

On the interface between experiment and computation, the experimentalists viewpoint



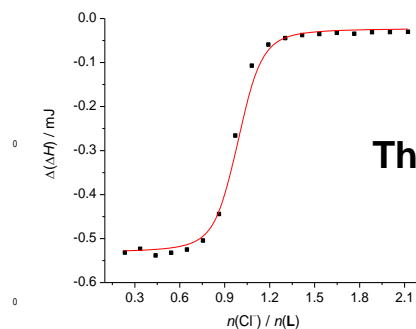
Gordan Horvat

Department of Chemistry, Faculty of Science,
University of Zagreb

Observation

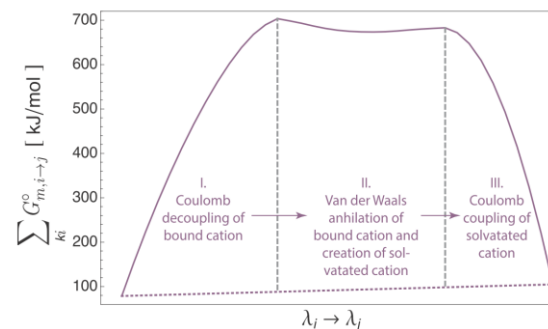


Microcalorimetry

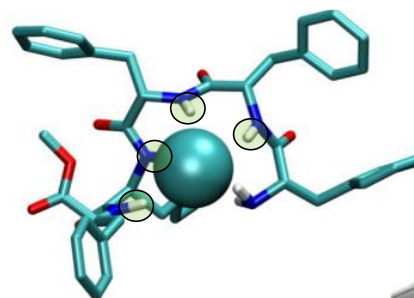
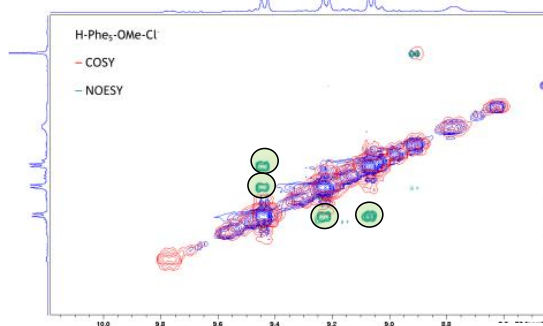


Thermodynamics

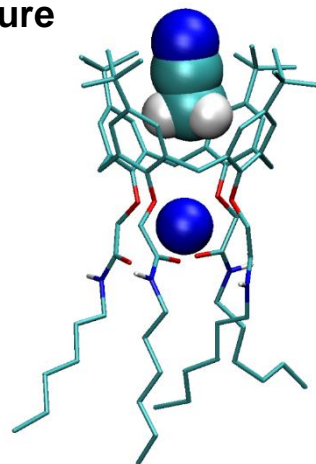
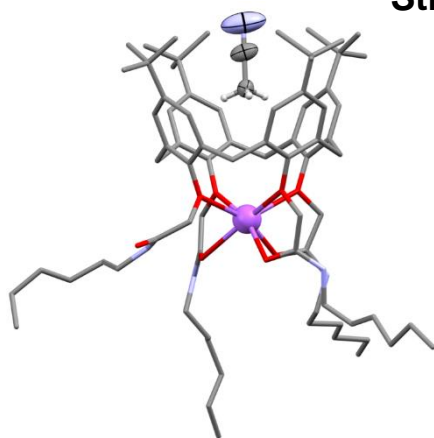
Modelling



NMR Spectroscopy



Structure



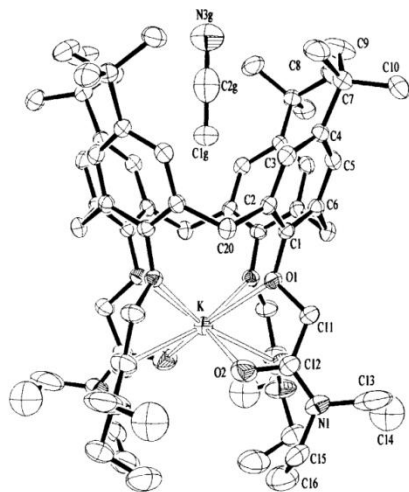
$$U(\mathbf{r}_N) = \sum_{bonds} K_r (r_i - r_{eq})^2 + \sum_{angles} K_\phi (\theta_i - \theta_{eq})^2 + E_{torsions} + \frac{1}{2} \sum_{i=1}^N \sum_{j=1, j \neq i}^N \frac{q_i q_j}{4\pi\epsilon_0 d_{ij}} + \frac{1}{2} \sum_{i=1}^N \sum_{j=1, j \neq i}^N 4\epsilon_{ij} \left(\left(\frac{\sigma_{ij}}{d_{ij}} \right)^{12} - \left(\frac{\sigma_{ij}}{d_{ij}} \right)^6 \right)$$

Classical MD simulations



X-ray Crystallography

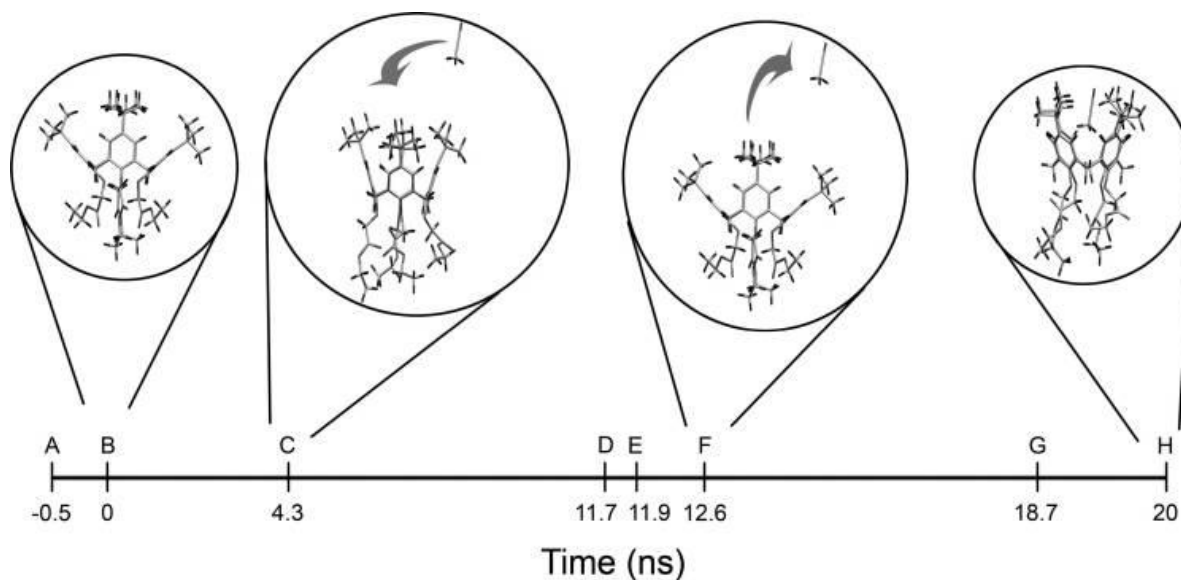
Calix[4]arene cation complexes, PhD thesis



Arduini et al., *Tetrahedron* **57** (2001) 2411–2417.

