

Inhibitory Zinc Ion Binding Site and The Metal Exchange Mechanism in Human DPP III

Sanja Tomić, Antonija Tomić, Hrvoje Brkić and Antonia Matić



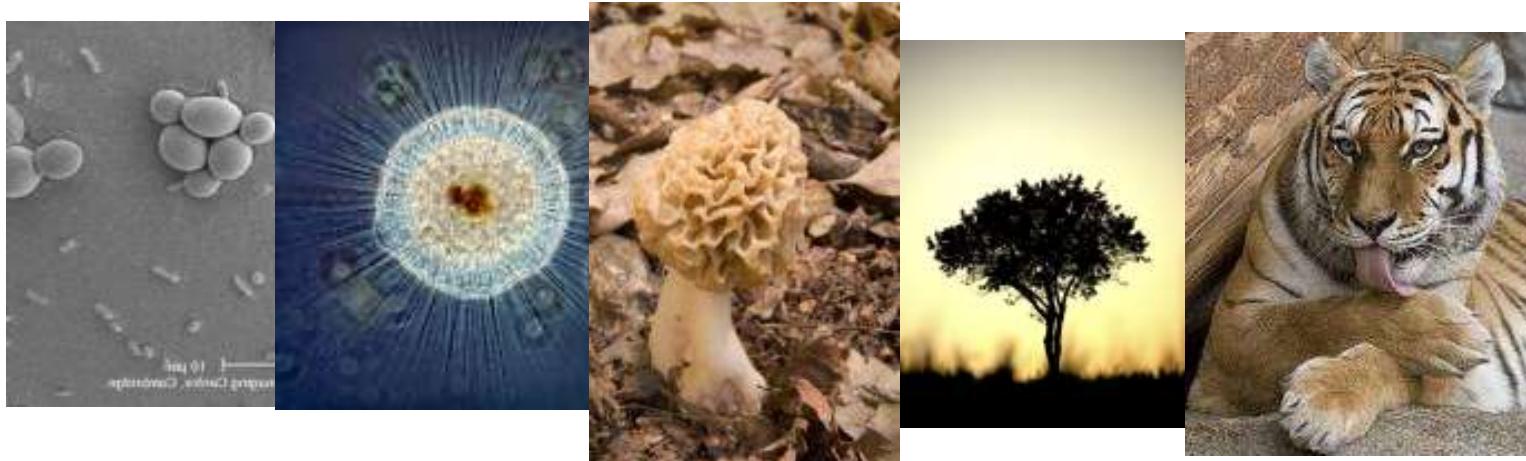
The work has been supported by Croatian Science Foundation

project IP-2018-01-2936

DPP III or M49 enzyme family

Zn dependent metalo enzymes of Molecular mass ~61-97 (103) kDa

widely spread 5 kingdoms: Eubacteria, Protista, Fungi, Plantae, Animalia



Zinc-exopeptidase hydrolyzes dipeptides from the N-terminal of its substrates, peptides with three AA and more

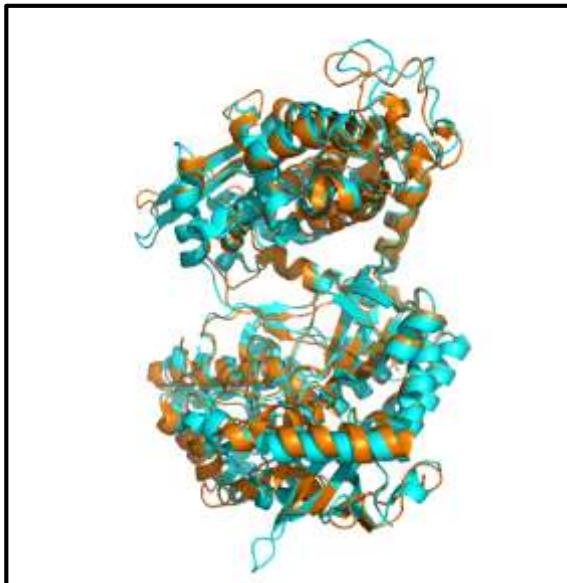


Human DPP III

2008. → yeast DPP III (PBD id 2CSK)

2009. → human DPP III (PBD id 3FVY)

2012. → E451A mutant of human DPP III with tynorphin (PBD id 3T6B)



Oxidative
stress

Cancer

Pain

Immune
Response

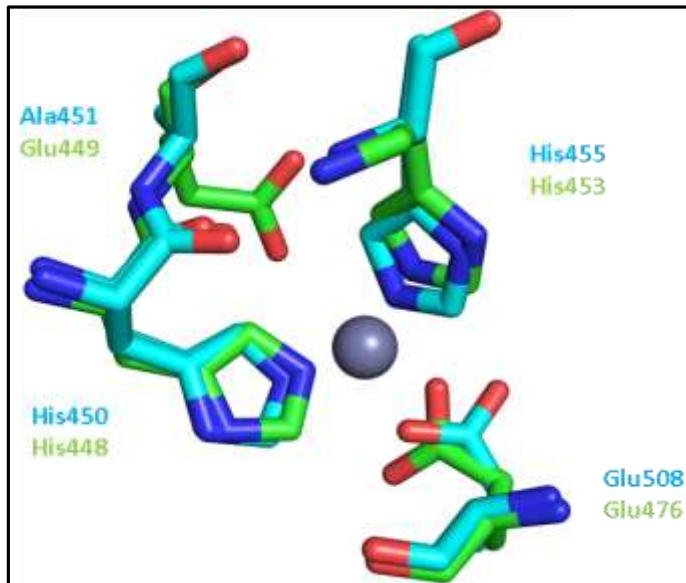
cardiac shock
biomarker

Influenza

Blood
pressure

Conserved motifs:

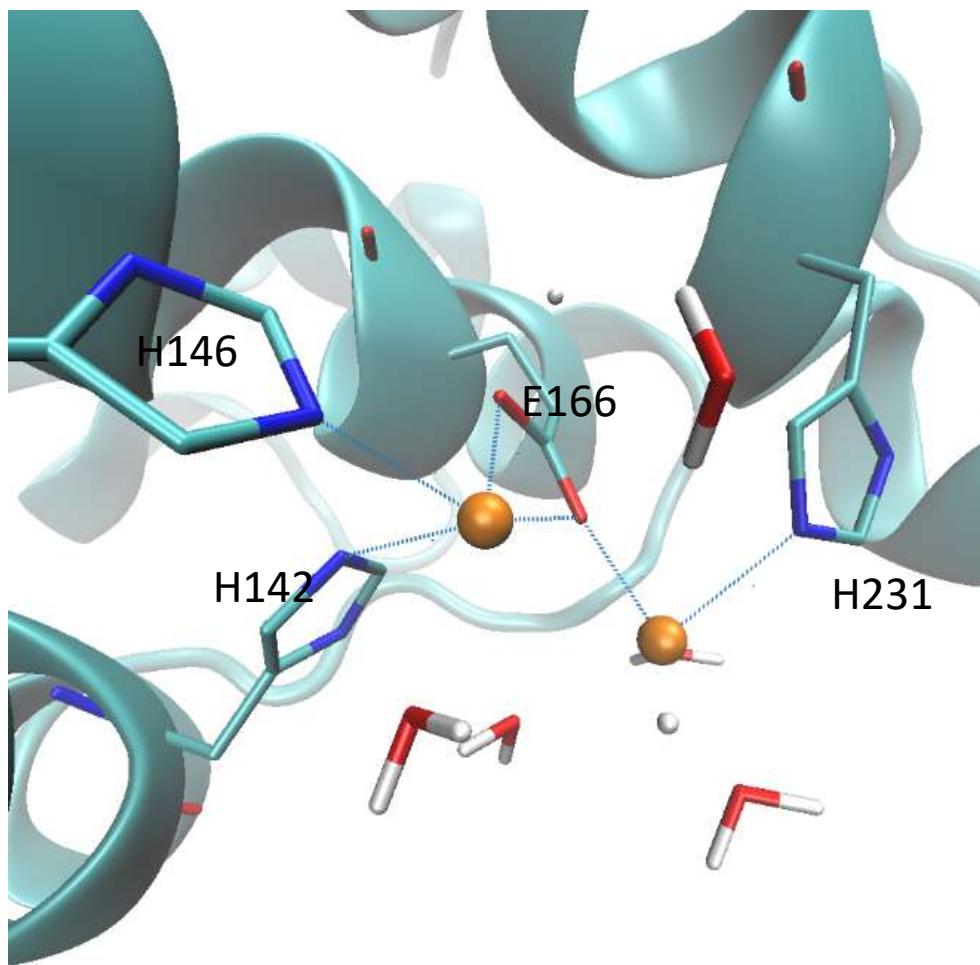
HEXX**G****H** and **E****E**X**R(K)****A****E(D)** coordinate Zn ion



Experiments: excess zinc inhibits hDPP III hydrolytic activity

Zn ion added as $\text{Zn}(\text{Ac})_2$ to the reaction mixture of pH 8.0, inhibited rat DPP III already at a $1\mu\text{M}$ level ($\text{IC}_{50} = 1.8\mu\text{M}$), while $10\mu\text{M}$ $\text{Zn}(\text{Ac})_2$ completely abolished its activity.

Thermolysin, PDB_id 1Ind (TML)

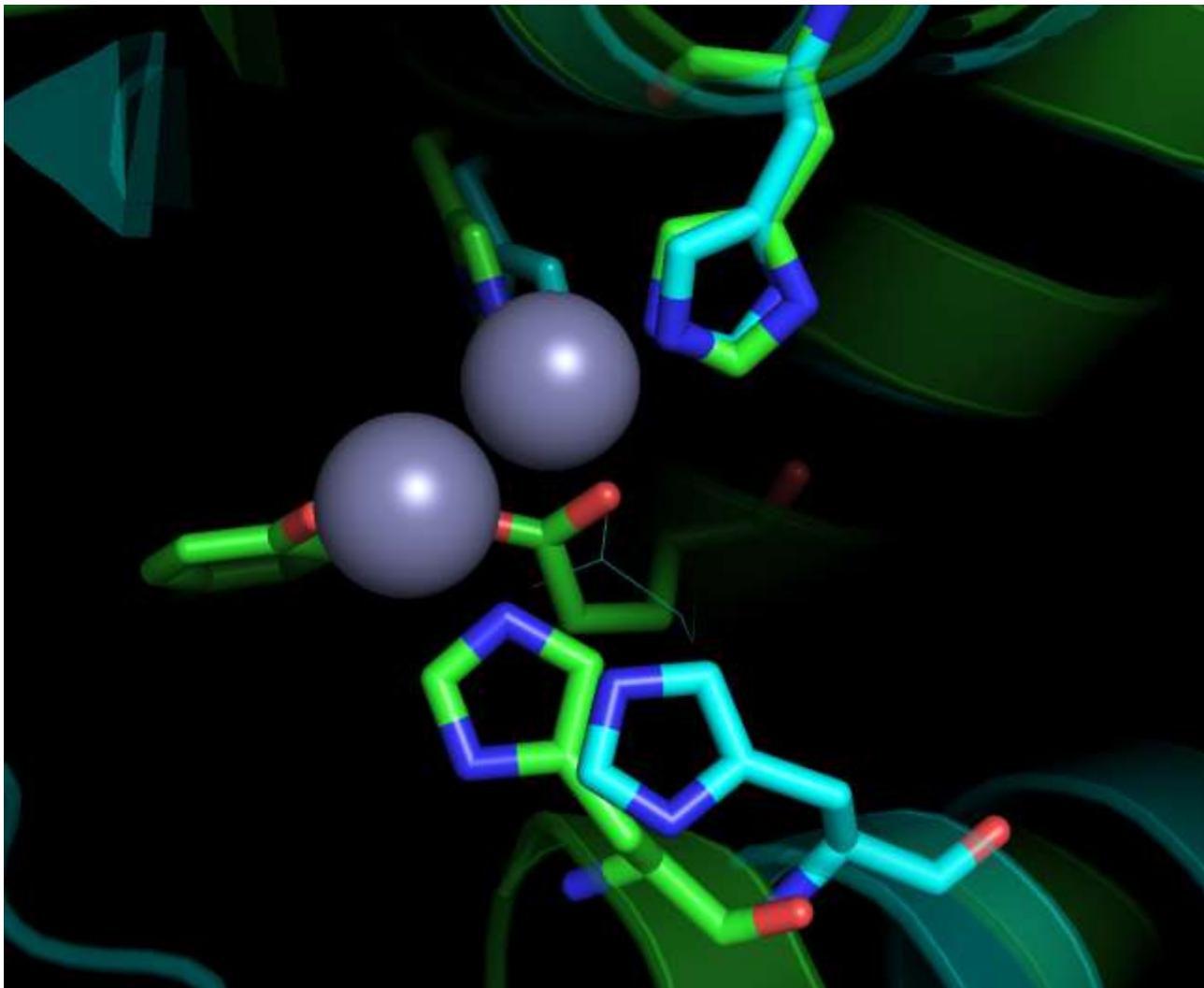


Excess zinc also inhibits other exo- and endometallopeptidases. One example is thermolysin where the inhibitory effect is explained by the binding of a second zinc ion to the catalytically important H231 within 3.2 Å of the zinc bound to native thermolysin.

