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# Kinetic and spectroscopic investigation of $Co^{II}$ , $Ni^{II}$ , and N-oxalylglycine inhibition of the $Fe^{II}/\alpha$ -ketoglutarate dioxygenase, TauD

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#### Abstract

 $Co^{II}$ ,  $Ni^{II}$ , and N-oxalylglycine (NOG) are well-known inhibitors of  $Fe^{II}/\alpha$ -ketoglutarate ( $\alpha KG$ )-dependent hydroxylases, but few studies describe their kinetics and no spectroscopic investigations have been reported. Using taurine/ $\alpha KG$  dioxygenase (TauD) as a paradigm for this enzyme family, time-dependent inhibition assays showed that  $Co^{II}$  and  $Ni^{II}$  follow slow-binding inhibition kinetics. Whereas  $Ni^{II}$ -substituted TauD was non-chromophoric, spectroscopic studies of the  $Co^{II}$ -substituted enzyme revealed a six-coordinate site (protein alone or with  $\alpha KG$ ) that became five-coordinate upon taurine addition. The  $Co^{II}$  spectrum was not perturbed by a series of anions or oxidants, suggesting the  $Co^{II}$  is inaccessible and could be used to stabilize the protein. NOG competed weakly ( $K_i \sim 290 \, \mu M$ ) with  $\alpha KG$  for binding to TauD, with the increased electron density of NOG yielding electronic transitions for NOG-Fe<sup>II</sup>-TauD and taurine-NOG-Fe<sup>II</sup>-TauD at 380 nm ( $\epsilon_{380}$  90–105  $M^{-1}$  cm<sup>-1</sup>). The spectra of the NOG-bound TauD species did not change significantly upon oxygen exposure, arguing against the formation of an oxygen-bound state mimicking an early intermediate in catalysis. © 2005 Elsevier Inc. All rights reserved.

Keywords: α-Ketoglutarate; Oxoglutarate; Ferrous; Hydroxylase; Dioxygenase; Metal-substituted; Spectroscopy; Inhibition

Fe<sup>II</sup>/ $\alpha$ -ketoglutarate ( $\alpha$ KG)-dependent dioxygenases couple the oxidative decarboxylation of  $\alpha$ KG to the oxidation of their primary substrates using a mononuclear, nonheme iron metallocenter [1]. These enzymes catalyze a wide variety of crucial chemical transformations including the repair of alkylation damage in DNA or RNA [2,3], sensing of hypoxia [4], modification of structural proteins [5], synthesis of various metabolites ranging from antibiotics [6] to plant hormones [7,8], and degradation of compounds such as herbicides [9] and phytanic acids [10]. The focus of this study is the archetype member of this enzyme family, *Escherichia coli* taurine/ $\alpha$ KG dioxygenase (TauD) that metabo-

lizes aminoethanesulfonate (Scheme 1) to produce sulfite as a cellular sulfur source [11].

Recent crystallographic and spectroscopic studies of TauD have confirmed many aspects of the general enzyme mechanism of Fe<sup>II</sup>/ $\alpha$ KG-dependent dioxygenases (Scheme 2) that was first proposed over two decades ago [12]. Structural studies [13,14] reveal that the Fe<sup>II</sup> center is ligated by three amino acid side chains on the same face of the metal: His99, Asp101, and His255. In the absence of substrate, three water molecules complete the six-coordinate environment (A). Two waters are displaced upon binding of  $\alpha$ KG (shown as RCOCOO<sup>-</sup>), which coordinates the Fe<sup>II</sup> center through its C-1 carboxylate and C-2 carbonyl moieties (B) producing a diagnostic metal-to-ligand charge-transfer transition with a  $\lambda_{\rm max}$  at 530 nm and  $\epsilon_{530}$  of 140–240 M<sup>-1</sup> cm<sup>-1</sup> [15,16]. Taurine (illustrated by R'—H) binds near the active site and promotes dissociation of the

