

# **Probing the Active Site of Thimet Oligopeptidase: A Study In Removing the Catalytic Metal**

## **Abstract**

Thimet Oligopeptidase(TOP, E.C. 3.4.24.15) is a metallopeptidase that cleaves various peptides involved in many physiological functions including the regulation of blood pressure and pain recognition. TOP is a zincin enzyme of clan MA<sup>1</sup> with the conserved HEXXH binding motif at the active site. Recent crystal structures have elucidated much of the structural information about TOP<sup>2</sup>; however, little is known about the specific coordinate geometry of Zn<sup>2+</sup> in the active site. In this study, the preparation of spectroscopically active metal substituted derivatives of TOP was attempted in order to better define this coordination and, if possible, extrapolate an explanation for the preferential binding of Zn<sup>2+</sup> as opposed to other divalent metals which have recently been discovered in the active sites of similar peptidases<sup>3</sup>. In addition, different metal substituted derivatives of TOP were prepared in order to determine the kinetic activity of said derivatives, with the intention of using this data to draw conclusions as to a “preferred” coordination of the metal.

## **Introduction**

Thimet Oligopeptidase(TOP, E.C. 3.4.24.15) is a zincin metalloendopeptidase which is ubiquitous through the body but is most highly concentrated in the brain and gonads. As a peptidase, its physiological function is the cleaving of short peptides(<17 AA)and plays in an important role in regulating blood pressure and has been implicated in the degradation of amyloid-β peptides which contribute to Alzheimer’s disease.<sup>4,5,6</sup> Most importantly for this study, TOP contains the HEXXH binding domain at the active site (see Fig. 1) which is homologous in many other zinc containing metalloenzymes. In the past, studies utilizing metal ion substitution showed that close homologues of TOP, such as thermolysin(E.C. 3.4.24.27 )which contains the HEXXH motif, actually become hyperactive with a cobalt substitution<sup>7</sup>. More recently other peptidases(including, interestingly, a carboxypeptidase) with the HEXXH binding motif have been discovered which natively utilize cobalt in their active site<sup>3,8</sup>. This raises questions as to why the majority of enzymes with this motif natively utilize zinc.

No spectrophotometric data can be gathered about native zinc enzymes, as  $Zn^{2+}$  has no absorbance in the visible region, therefore metal substituted derivatives play an important role in characterizing the coordination geometry of these zincin enzymes.

Spectrophotometric studies on metal substituted derivatives of peptidases similar to TOP have yielded interesting results. In a study of dipeptidyl peptidase III (DPP, E.C. 3.4.14.4), which contains the HEXXH motif, UV-vis showed that  $Co^{2+}$ -DPP III exhibited a pentacoordinate geometry. This is only slightly helpful however as the study did not observe coordination of the metal during catalysis. In a more thorough study of Thermolysin, an even closer homologue of TOP, the  $Co^{2+}$ -substituted form was observed to adopt a pentacoordinate geometry and was seen to exhibit flexibility in coordination during catalysis (i.e. adopting different coordinate geometries during catalysis).<sup>7</sup> According to the authors, the slight preference for a five-coordinate geometry, which is close to the transition state, along with the flexibility in coordination, explain the ability of  $Co^{2+}$ -thermolysin to exhibit hyperactivity.

In this study, metal substituted derivatives of TOP were produced and attempted to be analyzed by quenched fluorescence kinetic assays as well as U.V.-vis spectroscopy.

### **Materials and Methods**

Many methods were attempted in order to remove the divalent zinc ion from TOP; however, only one was successful. Still, it is illustrative to describe the unsuccessful methods in order to more thoroughly understand the nature of the  $Zn^{2+}$  as it is substituted in the enzyme. In addition, the method of substituting in new metals was refined as time went on. In all studies, TOP was prepared by Marc. J Glucksman of the Midwest Proteome Center and Department of Biochemistry and Molecular Biology, Rosalind Franklin University of Medicine and Science in Chicago, IL.

#### *Activity Assay*

Activity assay was performed by using a Perkin Elmer LS 50 B Luminescence Spectrometer 1980  $\mu$ L of buffer containing 25mM tris, 125mM KCl and 1mM TCEP was equilibrated with 10  $\mu$ L of MCA<sup>i</sup> substrate (of varying concentrations) at 23°C. Then, 10  $\mu$ L of .1  $\mu$ M enzyme were added to the chamber and fluorescence was detected for 150 seconds with an excitation of 325nm and an emission of 400nm. Note: For simple

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<sup>i</sup>7-methoxycoumarin-4-acetyl-Pro-Leu-Gly- Pro-Lys-dinitrophenol

binary tests of activity, enzyme concentration was noted(if possible) but not always .1 $\mu$ M. Often, a total of 2 $\mu$ mol of enzyme was present in the reaction chamber.

#### *Dialysis Against 2,6-Pyridinedicarboxylic Acid*

This method was adapted from a paper by Hirose et. al.<sup>9</sup> and involved the dialysis of the enzyme against 50mM tris buffer with 20mM 2,6 Pyridinedicarboxylic Acid(2,6-PCA) at pH 7.4 with various concentrations of Dithiolreitol(DTT) ranging from .5mM to 3mM. Dialysis was carried out at 4°C for 24 hours. After dialysis, the enzyme was concentrated using a Millipore centricon and activity was determined by the method described above.

#### *Dialysis Against Ethylenediaminetetraacetic Acid*

A similar method was attempted using Ethylenediaminetetraacetic Acid(EDTA). This was adopted from the method of Barrett and Brown<sup>10</sup> who worked with TOP. Using a dialysis buffer of 50mM tris, 3mM DTT and 100mM EDTA at pH 7.0. The dialysis was carried for approximately 24 hours at 4°C. The enzyme was then concentrated and activity was ascertained by quenched fluorescence assay by the method describe above.

