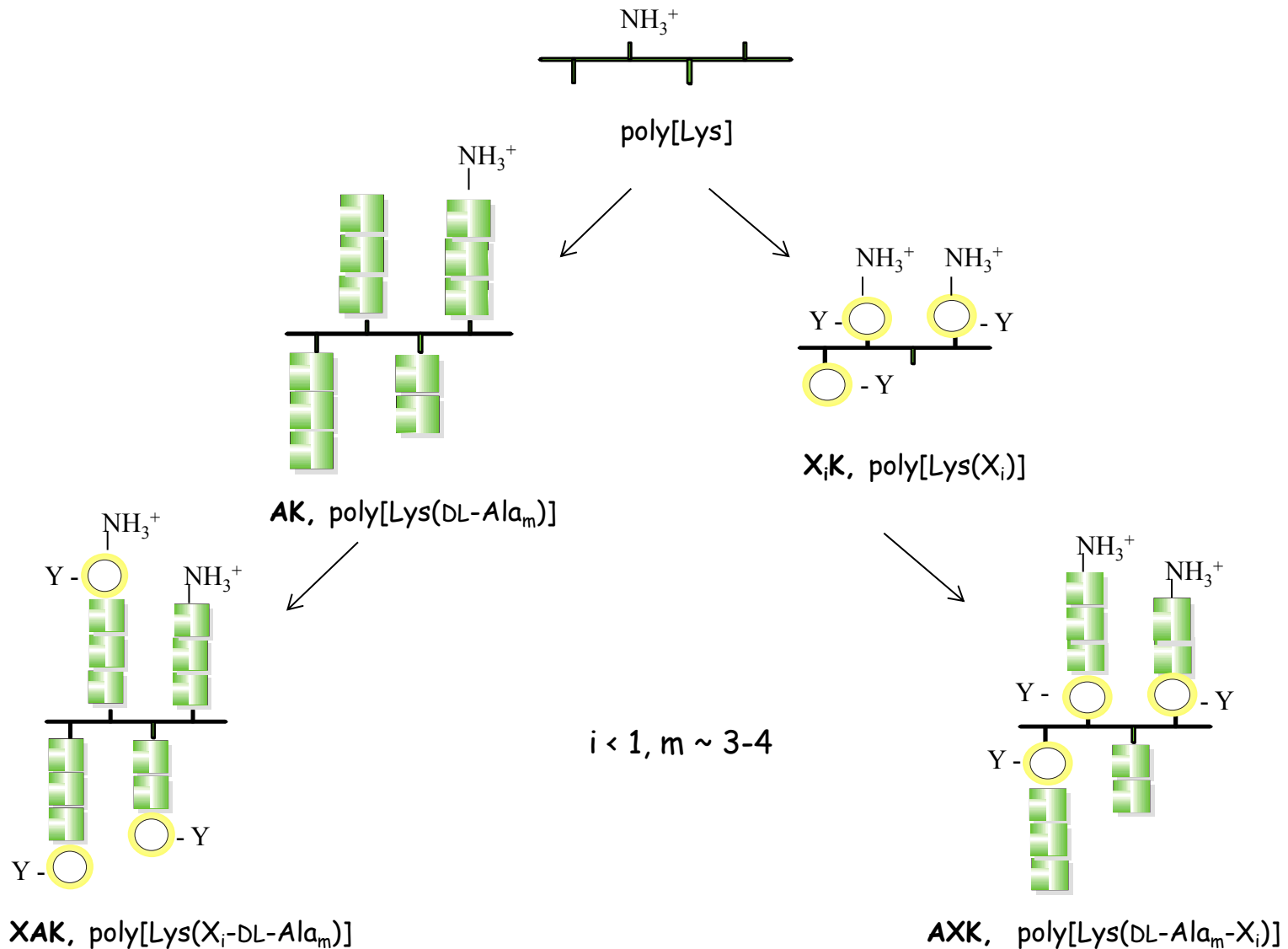
Hudecz, F., Kóczán, Gy., Reményi, J.: (Eds.: Kéri, Gy., Tóth, I.) Taylor and Francis Group (2003) pp. 553-578

Sebestyén, M., Szabó, R., Köhidai, L., Pállinger, É., Mező, G., Kóczán, Gy., Hudecz, F.:

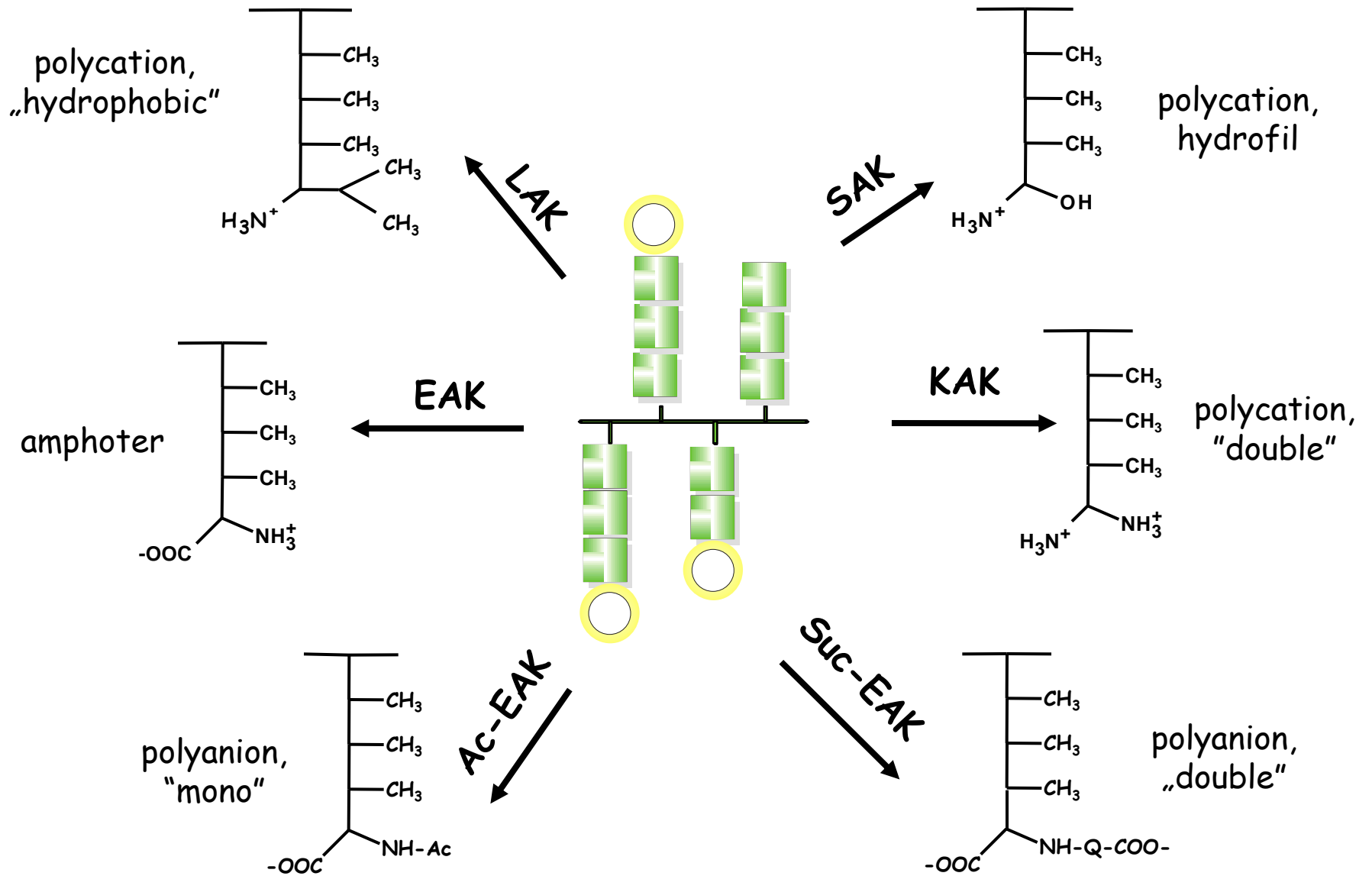
Structural Chemistry, 28: 527-536 (2017)

Szabó, R., Sebestyén, M., Kóczán, Gy., Orosz, Á, Mező, G., Hudecz, F.: ACS Combinatorial Science 19: 246-254(2017)

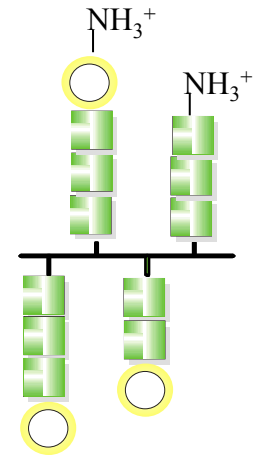
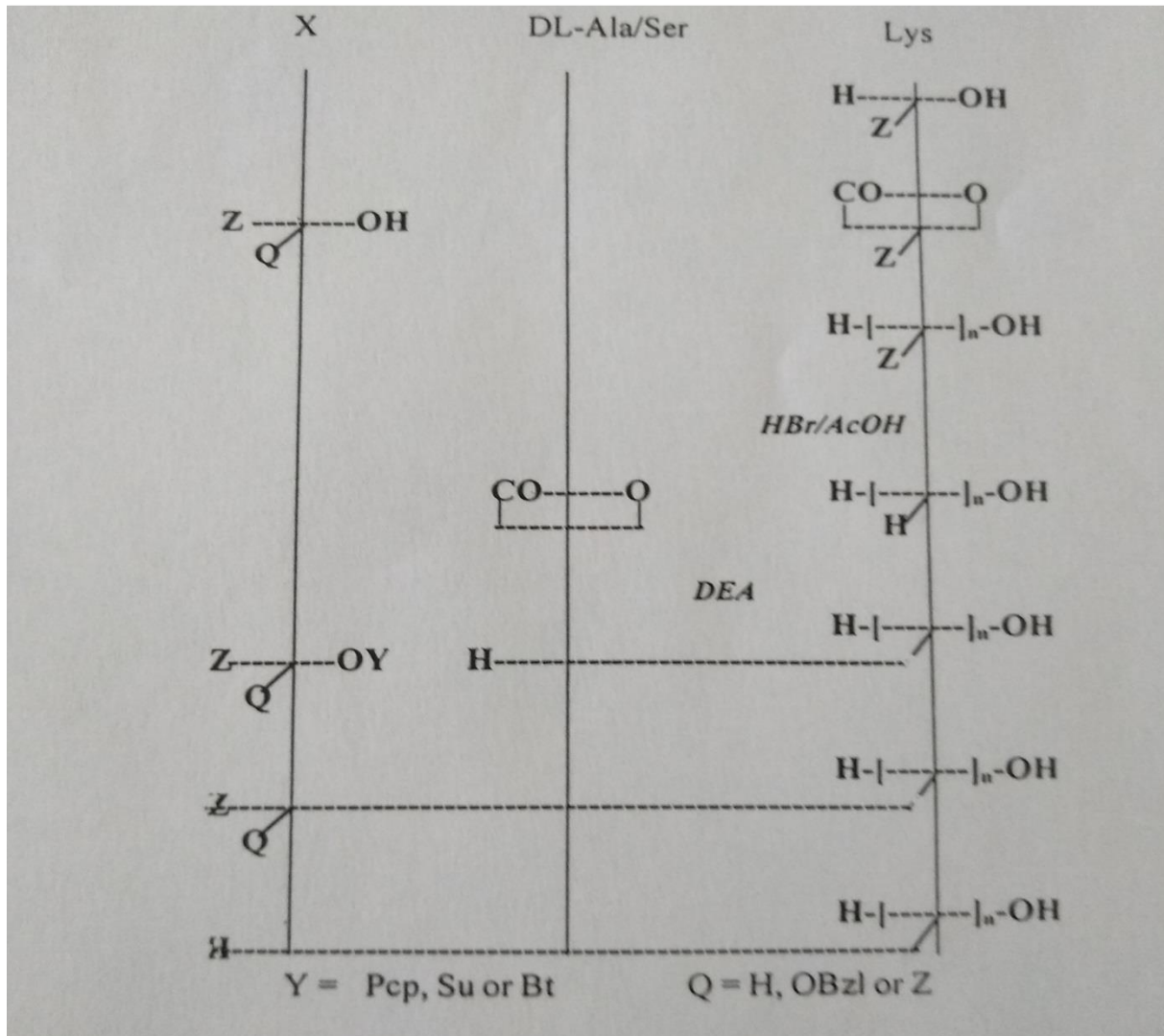
XAK, X_iK and AXK type branched polypeptides: synthesis



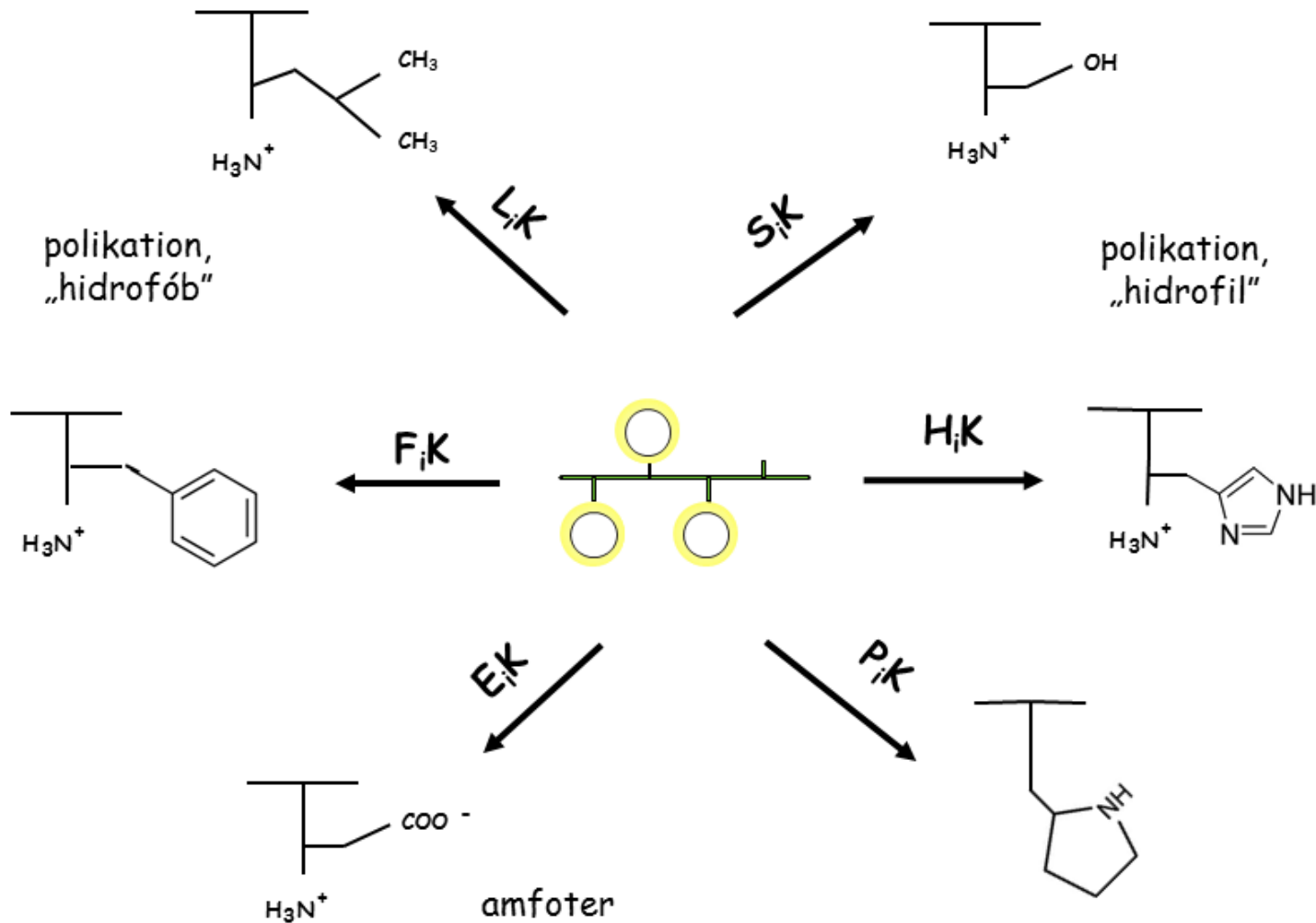
XAK type branched chain polypeptides



Preparation of XAK/XSK type branched chain polypeptides

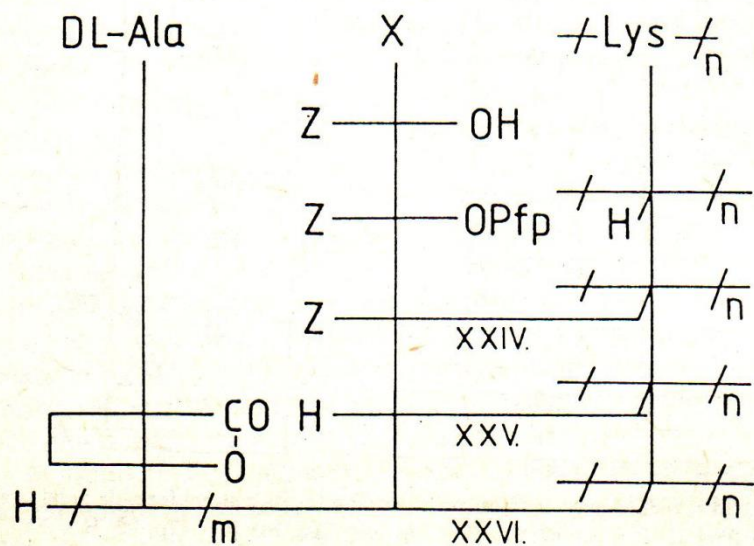
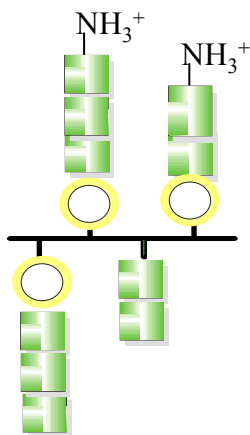
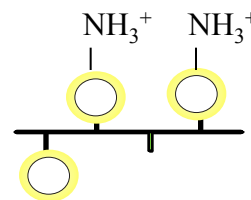
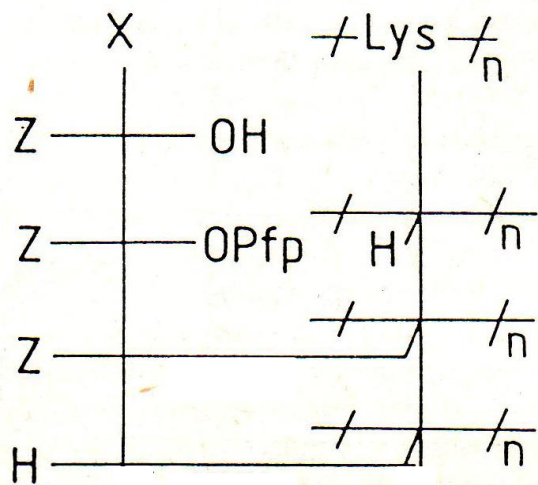


X_iK type branched chain polypeptides

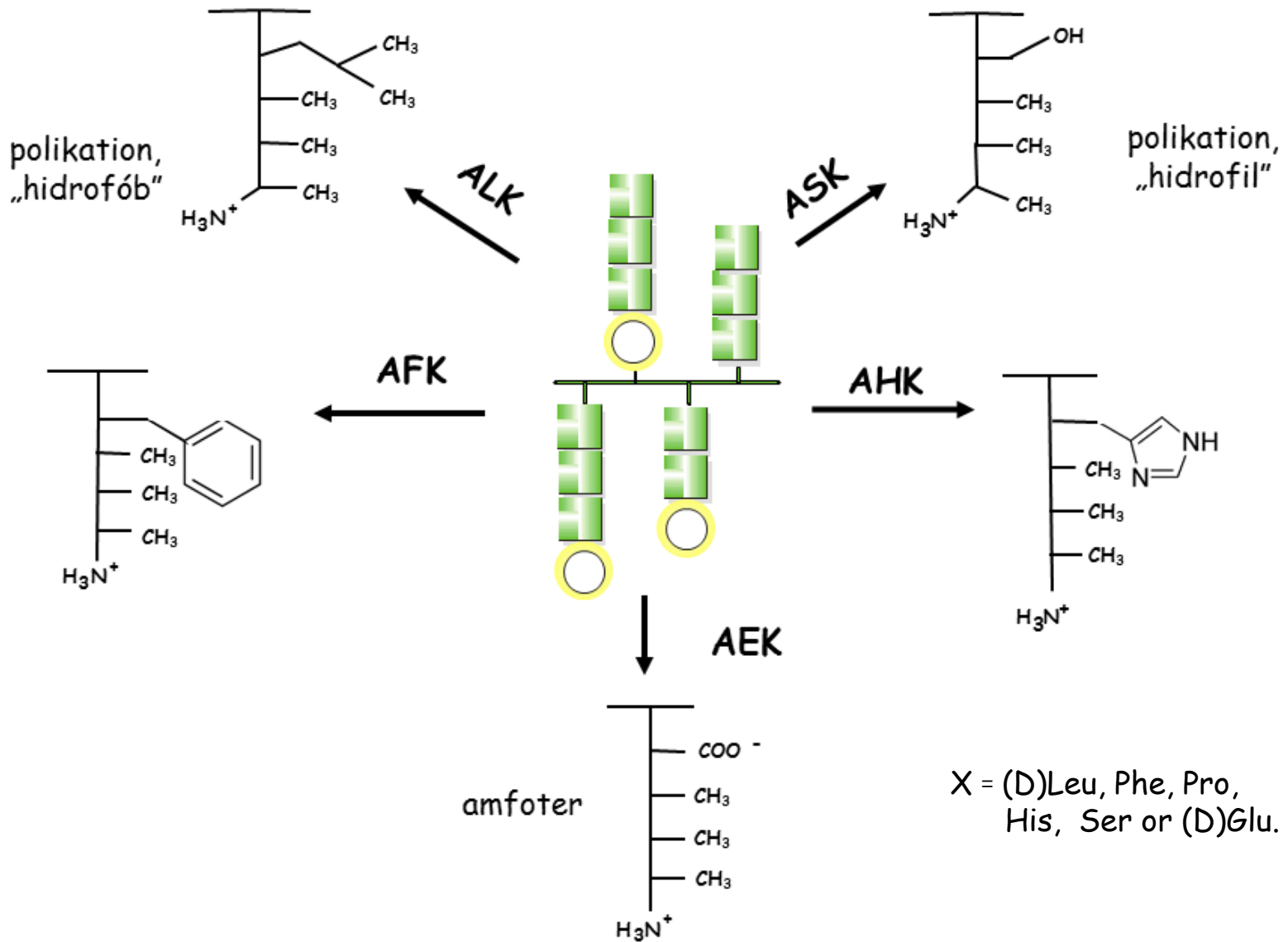


X = (D)Glu, (D)Leu, (D)Phe, (D)Ala, Ile, Pro, His or Ser.

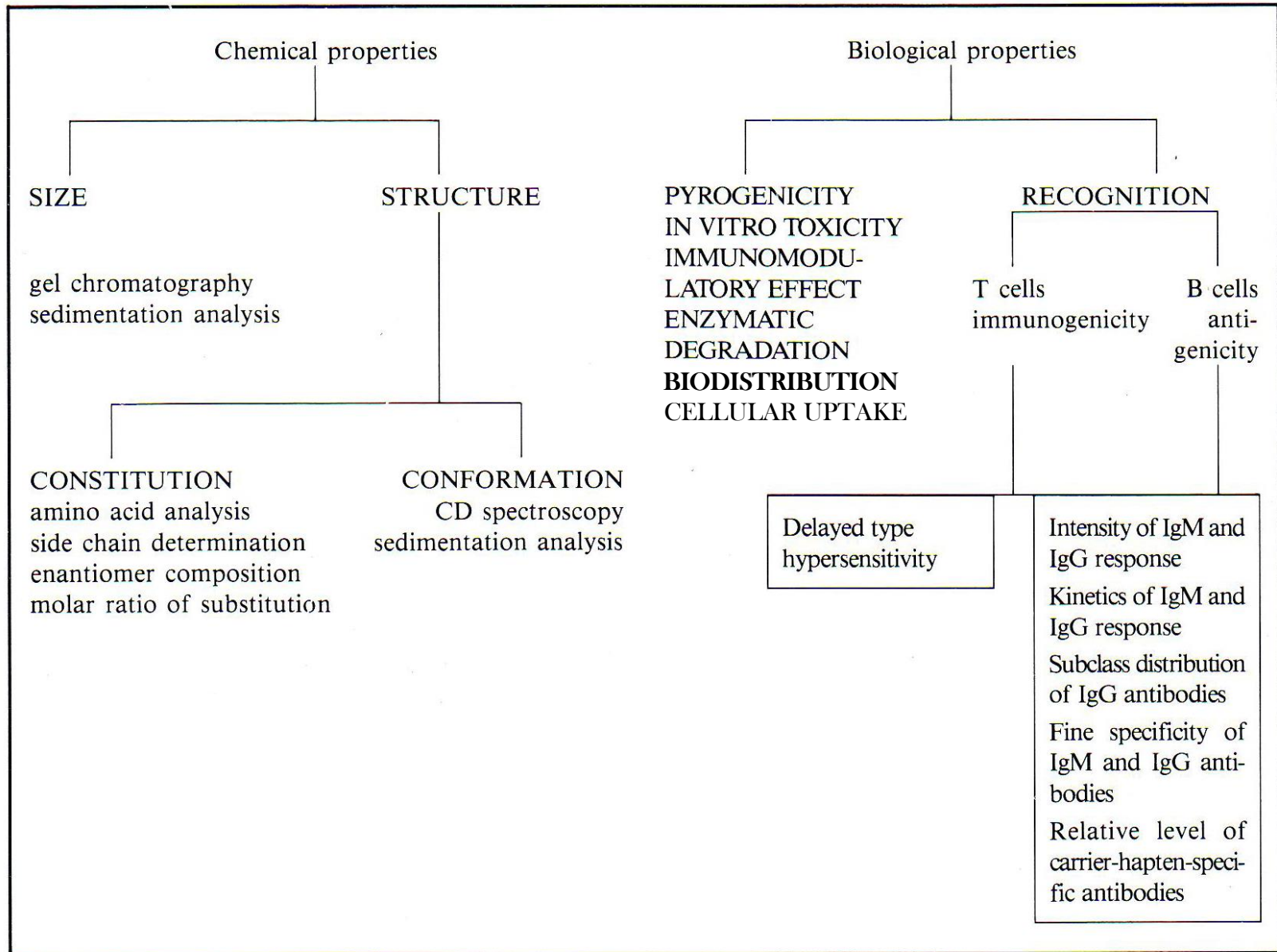
Preparation of X_iK and AXK type branched chain polypeptides



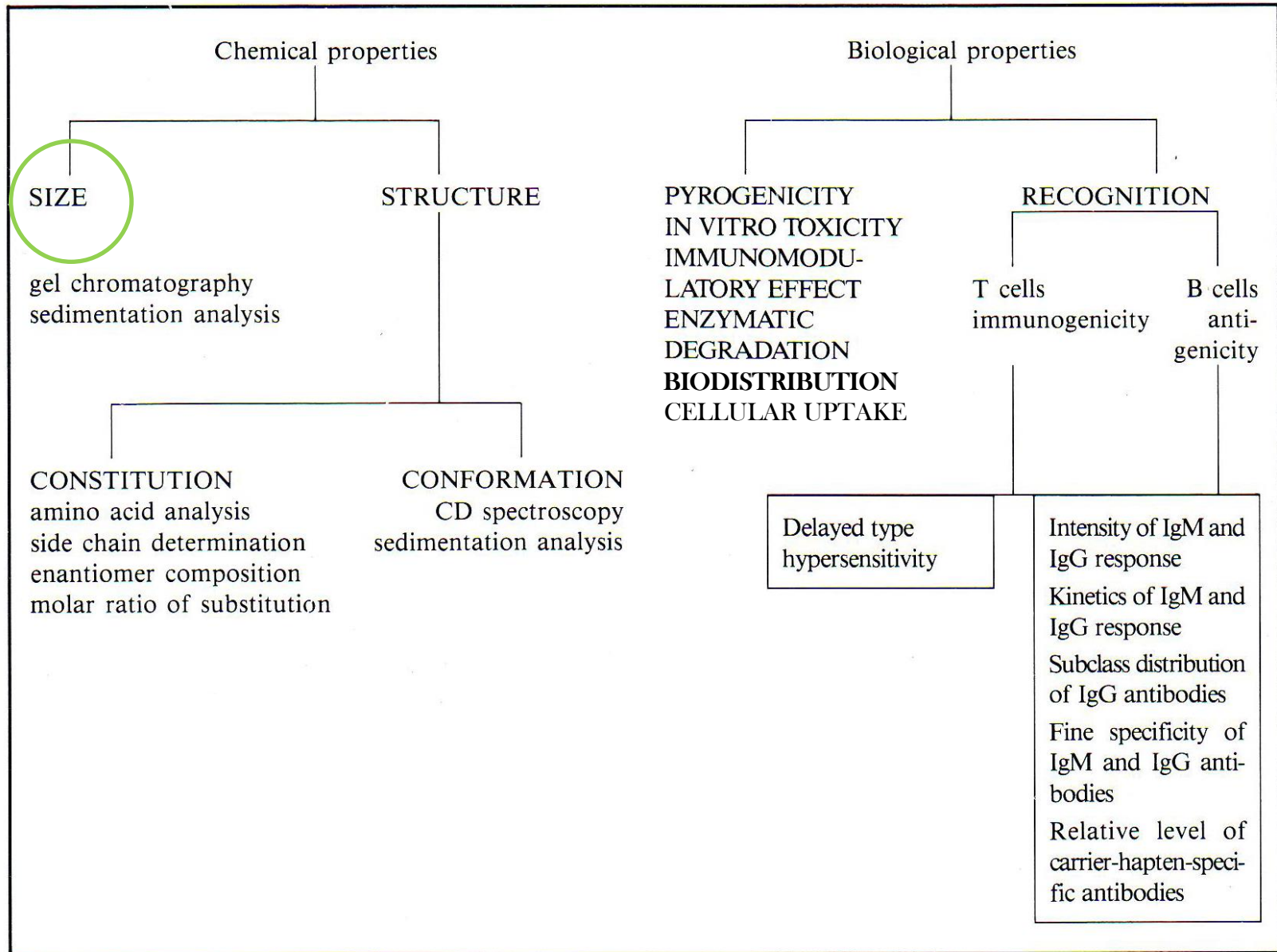
AXK type branched chain polypeptides



Characterization branched chain polypeptides



Characterization branched chain polypeptides

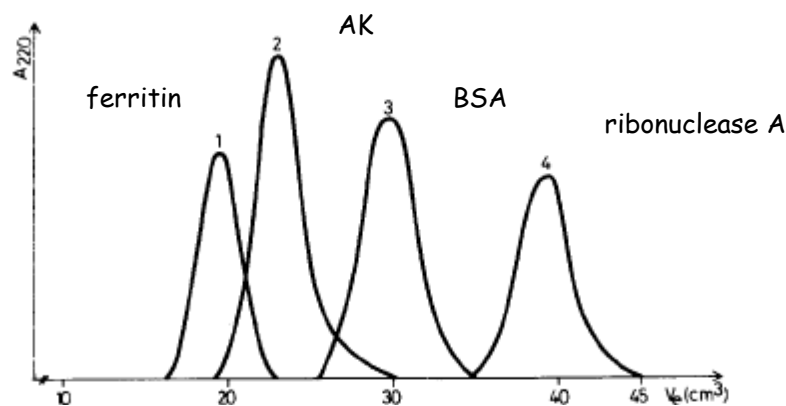


GPC, CD and sedimentation analysis of poly-Lys and branched chain poly-Lys–poly-DL-Ala polypeptides*)