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remaining water ligand to shift the  $\lambda_{max}$  to 520 nm with  $\epsilon_{520} = 180\text{--}270~\text{M}^{-1}~\text{cm}^{-1}~[15,16]$ , leaving the Fe<sup>II</sup> five-coordinate and primed to react with oxygen (C). Binding of oxygen produces a yet uncharacterized Fe<sup>III</sup>-superoxo or Fe<sup>IV</sup>-peroxo species (D) that attacks the  $\alpha$ KG carbonyl group, leading to decomposition of  $\alpha$ KG and heterolytic O–O bond cleavage. The resulting Fe<sup>IV</sup>-oxo species (E) inserts oxygen into the target C–H bond of the substrate by hydrogen atom transfer and oxygen rebound, as found in heme-type oxygenases, to restore the Fe<sup>II</sup> state of the enzyme. The Fe<sup>IV</sup>-oxo species has been identified on the basis of stopped-flow UV/visible spectroscopy, freeze-quench Mössbauer analyses, EPR spectroscopy of cryoreduced sample, cryogenic continuous-flow resonance Raman studies, and X-ray absorption spectroscopy [16–21].

Here, we examine the kinetics and spectroscopy of TauD interaction with three well-known inhibitors of this class of enzymes.  $\text{Co}^{\text{II}}$  and  $\text{Ni}^{\text{II}}$  inhibit prolyl and asparaginyl hydroxylases that target the hypoxia inducible factor (HIF), involved in oxygen sensing [22], so the metal ions lead to cellular gene expression changes mimicking those observed during hypoxia [23,24]. In addition,  $\text{Co}^{\text{II}}$  inhibits several other enzyme family members including TauD [11] and 1-aminocyclopropanecarboxylate oxidase, for which the Co-bound crystal structure is known [25]. Despite the importance of these inhibitory metal ions to the functioning of  $\text{Fe}^{\text{II}}/\alpha \text{KG}$ -dependent dioxygenases, few kinetic characterization studies and no spectroscopic investigations have been reported. Similarly, the  $\alpha \text{KG}$  analogue N-oxalylglycine (NOG) is an established inhibitor of procollagen

prolyl 4-hydroxylase [26,27] as well as HIF-specific prolyl and asparaginyl hydroxylases [28–30]. In the asparaginyl hydroxylase known as factor inhibiting HIF (FIH), the structure of the NOG-bound enzyme was determined and NOG was found to coordinate Fe in the same manner as the aKG shown in B of Scheme 2 [31]. Furthermore, inhibition of the iron-mediated degradation of the iron regulatory protein 2 by dimethyl-NOG (which is hydrolyzed to NOG by esterases within the cell) was cited as evidence for the participation of an Fe<sup>II</sup>/ $\alpha$ KG-dependent dioxygenase in this pathway [32,33]. As with the inhibitory metals, few NOG-related studies address the kinetics of inhibition and none examine the spectroscopic properties of the inhibited enzyme. We have chosen to study these aspects of Co<sup>II</sup>, Ni<sup>II</sup>, and NOG interaction with TauD as a paradigm for related systems. In addition, we sought to obtain new insights into the structures and properties of the early intermediates in TauD catalysis.

### Materials and methods

Purification of TauD apoprotein. Cultures of E. coli BL21(DE3) with pME4141 [11], containing tauD under the control of the T7 RNA polymerase promoter, were grown in TB medium (1 L) at 37 °C with stirring at 200 rpm. When cultures reached an  $A_{600}$  of  $\sim$ 0.4, tauD expression was induced by the addition of isopropyl-β-D-thiogalactopyranoside to a final concentration of 0.1 mM. Cells were harvested after ~12 h by centrifugation for 10 min at 10,000g and 4 ° C. The cell pellet was resuspended in 20 mL of 20 mM Tris with 1 mM EDTA (TE) buffer (pH 8.0). The sample was stored at -80 °C until disrupting the cells by using a French pressure apparatus. The lysate was clarified by centrifugation for 40 min at 150,000g and applied to a DEAE-Sepharose column (5.0 × 30 cm, Amersham Biosciences). The column was rinsed with two column volumes of TE buffer (pH 8.0) and eluted by using a linear 1500 mL gradient from 70 to 230 mM NaCl in the same buffer at a flow rate of 7 mL/min. Fractions containing TauD were concentrated in an Amicon stirred cell concentrator with a 30 kDa cutoff membrane. The concentrated eluant was loaded onto a high-performance phenyl-Sepharose column ( $2.5 \times 30 \, \text{cm}$ , Amersham Biosciences) equilibrated with TE buffer (pH 8.0) containing 500 mM (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub>. Proteins were eluted with a 1200 mL linear gradient from 500 to 0 mM (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub> in the same buffer at a flow rate of 5 mL/min. The TauD-containing fractions were concentrated and extensively dialyzed

