

**International Summer School for young scientists**

**NANOTECHNOLOGY: from fundamental research to innovations**

**August 26 - September 2, 2012, Bukovel, Ukraine**



# **Study of nanocomposites of amino acids and organic polyethers by means of mass spectrometry and molecular dynamics simulation**

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Nanoparticles of various materials are promising...

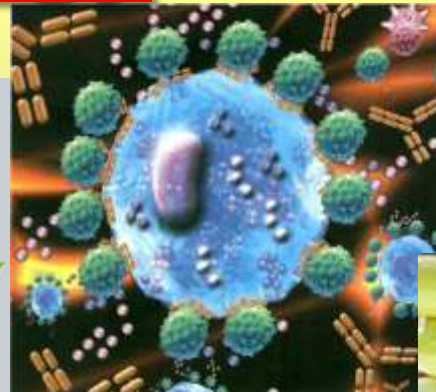
biophysics



cryobiology



nanobiotechnology



**GREEN CHEMISTRY**  
and Cradle to Cradle Product Design



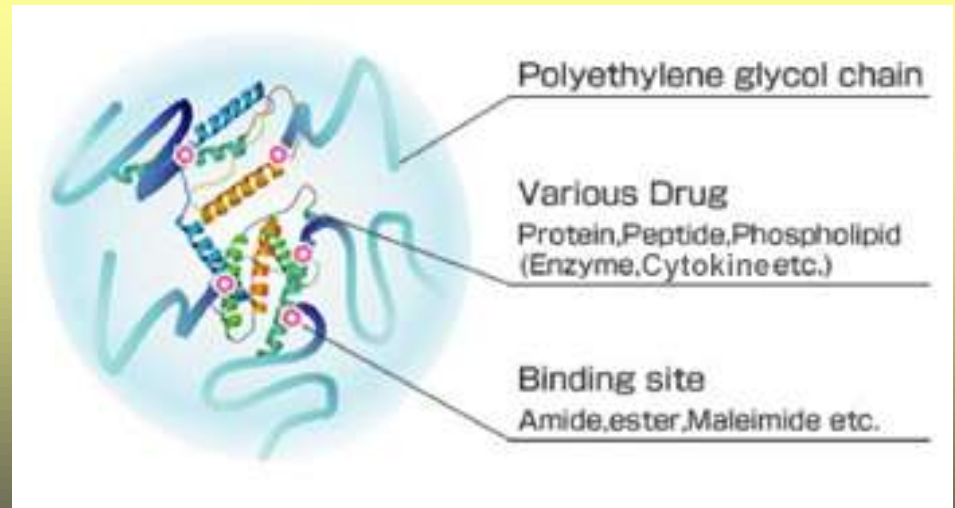
pharmacology



**OBJECTS: NANOPARTICLES** composed of biomolecules and organic polymers

## **Applications**

- **Pegylated proteins**
- **Nanocomposites based on organic polymers**
- **for drug delivery**



# AIM:

To understand the structure and self-assembling of **nanoparticles** determined by interactions of polyether chains with protein amino acids

# TASK:

To study **nanocluster interaction** on simple model of nanoparticles consisting of polyether oligomers and amino acids by means of a combined approach of experiment - **electrospray mass spectrometry** and computer modeling - **molecular dynamics calculations.**

