

HOW DO ENZYMES RECOGNIZE SUBSTRATES AND INHIBITORS: STRUCTURAL AND ELECTRON DENSITY ASPECTS

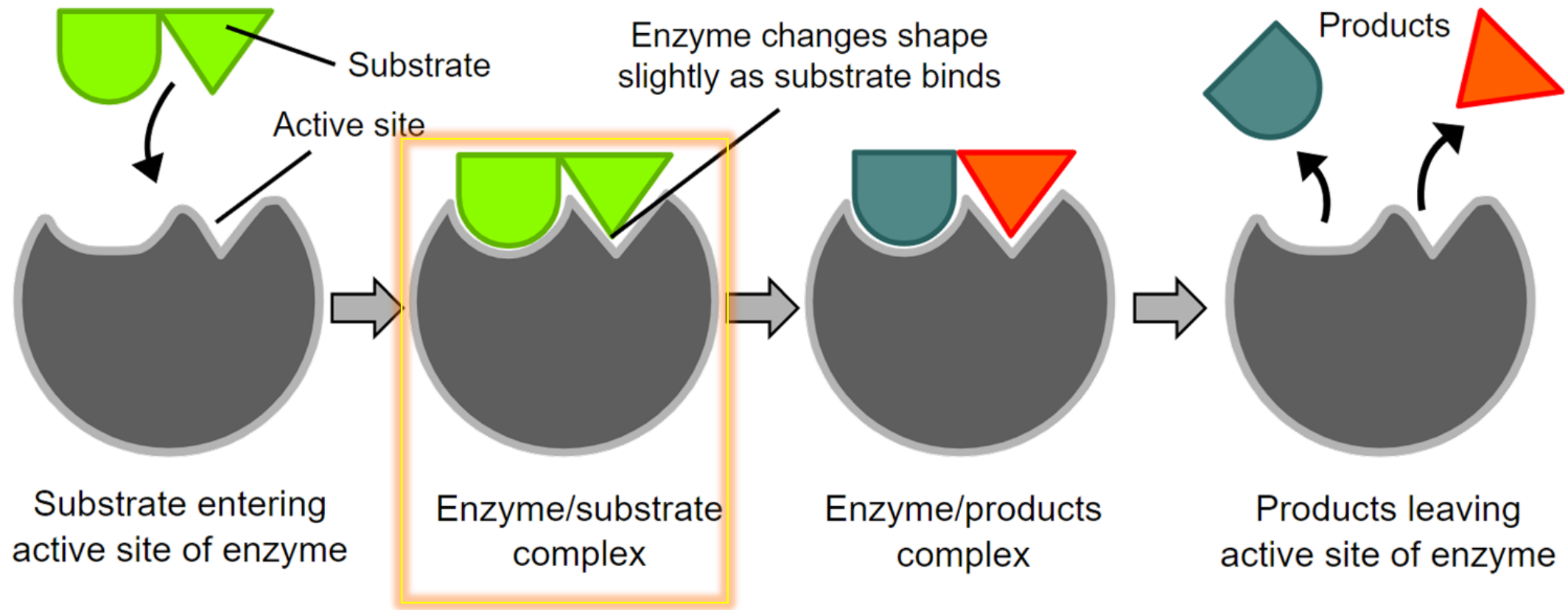
Maria G. Khrenova

¹ Lomonosov Moscow State University

² Federal Research Centre “Fundamentals of Biotechnology” of the Russian Academy of Sciences

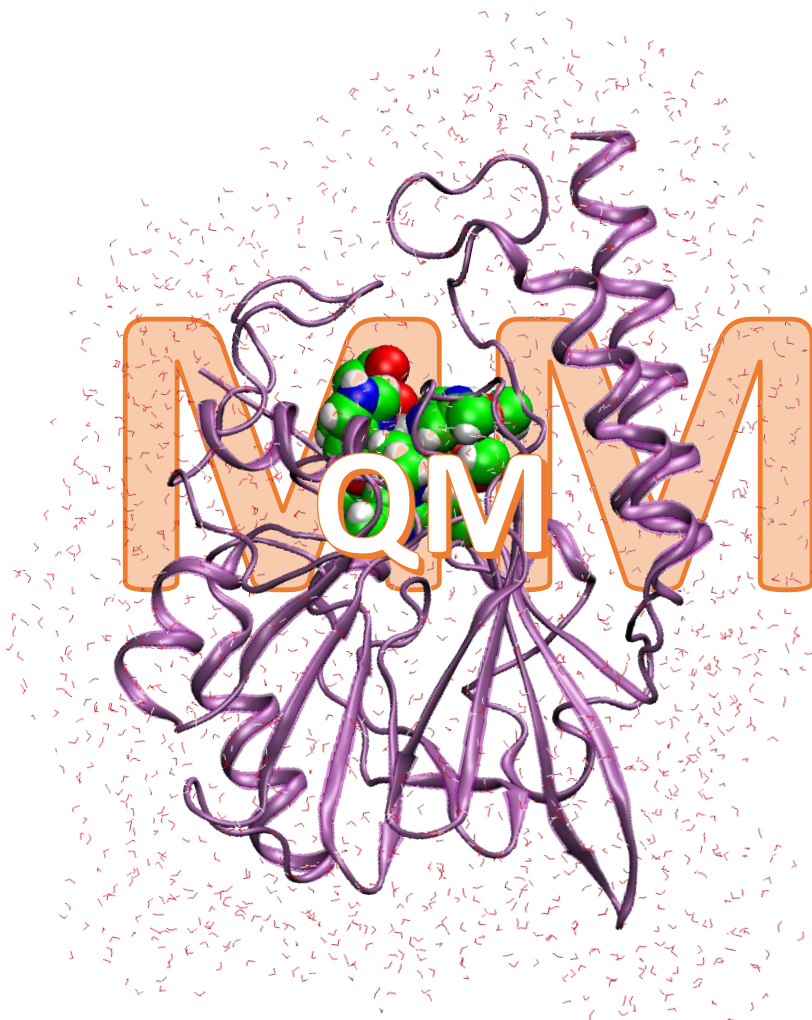
**XXVIII Symposium on Bioinformatics
and Computer-Aided Drug Discovery**

Enzyme-substrate interactions



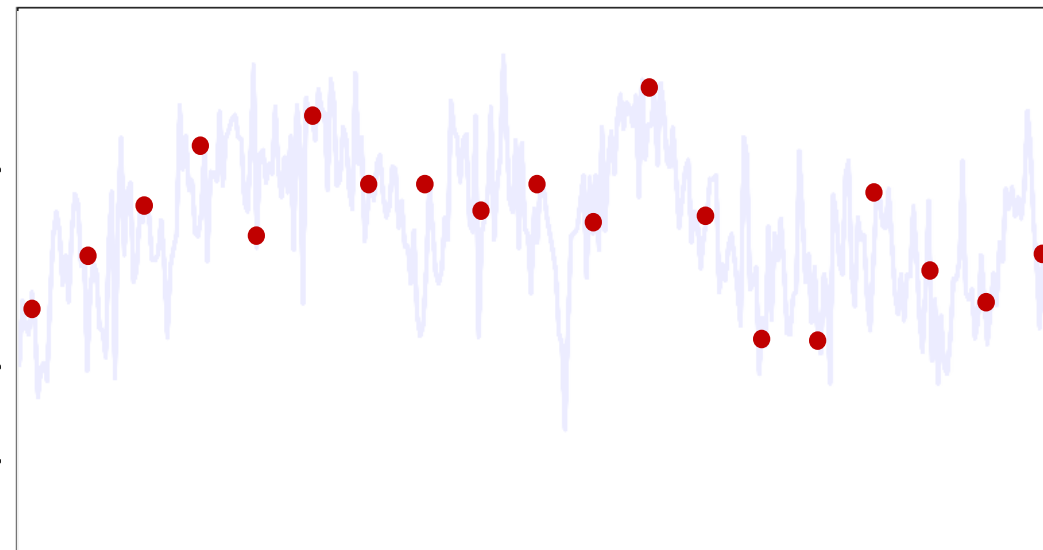
What is the substrate activation in the active site of the enzyme?

From local minima to ensembles of states

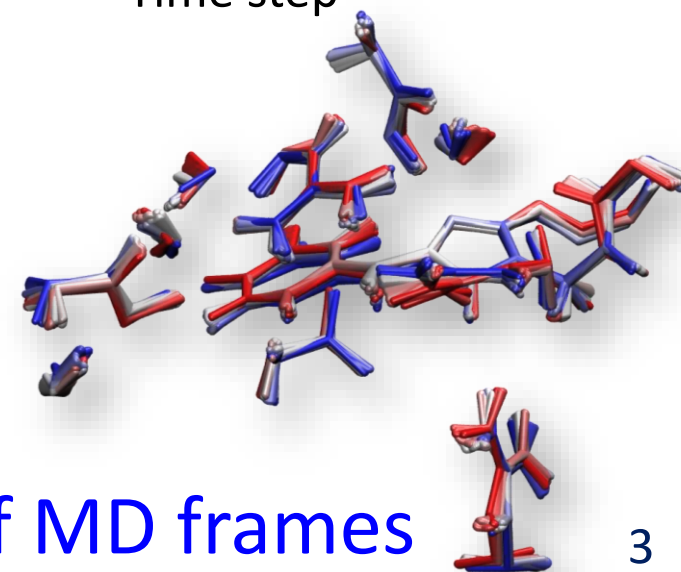


Molecular
dynamics

Property of the system



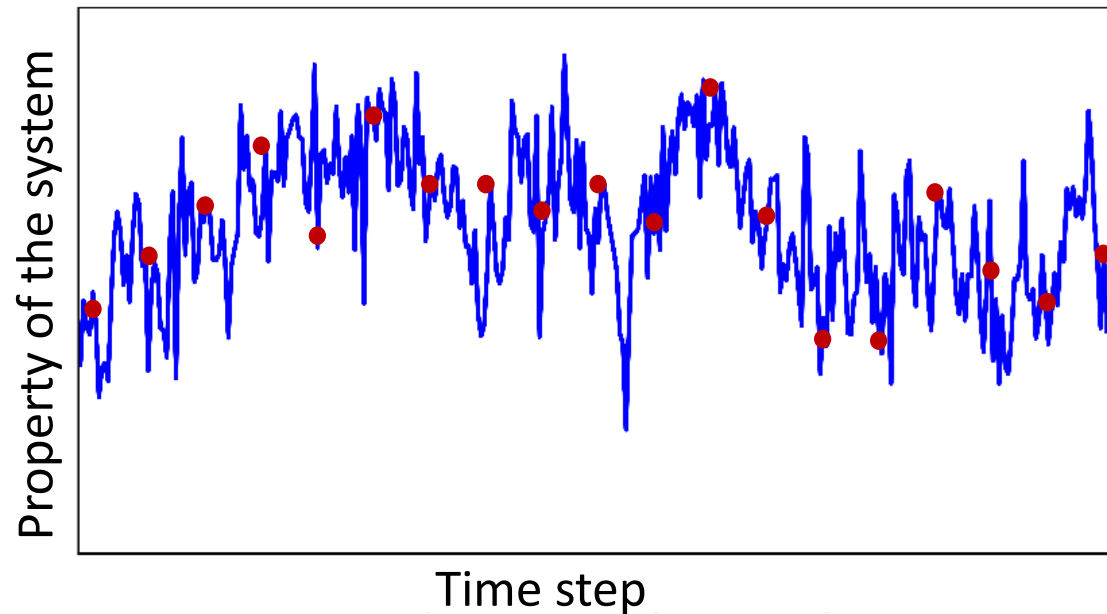
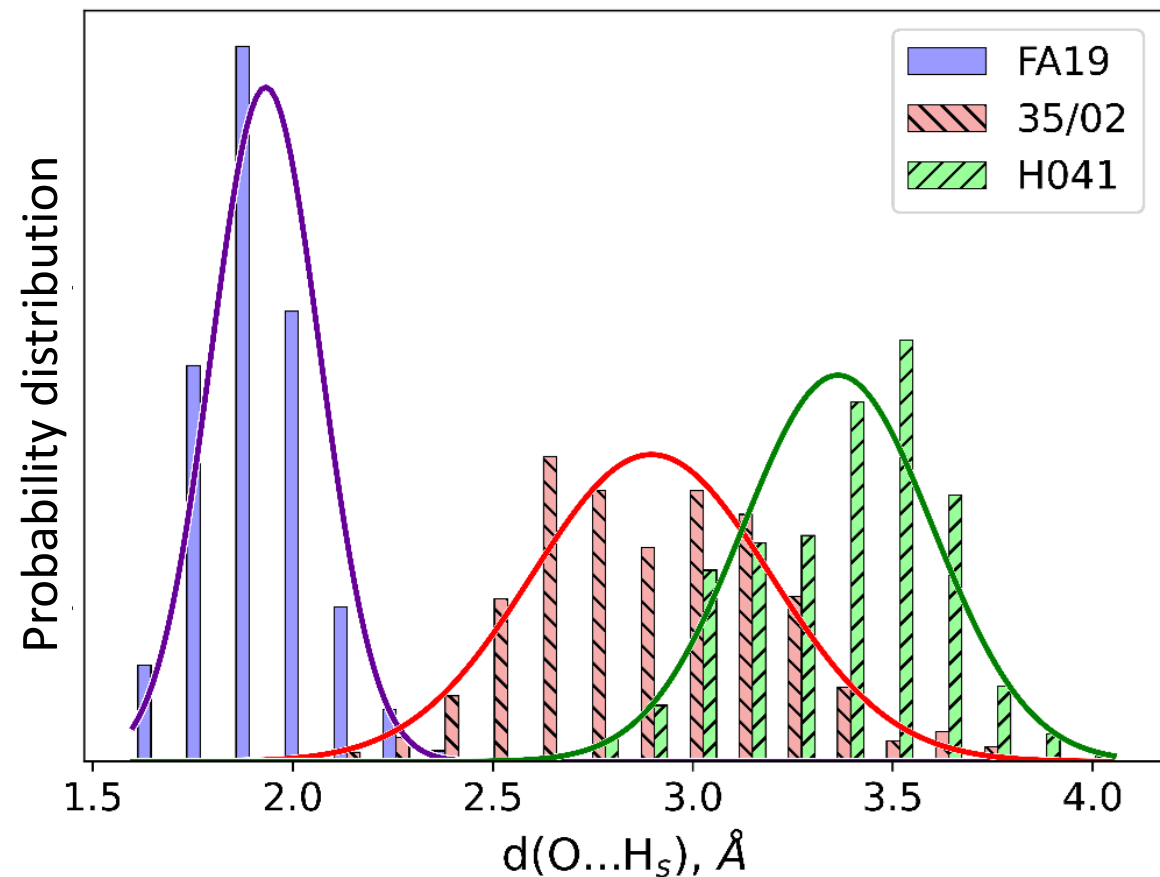
Time step



