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## Intermediates involved in the reduction of $\text{SO}_2$ : insight into the mechanism of sulfite reductases†

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Sulfite reductases (SiRs) catalyze the reduction of  $\text{SO}_3^{2-}$  to  $\text{H}_2\text{S}$  in biosynthetic sulfur assimilation and dissimilation of sulfate. The mechanism of the  $6\text{e}^-/6\text{H}^+$  reduction of  $\text{SO}_3^{2-}$  at the siroheme cofactor is debated, and proposed intermediates involved in this  $6\text{e}^-$  reduction are yet to be spectroscopically characterized. The reaction of  $\text{SO}_2$  with a ferrous iron porphyrin is investigated, and two intermediates are trapped and characterized: an initial  $\text{Fe(III)}-\text{SO}_2^{2-}$  species, which undergoes proton-assisted S–O bond cleavage to form an  $\text{Fe(III)}-\text{SO}$  species. These species are characterized using a combination of resonance Raman (with  $^{34}\text{S}$ -labelled  $\text{SO}_2$ ), EPR and DFT calculations. Results obtained help reconcile the different proposed mechanisms for the SiRs.

The reduction of sulfate to sulfide is a crucial step of the geochemical sulfur cycle, which controls biochemical sulfur assimilation and the respiration of sulfate-reducing bacteria.<sup>1,2</sup> The reduction of sulfate to sulfide is catalyzed by two key metalloenzymes, ubiquitous in microorganisms, including sulfate-reducing bacteria and archaea as well as in methanogens.<sup>3,4</sup> The reduction of sulfate ( $\text{SO}_4^{2-}$ ) requires its insertion into adenosine 5'-monophosphate to form adenosine 5'-phosphosulfate, which is then reduced to release sulfite ( $\text{SO}_3^{2-}$ ).  $\text{SO}_3^{2-}$  is then reduced in the active site of sulfite reductase (SiR). The reduction of sulfite is catalyzed by the siroheme cofactor (Fig. 1A) present in all the SiRs.<sup>5</sup> The siroheme cofactor is bridged to an  $\text{Fe}_4\text{S}_4$  cubane via one of its cysteine ligands.<sup>6–8</sup> The mechanism of the  $6\text{e}^-$  reduction of  $\text{SO}_3^{2-}$  to  $\text{S}^{2-}$  is debated. In its active form siroheme and  $\text{Fe}_4\text{S}_4$  clusters are reduced: i.e. the iron in the siroheme is in its  $\text{Fe(II)}$  state, and the  $\text{Fe}_4\text{S}_4$  cluster is reduced. Although there are crystal structures of substrate-bound enzyme (Fig. 1A, left), there is no clarity on the oxidation states of the siroheme- $\text{Fe}_4\text{S}_4$  unit or sulfur in these structures, and two different mechanisms have been proposed (Fig. 1B).<sup>9,10</sup> Initially, based on observation of  $2\text{e}^-$  and  $4\text{e}^-$

partially reduced species, trithionate and thiosulfate, during  $\text{SO}_3^{2-}$  reduction, three consecutive  $2\text{e}^-/2\text{H}^+$  steps were proposed (Fig. 1B).<sup>11–17</sup> Protons are provided by conserved arginine and lysine residues present in the active site.<sup>15</sup> In addition, a sulfur monoxide (SO)-bound siroheme intermediate was proposed based on the crystal structure obtained by the oxidation of an  $\text{S}^{2-}$ -bound siroheme- $\text{Fe}_4\text{S}_4$  site (Fig. 1A right).<sup>9</sup> Recently, the direct  $6\text{e}^-/6\text{H}^+$  reduction of  $\text{SO}_3^{2-}$  in the active site of an SiR (Fig. 1B) was called into question with the identification of a trisulfide formed between the two conserved cysteine residues of another dissimilatory sulfite reductase protein C (DsrC), which is encoded by all

