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Characterization of chloride-depleted human sulfite oxidase by EPR spectroscopy: experimental evidence for the role of anions in product release

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Abstract

The Mo(V) state of the molybdoenzyme sulfite oxidase (SO) is paramagnetic and can be studied by electron paramagnetic resonance (EPR) spectroscopy. Vertebrate SO at pH < 7 and pH > 9 exhibits characteristic EPR spectra that correspond to two structurally different forms of the Mo(V) active center referred to as the low-pH (*lpH*) and high-pH (*hpH*) forms, respectively. Both EPR forms have an exchangeable equatorial OH ligand, but its orientation in the two forms is different. It has been hypothesized that the formation of the *lpH* species is dependent upon the presence of chloride. In this work we have prepared and purified samples of wild type and various mutants of human SO that are depleted in chloride. These samples do not exhibit the typical *lpH* EPR spectrum at low pH, but rather show spectra that are characteristic of the *blocked* species that contains an exchangeable equatorial sulfate ligand. Addition of chloride to these samples results in the disappearance of the *blocked* species and the formation of the *lpH* species. Similarly, if chloride is added before sulfite, the *lpH* species is formed instead of the *blocked* one. Qualitatively similar results were observed for samples of sulfite oxidizing enzymes from other organisms that were previously reported to form a *blocked* species at low pH. However, the depletion of chloride has no effect upon the formation of the *hpH* species.

The sulfite-oxidizing enzymes (SOEs), represented by sulfite oxidase (SO) in vertebrates and plants and sulfite dehydrogenase (SDH) in bacteria, catalyze the oxidation of sulfite to sulfate as represented by generic Eq. 1 (1).

$$SO_3^{2-} + H_2O \rightarrow SO_4^{2-} + 2H^+ + 2e^-$$
 (1)

In humans SO is essential for normal neonatal neurological development, and inborn deficiencies in SO result in severe physical and neurological disorders and early death (2,3).

Reaction (1) is catalyzed by the square-pyramidal oxo-molybdenum active center, which has three equatorial sulfur ligands (one from the conserved cysteinyl side chain, and two from the molybdopterin cofactor), one axial oxo ligand, and an exchangeable equatorial oxo ligand in the solvent accessible pocket of the active site (4, 5). During the proposed catalytic cycle (6), sulfite initially reduces Mo(VI) to Mo(IV). Regeneration of the Mo(VI) state

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SUPPORTING INFORMATION: List of mutants, HYSCORE spectra of *wt* hSO enriched with ³³S-enriched sulfite at pH 5.8, and Kinetic analyses plots for determination of binding constant for Cl^- . This material is available free of charge via the Internet at http://pubs.acs.org.

involves two one-electron oxidations, as shown in Scheme 1, (only the exchangeable equatorial ligand of the Mo ion is shown). The intermediate Mo(V) state is paramagnetic and has been extensively studied by electron paramagnetic resonance (EPR) spectroscopy (7–14).

Unlike X-ray crystallography or extended X-ray absorption fine structure (EXAFS) spectroscopy, EPR can detect protons in the vicinity of a paramagnetic center and is able to unequivocally identify specific nuclei through using substitutions by or permutations of magnetic isotopes (e.g., ${}^{16}\text{O} \rightarrow {}^{17}\text{O}$, ${}^{35}\text{Cl} \rightarrow {}^{37}\text{Cl}$, ${}^{14}\text{N} \rightarrow {}^{15}\text{N}$, etc.). Both, continuous wave (CW) and pulsed EPR spectroscopic approaches have been used to establish the effects of pH, anions in the media, and mutations near the active site on the identity and structure of the exchangeable equatorial ligand of the Mo(V) ion. It was found that in the absence of inhibiting anions (e.g., PO_4^{3-} , AsO_4^{3-}), wild type (wt) vertebrate SO can show two distinct types of EPR signals, high-pH (hpH) and low-pH (lpH), corresponding to two different structural forms of the Mo active center. For both of these forms the exchangeable equatorial ligand is OH (see Stage 3 of Scheme 1). However, in some mutant forms of vertebrate SO (Y343F, R160Q hSO), as well as in the R55Q variant of bacterial SDH, and wt plant SO (At-SO from Arabidopsis thaliana), two types of low-pH blocked forms have recently been observed, in which the exchangeable equatorial ligand is sulfate (bound product) rather than OH (Stage 3A of Scheme 1) (15-17). The lethality of the pathogenic R160Q mutation of hSO has been attributed to the occurrence of the *blocked* form (16), which may represent a catalytically dead end, and inefficient intramolecular electron transfer (18).