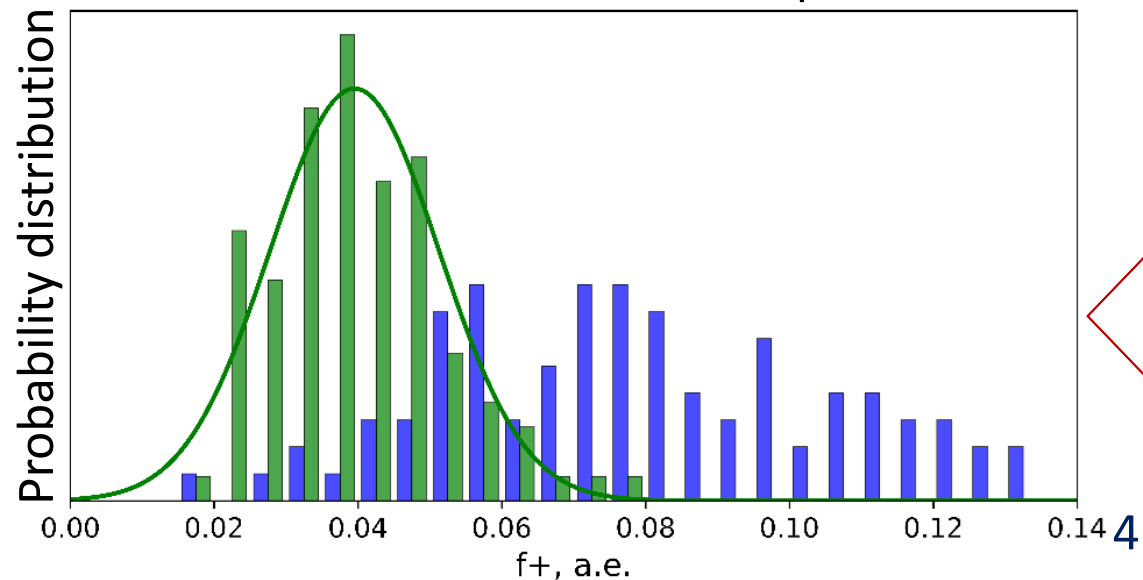
Set of MD frames

Distributions and average values

Geometry parameter



Electron density based descriptors



Breakout: GPU-based DFT code

Terachem:

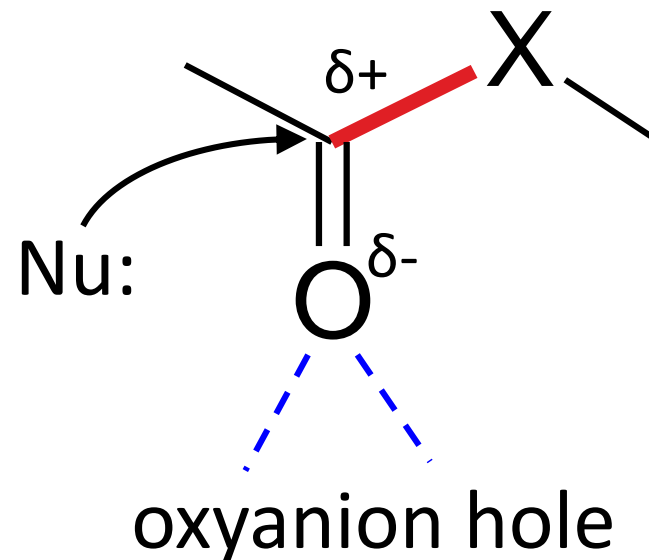
- QM subsystem: DFT(hybrid functional/6-31G**), ~100 atoms
- Benchmark (energy + gradient)
 - NVIDIA 1070 TI – 2 min
 - NVIDIA 3070 TI – 1 min.



Nucleophilic attack in enzymatic reactions

EC 3 Hydrolases:

- EC 3.1 Acting on ester bonds;
- EC 3.4 Acting on peptide bonds;
- EC 3.5 Acting on C-N bonds, other than peptide bonds;
- EC 3.7 Acting on C-C bonds.

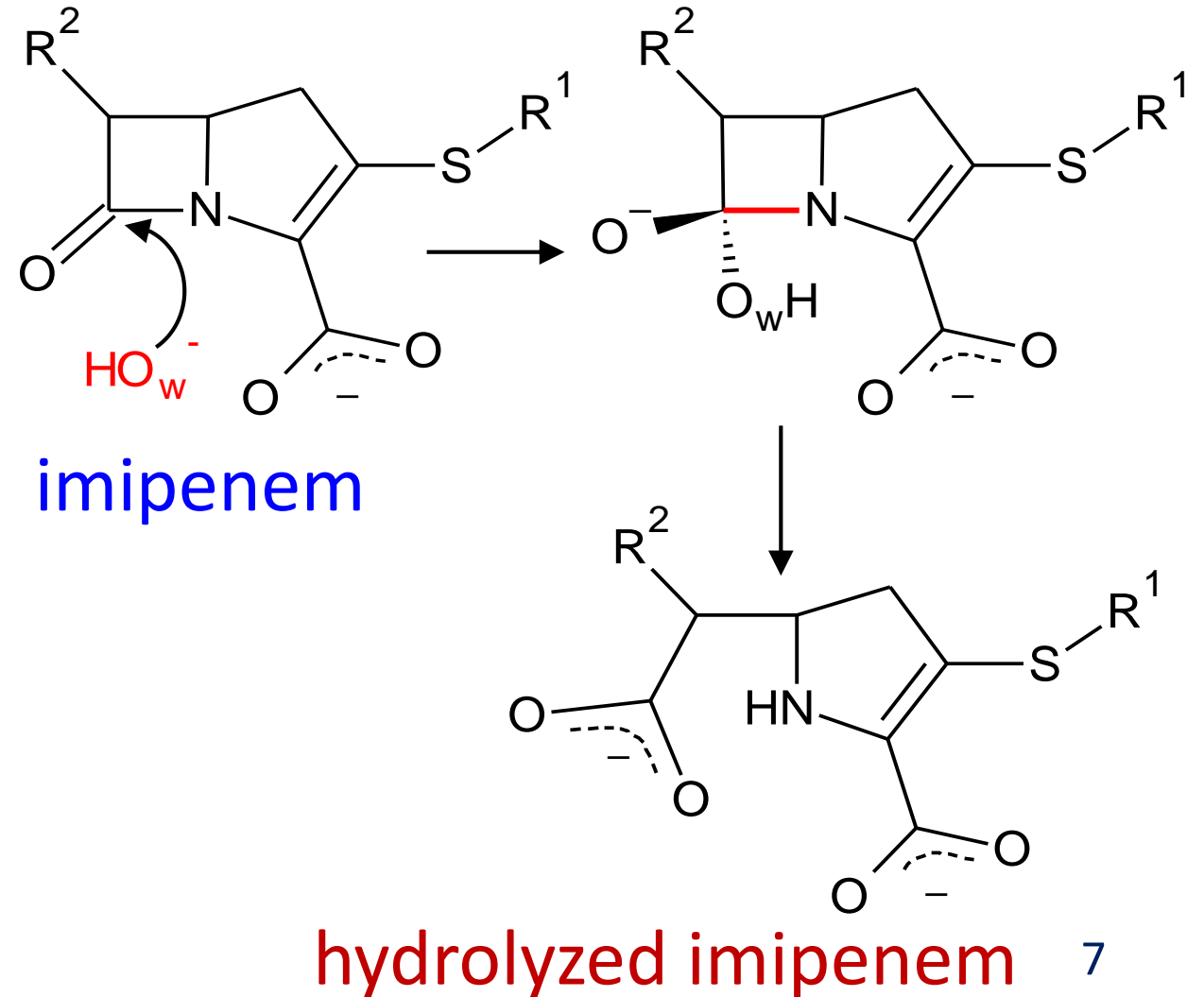
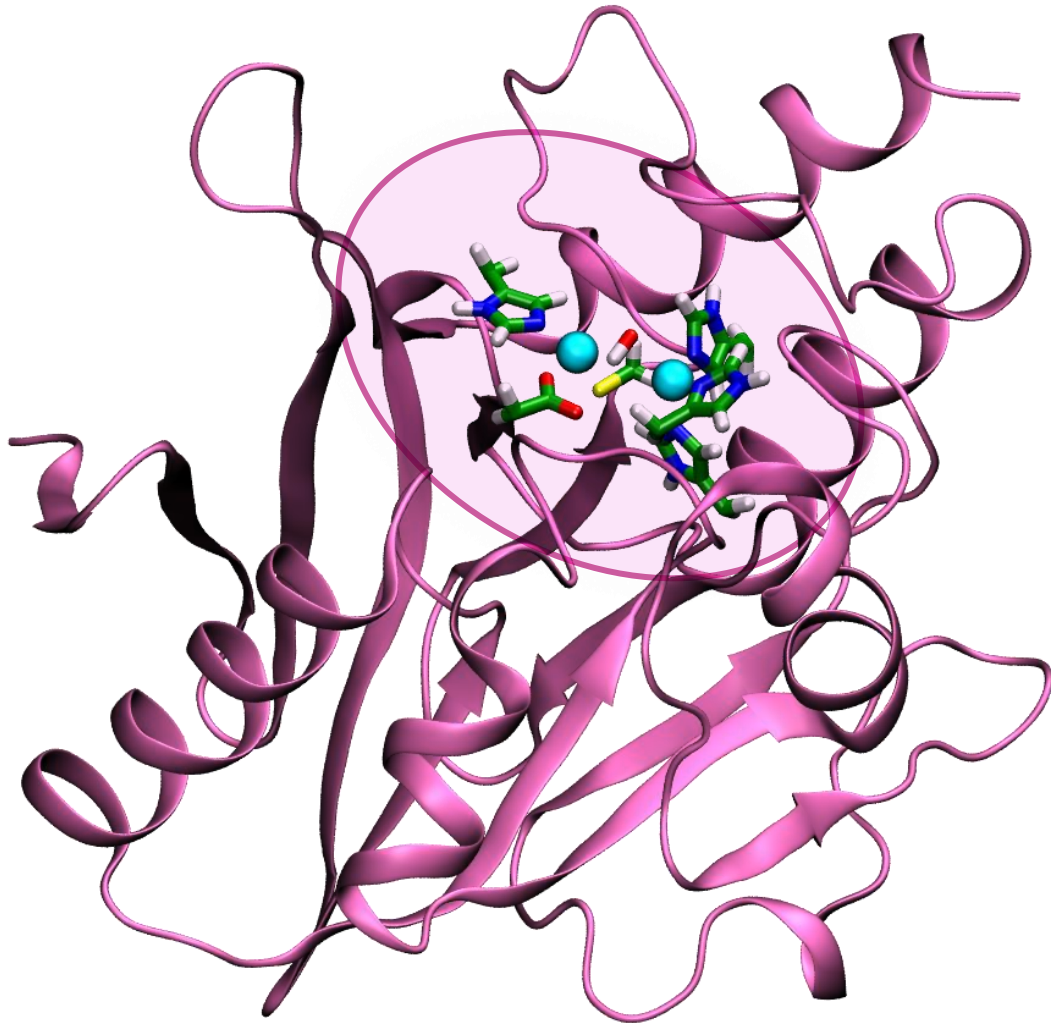