$$\mathbf{B} \xrightarrow{\text{Asp.}} Fe(|||) \xrightarrow{\text{R}} H_{20}$$

$$\mathbf{A} \xrightarrow{\text{Asp.}} Fe(|||) \xrightarrow{\text{His}} H_{20}$$

$$\mathbf{A} \xrightarrow{\text{Asp.}} Fe(||||) \xrightarrow{\text{His}} H_{20}$$

$$\mathbf{A} \xrightarrow{\text{Asp.}} Fe(||||) \xrightarrow{\text{His}} H_{20}$$

$$\mathbf{B} \xrightarrow{\text{Asp.}} Fe(||||) \xrightarrow{\text{His}} H_{20}$$

$$\mathbf{C} \xrightarrow{\text{Asp.}} Fe(||||) \xrightarrow{\text{His}} H_{20}$$

$$\mathbf{R} \xrightarrow{\text{His}} Fe(||||) \xrightarrow{\text{His}} H_{10}$$

$$\mathbf{R} \xrightarrow{\text{Coo}_{2}}$$

$$\mathbf{R} \xrightarrow{\text{Asp.}} Fe(||||) \xrightarrow{\text{His}} H_{10}$$

$$\mathbf{R} \xrightarrow{\text{Coo}_{2}}$$

$$\mathbf{R} \xrightarrow{\text{Asp.}} Fe(||||) \xrightarrow{\text{His}} H_{10}$$

$$\mathbf{R} \xrightarrow{\text{Coo}_{2}}$$

Scheme 2.

against 25 mM Tris buffer (pH 8.0) at 4 °C. Purified TauD apoprotein exhibited a single 32.2 kDa band when examined by denaturing polyacrylamide gel electrophoresis. Protein concentrations were estimated by using  $\varepsilon_{280}$  46,400 M<sup>-1</sup> cm<sup>-1</sup>. The dialyzed apoprotein was stored frozen at -80 °C.

Enzyme assays. TauD activity was measured by using Ellman's reagent to quantify sulfite, as previously described [11]. One unit (U) of enzyme activity is defined as the amount of enzyme that releases 1  $\mu$ mol of sulfite per minute at 30 °C in assay buffer containing 25 mM Tris (pH 8.0), 50 or 100  $\mu$ M Fe<sup>II</sup>, 50 or 100  $\mu$ M ascorbate, 100 or 500  $\mu$ M  $\alpha$ KG, and 1 mM taurine. The TauD used in these studies had a specific activity ranging from 3 to 6.4 U (mg of protein) $^{-1}$ .

Characterization of inhibition kinetics. Steady-state inhibition assays were carried out during 5 min incubations using the typical assay conditions (sometimes with one component varied in concentration) and amended with the indicated concentrations of inhibitor. The assays were initiated either by adding enzyme or by adding Fe<sup>II</sup> to the assay buffers. In addition, the time dependence of inhibition was assessed in some cases by removing timed aliquots to an EDTA quench solution.

Electronic spectroscopy. Spectra were recorded at room temperature on a Shimadzu UV-2401 UV/visible spectrophotometer. All stock solutions for anaerobic binding studies were prepared inside serum vials sealed with butyl rubber stoppers and purged of oxygen by several rounds of vacuum degassing and flushing with argon using a vacuum manifold. Stock solutions of  $\alpha KG$  (50 or 100 mM), taurine (50 or 100 mM), and NOG (100 mM) were prepared in 25 mM Tris buffer (pH 8.0). Ferrous ammonium sulfate stock solutions (25 mM) containing 5 mM ascorbate were prepared by several rounds of degassing and argon flushing of the solids, followed by addition of the desired volume of H<sub>2</sub>O. CoCl<sub>2</sub> and NiCl<sub>2</sub> stock solutions (25 mM) were prepared by several rounds of degassing and flushing with argon inside a sealed serum vial. TauD apoprotein (0.25 or 0.5 mM subunit in 25 mM Tris buffer, pH 8.0) was placed into a 1 cm path length, 1 mL quartz cuvette fitted with a stopper and purged with argon. Other components were added by using gastight syringes (Hamilton) that had been flushed with anaerobic buffer. Selected samples were mixed with an equal volume of buffer sparged with 100% O2, and spectral changes were monitored over time. The effect of added H<sub>2</sub>O<sub>2</sub> was examined in one sample. All spectra were corrected to account for sample dilutions.