Previous pH titrations of vertebrate SO have shown that the equilibrium between the lpH and hpH forms depends on the amount of Cl⁻ added to the buffer in the form of NaCl (12). From these experiments it was hypothesized that Cl⁻ is an integral part of the lpH form, likely to be coordinated to the Mo ion. Attempts to detect the hyperfine interaction (hfi) with this chloride were later undertaken by Doonan *et al.* using CW EPR and ³⁵Cl or ³⁷Cl- enriched NaCl (19). While some differences between the ³⁷Cl- and ³⁵Cl-enriched samples were observed, they were at the limit of the experimental accuracy. The weakness of the experimental CW EPR evidence for chloride interaction was, however, alleviated by the fact that when fluoride, bromide or iodide were used instead of chloride, clearly observable EPR splittings due to the halogen nucleus hfi were detected (12,19). Based on these data, the chloride ligand was suggested to be weakly coordinated at the axial position of the Mo(V) center *trans* to the axial oxo ligand (19).

More conclusive and detailed information about the interaction between Mo and Cl in lpH SO was obtained recently using pulsed EPR and DFT calculations (17). The electron spin echo envelope modulation (ESEEM) caused by ³⁵Cl and ³⁷Cl was unequivocally detected, and the *hfi* and nuclear quadrupole interaction (*nqi*) parameters of the nearby chlorine nucleus were determined. DFT calculations performed for various structural models for incorporation of Cl⁻ into the active site established that Cl⁻ does not coordinate in the axial position, but rather is hydrogen bonded to the equatorial OH ligand, to the hydrogen atoms of the surrounding OH groups, and to amino acid residues of the binding site. Interestingly, even a sample of hSO purified elsewhere and prepared without added chloride showed the CW EPR spectrum characteristic of the *lpH* form (12,19), as well as a ^{35,37}Cl ESEEM (17). Thus, it was hypothesized that Cl⁻ may be essential for the formation of the *lpH* form, and that *wt* vertebrate SO has a high affinity for Cl⁻ at low pH (17).

Here we report that an SOE apparently *lacking* chloride near the Mo active site can be prepared when the enzyme is purified using an initial desalting column about twice as large as that employed in earlier preparations (20,21). The *wt* hSO purified in this manner does *not* exhibit the typical *lpH* species at pH ~6, but shows instead formation of the *blocked*

species, as confirmed by detecting characteristic ³³S ESEEM in a sample prepared with ³³Slabeled sulfite. A similar result was also obtained for various mutated hSO enzymes. Addition of chloride to these samples converted their EPR spectra to the typical *lpH* form (transition between Stages 3A and 3 in Scheme 1) with the exception of the lethal R160Q mutant of hSO. Similarly, adding chloride before the reduction with sulfite resulted in formation of the *lpH* species instead of the *blocked* one.

MATERIALS AND METHODS

Recombinant *wt* hSO and mutant forms of hSO were expressed and purified as previously described (20,21). The chloride content of the enzyme samples was analyzed using QuantiChromTM Chloride assay kit (DICL-250) from BioAssay Systems. Steady-state kinetic measurements were performed using the sulfite/cytochrome c assay reaction for SO and monitoring the reduction of cytochrome *c* at 550 nm. Concentrations of chloride and sulfite were varied and plots of l/rate *vs*. l/substrate were made for a series of chloride concentrations. The binding constant for chloride was determined from the **x** intercept of a replot of the slopes of the lines (l/rate *vs*. l/substrate) *vs*. chloride concentration (see Supporting Information).