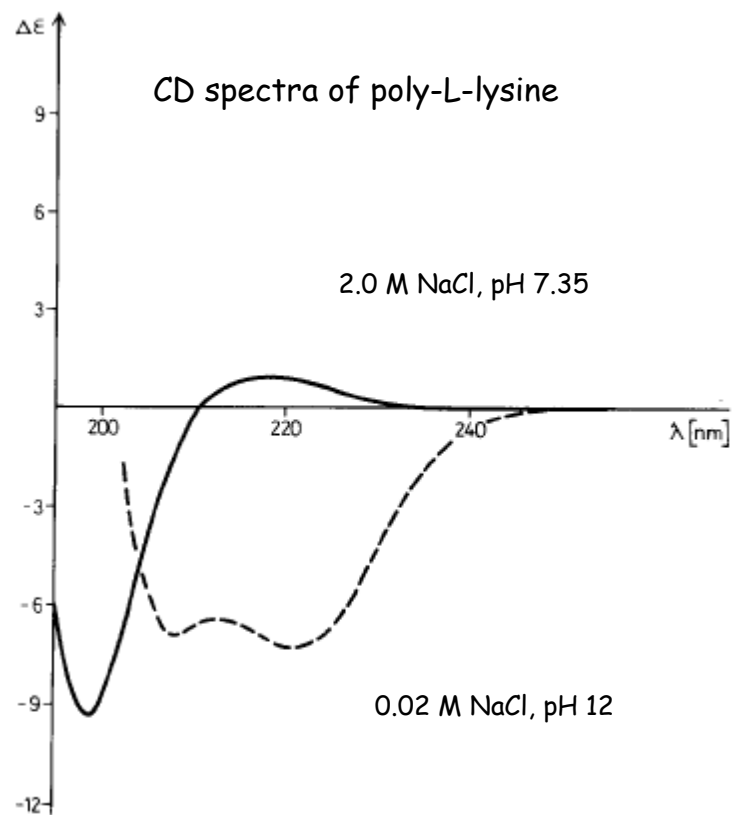
F. Hudecz, P. Kovács*), S. Kutassi-Kovács, and J. Kajtár**)

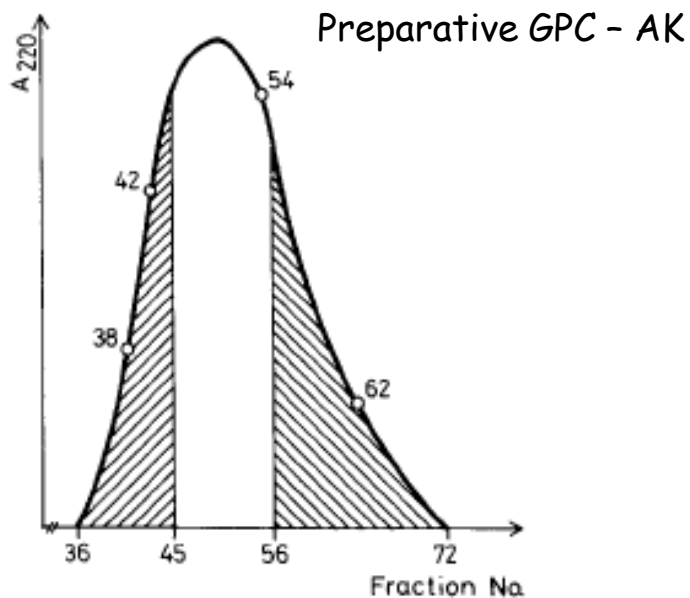
Table 1. Data on sedimentation analysis of poly-L-lysine samples and poly-L-Lys–poly-DL-Ala (AK)

	pLys		AK	
\bar{M}_w	30000	41000	94000	67000
\bar{M}_z	36000	45000	101000	77000
\bar{M}_z/\bar{M}_w	1.2	1.1	1.07	1.15
\bar{M}_n	25000	37000	88000	58000
\overline{DP}_n	120	180	430	174

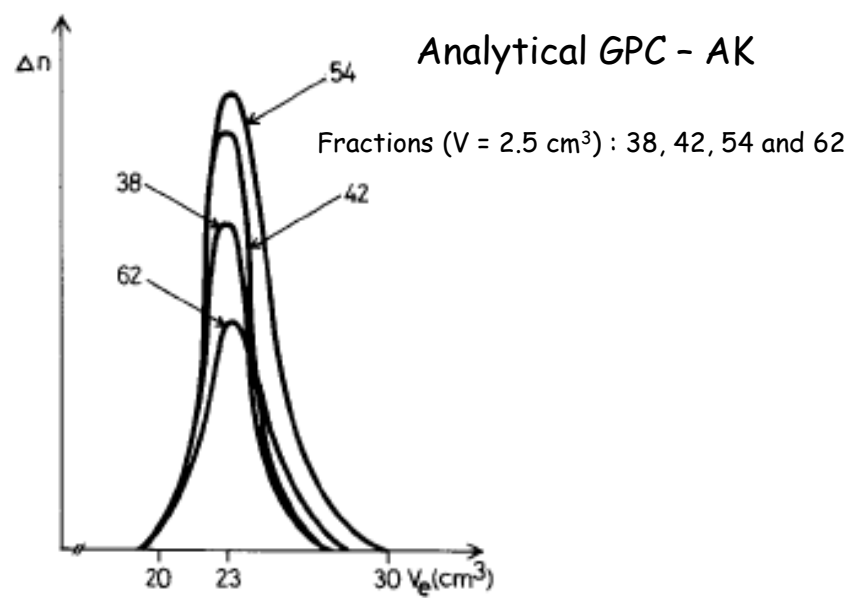


Sephadex G-200, 0.9 x 40 cm, 0.05 M NaHCO₃, 8 cm³/h



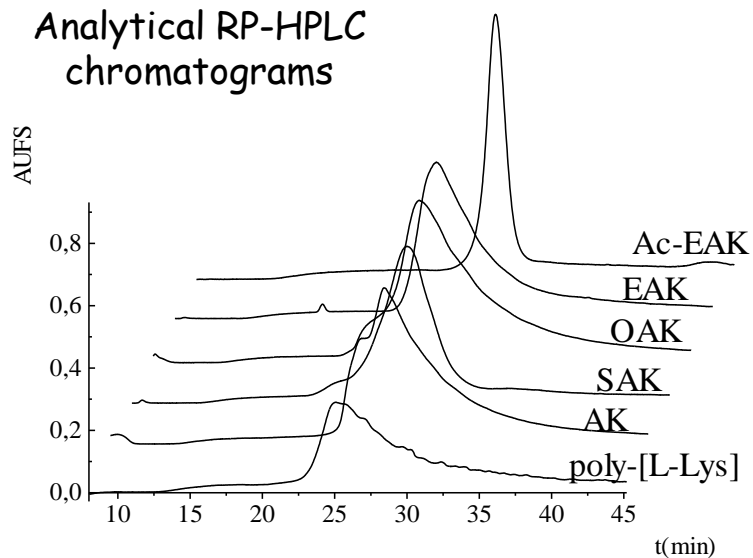


Sephadex G-200, 4 x 66 cm, 0.05 M NaHCO₃



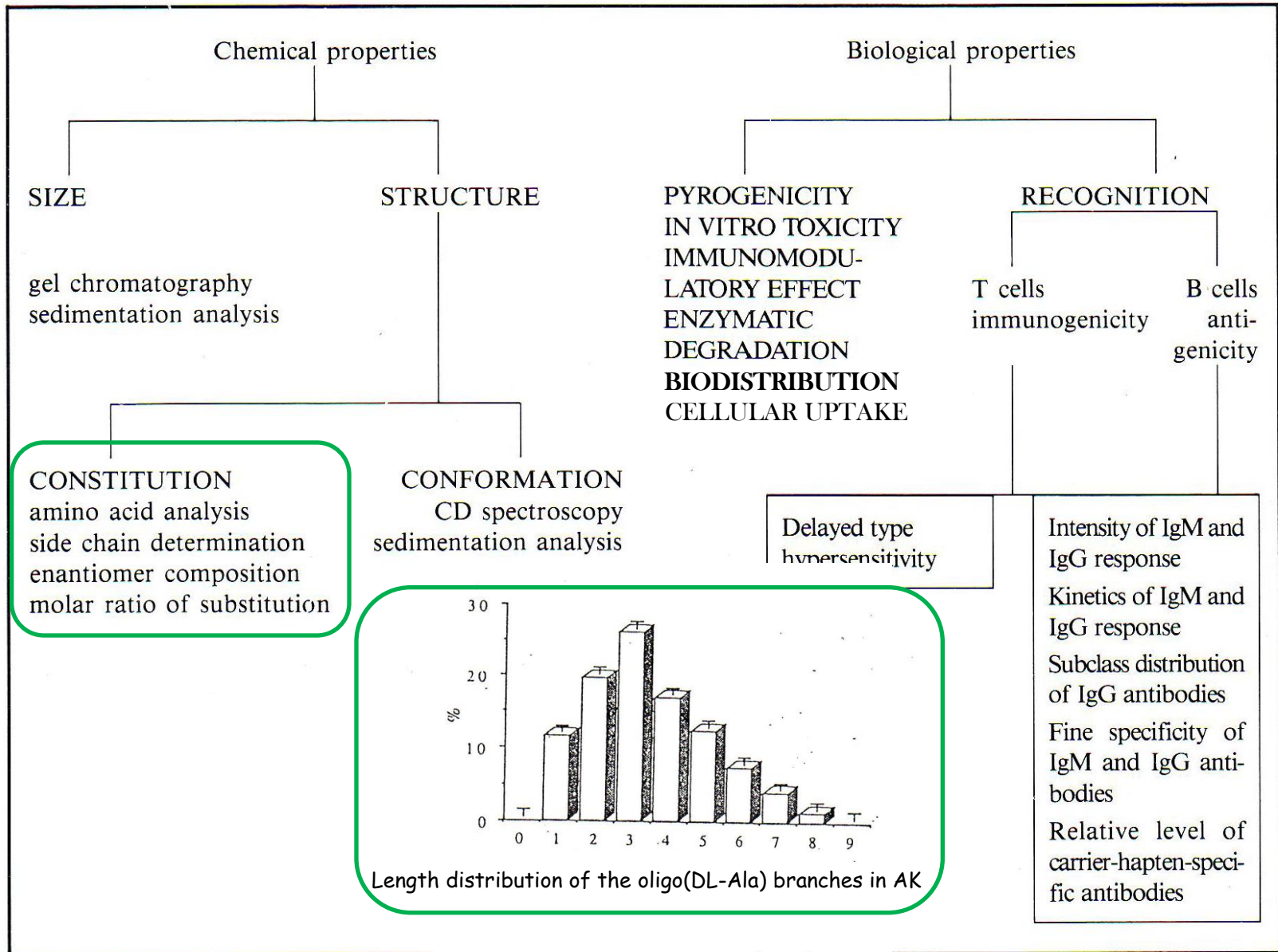
Sephadex G-200, 0.9 x 40 cm, 0.05 M NaHCO₃, 8 cm³/h

Analytical RP-HPLC chromatograms

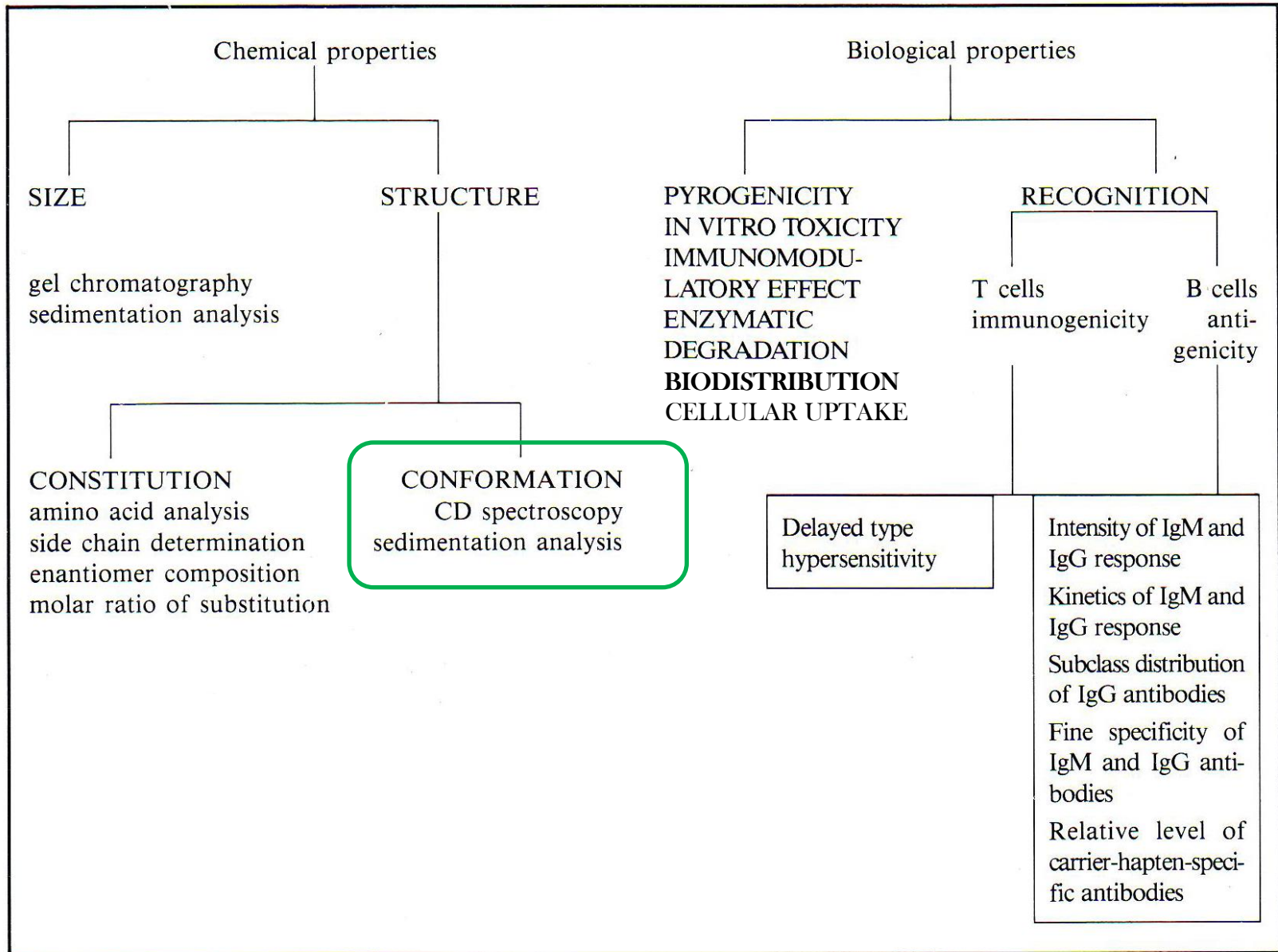


Waters (Milford, USA),
 Delta Pak C₁₈ column (300 x 3.9 mm I.D.)
 with 15 mm silica (300 Å pore size)
 eluent A: 0.1 % TFA
 eluent B: 0.1% TFA in AcN-water (80:20, v/v %)
 linear gradient 10% - 65% v/v of B in 45 min.
 flow rate: 2 ml/min
 UV detection at $\lambda = 208$ and 214 nm
 polymer samples: 3mg/ml

Characterization branched chain polypeptides



Characterization branched chain polypeptides



Primary structure - solution conformation

Amino acid X

- identity, character, charge
- position in the side the side chain
- number

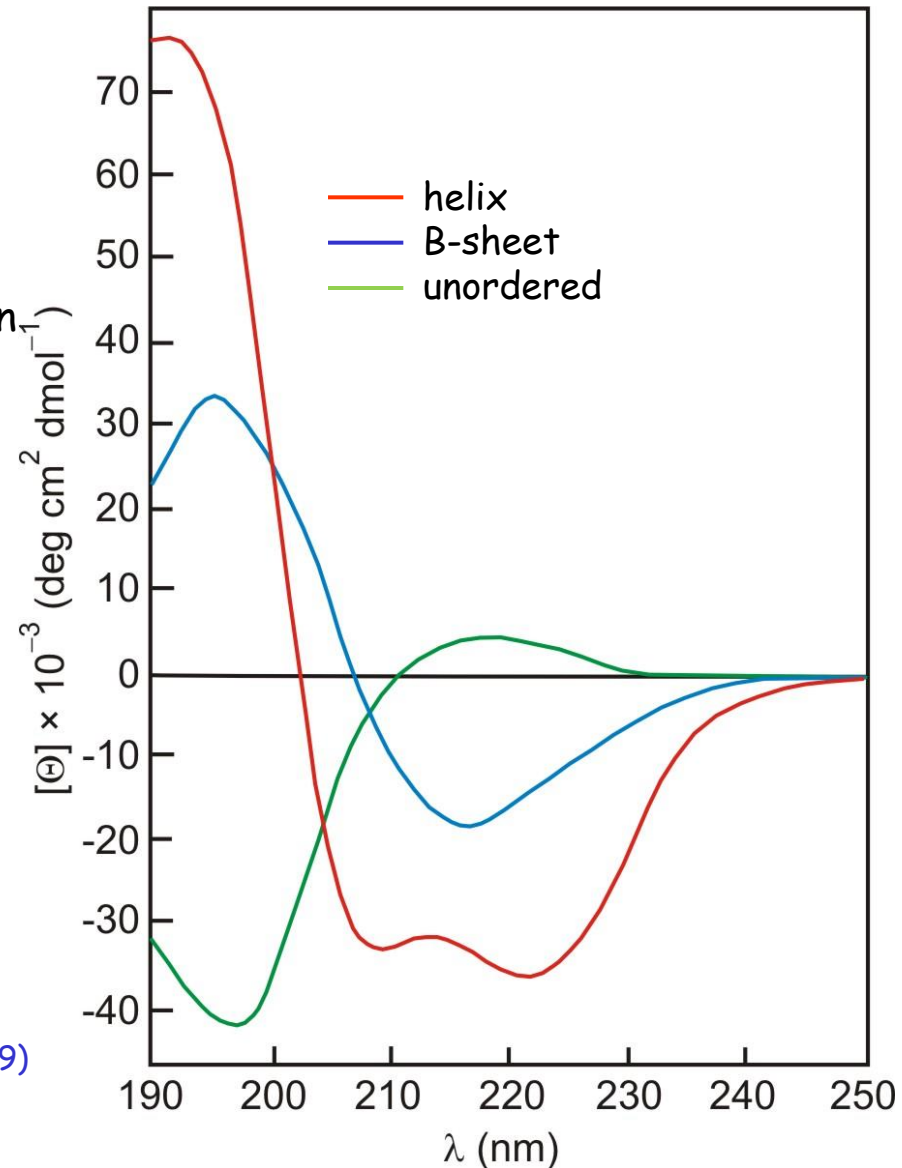
could influence ordered structure formation

Method: ECD spectroscopy

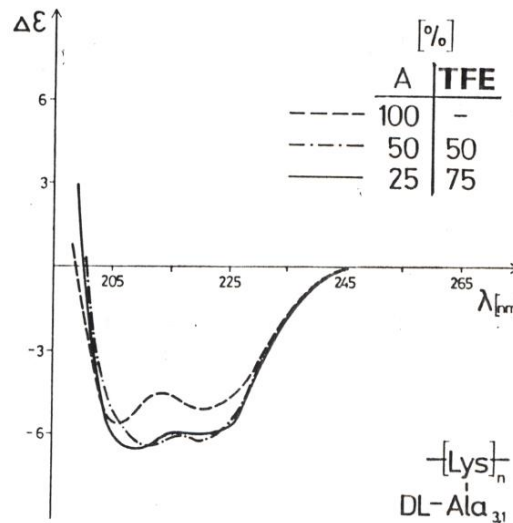
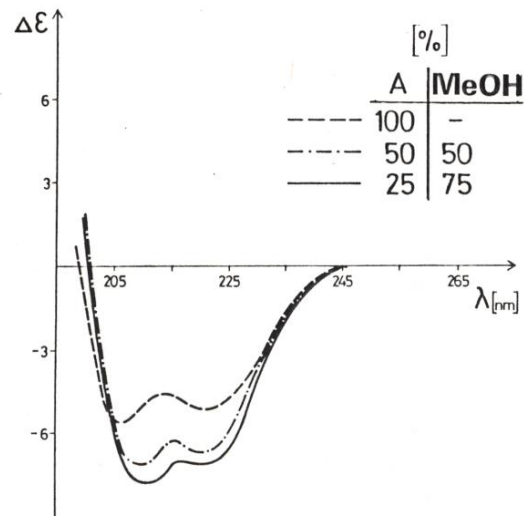
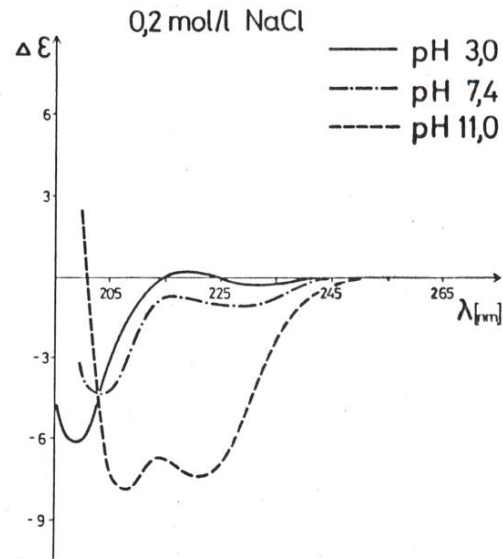
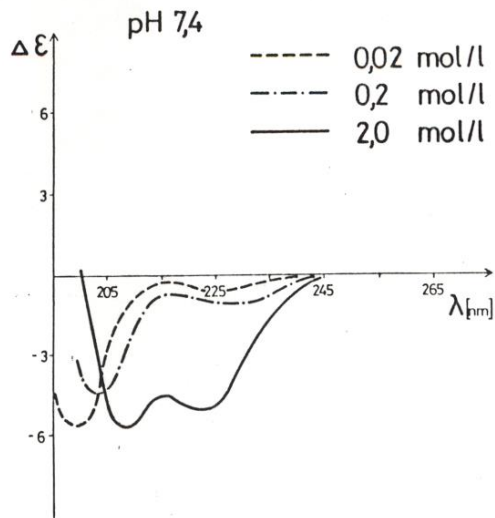
Conditions:

- pH: 3, 7.4 and 12
- Ionic strength:
0.02, 0.2, 2.0 mol/dm³ NaCl
- solvent:
water, MeOH/water or
TFE/water (25,50,75%)

N. Greenfield, G.D. Fasman: *Biochemistry* 8, 4108-4116 (1969)
N. Greenfield: *Nature Protoc.* 1, 2876-2880 (2006)



ECD spectrum - solution conformation

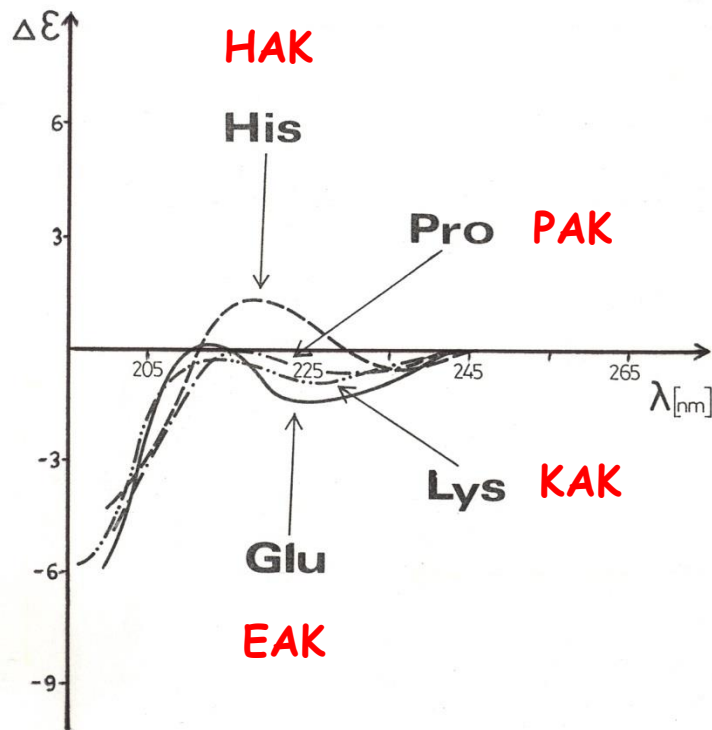


Conditions:

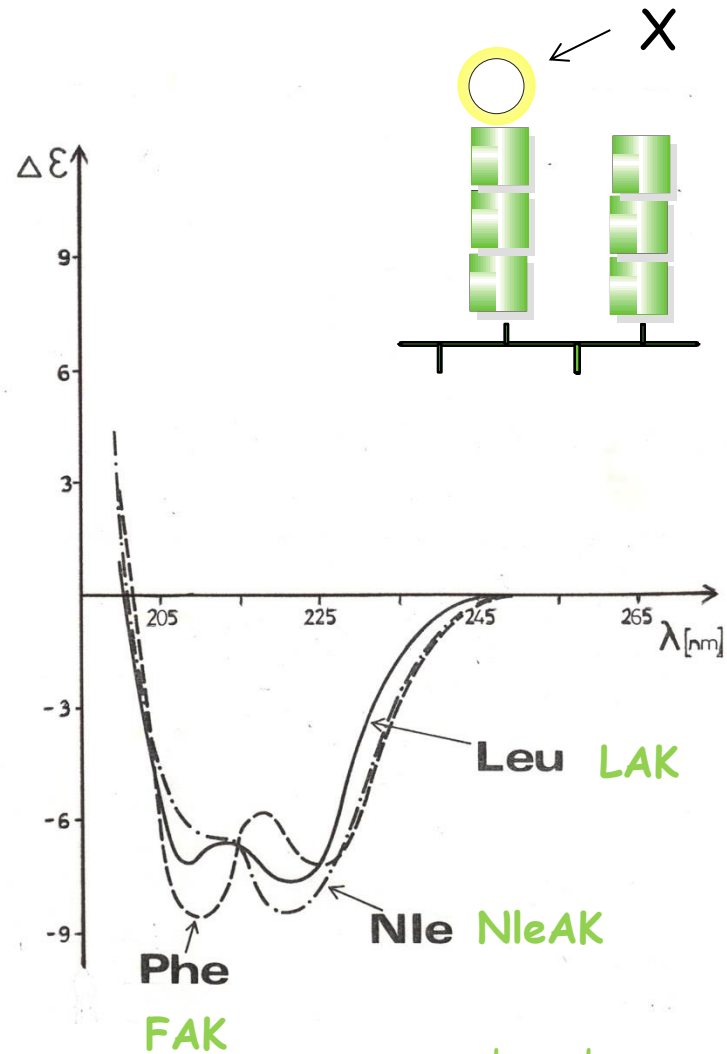
- pH: 3, 7.4 and 12
- ionic strength:
0.02, 0.2, 2.0 mol/dm³ NaCl
- solvent:
water, MeOH/water or
TFE/water (25,50,75%)