Nucleophile:

- H_2O
- ☆ OH^-
- ☆ OH of Ser
- OH of Thr
- ☆ SH of Cys

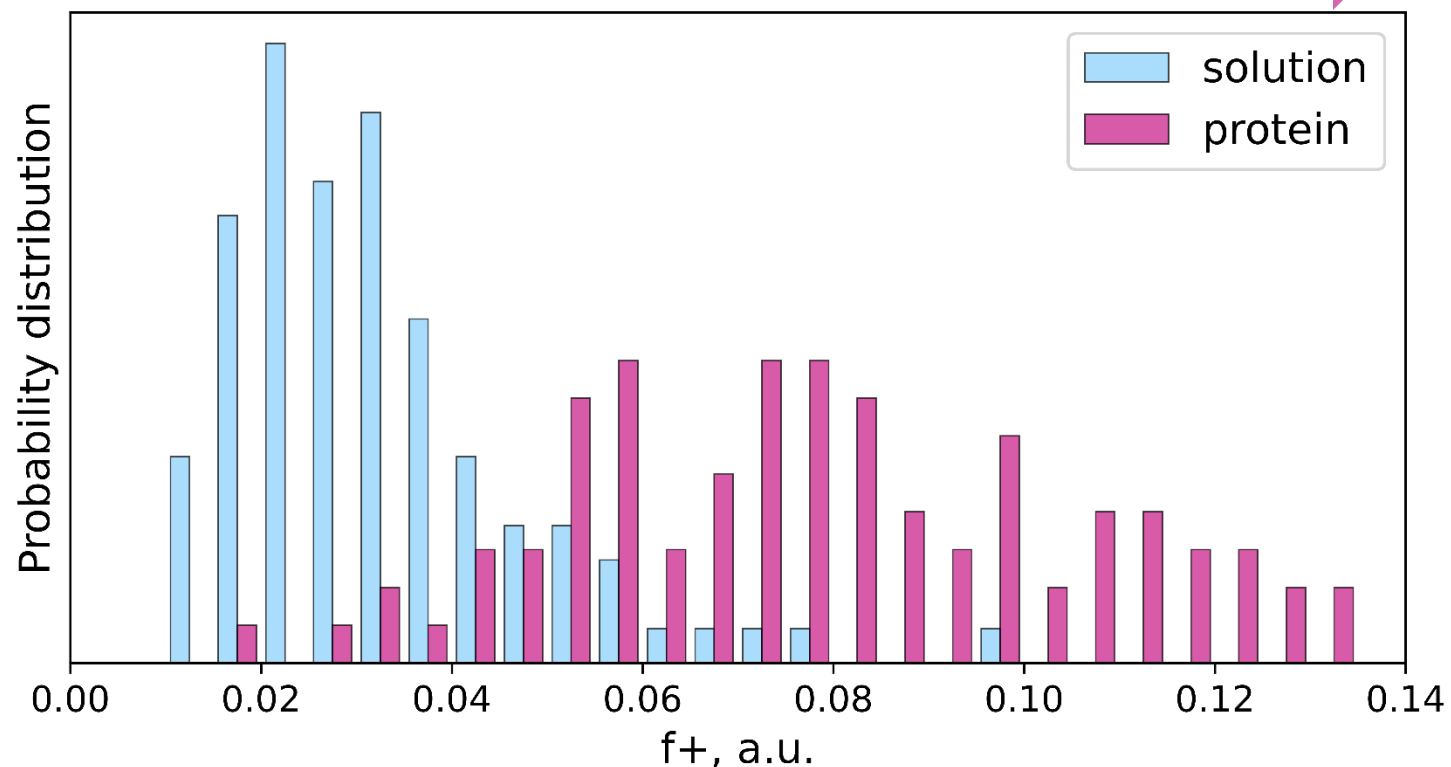
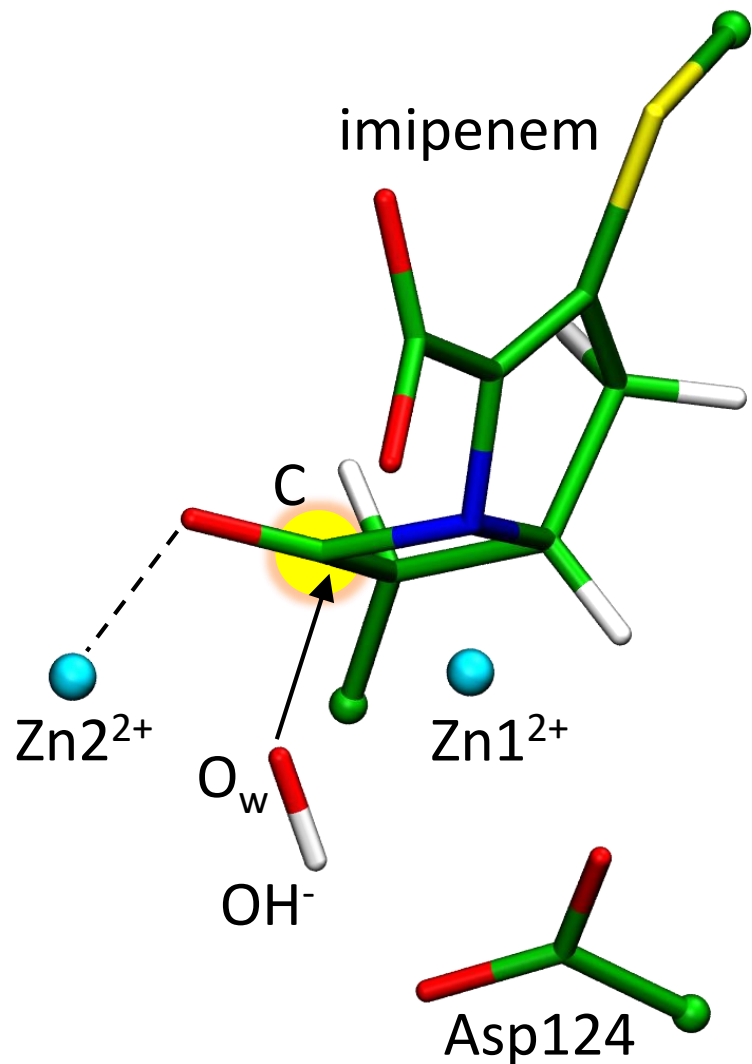
Case 1: NDM-1 metallo- β -lactamase

Hydrolysis of antibiotics related to the drug resistance



Case 1: NDM-1 metallo-beta lactamase

Protein activates a substrate



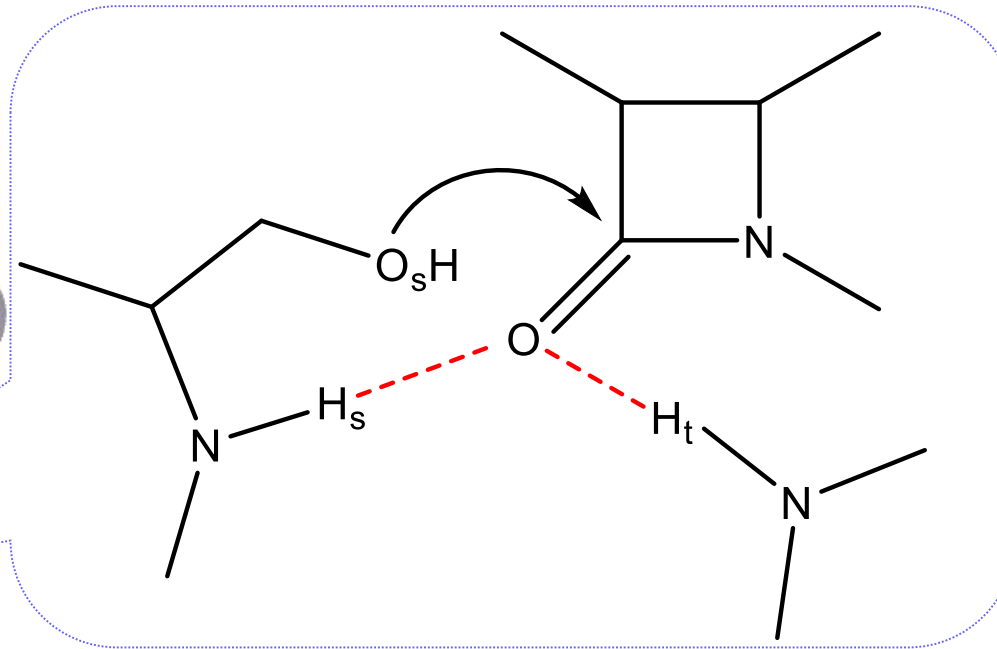
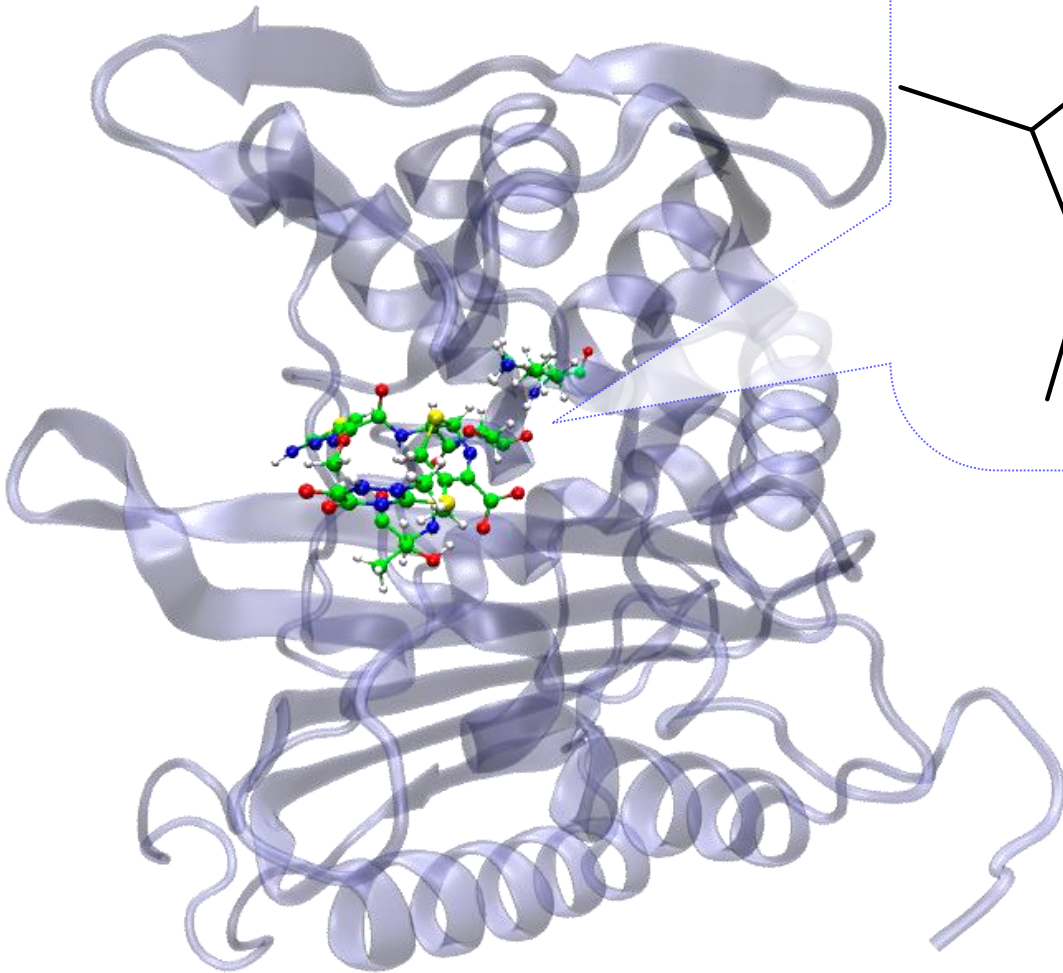
Fukui electrophilicity index of a carbonyl carbon atom