The EPR samples at pH 5.8 were prepared in 100 mM Bis-Tris buffer. The samples at pH 7 and 9 were prepared in 100 mM Bis-Tris propane buffer. The enzyme was reduced with a 20-fold excess of sodium sulfite under argon and immediately frozen in liquid nitrogen. The same buffer system and procedure were used for reduction with ³³S-labeled sulfite, prepared as previously described (15). After the *blocked* form was detected by EPR measurements of the samples at low pH, the samples were thawed at 0°C, 10 or 100 mM NaCl was added, the samples were frozen again, and the EPR measurements were repeated. In a parallel set of experiments, 100 mM NaCl was added to chloride-depleted enzymes prior to adding Na(SO₃)₂.

The X-band (~9 GHz) CW EPR experiments were performed on a Bruker ESP300 spectrometer at 77 K. The ESEEM measurements were done on a homebuilt K_{a} -band (26–40 GHz) pulsed EPR spectrometer (22). The measurement temperature was about 21 K.

The concentrations of Mo(V) in the SO enzyme samples were determined by comparing the double integrals of the X-band CW EPR spectra recorded at 77 K with that of the 5 mM Cu(NO₃)₂ in glassy water/glycerol solution. For double intergation, only the well-defined EPR features of the active centers containing non-magnetic (I = 0, ~ 75% natural abundance) molybdenum isotopes were used. The broad and rather featureless EPR lines of the Mo(V) centers containing magnetic molybdenum isotopes (95 Mo, 97 Mo, both I = 5/2, total natural abundance ~ 25%) were suppressed by the baseline subtraction preceding the double integration. Therefore, to account for the magnetic isotopes and obtain the total concentration of the Mo(V) active centers, the estimated concentrations of Mo(V) centers relative to the enzyme concentrations were about 40% – 50%. Thawing the samples to add NaCl (overall handling time at 0 °C about 2 min) resulted in a decrease of the Mo(V) concentration to about 30% – 40%.

RESULTS AND DISCUSSION

Numerous previous EPR experiments on *wt* vertebrate SO at low pH (< 7) performed by us and other authors (7–10,12,17,19) always showed formation of the *lpH* EPR-active Mo(V) species with characteristic hyperfine coupling from an exchangeable proton. Recently, we began preparing and purifying recombinant *wt* hSO in our own laboratory and were initially

surprised to find that the EPR signal we systematically obtained at low pH (typically, 5.8) was similar to that characteristic of the *blocked* form of SO (trace 1 in Figure 1) with coordinated sulfate (product). Similar to the *blocked* forms observed earlier in *wt* At-SO (15) and Y343F hSO (16), this EPR signal is characterized by the principal g-values (g_1, g_2, g_3) \approx (1.999, 1.972, 1.963) and does not show any hyperfine splittings because the exchangeable equatorial ligand, sulfate, does not have any magnetic nuclei (the natural abundance of ¹⁶O is 99.757%, and that of ³²S+³⁴S is 99.2%).

In order to verify that the EPR signal we detected for *wt* hSO indeed belongs to the *blocked* form, the enzyme was reduced with ³³S-enriched sulfite, and K_a-band ESEEM experiments were performed similar to those described elsewhere (15,16). The ESEEM spectra (see Figure 2) confirm that the observed EPR signal belongs to the *blocked* form of SO. The ³³S *hfi* constant ($a_{iso} \approx 2.6$ MHz) and quadrupole coupling constant ($e^2Qq/h \approx 36$ MHz) estimated from hypefine sublevel correlation (HYSCORE) spectra (see Supporting Information) are similar to those found for the *blocked* species in other SOEs (15,16). The samples prepared at high pH (~ 9.0) show the formation of the usual *hpH* species (trace 3 in Figure 1) characterized by the principal g-values (g_1, g_2, g_3) \approx (1.987, 1.964, 1.953) (10). At intermediate pH (*e.g.*, pH 7) a mixture of the *blocked* and *hpH* forms was observed (trace 2 in Figure 1). The typical *lpH* form was *not* generated at any pH within the studied range from 5.8 to 9.