Identifikacation of the inhibitory Zn binding site hDPP III



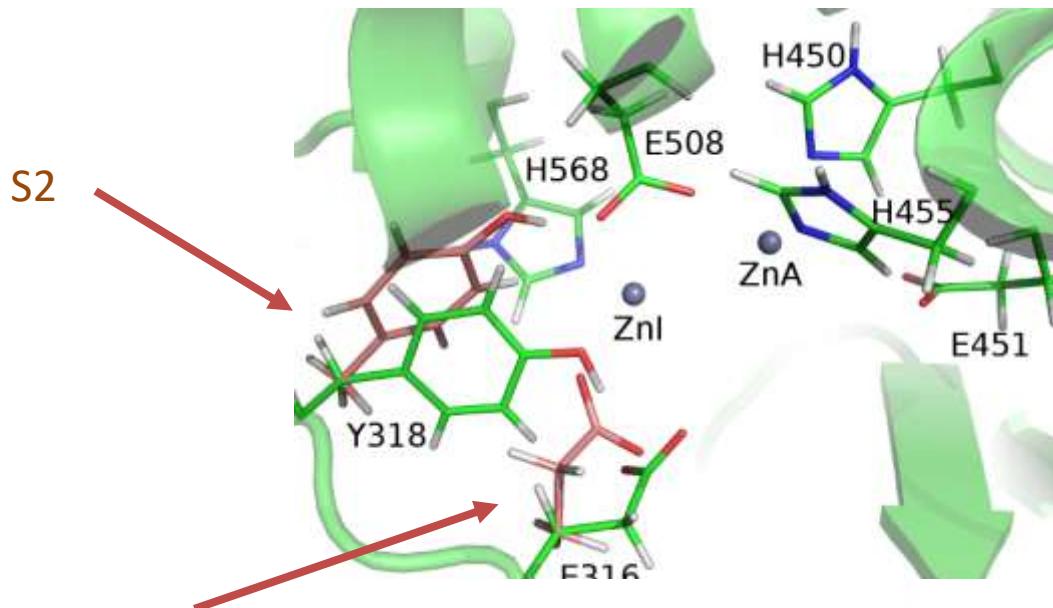
hDPP III – TML 3D structures - alignment



hDPP III – TML hDPP III -sequences - alignment

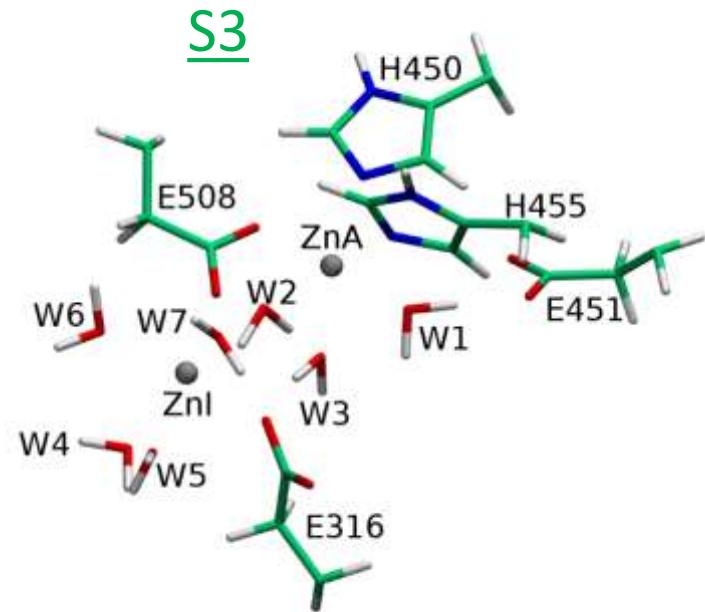
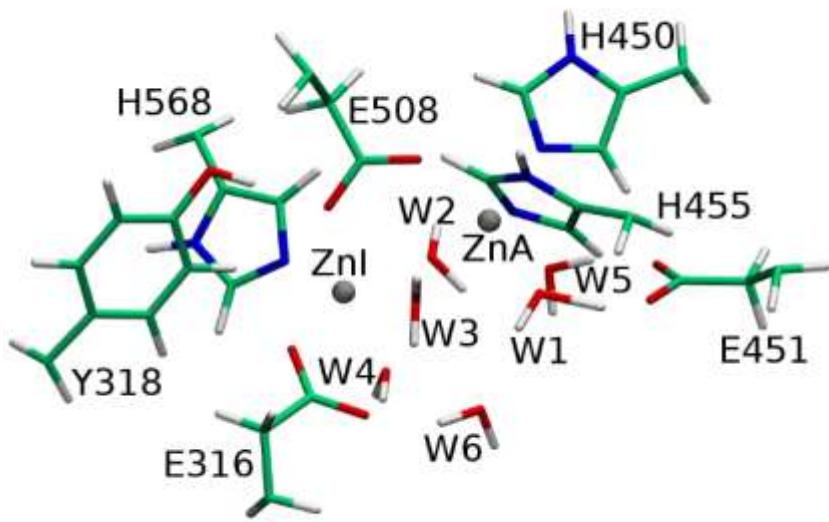
| | |
|------------|---|
| 419 99 | YATQREKLTFLEEDDKDLYILWKGPSFDVQVGLHELLGHGS GKL FVQDEKGAFNFDQETV --AIRSSVHYS---QGYNNNAFWNGSE-----MVYGDGDGQT FI PLSGGIDVVAHEL- : *...: : : * : * . : *.*.*: *: . * . . |
| 479 145 | INPETGEQIQSWYRSGETWDSKFSTIASSYEECRAESVG LYLCLHP QVLEIFGFEGADE ----THA-VTDYTAGLI YQNE SGAIN-----EAISDIFG- - : * :* - .: .: * - : : *** |
| 539 174 | DVIYVNWLNMVRAGLLALEFYTPEAFNWR QAHM QARFVILRV L EAGEGLVTITPTTGSD -----TLVEFYANKNP DWEIGEDVYTPGI -----SGDSLRSMS-DPAKY :****: : :*.. . * :*:.* : : .. |
| 599 212 | GRPDARVRLDRSKIRSVGKP ALERFLRR LQVLKSTGD-----VAGGRALYEGY GDPDHYSKRYTGT QDN GGVHINS GIINKAAYL I SQGGTHYGVSVVG IGRD KIFYRAL * ** : * - . : .: * * *. *: :* .. |
| 647 272 | ATVTDAPPECFLTLRDTVLLRK ESRKL IVQPNT RLEGSDVQL LEYEASAAGLIRSF SERF TQ-YLTPTSNFSQLRA-----AAVQSATDLYGSTSQEV-----ASVKQAF : * * ** : ** * * * * : * .. * |

QMMM calculations (S_1 , S_2 , $\underline{S_3}$)



ZNI coordination:

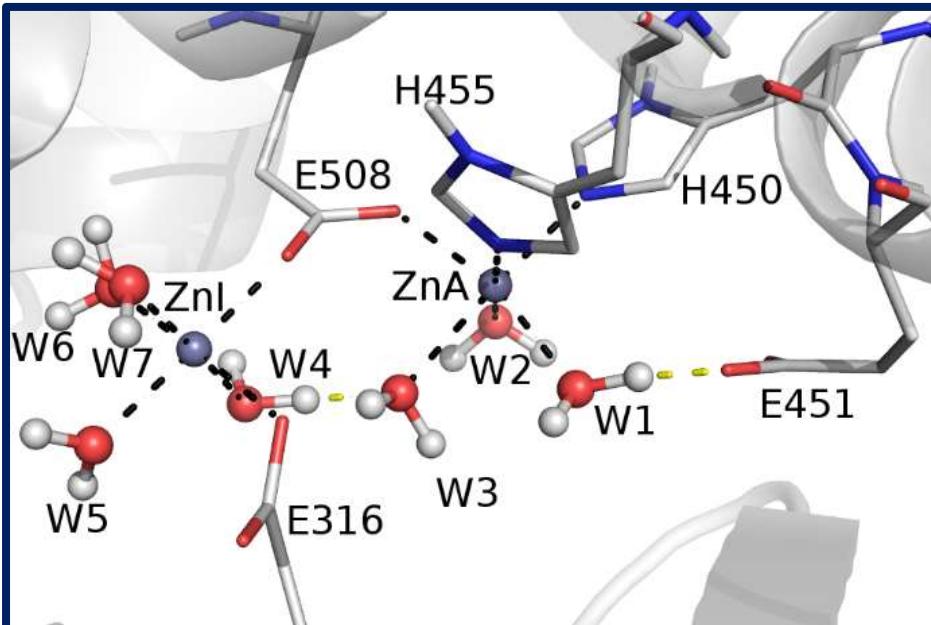
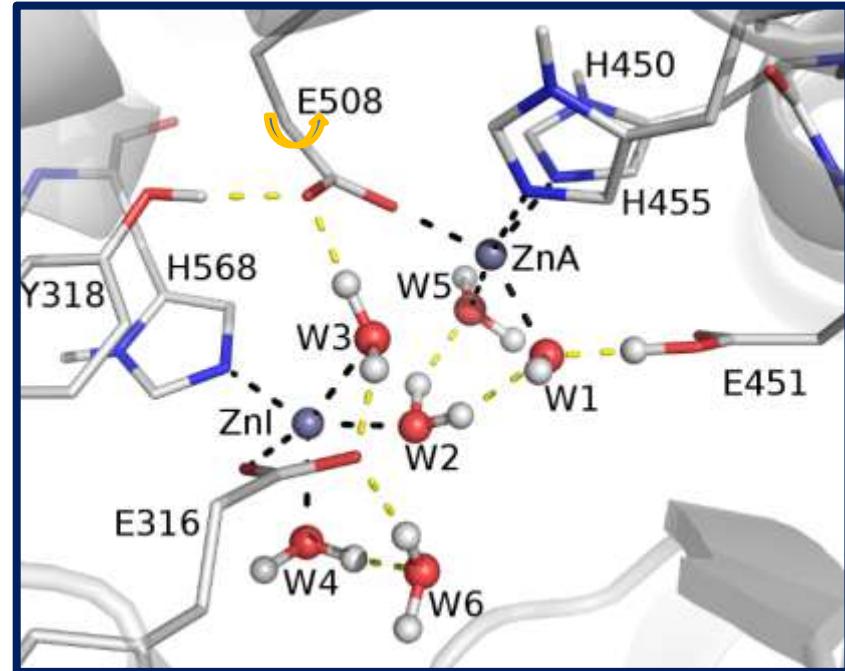
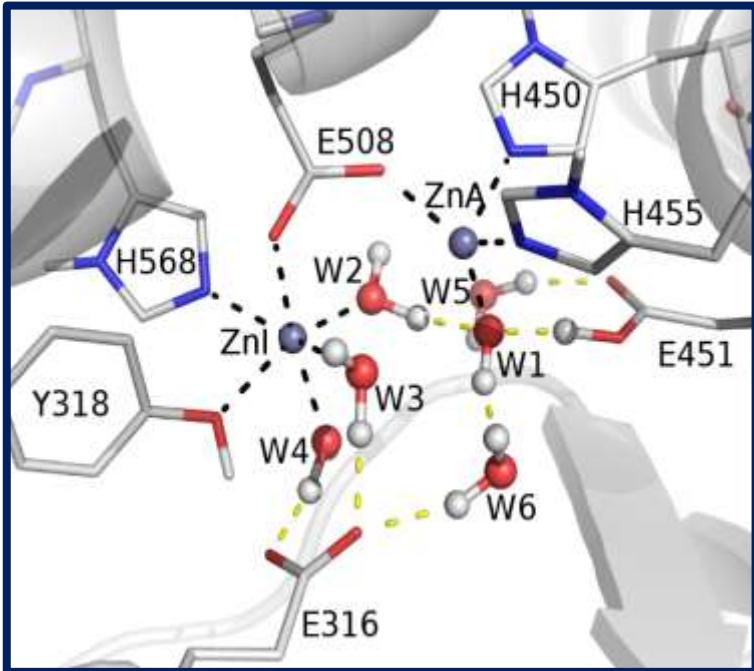
E508, H568 , Y318
E508, H568 , E316
E508, E316,
+ 3W



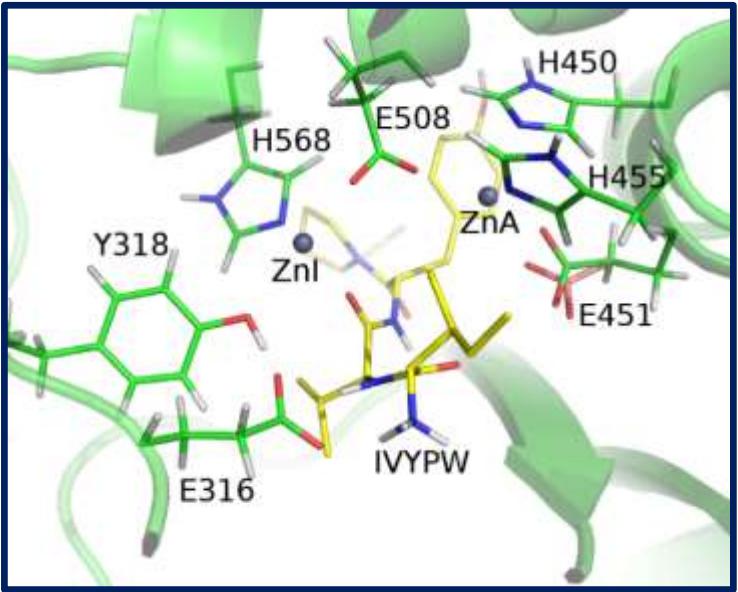
S1 / S2 / S3 initial and QMMM optimized

| | $d/\text{\AA}$ | | | | |
|---------------|----------------|------|------|------|------|
| | I | S1 | S2 | I | S3 |
| ZnA-ZnI | 3.45 | 3.78 | 4.31 | 5.05 | 4.95 |
| ZnA-H450(ne2) | 2.34 | 2.11 | 2.14 | 2.29 | 2.19 |
| ZnA-H455(ne2) | 2.45 | 2.10 | 2.21 | 2.39 | 2.16 |
| ZnA-E508(oe2) | 2.18 | 2.03 | 2.06 | 2.05 | 2.14 |
| ZnI-E508(oe1) | 2.17 | 2.16 | 3.98 | 2.06 | 2.09 |
| ZnI-H568(ne2) | 2.35 | 2.20 | 2.18 | 9.59 | 9.57 |
| ZnI-Y318(oh) | 2.53/4.19 | 2.22 | 4.79 | 9.28 | 9.98 |
| ZnI-E316(oe2) | 3.89 /2.75 | 4.03 | 2.10 | 2.07 | 2.11 |