# Object:

**NANOPARTICLES =**

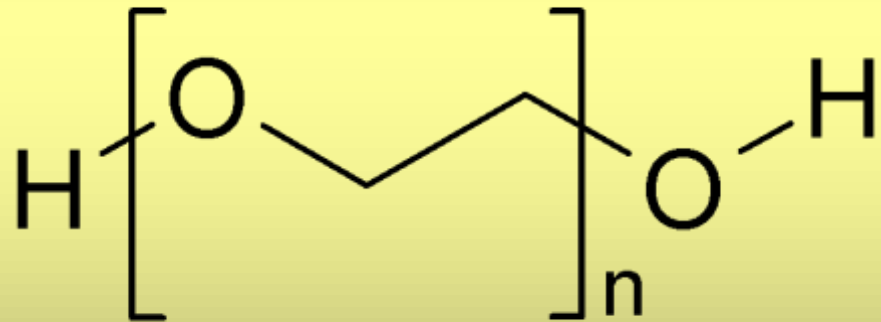
**organic polymers  
polyethers**

**Polyethylene glycol  
and its derivatives**

**+**

**Amino acid**

**PEG-400**

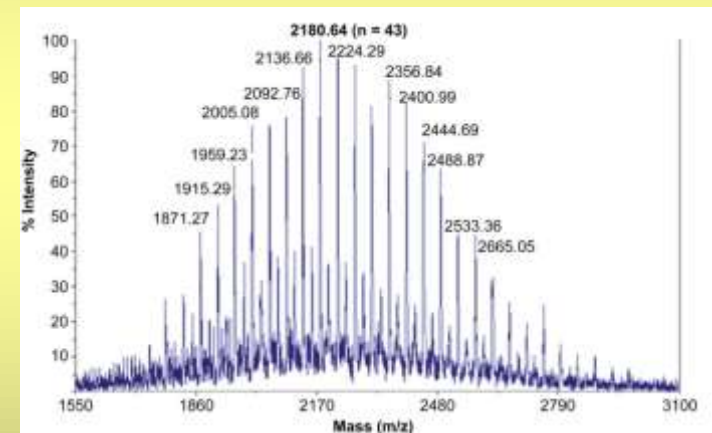


**Dispersity ( $\mathcal{D}$ )** is defined as  $\mathcal{D} = M_m/M_n$  where  $M_m$  is the mass-average molar mass and  $M_n$  is the number-average molar mass.

Polymers have no an individual mass number, it is not a single molecule, but a set of olygomers

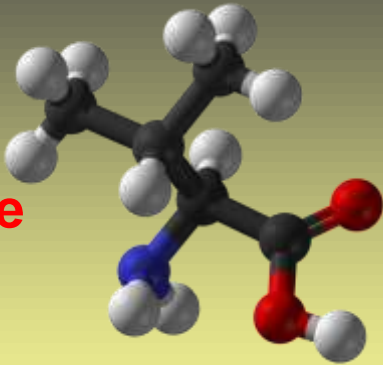
This property is called polydispersity

The name of polymer is given for its average mass (n =polymeryzation degree)