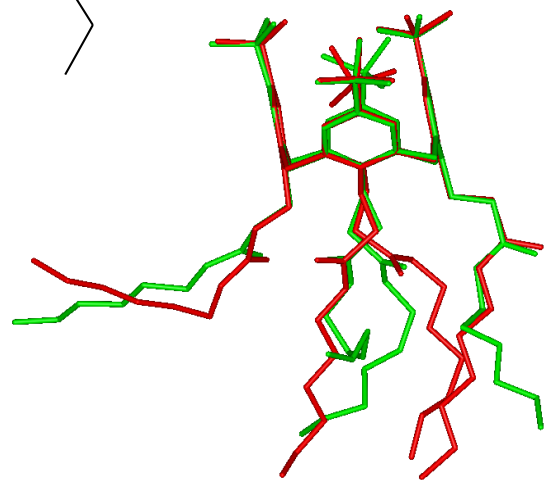
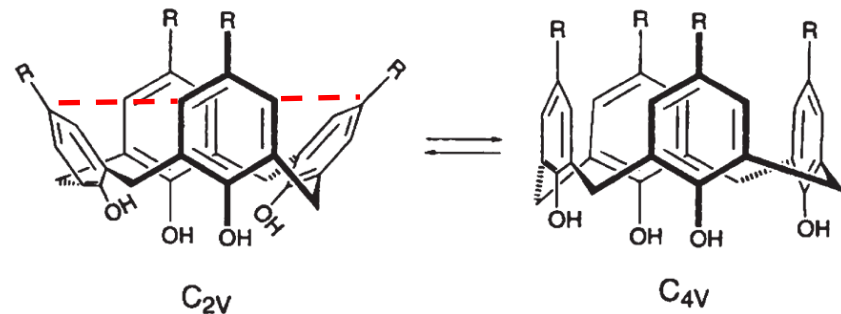
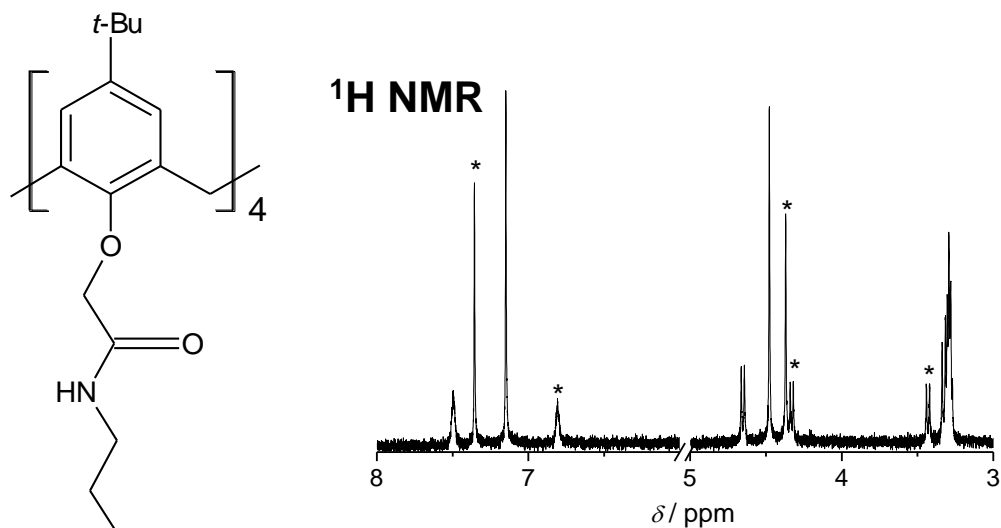


Arena et al., *New J. Chem.* **28** (2004) 56–61.