#### *Denaturation with an EDTA Buffer*

A stock buffer containing 25mM tris, 50mM EDTA, 125mM KCl and 9M urea with 10% glycerol at pH 7.8 was prepared. At the time of use, solid DTT was added to this buffer to 5mM. 150 $\mu$ L of this buffer was then combined with 50 $\mu$ L of concentrated enzyme stock in a microfuge tube. The tube was allowed to sit at room temperature for approximately 1 hour and then was moved to a 4°C where it incubated for 18 hours. The enzyme was then purified of excess urea and EDTA by using an Amersham Biosciences PD-10 size exclusion column(PD-10). The new buffer which the enzyme was transferred to contained 25mM tris and 5mM DTT at pH 7.4. The enzyme concentration was obtained via -U.V. vis and then activity assay run using the same method described above. This procedure was also repeated substituting 50mM 2,6-PCA for EDTA.

#### *Incubation with 1,10-Phenanthroline*

A buffer containing 25mM tris and approximately 8mM 1,10-Phenanthroline was prepared. DTT was then added to 200 $\mu$ L of this buffer to 5mM. 150 $\mu$ L of the new buffer was then combined with 50 $\mu$ L of a concentrated enzyme stock and allowed to sit at room temperature for 1 hour and then transferred to a 4°C environment for 12 hours. The

sample was then loaded onto a PD-10 column and transferred to a buffer containing 25mM tris and 5mM DTT at pH 7.4. Activity assay was then run using same the method as described above. This method was also attempted with the chelating agents 2,6-PCA and EDTA.

#### *General Test for Loss of Metal*

If a sample of enzyme was observed to be inactive after a chelation or denaturation, loss of the catalytic metal was tested by taking an aliquot of the inactive sample and adding equimolar amounts of  $ZnCl_2$  to the enzyme. Activity assay was then performed on the sample with  $Zn^{2+}$  and without added  $Zn^{2+}$  at regular time intervals. If the activity of the sample without  $Zn^{2+}$  remained low, and the sample with  $Zn^{2+}$  regained activity, the sample was determined to be actual apo-enzyme.

#### *Substitution of Metals*

Metal substitution was attempted by three methods, dialysis, incubation and exchange of buffer via a PD-10 column.

Dialysis was performed by transferring the enzyme onto a dialysis chamber with a metal at .5mM, 50mM tris and 3mM DTT at pH 7.4. The sample was then concentrated. Incubation was performed by adding concentrated dissolved divalent metals to enzyme buffered solutions.(Note: inactivation of enzyme was seen with  $Zn^{2+}$  concentrations higher than 5x of the enzyme concentration). Finally, for the exchange of buffer method, metal buffers were prepared with .1mM metal, 50mM tris, 3mM TCEP, 125mM KCl at pH 7.0 were prepared. A PD-10 column was equilibrated to this buffer and the enzyme was passed down the column in order to exchange the buffer. (Note: DTT could not be used as it reduced the metals and precipitated them).

### **Results**

As noted above, not all of these methods for removal of the  $Zn^{2+}$  ion were successful. As such, the results have been divided between the successful and unsuccessful attempts.

#### *Unsuccessful Removal of the $Zn^{2+}$ Ion*

Dialysis of TOP against 2,6-PCA resulted in no decrease in activity of the enzyme. Initially, this was difficult to determine as 2,6-PCA obscured the 280nm protein peak(as 2,6-PCA absorbs very strongly at 271nm with a distinctive triple peak) and therefore made it difficult to find the concentration of the enzyme. This problem was overcome

using Low Pressure Chromatography and, later, simple PD-10 columns to separate out the excess 2,6-PCA

Dialysis for TOP against EDTA initially looked promising. After 24 hours of dialysis against 100mM EDTA, the TOP was completely inactive(0% activity of holoenzyme), however, upon an exchange of buffer which occurs as a result of the concentration process, the enzyme regained nearly full activity. To combat against contamination, two new samples were dialyzed under the same conditions to inactivation, then one was transferred directly to a plain buffer(50mM tris, .5mM DTT, pH 7.4) and the other to a buffer with metal(50mM tris , .5mM TCEP, .5mM CoCl<sub>2</sub> pH7.4). After 6 hours, the enzyme on the buffer without metal had regained activity to 10% of holoenzyme while the sample on the buffer with metal had regained only 2.6% activity.(See Fig. 2)

Incubation of the enzyme with any chelating agent(i.e. EDTA, 2,6-PCA and 1,10-Phenanthroline) was unsuccessful. Neither 2,6-PCA or EDTA inactivated the enzyme at all after 24 hours of incubation at 4°C. Incubation with 1,10-Phenanthroline did decrease the activity of the enzyme to 12% of the t=0 sample. Upon purification and addition of Zn<sup>2+</sup> to the solution, the enzyme did not regain any activity.

Finally, denaturation in the presence of high concentrations of 2,6-PCA was unsuccessful. Though the enzyme lost activity during the denaturation, it was regained as soon as the solution was purified of Urea.

#### *Successful Removal of the Zn<sup>2+</sup> Ion*

The procedure of denaturing the enzyme with urea in the presence of high concentrations of EDTA was the only one which exhibited success. After 16 hours of incubation at 4°C in the buffer described above, the enzyme was completely inactive. Upon purification with a PD-10 column the enzyme remained inactive. Two samples were then created, one with and one without Zn<sup>2+</sup> present. After only 4 hours of incubation at 4°C, the sample with Zn<sup>2+</sup> exhibited full activity, while the sample without was completely inactive(See Fig. 3). This result, however, was difficult to reproduce.