$$f+(C) = q_{N+1}(C) - q_N(C)$$

$q_N(C)$ – Hirshfeld charge of C atom

$q_{N+1}(C)$ – Hirshfeld charge of C atom in a system with an extra electron

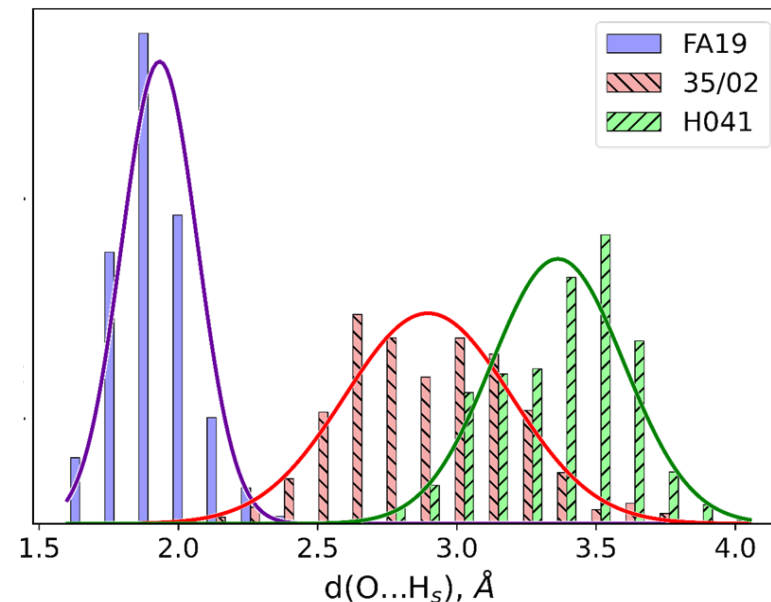
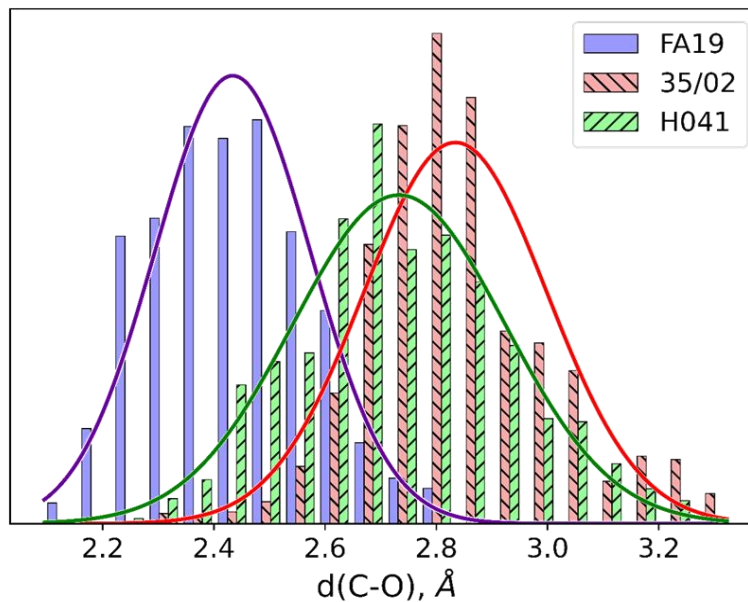
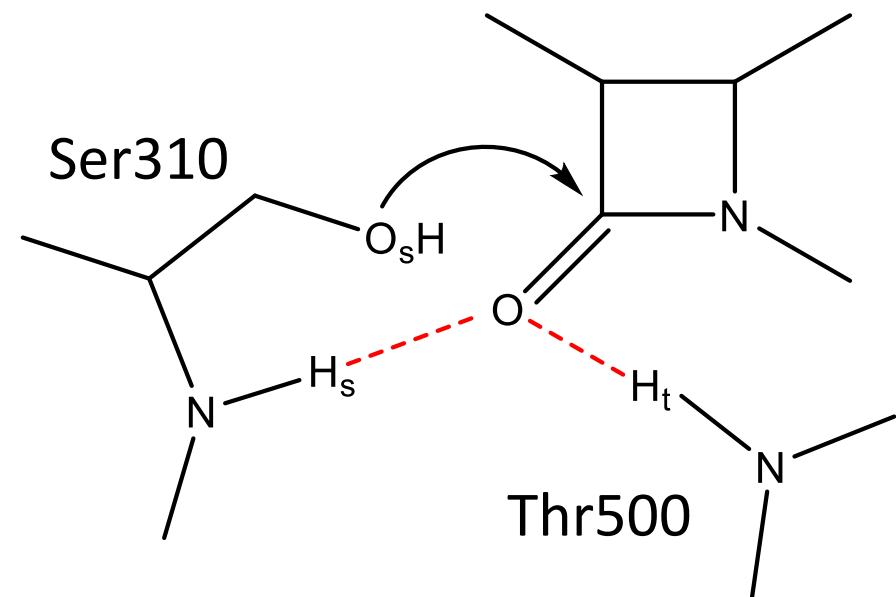
Case 2: Penicillin binding protein 2



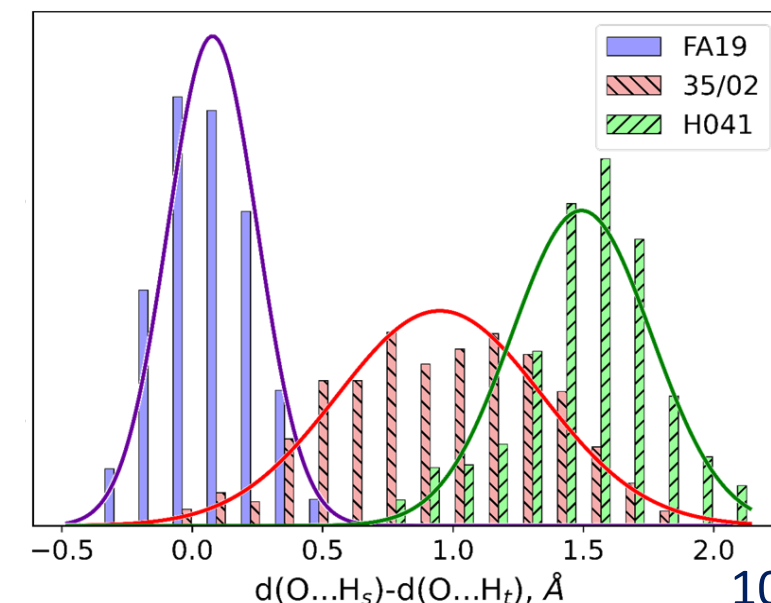
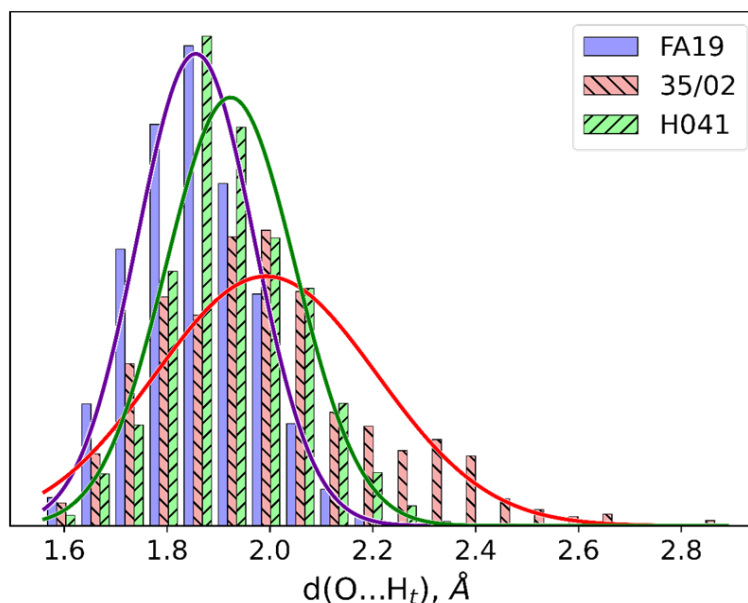
k_2/K_s for PBP2 from different *Neisseria gonorrhoeae* strains for ceftriaxone

strain	k_2/K_s , $\text{mM}^{-1}\text{s}^{-1}$
FA19	1710 ± 90
35/02	11.3 ± 0.4
H041	0.74 ± 0.03

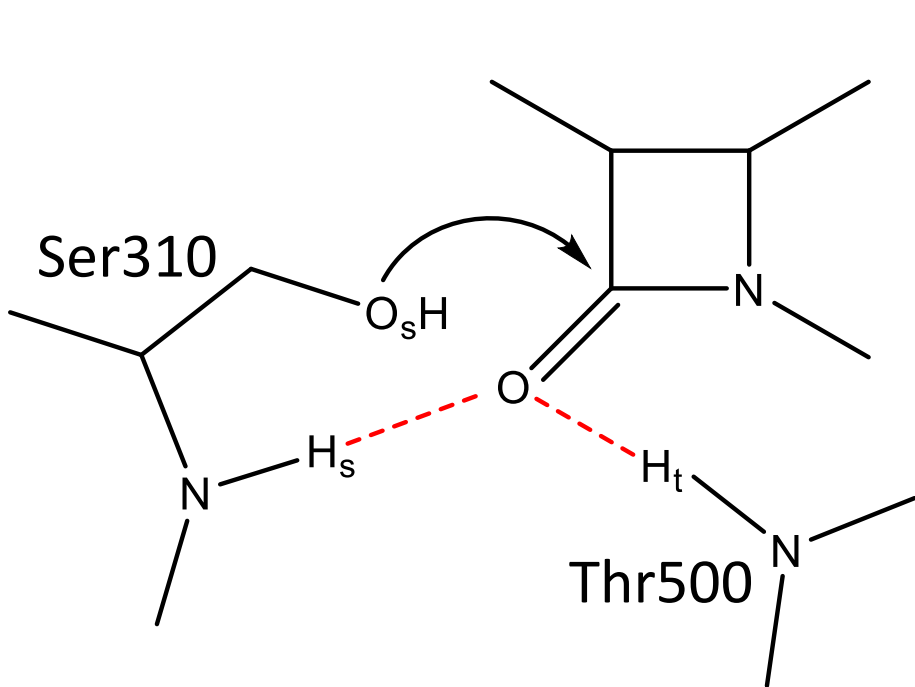
QM/MM MD simulations of the ES complexes