The effect of the identity of amino acid X

0.2 M NaCl, pH 7.3

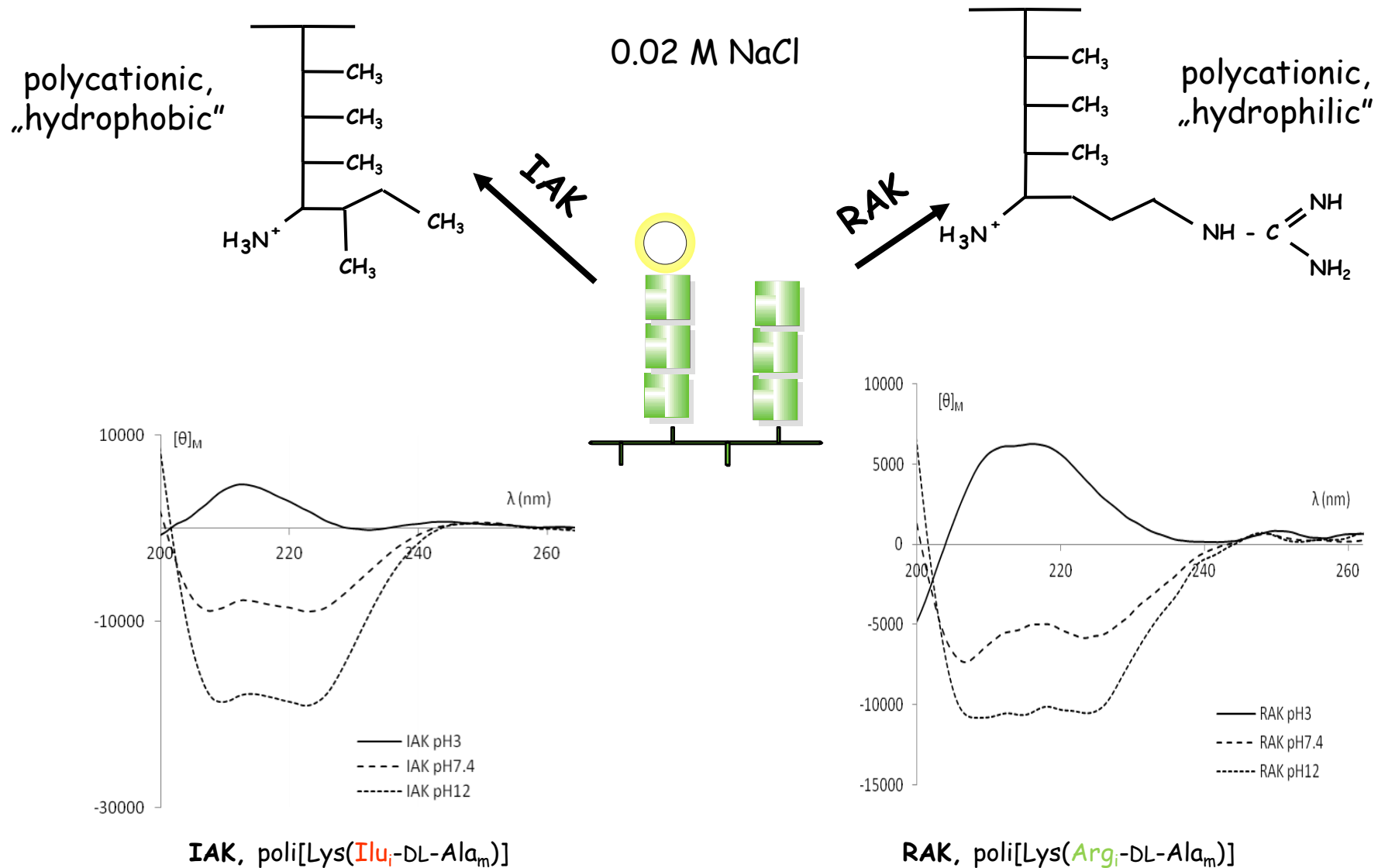


unordered

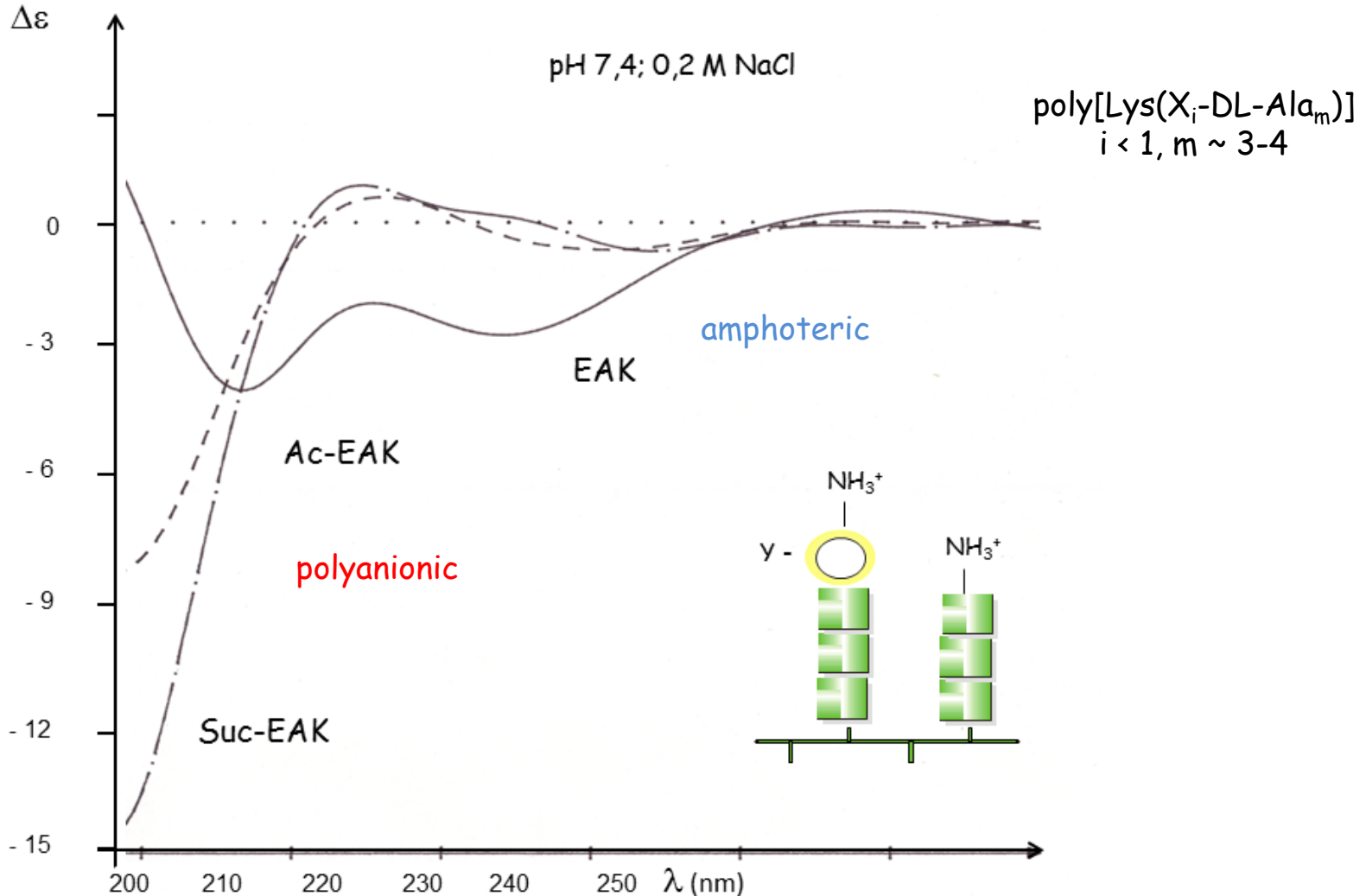


ordered

The effect of the identity of amino acid X : Ile or Arg

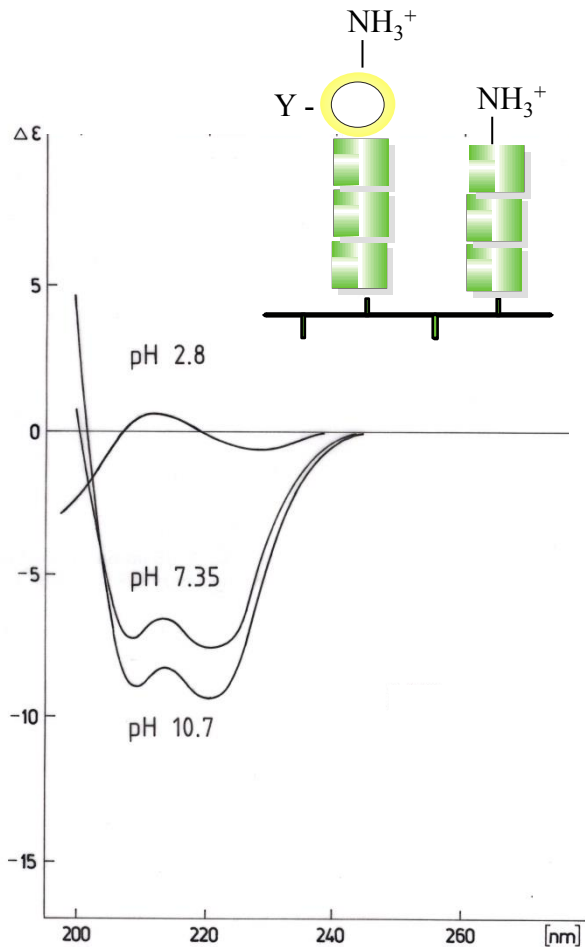


The effect of the charge of the terminal amino acid X : Glu (EAK), Ac-EAK or Suc-EAK

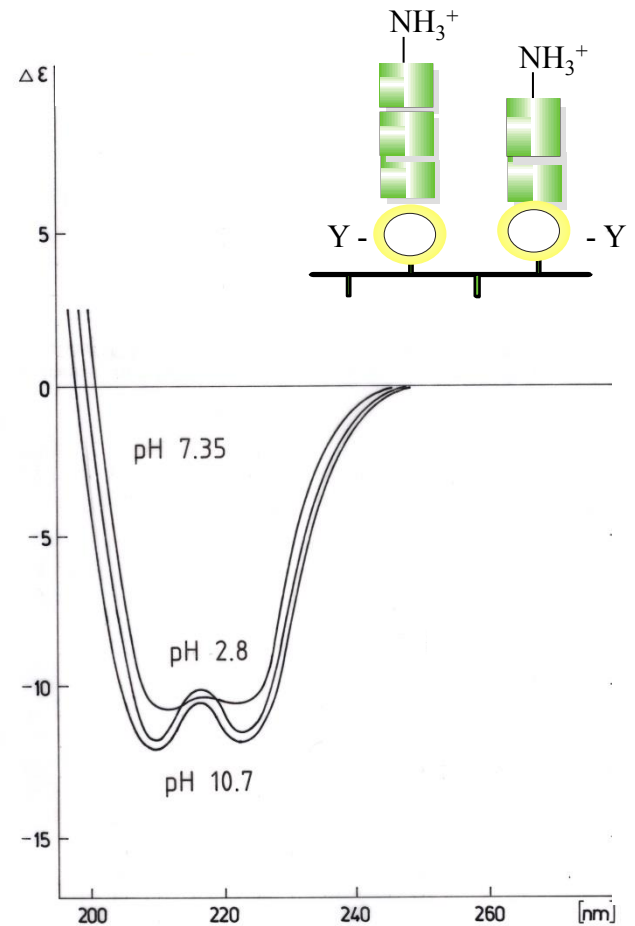


The effect of the position of hydrophobic amino acid X

0.2 M NaCl



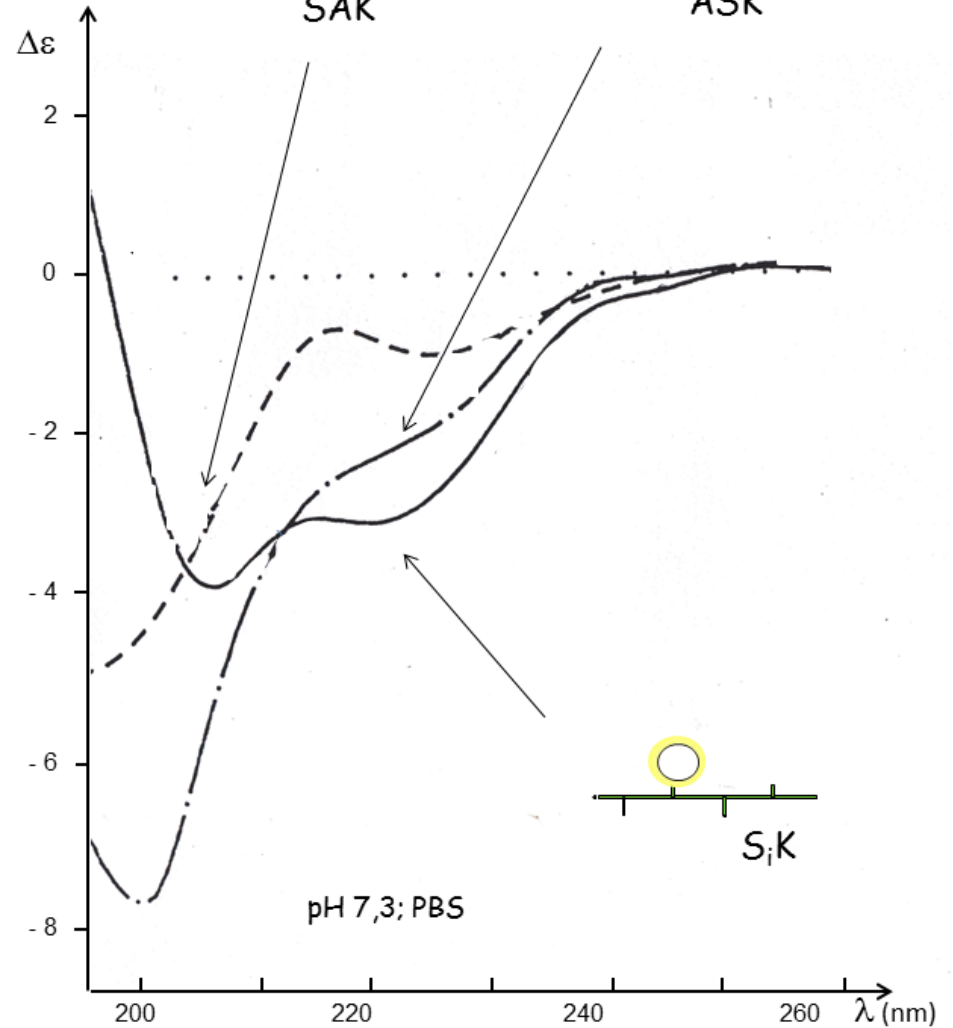
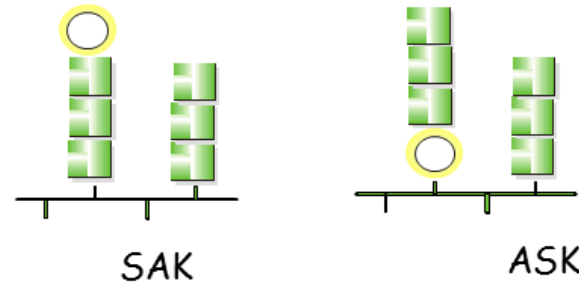
LAK, poly[Lys(Leu_i-DL-Ala_m)]



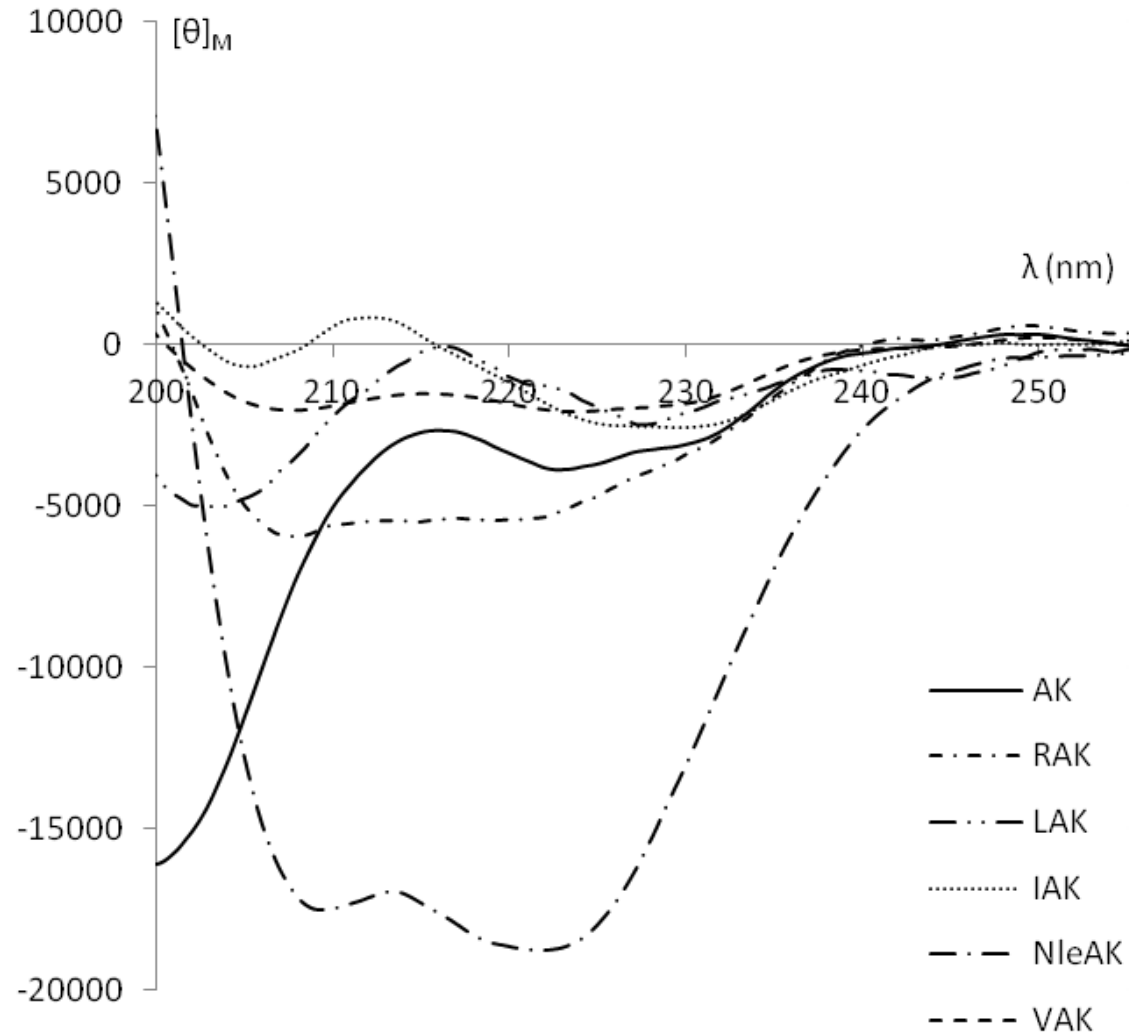
ALK, poly[Lys(DL-Ala_m-Leu_i)]

The effect of the position of hydrophilic amino acid X

X = Ser

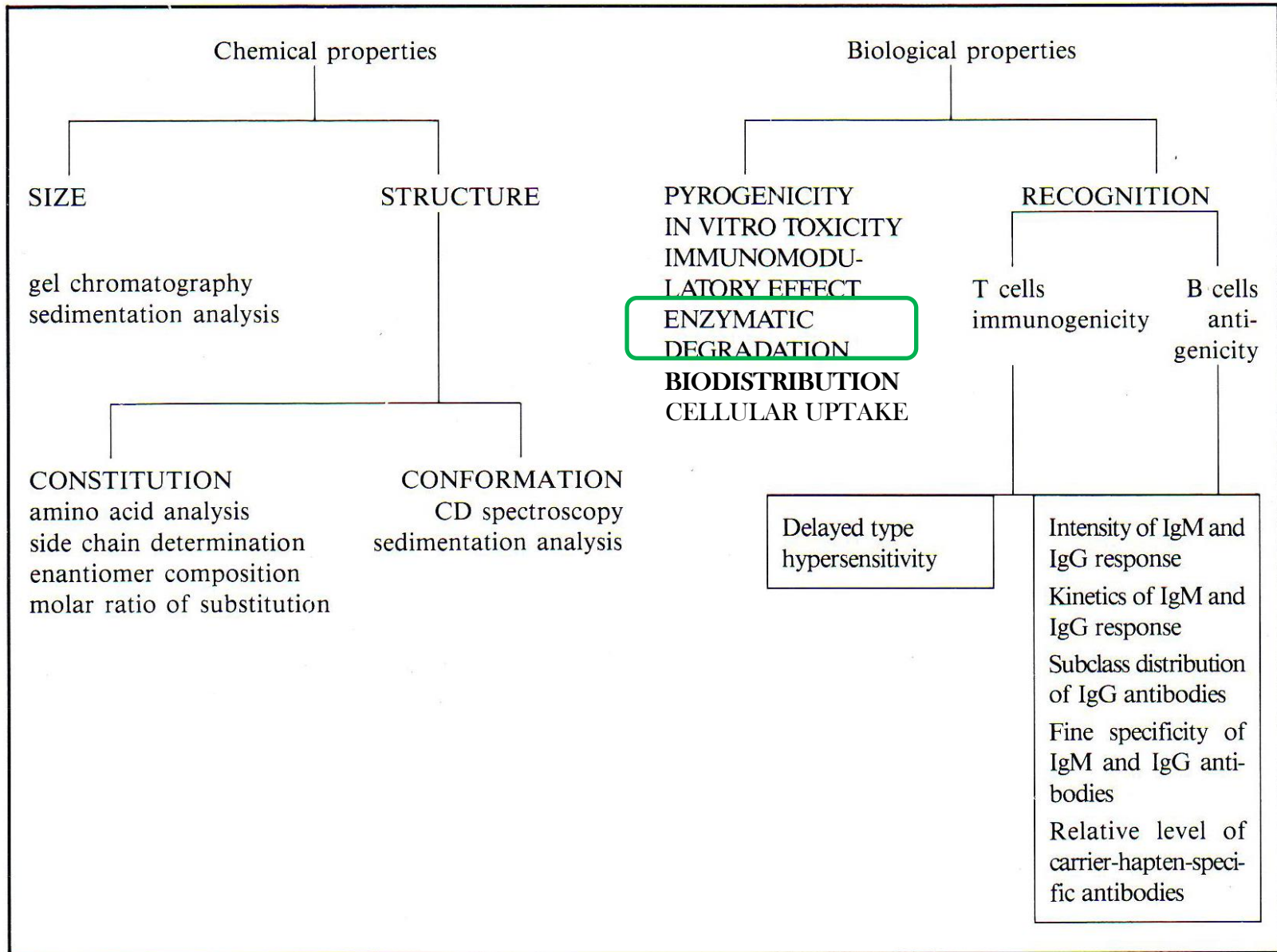


ECD spectra of branched polypeptides under nearly physiological conditions - a summary

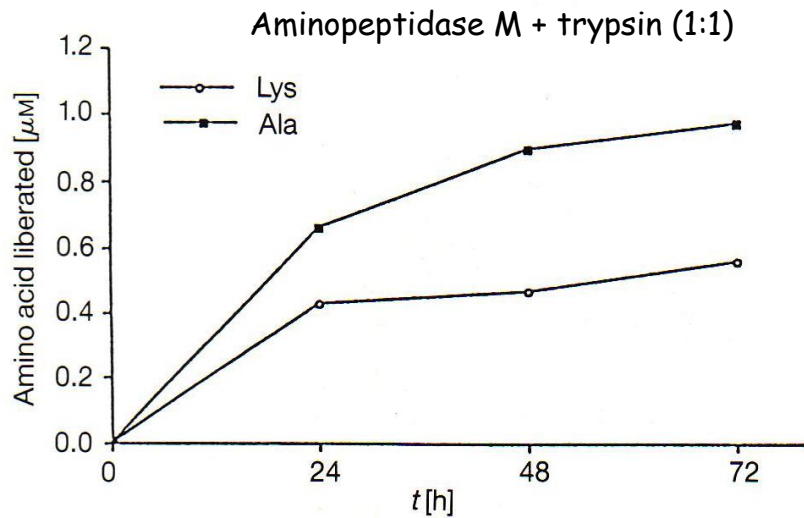
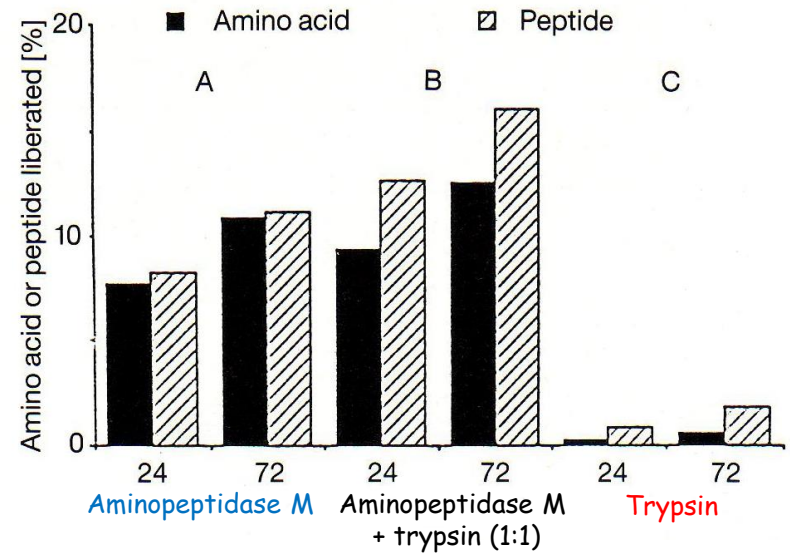
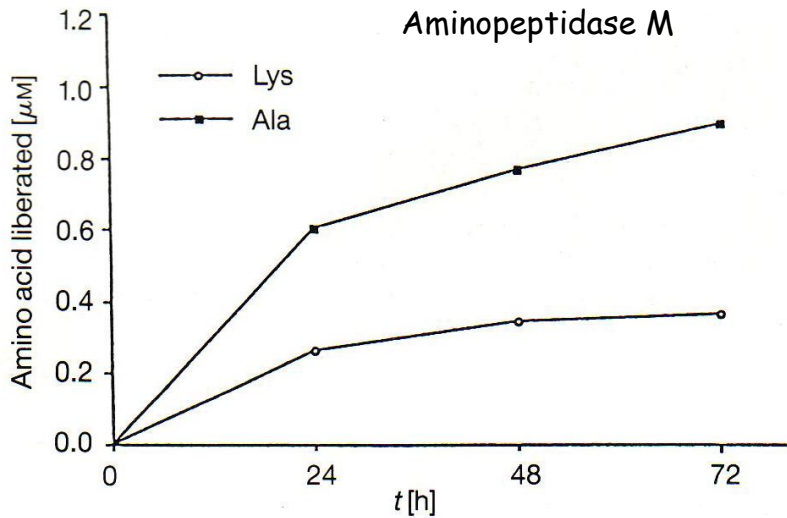


0.2 M NaCl
pH ~7.4
room temperature

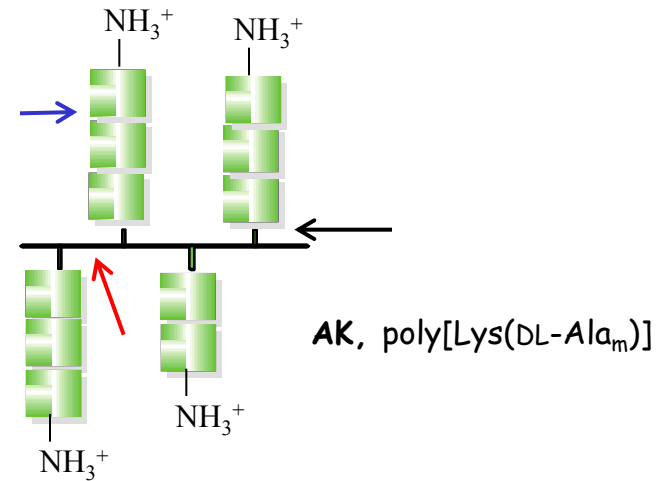
Characterization branched chain polypeptides