After apo-enzyme was confirmed to have been prepared, different metals were substituted. In this way, Co-TOP, Cu-TOP and Zn-TOP(i.e. TOP with Zn<sup>2+</sup> reintroduced to the active site after the loss of the metal) were prepared. After these derivatives were prepared, a full activity assay was run on each of them and the K<sub>M</sub> and K<sub>CAT</sub> determined

by using the Michaelis-Menton plots of the data. These methods yield the results listed in the table below and shown in Figs. 4-6. A graphical representation of this data may be found in Figures 7-8.

<b>Kinetic Parameters</b>	<b><u>WT-TOP</u><sup>ii</sup></b>	<b><u>WT-TOP (New)</u><sup>iii</sup></b>	<b><u>Co-TOP</u></b>	<b><u>Cu-TOP</u></b>	<b><u>Zn-TOP</u></b>
<b><math>K_{CAT}(s^{-1})</math></b>	.44	1.37	.12	.74	.38
<b><math>K_M(\mu M/s)</math></b>	7.88	37.67	3.27	12.65	10.64
<b><math>K_{CAT}/K_M(\mu M/s)</math></b>	.05	.036	.04	.06	.04

### **Discussion**

As is apparent from the results, many of the methods attempted in this study were unsuccessful, but still informative. Though we gained little information about the nature of the active site by the initial methods proposed, we did learn a great many ways in which we are unable to deplete TOP of its catalytic metal ion. In this way, the specific reason for each of these failures helps us to understand the nature of the active site from a different perspective.

#### *Problems with 2,6-PCA*

As shown above, 2,6-PCA was unable to chelate the metal in any situation. Though it was initially thought that 2,6-PCA would be a more potent chelating agent than EDTA, this seems to have not been the case. The fact that it did not even bind to metals in solution during a denaturation protocol raises questions as to the purity of the 2,6-PCA utilized in this study. Though it makes sense that 2,6-PCA may not be able to chelate to metal within TOP, it still should have bound to all free, divalent, metals dissolved in solution. Other explanations, however, are possible, such as metal contaminated buffer after purification through the PD-10 column.

#### *Inactivation but not Chelation with Dialysis Against EDTA*

One of the more confounding results deals with the inactivation of the enzyme when dialyzed against high concentrations of EDTA. Initially, it seemed that the EDTA was

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<sup>ii</sup> Courtesy of Jeffrey Sigman, Summer Research Program Mentor

<sup>iii</sup> Courtesy of Zhe Christopher Lin, Summer Research Program Participant

able to chelate the metal, and thus, produce apo-TOP. The experiment in which TOP was dialyzed against EDTA and then immediately moved to a dialysis chamber which contained either a “plain” buffer or a metal buffer was inconclusive as both regained activity. The fact that the sample on a “plain” buffer regained activity more quickly is most likely because high concentrations of metal often inactivate enzymes, leading to a more active enzyme on “plain” buffer. A final experiment was performed in which TOP was dialyzed against 100mM EDTA and then concentrated in a buffer with 10 $\mu$ M EDTA, a sufficient concentration to get rid of most trace metals in the water. This sample regained activity to 100% of the holoenzyme.

These results lead to the conclusion that EDTA inactivates the enzyme, but does not chelate the metal. The mechanism by which this happens is most likely solvation of the active site. In past experiments, Sigman observed, anecdotally, that high concentrations of DTT would inactivate the enzyme. This is due to the crowding of the active site by a large number of DTT molecules. As DTT and EDTA have similar structures, it is likely that much the same thing is happening. Also, this is consistent with the lack of inactivation seen when the enzyme was dialyzed against 2,6-PCA. Where DTT and EDTA have structure such that they may solvate the active site, 2,6-PCA has more of a winged structure which may coordinate with the metal by a “closing of its wings” over the metal, but ultimately, block other molecules from getting too close.

#### *Ineffectiveness of Incubation with Chelating Agents*

In all experiments in which TOP was incubated with various chelating agents(EDTA, 2,6-PCA, and 1,10-Phenanthroline) inactivation did not occur at all. This is intriguing because of the results seen with dialysis of EDTA. This could be because, during dialysis, as EDTA solvates the active site of TOP, the concentration gradient changed and more EDTA moved into the enzyme chamber, making the concentration around the enzyme much greater than in an incubation chamber, in which the concentration remains constant. In addition, concentrations of EDTA used in incubation experiments were 20mM, which may not have been high enough to cause inactivation.

Also interesting was the loss of activity when TOP was incubated with 1,10-phenanthroline. As it did not regain activity with incubation with Zn<sup>2+</sup>, this could be because the high concentration of 1,10-phenanthroline (8mM, which is saturation in

water) actually damaged the enzyme, inactivating it completely, instead of removing the metal from TOP.

#### *Problems with 2,6-PCA*

2,6-PCA failed to remove the metal from TOP by all methods attempted, dialysis, incubation and incubation with the denaturant urea. This is a very odd result as in other metallopeptidases, 2,6-PCA was utilized quite successfully to chelate the metal ion by simple dialysis<sup>9</sup>. Most frustrating was the fact that 2,6-PCA seemed incapable of binding to the metal even when it had “fallen-out” of TOP during denaturation. Because of this, two possibilities arise. First, the 2,6-PCA, prepared by Sigma-Aldrich, was impure. Alternatively, the enzyme may have a higher affinity for the metal than the 2,6-PCA. The latter possibility will be discussed further later.

#### *Successful Preparation of apo-TOP by Denaturation in the Presence of EDTA*

As stated earlier, the only method by which apo-TOP was created was by denaturation in the presence of EDTA. Combined with all of the other results obtained, a fairly clear mechanistic picture of what occurred may be described. The denaturation by urea unfolded the enzyme enough for the metal to be lost by TOP. Next, the EDTA solvated the newly released  $Zn^{2+}$ , making it impossible for TOP to re-acquire the metal. Finally, when the solution was purified down a PD-10 column, the apo-TOP was cleansed of urea, allowing it to refold without a metal present. The idea that the EDTA solvated the metal instead of simply binding to it is important, because it bypasses all problems that may occur because TOP has a high affinity for  $Zn^{2+}$  than EDTA.