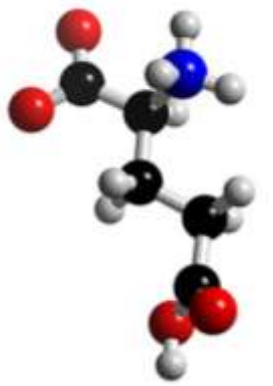


# Amino acids

**Valine**

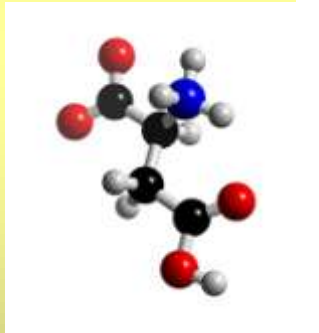


**Glutamic acid**

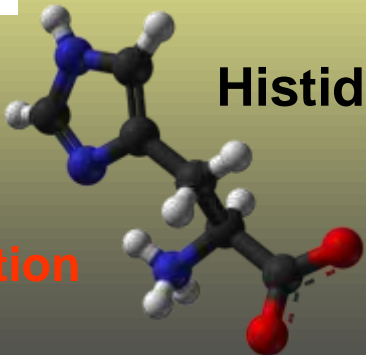


**Anion**

**Aspartic acid**



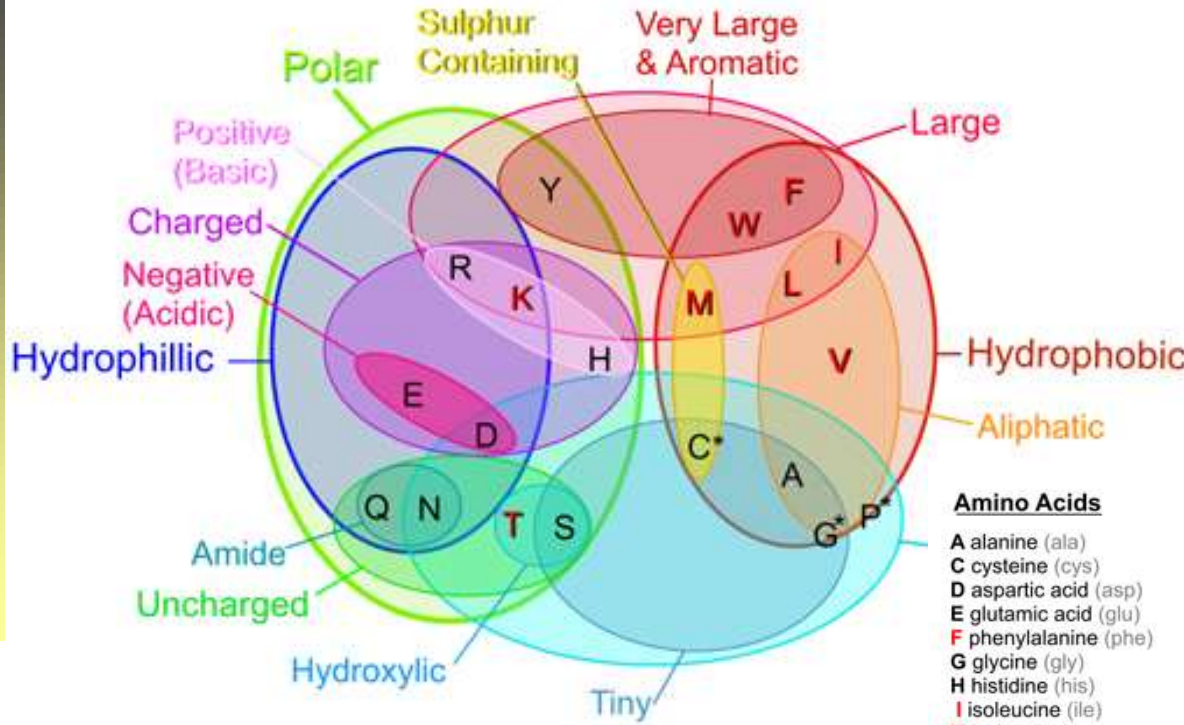
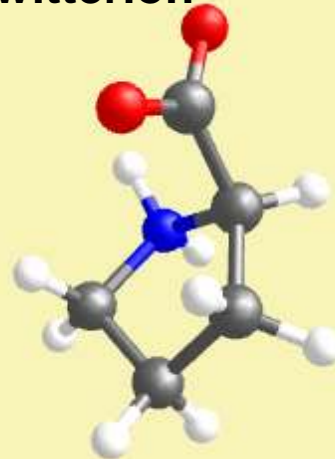
**Cation**



**Histidine**

**Proline**

**Zwitterion**

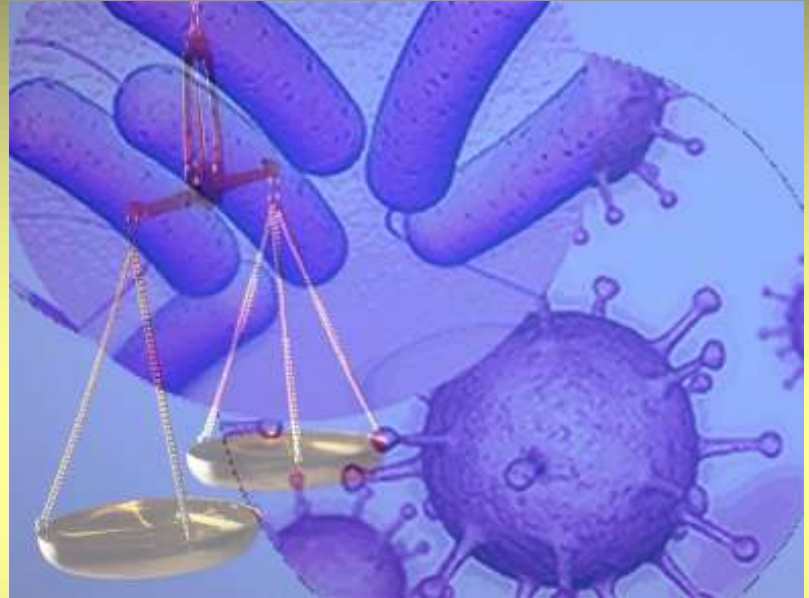


## Amino Acids

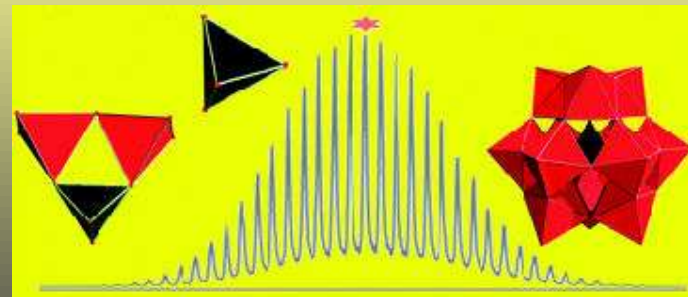
- A alanine (ala)
- C cysteine (cys)
- D aspartic acid (asp)
- E glutamic acid (glu)
- F phenylalanine (phe)
- G glycine (gly)
- H histidine (his)
- I isoleucine (ile)
- K lysine (lys)
- L leucine (leu)
- M methionine (met)
- N asparagine (asn)
- P proline (pro)
- Q glutamine (gln)
- R arginine (arg)
- S serine (ser)
- T threonine (thr)
- V valine (val)
- W tryptophan (trp)
- Y tyrosine (tyr)