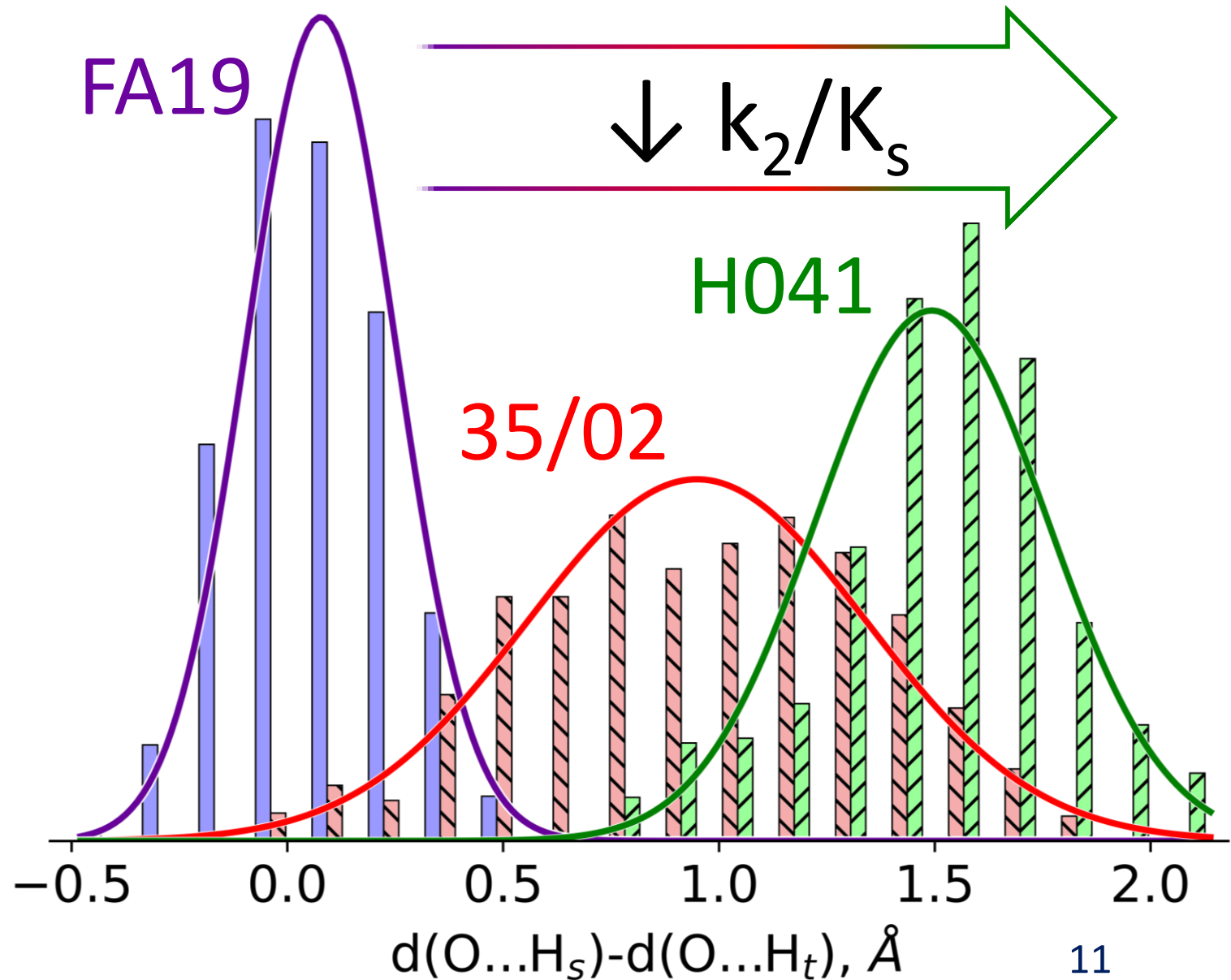
➤ Hydrogen bonds in the oxyanion hole are responsible for the substrate activation



QM/MM MD simulations of the ES complexes



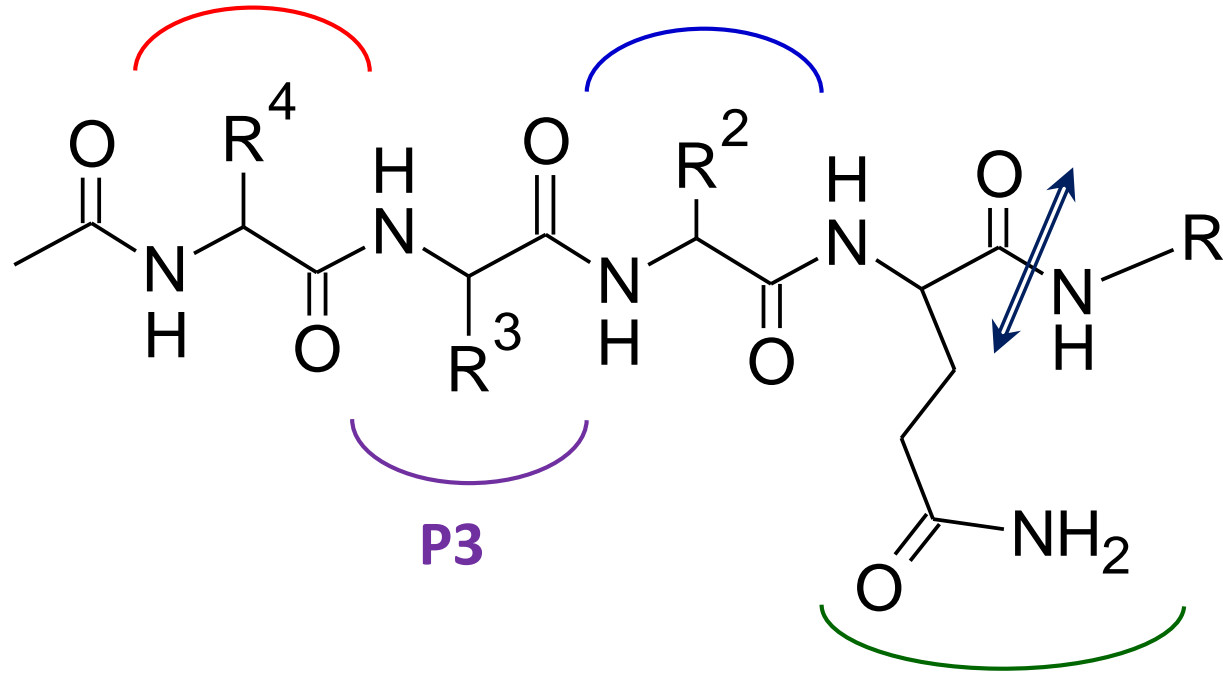
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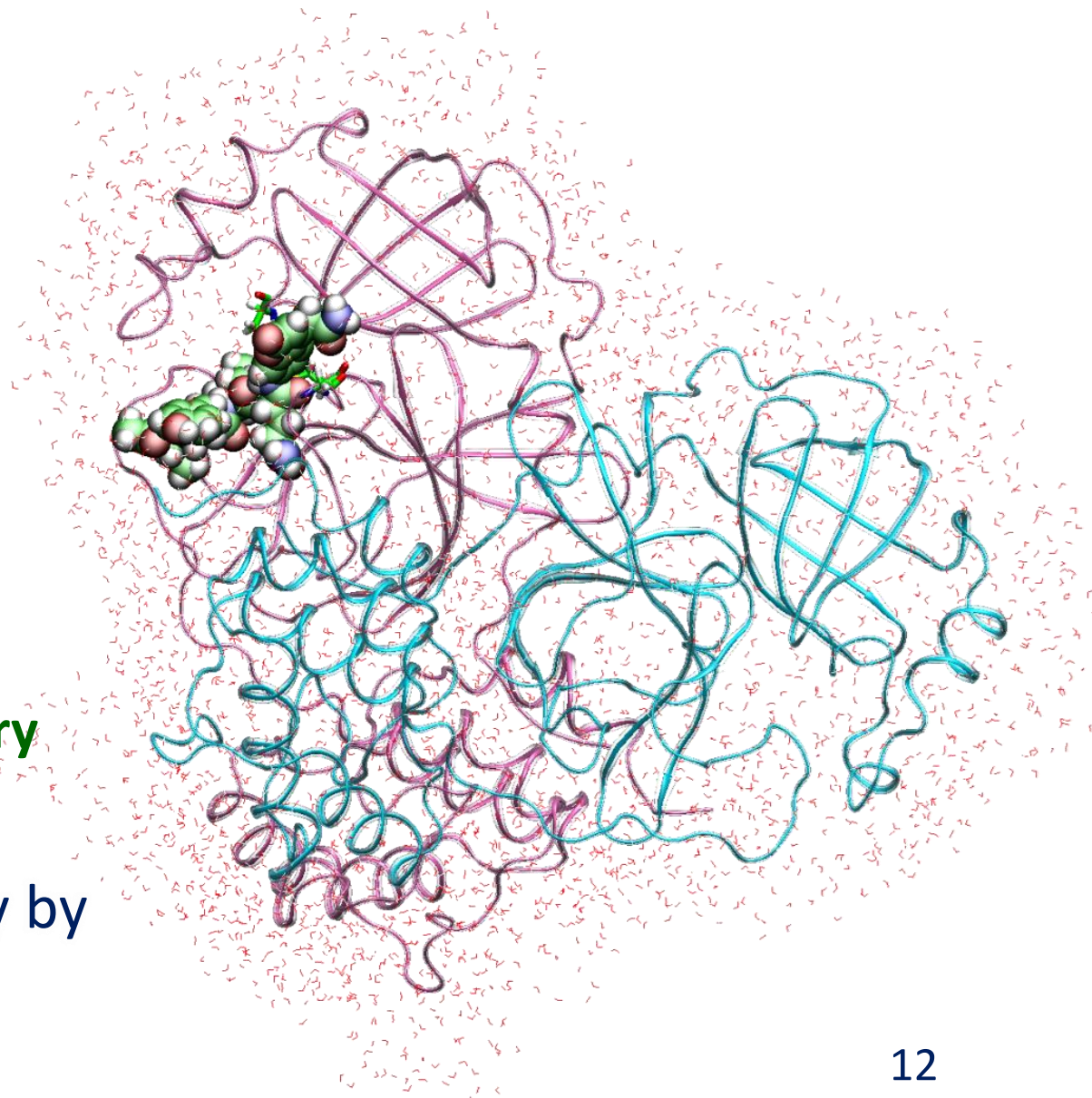
Case 3: Main protease M^{Pro} from SARS-CoV-2