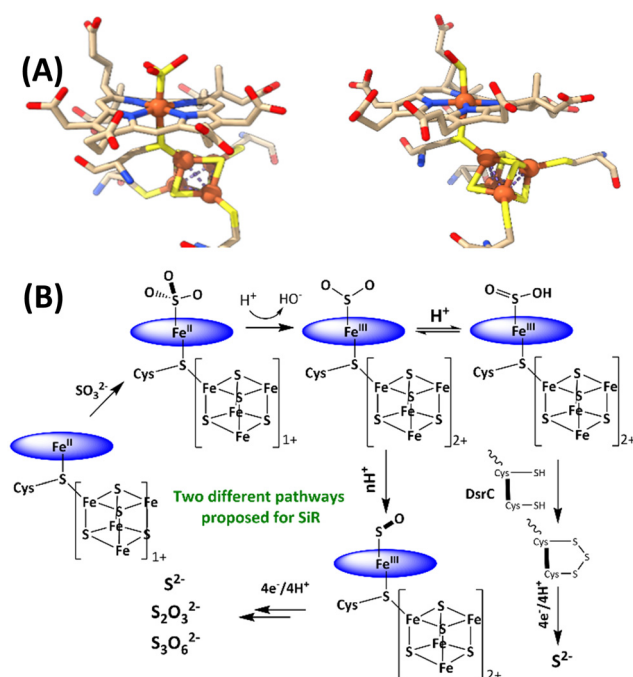


Fig. 1 (A) active sites of  $\text{SO}_3^{2-}$ -bound SiR and  $\text{SO}$ -bound SiR (pdb id: 7GEP) and (B) proposed mechanisms of the SiRs (direct  $6\text{e}^-$  reduction and trisulfide pathways are shown). The charge of the  $[\text{Fe}_4\text{S}_4]$  cluster (excluding the cysteines) is indicated next to the cubane.

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genomes that contain the genes of the catalytic A and B domains of an SiR and binds the DsrAB complex, as an end product of  $\text{SO}_3^{2-}$  reduction. The rate of  $\text{SO}_3^{2-}$  reduction by the SiR is first order with respect to the DsrC and about 15 times higher than the rate in the absence of the DsrC.<sup>18</sup> The proposed mechanism invoked the attack on an  $\text{Fe(III)}\text{-SO}_2^-/\text{Fe(III)}\text{-SO}_2\text{H}$  species (Fig. 1B), formed after the initial  $2e^-$  reduction of  $\text{SO}_3^{2-}$ , by the two conserved cysteines of the DsrC to form a trisulfide.<sup>19</sup> This trisulfide is then reduced to release sulfide, avoiding generation of trithionate and thiosulfate and resulting in sulfide as the only product of  $\text{SO}_3^{2-}$  reduction.

A synthetic analogue of the siroheme could be useful in understanding the mechanism.<sup>20</sup> Unfortunately, the synthesis of this cofactor is not trivial and has not yet been achieved.<sup>21,22</sup> Alternatively, simpler porphyrins such as iron tetraphenyl porphyrin (FeTPP), while not an exact model of a siroheme- $\text{Fe}_4\text{S}_4$  active site, has a reduction potential similar to that of the siroheme from different enzymes (which vary between  $-0.19$  and  $-0.29$  V at pH 7, and for FeTPP is  $-0.20$  V at pH 7) and hence can be used to gain insight into this intriguing reaction.<sup>23,24</sup> Initial attempts to investigate the reaction of  $\text{SO}_2$  with ferrous porphyrin in non-polar organic solvents inevitably resulted in the formation of sulfate-bound ferric porphyrin.<sup>21,25–27</sup> Recently, the reaction of ferrous tetraphenyl porphyrin with  $\text{SO}_2$  was investigated in a protic organic solvent at room temperature (RT). The reaction proceeded to result in the  $2e^-$  reduction of  $\text{SO}_2$  to  $\text{SO}$ , and the released  $\text{SO}$  could be trapped using 2,3-dimethylbutadiene, resulting in the formation of a cyclic sulfoxide.<sup>28</sup> An  $\text{Fe(III)}\text{-SO}$  intermediate was identified and was characterized using Mössbauer and EPR spectroscopy as a low-spin ferric porphyrin antiferromagnetically coupled to a triplet  $\text{SO}$ , resulting in an  $S = 1/2$  species. Although the presence of the Fe-S bond could not be established experimentally, the observed S-O vibrations of this species were consistent with theoretically predicted vibrations for an  $\text{Fe(III)}\text{-SO}$  species.<sup>28</sup> One of the  $2e^-$  needed is derived from iron porphyrin to which  $\text{SO}$  is bound, and the other comes from a free  $\text{Fe(II)TPP}$ . The 1:1 ratio of  $[\text{Fe(III)TPP}]^+$  and  $[\text{Fe(III)TPP-SO}]^+$  species was confirmed by Mössbauer spectroscopy.<sup>24,26</sup> The  $\text{Fe(III)}\text{-SO}_2^-/\text{Fe(III)}\text{-SO}_2\text{H}$  species, proposed in the SiR mechanism, has not however been observed either in the protein or in synthetic systems.