# Mass spectrometry

is an **efficient tool** in studies of noncovalent complexes of organic and biological molecules



monitoring of self-assembling of nanoparticles



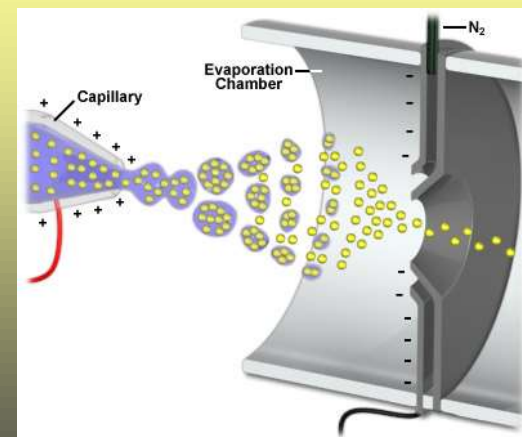
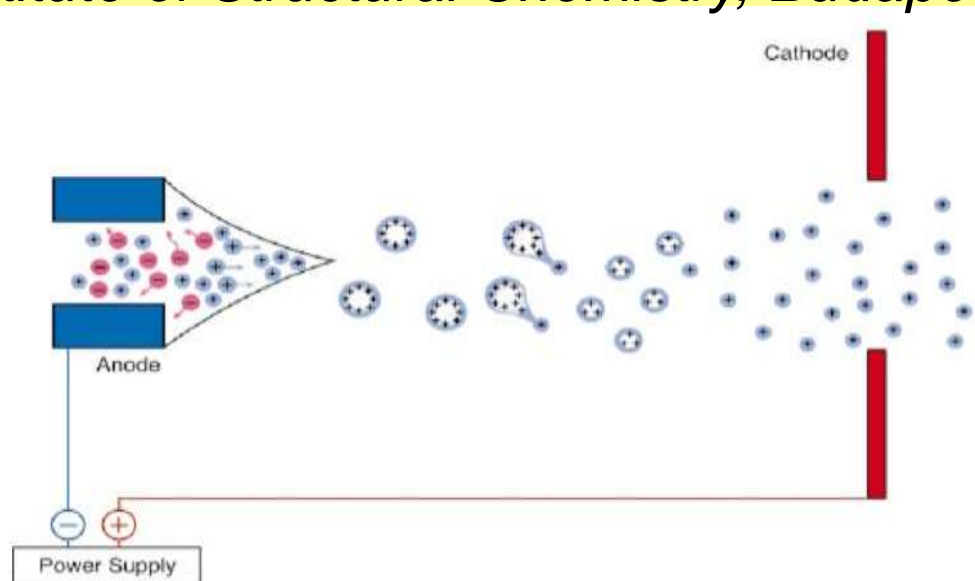
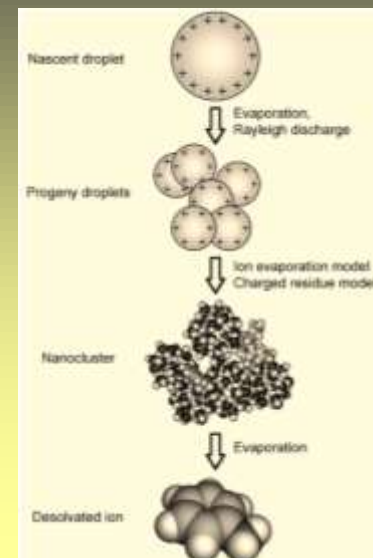


# METHODS:

## Mass spectrometry

### Mass spectrometry with electrospray ionization (ESI)

PE Sciex API 2000 Triple Quadrupole MS  
(PE Sciex, Canada) *in collaboration with  
Institute of Structural Chemistry, Budapest*