P4

P2: preferred residue is Leu



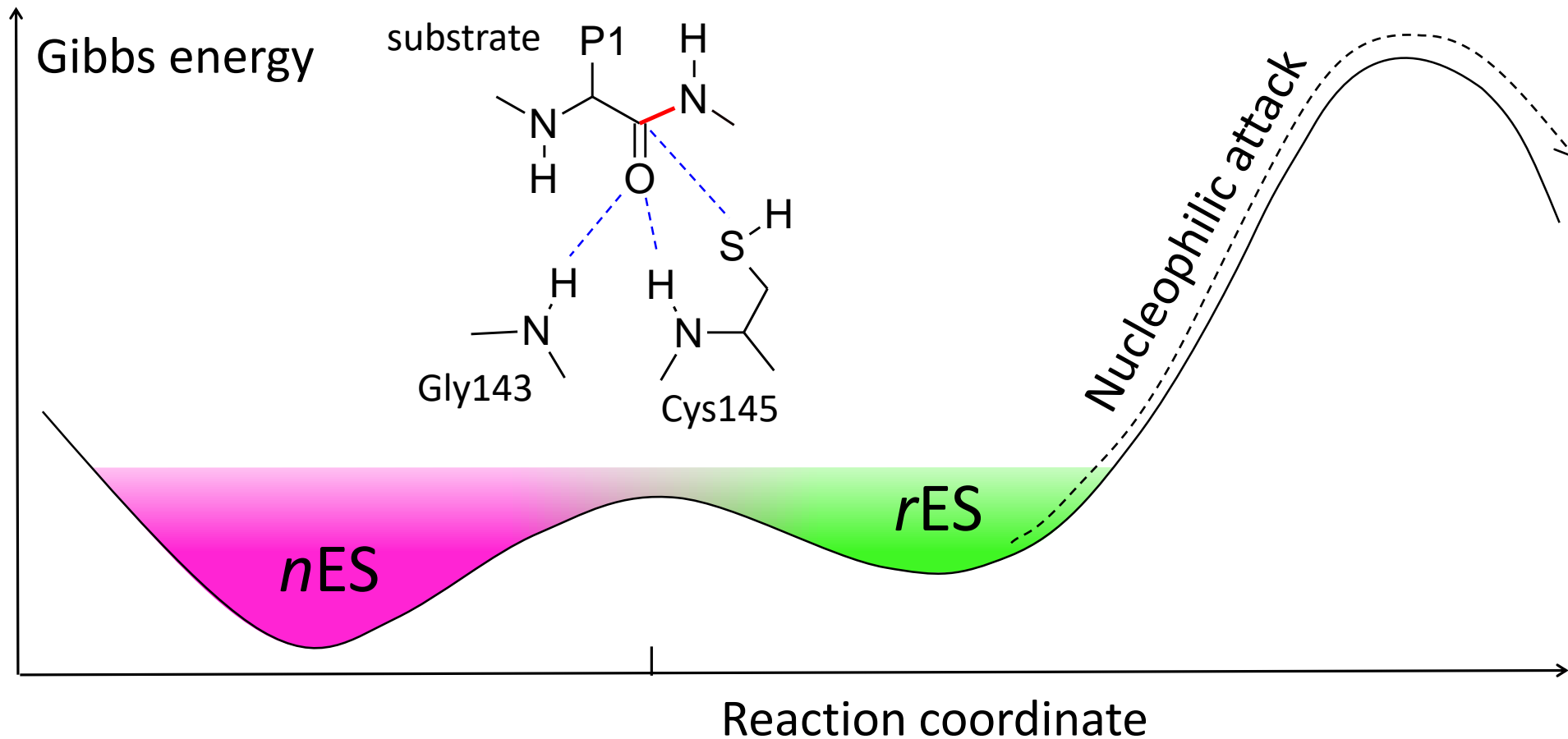
P1: Gln is obligatory



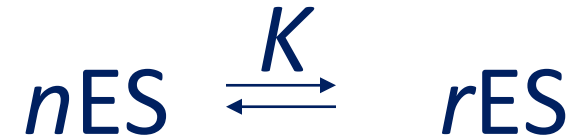
High substrate specificity:
Replacement of Leu at P2 decreases reactivity by
2 – 50 fold.

What is the origin of substrate specificity?

Efficiency of the substrate activation might be the reason



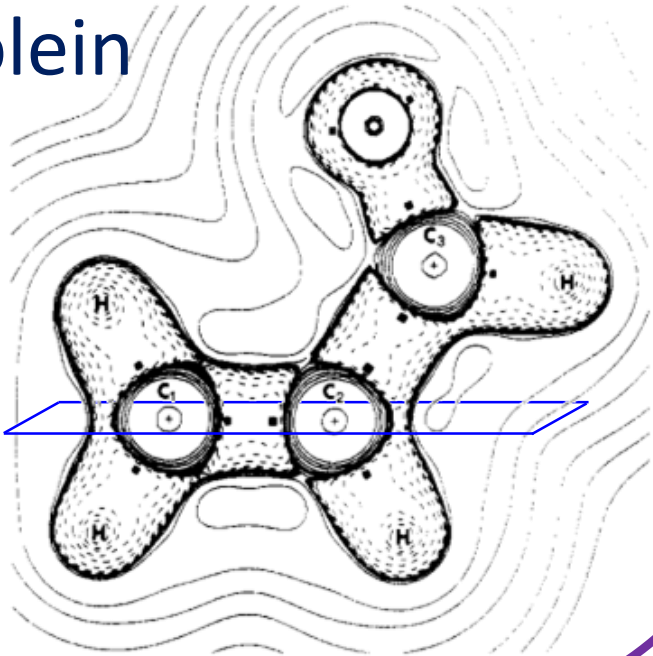
How to evaluate substrate activation?



- ✓ Criteria of assignment of conformations to either reactive or nonreactive

Laplacian of electron density

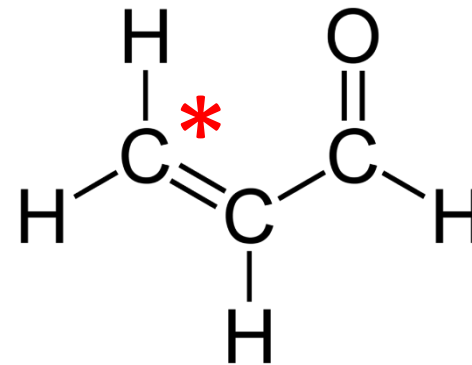
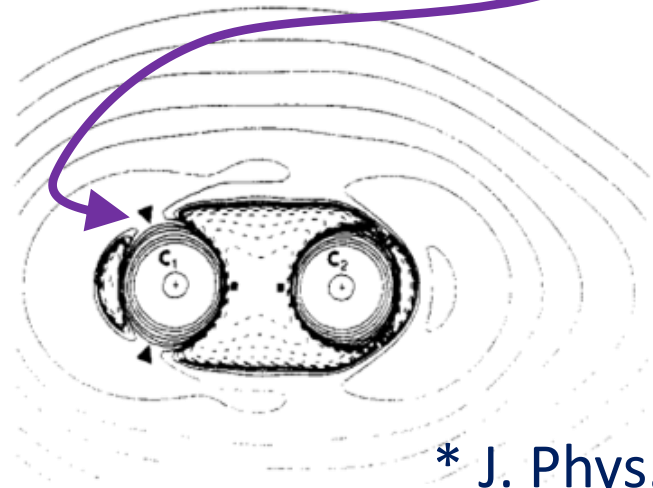
Acrolein



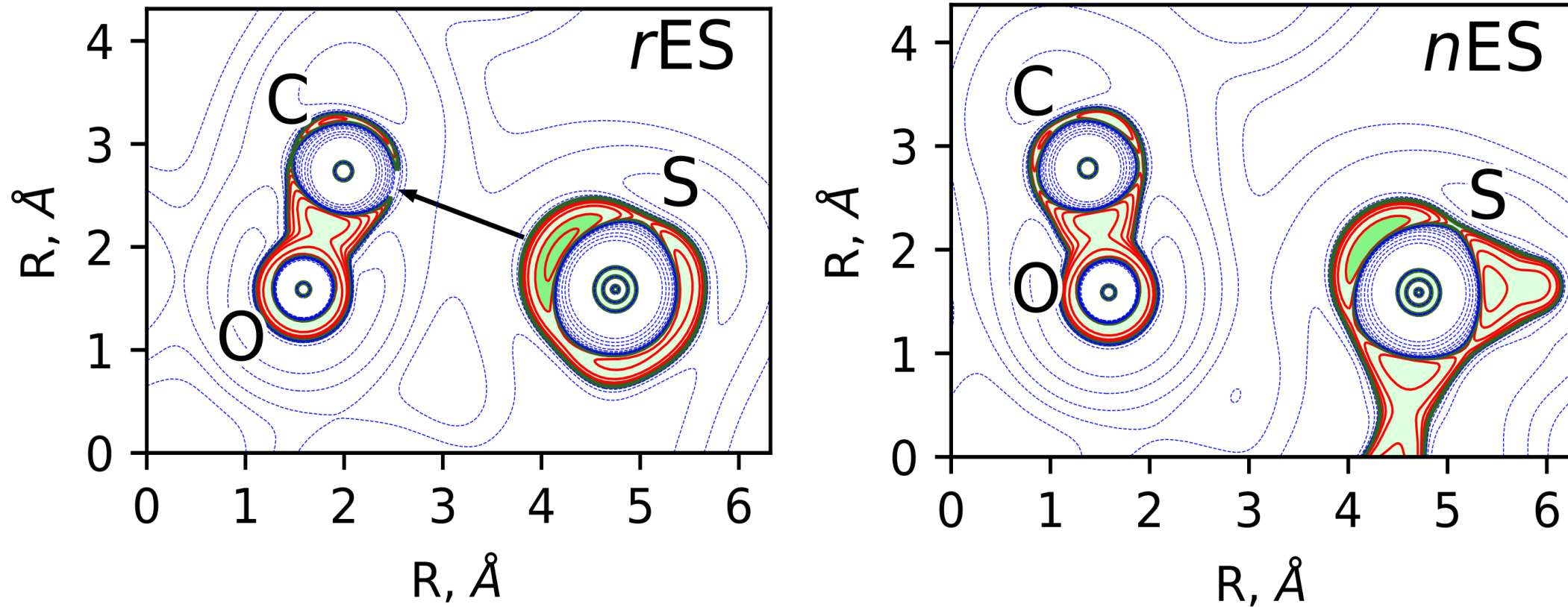
$$\nabla^2 \rho(\mathbf{r}) = \frac{\partial^2 \rho(\mathbf{r})}{\partial x^2} + \frac{\partial^2 \rho(\mathbf{r})}{\partial y^2} + \frac{\partial^2 \rho(\mathbf{r})}{\partial z^2}$$

$\nabla^2 \rho(\mathbf{r}) > 0$ – electron density depletion regions

$\nabla^2 \rho(\mathbf{r}) < 0$ – electron density concentration regions



Criterion to discriminate reactive and nonreactive species



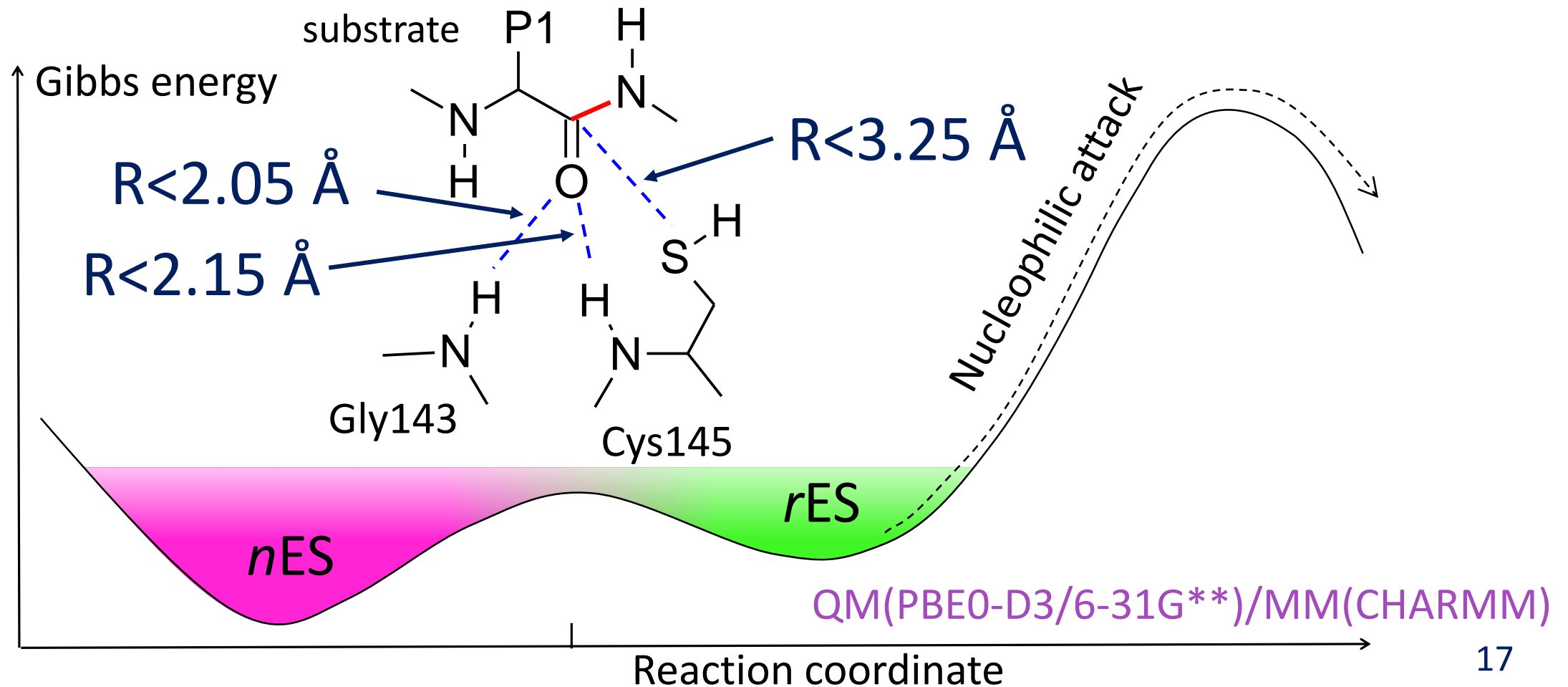
$\nabla^2\rho(\mathbf{r})$ maps in the S (Cys145) and C=O (substrate) plane