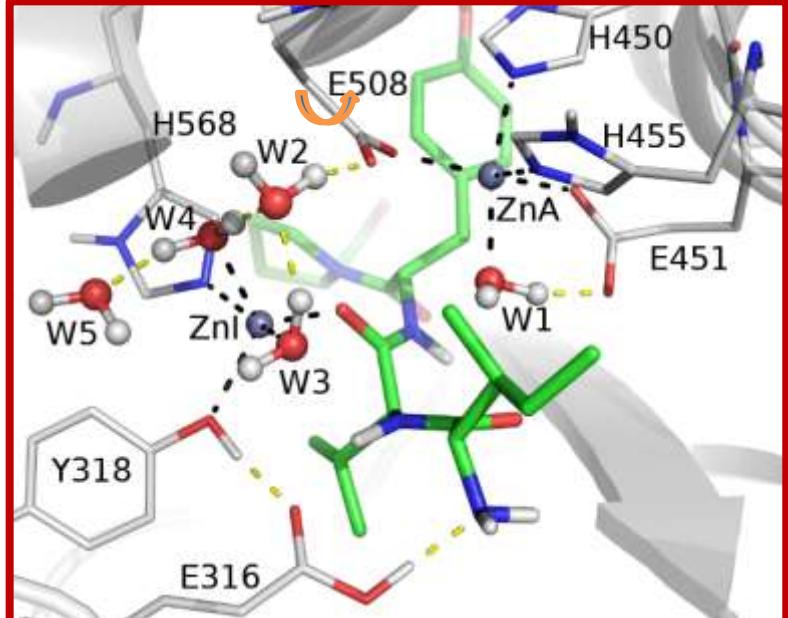
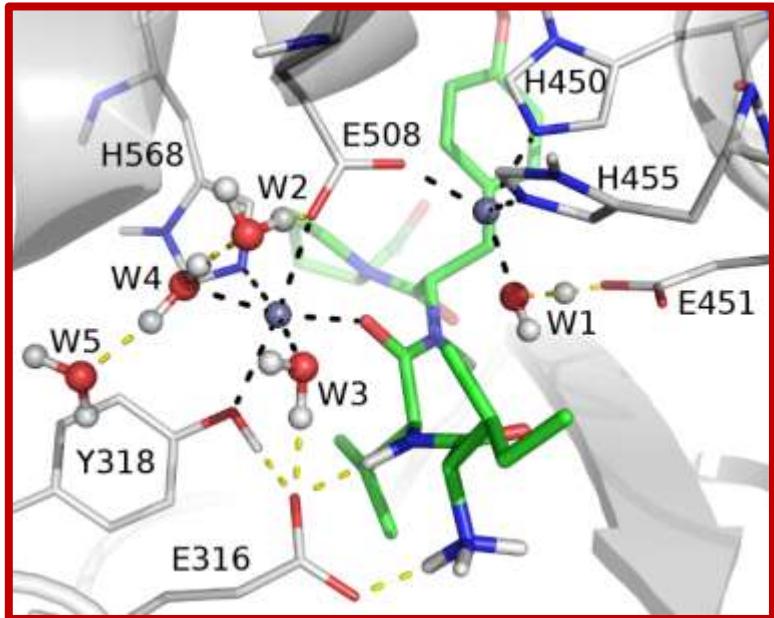
S1 / S2 / S3 QMMM optimized



QMMM calculations (CPLX1 / CPLX2)



| | <i>d</i> /Å | | |
|----------------------|-------------|-------|-------------|
| | I | CPLX1 | CPLX2 |
| ZnA-ZnI | 5.30 | 4.75 | 5.54 |
| ZnA-H450(ne2) | 2.19 | 2.11 | 2.15 |
| ZnA-E451(oe2) | 2.93 / 2.19 | 3.67 | 2.18 |
| ZnA-H455(ne2) | 2.29 | 2.14 | 2.17 |
| ZnA-E508(oe2) | 2.17 | 2.02 | 2.10 |
| ZnI-E508(oe1) | 2.20 | 2.29 | 4.26 |
| ZnI-H568(ne2) | 2.30 | 2.23 | 2.19 |
| ZnI-Y318(oh) | 2.60 | 2.28 | 2.17 |



Zn nonbonding parameters models

| Model | # | charge/e | | | | E451 | |
|------------|-----------------|------------------|--|--|--|-------|--|
| | | Zn | | | | | |
| Dummy atom | D1 ¹ | Total | central point (carry vdw parameters) | δ points at z- axis of octahedron | points at x and y axes of octahedron | -0.65 | |
| | D2 | 2 | -1 | 0.5 | 0.5 | | |
| | | 1.325 | -0.475 | 0.1 | 0.4 | | |
| 6-12 | 1 ² | 2.0 | | | | -0.75 | |
| | 2 | 1.375 | | | | | |
| | 3 | ZnA 1.1, ZnI 0.9 | | | | | |
| | 3' | 1.0 | | | | | |
| | 3r | ZnA 0.9, ZnI 1.1 | | | | | |

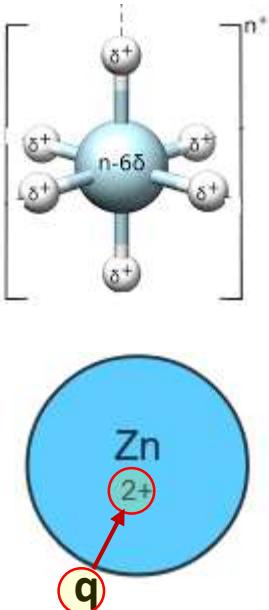


$$V(r^N) = \sum_{veze} \frac{k_l}{2} (l_i - l_{i,0})^2 + \sum_{kutovi} \frac{k_\theta}{2} (\theta_i - \theta_{i,0})^2 + \sum_{torzije} \frac{V_n}{2} (1 + \cos(n\omega - \gamma))$$

$$+ \sum_i^N \sum_j^N \left(4\epsilon_{ij} \left[\left(\frac{\sigma_{ij}}{r_{ij}} \right)^{12} - \left(\frac{\sigma_{ij}}{r_{ij}} \right)^6 \right] + \frac{q_i q_j}{4\pi\sigma_{ij}} \right)$$

Lorentz-Berthelot rules

$$\sigma_{ab}^{\text{LB}} = \frac{\sigma_{aa} + \sigma_{bb}}{2}, \quad \epsilon_{ab}^{\text{LB}} = \sqrt{\epsilon_{aa}\epsilon_{bb}}.$$



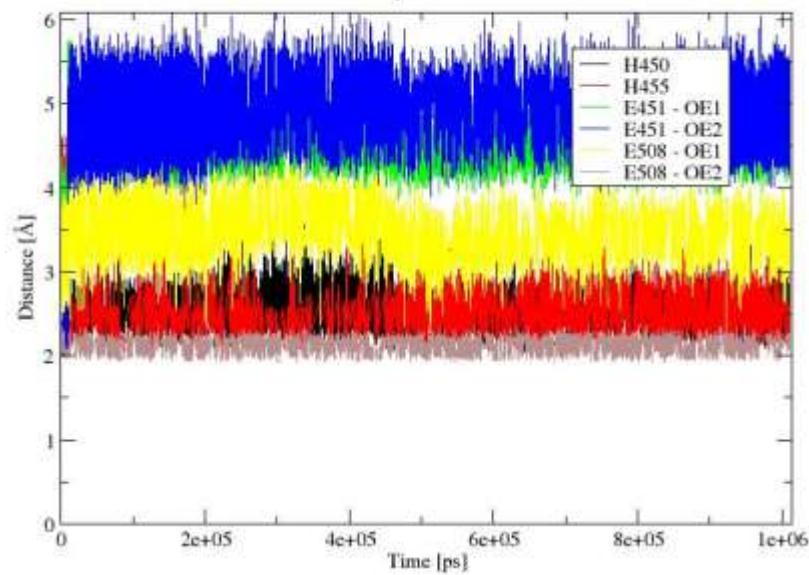
Dummy atom Zn parameter models

| Model # | SYSTEM Force field | t _r /t _{total} (μs) | <ZnA–ZnI> (Å) | ZnA ligands | ZnI ligands | MMGBSA(kcal/mol) | | |
|-----------|------------------------|--|--|--|---|--|----------------------------------|-------------------------------------|
| | | | | | | Receptor: DPP III Ligand: ZnA + ZnI | Receptor: DPP III Ligand: ZnA | Rec: DPP III+ ZnA Ligand: ZnI |
| D1 | S1 ff14SB | 0.099/0.1 ^{ZnA} 0.095/0.1 ^{ZnI} | <Zn _A –Zn _I > (Å) | H450, H455, E508^M, E451^M 2 W | H568, E316^B, E508^M, Y315 2W | -292±11 | -164±6 | -108±7 |
| | S1 ff03 | 0.1/0.1 | 5.4±0.1 51% 5.1±0.1 49% | H450, H455, E508^M, E451^M 2W | E316^M, E508^M, H568 3W | -320±8 | -187±6 | -121±6 |
| D2 | S1 ff03 | 0.085/0.1 ^{ZnA} /0.1/0.1 ^{ZnI} | 3.5±0.2 | E508^M, H450, H455 mostly 3W | E316^M, E508^M mostly 3W | -144±6 | -80±5 | -47±7 |
| | S1 ff03 | 0.066/0.1 ^{ZnA} 0.056/0.1 ^{ZnI} | 4.9±0.5 | E508^M, H450, H455 3W | E316^M, E508^M, H568 3W | -166±8 | -81±5 | -50±10 |
| | S1 ff14SB | 0.58/1 ^{ZnA} 0.88/1 ^{ZnI} | 5.4±0.3 | H450, H455, E508^M 3-4W | E316^M, E508^M mostly 4 W | -122±8 | -67±5 | -48±7 |
| | S2 ff14SB | 0.5/1 ^{ZnA} 0.99/1 ^{ZnI} | 4.7±0.5 | H450, H455, E508^M 3-4W | E316^M, E508^M 4-5W | -116±7 | -68±5 | -43±5 |
| | S1 ff14SB | 0.062/0.1 ^{ZnA} 0.1/0.1 ^{ZnI} | 4.1±0.3 | H450, H455, E508^M 3W | E508^M 4-5 W | -102±9 | -69±4 | -23±6 |
| | S1 ff14SB | 0.58/0.25 ^{ZnA} 0.88/0.25 ^{ZnI} | 4.7±0.1 | H450, H455, E508^M, E451^M 1W | H568, E508^M, V730 2-3W | -160±7 | -96±5 | -57±5 |
| | CPLX1 ff14SB | 0.5/0.5 ^{ZnA} 0.5/0.5 ^{ZnI} | 5.2±0.3 | H450, H455, E508^{M,B}, E451 mostly 1 W | E508^M mostly 4 W | -114±7 | -93±5 | -18±4 |
| | CPLX2 ff14SB | 0.5/0.5 ^{ZnA} 0.44/0.5 ^{ZnI} | 5.4±0.3 | H450, H455, E508^{M,B} 1W | H568, E508^{M,B} cHM-O2 nd mostly 2 W | -150±8 | -85±5 | -39±4 |