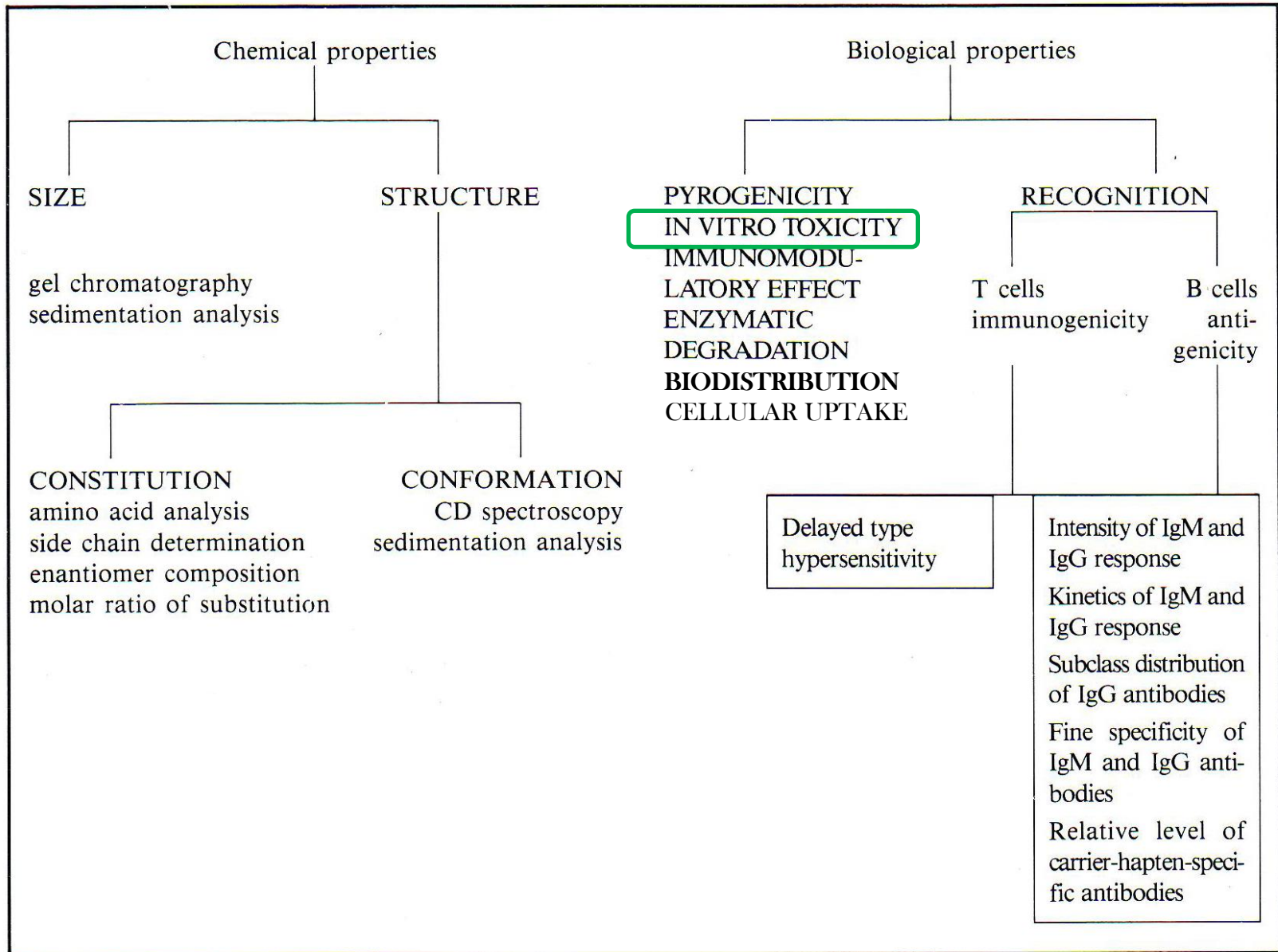
Branched polypeptides - biodegradation



0.02 M phosphate, pH 8

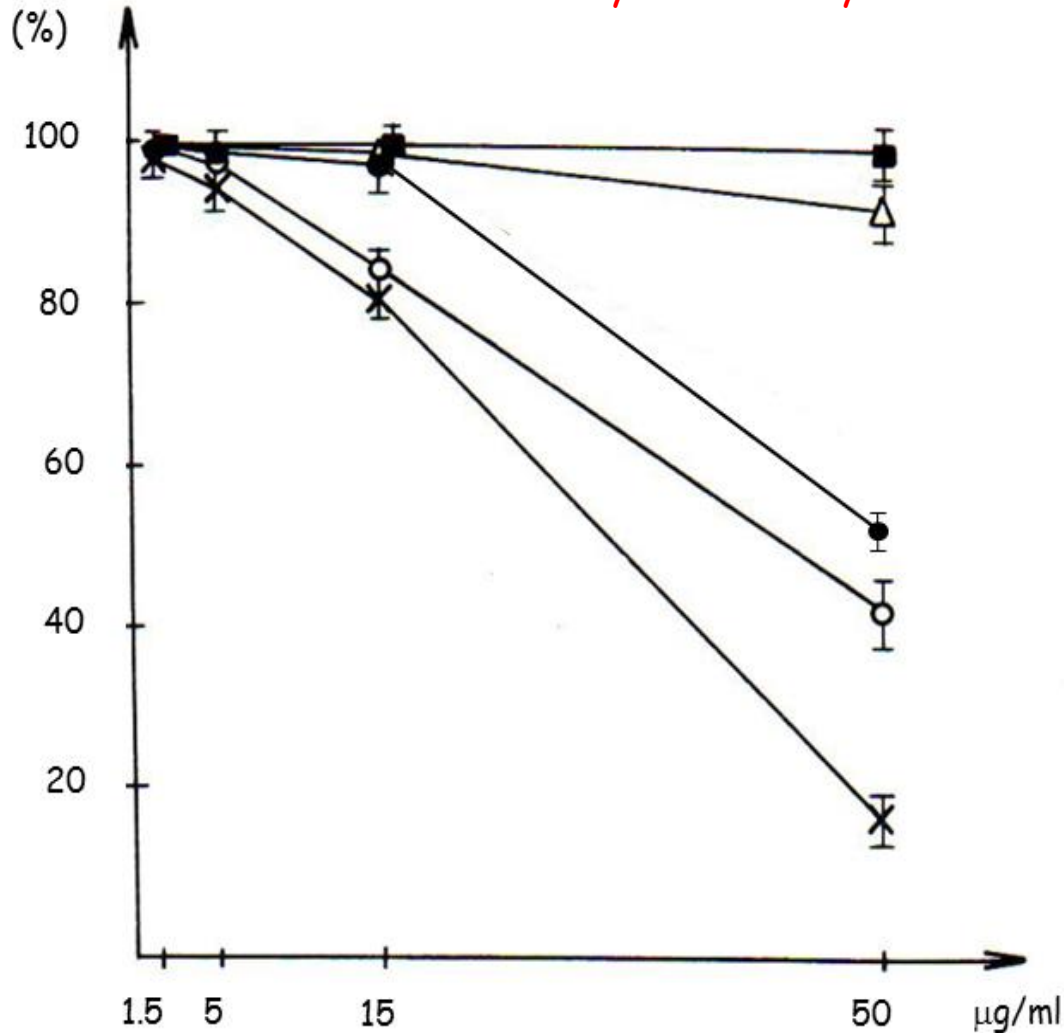


Characterization branched chain polypeptides



The effect of position and identity of amino acid

in vitro cytotoxicity on isolated rat liver cells

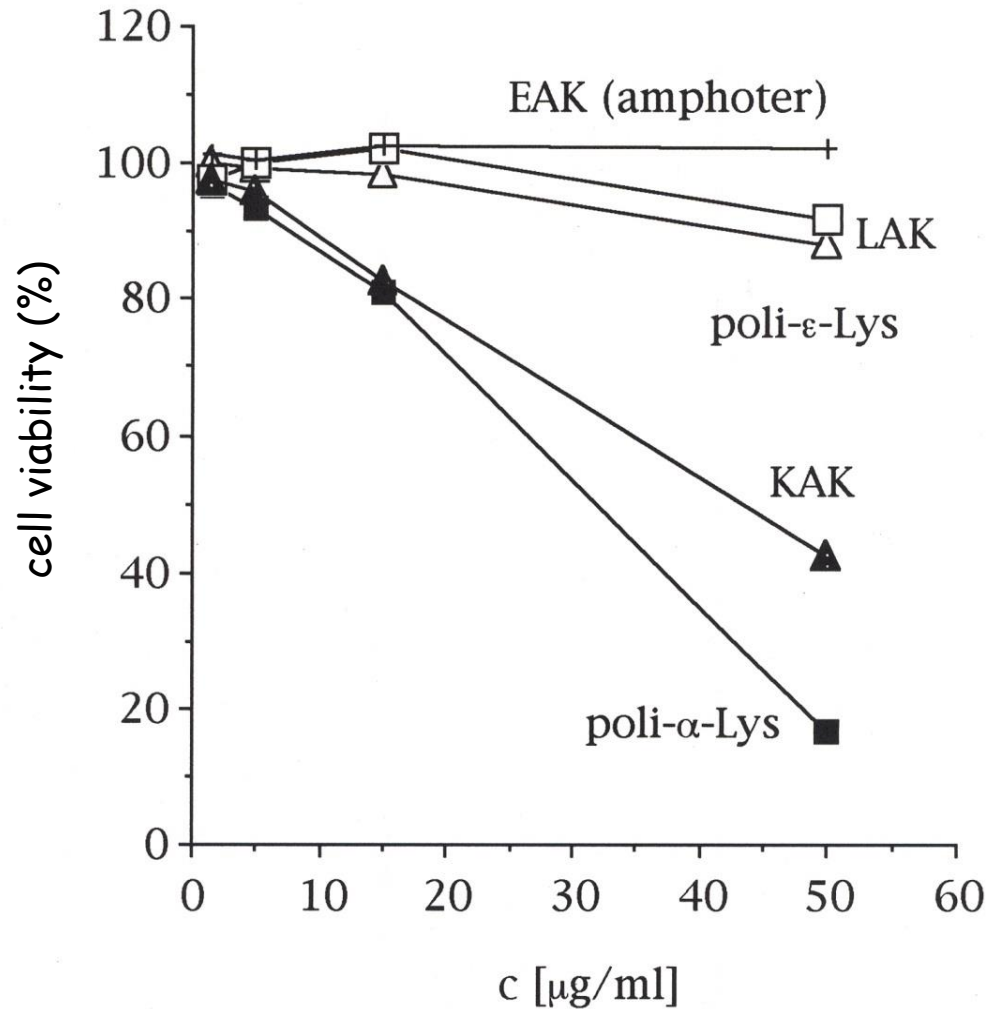


- = EAK, AEK, SAK, ASK
- Δ = LAK, ALK, D-LAK
FAK, D-FAK
HAK, D-HAK, PAK,
D-EAK
- = AK (m = 3)
- = KAK, D-KAK
- × = poly[Lys]

Incubation: 1 hr, 37° C

The effect of charge - the identity of amino acid X:

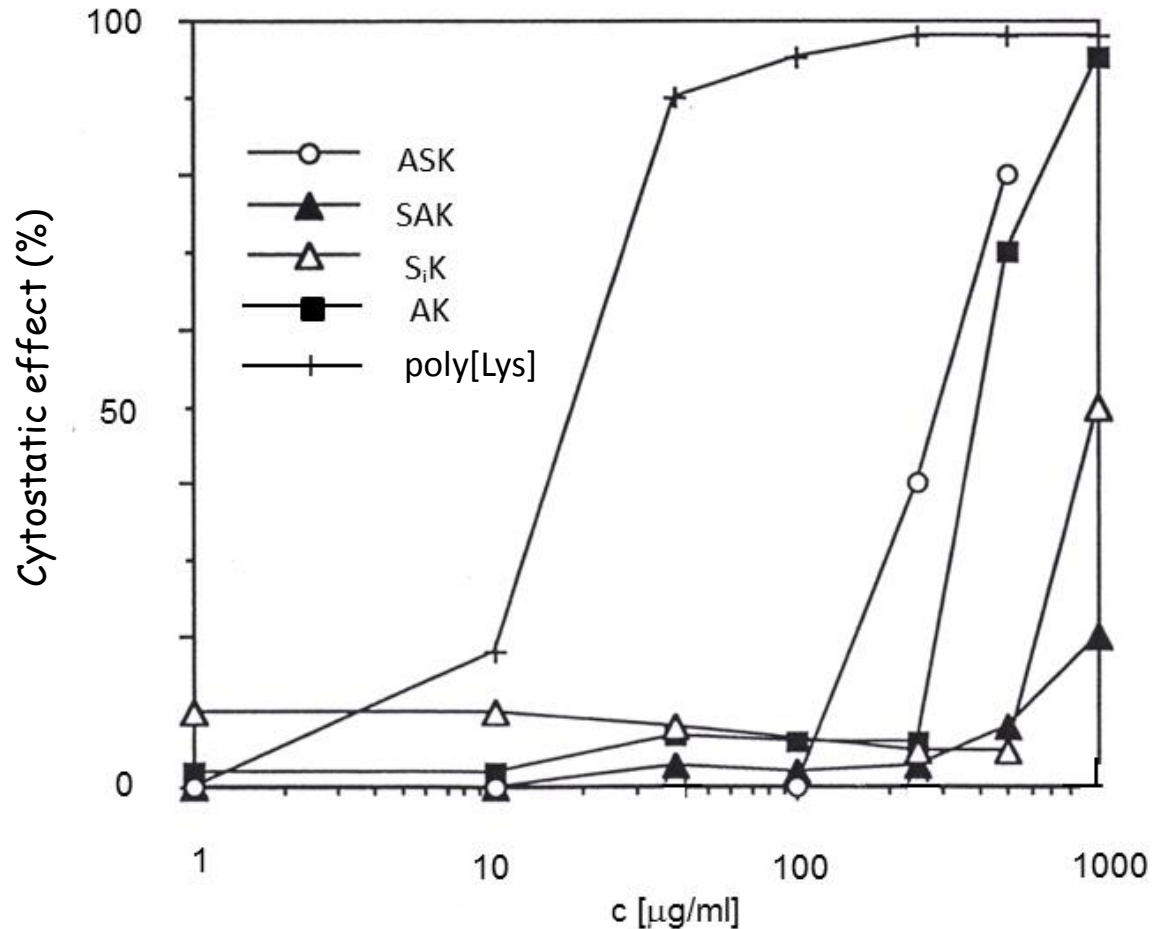
in vitro cytotoxicity HeLa cells



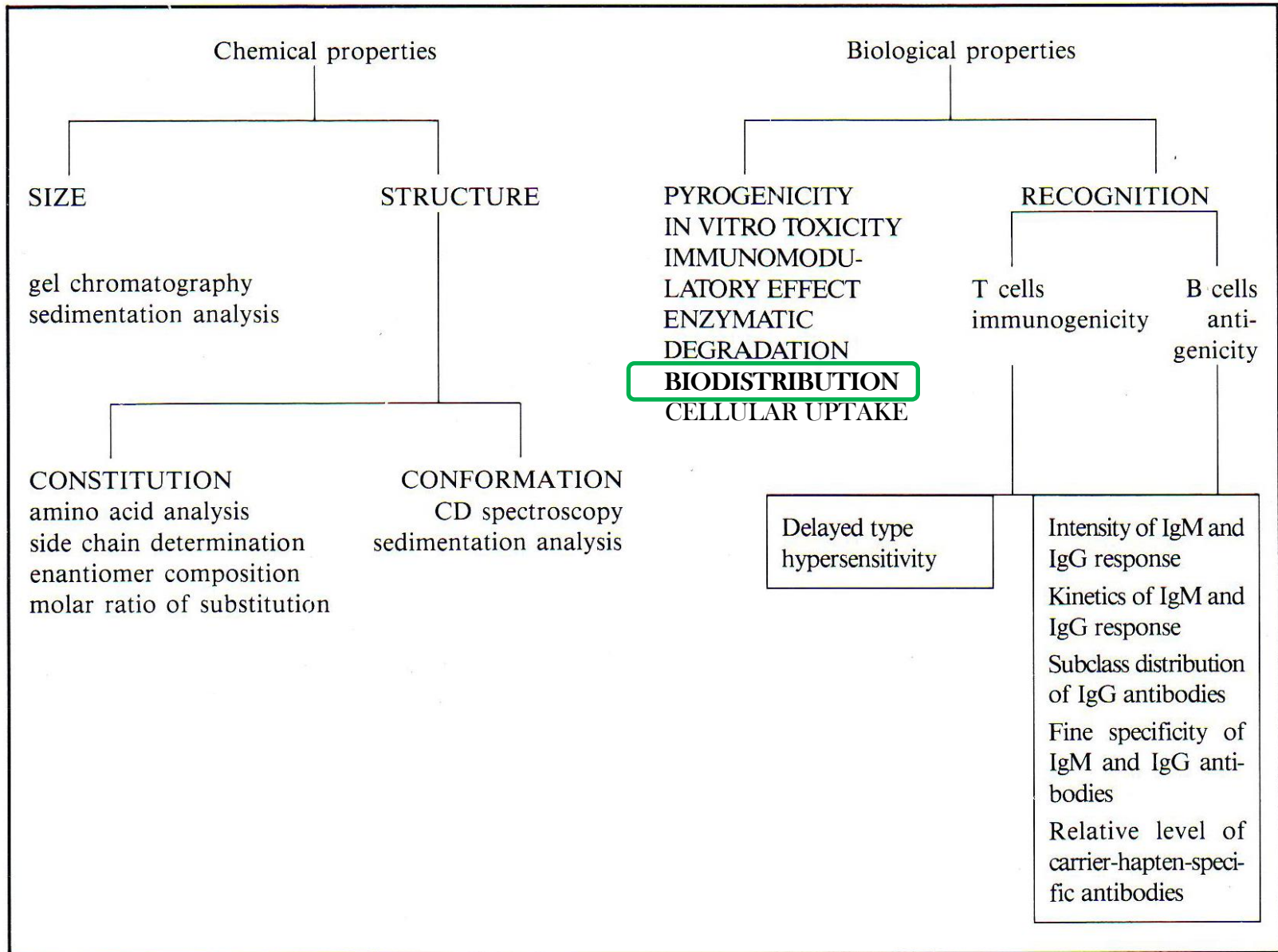
The effect of the position of hydrophilic amino acid (X = Ser) on *in vitro* cytostatic activity

C26 mouse carcinom cells

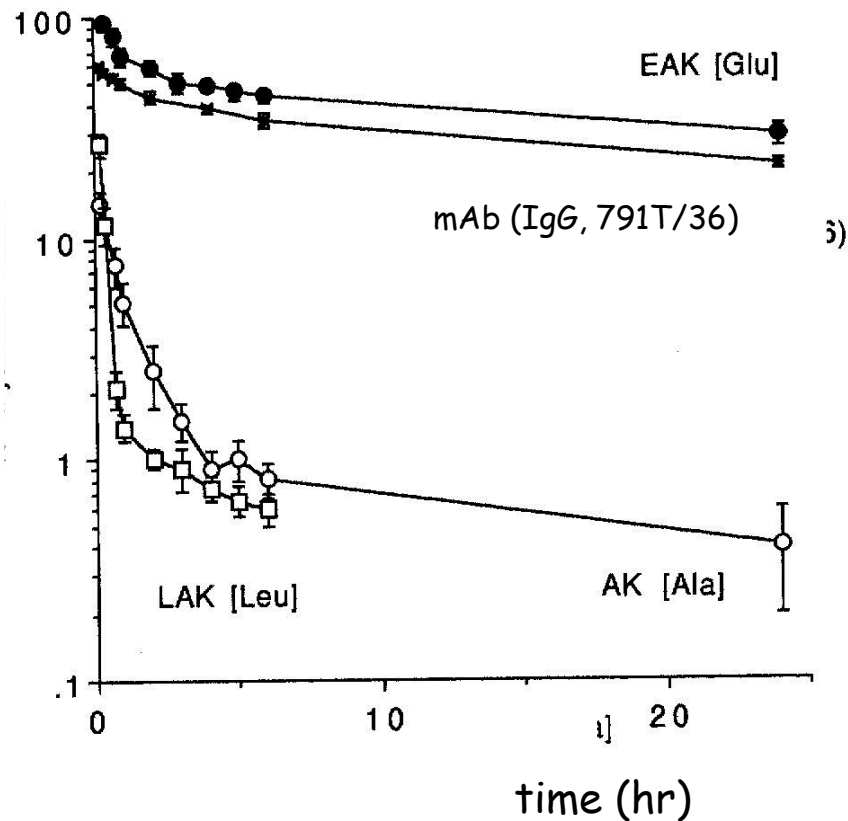
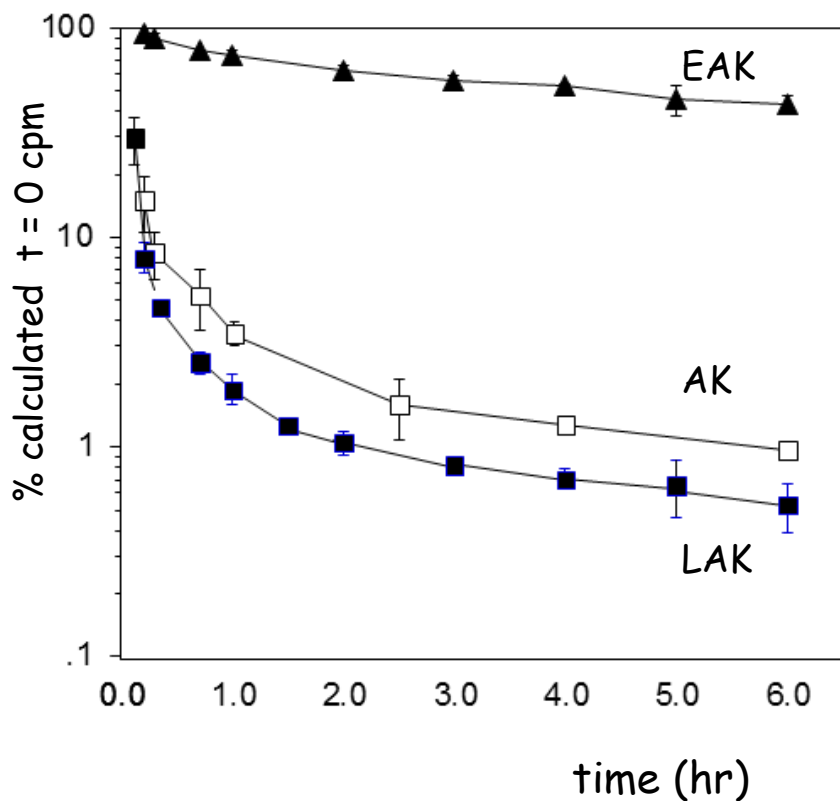
Incubation: 3 hrs, 37° C



Characterization branched chain polypeptides

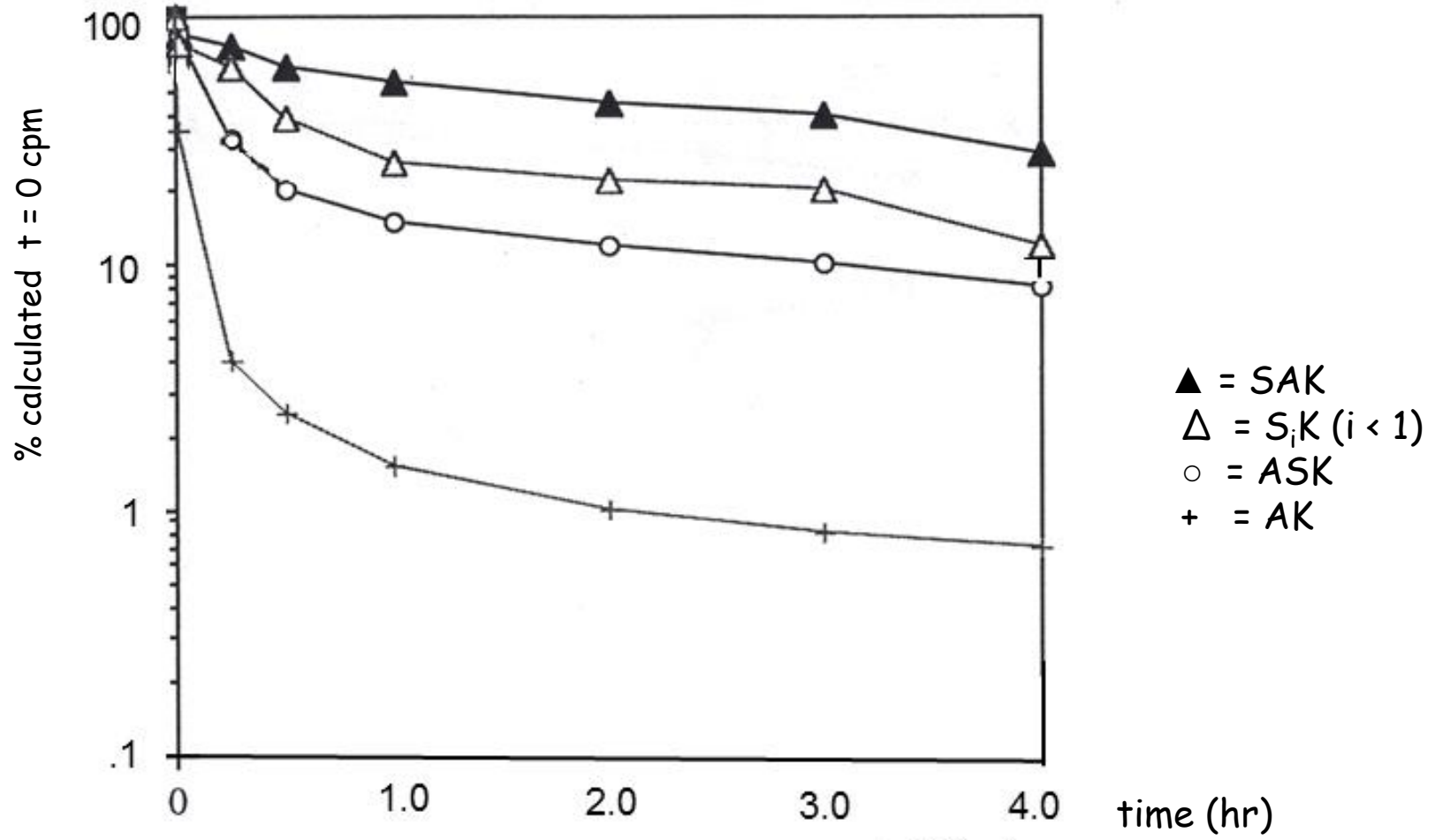


Blood clearance: the effect of the identity of amino acid X comparison with mAb



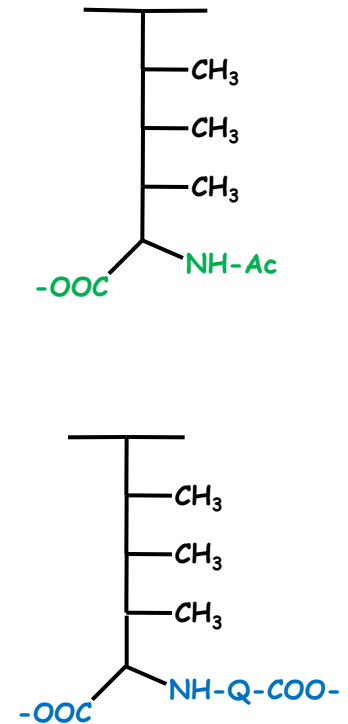
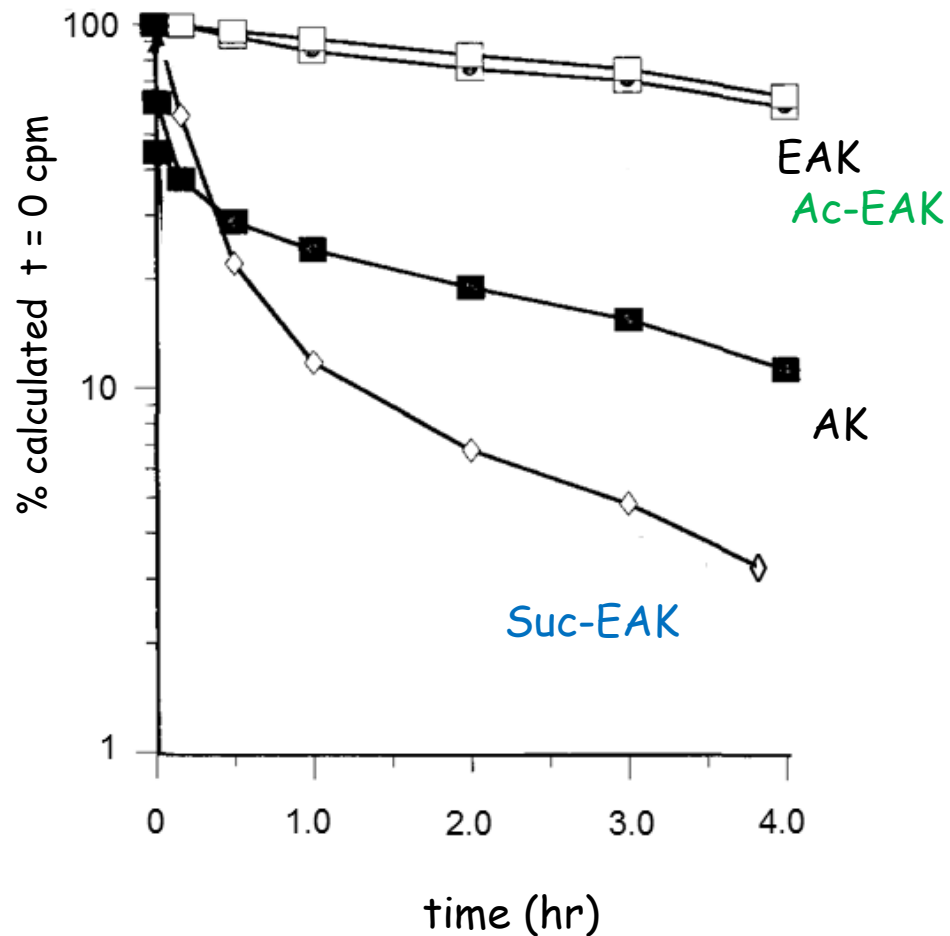
I^{125} isotope labelled polypeptides/protein i.v. injection

The effect of the position of hydrophilic amino acid (X = Ser) on blood clearance in Balb/c mice