To assess the involvement of any intermediate species prior to the  $[\text{Fe(III)TPP-SO}]^+$  species, the reaction of  $\text{Fe(II)TPP}$  with  $\text{SO}_2$  is allowed to proceed at  $-80^\circ\text{C}$  (MeOH-liq.  $\text{N}_2$  bath) for 10 min, and then, the reaction mixture is frozen in liq.  $\text{N}_2$ . New EPR signals (Fig. 2A, green) are obtained with  $g = 2.36$ ,  $2.27$ , and  $1.89$ , indicating the formation of another  $S = 1/2$  low-spin intermediate species (henceforth referred to as Int-I) prior to the formation of  $[\text{Fe(III)}\text{-SO}]^+$  species (henceforth referred to as Int-II). In parallel to the Int-I signals, there is a  $g = 6.0$  signal corresponding to a high-spin  $[\text{Fe(III)TPP}]^+$  species, suggesting that Int-I also results from the  $2e^-$  reduction of  $\text{SO}_2$ , where one electron is derived from  $\text{Fe(II)TPP}$  to which  $\text{SO}_2$  to, and the other electron is derived from a free  $\text{Fe(II)TPP}$ , which gets oxidized to a high-spin  $[\text{Fe(III)TPP}]^+$  species with a  $g = 6.0$  EPR signal. Mössbauer data suggests an equal population of these two species in the sample (Fig. S1, ESI†). When the solution is warmed up to  $-40^\circ\text{C}$  (10 min), the EPR