D2 model

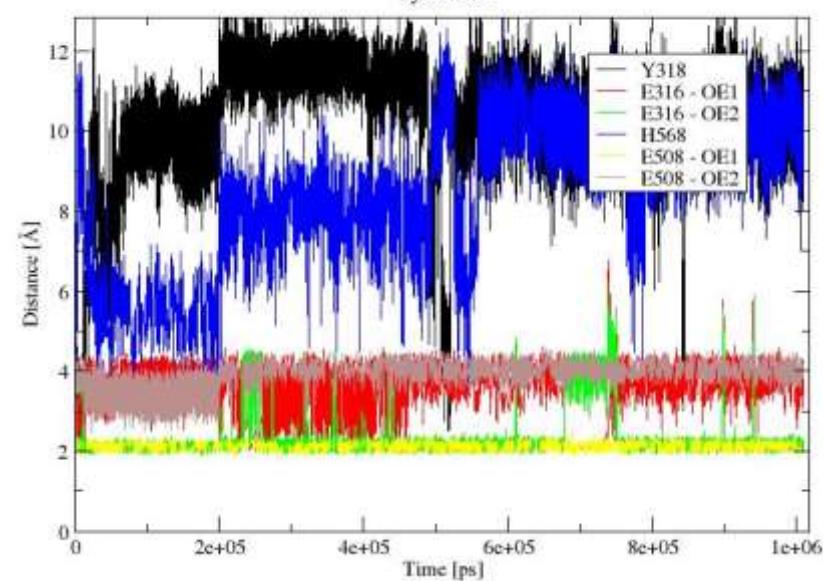
ZnA coordination

System S1



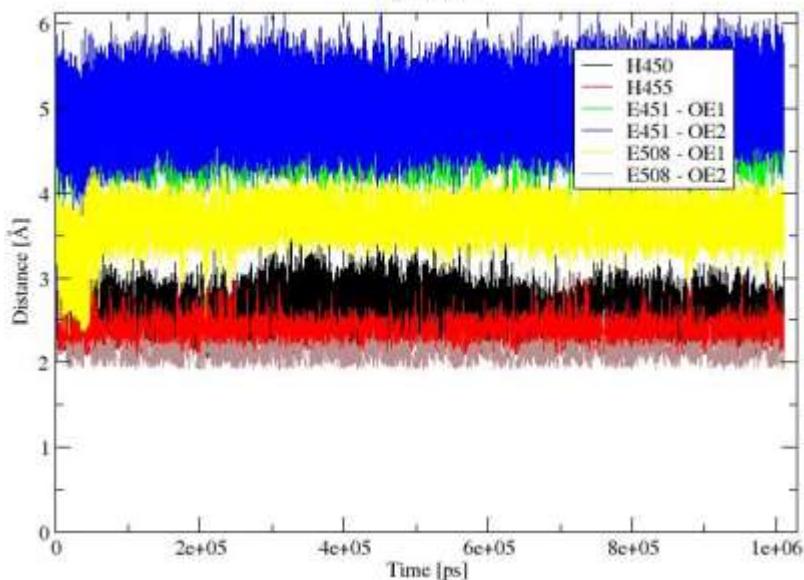
ZnI coordination

System S1



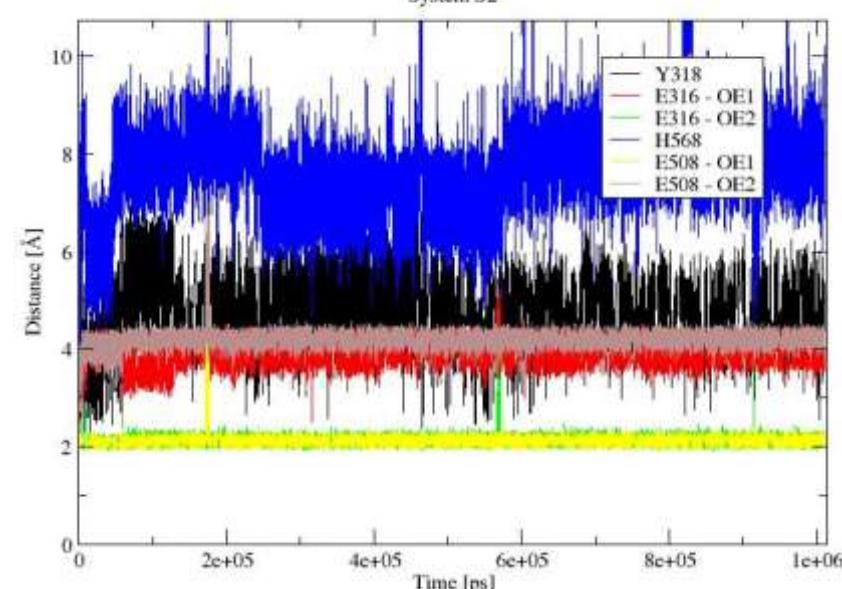
ZnA coordination

System S2

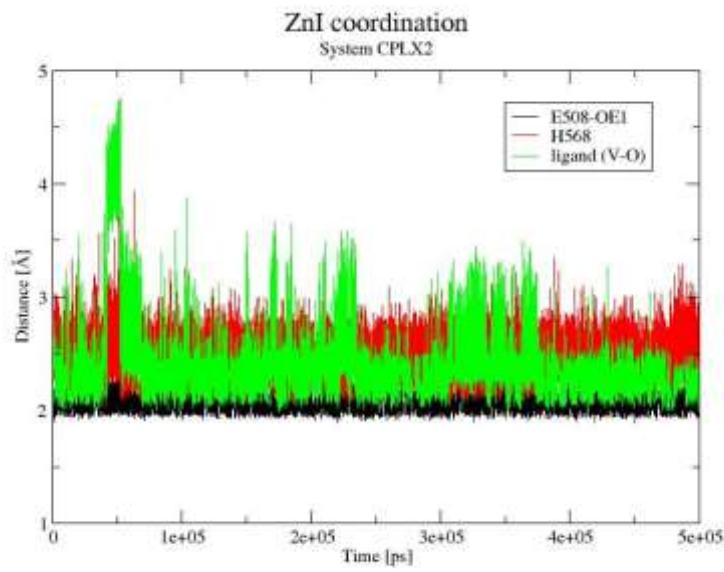
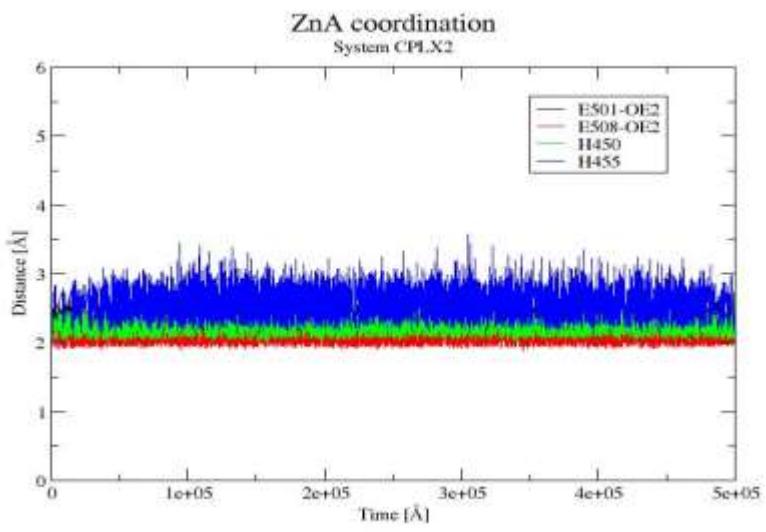
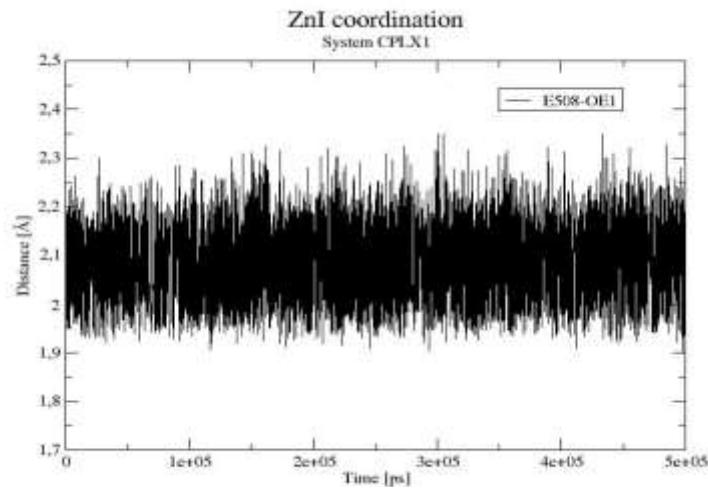
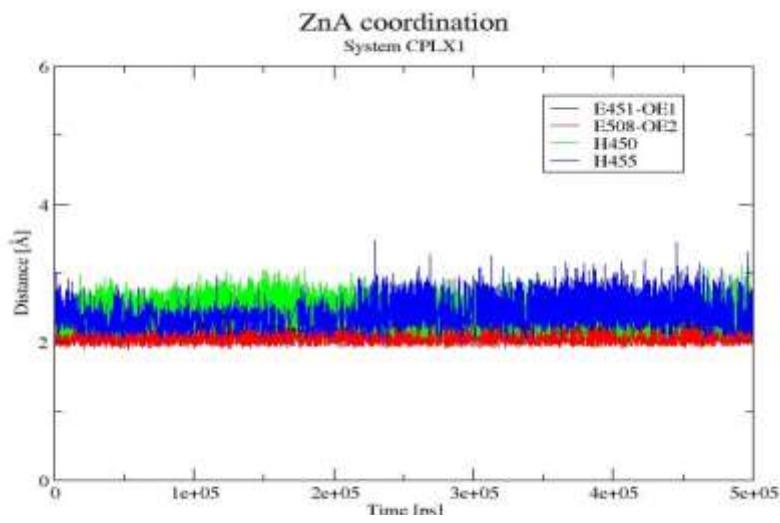


ZnI coordination

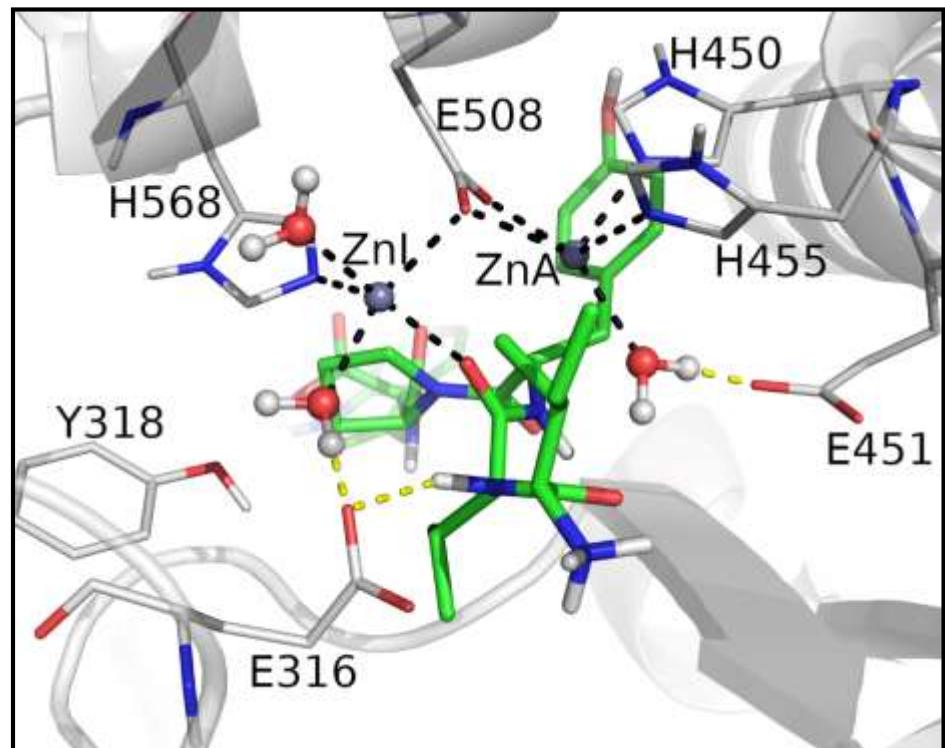
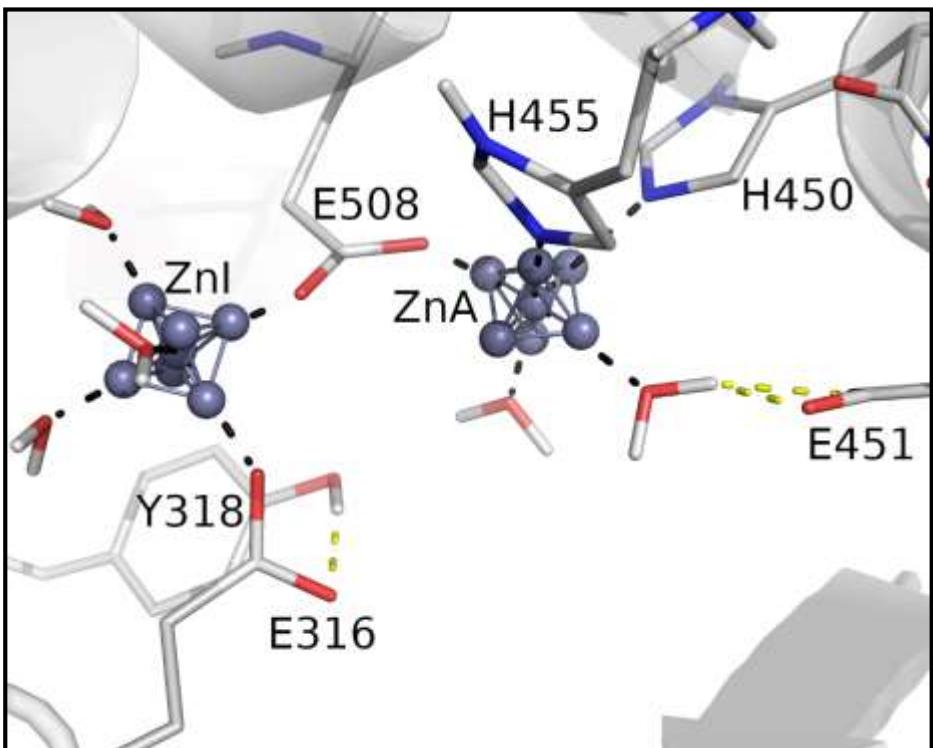
System S2



D2 model



D2 model, structures of S2 and CPLX2 obtained after 1μs and 500 ns, respectively, of MD simulations