# Results and discussion

Kinetics of inhibition of TauD by Co<sup>II</sup> and Ni<sup>II</sup>

The effects of Co<sup>II</sup> and Ni<sup>II</sup> on TauD activity were examined by using two methods to initiate the assays. First, TauD apoprotein was added to standard assay mixes con-

taining Fe<sup>II</sup> and varied concentrations of the inhibitory metal ions, incubated for 5 min, and the total sulfite product was measured (Fig. 1, solid lines). The data were fit to Eq. (1) (where [M] is the inhibitory metal ion concentration) to determine the IC<sub>50</sub> (inhibitor concentration resulting in 50% inhibition, with the approximate range of this value shown in parentheses). The  $IC_{50}$  of  $\text{Co}^{\text{II}}$  was 41  $\mu\text{M}$  $(30-70 \mu M)$  while that of Ni<sup>II</sup> was  $32 \mu M$   $(20-40 \mu M)$ . When the TauD inhibition assays were repeated using Fe<sup>II</sup> addition to initiate the reaction (Fig. 1, dashed lines), the observed  $IC_{50}$  values were 1.9  $\mu M$  (1–3.5  $\mu M$ ) and 0.71  $\mu M$  (0.60–1.0  $\mu M$ ) for  $Co^{II}$  and  $Ni^{II}$ , respectively. The large differences in IC<sub>50</sub> values observed when using the distinct methods to initiate the reaction suggest that Fe<sup>II</sup> does not readily displace the inhibitory metal ions previously bound to the protein. These differences also highlight the fact that steady-state assays that assume rapid equilibrium kinetics are inadequate for defining the true kinetic inhibition mechanism of metal ions. Despite this caveat, the IC<sub>50</sub> values obtained for TauD were compared to those reported for three human HIF-specific prolyl 4-hydroxylase isozymes (38  $\pm$  8, 100  $\pm$  15, and 9  $\pm$  4  $\mu$ M for  $Co^{II}$ ; 130 ± 76, >1000, and 120 ± 49  $\mu M$  for  $Ni^{II}$ ) and a collagen-specific prolyl 4-hydroxylase (14  $\pm$  3  $\mu M$  for Co<sup>II</sup> and 37  $\pm$  11  $\mu M$  for Ni<sup>II</sup>) along with the  $K_i$  values estimated for FIH inhibition (1.0  $\pm$  0.4  $\mu M$  for Co<sup>II</sup> and 4  $\pm$  1  $\mu M$ for Ni<sup>II</sup>) [22]. Curiously, Co<sup>II</sup> and Ni<sup>II</sup> inhibition of the HIF prolyl 4-hydroxylases was incomplete with up to 50% activity remaining at  $0.5 \text{ mM Co}^{\text{II}}$  and up to 55%activity remaining at 1 mM Ni<sup>II</sup>. We attribute the incomplete inhibition of the prolyl 4-hydroxylases to their purification as partial holoproteins, compared to TauD that was purified as the apoprotein.

% activity remaining = 
$$100 - 100[M]/(IC_{50} + [M])$$
. (1)

To better define the kinetic mechanism of inhibition by metal ions, the time-dependence of Co<sup>II</sup> and Ni<sup>II</sup> inhibition was determined (Figs. 2A and B). In the absence of inhibitory metal ion, sulfite production began immediately (i.e., TauD apoprotein binds Fe<sup>II</sup> rapidly) and increased steadily. When

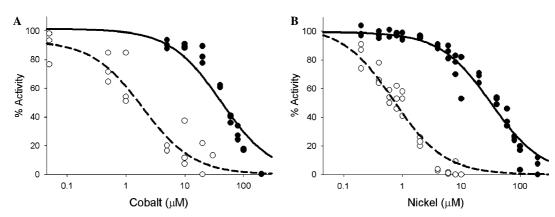


Fig. 1. Inhibition of TauD by  $Co^{II}$  and  $Ni^{II}$ . The concentrations of sulfite produced during 5 min incubations were used to assess the percent activity of TauD in standard assay conditions containing the indicated concentrations of  $Co^{II}$  (A) and  $Ni^{II}$  (B). The solid circles represent data associated with assays initiated with TauD apoprotein, whereas the open circles represent data for assays initiated by  $Fe^{II}$  addition to samples exposed to inhibitory metal ions for 2 min. The data were fit to Eq. (1) to calculate  $IC_{50}$  values.