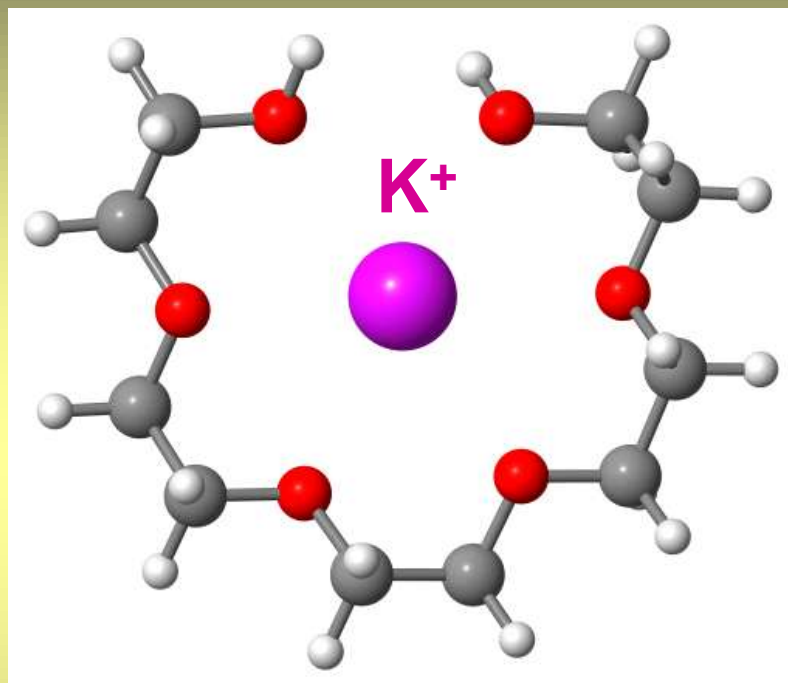
# Molecular dynamics simulation

To establish structures of complexes registered in mass spectrometric experiment

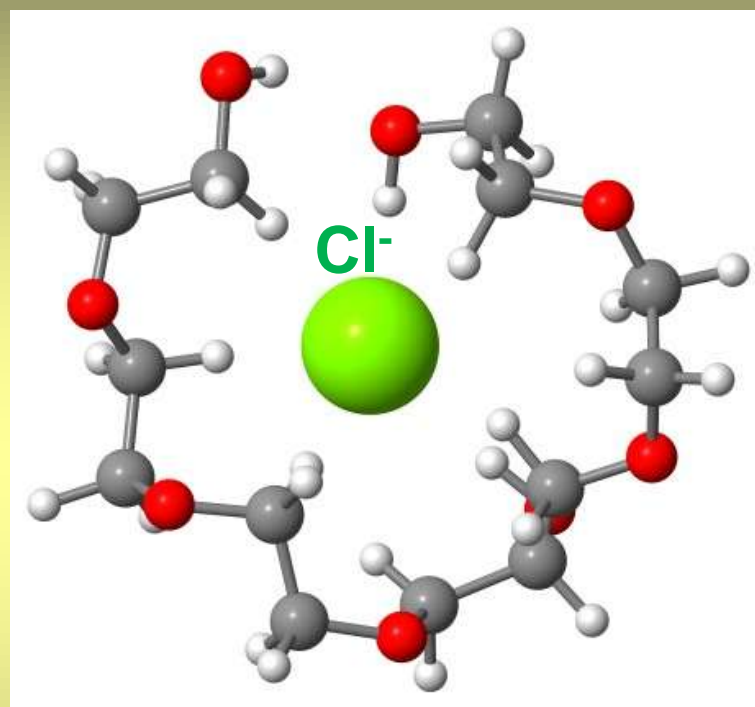
The screenshot displays the VMD 1.8.5 OpenGL Display window. The main view shows a 3D ball-and-stick model of a protein structure. The interface includes several panels and windows:

- Graphical Representations:** A panel showing the selected molecule '0: pro-h\_og\_out.coor' with buttons for 'Create Rep' and 'Delete Rep'. It also has columns for 'Style', 'Color', and 'Selection'.
- VMD Main:** A menu bar with options: File, Molecule, Graphics, Display, Mouse, Extensions, Help.
- Table:** A table with columns: ID, T, A, D, F, Molecule, Atoms, Frames, Vo. The first row is highlighted in yellow: 0, T, A, D, F, pro-h\_og\_out.coor, 100, 2, 0.
- Draw style:** A panel with a 'Name' field and a 'Drawing Method' dropdown set to 'CPK'.
- Control Panel:** Includes 'zoom', 'Loop', 'step', '1', and 'speed' controls.
- System Tray:** Shows the time as 9:15 and the user as cpe66.