#### *Kinetic Parameters of Metal Substituted TOP*

Co-TOP, Cu-TOP and Zn-Top(i.e. apo-TOP which then had  $Zn^{2+}$  added to its active site) were prepared. All of these metal derivatives had overall activities(i.e.  $K_{CAT}/K_M$ ) that were very similar to the holo enzyme with WT-TOP=.05 $\mu$ M/s, WT-TOP(New)=.036 $\mu$ M/s, Co-TOP= .04 $\mu$ M/s, Cu-TOP=.06 $\mu$ M/s, and Zn-TOP=.04 $\mu$ M/s. The fact that Cu-TOP is the most active is not altogether surprising as in it's closest homologue, thermolysin, the  $Cu^{2+}$  substituted form is most active<sup>7</sup>. In addition, this result is in direct contradiction to Barrett and Brown<sup>7</sup> who saw very slight activation with  $Co^{2+}$  and no activation with  $Cu^{2+}$ .<sup>10</sup> The reasons for this will be discussed more thoroughly in the conclusion.

While all of the metal derivatives had similar overall activities, their specific kinetic parameters were very different. Co-TOP( $K_M=3.27\mu\text{M/s}$ ) had a much higher binding affinity than Cu-TOP and Zn-TOP( $K_M=12.65\mu\text{M/s}$  and  $10.64\mu\text{M/s}$  respectively). This result is difficult to interpret but may be due to different coordinate geometries of the metals. In addition,  $K_{CAT}$  for Cu-TOP( $.74\text{s}^{-1}$ ) was nearly 5 times larger than Co-TOP( $.12\text{s}^{-1}$ ) and nearly twice as large as Zn-TOP( $.38\text{s}^{-1}$ ). As Cu-TOP has the highest overall kinetic rate this result, along with its low substrate affinity, probably mean that Cu-TOP binds weakly (relative to other metal substituted derivatives) to the substrate and but acts very quickly.

In addition, it is of interest that the two WT-TOP samples are so different. The kinetic parameters for WT-TOP were measured quite some time ago, while the parameters for WT-TOP(New) were measured for this specific batch of enzyme. The much lower binding affinity seen for WT-TOP(New) is important because it shows that the binding affinities for all three of the metal substituted enzymes were very much higher than that of the WT from which they were prepared. In addition, the  $K_{CAT}$  for WT-TOP(New) is an order of magnitude larger than any of the metal substituted samples. This means that the WT enzyme(for this batch of enzyme) has a much higher turnover rate than that of the metal substituted derivatives (and the older WT-TOP sample), but the derivatives more than compensate for this with much higher binding affinities. This difference seems to imply some sort of change in catalytic mechanism that occurs as a result of metal substitution, as the  $K_M$  for Zn-TOP is so different from that for WT-TOP(New). Additionally, the difference could also be caused by a structural change at the active site due to re-uptake of metal.

### **Conclusion**

The amount of difficulty experienced in preparing apo-TOP is, at this point, apparent. Many different methods were attempted and only a very few were successful. One of the greatest confounding factors in the process may have been an impure water system. Since such low concentrations of TOP were used, even a slight contamination with any one of a myriad of divalent metal ions could have caused the enzyme to reactivate during the denaturation protocols.

In addition to this problem, there is a good probability that TOP has a high affinity for metal. This can be seen in the complete inefficacy of 2,6-PCA to chelate the metal. In addition, when Sigman was preparing a denaturation curve, which was measured using a chelating agent which has a color shift between the bound and unbound forms (i.e. once the  $Zn^{2+}$  has “fallen out” of TOP due to unfolding, the chelating agent would bind to the metal and change color) he experienced a great amount of difficulty in achieving consistent results. One of the reasons for this may have been this high affinity for metal by the TOP, meaning that TOP could outcompete the chelating agent, even when partially unfolded. Another explanation is water that was contaminated with metals. This would change the total amount of metal in solution and give the chelating agent ions to bind to which were not from TOP. This second explanation also was a likely confounding factor in this experiment.

After attempting multiple times, the methods of Barrett and Brown<sup>10</sup>, it can be fairly well established that they were not able to produce apo-TOP. The reactivation they saw may have simply been a result of which metals inhibited the enzyme least in solution, as all of their activity assays were carried out with metals in solution<sup>10</sup>. Further work must be done to confirm this. An experiment with a well-studied metalloprotease (e.g. Carboxypeptidase A) would allow for the testing of all methods in this paper. If the *method*, described by Barrett and Brown and replicated in this paper, is successful then the *results* of Barrett and Brown must be inaccurate. That is, if the method leads to a preparation of apo-Carboxypeptidase A, then it will be known that apo-TOP cannot be prepared by this procedure for some reason other than a methodological error.

Other further studies that might be attempted are the repetition of all chelation and denaturation protocols with more pure water. This would eliminate any need problems that may be associated with metal contamination of the water. In addition, it would be beneficial to submit a portion of the metal substituted TOP for mass spectroscopy or elemental analysis, in order to confirm that the substitution worked, as all of the catalytic data is relatively similar. Finally, after a consistent method is discovered and more metal substituted TOP is prepared, it will be beneficial to do more activity assays on Co, Cu, and Zn-TOP in order to verify the results presented in this paper. Also, U.V. –visible light spectrophotometry can be used on this newly prepared Co-TOP in order to

determine the conformation of ligands around the metal ion, possibly revealing why the metal is so hard to chelate. Finally, UV-visible spectra of bound forms of TOP (i.e. TOP which has MCA in the active site and has undergone a conformational change) should be obtained in order to study the coordination of the metal during catalysis.