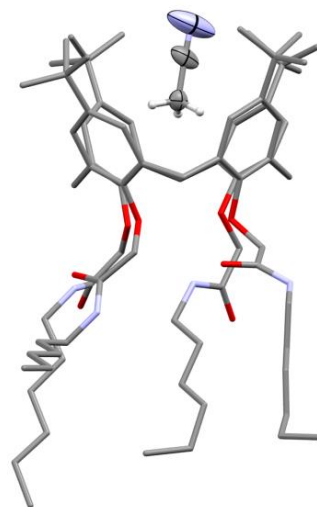


A. S. de Araujo, O. E. Piro, E. E. Castellano, A. F. Danil de Namor, *J. Phys. Chem. A* **112** (2008) 11885–11894.

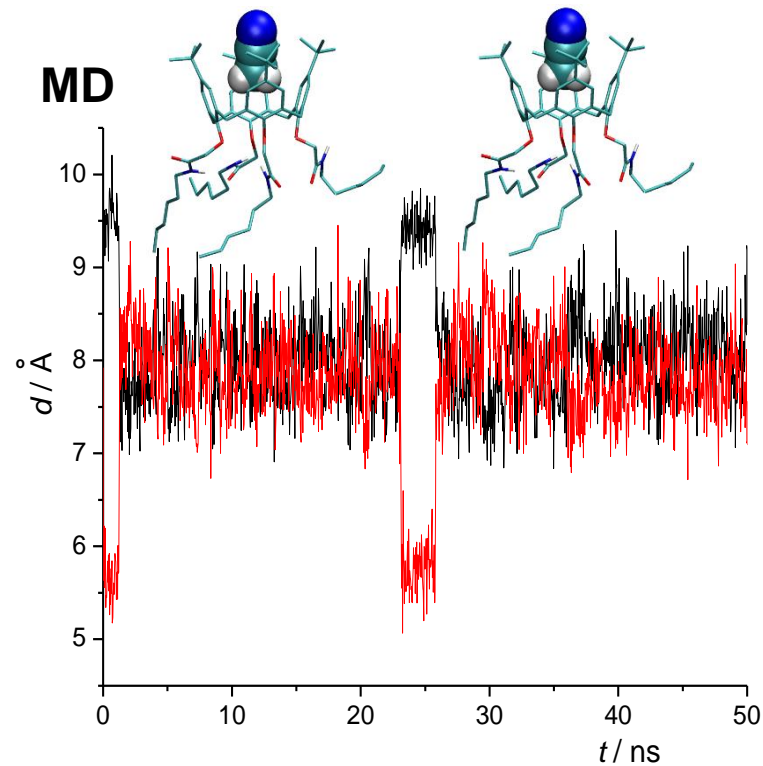
Free calix[4]arene in acetonitrile



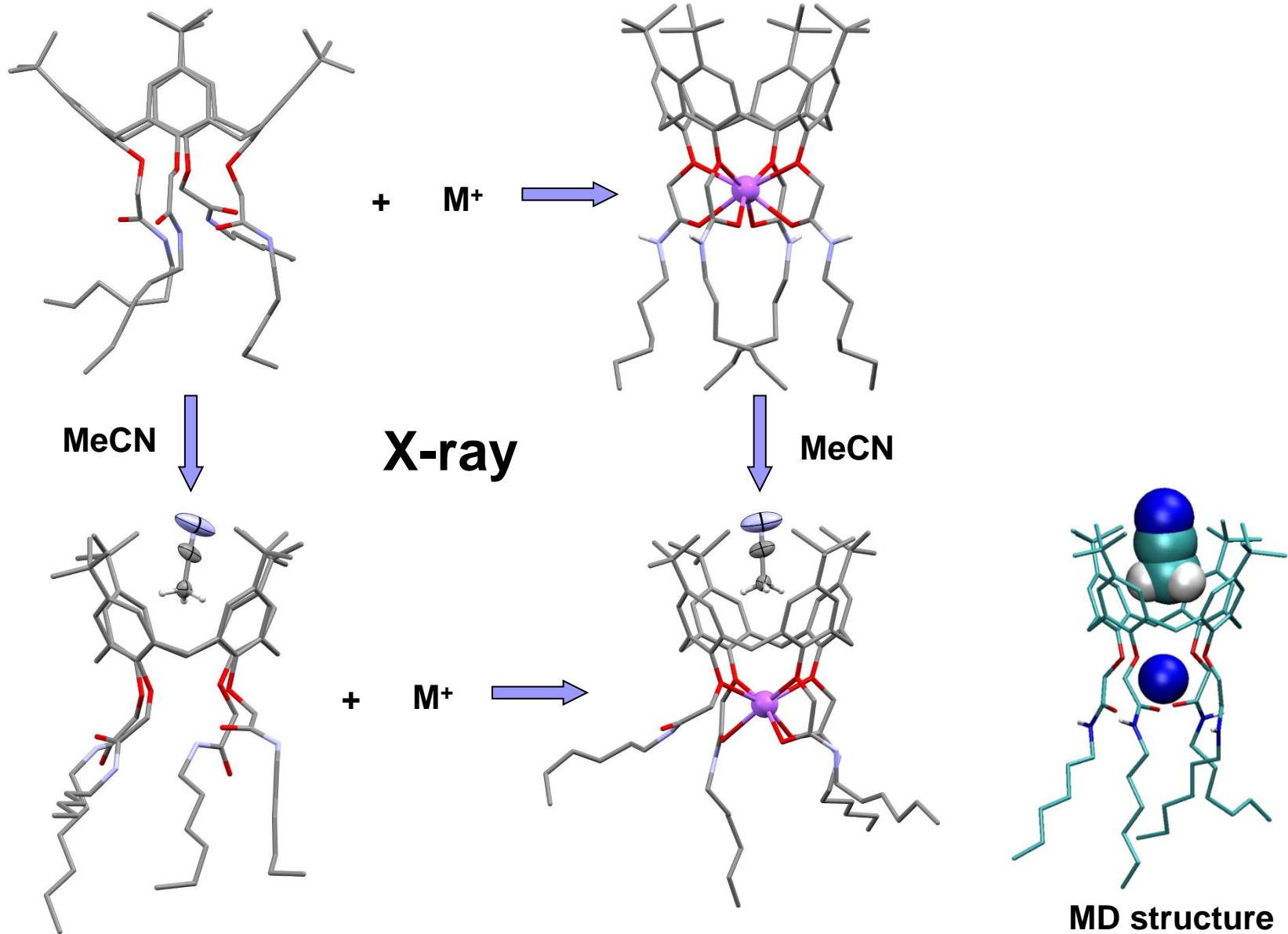
Free: X-ray vs MD



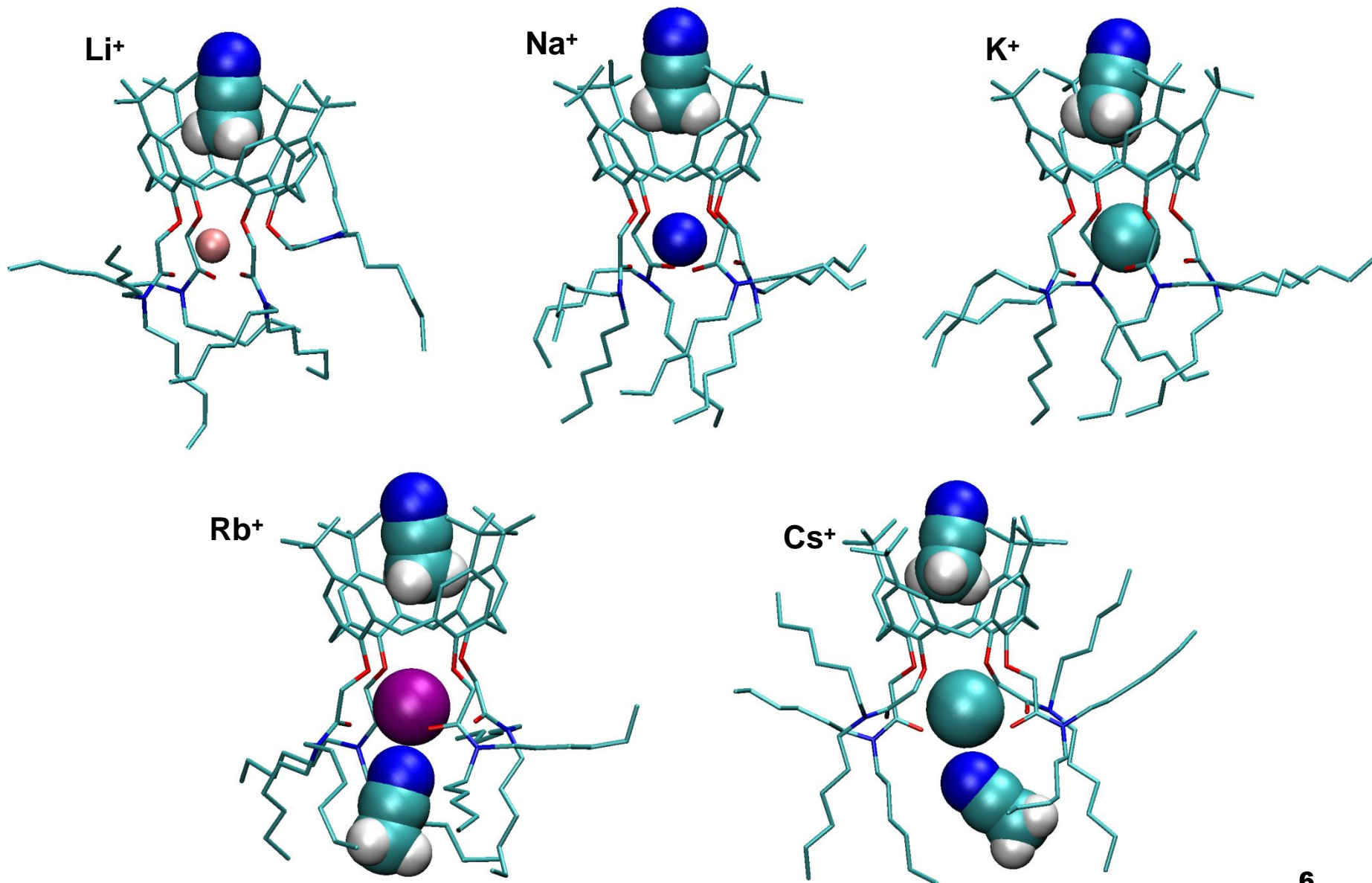
MeCN adduct: X-ray



Crystal structure vs MD

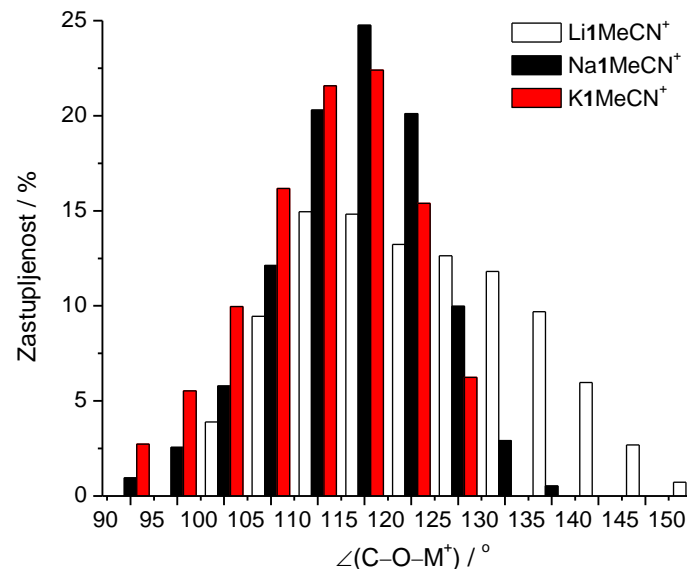
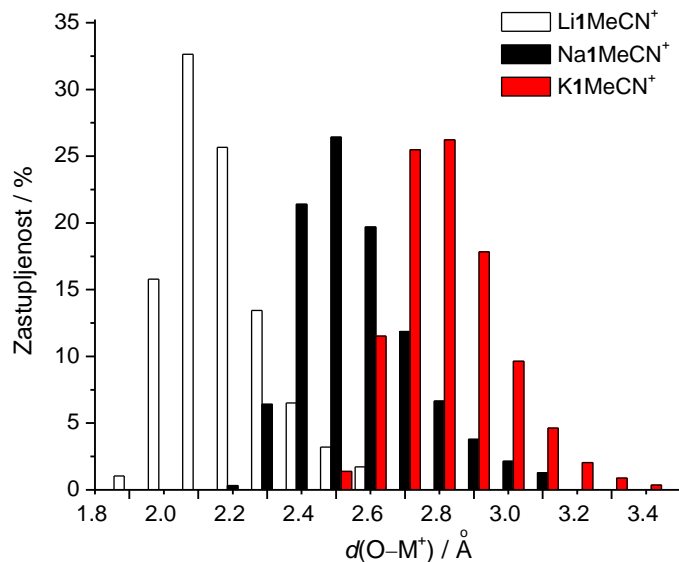