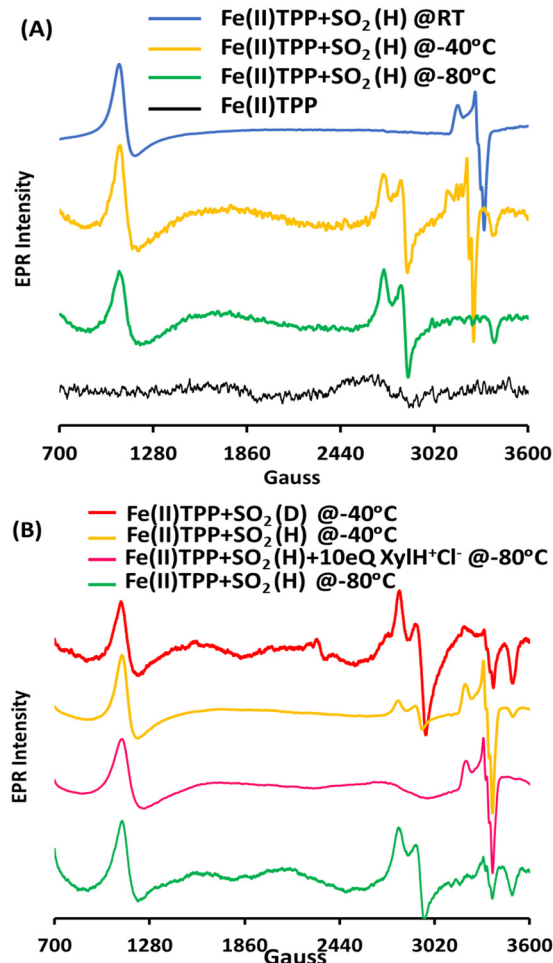


Fig. 2 (A) X-band EPR data collected at 77 K for a frozen solution of  $\text{Fe(II)TPP}$  (black), reaction mixture of  $\text{Fe(II)TPP}$  and  $\text{SO}_2$  in THF with 1%MeOH (v/v) kept at  $-80^\circ\text{C}$  for 10 min and later frozen in liq.  $\text{N}_2$  (green), kept at  $-40^\circ\text{C}$  for 10 min and later frozen in liq.  $\text{N}_2$  (yellow) and kept at RT for 10 min and further frozen in liq.  $\text{N}_2$  (blue, from reference 27). (B) Reaction mixture of  $\text{Fe(II)TPP} + \text{SO}_2$  in THF with 1%MeOH (v/v) kept at  $-80^\circ\text{C}$  for 10 min and later frozen in liq.  $\text{N}_2$  (green); reaction mixture of  $\text{Fe(II)TPP} + \text{SO}_2$  in THF with 1%MeOH (v/v) and 10 eq. of  $\text{XylH}^+\text{Cl}^-$  kept at  $-80^\circ\text{C}$  for 10 min and later frozen in liq.  $\text{N}_2$  (pink), kept at  $-40^\circ\text{C}$  for 10 min and further frozen in liq.  $\text{N}_2$  (yellow); and  $\text{Fe(II)TPP} + \text{SO}_2$  reaction mixture in THF with 1%MeOD (v/v) kept at  $-40^\circ\text{C}$  for 10 min and later frozen in liq.  $\text{N}_2$  (red).

signals originating from Int-I decrease, and the EPR signals from Int-II emerge (Fig. 2A, yellow), indicating that Int-I decays to produce Int-II (Fig. 2A, blue), i.e., Int-I is a species formed prior to Int-II in the reaction.

The transition of Int-I to Int-II depends on the availability of the protons. When the reaction of  $\text{SO}_2$  and  $\text{Fe(II)TPP}$  is performed at  $-80^\circ\text{C}$  (10 min) but in the presence of 10 eq. of xylydinium chloride ( $\text{XylH}^+\text{Cl}^-$ ) as a proton source, the EPR signals from Int-I (Fig. 2B, green) are no longer observed and only those of Int-II are observed (Fig. 2B, pink). Similarly, when the MeOH proton source in the solution is deuterated (1%  $\text{CD}_3\text{OD}$  instead of 1% MeOH,  $\text{CD}_3\text{OD}$  represented as MeOD) and instead used as the proton source, the EPR signal of the reaction at  $-40^\circ\text{C}$  (Fig. 2B, red) shows Int-I and Int-II in a 5:1 ratio relative to MeOH where this ratio is 1:2, indicating that there is an H/D isotope effect in the

conversion of Int-I to Int-II, consistent with a protonation step being involved in the reaction.

Int-I trapped at  $-80^{\circ}\text{C}$  is further characterized using resonance Raman spectroscopy of samples prepared with  $\text{SO}_2$  and comparing the vibrations with those observed when  $^{34}\text{SO}_2$  is used. In Int-I, vibration is observed at  $984\text{ cm}^{-1}$ , which shifts to  $970\text{ cm}^{-1}$  on  $^{34}\text{S}$  substitution (Fig. 3A, top). This is consistent with an S–O stretching mode from an  $\text{SO}_2$ -derived axial ligand.<sup>29,30</sup> In the lower energy region, vibration is observed at  $340\text{ cm}^{-1}$ , which shifts to  $337\text{ cm}^{-1}$  on  $^{34}\text{S}$  labelling (Fig. 3B, bottom). The energy of this vibration and isotope shift indicate that it is an Fe–S stretching mode.<sup>31,32</sup>

In the past, Int-II had been tentatively assigned as a solvent-bound  $[\text{Fe(III)TPP-SO}]^+$  species with the S–O vibration at  $1014\text{ cm}^{-1}$ , which falls on the shoulder of a porphyrin band at  $1004\text{ cm}^{-1}$ . Accordingly, the  $1014\text{ cm}^{-1}$  vibration shifts to  $1005\text{ cm}^{-1}$  (Fig. 4A, top) on  $^{34}\text{S}$  substitution, confirming it to be an S–O stretching vibration, as previously assigned. The Fe–S vibration of Int-II is observed at  $382\text{ cm}^{-1}$ , which shifts to  $377\text{ cm}^{-1}$  with  $^{34}\text{S}$  substitution (Fig. 4B, bottom). There is also another vibration at  $206\text{ cm}^{-1}$ , which shifts to  $204\text{ cm}^{-1}$  with  $^{34}\text{S}$  (Fig. 4B, bottom) which may very well result from mixing of the Fe–S mode with a porphyrin vibration. There is some residual Fe–S vibration at  $340\text{ cm}^{-1}$  from Int-I (Fig. 4B, bottom). Thus, the Fe–S and S–O vibrations, confirmed with  $^{34}\text{S}$  labelling, clearly indicate that Int-I and Int-II have Fe–S and S–O bonds. Samples prepared with MeOD did not show any shift in any of the S–O or Fe–S vibrations, suggesting that none