**Figure 1:**

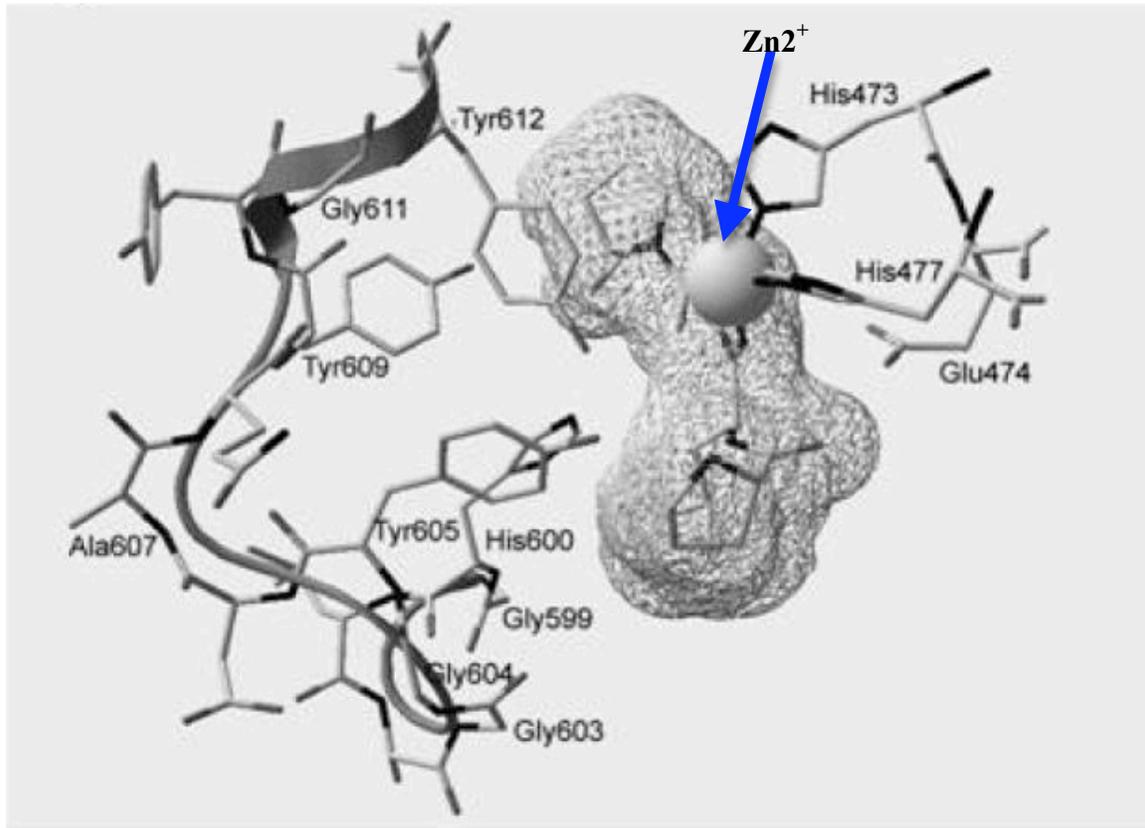
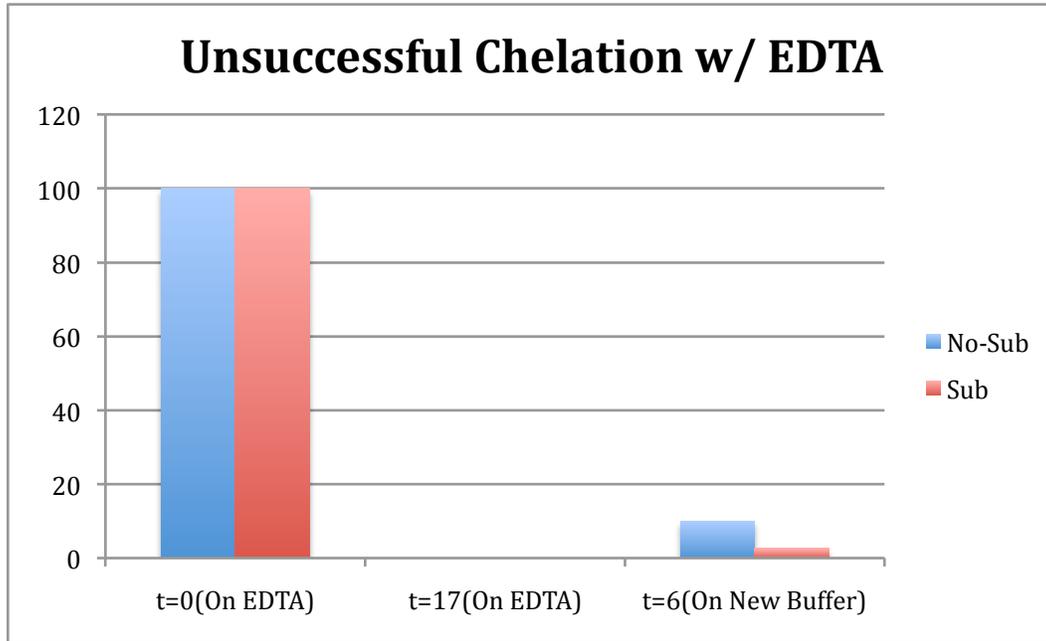
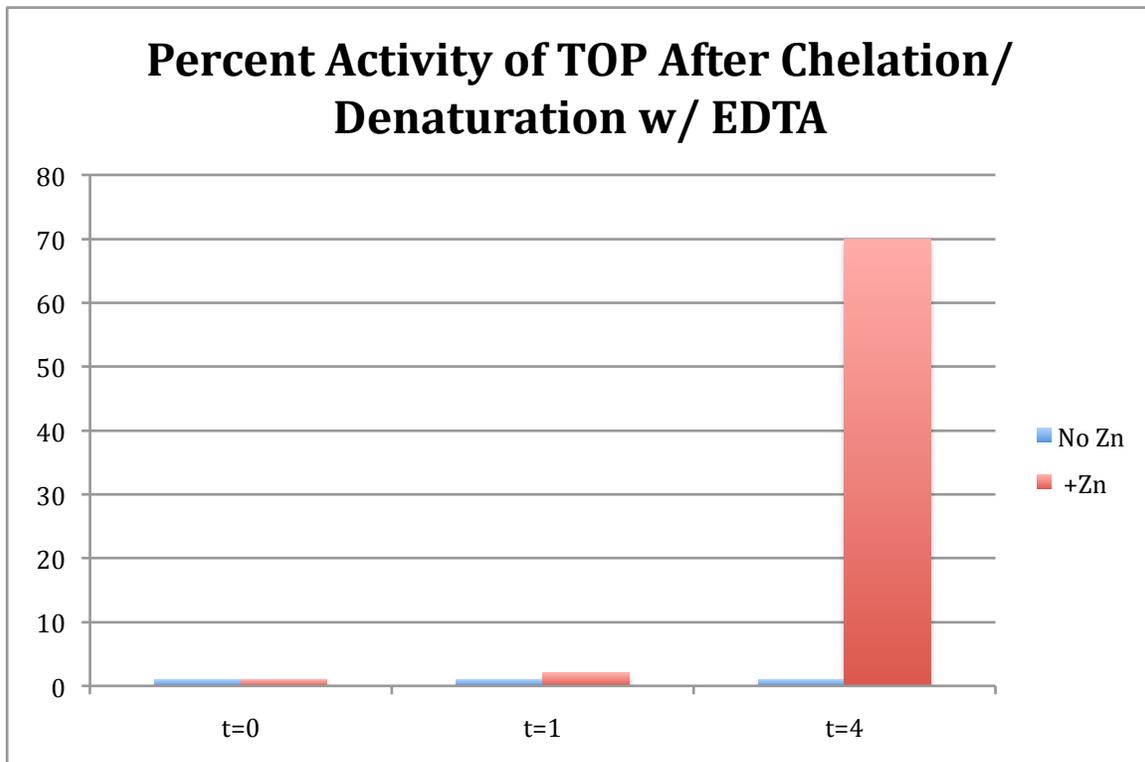