MD structures of M^+ calix[4]arene complexes

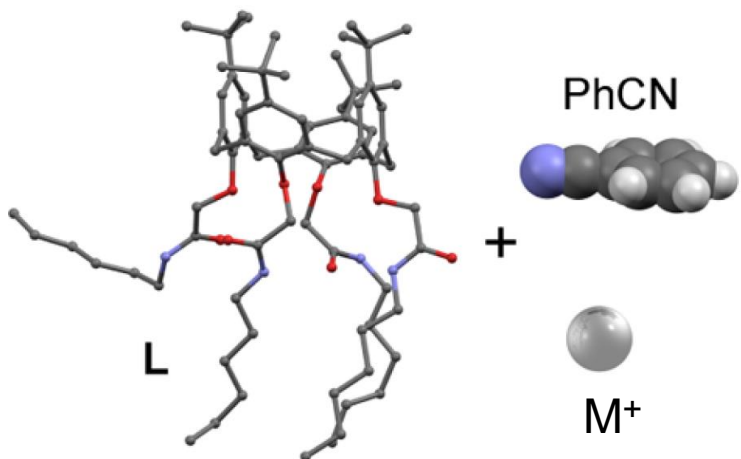


Interactions and structure

	Li ⁺		Na ⁺		K ⁺	
	Li1 ⁺	Li1MeCN ⁺	Na1 ⁺	Na1MeCN ⁺	K1 ⁺	K1MeCN ⁺
$E(M^+-L) / \text{kJ mol}^{-1}$	-492	-507	-	-431	-353	-360
$E(L-\text{MeCN}) / \text{kJ mol}^{-1}$	-505	-561	-	-578	-508	-594
$E(L-\text{MeCN}_{\text{inkl}}) / \text{kJ mol}^{-1}$	-	-52	-	-51	-	-50
$E(M^+-\text{MeCN}) / \text{kJ mol}^{-1}$	-18	-15	-	-6	-3	1
$E(M^+-\text{MeCN}_{\text{inkl}}) / \text{kJ mol}^{-1}$	-	8	-	8	-	7
$t_{\text{total}} / \text{ns}$		50		50		50
t / t_{kupno}	0,009	0,991	0	1	0,001	0,999
$N(\text{carbonyl})$	2,3	2,8	-	3,3	3,7	3,9
$N(\text{hydrogen bonds})$	0,56	0,65	-	0,43	0,13	0,07
$N(\text{MeCN}_{\text{inkl}})$	-	6	-	1	-	4
$\bar{d} / \text{\AA}$	7,53	7,85	-	7,85	7,60	7,80
	8,13	7,93	-	7,85	7,80	7,81
$ d - d_{\text{ref}} / \text{\AA}$	0,41	0,22	-	0,21	0,38	0,20
	0,46	0,23	-	0,21	0,45	0,21
$\sigma(d) / \text{\AA}$	0,42	0,28	-	0,26	0,48	0,25
	0,50	0,28	-	0,26	0,51	0,25

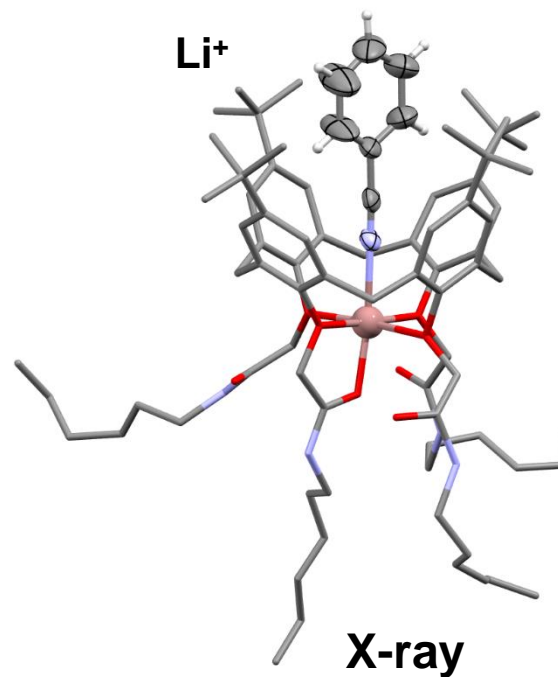
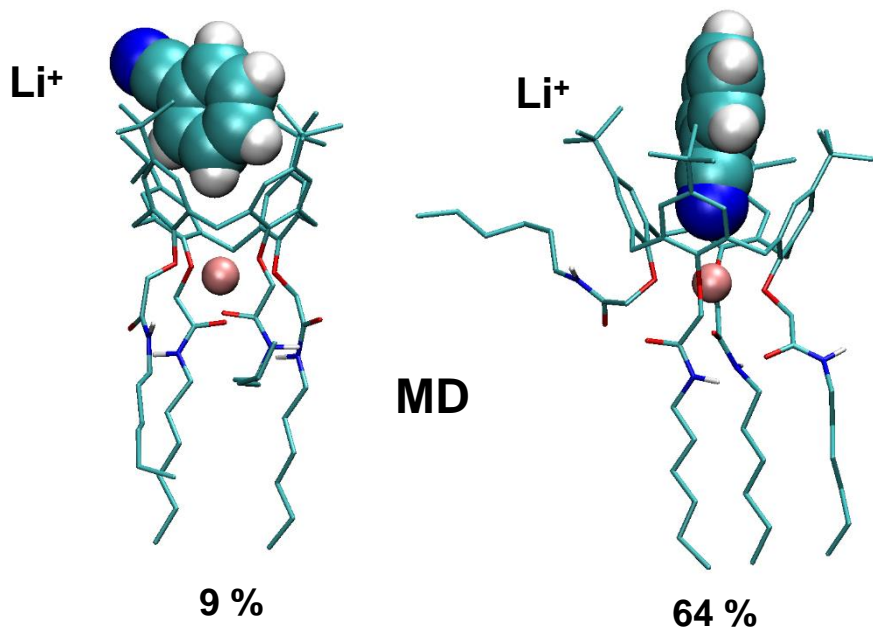


Calix[4]arene-cation complexes in benzonitrile



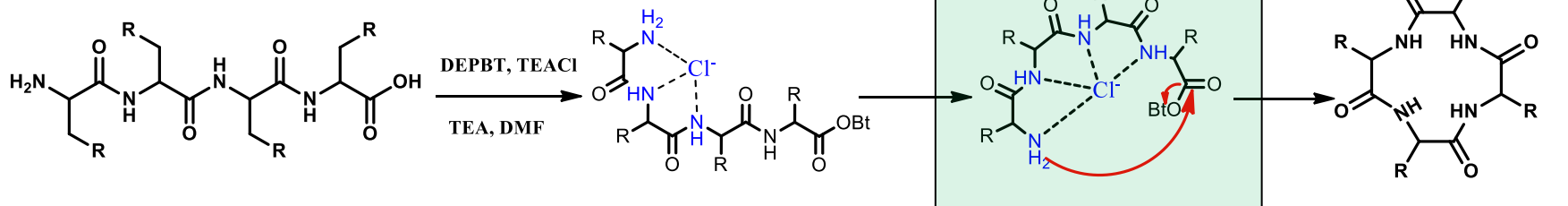
ITC

Kation	$\log\left(\frac{K}{\text{dm}^3 \text{mol}^{-1}}\right) \pm \text{SE}$	$\frac{(\Delta_r H^\ominus \pm \text{SE})}{\text{kJ mol}^{-1}}$	$\frac{(\Delta_r S^\ominus \pm \text{SE})}{\text{J K}^{-1} \text{mol}^{-1}}$
Li ⁺	$6,17 \pm 0,01$	$-8,9 \pm 0,1$	$88,1 \pm 0,4$
Na ⁺	$5,54 \pm 0,01$	$-16,6 \pm 0,1$	$50,4 \pm 0,5$
K ⁺	--_a	--_a	--_a