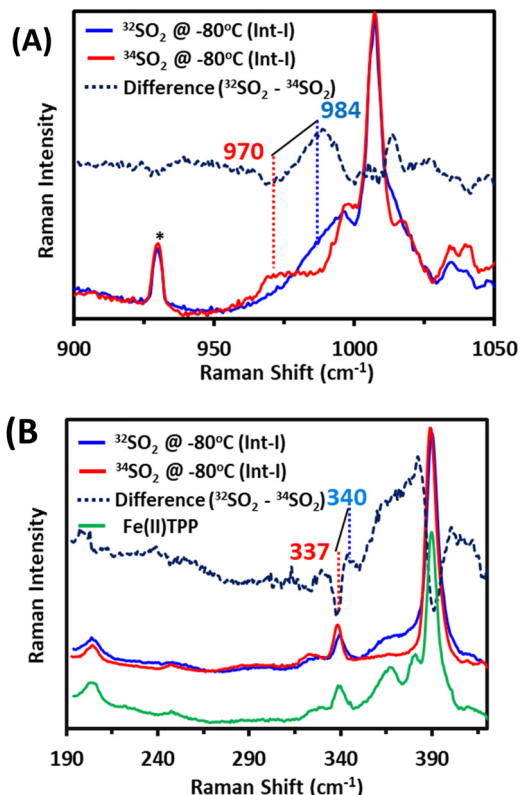


Fig. 3 Resonance Raman data on a frozen solution of Int-I (A) S–O region and (B) Fe–S region. 413.1 nm excitation and 10 mW laser power, prepared with  $^{32}\text{SO}_2$  (blue traces) and  $^{34}\text{SO}_2$  (red traces). The  $^{32}\text{SO}_2$  sample contains some unreacted  $\text{Fe(II)TPP}$  (green). \*Indicates a solvent peak.

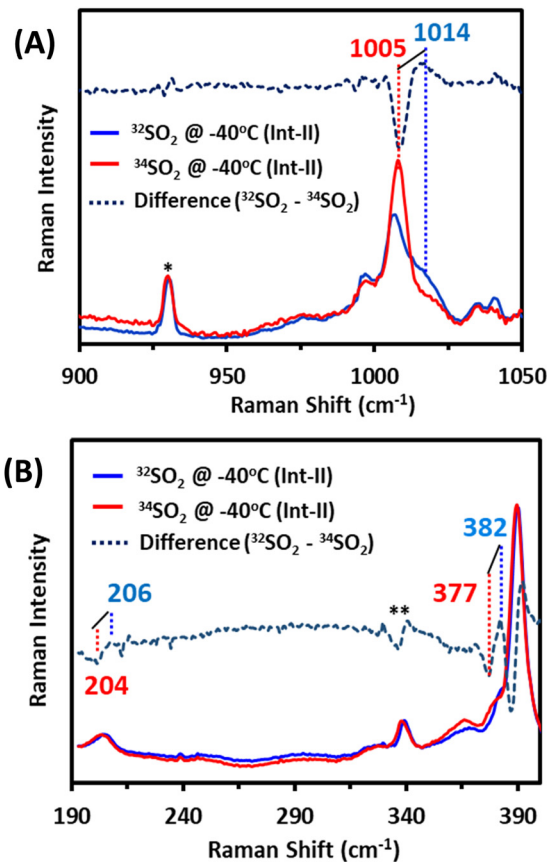
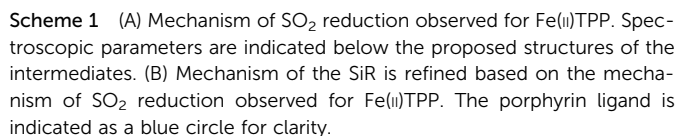


Fig. 4 Resonance Raman data on a frozen solution of Int-II (A) S–O region and (B) Fe–S region. 413.1 nm excitation and 10 mW laser power, prepared with  $^{32}\text{SO}_2$  (blue traces) and  $^{34}\text{SO}_2$  (red traces). \*Indicates a solvent peak and \*\* represents residual Int-I.

of the species are likely to be protonated. Density functional theory (DFT) calculations are used to gain further insight.