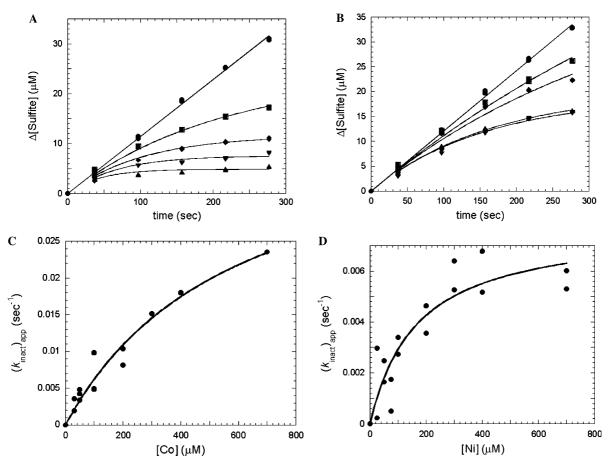


Fig. 2. Time-dependent inhibition of TauD by  $Co^{II}$  and  $Ni^{II}$ . TauD apoprotein was added to standard assay conditions and varied concentrations of  $Co^{II}$  or  $Ni^{II}$  (including 0  $\mu$ M,  $\bullet$ ; 50  $\mu$ M,  $\blacksquare$ , 100  $\mu$ M,  $\blacklozenge$ ; 300  $\mu$ M,  $\blacktriangledown$ , and 700  $\mu$ M,  $\blacktriangle$ ) were added after 23 s (equivalent to the zero time point in the subset of studies illustrated in (A,B)). Subsequent productions of sulfite were analyzed according to Eq. (2) to determine the effects of inhibitory metal concentrations on  $k_{\text{inact}}$ , the apparent rates of inactivation, as illustrated in (C,D).

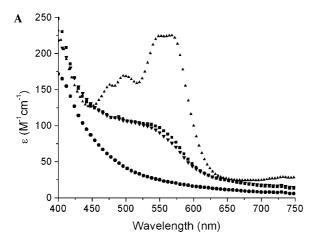
various concentrations of  $\mathrm{Co^{II}}$  or  $\mathrm{Ni^{II}}$  were added at 23 s into the assays, the rates of sulfite production were observed to decrease over time. The apparent first-order rate constants of enzyme inactivation were calculated for each assay according to Eq. (2) (where  $P_t$  is the amount of product at time t,  $v_i$  is the initial rate, and  $k_{\mathrm{inact}}$  is the apparent inactivation rate constant), and the values were replotted as a function of inhibitory metal ion concentrations (Figs. 2C and D). The apparent  $k_{\mathrm{inact}}$  values were observed to saturate at high concentrations of metal ions, consistent with slow-binding inhibition kinetics (Scheme 3) [34]. The initial dissociation constant  $K_i(k_{-1}/k_1)$  was estimated to be  $600 \pm 180 \ \mu\mathrm{M}$  for  $\mathrm{Co^{II}}$  and  $166 \pm 65 \ \mu\mathrm{M}$  for  $\mathrm{Ni^{II}}$ , and  $k_3$  (equivalent to  $(k_{\mathrm{inact}})_{\mathrm{max}}$ ) was estimated as  $0.044 \pm 0.008 \ \mathrm{s^{-1}}$  for  $\mathrm{Co^{II}}$  and  $0.078 \pm 0.0012 \ \mathrm{s^{-1}}$  for  $\mathrm{Ni^{II}}$ .

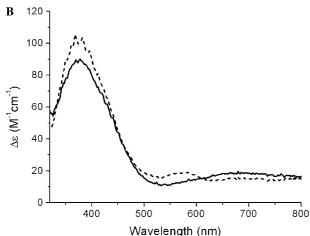
$$P_t = v_i(1 - \exp(-k_{\text{inact}}t))/k_{\text{inact}}.$$
 (2)

$$E + I \xrightarrow{k_1} EI \xrightarrow{k_2} EI^*$$
Scheme 3.