Cyclopeptide anion receptors

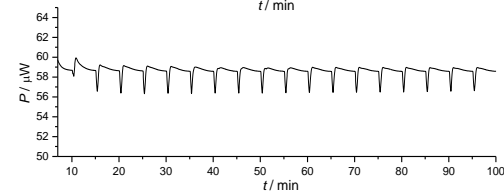
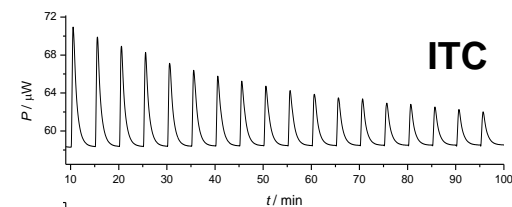
Synthesis



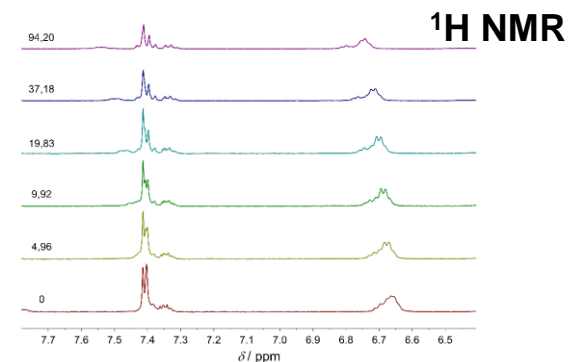
N. Vidović, G. Horvat, D. Riva, T. Rinkovec, N. Cindro, V. Tomišić, G. Speranza, *Org. Lett.* **2020**, *22*, 2129.

Cyclization yields

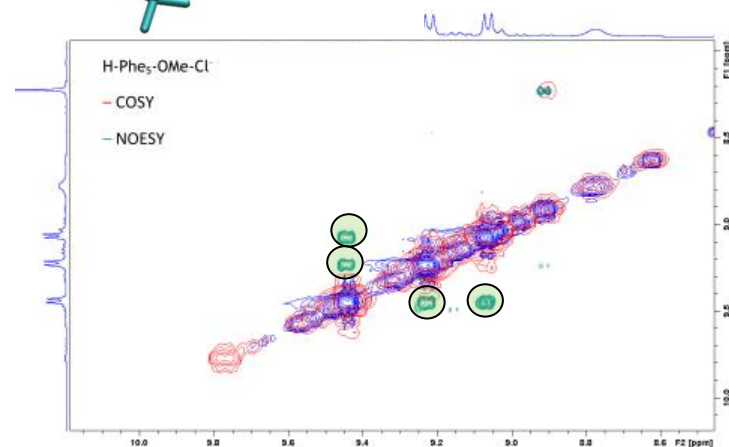
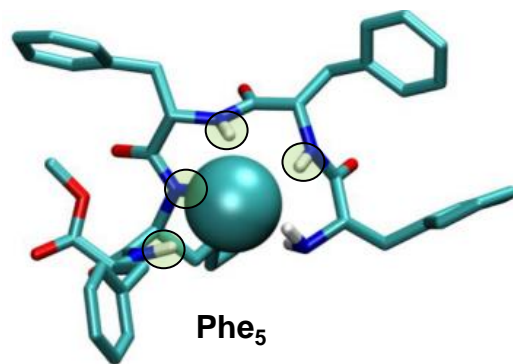
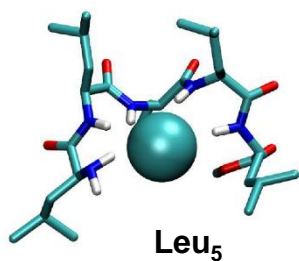
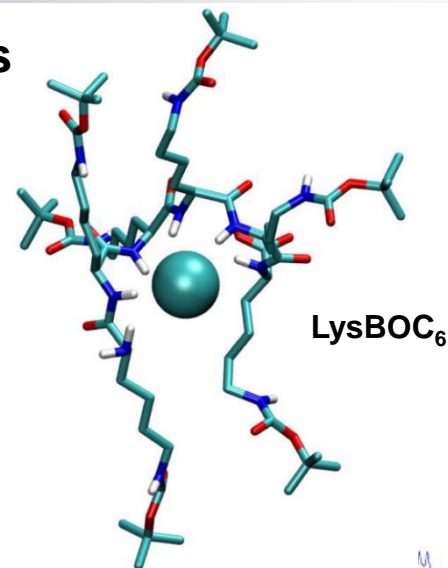
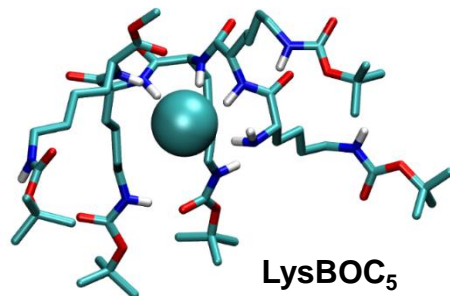
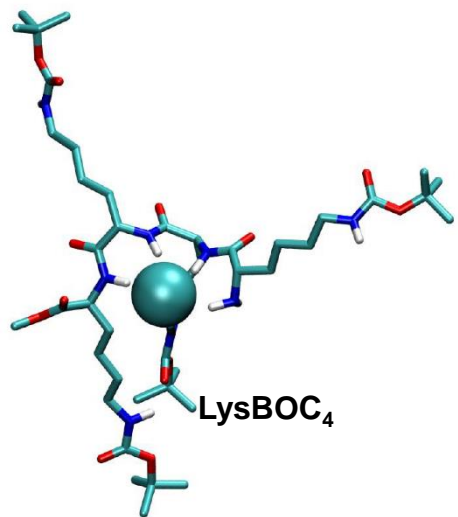
linear peptide ^a	Yield (%)			
	LiCl/NaCl	NaTPB	TEACl	NaClO ₄
K4 (1)	21 ^b	8	47	0
K5 (2)	35 ^c	26	43	0
K6 (3)	15 ^c	10	17	0
L5 (7)		15	52	0
(FLL) ₂ (8)	<5 ^c	6	23	<5
I4 (11)	<5 ^b	<5	18	0
I5 (12)	11 ^c	<5	26	0
S5 (13)	21 ^c		29	
F5 (14)			46	



log K around 1.5



Structures of chloride-linear peptide complexes

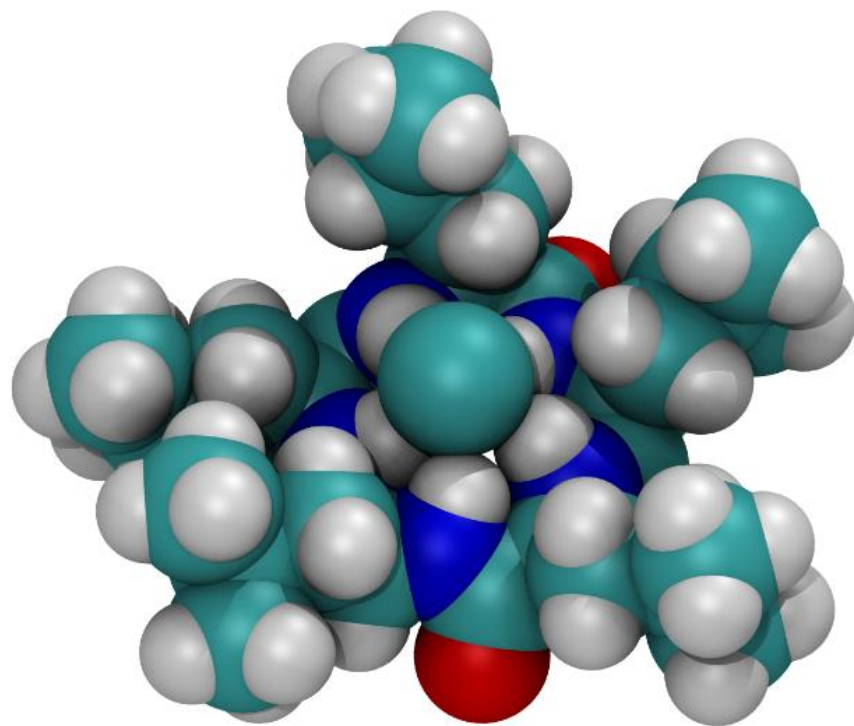


RESEARCH HIGHLIGHTS

Nature Reviews Chemistry | <https://doi.org/10.1038/s41570-020-0185-0> | Published online 15 April 2020