Blue isolines correspond to the ED depletion regions, $\nabla^2\rho(\mathbf{r}) > 0$

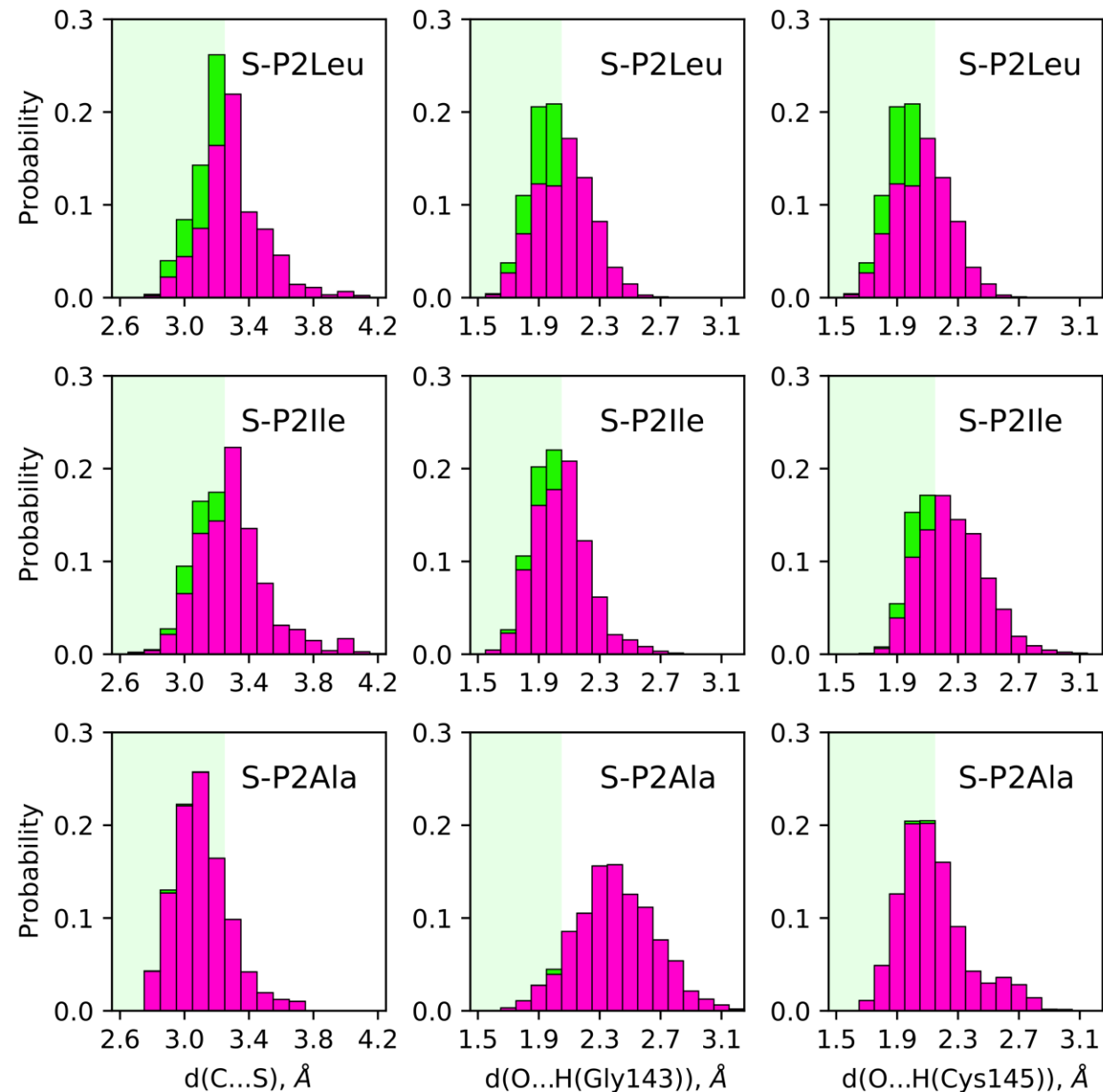
Red isolines correspond to the ED concentration regions, $\nabla^2\rho(\mathbf{r}) < 0$

Criteria to discriminate reactive and nonreactive species

All three geometry criteria should be satisfied together



Substrate specificity and rES \leftrightarrow nES equilibrium



Substrate	χ , %	k_{cat} (calc.)	k_{cat} (exp.)
S-P2Leu	22.4	1	1
S-P2Ile	10.2	0.46	0.45
S-P2Ala	0.6	0.03	<0.1

* values relative to S-P2Leu

$$k_{\text{cat}}(\text{AA}) = k_{\text{cat}}(\text{Leu}) \chi(\text{AA}) / \chi(\text{Leu})$$

Results obtained at the

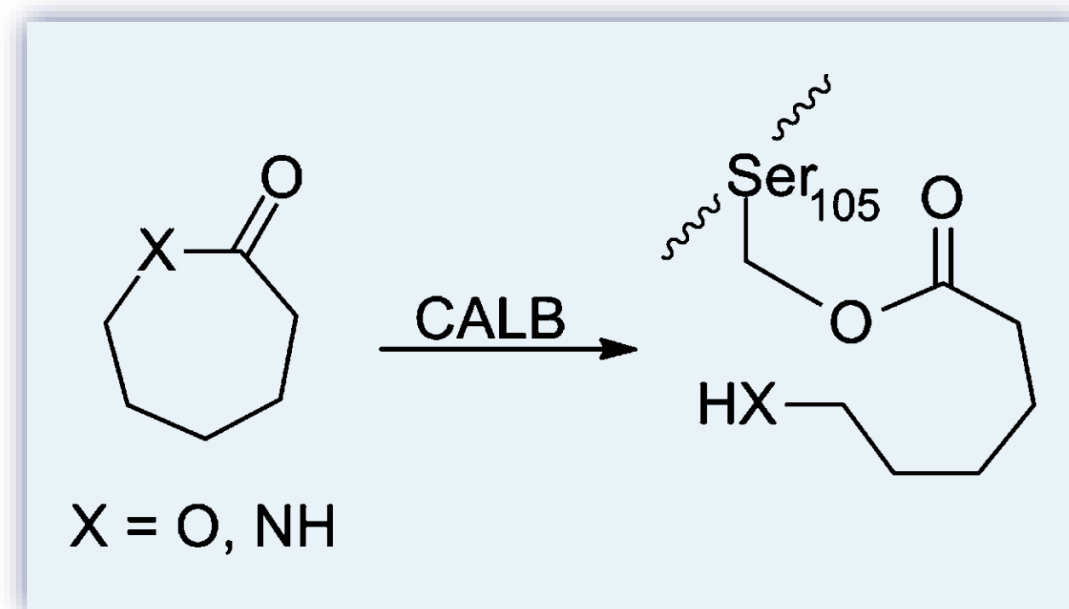
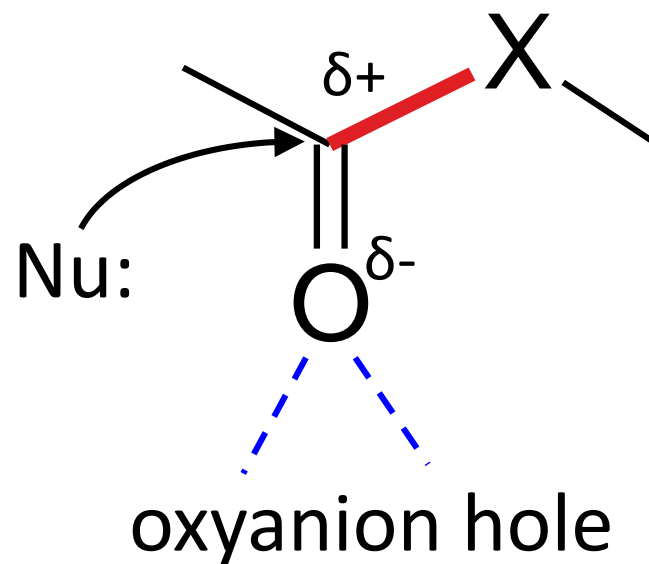
QM(PBE0-D3/6-31G**)/MM(CHARMM)

Example from the literature data

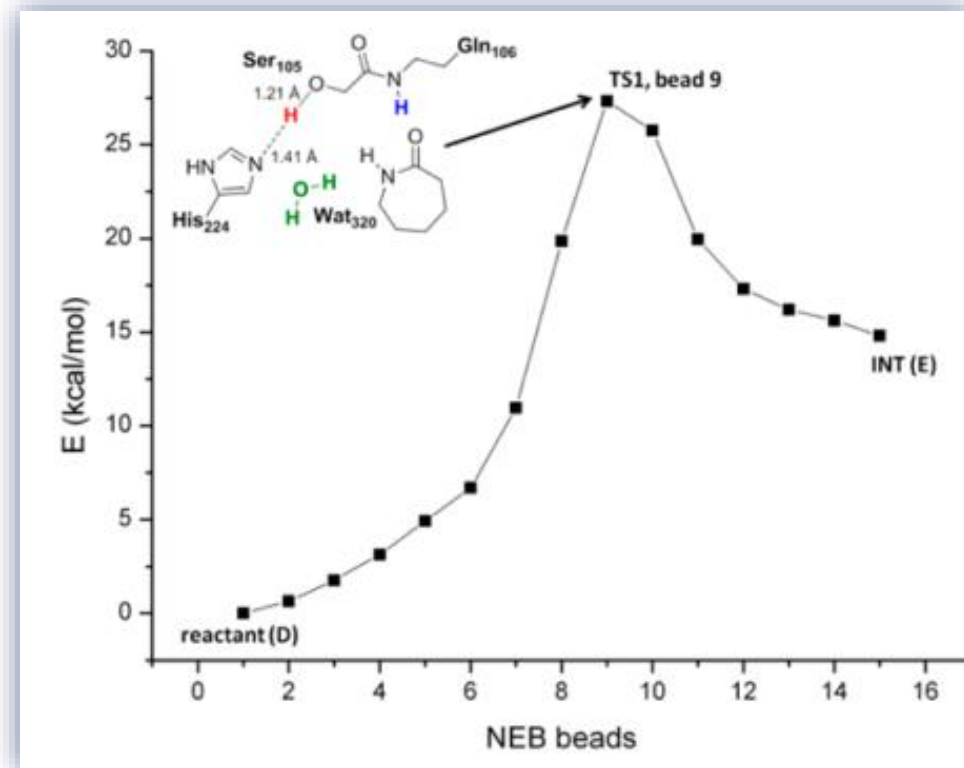
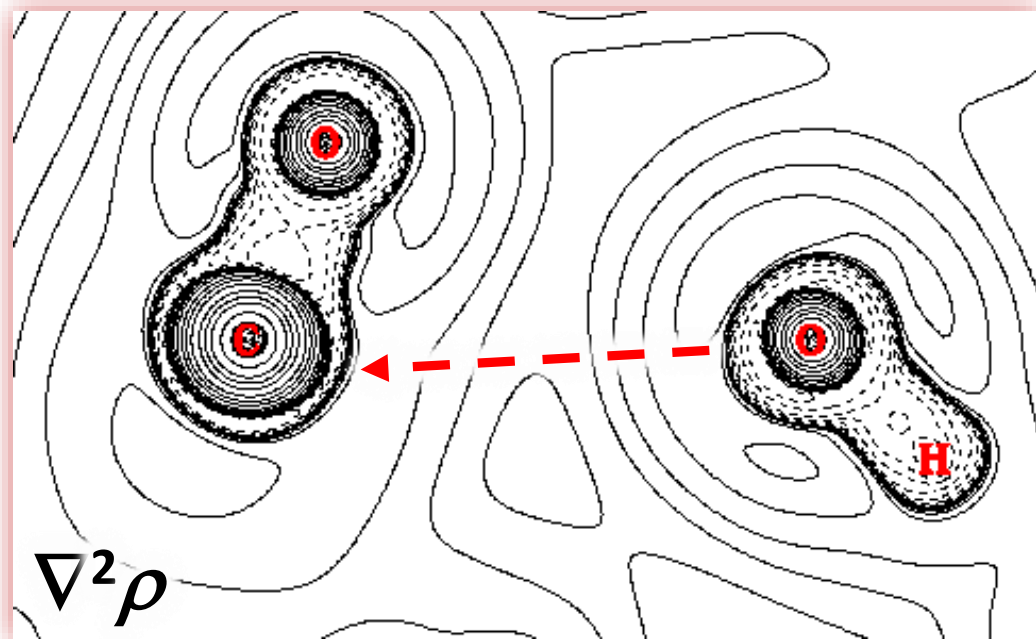
Comparative Theoretical Study of the Ring-Opening Polymerization of Caprolactam vs Caprolactone Using QM/MM Methods