Figure Credit: Bruce L, Sigman J, Randall D, Rodriguez S, Song M, Dai Y, Elmore D, Pabon A, Glucksman M, Wolfson A (2008) Hydrogen bond residue positioning in the 599-611 loop of thimet oligopeptidase is required for substrate selection *FEBS J* 5607–5617 doi: 10.1111/j.1742-4658.2008.06685.x

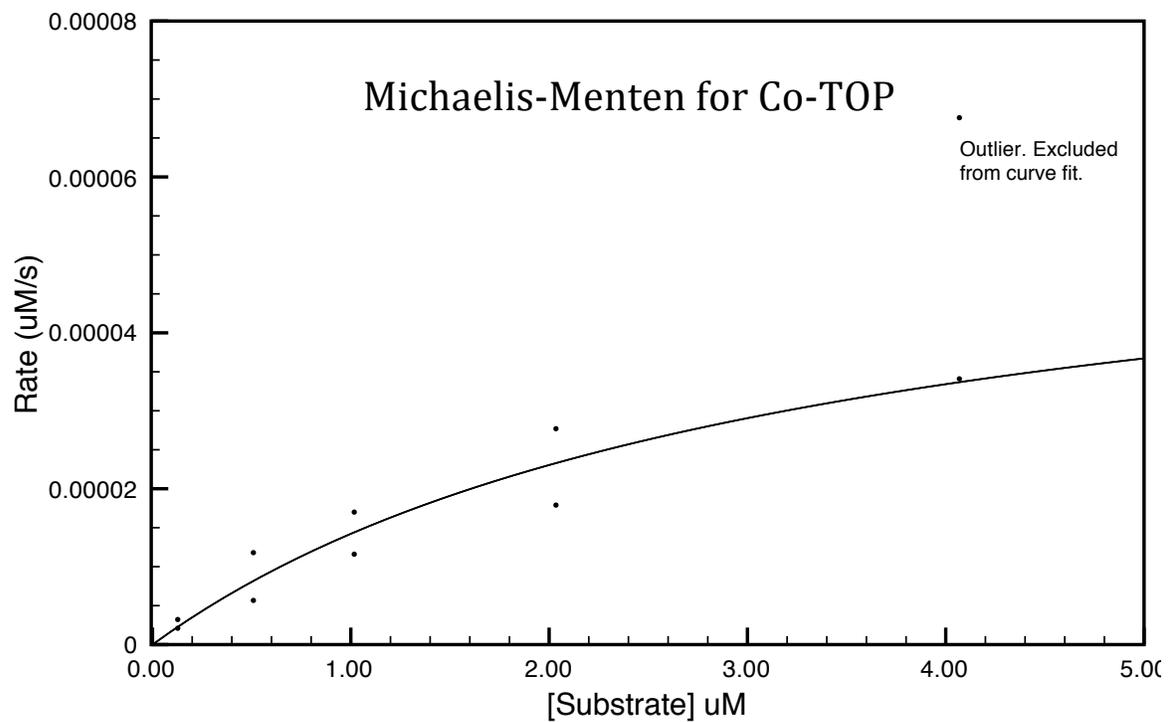