## POLYMERIC CHANE WRAPS AROUND THE IONS IN Polymer •inorganic ion COMPLEXES



PEG<sub>5</sub>•K<sup>+</sup>



OEG<sub>5</sub>•Cl<sup>-</sup>

Assumption: similar structures can be formed with organic ions

Observation of poly (ethylene glycol) clusters with the chlorine anion in the gas phase under electrospray conditions // Rapid Communications in Mass Spectrometry, 2011, V.25, N. ,P. 713-718

# Experimental results

**Valine and proline –  
representatives of  
hydrophobic amino acids**

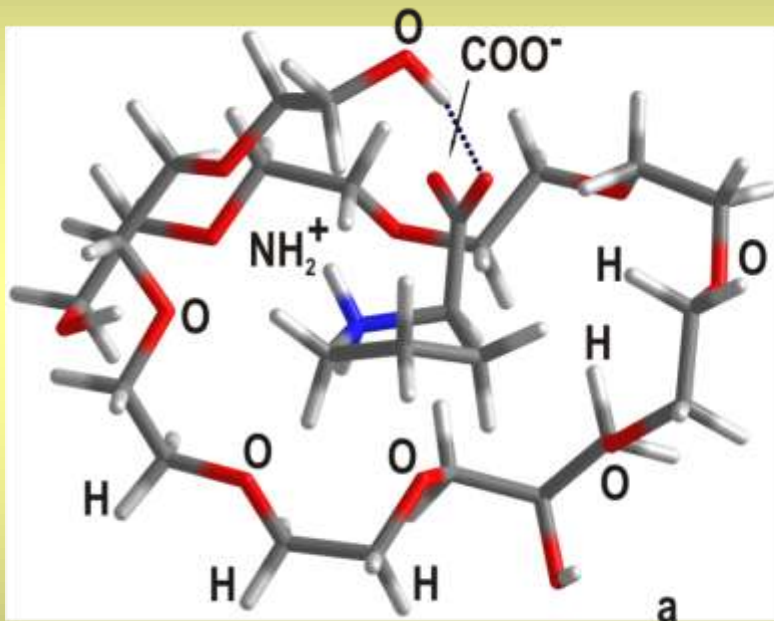
# Experimental results

## ESI mass spectra of “Valine – OEG” nanoclusters

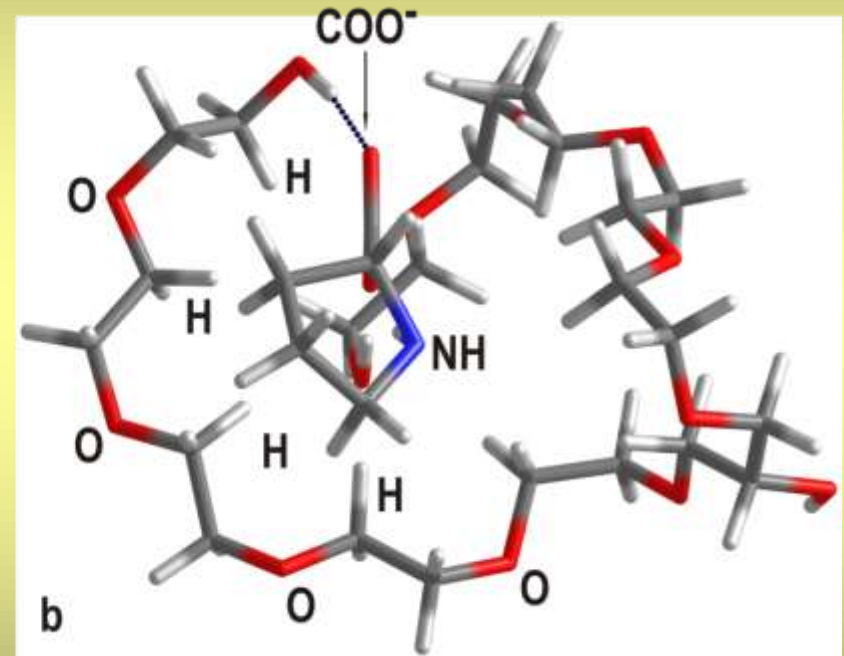
Косевич М.В., Зобнина В.Г., Боряк О.А., Шелковский В.С., Гомори А., Векей К. Исследование раствора аминокислоты валина в криопротекторе этиленгликоле при низких температурах методом вторично-ионной масс-спектрометрии // Масс-спектрометрия, 2006, Т. 3, N 1, С. 33-42.