Careful examination of the enzyme purification and sample preparation procedures has revealed that the only difference between the published procedure (20,21) and our current procedure is in the larger size of the G25 desalting column (500 mL vs. 250 mL bed volume) used in our initial chromatographic purification step. Taking into account the recent suggestion that Cl⁻ may be indispensable for the formation of the *lpH* species (17), we hypothesized that our EPR results can be explained if we assume that our purification procedure sufficiently reduces the Cl⁻ concentration to preclude formation of the usual *lpH* form of SO. Indeed, the measurement of the chloride content of our enzyme samples prepared for EPR studies at pH 5.8 yielded the ratio [Cl⁻]/[SO] to be about 20. For comparison, the chloride concentration in samples of hSO purified elsewhere with the smaller desalting column was significantly larger, [Cl⁻]/[SO] \approx 30–40.

To test if this low chloride concentration was responsible for the formation of the *blocked* species instead of the *lpH* form, Cl⁻ was added to the low-pH sample of *wt* hSO that was purified in our laboratory as described above. Adding 10 mM NaCl resulted in formation of a substantial amount of the *lpH* species (characterized by the principal g-values (g_1, g_2, g_3) \approx (2.004, 1.973, 1.966) and showing the hyperfine splittings (A_1, A_2, A_3) \approx (0.8, 0.8, 1.3) mT at the EPR turning points caused by the *hfi* with the OH ligand proton (10), and a significant decrease in the amount of the *blocked* form (compare traces 1 and 2 in Figure 3). An EPR sample with 100 mM NaCl gave entirely the typical *lpH* form (trace 3 in Figure 3). When 100 mM NaCl was added to an EPR sample before the reduction by sulfite, the same end result was obtained: the usual *lpH* species was generated instead of the *blocked* one.

The effect of added Cl⁻ on various preparations of SO and SDH that exhibited the *blocked* form was also studied for the purpose of comparison and generalization. With one exception, all of the hSO mutants purified in our lab (see Supporting Information) that exhibited the *blocked* form when prepared at low pH without added chloride, were completely converted to the typical *lpH* form by adding 100 mM NaCl (as exemplified in Figure 3 for *wt* hSO). The exception is the pathogenic R160Q mutant (16), which was not affected by added Cl⁻ (up to 300 mM NaCl) and remained in the *blocked* form. On the other hand, for the analogous mutation (R55Q) of bacterial SDH the *blocked* form was converted to the *lpH* form by adding 100 mM NaCl (see Figure 4) (23). For At-SO and Y343F hSO at

pH 6 adding 100 mM NaCl resulted in formation of measurable amounts of the typical *lpH* species, although the EPR spectra were still dominated by the *blocked* form (as an example, see Figure 5 for the results obtained for At-SO).

The formation of the *blocked* form of hSO in chloride-depleted samples is consistent with the earlier proposal that multiple mechanisms may be possible for SO (24). The *blocked* form results from one-electron oxidation of the molybdenum center prior to product release to give a Mo(V)-OSO₃ moiety (2 \rightarrow 3a of Scheme 1). However, the weakly coordinating sulfate ion is readily replaced in the presence of other anions at low pH. Addition of chloride gives the well-known *lpH* form (3 in Scheme 1); other halogens (F⁻, Br⁻, I⁻) produce the same effect (12,19). For those anions, however, the EPR splittings due to the halogen nucleus *hfi* are directly observed by CW EPR because the *hfi* constants of ¹⁹F, ⁷⁹Br and ¹²⁷I per unit unpaired electron spin density are several fold greater than that of ^{35,37}Cl (25). Distinct EPR forms with coordinated phosphate (11,26) and arsenate (9) have also been observed at low pH. It has also been suggested that the *blocked* form actually contains bound sulfite, the reactant, which is present in large excess in the EPR samples (27,28).