**Figure 2:**



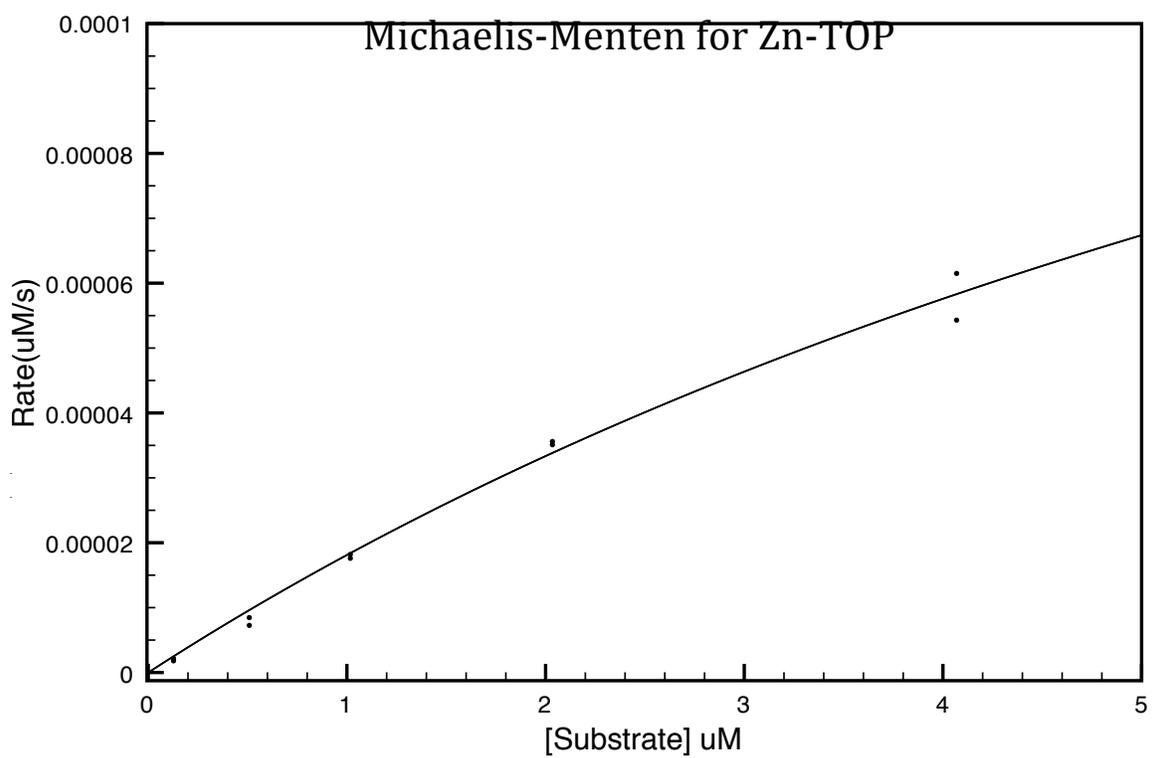
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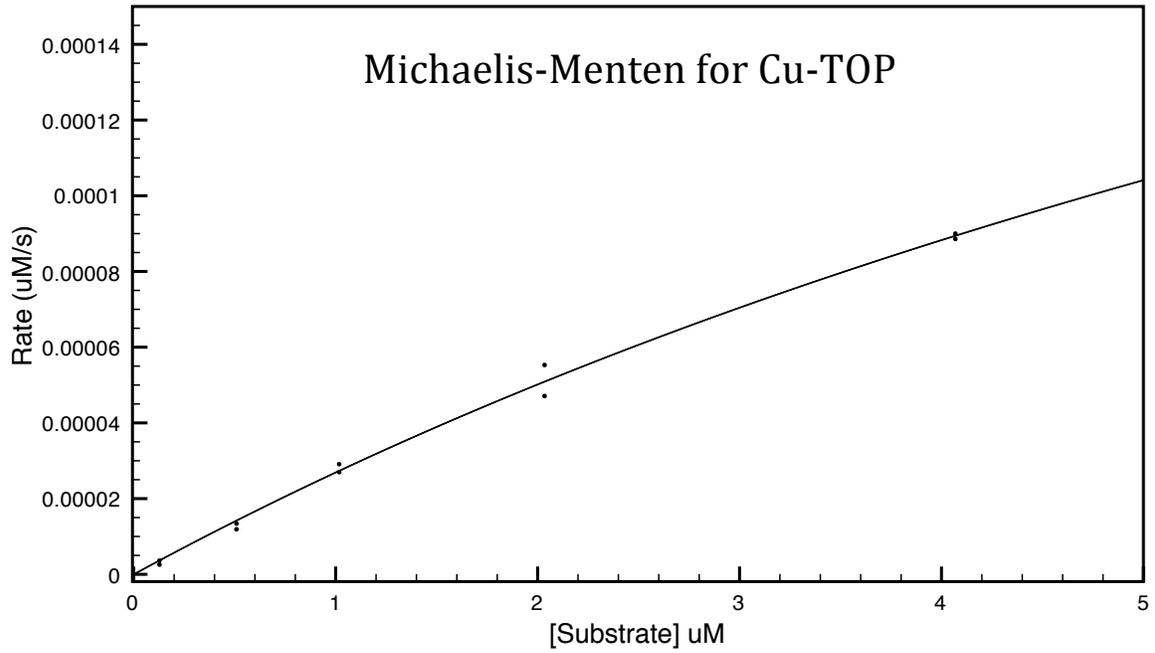
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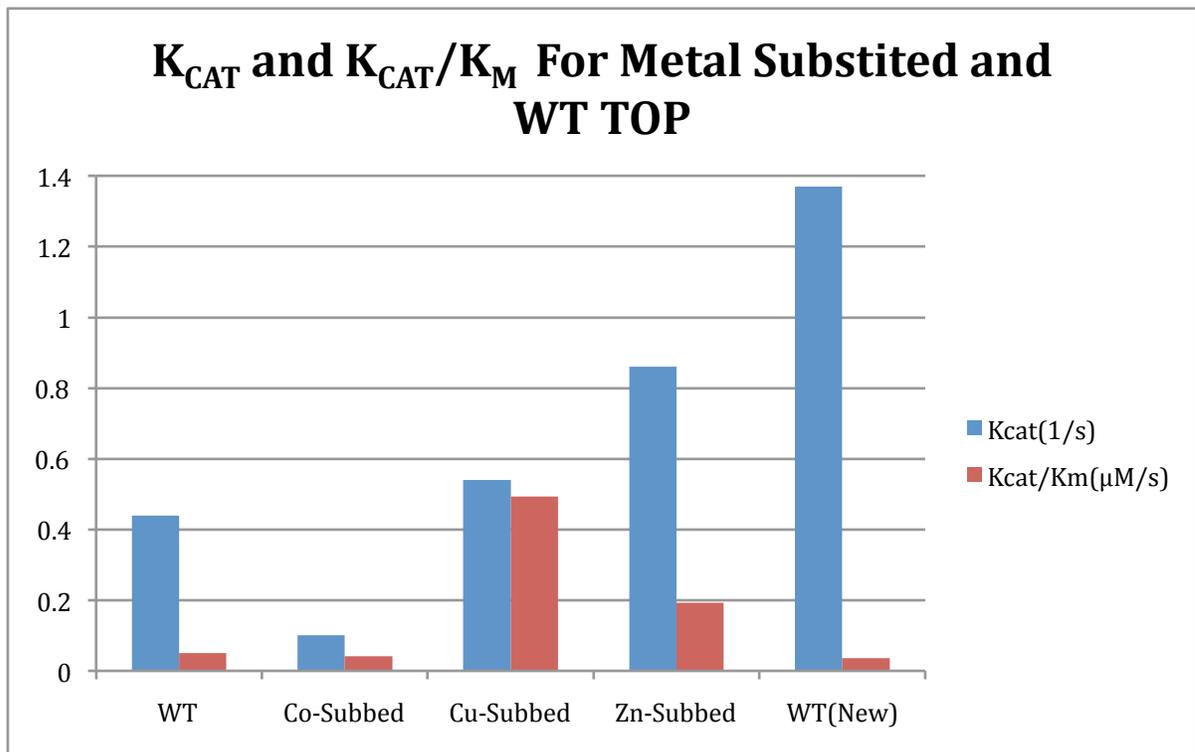
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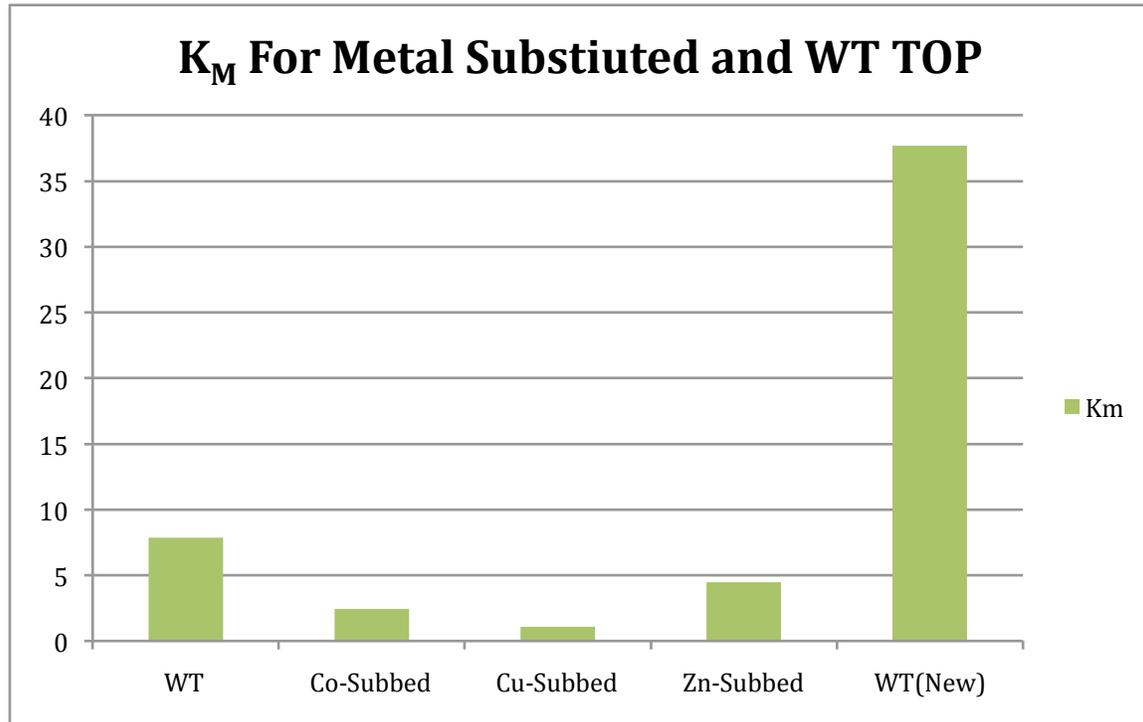
**Figure 6:**



**Figure 7:**



**Figure 8:**



<sup>1</sup> Rawlings ND & Barrett AJ (1995) EVOLUTIONARY FAMILIES OF METALLOPEPTIDASES. In *Proteolytic Enzymes: Aspartic and Metallo Peptidases*, pp. 183-228. Academic Press Inc, San Diego.

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