# Enzymatic reaction – computational approach

Antonija Tomić & Borislav Kovačević

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# QM calculations



- Minima (ES, INT, P)  $\rightarrow$  scan method
- Maxima (TS) → Quadratic Synchronous Transit method (QST)
- Vibrational analysis
- Intrinsic reaction coordinate, IRC
- Levels of theory:
  - 1. opt.: B97D/[6-31G + LanL2DZ]
  - 2. opt: B97D/[6-31G(d) + LanL2DZ]
  - SP: B97D/[6-311++G(d,p) + LanL2DZ]



		Δ <i>E</i> / kJ mol⁻¹		
	system	model 1	model 2	
L.	ES	0.0	0.0	
	TS1	91.8	85.7	
	INT	45.2	47.4	
	TS2	79.7	37.3	
	Р	-21.5	-28.3	
	mode	el 1 model 2 01,8 ks mol	-1 TS2	

reaction coordinate

model 1: B97D/[6-31G + LanL2DZ] + *ZPVE*<sub>B97D/[6-31G + LanL2DZ]</sub> model 2: B97D/[6-311++G(d,p) + LanL2DZ]//B97D/[6-31G + LanL2DZ] + *ZPVE*<sub>B97D/[6-31G + LanL2DZ]</sub>





















# **QMMM** calculations





- SYSTEM = enzyme + Leu-enkephalin + water
- 105/6 QM atoms
- ONIOM calculations:

 $E^{ONIOM} = E_{MM}(S) + E_{QM}(SM) - E_{MM}(SM)$ 



REACTION MECHANISM with two ES structures: MD MD + X-ray



MD – X-ray ES structure

























#### MD – X-ray ES str. vs. MD ES str.



O<sub>s</sub>-His568



Zn-Glu508(Oe1)

# THANK YOU!



			Δ <i>E</i> / kJ mol⁻¹	
		system	model 1	model 2
100 -	model 1model 2	ES	0.0	0.0
80 -		TS1	60.8	76.5
80 -		INT1	59.9	77.1
60 -		TS2	66.5	83.7
lom		INT2	46.8	61.0
3 40 -		TS3	40.9	55.4
20 -		INT3	35.9	51.1
√ 20		TS4	30.0	47.1
0 -		INT4	8.3	23.5
	ES TS1 INT1 TS2 INT2 TS3 INT3 TS4 INT4 TS5 INT5 TS6 P	TS5	25.2	41.1
-20 -	reaction coordinate	INT5	2.9	14.3
		TS6	10.0	23.1
		Ρ	-4.6	13.1
		Ρ'	-33.6	-30.2

model 1: B97D/[6-31G (d)+ LanL2DZ] +  $ZPVE_{B97D/[6-31G(d) + LanL2DZ]}$ model 2: B97D/[6-311++G(d,p) + LanL2DZ]//B97D/[6-31G(d) + LanL2DZ] +  $ZPVE_{B97D/[6-31G(d) + LanL2DZ]}$ 

