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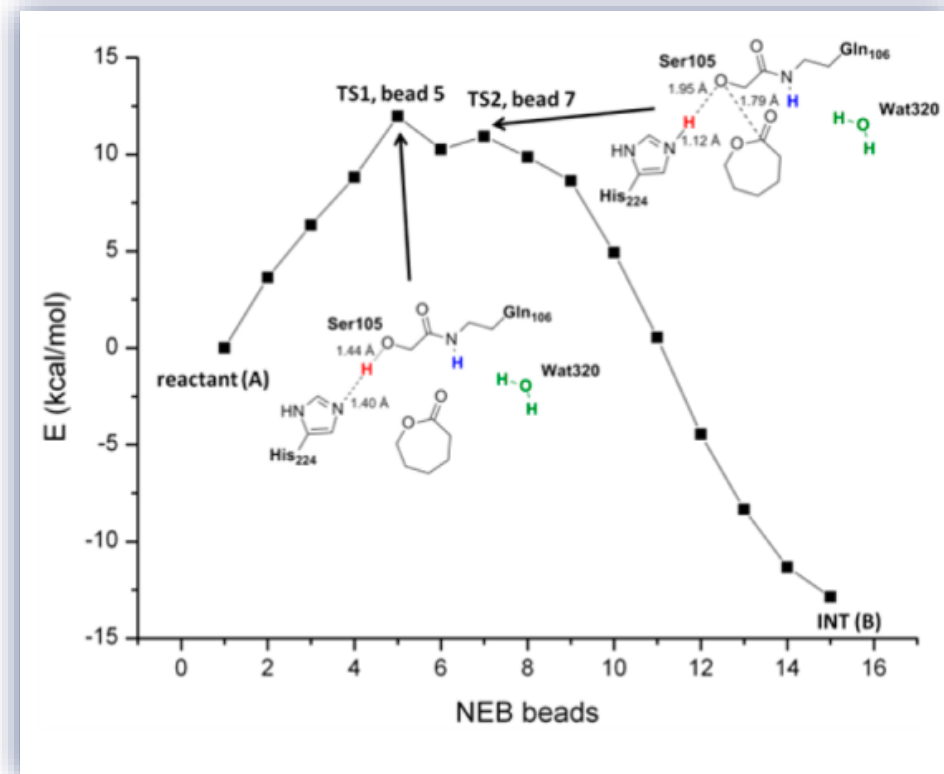
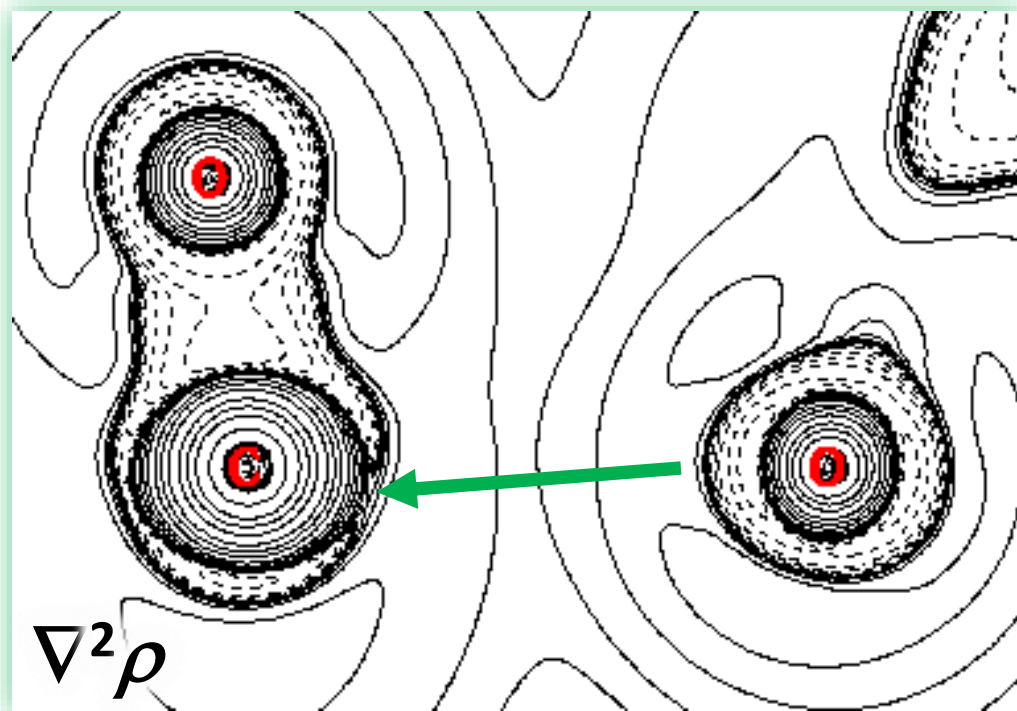


Caprolactam



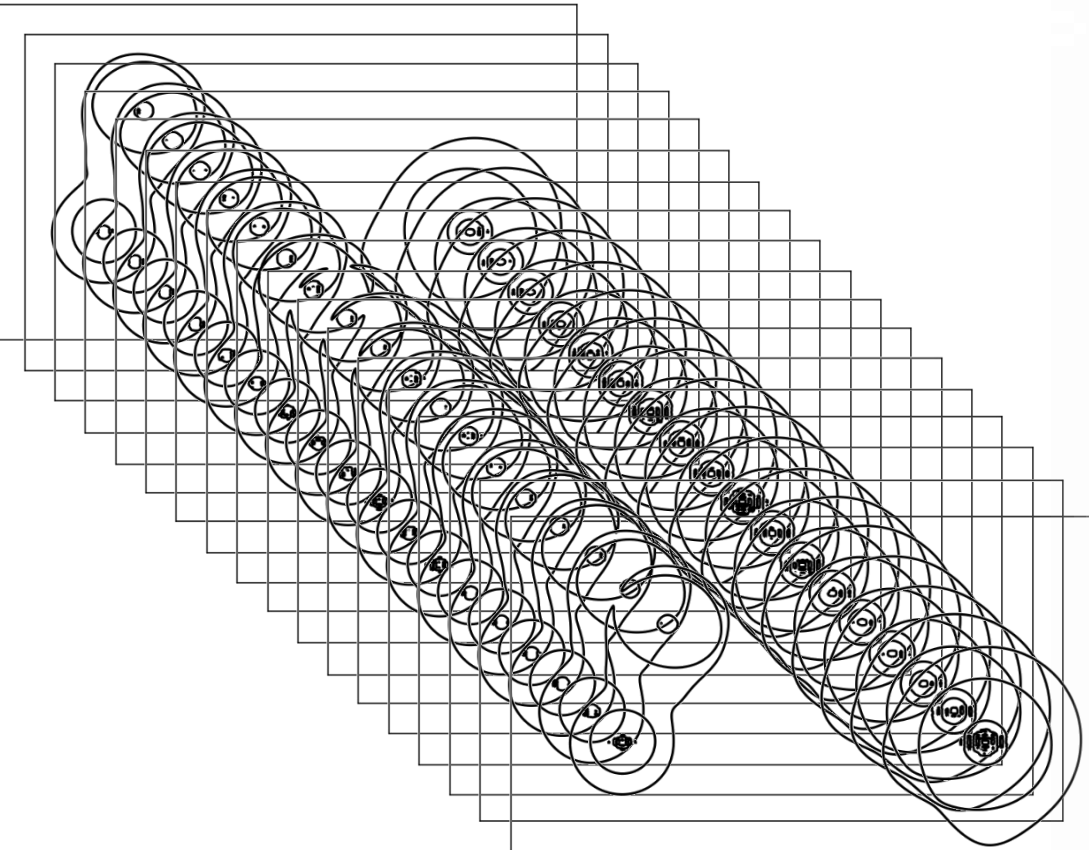
QM(PBE0/cc-pvdz)/MM

Caprolactone



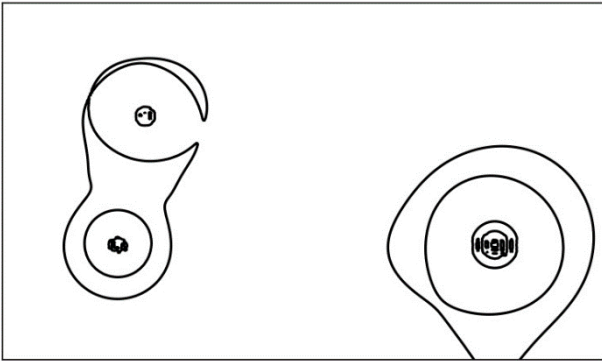
QM(PBE0/cc-pvdz)/MM

On-the-fly identification of the reactive and non-reactive species from MD trajectories

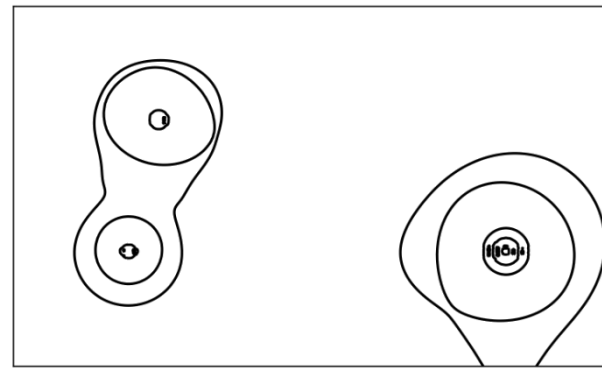


or
neural network

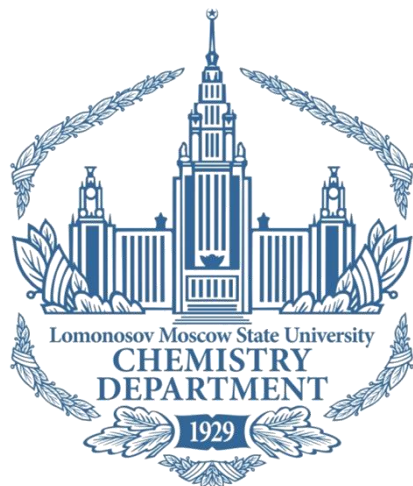
reactive



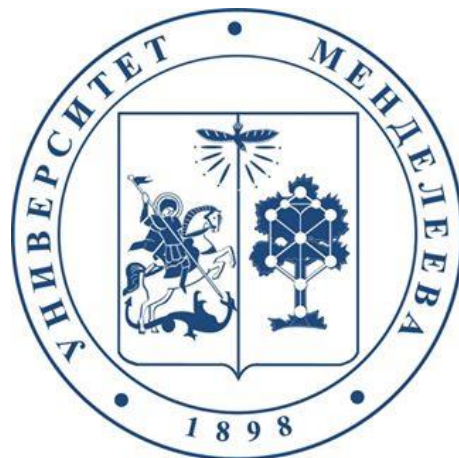
nonreactive



Scientific collaborations



Prof. A.V. Nemukhin
and members of
Laboratory of Quantum
Chemistry and Molecular
Modeling



Prof. V.G. Tsirelson
and members of
Quantum Chemistry
Department



FEDERAL RESEARCH CENTRE
«FUNDAMENTALS OF
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OF THE RUSSIAN ACADEMY
OF SCIENCES

Prof. M.G. Khrenova
and members of
Group of Molecular
Modeling

Financial support



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