I^{125} isotope labelled polypeptides i.v. injection

The effect of the position of hydrophilic amino acid X on blood clearance

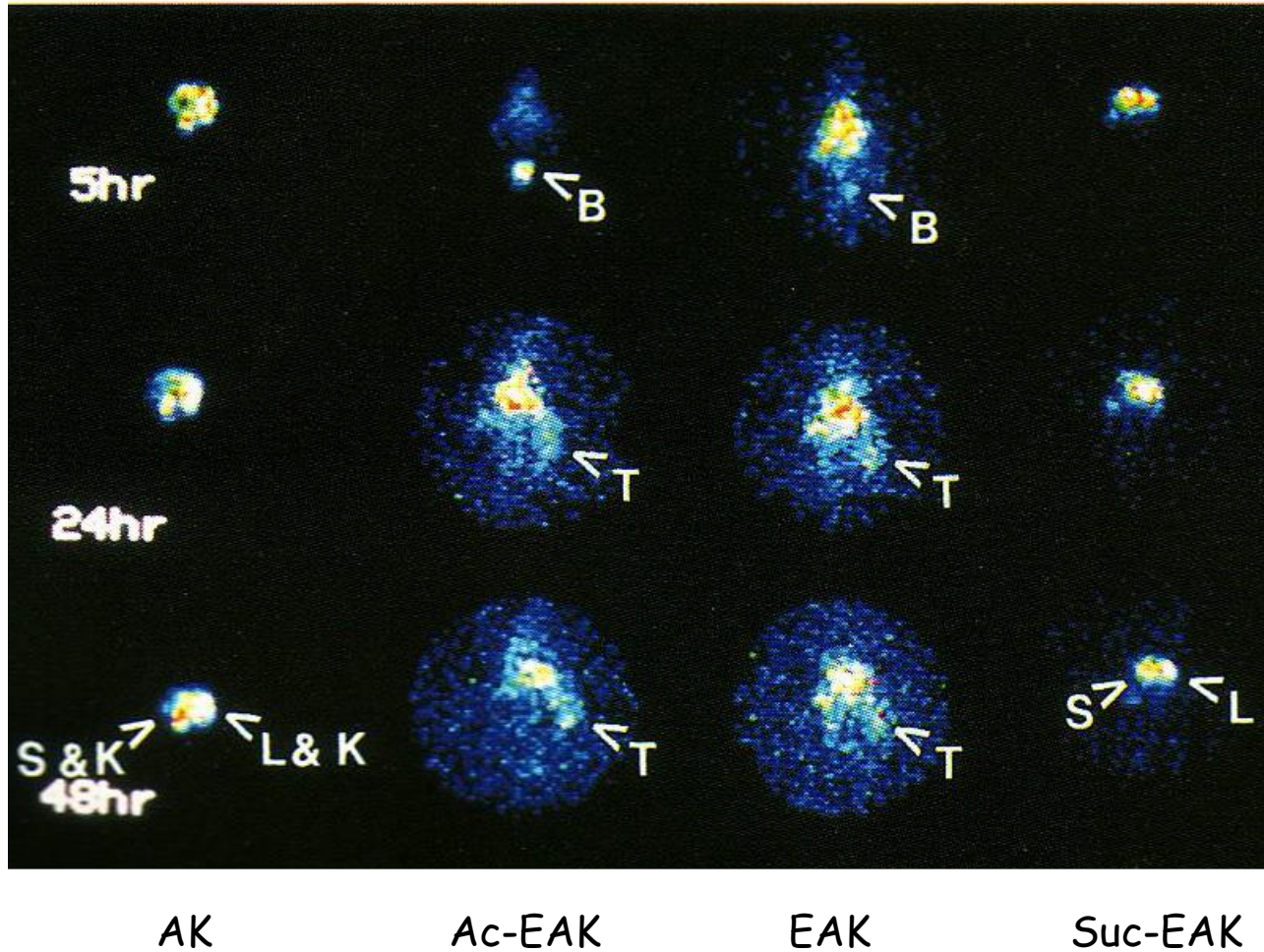


In^{111} isotope labelled polypeptides i.v. injection

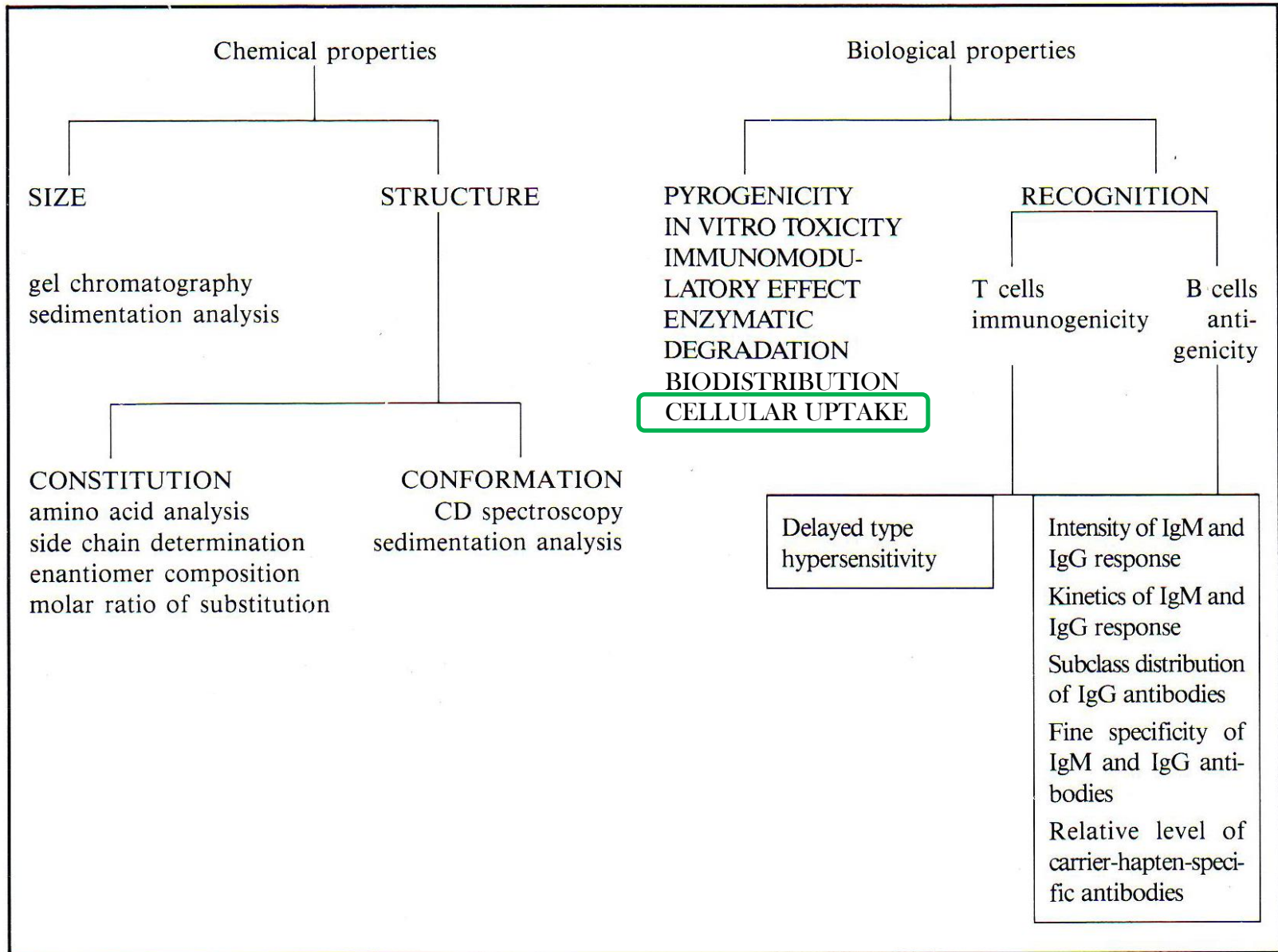
The effect of the charge of amino acid X in XAK branched polypeptide on tissue distribution in mice with mammary carcinoma

carcinoma graft size:
1.8 x 1.2 cm

B = urinary bladder
T = tumour
S = spleen
K = kidney



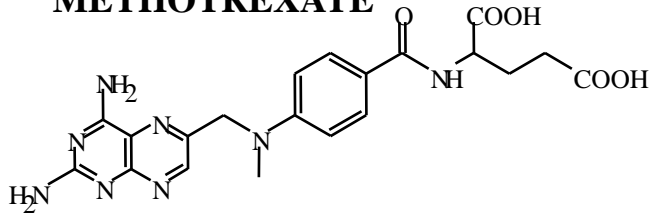
Characterization branched chain polypeptides



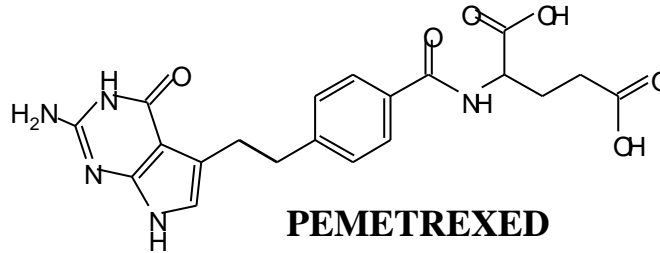
The effect of the identity and position of amino acid X
on cellular uptake

Drug, epitope, reporter molecule

METHOTREXATE

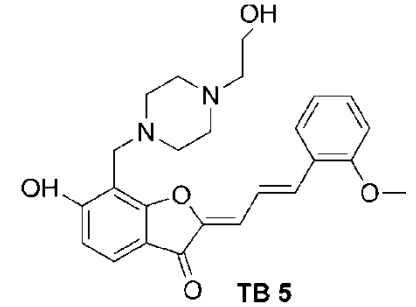


Hudecz F. et al. *Bioconjugate Chem.* **4**: 25 (1993)
Kóczán Gy. et al. *Bioconjugate Chem.* **13**: (2002)



PEMETREXED

Miklán Zs. et al.
J. Peptide Sci. **17**: 805 (2011)



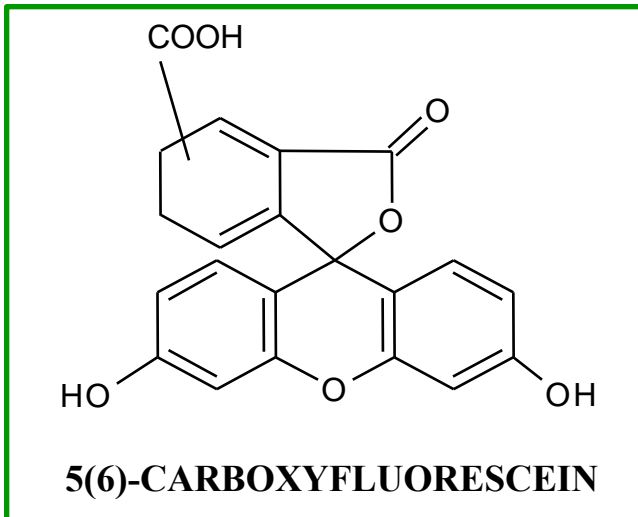
TB 5

Horvati, K.. et al.
Bioconjugate Chem. **22**:981 (2012)

D-Trp-*D*-Cpa-*D*-Trp-Ser-Tyr-*D*-Lys-Leu-Arg-Pro-*D*-Ala

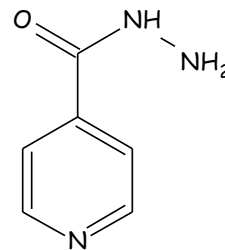
GN-RH ANTAGONIST, MI-1544

Mező, G. et al. *Bioconjugate Chem.* **7**: 642 (1996)
Vincze, B. et al. *J. Cancer Res. Clin. Onc.* **120**: 578 (1994)



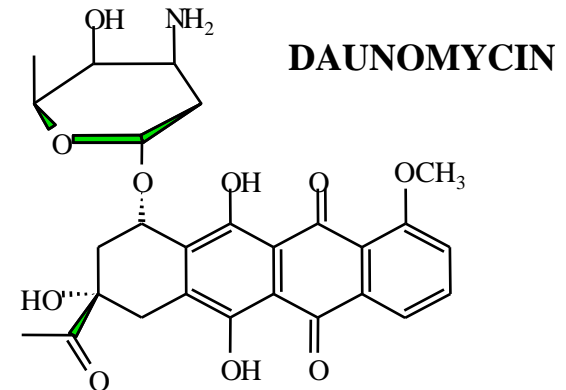
5(6)-CARBOXYFLUORESCIN

Szabó R. et al. *Bioconjugate Chem.* **19**: 1078 (2008)
Bánóczy Z. et al. *Bioconjugate Chem.* **19**: 1375 (2008)



ISONIAZID

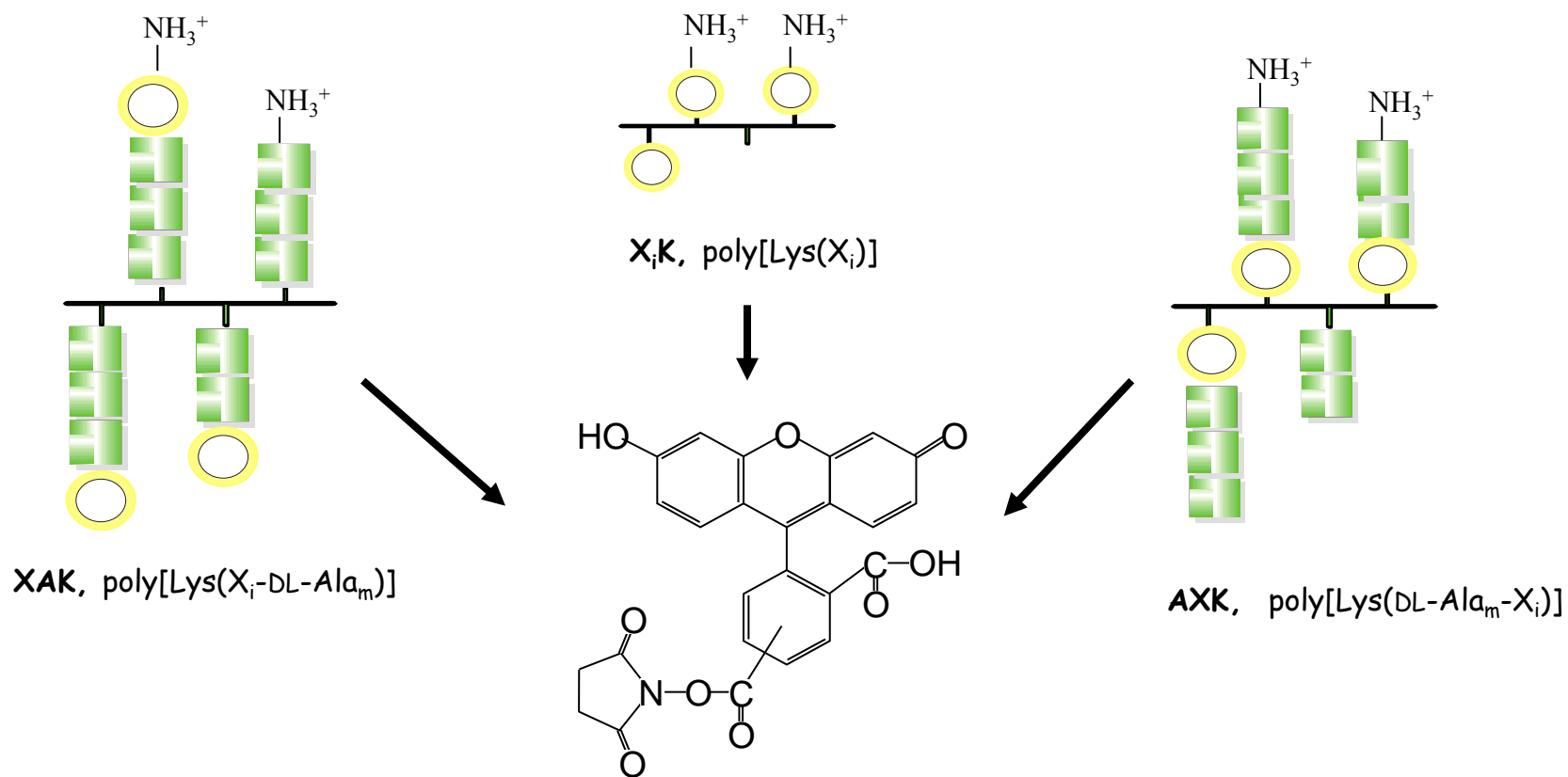
Horvati, K.. et al.
J. Peptide Sci. **15**:385 (2009)



DAUNOMYCIN

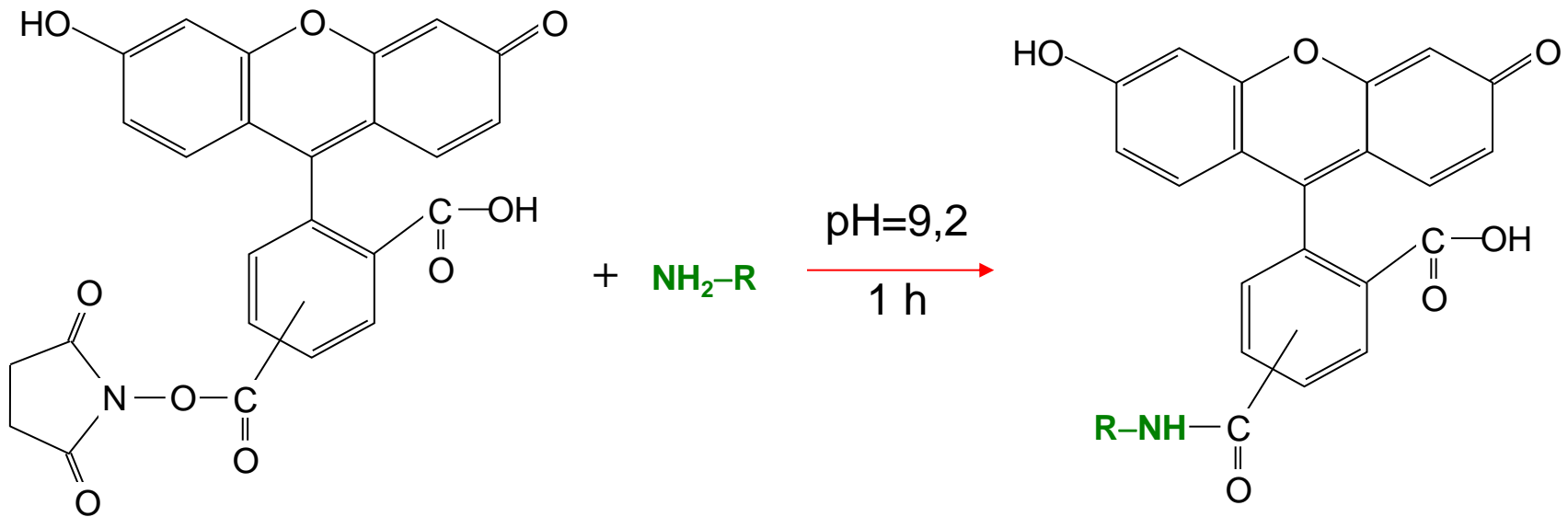
Bánóczy Z. et al. *Archivoc* **143** (2008)
Miklán Zs. et al. *Biopolymers* **92**:489 (2009)
Szabo R. et al. *BBA* **1798**: 2209 (2010)

The effect of the identity and position of amino acid X on cellular uptake



Hudecz F. et al. *J. Mol. Recognition* **16**: 288 (2003)
Szabó R. et al. *Bioconjugate Chemistry* **16**: 1442 (2005)
Szabó R. et al. *Bioconjugate Chem.* **19**: 1078 (2008)
Szabó, R., Sebestyén, M. et al. *ACS Combinatorial Science* **19**: 246-254(2017)

Labelling branched polypeptides with 5(6)-carboxyfluorescein (CF)

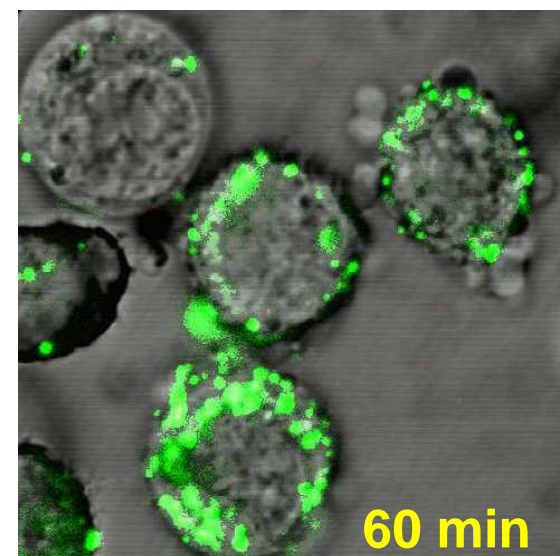
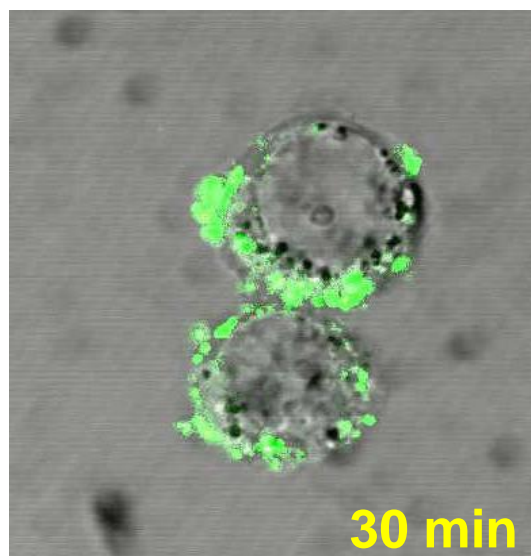
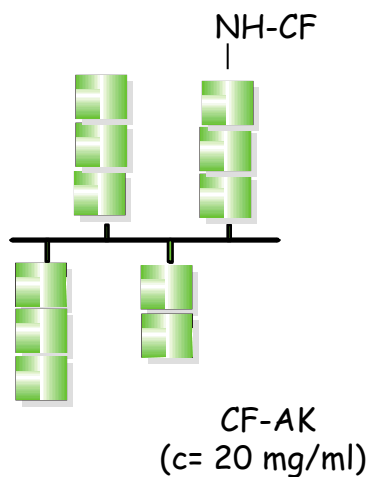
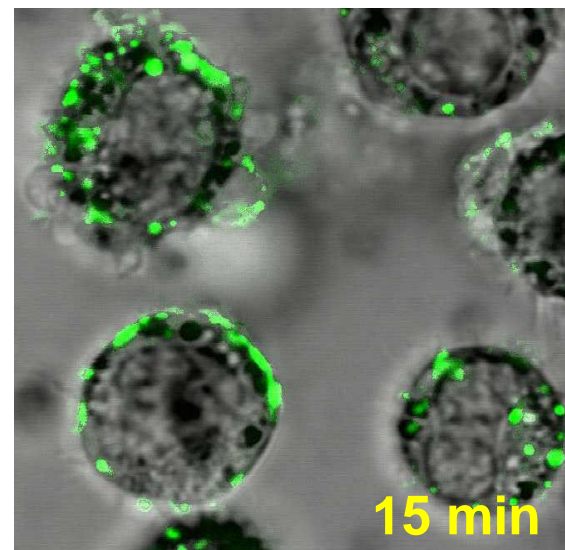
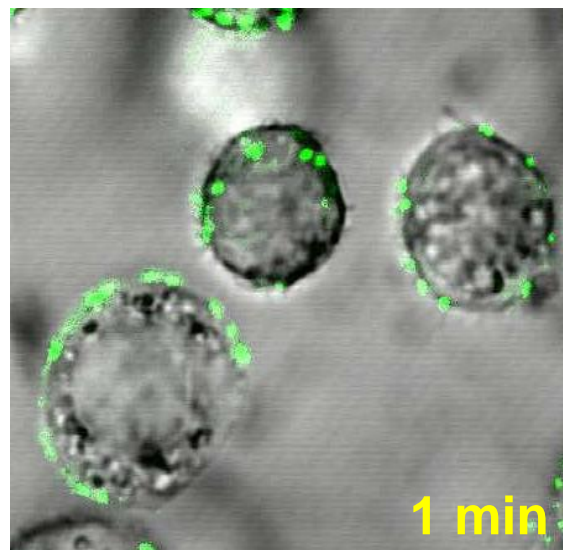
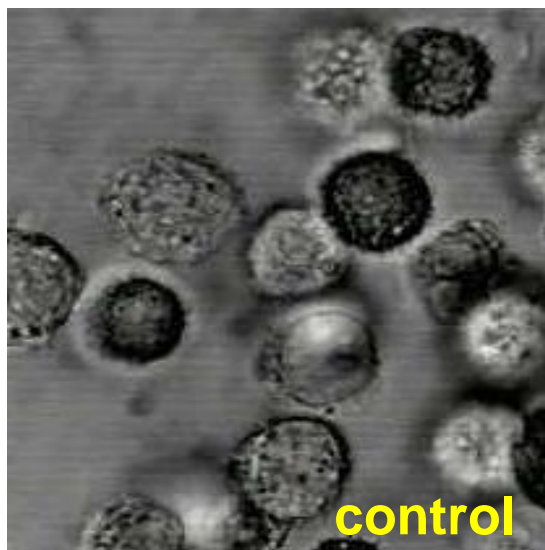


5(6)-carboxyfluorescein-succinimid-ester

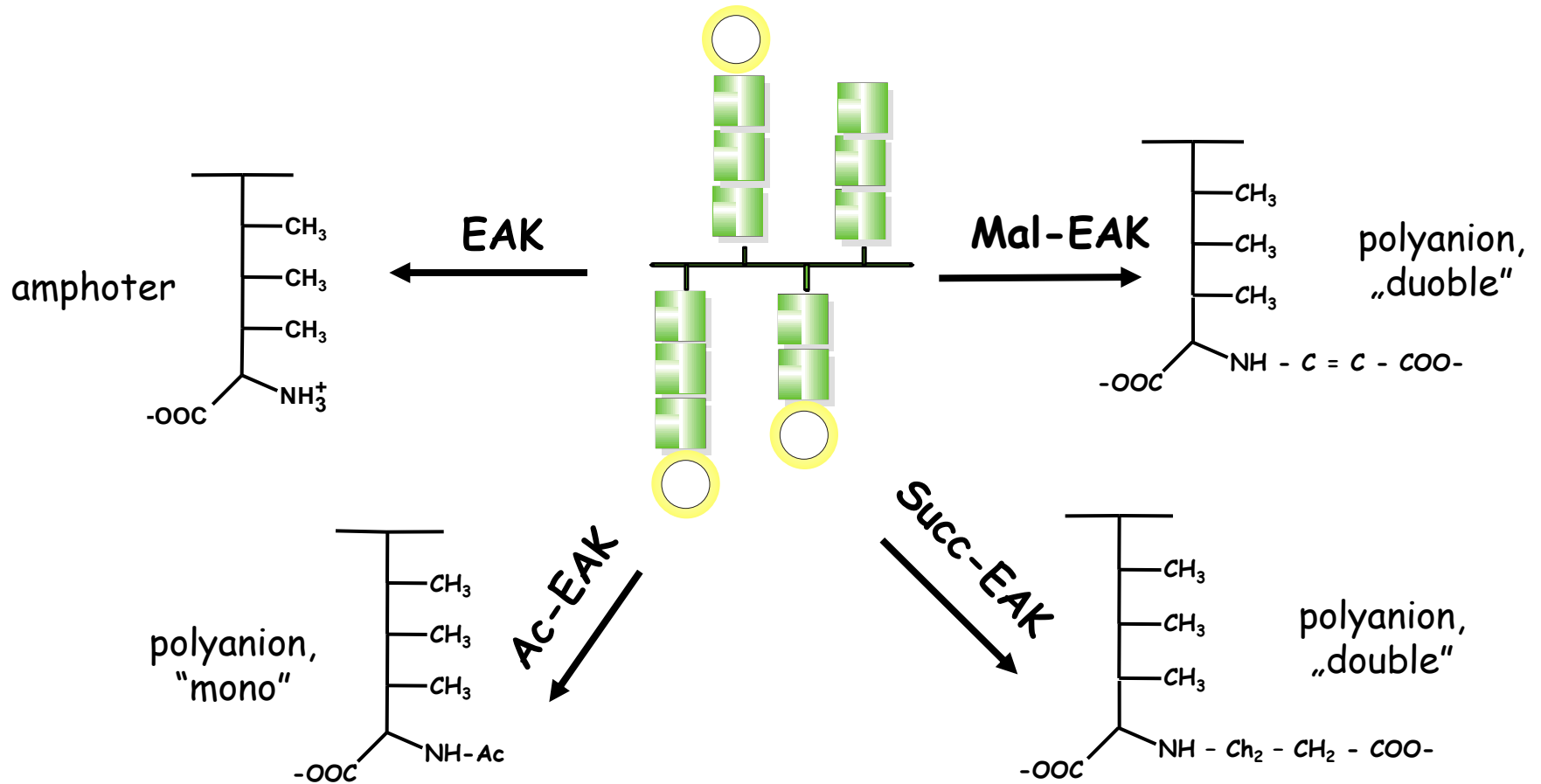
CF-polypeptide

Time dependent cellular uptake of polypeptide AK

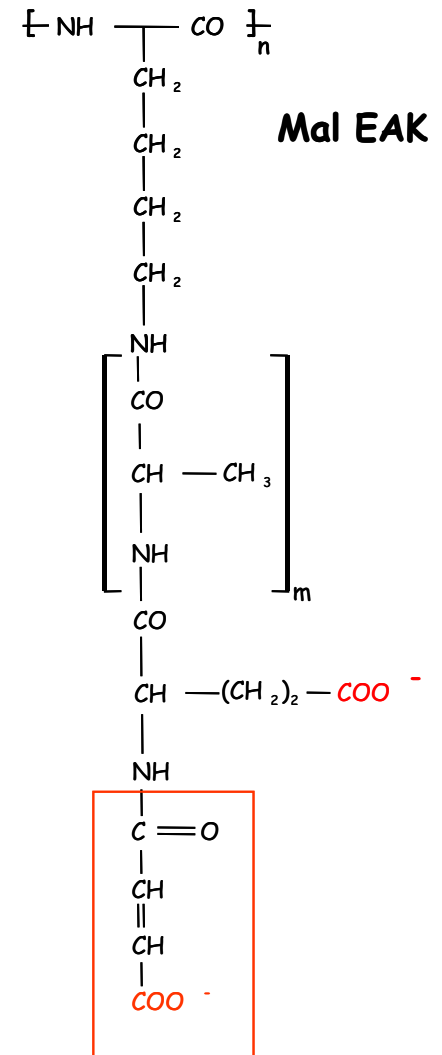
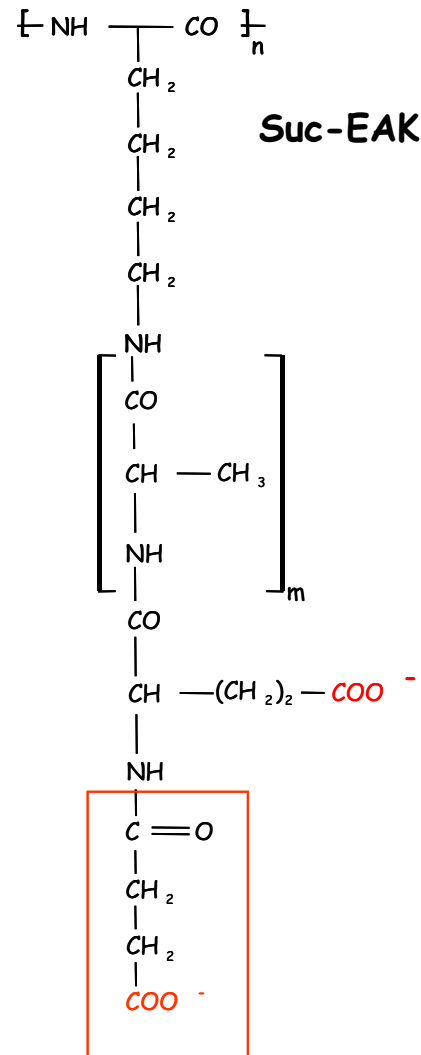
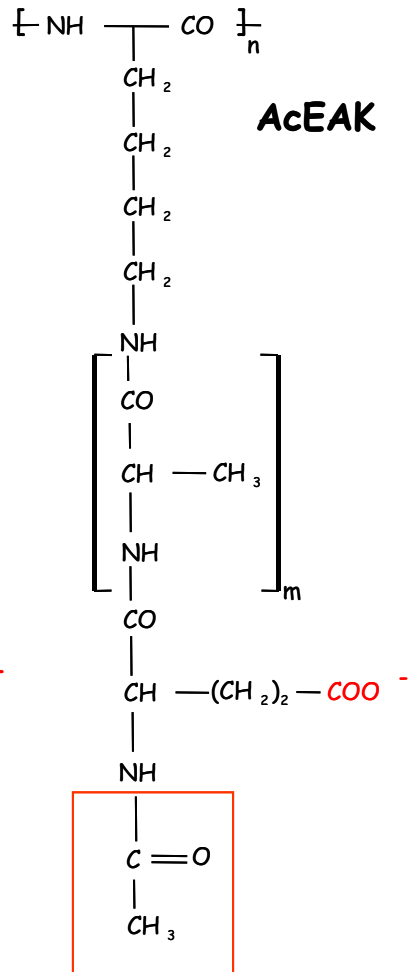
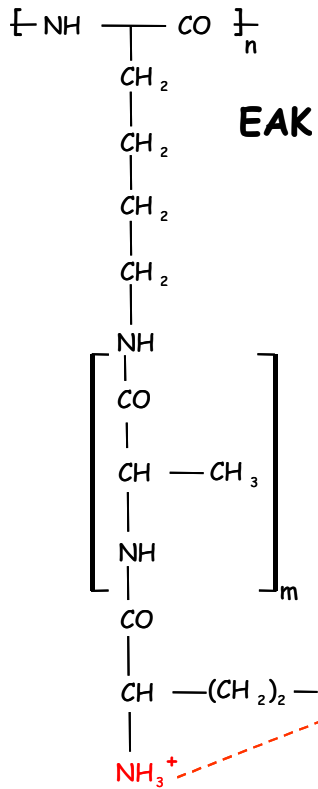
(J774 macrophage cells)