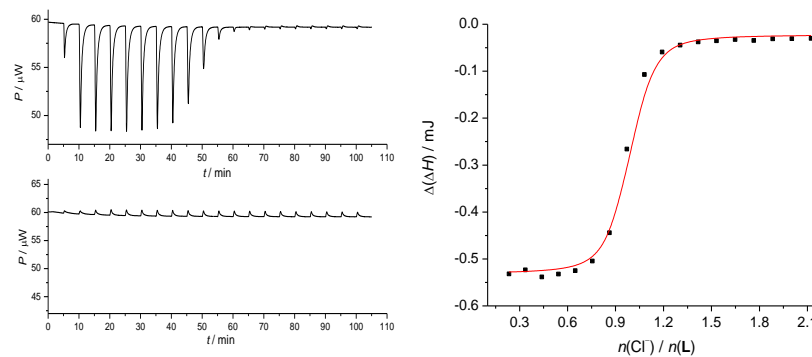
Running rings around chloride

Thermodynamic and MD studies of anion complexation by cyclopentaleucine in MeCN and DMSO



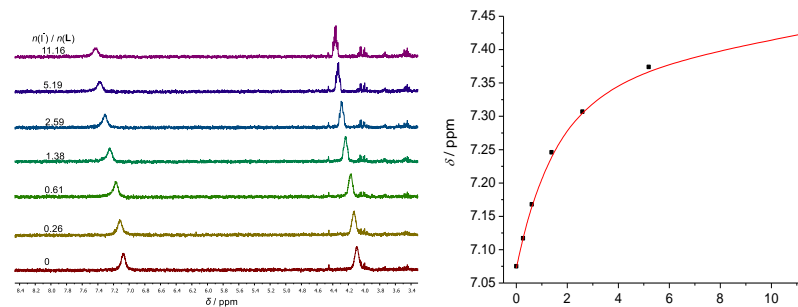
MD simulations

ITC



Leu₅ titration with Cl⁻ in MeCN

¹H NMR



Leu₅ titration with I⁻ in MeCN

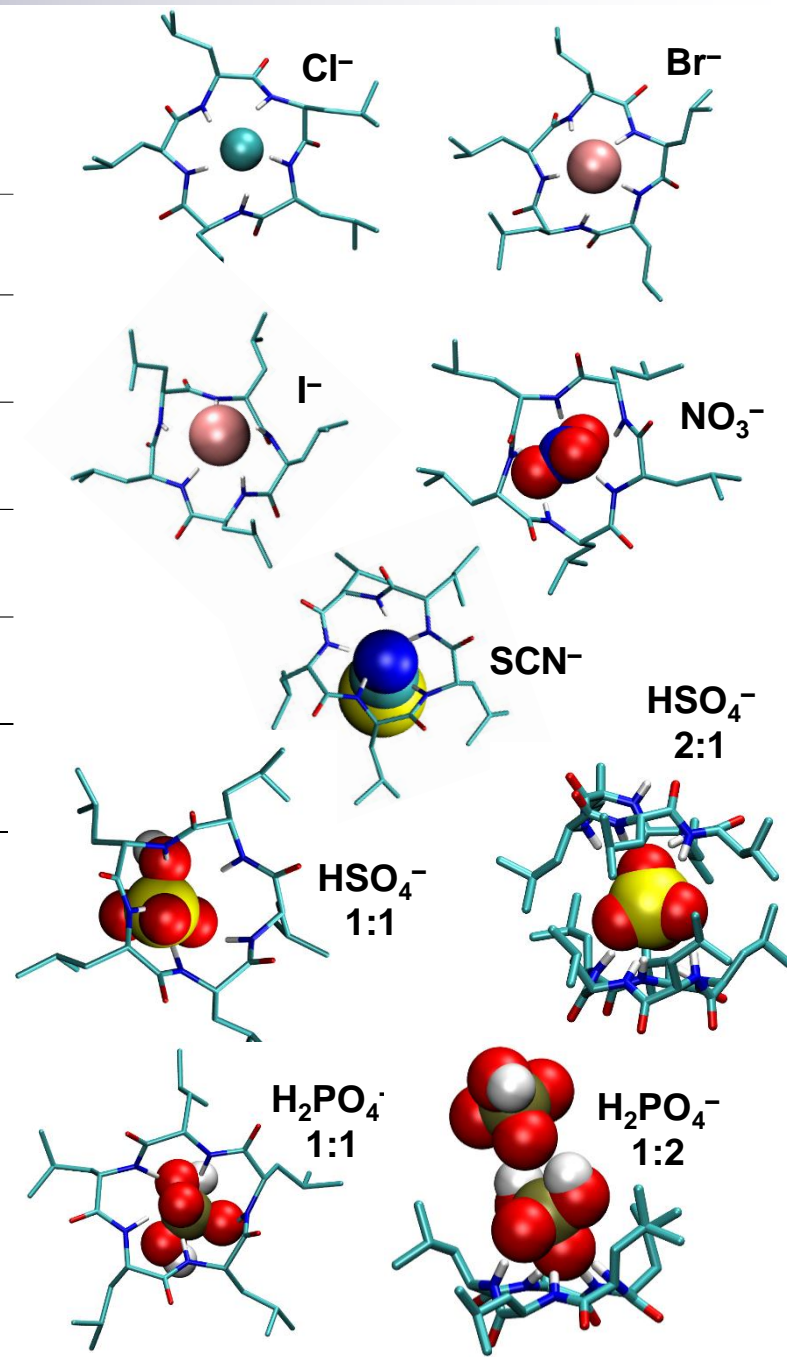
Cyclopeptide-anion complexation in MeCN

1:1 complexes

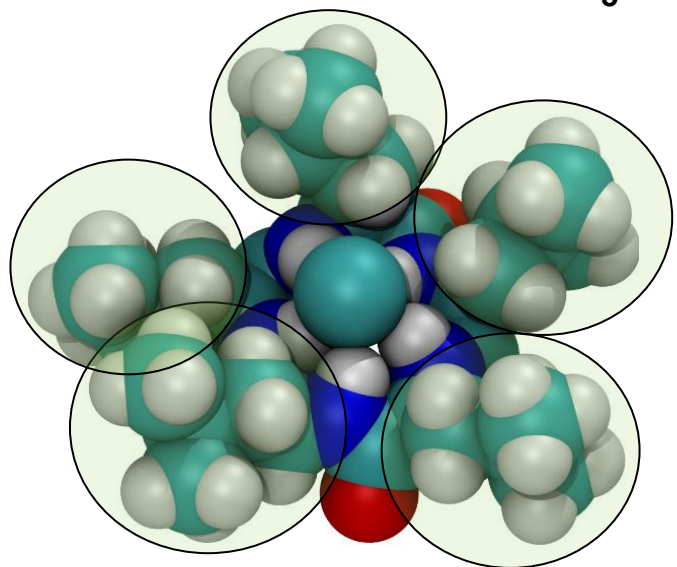
Anion	$\log\left(\frac{K}{\text{dm}^3\text{mol}^{-1}}\right) \pm \text{SE}$	$(\Delta_r G^\circ \pm \text{SE})$ kJ mol ⁻¹	$(\Delta_r H^\circ \pm \text{SE})$ kJ mol ⁻¹	$(\Delta_r S^\circ \pm \text{SE})$ J mol ⁻¹ K ⁻¹
Cl ⁻	5.84 ± 0.03 ^a	-33.33 ± 0.11	-10.68 ± 0.08	75.9 ± 0.6
Br ⁻	4.70 ± 0.01 ^a	-26.82 ± 0.06	-4.84 ± 0.02	73.7 ± 0.2
I ⁻	3.20 ± 0.03 ^a 3.12 ^b	-18.27 ± 0.15	1.91 ± 0.08	67.7 ± 0.3
SCN ⁻	2.90 ± 0.02 ^a 2.77 ^b	-16.5 ± 0.1	-4.72 ± 0.095	39.7 ± 0.7
NO ₃ ⁻	3.18 ± 0.01 ^a 3.29 ^b	-18.16 ± 0.04	-6.08 ± 0.04	40.5 ± 0.3