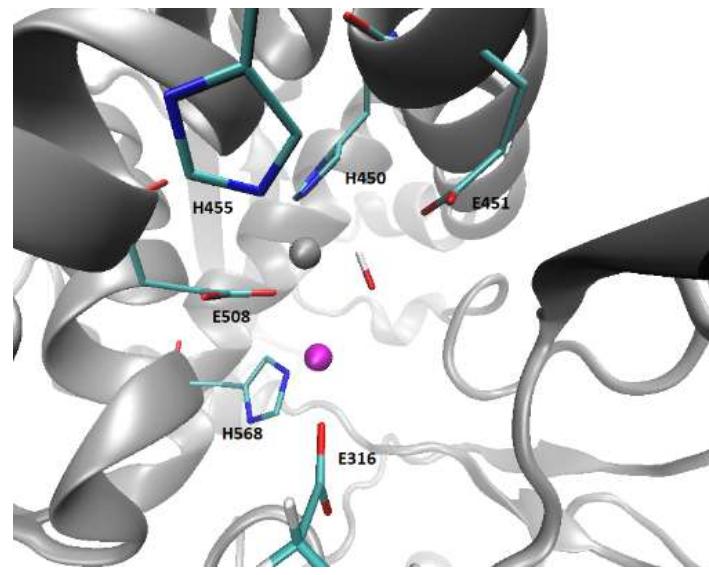
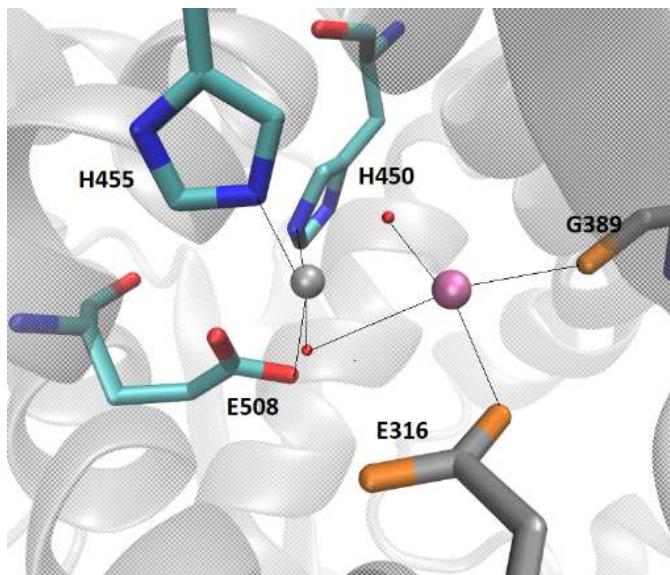
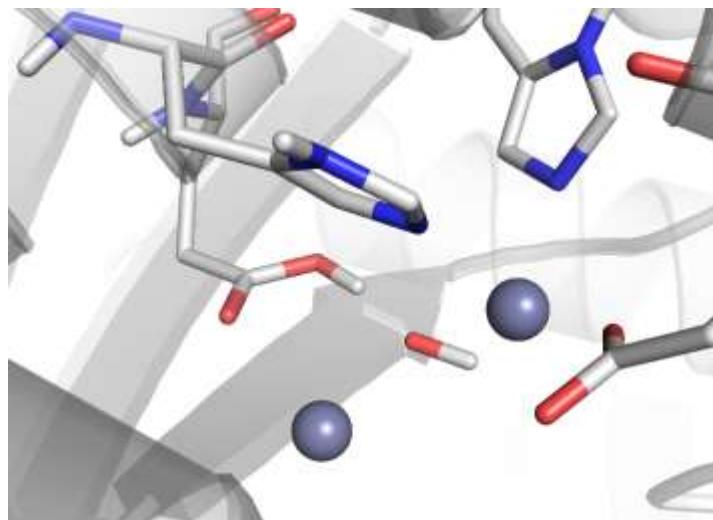


12-6 ZN parameters models

| Model # | SIMULATED SYSTEM Force field | t _i /t _{total} ^a (μs/μs) | < ZnA –ZnI > (Å) | ZnA ligands | ZnI ligands (coordinated with) | MMGBSA(kcal/mol) ^b | | |
|---------|---------------------------------|--|---------------------------|--|---|---|---|-----------------------------|
| | | | | | | Rec: DPP III Ligand: Zn _A + Zn _I | Receptor: DPP III + ZnA Ligand: Zn _I | Rec: DPP III Ligand: ZnA |
| 2 | S2 ff03 | 0.85/1 | 4.0±0.4 | H450, H455, E508 ^M , E451 ^M W 1-4 | E508 ^B , E316 ^B W 1-4 | -31±4 | -2±2 | -27±3 |
| | S1 ff03 | 0.99/1 ^{ZnA} 0.81/1 ^{ZnI} | 3.8±0.2 | H450, H455, E508 ^M , E451 ^M mostly 2 W | E508 ^M , E316 ^B mostly 3 W | -32±4 | 1±3 | -26±3 |
| | S1-OH ⁻ ff03 | 0.25/0.35 ^{ZnA} 0.21/0.35 ^{ZnI} | 3.5±0.3 | H450, H455, E508 ^M , E451 ^M , OH ⁻ | E316 ^B E508 ^B , OH ⁻ mostly 3 W | -24±6 | 13±5 | -27±4 |
| 3 | S1-OH ⁻ ff03 | 0.40/0.44 ^{ZnA} 0.17/0.44 ^{ZnI} | 7.0±3.2 | H450, H455, E508 ^{B,M} , OH ⁻ 0-1 W | E316 ^{B,M} , 2-3 W | -52±5 | -4±3 | -43±3 |
| | S1- OH ⁻ ff14SB | 0.69/0.7 ^{ZnA} 0.55/0.7 ^{ZnI} | 3.5±0.2 (last 590 ns) | H450, H455, E508 ^M , OH ⁻ 1-2 W | E451 ^{M,B} , OH ⁻ 2-3 W | -49±4 | -11±4 | -35±3 |
| | S1- OH ^{- exc} ff14SB | 0.5/1.0 ^{ZnA} 0.87/1.0 ^{ZnI} | 10.0±0.7 (last 740 ns) | N294, E316 ^M mostly 2-3 W | H450, H455, E508 ^M mostly 1-2 W | -27±4 | -14±4 | -12±2 |
| | CPLX1 ff14SB | 1.1/1.1 ^{ZnA} 1.1/1.1 ^{ZnI} | 4.0±0.2 | H450, E451 ^{M,B} H455, E508 ^M , 1 W | H568, E508 ^{M,B} cHM- O2 nd mostly 1 W | -34±4 | -1±3 | -23±3 |
| 3' | CPLX2 ff14SB | 0.94/1.0 ^{ZnA} 1.0/1.0 ^{ZnI} | 3.8±0.2 | H450, H455, E508 ^{M,B} mostly 1 W | H568, E508 ^{M,B} cHM- O2 nd mostly 2 W | -11±4 | 4±3 | -8±3 |
| 3r | CPLX2 ^{exc} ff14SB | 0.44/0.98 ^{Zn} _A 0.41/0.98 ^{ZnI} | 4.5±0.4 | H450, H455, E451 ^{M,B} mostly 1W | H450, H455, , E508 ^{M,B} cHM-O2 nd mostly 1 W | -23±2 | -2±1 | -21±2 |

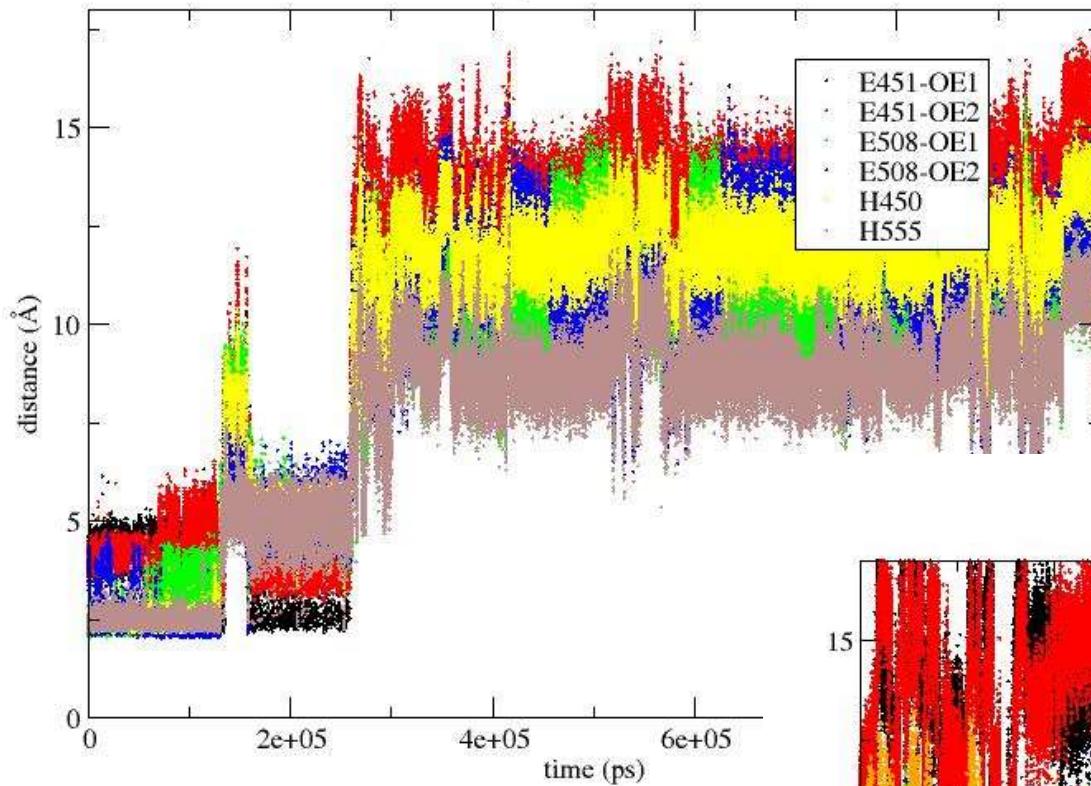
LJ nonbonding model 3

S1-OH⁻

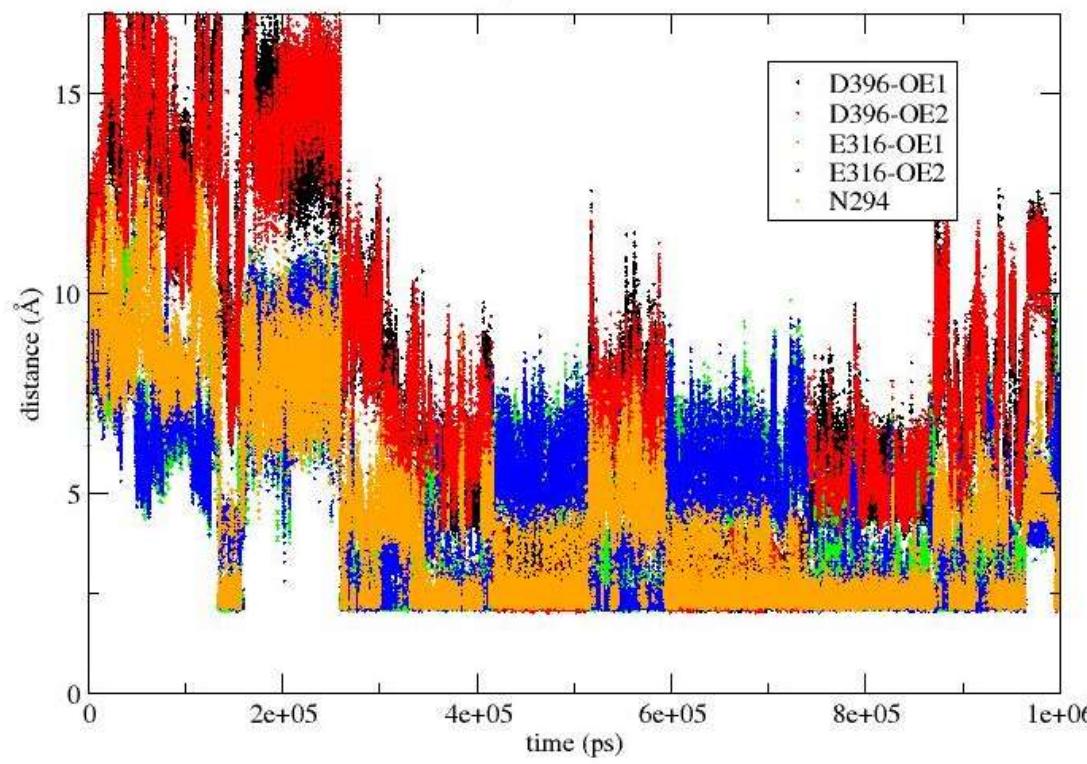


ZNA coordination
system S1-OH

MD-S1

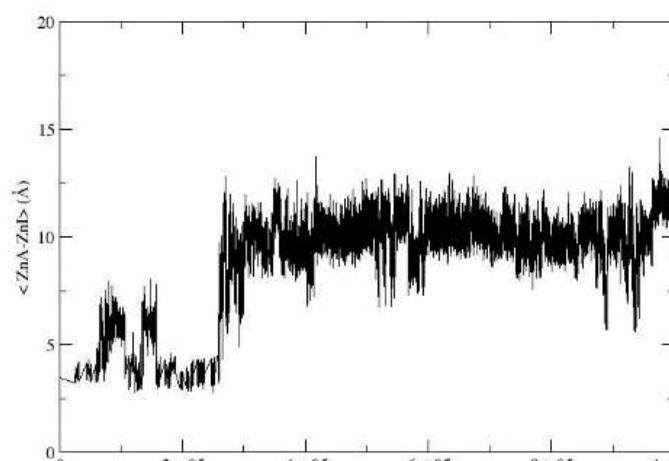
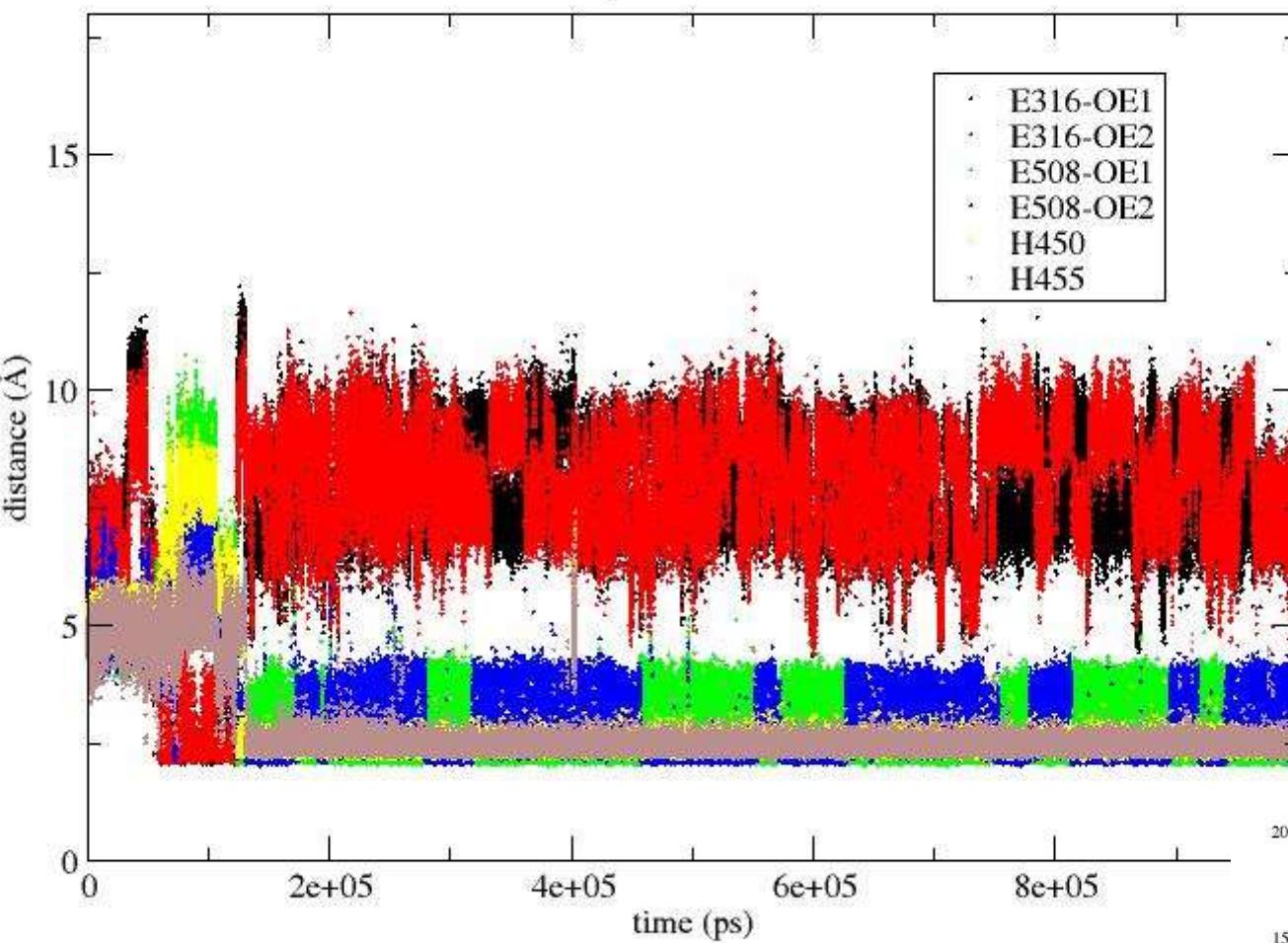