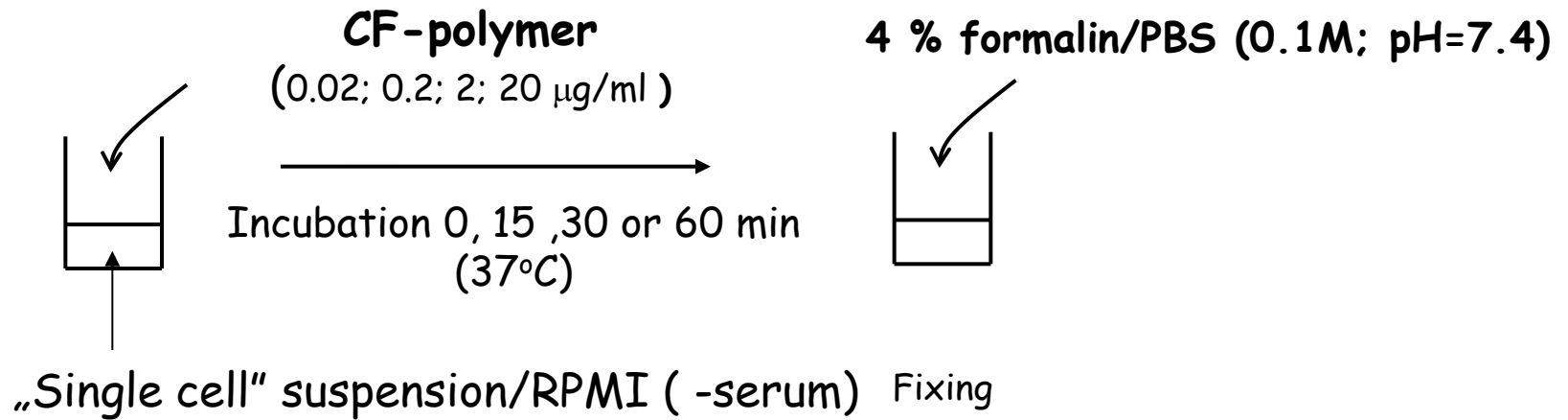
The effect of the identity and position of amino acid X on cellular uptake: **amphotheric and polyanionic** polypeptides



The structure of polypeptides



Protocol of the uptake studies

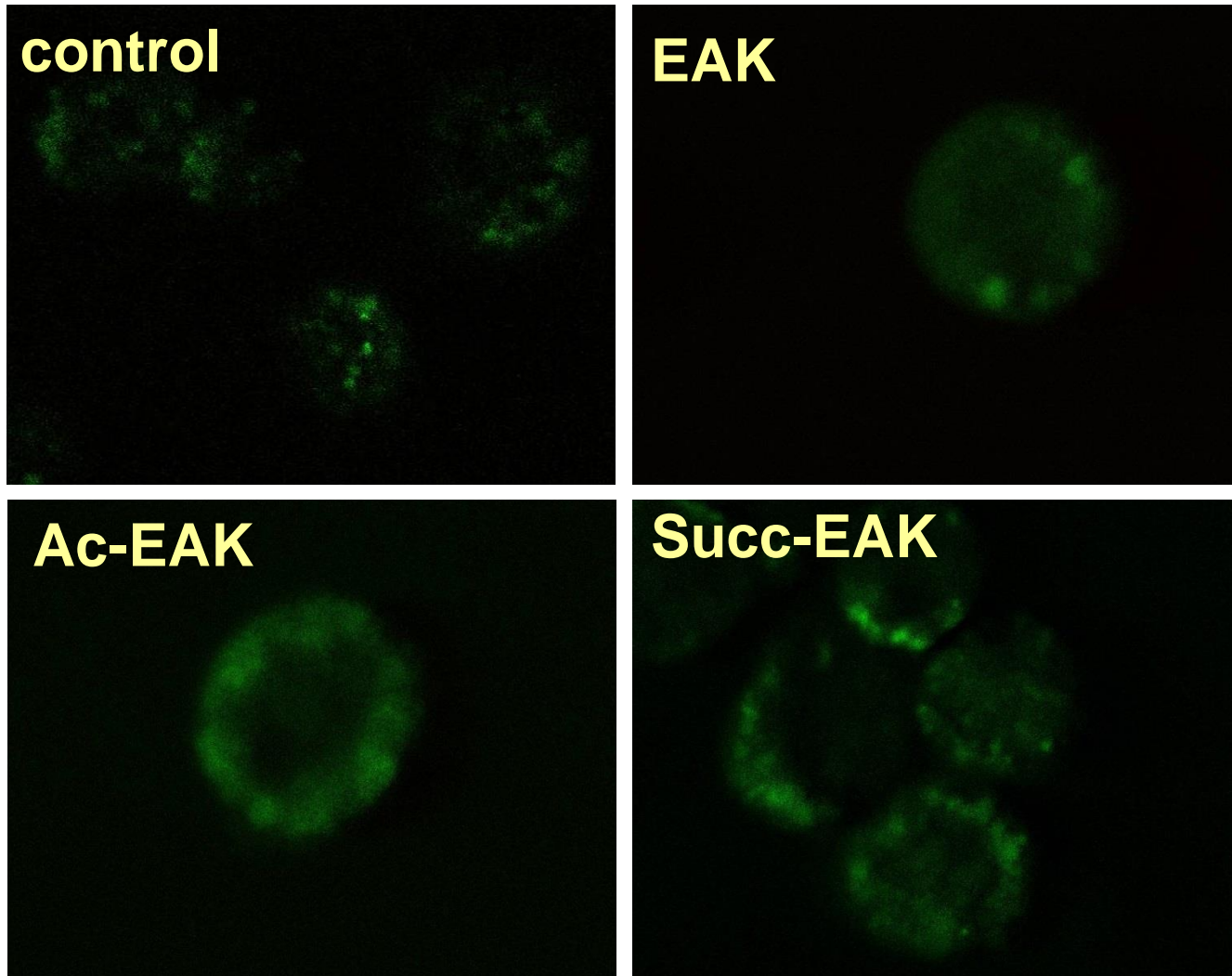


1. 0.5 % tryphan blue, 5 min
2. washing, PBS

Fluorescence intensity (FACS)
(Beckton Dickinson, FACSCalibur, FL1)

Confocal microscopy
(BIO RAD MRC 1024, krypton-argon laser ($\lambda_{ex} = 419 \text{ nm}$, $\lambda_{em} = 519 \text{ nm}$))

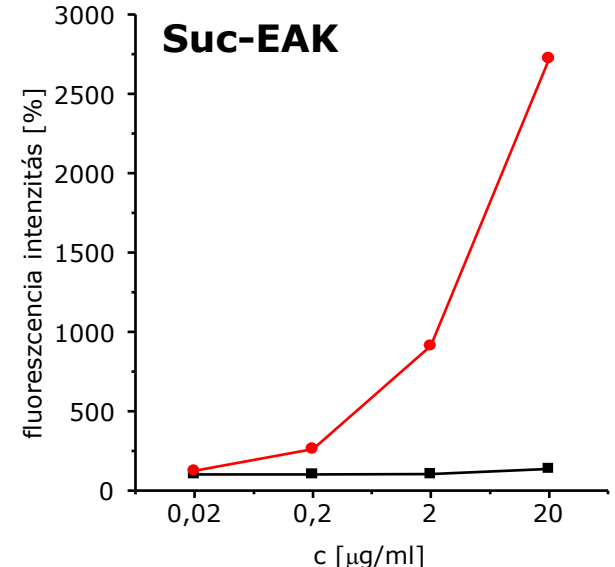
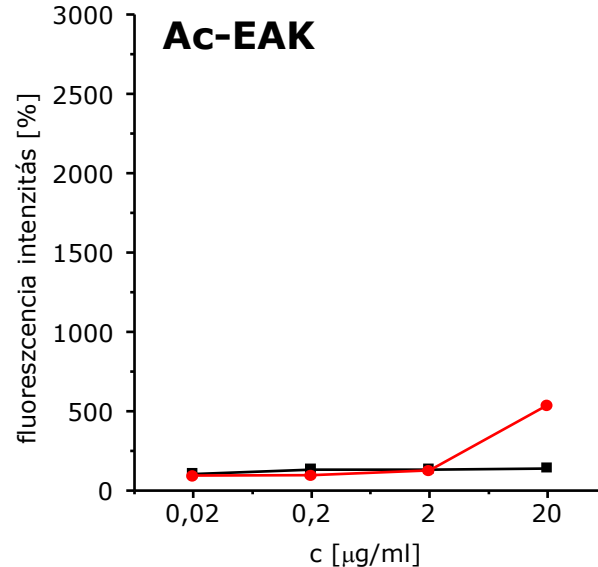
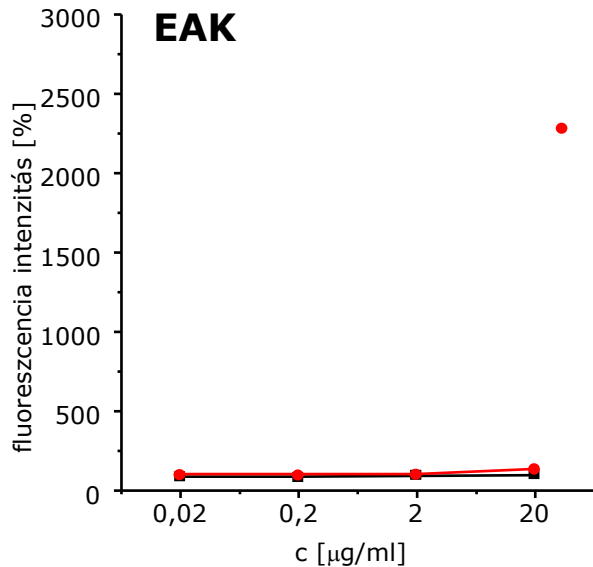
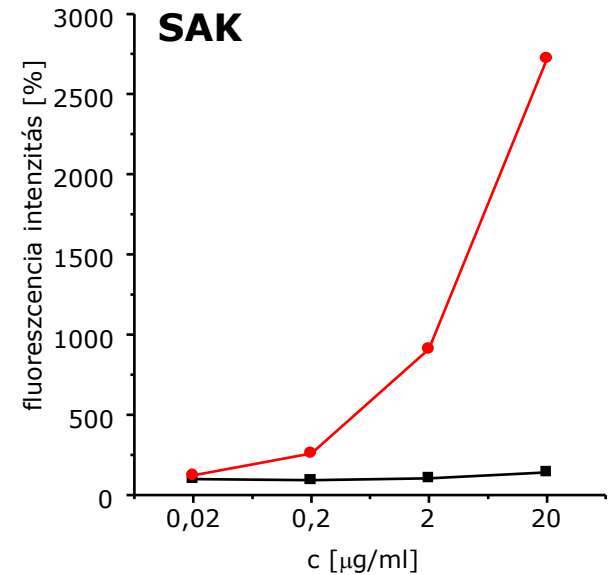
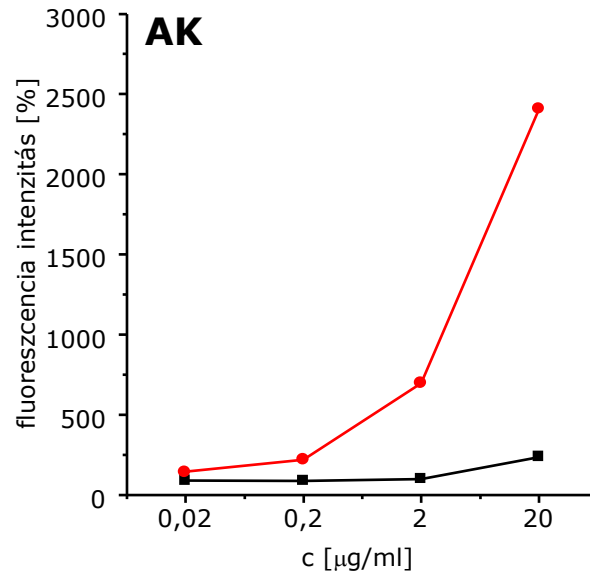
Uptake of amphoteric (EAK) and polyanionic branched polypeptides by J774 cells



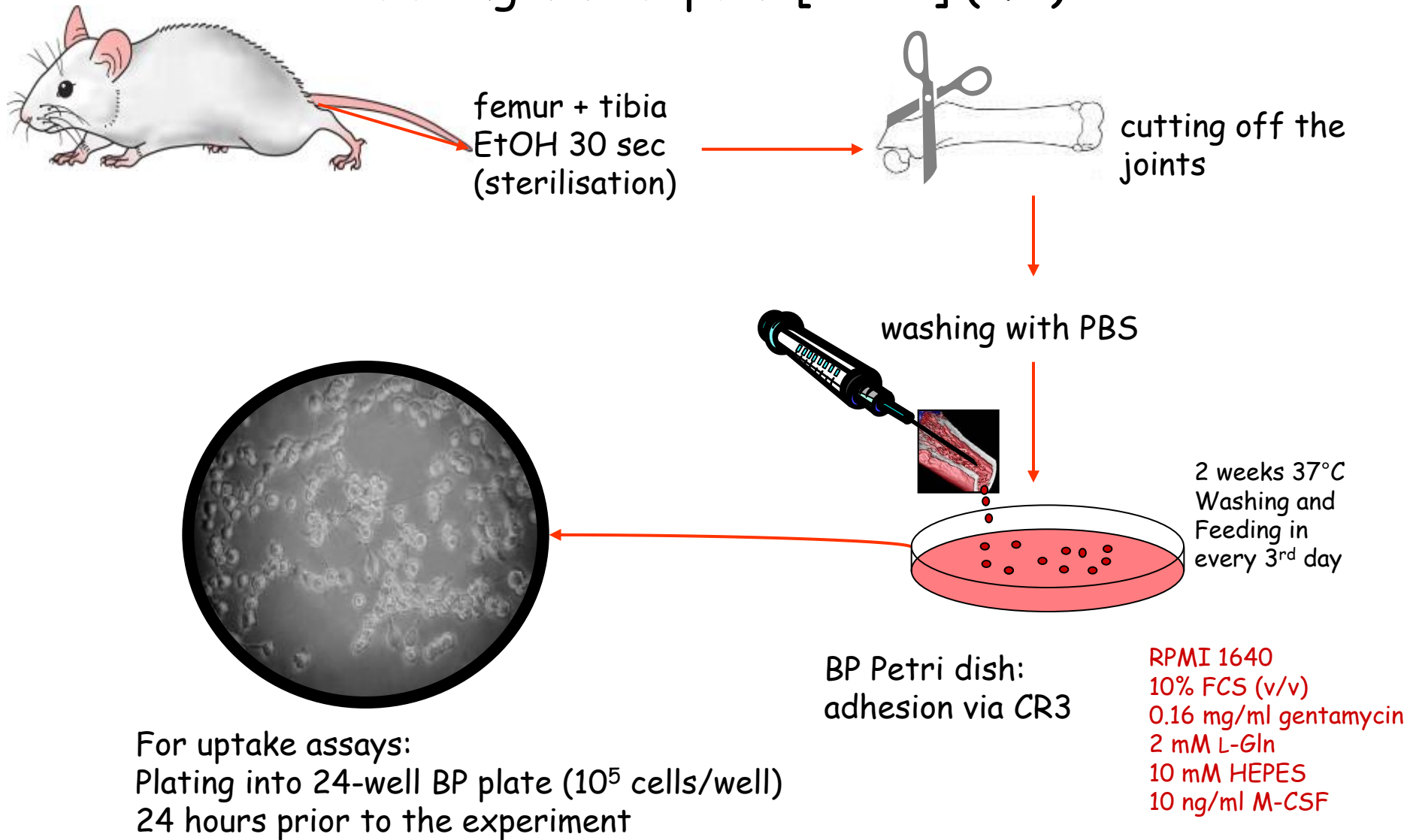
(20 $\mu\text{g/ml}$, 60 min)

Uptake kinetics of polymers by fixed and living J774 cells

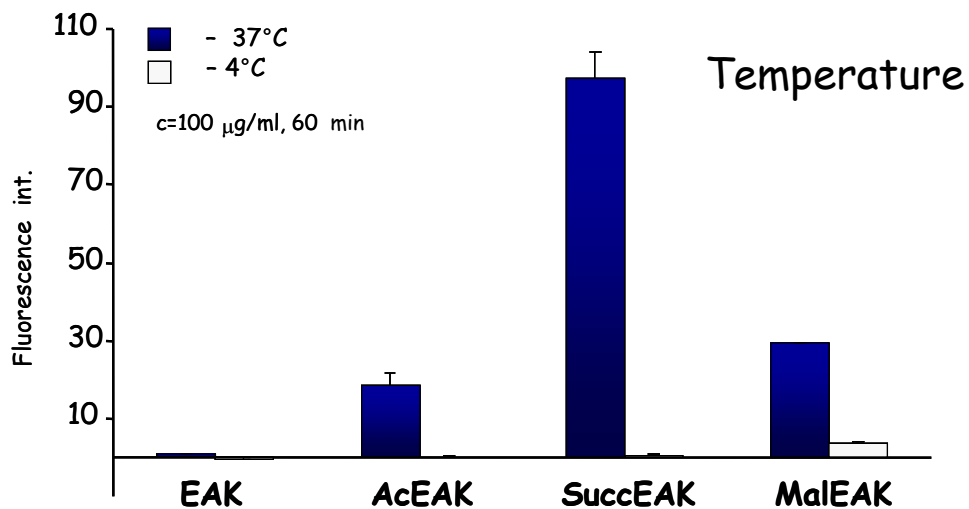
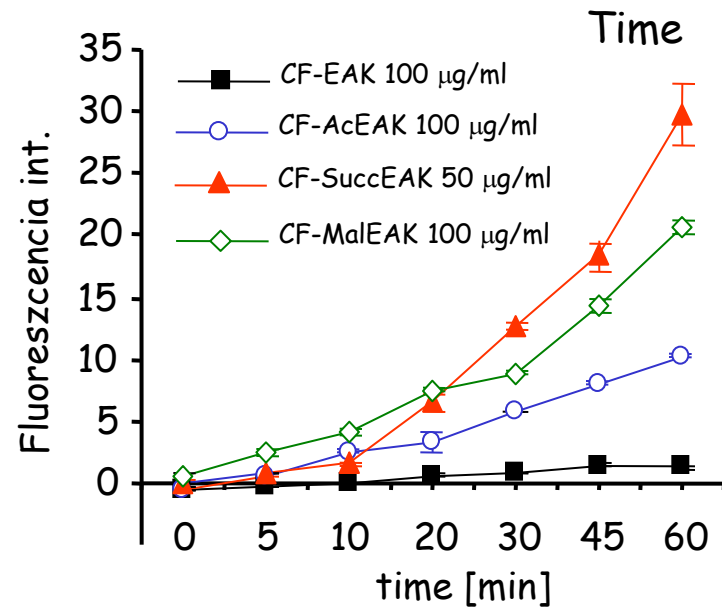
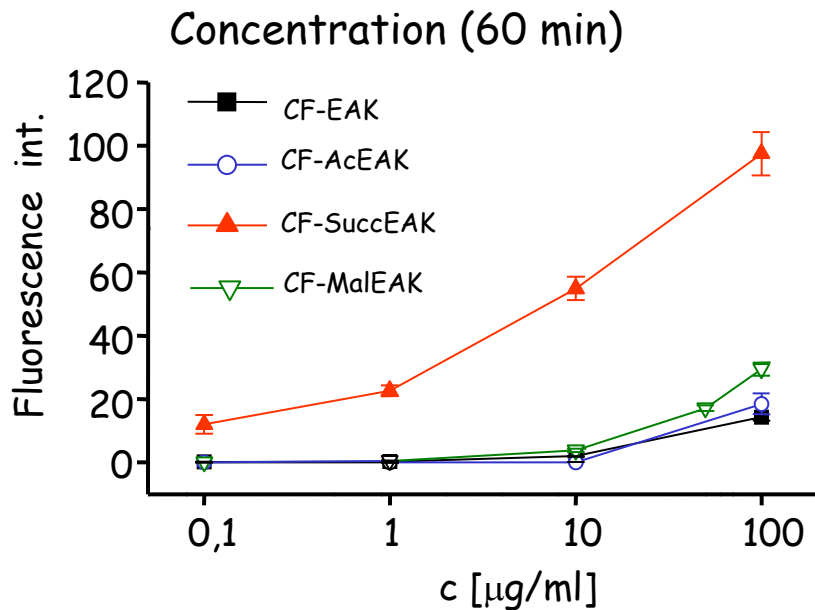
—■— living cells
—●— fixed cells by 4 % formalin



Isolation and differentiation of BMDM scavenger receptor [SR-A] (+/-)

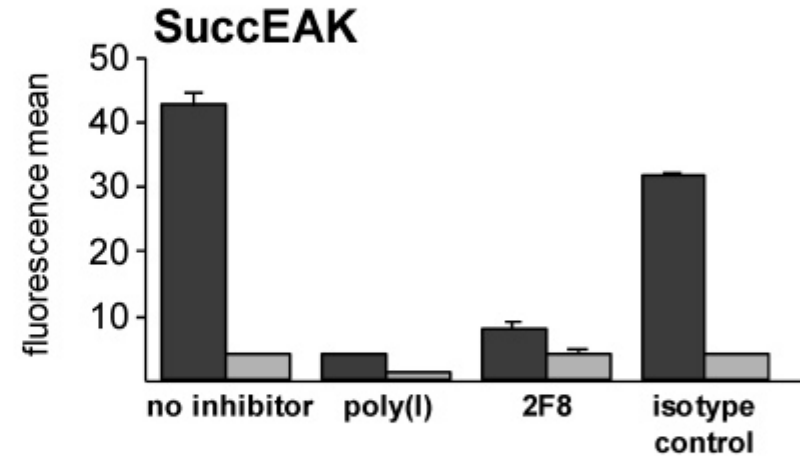
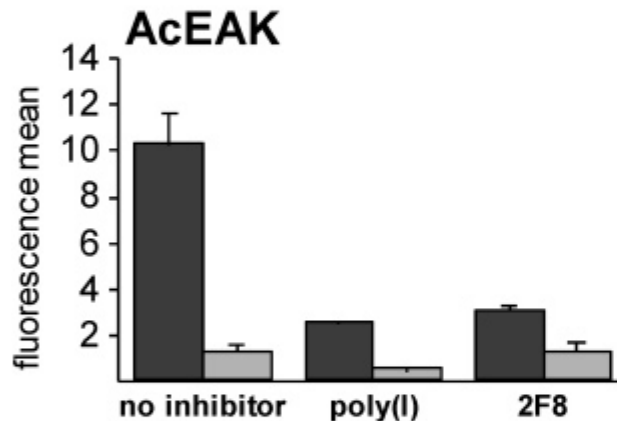
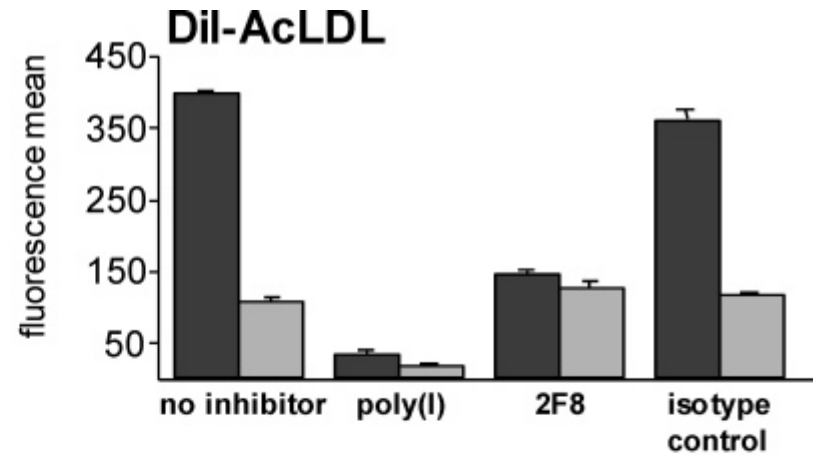
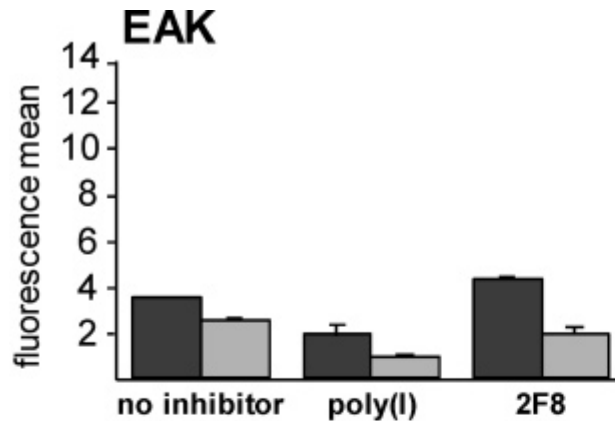
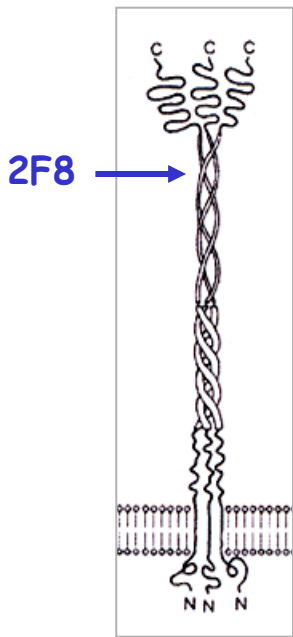


Uptake of CF-polyanionic polypeptides by BMDM macrophages



Inhibition of the SR-A mediated uptake of CF-polypeptides by receptor specific mAb or poly(I) in macrophages

129/ICR
 SR-A -/-



Pre treatment with poly(I) (c = 50 μ g/ml) or MAb 2F8 (c = 15 μ g/ml) for 30 min
 Cells were fixed with 4% formaldehyde

SR-A

Targeting macrophages via scavenger receptor

Ligands

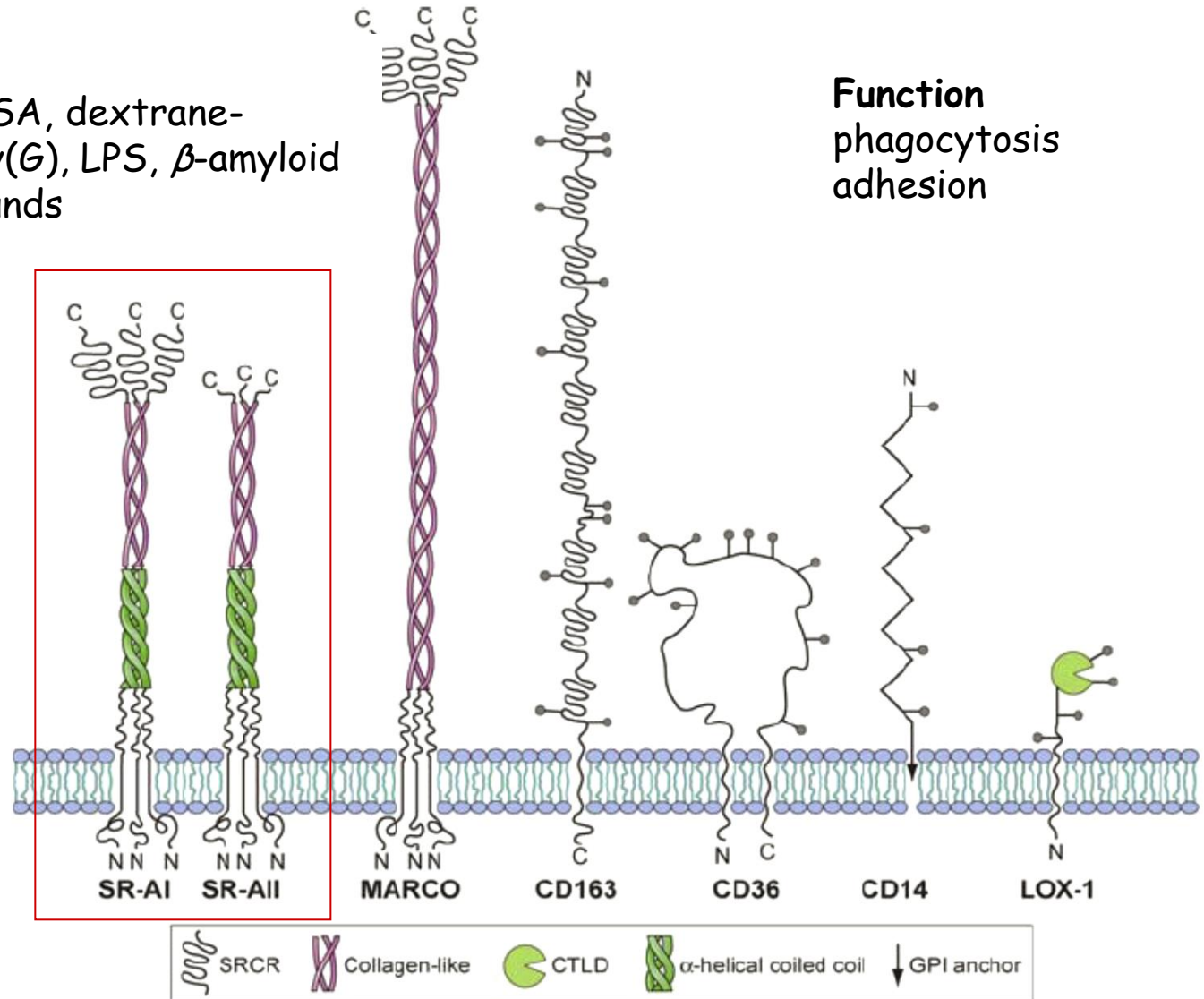
OxLDL, AcLDL, malBSA, dextrane-sulphate, poly(I), poly(G), LPS, β -amyloid
 ⇒ polyanionic compounds