Complexes of higher stoichiometry

Anion	$\log\left(\frac{K}{\text{dm}^3\text{mol}^{-1}}\right) \pm \text{SE}$	$(\Delta_r G^\circ \pm \text{SE})$ kJ mol ⁻¹	$(\Delta_r H^\circ \pm \text{SE})$ kJ mol ⁻¹	$(\Delta_r S^\circ \pm \text{SE})$ J mol ⁻¹ K ⁻¹
HSO ₄ ⁻	4.38 ± 0.02 (1:1)	-25.0 ± 0.1	-16.0 ± 0.1	30.2 ± 0.7
	2.96 ± 0.05 (2:1)	-16.9 ± 0.3	0.4 ± 0.3	58 ± 2
H ₂ PO ₄ ⁻	4.56 ± 0.02 (1:1)	-26.0 ± 0.1	-12.6 ± 0.3	45 ± 1
	4.18 ± 0.02 (1:2)	-23.8 ± 0.1	-79.7 ± 0.6	-187 ± 2



Structures of anion-Leu₅ complexes

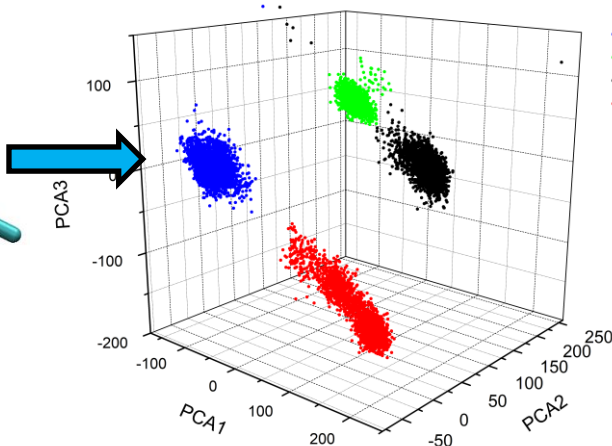
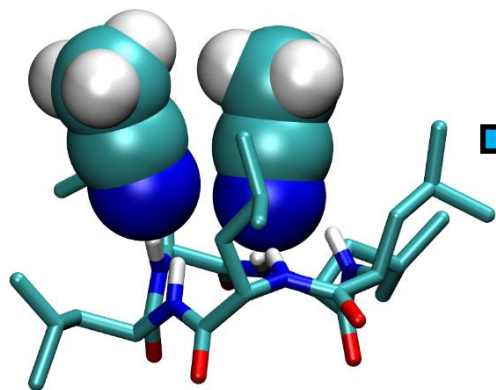


endo complexes

Average number of coordinated amide protons in MeCN

anion	Cl ⁻	Br ⁻	I ⁻	NO ₃ ⁻	SCN ⁻	HSO ₄ ⁻	H ₂ PO ₄ ⁻
N(-H)	4.98	4.98	4.88	4.95	4.77	4.75	4.97

Structure of free Leu₅ in MeCN



- Cluster 1
- Cluster 2
- Cluster 3
- Cluster 4

Free energy calculations by MD

Molecular Dynamics Simulations of Selected Homocyclopentapeptides with Chloride, Bromide and Iodide Anions

Ivan Petranović,¹ Gordana Horvat¹

¹Department of Chemistry, Faculty of Science, Zagreb, Croatia
*E-mail: ipetranovic@chem.pmf.hr

Introduction

Introduction of a free energy calculation in molecular dynamics simulation is a challenging task. The aim of these calculations is to obtain the relative free energy of different states of a system. The aim of these calculations is to obtain the relative free energy of different states of a system. The aim of these calculations is to obtain the relative free energy of different states of a system.

Ion exchange

The aim of the present study is to determine the relative free energy of different states of a system. The aim of the present study is to determine the relative free energy of different states of a system. The aim of the present study is to determine the relative free energy of different states of a system.

Complexation

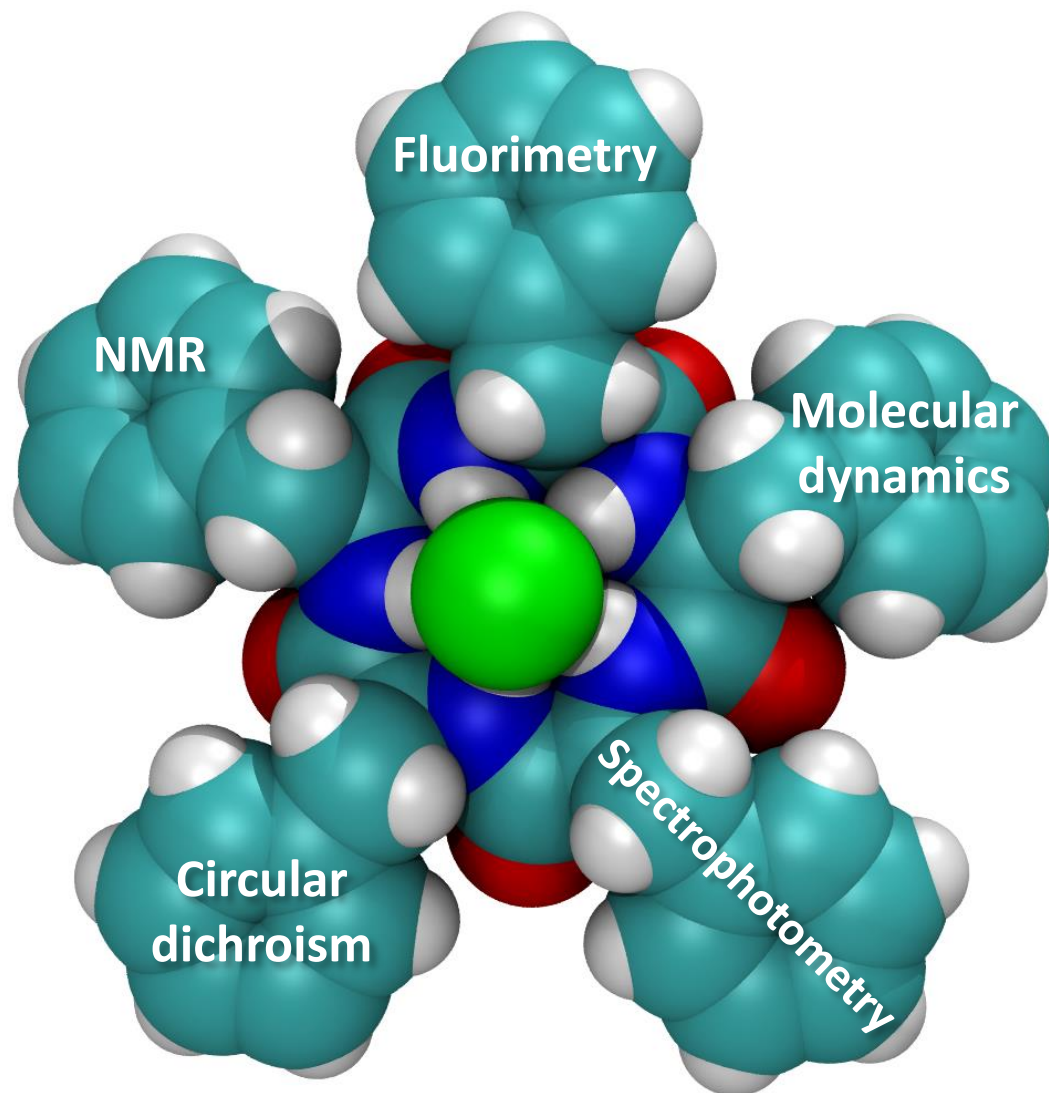
The aim of the present study is to determine the relative free energy of different states of a system. The aim of the present study is to determine the relative free energy of different states of a system. The aim of the present study is to determine the relative free energy of different states of a system.