ZnA ligands
System S1-OH

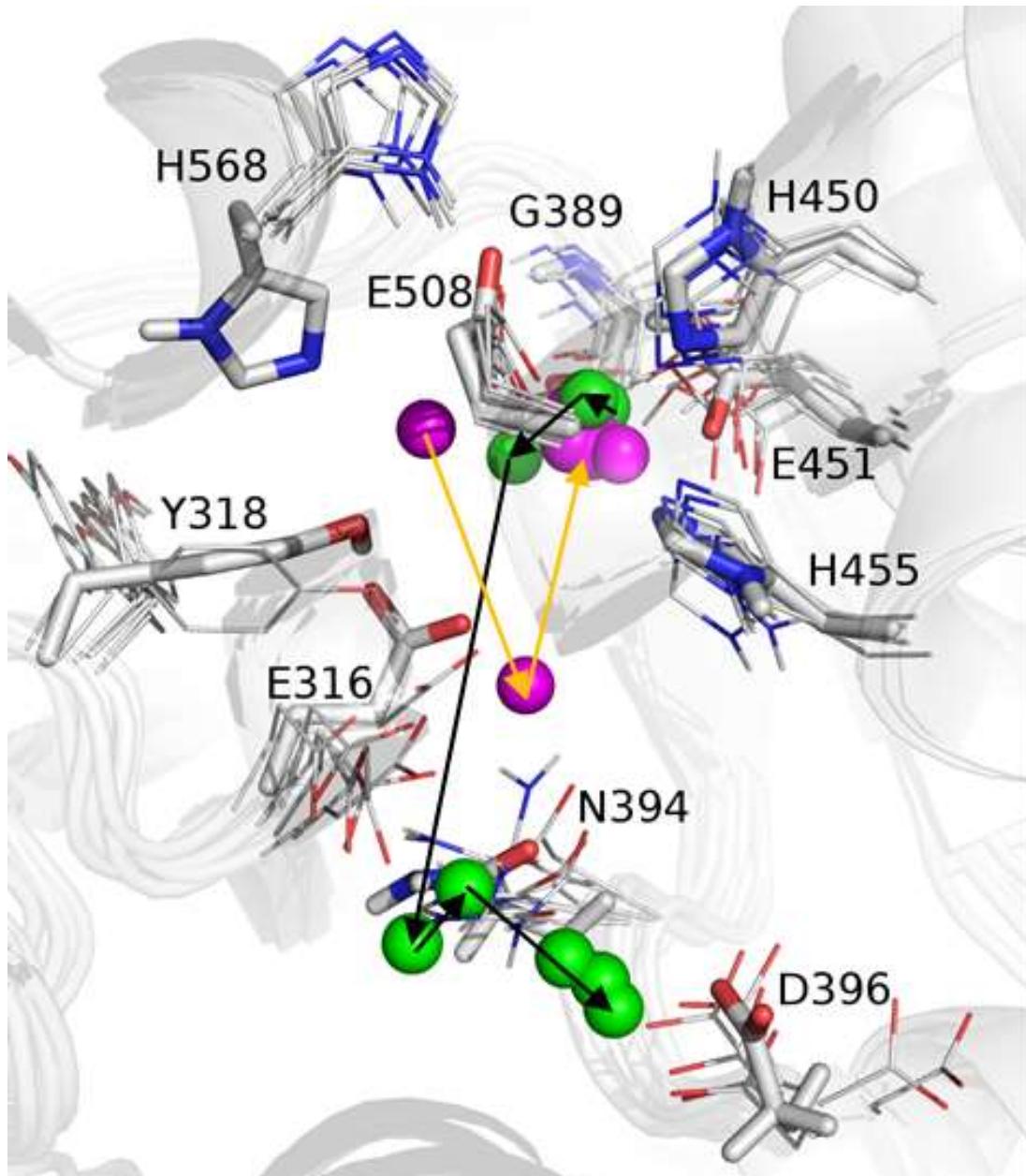


ZNI coordinatio
System S1-OH

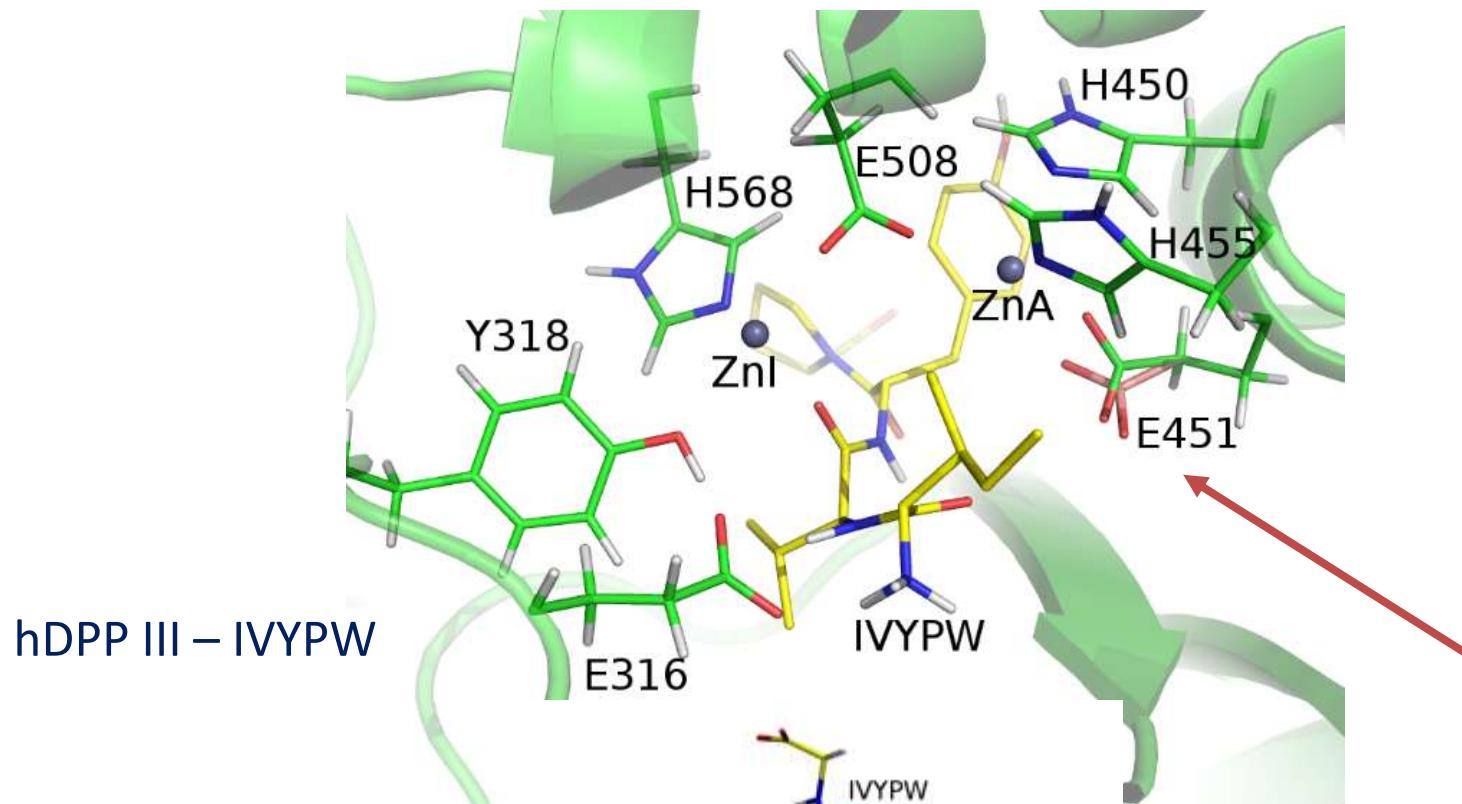
MD-S1



Izmjena ZNA i ZNI tijekom MD

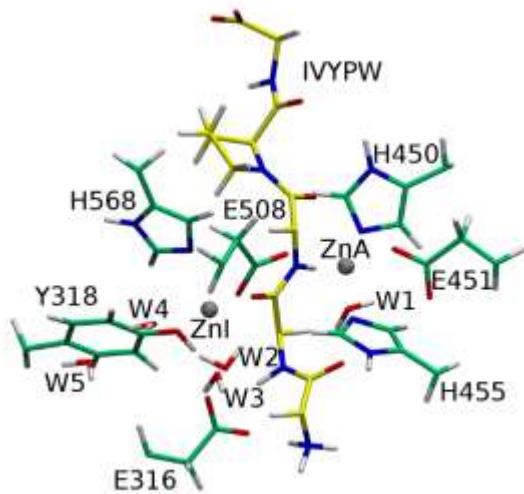


QMMM račun CPLX1 i CPLX2



ZnA coordination:

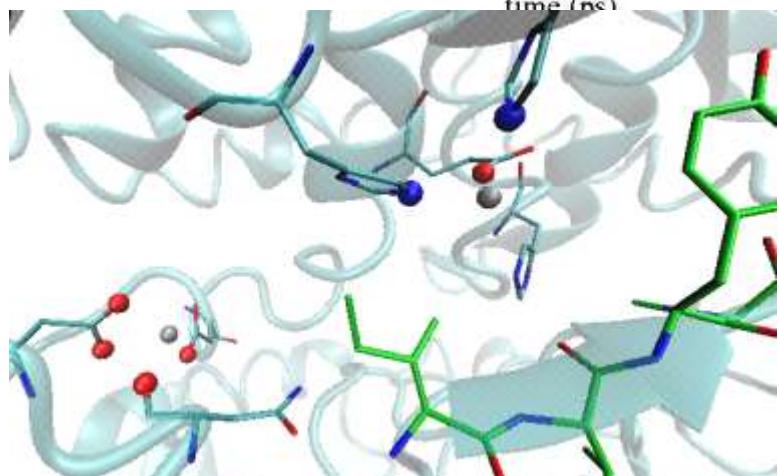
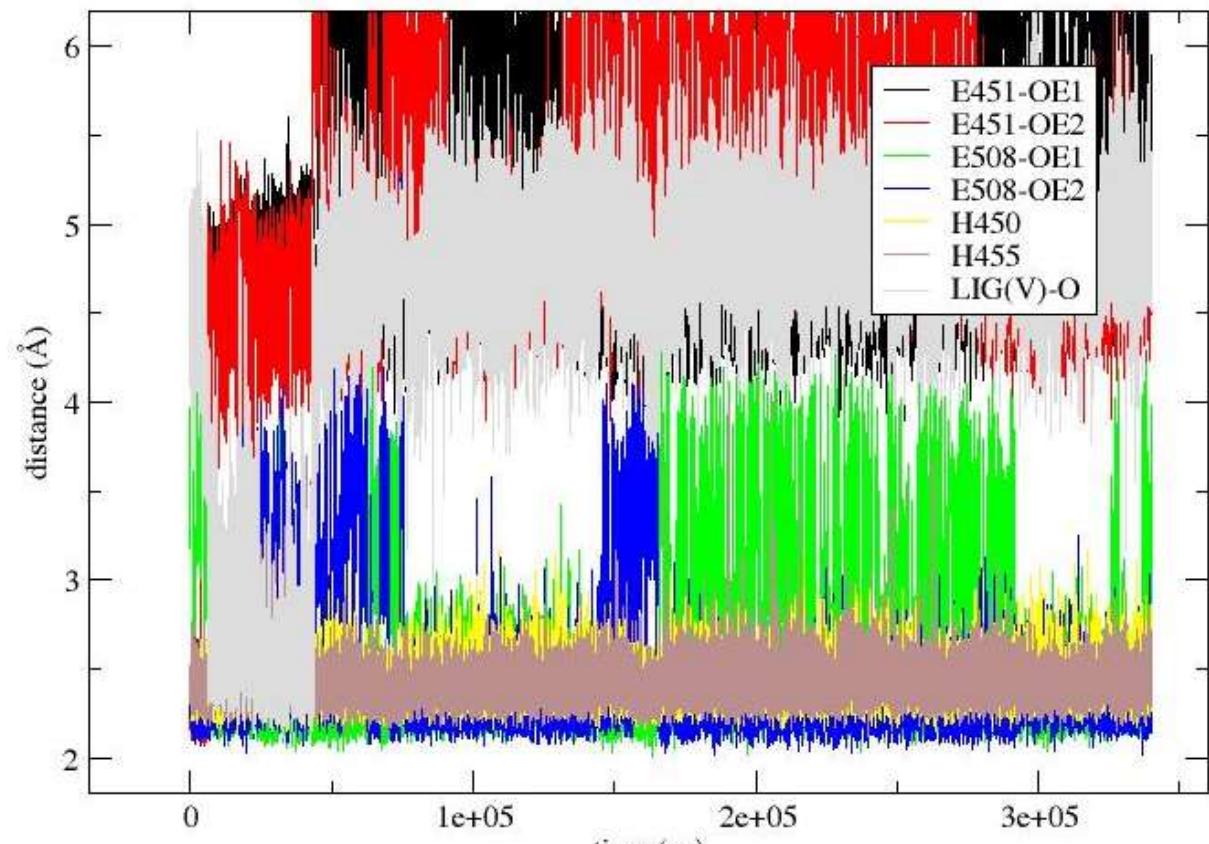
E508, H450, H455
E508, H450, H455, E451