Recently we used 35,37 Cl ESEEM spectroscopy and DFT calculations to obtain information about the chloride binding site in the vicinity of the molybdenum active center (17). In contrast to earlier suggestions (19) that Cl⁻ may be coordinated in the axial position to Mo(V), *trans* to the oxo ligand, our data have shown that in *lpH* SOEs Cl⁻ is most likely hydrogen-bonded to the proton of the equatorial OH ligand as shown in Figure 6a. Other details of the Cl⁻ binding site are less definitive, although it is clear the Cl⁻ is stabilized in place by several hydrogen bonds, and the overall environment is fairly spherical. This is suggested by a relatively weak nuclear quadrupole interaction of this chloride ion. Based on X-ray structures (PDB 2A99 (4)), one of the amino acid residues involved in the Cl⁻ binding site in chicken SO (cSO) was thought to be W204, which corresponds to W226 in hSO. However, mutations of W226 residue to Ala or Phe did not affect the formation of the typical *lpH* form with added 100 mM NaCl at low pH. Therefore, the role of W226 in stabilization of Cl⁻ in the enzyme in solution is questionable.

One possible qualitative explanation of the chloride effect on hydrolysis of sulfate from the Mo active center is that Cl⁻ competes with one of the distal oxygens of SO_4^{2-} for the same binding site (see Figure 6b), so that when Cl⁻ displaces this oxygen, SO_4^{2-} is hydrolyzed more easily. This model is indirectly supported by the fact that neither the *blocked* species detected in this work nor those studied earlier (15, 16), exhibit ^{35,37}Cl ESEEM spectra.

At high pH, the hydrolysis of sulfate and the complete catalytic turnover occur without the presence of halogens, with the Mo(V) *hpH* species being formed at Stage 3 of Scheme 1. The results of recent ¹H and ¹⁷O ESEEM measurements suggest that in this case the exchangeable equatorial OH ligand is hydrogen-bonded to a second-sphere OH⁻ (29,30). We hypothesize, therefore, that at high pH the OH⁻ anion plays the same role in facilitating the hydrolysis of sulfate as the halogen anions do at low pH. The provisional structure for an *hpH* SOE based on this hypothesis is shown in Figure 6c.

It would be tempting to interpret the EPR results described above as suggesting that chloride plays a physiological role in SO by facilitating the hydrolysis of the product from the molybdenum active center. The EPR experiments, however, do not represent a sufficient ground for such interpretation because the processes in the EPR samples of SO are limited by a single electron transfer event, and the catalytic cycle is incomplete. In order to assess the effects of chloride ion concentration on the overall catalytic cycle, we have carried out steady-state activity measurements on *wt* hSO at pH 5.8 and 8.0 (Table 1). At low pH (5.8), high concentrations of chloride greatly decrease the catalytic activity of recombinant hSO

and exhibit "mixed" type inhibition. This result is in agreement with previous studies of the effect of chloride on the activity of native chicken SO (31). In addition, the presence of high concentrations of chloride (and other small anions) have an inhibitory effect on the rate constant for intramolecular electron transfer (IET) between the Mo(VI)/Fe(II) and Mo(V)/Fe(III) forms of SO (32). It was proposed that small anions which can fit into the Mo active site will weaken the electrostatic attraction between the Mo and heme domains and disfavor IET. The dissociation constant for Cl⁻ from hSO, measured under steady-state conditions using the sulfite/cytochrome *c* assay reaction at pH 5.8, was 13 ± 3 mM (Table 1). From this result it follows that the chloride concentration in the EPR sample of trace 1 of Figure 1 (~5 mM) is insufficient to produce the *lpH* EPR signal. Hence, we observe the *blocked* form. The subsequent addition of chloride results in sufficient population of Cl in the active site to observe the *lpH* signal (traces 2 and 3 of Figure 1).