# Computer Simulation

Structures of complexes formed by oxyethylated glycerol oligomers OEGn and PEGn with amino acid Proline



Protonated Proline

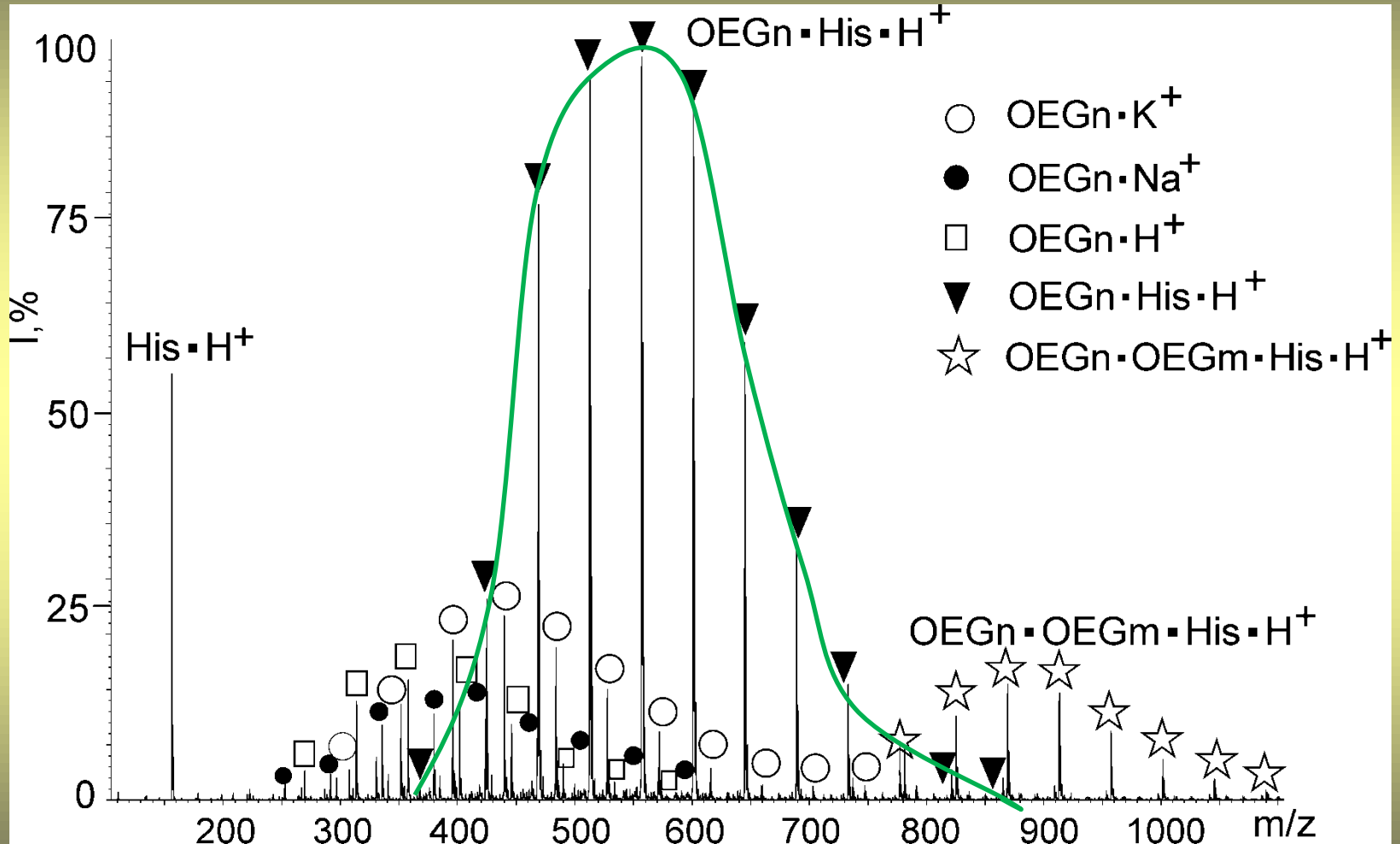


Deprotonated Proline

**Histidine – representative of  
ionic positively charged amino  
acids**

# Experimental results

## ESI mass spectra of "Histidine – OEG" nanoparticles

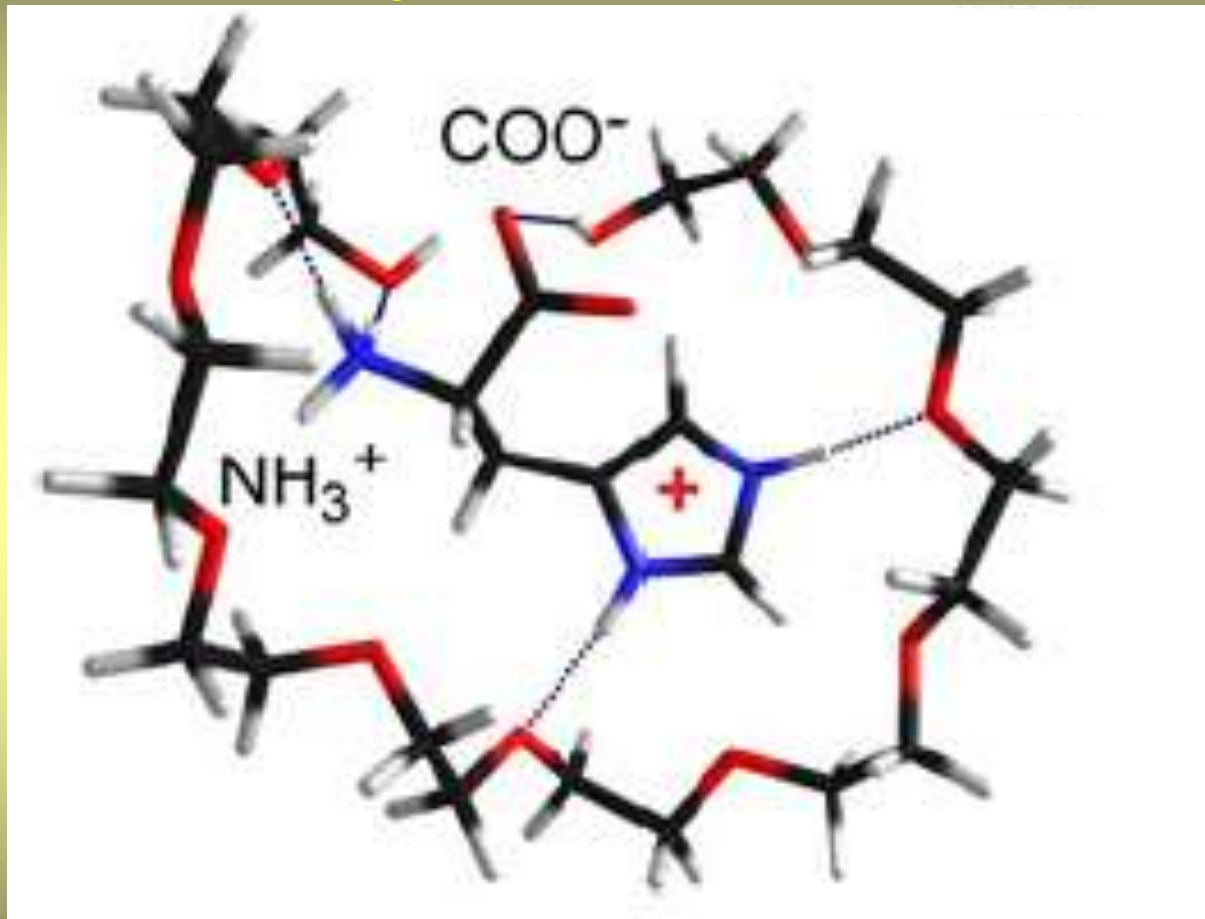


Interactions of oligomers of organic polyethers with histidine amino acid // Rapid Communications in Mass Spectrometry, 2012, V. 26, N 5. P. 532-540



# Molecular dynamics simulation

$\text{PEG}_{10}\cdot\text{His}\cdot\text{H}^+$



His in cationic zwitterionic form

**Aspartic acid,  
Glutamic acid – representatives  
of ionic negatively charged  
amino acids**

# CONCLUSION

Structural assembling and organization in nanoparticles are realized owing to the wrapping of polyether chain around amino acids charged groups

**The results obtained for the self-organization and structure of the studied model nanoparticles can be applied for modeling of the structure of the larger assemblies of proteins with polyethers using the same structural motifs**