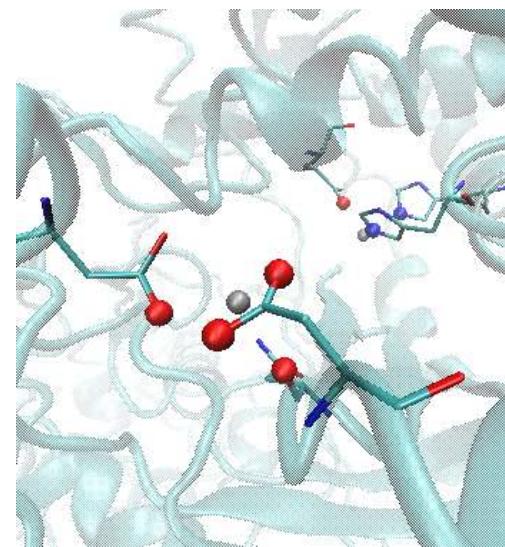
MD- cplx

| Model # | SYSTEM Force field | t _i /t _{total} (μs) | <ZnA–Znl> (Å) | ZnA ligands | Znl ligands | MMGBSA(kcal/mol) | | |
|---------|-----------------------|---|------------------|---|---|--|----------------------------------|-------------------------------------|
| | | | | | | Receptor: DPP III Ligand: ZnA + Znl | Receptor: DPP III Ligand: ZnA | Rec: DPP III+ ZnA Ligand: Znl |
| D2 | CPLX1 ff14SB | 0.5/0.5 ^{ZnA} 0.5/0.5 ^{Znl} | 5.2±0.3 | H450, H455, E508 ^{M,B} mostly 1 W | E508 ^M mostly 4 W | -114±7 | -93±5 | -18±4 |
| | CPLX2 ff14SB | 0.5/0.5 ^{ZnA} 0.44/0.5 ^{Znl} | 5.4±0.3 | H450, H455, E508 ^{M,B} 1W | H568, E508 ^{M,B} ^c HM-O2 nd mostly 2 W | -150±8 | -85±5 | -39±4 |
| 2 | CPLX1 ff03 | 0.24/0.25 ^{ZnA} 0.24/0.25 ^{Znl} | 5.2±0.3 | H450, H455, E508 ^M , E451 ^M 1-2 W | H565, E508 ^M mostly 2 W | -26±2 | 0±1 | -22±2 |
| 3 | CPLX1 ff14SB | 1.1/1.1 ^{ZnA} 1.1/1.1 ^{Znl} | 4.0±0.2 | H450, E451 ^M H455, E508 ^M , 1 W | H568, E508 ^{M,B} ^c HM-O2 nd mostly 1 W | -34±4 | -1±3 | -23±3 |
| | CPLX1 ff14SB | 1.1/1.2 ^{ZnA} 0.5/1.2 ^{Znl} | 4.0±0.2 | H450, E451 ^M H455, E508 ^M , 1 W | H568, E508 ^M ^c HM-O2 nd 1 W | -31±3 | -2±2 | -22±3 |
| | CPLX2 ff14SB | 0.32/0.34 ^{ZnA} 0.32/0.34 ^{Znl} | 11.6±0.9 | H450, H455, E508 ^M , mostly 2 W | D396 ^{M,B} , D496 ^{M,B} 1-2 W | -25±3 | -12±2 | -12±3 |
| | CPLX2 ff14SB | 0.28/0.28 ^{ZnA} 0.08/0.028 ^{Znl} | 18.6±12.7 | H450, H455, E508 ^M , 1 W | E508 ^M ^c HM-O2 nd 1 W | -21±5 | -4±3 | -17±3 |
| 3' | CPLX2 ff14SB | 0.94/1.0 ^{ZnA} 1.0/1.0 ^{Znl} | 3.8±0.2 | H450, H455, E508 ^{M,B} mostly 1 W | H568, E508 ^{M,B} ^c HM-O2 nd mostly 2 W | -11±4 | 4±3 | -8±3 |
| 3r | CPLX2 ff14SB | 0.44/0.72 ^{ZnA} 0.21/0.72 ^{Znl} | 4.5±0.4 | H450, H455, E451 ^{M,B} mostly 1W | H568, E508 ^{M,B} ^c HM-O2 nd mostly 1 W | -21±3 | -2±2 | -14±5 |