At high pH (8.0) the activity decreases by ~30% as the chloride concentration is increased to 100 mM, substantially less than the seven-fold decrease at pH 5.8. The steady-state results at high pH are consistent with the fact that these samples do not exhibit 35,37 Cl ESEEM spectra (15,16) and give a typical *hpH* EPR spectrum (29,30).

The EPR results indicate that reduction of hSO with sulfite at low chloride concentrations and at low pH involves the one-electron oxidation of the Mo^{IV}-OSO₃ center to Mo^V-OSO₃ (pathway $2 \rightarrow 3A$ in Scheme 1) to give the *blocked* form. However, the predominant Mo(V) species observed by EPR in the reductive half-reaction may not necessarily be the most catalytically efficient one. The steady-state experiments on SOEs involve a second oneelectron oxidation to regenerate the Mo^{VI} form of the enzyme, and the possibility of multiple kinetic pathways for SOEs has been discussed previously (24). If the steady-state kinetics experiments at low chloride and low pH presented here for hSO also involve the blocked Mo(V) form (3A of Scheme 1), then a second one-electron oxidation to Mo(VI) could generate a catalytically competent species that could be hydrolyzed to product. Consequently, increasing the chloride concentration can have two competing effects. It can facilitate the dissociation of the product/substrate from Mo(V) state (Scheme 1, pathway 3A \rightarrow 3) to give the typical *lpH* EPR form, and the amount of this form that is observed depends upon the SOE and the mutation. However, large concentrations of chloride simultaneously decrease the IET rates (32) and have an inhibitory effect on the catalytic activity (Table 1). These observations are consistent with a previous extensive microcoulometric study of the effects of anions and pH on the potentials of SO (33).

CONCLUSIONS

In contrast to all of the earlier experimental observations (7–10,12,17,19), in this work we detected, for the first time, the formation of the *blocked* Mo(V) species in *wt* hSO at low pH (pH < 7). The formation of the *blocked* species was traced to the depletion of chloride from the purified enzyme by employing a large (compared to those used in other laboratories) desalting column. This observation supports the formation of Mo-OSO₃ as the first step of the catalytic cycle of SO (Scheme 1). Adding NaCl to the *wt* hSO samples where the *blocked* species was formed results in hydrolysis of the product, sulfate, from the Mo active center and the formation of the usual *lpH* species. Similar experiments on several mutant forms of hSO have shown that the formation of an irreversible *blocked* species for the R160Q mutant of hSO supports the earlier hypothesis for the lethality of this mutation (16). Steady-state assays as a function of pH and chloride concentration show that *k*_{cat} for *wt* hSO decreases with increasing chloride concentration, but this effect is much smaller at high pH. The combined EPR and kinetic data presented here are a further indication that the results of specific active site mutations on the chemical and spectroscopic properties of SOEs reflect a

Supplementary Material

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Abbreviations

SO	sulfite oxidase		
SOEs	sulfite oxidizing enzymes		
SDH	sulfite dehydrogenase		
EPR	electron paramagnetic resonance		
lpH	low-pH		
hpH	high-pH		
DFT	density functional theory		
ESEEM	electron spin echo envelope modulation		
HYSCORE	hypefine sublevel correlation		
hfi	hyperfine interaction		
wt	wild type		

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Figure 1.

Traces 1, 2, and 3, X-band CW EPR spectra of *wt* hSO at pH 5.8, 7.0 and 9.0, respectively. No NaCl was added to the samples. Experimental conditions: microwave (mw) frequency, 9.455 GHz; mw power, 2 mW; magnetic field modulation amplitude, 0.1 mT; temperature, 77 K, enzyme concentrations, 430, 570, and 500 μ M for traces 1, 2, and 3, respectively. Estimated relative concentrations of Mo(V) centers are 51%, 47% and 48% for traces 1, 2, and 3, respectively.



Figure 2.