Side chain mutation

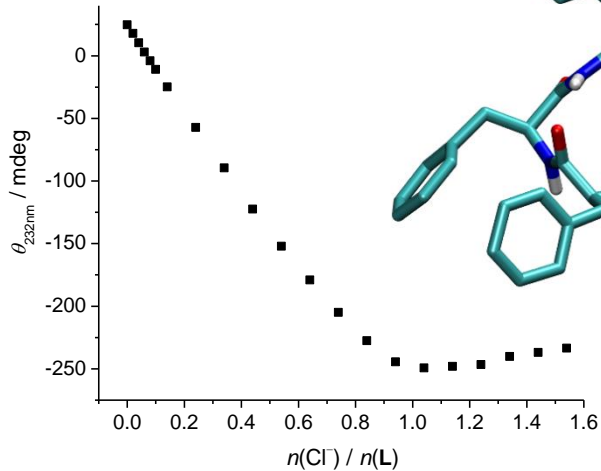
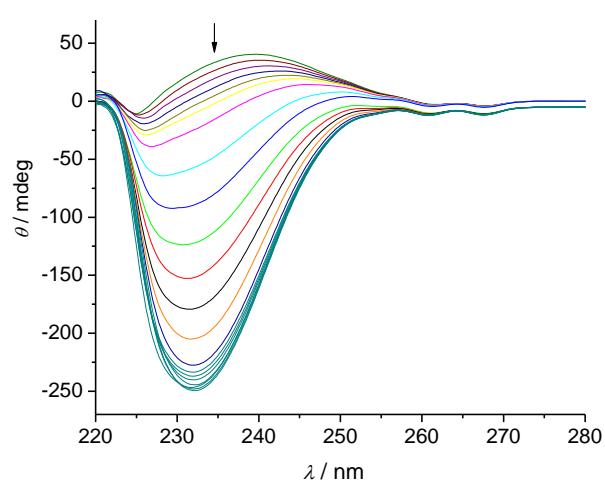
The aim of the present study is to determine the relative free energy of different states of a system. The aim of the present study is to determine the relative free energy of different states of a system. The aim of the present study is to determine the relative free energy of different states of a system.

CCD2023 poster: Ivan Petranović

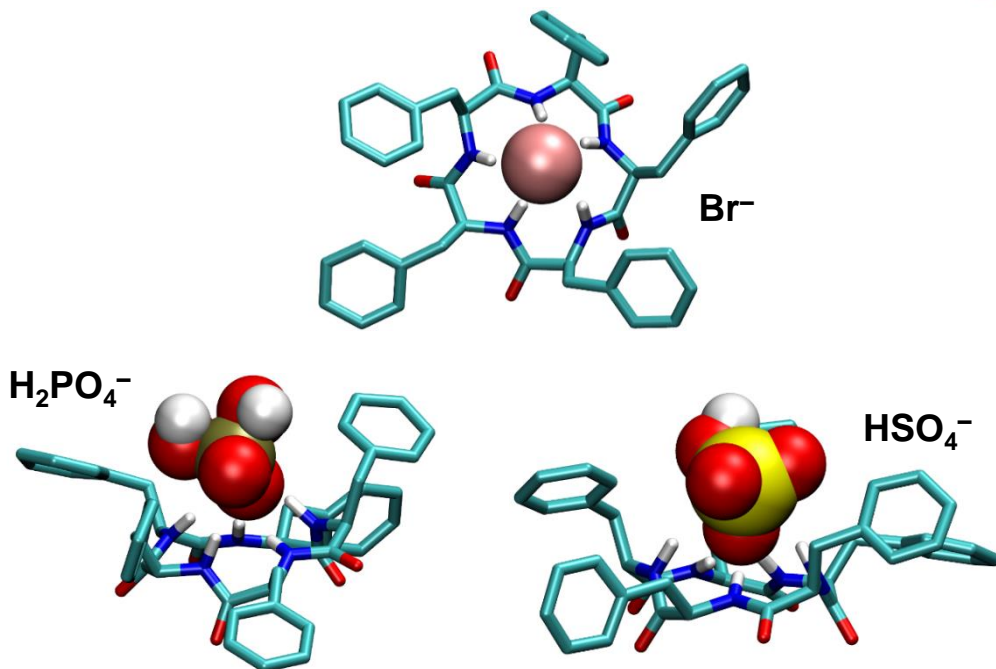
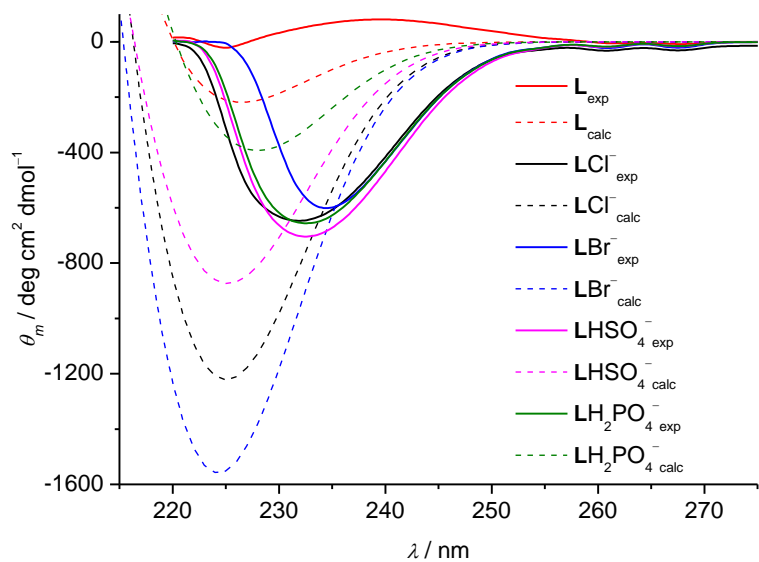
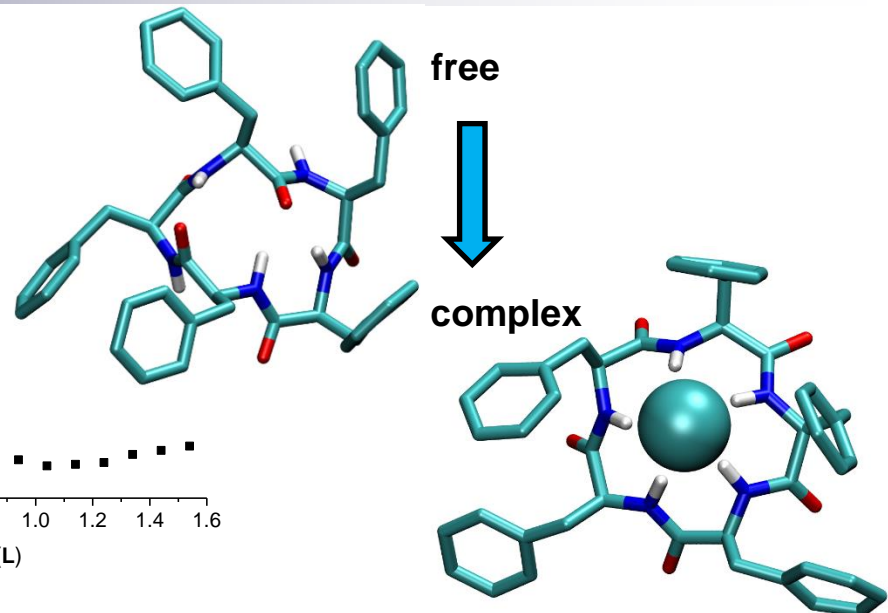
Anion-Sensing Properties of Cyclopentaphenylalanine



Circular dichroism titrations in MeCN



Phe₅ titration with Cl⁻ in MeCN



Experimental vs calculated CD spectra

Dichrocalc webserver + our MD data

Acknowledgements

Nikolina Vidović

Nikola Cindro

Giovanna Speranza

Matija Modrušan

Tamara Rinkovec

Siniša Tarana

Ivan Petters

Ivan Petranović

Tomislav Piteša

Vladimir Stilinović

Ivo Crnolatac

Ivo Piantanida

Vladislav Tomišić

MACRO SOL IP-2019-04-9560

SupraCAR IP-2014-09-7309