Expression

on macrophages,
 dendritic cells and
 endothelial cells



cell specificity

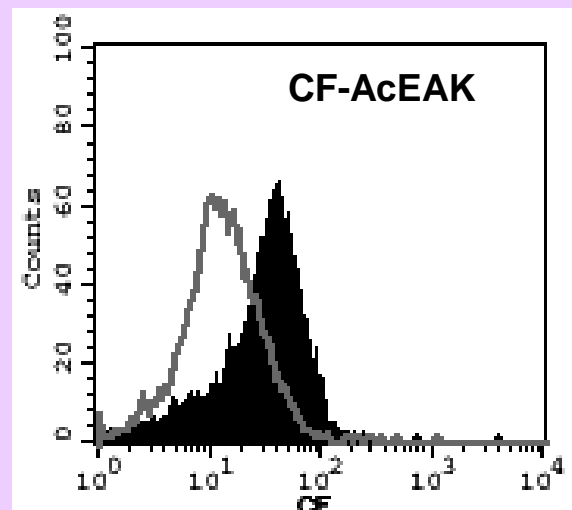
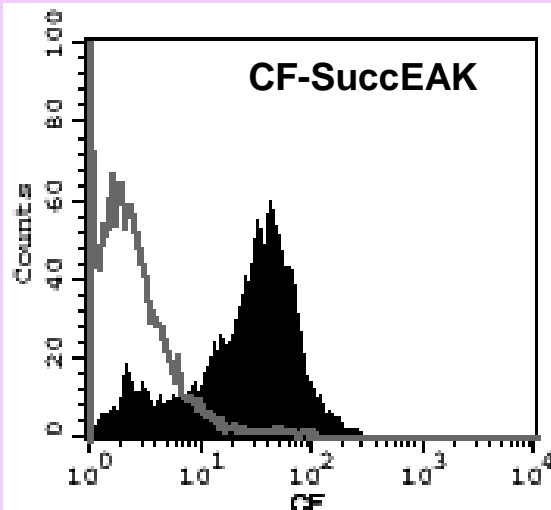
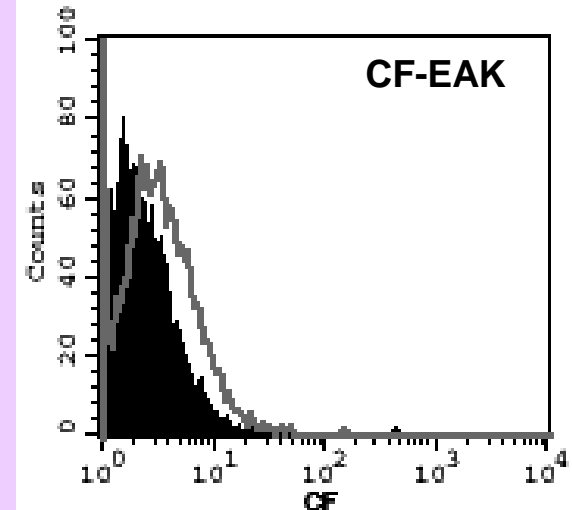
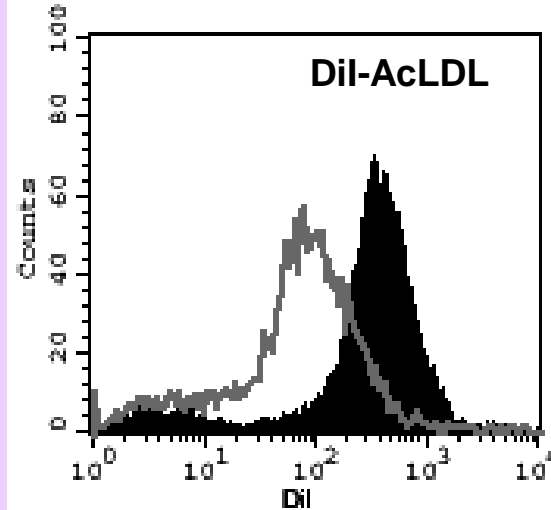
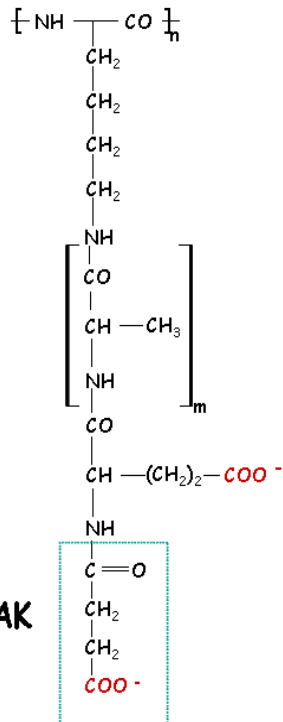


Function

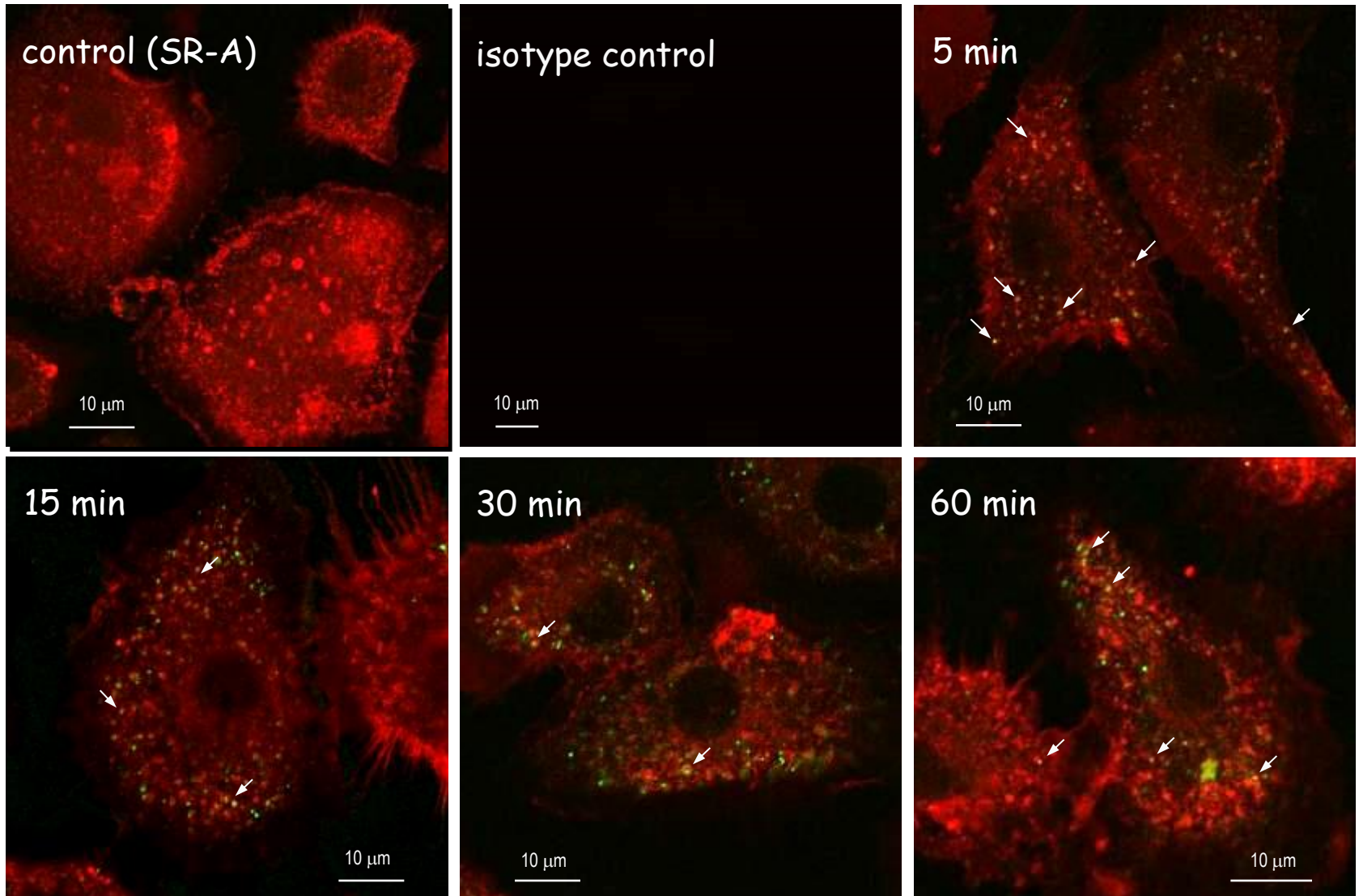
phagocytosis
 adhesion

Uptake of polyanionic polypeptides by macrophages via SR-A

wt (129/ICR)
 SR-A ^{-/-}

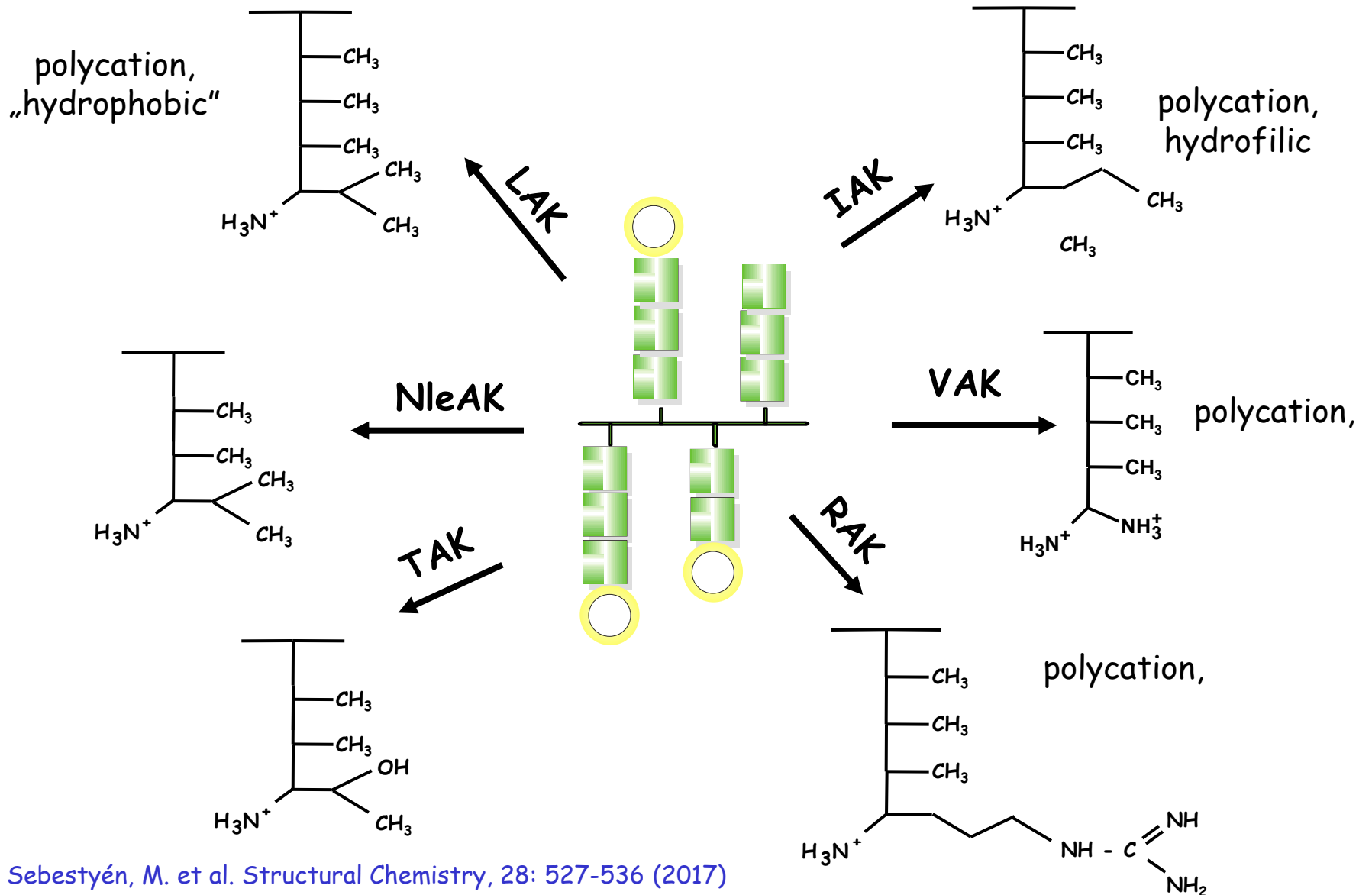


Uptake of polyanionic polypeptides by bone marrow derived macrophages via SR-A

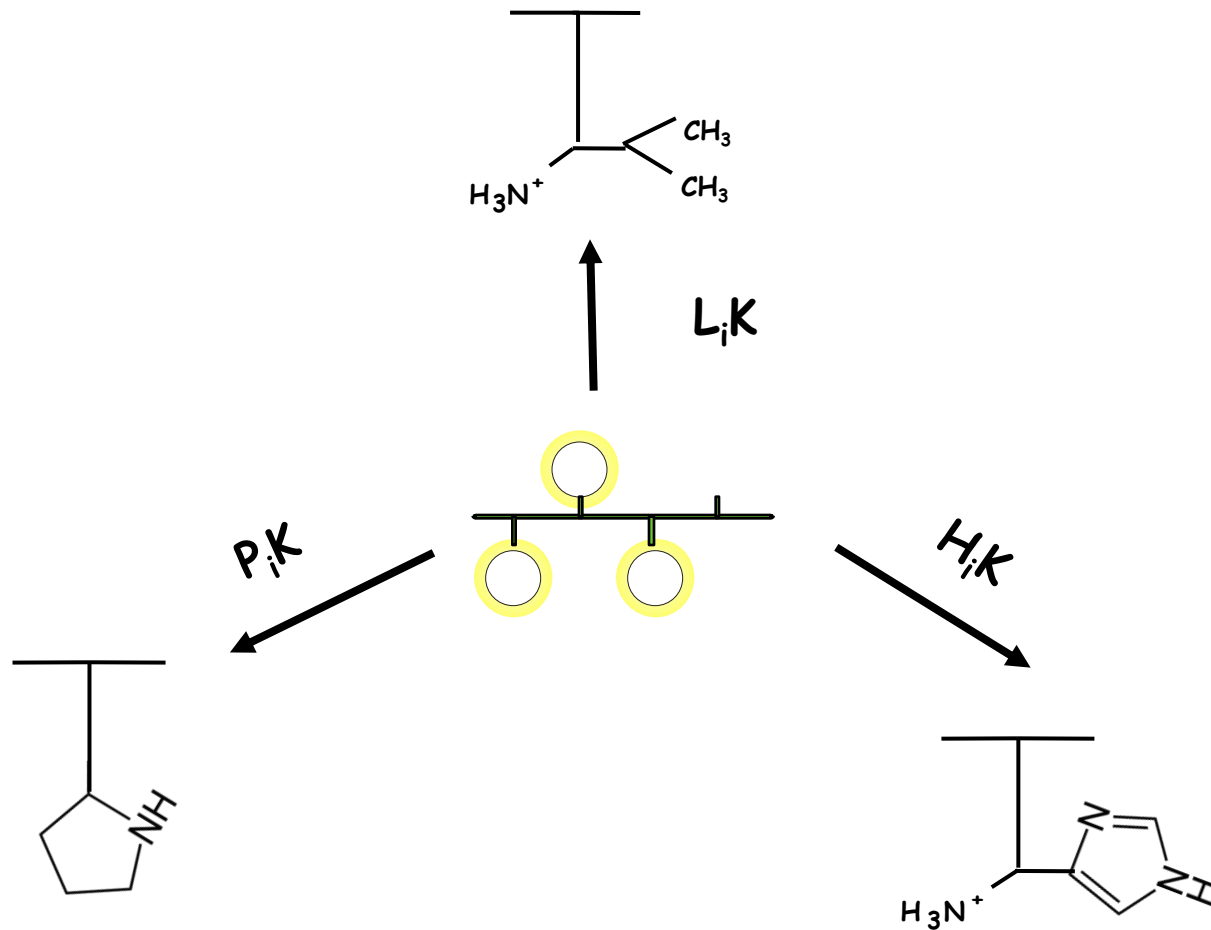


Colocalisation of **SR-A (2F8)** with **CF-SuccEAK**

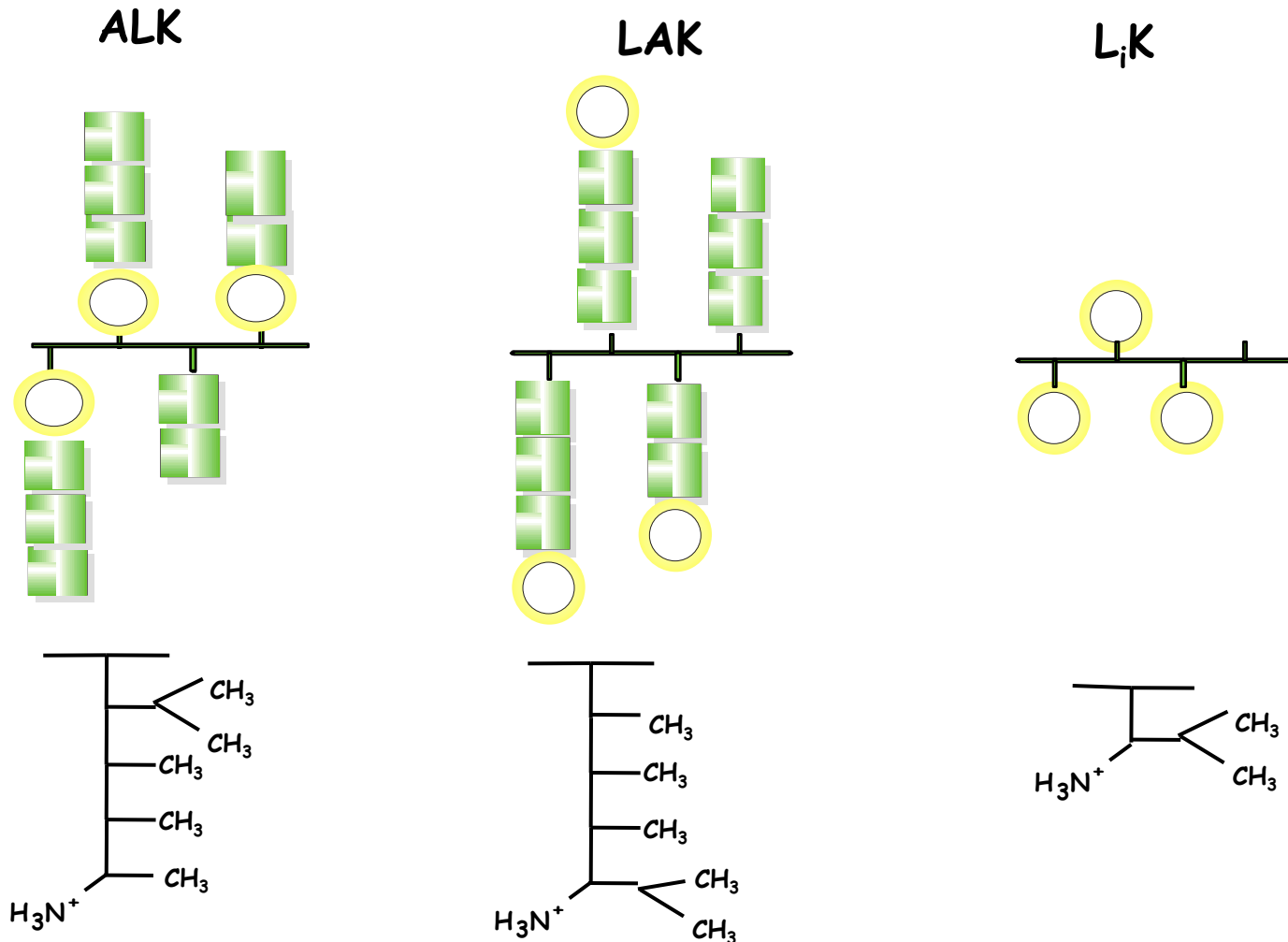
The effect of the **identity** of amino acid X on cellular uptake of **polycationic** XAK polypeptides



The effect of the **identity** of amino acid X on cellular uptake: polycationic X_iK polypeptides



The effect of the **position** of amino acid X on cellular uptake: polycationic polypeptides

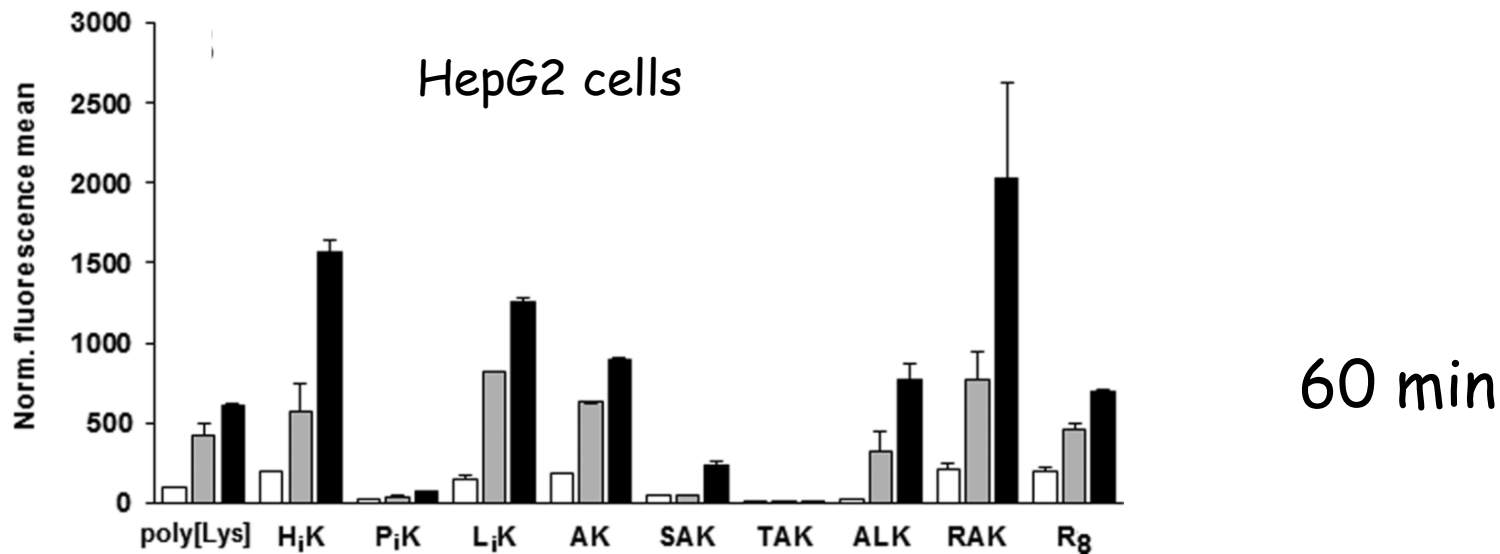
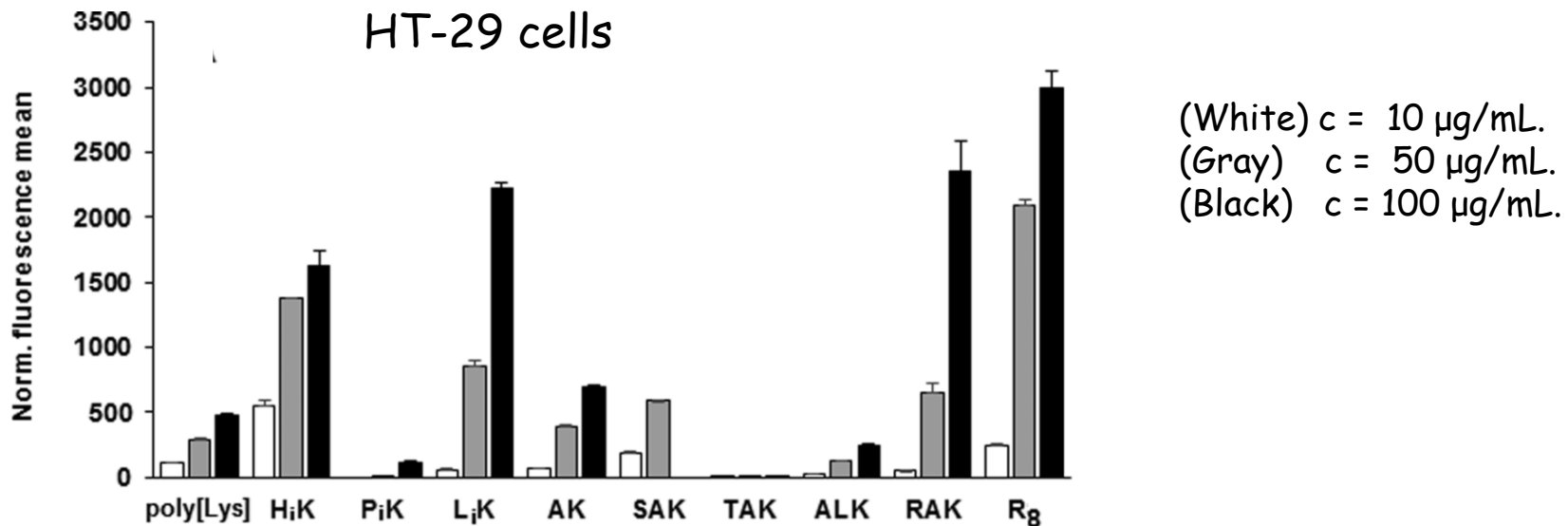


The effect of the **identity** of amino acid X and of cell type on cytotoxicity: polycationic polypeptides

LC₅₀ [μg/ml]

polypeptide	HT-29	HepG2
poly[Lys]	44.1±4.6	32.3±10.8
AK	>100	>100
SAK	>100	>100
TAK	>100	>100
H _i K	99,9	>100
P _i K	>100	>100
L _i K	>100	>100

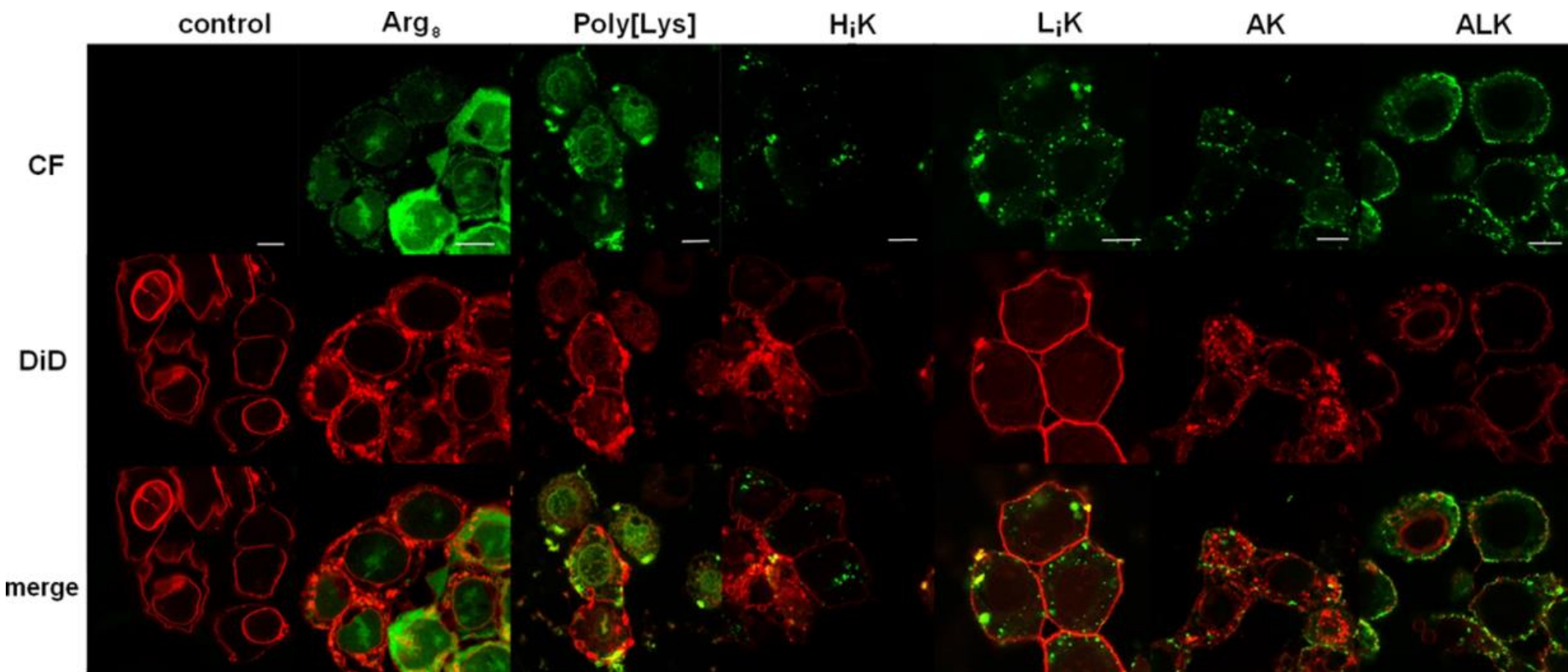
Concentration dependent uptake of the CF-polypeptides



Data were normalized to carboxyfluorescein content of each Cf-(poly)peptide.