Amplitude Fourier transformation spectra of normalized two-pulse ESEEM spectra of ³³S-labeled (*i.e.*, reduced with ³³S-enriched sulfite, solid trace) and unlabeled (dashed trace) chloride-depleted *wt* hSO at pH 5.8 obtained at g_Y EPR turning point. Experimental conditions: mw frequency, 29.51 GHz; $B_0 = 1074.1$ mT; mw pulses, 9 and 15 ns; temperature, 21 K, enzyme concentrations, 600 μ M for trace 1 and 430 μ M for trace 2. Since the only difference between the samples was in the enrichment with magnetic isotope of sulfur (³³S, natural abundance is 0.76%), the lines observed in the spectrum shown by the black trace are attributed to ³³S.

Rajapakshe et al.



Figure 3.

X-band CW EPR spectra of *wt* hSO at pH 5.8 without adding NaCl (trace 1, same as trace 1 in Figure 1), with 10 mM NaCl (trace 2), and with 100 mM NaCl (trace 3). Experimental conditions: mw frequency, 9.455 GHz; mw power, 2 mW; magnetic field modulation amplitude, 0.1 mT; temperature, 77 K, enzyme concentrations, 430 µM for traces 1 and 3, and 270 µM for trace 2. Estimated relative concentrations of Mo(V) centers are 51%, 30%, and 40% for traces 1, 2, and 3, respectively.



Figure 4.

X-band CW EPR spectra of R55Q SDH at pH 4.9 without added NaCl (trace 1) and with 100 mM NaCl (trace 2). Experimental conditions: mw frequency, 9.470 GHz; mw power, 2 mW; magnetic field modulation amplitude, 0.1 mT; temperature, 77 K, enzyme concentration, 300 μ M. The asterisks over trace 1 show the EPR turning points of the *blocked* species. The minor features in trace 1 belong to the *hpH* form. Estimated relative concentrations of Mo(V) centers are 42% and 30% for traces 1 and 2, respectively.



Figure 5.

X-band CW EPR spectra of *wt* At-SO at pH 6.0 with added 100 mM NaCl (trace 1) and without NaCl (trace 2). Trace 3 represents the difference between traces 1 and 2 showing the contribution of the *lpH* SO spectrum obtained after addition of chloride. Experimental conditions: mw frequency, 9.483 GHz; mw power, 2 mW; magnetic field modulation amplitude, 0.1 mT; temperature, 77 K, enzyme concentration, 460 μ M. Estimated relative concentrations of Mo(V) centers are 35% and 42% for traces 1 and 2, respectively.

Rajapakshe et al.



Figure 6.

Putative structures of the substrate-accessible side of the Mo active center in (a) lpH, (b) *blocked*, and (c) hpH forms

Rajapakshe et al.



Scheme 1.

Simplified catalytic cycle of SO and SDH. Stage 1 corresponds to the resting state of the enzyme. The pathway $1\rightarrow 2\rightarrow 3\rightarrow 1$ corresponds to the normal catalytic turnover. Depending on the buffer pH, Stage 3 corresponds to the *hpH* or *lpH* form of the active center. Stage 3A (typically observed at low pH < 7) corresponds to the *blocked* form of the active center. In *wt* vertebrate SO, Stage 3A is generated in chloride-depleted samples at low pH. Addition of chloride results in hydrolysis of sulfate (transition $3A\rightarrow 3$).

Table 1

Steady-state data for wt-hSO with various amounts of NaCl added

pН	Amount of added chloride, mM	$k_{\text{cat}}, \text{s}^{-1}$	<i>K</i> _M (sulfite), μM	Cl ⁻ dissociation constant, mM
5.8	0	23 ± 2	2 ± 1	13 ± 3
	2.5	26 ± 1	4 ± 1	
	5	28 ± 3	6 ± 1	
	15	15 ± 1	6 ± 1	
	30	17 ± 1	5 ± 1	
	100	4 ± 1	4 ± 1	
8.0	0	27 ± 2	9 ± 3	14 ± 3
	2.5	25 ± 1	9 ± 2	
	5	25 ± 1	7 ± 1	
	10	18 ± 1	8 ± 1	
	30	15 ± 1	8 ± 1	
	100	19 ± 1	12 ± 1	