DFT calculations are used to compute the hypothetical structure of possible intermediates in the reduction of  $\text{SO}_2$  to  $\text{SO}$  by  $\text{Fe(II)TPP}$ .<sup>33,34</sup> The DFT method being used was reported to reproduce the Mössbauer and vibrational spectroscopy data of the  $[\text{Fe(III)-SO}]^+$  species quite well.<sup>28</sup> EPR data indicates that Int-I and Int-II are  $S = 1/2$   $\text{Fe(III)}$  species, and the presence of a free  $[\text{Fe(III)TPP}]^+$  indicates that  $\text{SO}_2$  is reduced by  $2e^-$  to its formal +2 oxidation state in both these species. Additionally, conversion of Int-I to Int-II requires a proton, as indicated by the EPR data obtained with MeOD and  $\text{XylH}^+\text{Cl}^-$ . The most likely description of Int-I is  $[\text{Fe(III)TPP-SO}_2]^-$  and Int-II has already been proposed to be  $[\text{Fe(III)TPP-SO}]^+$ . The possibility of the formation of  $\text{Fe(III)TPP-SO}_2\text{H}$  is removed by the lack of an H/D isotope effect in the S–O vibration. The DFT-calculated structure of an  $S = 1/2$   $[\text{Fe(III)TPP-SO}_2]^-$  with an axial MeOH (from the solvent) shows an Fe–S bond length of  $2.33\text{ \AA}$  and an S–O bond length of  $1.50\text{ \AA}$  (Table S1, ESI†). The computed S–O bond length shows a substantial increment from the reported  $1.44\text{ \AA}$  bond length of  $\text{SO}_2$  and is closer to that computed for free  $\text{SO}_2^-$  (one-electron-reduced  $\text{SO}_2$ ) consistent with the reduction of  $\text{SO}_2$ .<sup>35,36</sup> The Fe–S stretching vibration is computed to be at  $339\text{ cm}^{-1}$ , which shifts to  $337\text{ cm}^{-1}$  on  $^{34}\text{S}$  substitution. The



The mechanism of  $\text{SO}_2$  reduction by  $\text{Fe(II)TPP}$  (Scheme 1A) thus involves a  $2e^-$  step, resulting in  $\text{SO}_2^{2-}$  species bound to  $[\text{Fe(III)TPP}]^+$  to form  $[\text{Fe(III)TPP-SO}_2]^-$  Int-I, like the mechanism proposed for the SiR by Pereira and co-workers.<sup>18</sup> No  $1e^-$ -reduced intermediate could be observed even at  $-80^\circ\text{C}$ , rather an additional electron is derived from a free  $\text{Fe(II)TPP}$  to result in a  $2e^-$  reduction, which substitutes for the reduced  $[\text{Fe}_4\text{S}_4]^+$  cluster at the SiR active site. Protonation of this species leads to the cleavage of the S-O bond and formation of the  $[\text{Fe(III)TPP-SO}]^+$  species, Int-II. The reaction can proceed even at  $-40^\circ\text{C}$  with a weak proton source such as MeOH, albeit it is accelerated in the presence of  $\text{XylH}^+\text{Cl}^-$ . This indicates that the  $\text{pK}_a$  of  $[\text{Fe(III)TPP-SO}_2]^-$  is higher than that of MeOH in THF. Thus, the  $2e^-$ -reduced intermediate in the active site of an SiR, which has several arginine and lysine residues, is most likely to be  $\text{Fe(III)-SO}_2\text{H}$ . The reaction mechanism observed here suggests that the most likely mechanism of an SiR (Scheme 1B) involves the initial formation of the  $\text{Fe(III)-SO}_2\text{H}$  species (Scheme 1B) after the  $2e^-$  reduction of  $\text{SO}_2$ .<sup>18</sup> The structure of the DsrC bound to the SiR shows that cysteinyl sulfur from the DsrC is less than  $2\text{ \AA}$  away from the siroheme, and the faster step is likely to be the attack of the cysteines of the DsrC protein resulting in the trisulfide.<sup>18,19</sup>

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