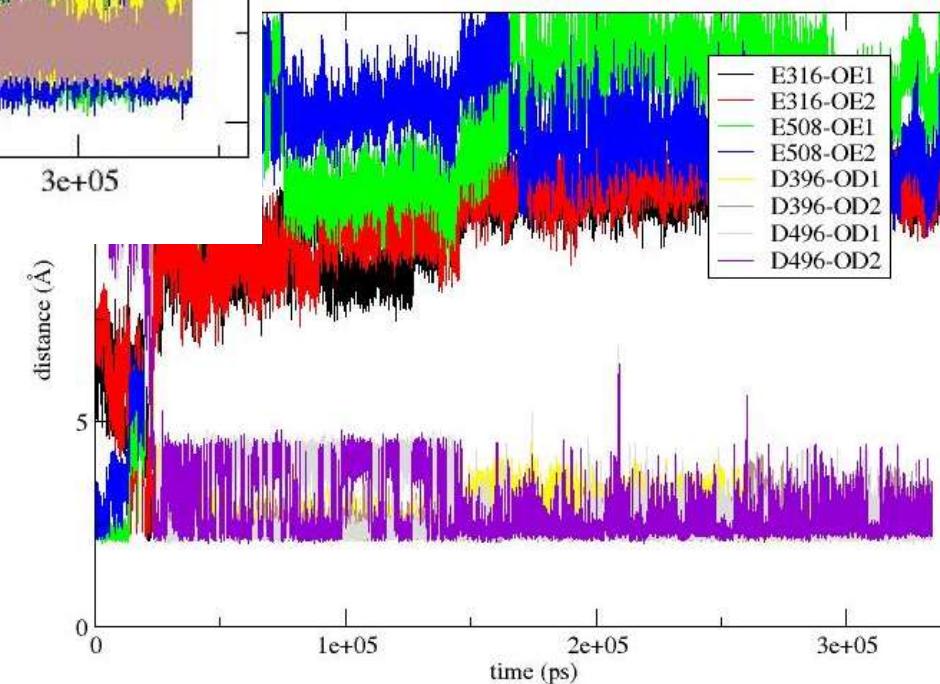
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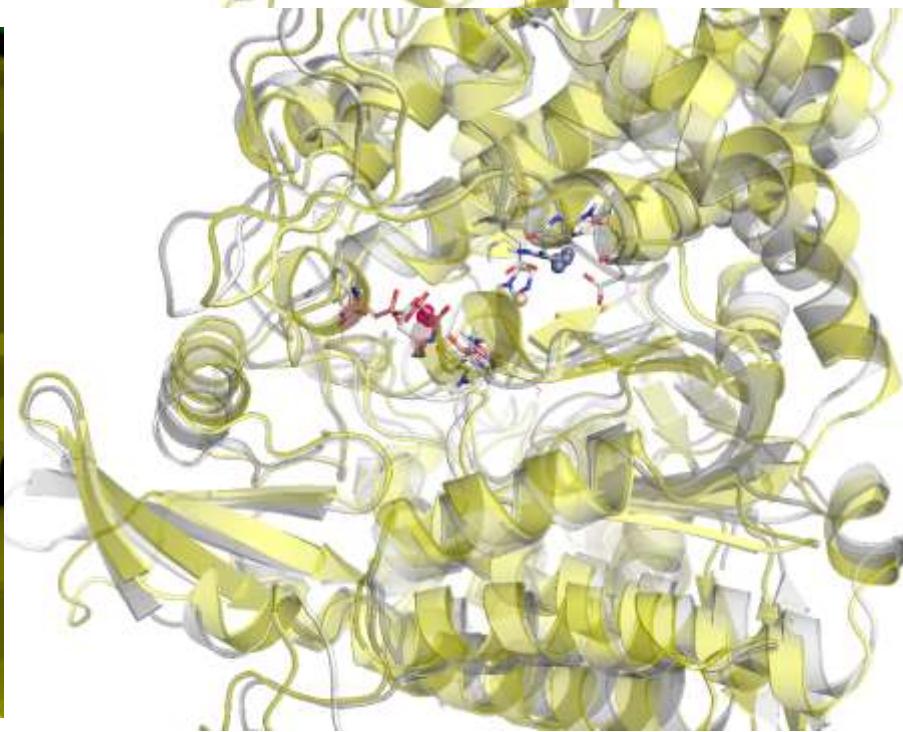
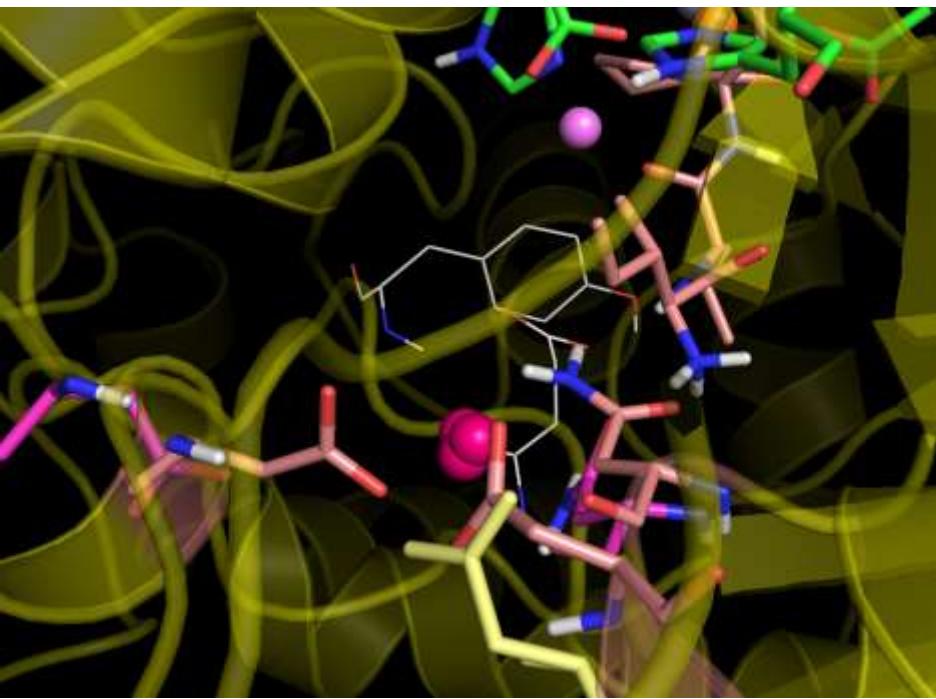
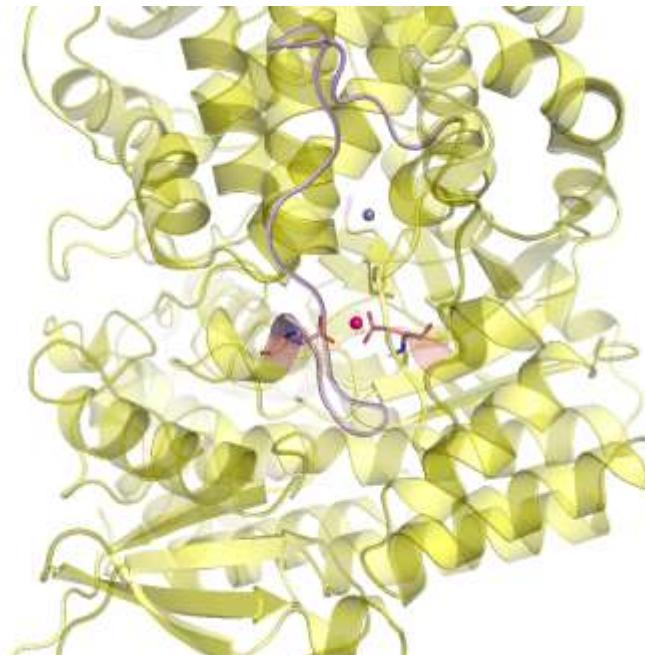
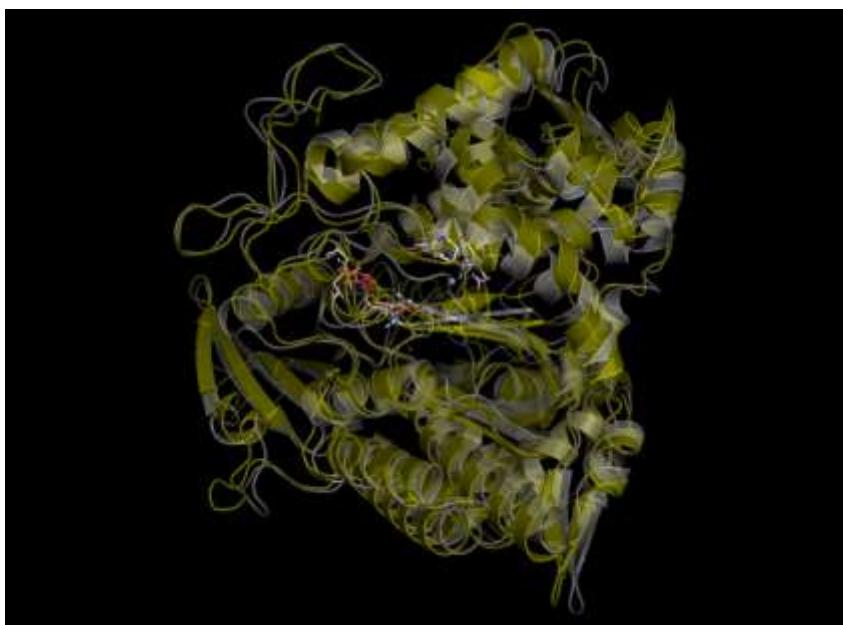
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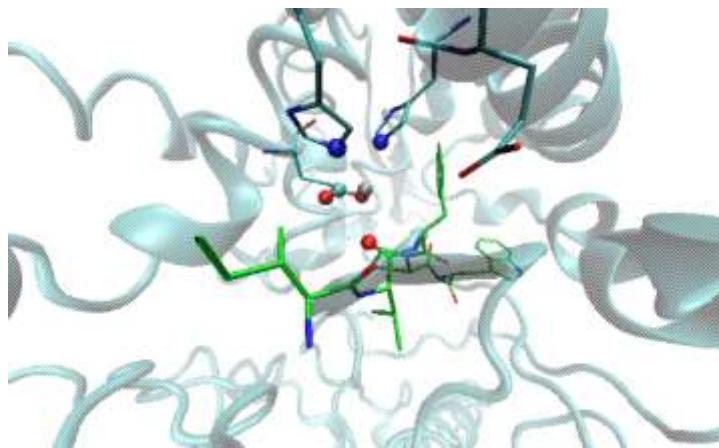
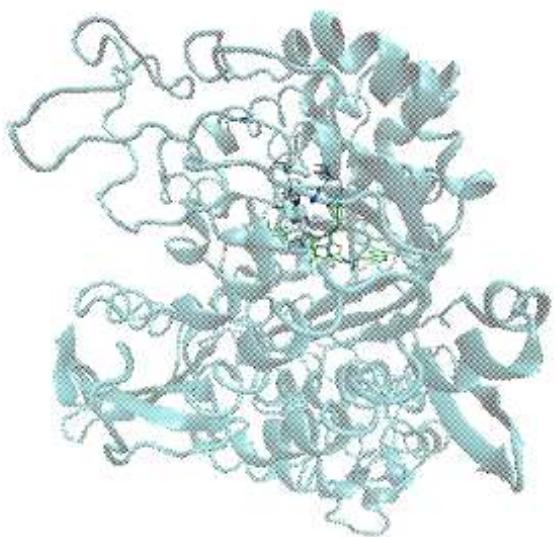
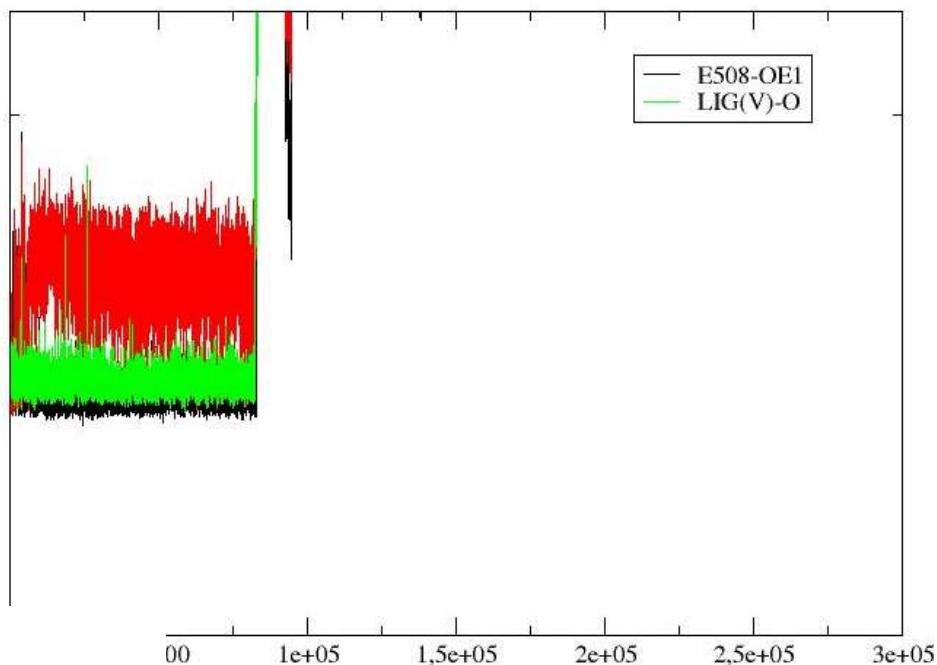
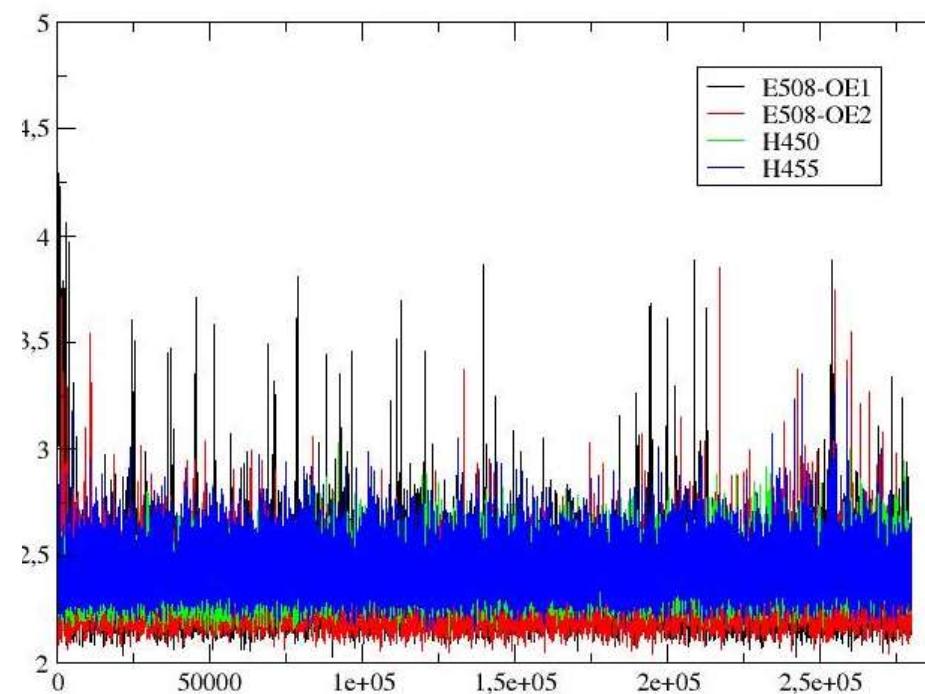
ZNI
CPLX2



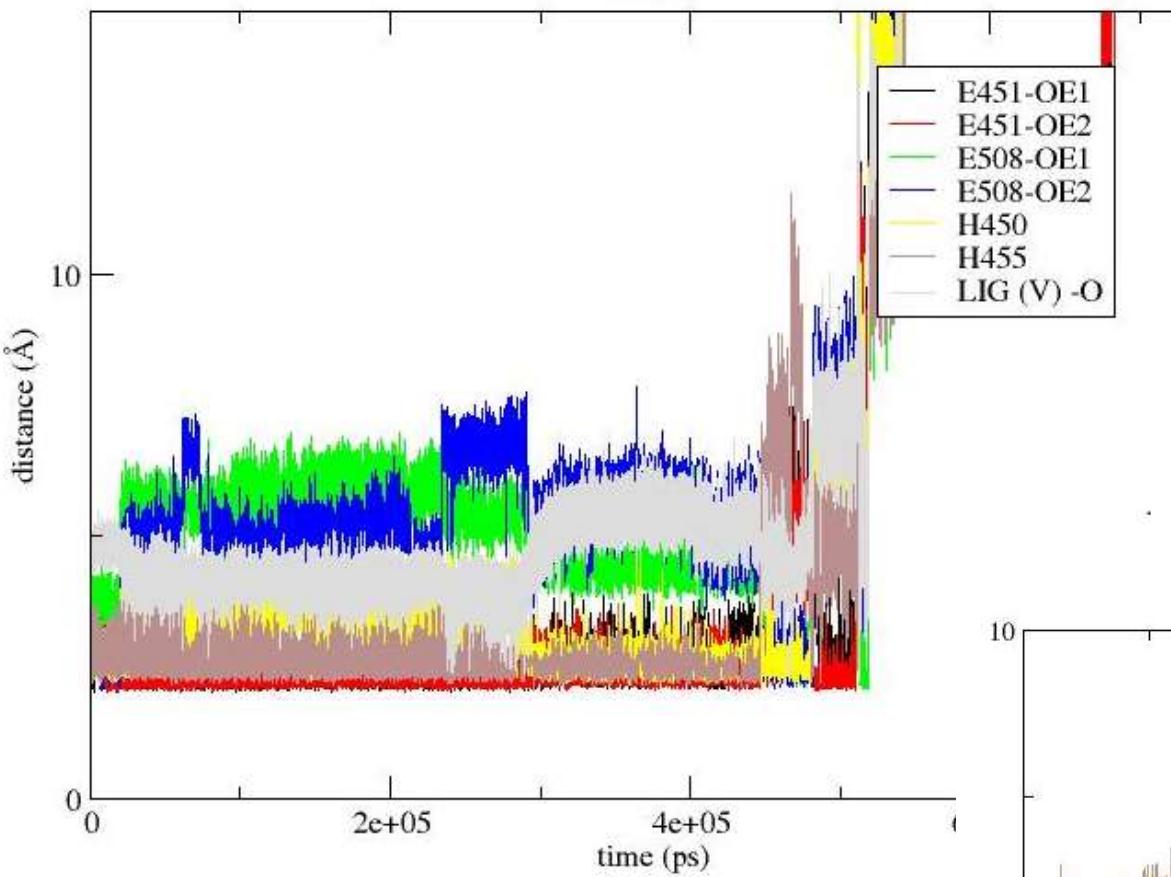
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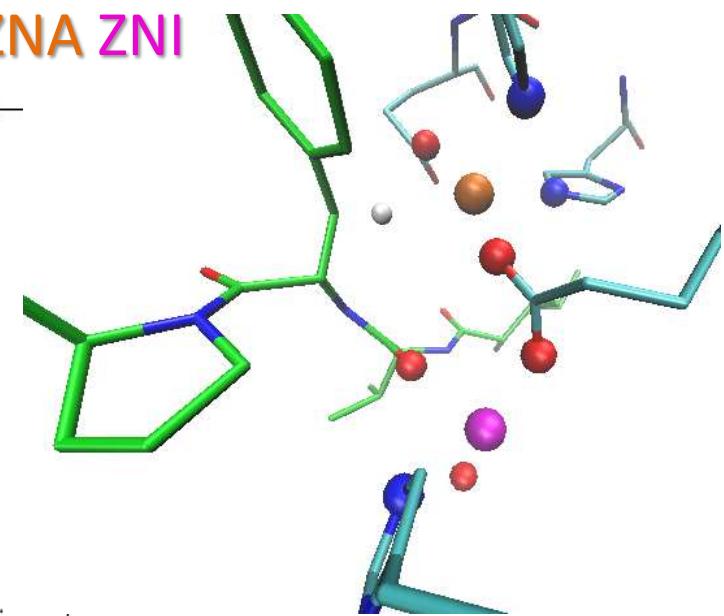
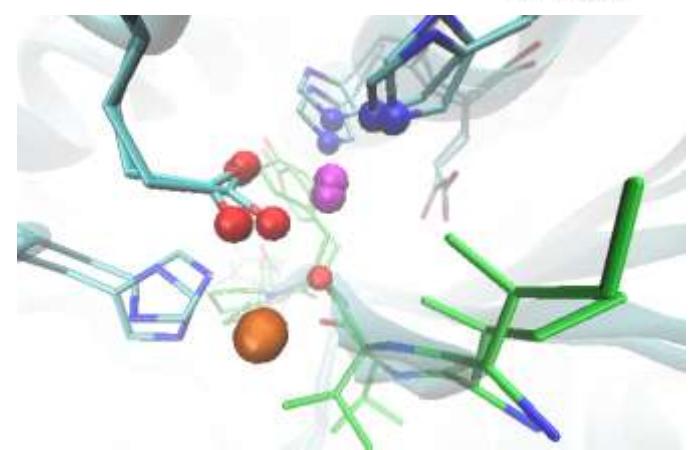
MD-CPLX2-m3



MD-CPLX2-m3r ZNA coodrnation CPLX2



IZMJENA ZNA ZNI



ZNI
CPLX2