Spectroscopy of Co<sup>II</sup> and Ni<sup>II</sup> interaction with TauD

The addition of Co<sup>II</sup> to an anaerobic sample of TauD (Fig. 3A) resulted in formation of a broad peak between 450 and 600 nm, with  $\lambda_{\rm max}$  at 530 nm and  $\epsilon_{530}$  of 70 M<sup>-1</sup> cm<sup>-1</sup> (calculated on the basis of the difference spectrum of Co<sup>II</sup>-TauD minus TauD). The addition of  $\alpha$ KG had little effect on the spectrum, whereas the further addition of taurine resulted in features at 565, 552, and 500 nm with extinction coefficients of 204, 200, and 127 M<sup>-1</sup> cm<sup>-1</sup> (calculated on the basis of the difference spectrum for taurine-αKG-Co<sup>II</sup>-TauD minus TauD). The magnitude of the Co<sup>II</sup> extinction coefficient has been shown empirically to correlate with the coordination number in Co<sup>II</sup> proteins: six-coordinate sites have extinction coefficients of about 50 M<sup>-1</sup> cm<sup>-1</sup>, five-coordinate sites have values between 50 and 300 M<sup>-1</sup> cm<sup>-1</sup>, and four-coordinate sites possess a coefficient of more than  $300~M^{-1}~cm^{-1}$  [35]. We conclude that the  $Co^{II}$  sites in  $Co^{II}$ -TauD and  $\alpha KG$ - $Co^{II}$ -TauD are six-coordinate, whereas that in taurine-αKG-Co<sup>II</sup>-TauD is five-coordinate. Thus, the binding of substrate most likely leads to dissociation of a water molecule in this metal-substituted protein just as in the active, Fe<sup>II</sup>-containing enzyme.





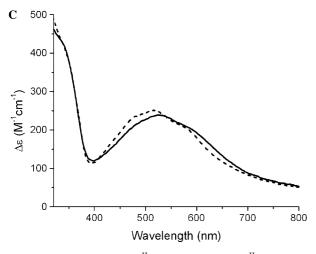


Fig. 3. Electronic spectra of  $Co^{II}$ -substituted,  $NOG\text{-}Fe^{II}$ , and  $\alpha KG\text{-}Fe^{II}$  forms of TauD. (A) An anaerobic solution of TauD apoprotein (550  $\mu M$  subunit, circles) was adjusted to contain near stoichiometric amounts of  $Co^{II}$  (inverted triangles), 2 mM  $\alpha KG$  (squares), and 2 mM taurine (triangles) in 25 mM Tris buffer, pH 8.0. (B) Analogous spectra were collected for 250  $\mu M$  TauD while substituting  $Fe^{II}$  for  $Co^{II}$  and 10 mM NOG for  $\alpha KG$ , with difference spectra shown for the taurine-NOG-Fe<sup>II</sup>-TauD minus  $Fe^{II}$ -TauD samples (solid line). (C) Analogous spectra were collected using  $Fe^{II}$  and  $\alpha KG$ , again showing difference spectra for the taurine- $\alpha KG\text{-}Fe^{II}$ -TauD minus  $Fe^{II}$ -TauD (dashed line) and the  $\alpha KG\text{-}Fe^{II}$ -TauD minus  $Fe^{II}$ -TauD (dashed line) and the  $\alpha KG\text{-}Fe^{II}$ -TauD minus  $Fe^{II}$ -TauD (solid line) samples.

We examined the accessibility of TauD-bound Co to exogenous ligands using both taurine- $\alpha$ KG-Co<sup>II</sup>-TauD and  $\alpha$ KG-Co<sup>II</sup>-TauD. No spectral changes were detected for either species upon addition of CN<sup>-</sup>, OCN<sup>-</sup>, SCN<sup>-</sup>, or ClO<sup>-</sup> at 4 mM concentrations, or in the presence of 200 mM NaCl. Although the metal coordination environment of taurine- $\alpha$ KG-Co<sup>II</sup>-TauD appears to closely resemble that of taurine- $\alpha$ KG-Fe<sup>II</sup>-TauD, the spectrum of the Co<sup>II</sup>-containing species remained unchanged when exposed to oxygen. Furthermore, despite precedence for the transformation of Co<sup>II</sup> species to Co<sup>III</sup>-OOH model compounds [36,37], no spectral perturbations were observed when H<sub>2</sub>O<sub>2</sub> (up to 750  $\mu$ M) was added to taurine- $\alpha$ KG-Co<sup>II</sup>-TauD.