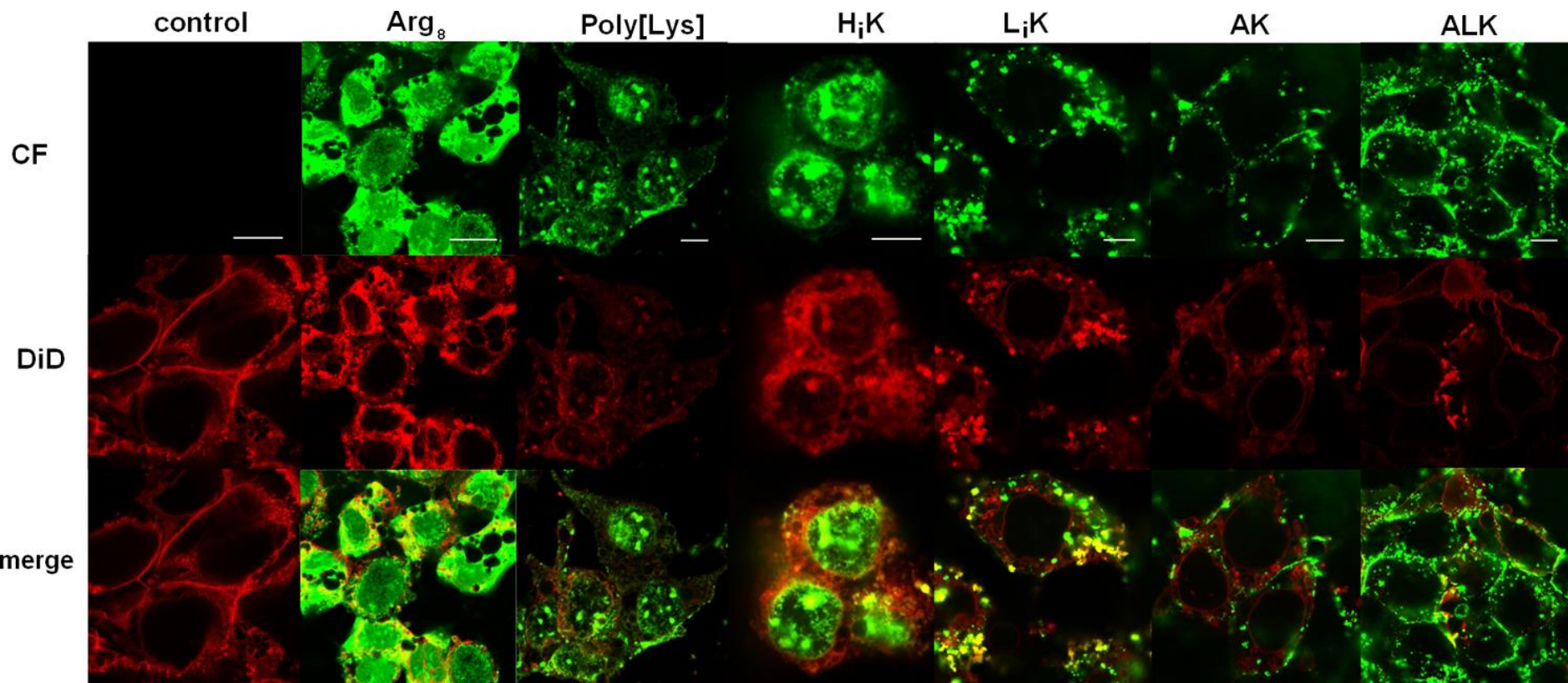
Intracellular localization of CF-polypeptides and Arg₈ in HT-29 human colon carcinoma cells

c = 100 µg/mL after 60 min



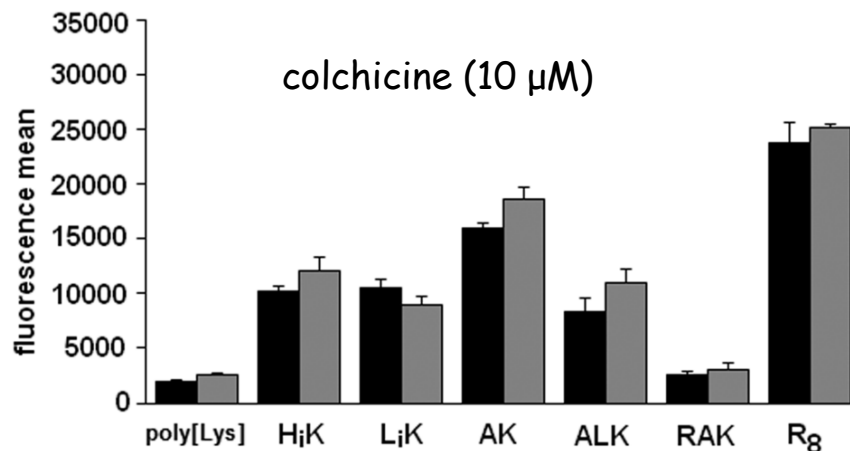
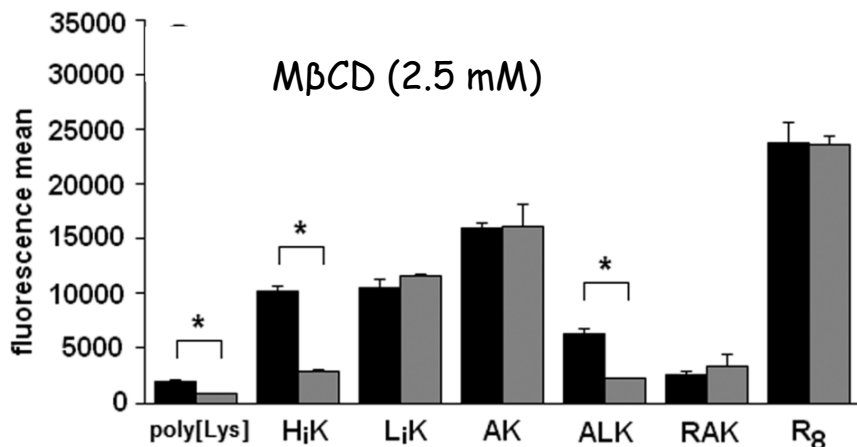
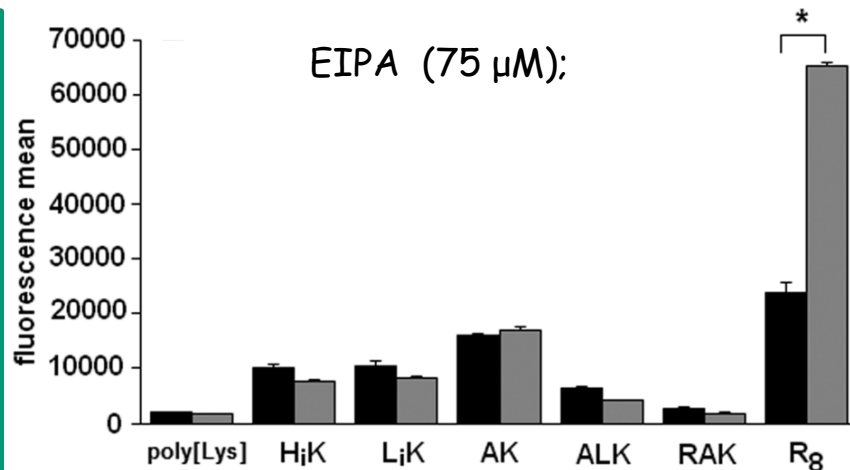
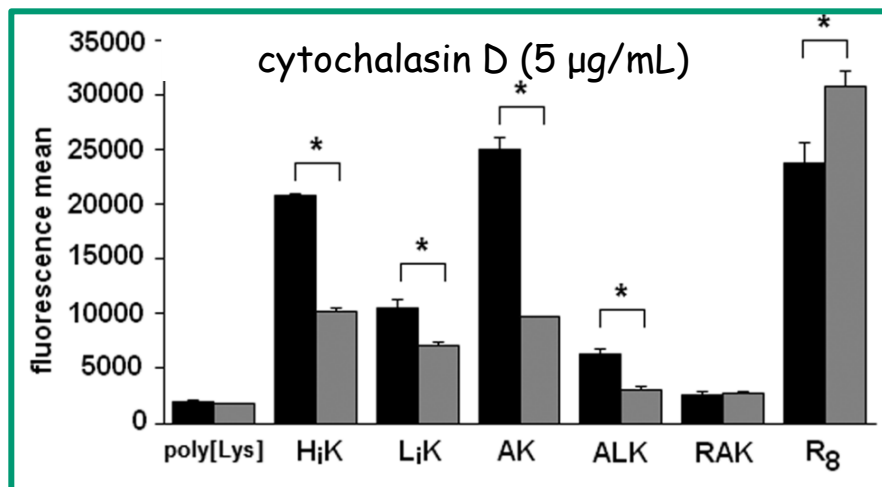
Cells were fixed with 4% paraformaldehyde in PBS (pH 7.4).
Membrane structures were labeled with DiD perchlorate (red). Each bar represents 10 µm.

Intracellular localization of CF-polypeptides and Arg₈ in HepG2 human hepatocarcinoma cells



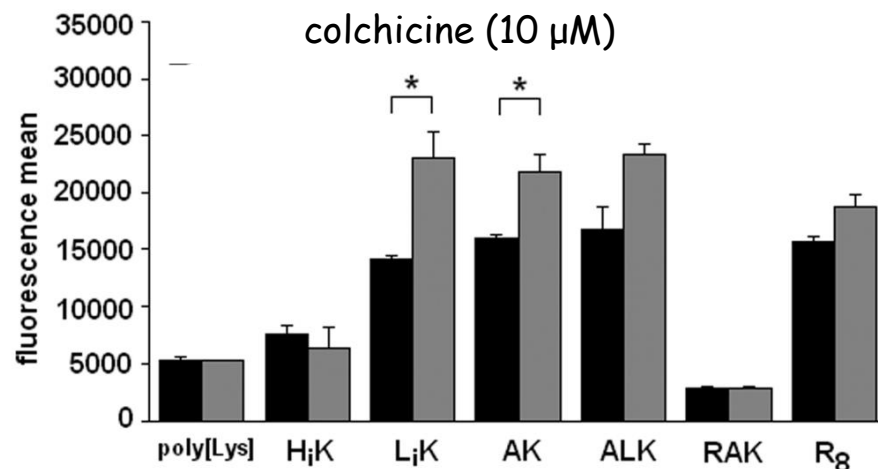
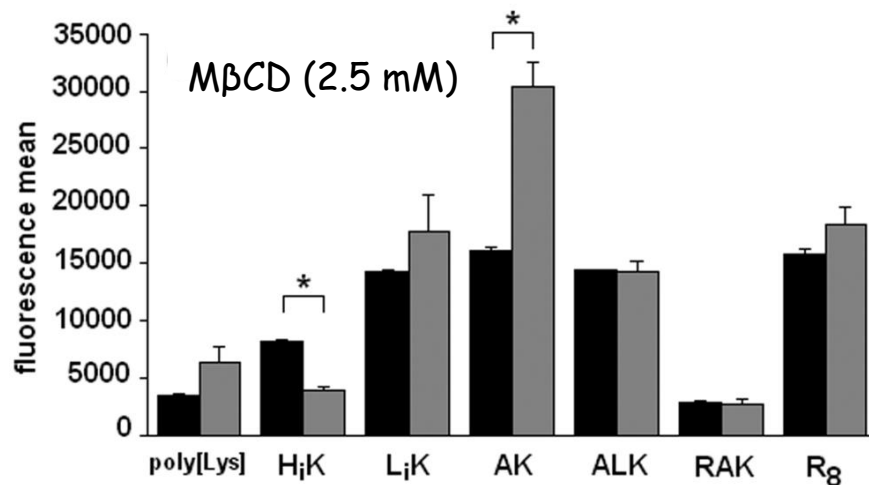
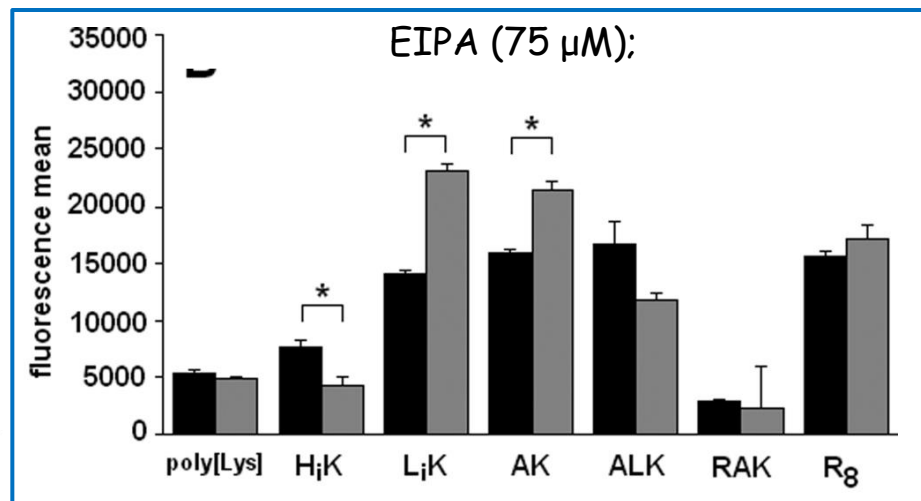
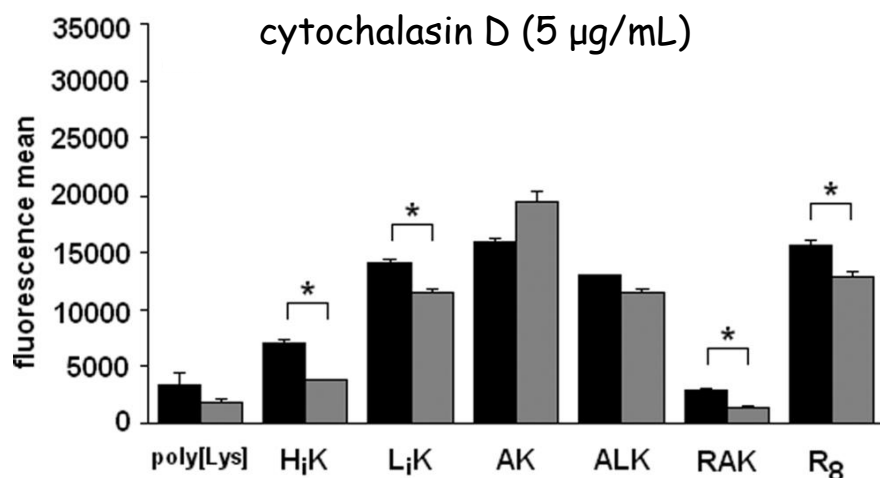
Cells were fixed with 4% paraformaldehyde in PBS (pH 7.4).
Membrane structures were labeled with DiD perchlorate (red). Each bar represents 10 μm .

Uptake of the CF labeled polypeptides and Arg₈ by HT-29 cells in the presence of endocytosis inhibitors



Black: CF-polypeptides (c = 100 µg/mL; CF-poly[Lys], c = 50 µg/mL) and Cf-Arg₈ (R₈, c = 50 µg/mL) without inhibitor. Average of mean fluorescence values of a representative experiment of two independent assays ± SD after subtracting the control. Statistical analysis by Student's t test; *p < 0.05

Uptake of the CF labeled polypeptides and Arg₈ by HepG2 cells in the presence of endocytosis inhibitors



Black: CF-polypeptides (c = 100 µg/mL; CF-poly[Lys], c = 50 µg/mL) and CF-Arg₈ (R₈, c = 50 µg/mL) without inhibitor.

Average of mean fluorescence values of a representative experiment of two independent assays ± SD after subtracting the control.

Statistical analysis by Student's t test; *p < 0.05

Mechanism of uptake of the CF labeled polypeptides in the presence of endocytosis inhibitors

Summary

Compound	Colchicin ⁴		EIPA ^{1,2,3}		Cytochalasin D ^{1,2}		Metil-β-ciklodextrin	
	HT-29	HepG2	HT-29	HepG2	HT-29	HepG2	HT-29	HepG2
Poly[Lys]	-	-	-	-	-	-	+	-
H _i K	-	-	-	+	+	+	+	+
AK	-	-	-	-	+	-	-	-
ALK	-	-	-	-	+	-	+	-

HepG2 Pinocytosis (poly[Lys], H_iK, AK, ALK)
 Macropinocytosis (H_iK)
 Lipid raft/caveola (H_iK)

HT-29 Pinocytosis (poly[Lys], H_iK, AK, ALK)
 Macropinocytosis (poly[Lys], H_iK, ALK)
 Lipid raft/caveola (poly[Lys], H_iK, ALK)

¹Nakase, I. et al, Mol. Ther. (2004) 10: 1011-1022

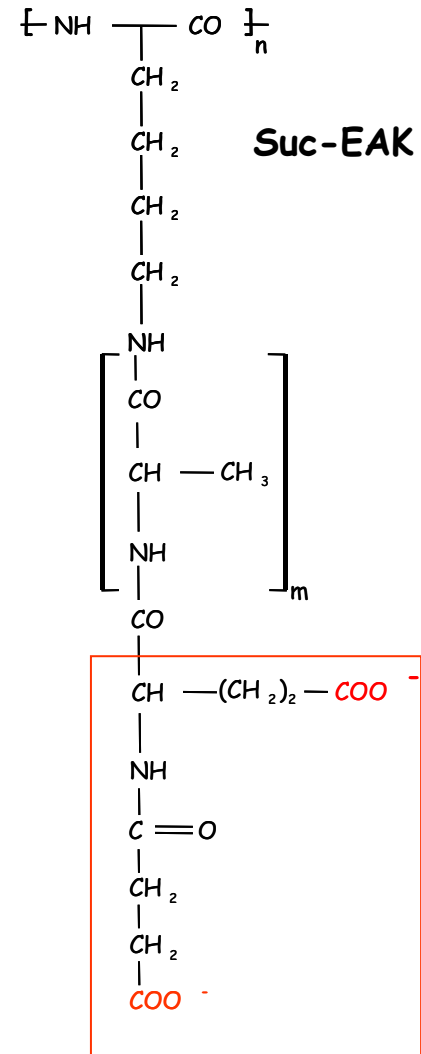
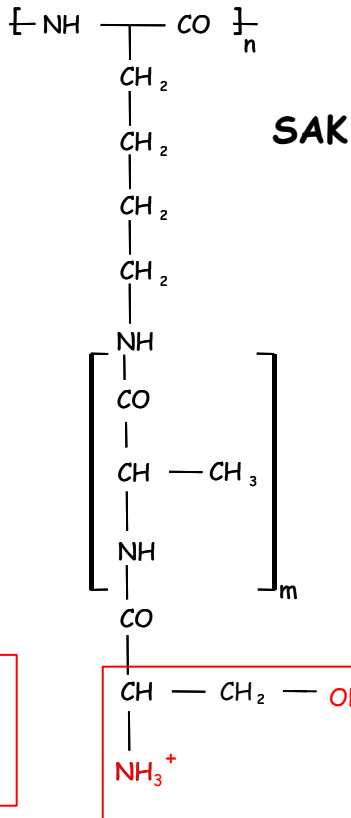
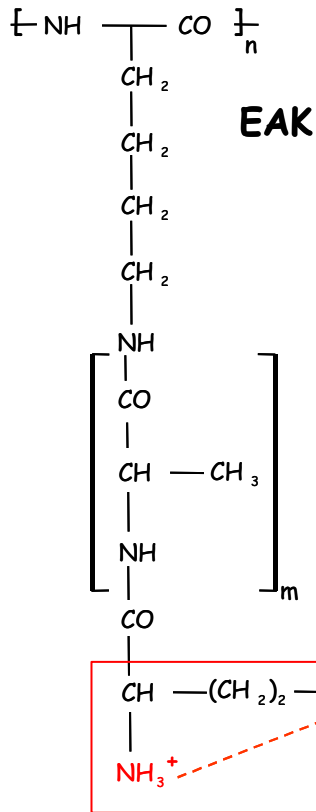
²Delwig, A. et al, Arthr. Res Ther. (2006) 312: 1345-1360

³Heikkilä, O. et al, J. Virol. (2010) 84: 3666-3681

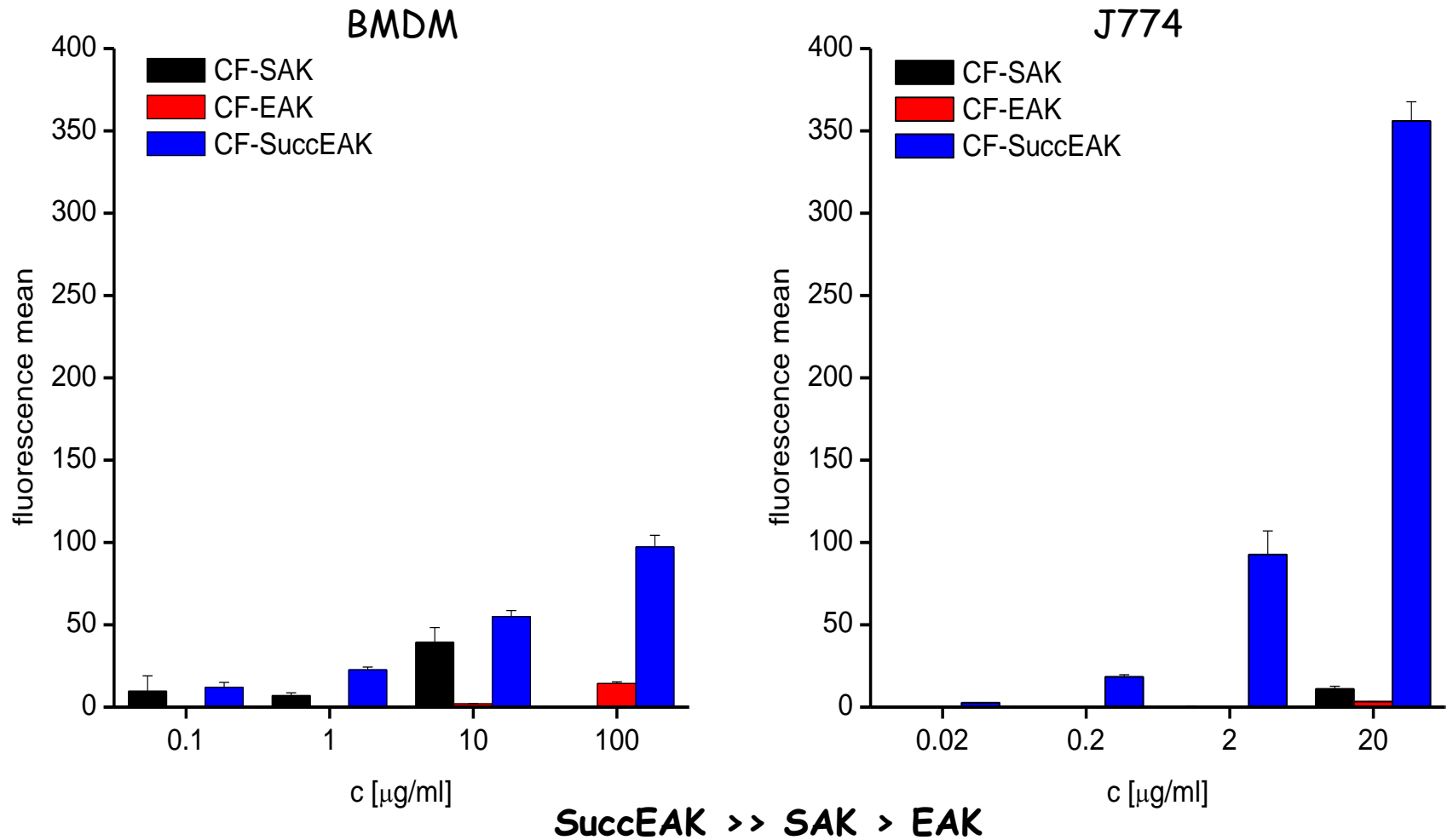
⁴Piasek, A. et al, hematol. Blood Transf. (1985) 29: 511-513

⁵Rodal, S.K. et al, Mol. Biol. Cell (1999) 10: 961-974

The structure of polypeptides



Uptake of polycationic, amphoteric and polyanionic polypeptides



Conclusions

1. Branched polypeptides are taken up by J774 cells. This process is **time and concentration** dependent.
2. The structure (e.g. charge properties) influence greatly the uptake. The following order could be established: **polycationic >> polyanionic >> amphoteric** polypeptides.
3. Polypeptides could be detected in the **cytoplasm**.
4. Higher level of uptake was observed by fixed than living cells.



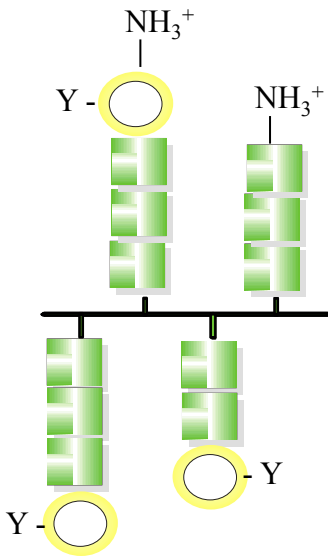


Conclusion

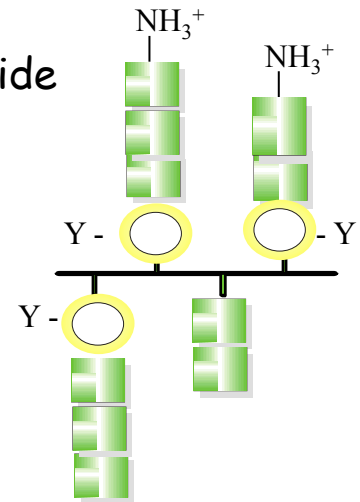


Solution conformation, cytotoxicity and cellular uptake of the XAK/AXK type branched polypeptide are influenced by the

- identity, character (hydrophobic, hydrophilic), charge
- position in the side the side chain
- number of amino acid X.



XAK, poli[Lys(X_i-DL-Ala_m)]



AXK, poli[Lys(DL-Ala_m-X_i)]

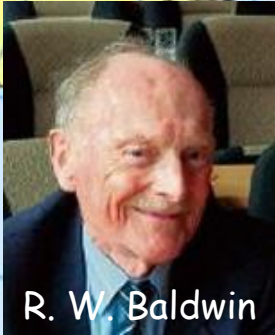
Thanks a lot!



Partners around...



E. Heber-Katz



R. W. Baldwin



Karel Blaha

Philadelphia

S. Gordon

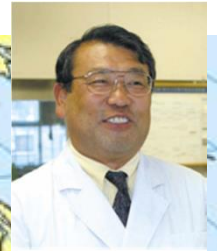
Nottingham

Prága

Budapest



Cooperation - multidisciplinary



Oxford

Konstanz

Saclay

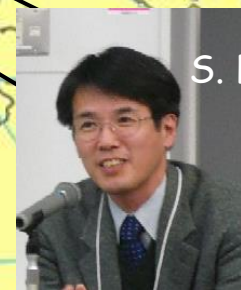
Barcelona

Palermo

Budapest



A.C. Ghose



S. Futaki

Tokyo

Kyoto

Kumamoto



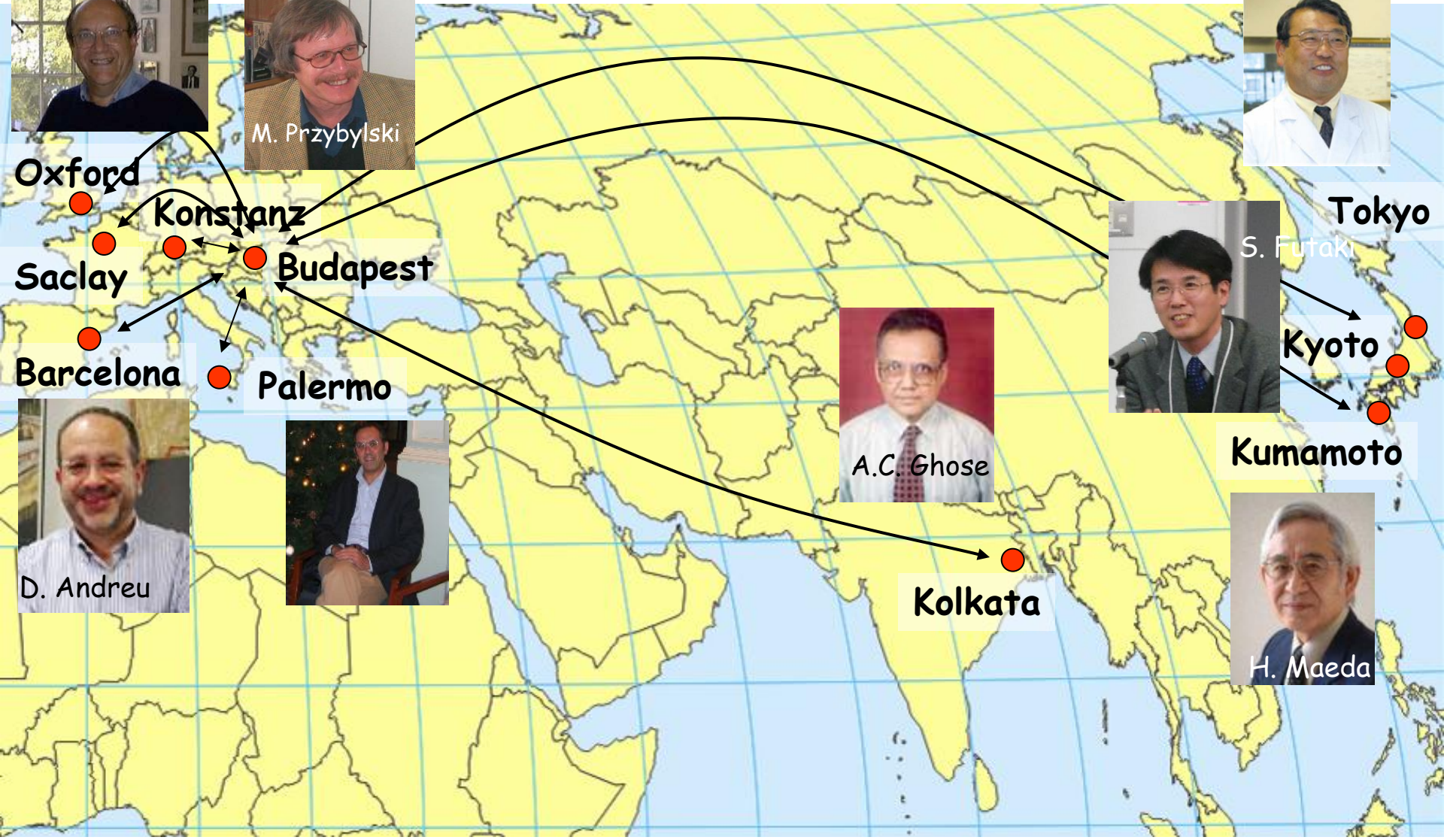
D. Andreu



Kolkata



H. Maeda



Acknowledgements

OTKA



375
E · L · T · E



Me@chem



Agence Nationale de la Recherche
ANR GIP



BÉRES
GYÓGYSZERGYÁR RT.