In contrast to the situation for  $Co^{II}$ , the addition of  $Ni^{II}$  did not affect the spectra of TauD, protein plus  $\alpha KG$ , protein with both substrates, or for these components plus oxygen (data not shown).

## Kinetics of Inhibition of TauD by NOG

Steady-state kinetic inhibition studies (data not shown) were used to establish that NOG competed with aKG for binding to TauD, with a  $K_i$  of 290  $\pm$  90  $\mu$ M. N-Oxalyglycine should be able to form all of the ionic interactions of αKG and it contains an oxamic acid moiety as a potentially better bidentate ligand for the enzyme-bound Fe<sup>II</sup>, so we were surprised by the weak inhibition of TauD by this compound. For comparison, NOG was shown to be a competitive inhibitor of collagen prolyl 4-hydroxylase and FIH with markedly different  $K_i$  values of 1.9–7.0  $\mu$ M [26] and  $1.2 \pm 0.3$  mM [30]. The reasons for this wide range of  $K_i$ values for these enzymes remain unclear, but interactions with the protein side chains probably play a role in determining this value. Studies with the structurally related compound N-oxalyl-D-phenylalanine provided a K<sub>i</sub> of  $83 \pm 18 \,\mu\text{M}$  for FIH,14-fold smaller than that of NOG [30].

## Spectroscopy of NOG interaction with TauD

The spectra of NOG-Fe<sup>II</sup>-TauD and taurine-NOG-Fe<sup>II</sup>-TauD (Fig. 3B) differ significantly from the previously described spectra of  $\alpha$ KG-Fe<sup>II</sup>-TauD and taurine- $\alpha$ KG-Fe<sup>II</sup>-TauD (shown for comparison in Fig. 3C). We interpret this result in terms of the excess electron density of NOG compared to  $\alpha$ KG (note the trianionic form of NOG versus the

Scheme 4.

dianionic αKG, Scheme 4) which causes a shift of the metal-to-ligand charge-transfer transition to higher energy. While  $\alpha KG$ -Fe<sup>II</sup>-TauD is known to react with oxygen resulting in aKG decomposition, protein self-hydroxylation, and the generation of a 550 nm chromophore [38], the spectra of NOG-Fe<sup>II</sup>-TauD and taurine-NOG-Fe<sup>II</sup>-TauD did not change at a significant rate upon exposure to oxygen. The reduced O2 reactivity for these species also contrasts with that observed for the product complex (succinate-Fe<sup>II</sup>-TauD) that generates a 720 nm chromophore when exposed to  $O_2$  [39]. The  $\alpha$ KG- and succinate-derived ligand-to-metal charge-transfer transitions arise from chelation of Fe<sup>III</sup> by the catecholate produced by hydroxylation of Tyr73, with or without bound bicarbonate ligand [39]. We conclude that hydroxylation of Tyr73 is greatly reduced using NOG-Fe<sup>II</sup>-TauD and that NOG does not undergo oxidative decarboxylation.

## **Conclusions**

Steady-state kinetic approaches are inappropriate for determining the kinetics of inhibition of Fe<sup>II</sup>/αKG-dependent hydroxylases by metal ions; rather, slow-binding kinetic inhibition methods must be utilized. The chromophore generated upon binding of both substrates to Co<sup>II</sup>-substituted protein might serve as a useful diagnostic marker for this enzyme family. While the resulting Co<sup>II</sup> center is most likely five-coordinate in TauD, it does not react with added ligands including oxidants. The low reactivity of the Co<sup>II</sup>- and Ni<sup>II</sup>substituted enzymes could be exploited during purification and crystallization efforts to stabilize related proteins against self-hydroxylation reactions. Despite the close structural similarity of NOG to  $\alpha$ KG, this is a weak competitive inhibitor of TauD and the  $K_i$  ranges widely for different enzymes. The NOG-bound state of the enzyme does not react with oxygen to catalyze oxidative decarboxylation.

## Acknowledgments

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