ChemComm

COMMUNICATION

Substrate specificity of an oxygen dependent sulfoxide synthase in ovothiol biosynthesis†

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OvoA is an iron(II) dependent sulfoxide synthase which catalyzes the first step in ovothiol A biosynthesis. This enzyme sulphurizes the C_5 position of the imidazole side chain of L-histidine. We report the substrate specificity profile of this catalyst and present data which indicate that OvoA catalysis follows an thiol-ene type mechanism.

Ovothiol A (**1**, Fig. 1) is a thiohistidine derivative which has been discovered in sea urchin eggs[1–3] and human pathogens such as *Leishmania major* and *Trypanosoma cruzi*.[4–7] Because of its thiol function ovothiol A is characterized by a remarkably low pK_a of 1.4,[8] and an increased redox potential (-0.09 V *vs.* SHE)

compared to glutathione (-0.26 V)[8] or trypanothione (-0.24 V).[9] These distinct parameters suggest that ovothiol A occupies functional niches in cellular redox homeostasis but its precise physiological roles are unknown.[4–7,10–12]

Ovothiol A is biosynthesized from L-cysteine, L-histidine, molecular oxygen (O_2) and *S*-adenosyl methionine (SAM) (Fig. 1).[4–6] The key step in this pathway is oxidative insertion of a sulphur atom into the C_5–H bond on the histidine side chain. Subsequent elimination of the L-cysteine derived carbon scaffold from intermediate **2** (Fig. 1) and reduction of the oxidized 5-thiohistidine complete the transfer of a sulphur atom from L-cysteine to L-histidine. Recently we characterized an iron(II) dependent enzyme, OvoA, from *Erwinia tasmaniensis* which mediates this unusual oxidative sulphur transfer.[13] We also described an OvoA homolog from *Mycobacterium smegmatis*, EgtB, which is involved in ergothioneine biosynthesis (**3**, Fig. 1).[14] The mycobacterial enzyme inserts a sulphur atom into the C_2–H bond on the imidazole ring of α-*N,N,N*-trimethyl-L-histidine (**4**, Fig. 1).

The catalytic mechanism and the substrate scope of this novel class of non-heme iron enzymes are poorly understood. In the present report we demonstrate that OvoA catalyzes efficient *in vitro* sulphurization of L-histidine, D-histidine, 2-fluoro-L-histidine and compounds other than amino acids. In addition, we discuss indications that OvoA may catalyze C–S bond formation by a thiol-ene reaction mechanism,[15] in which an OvoA generated L-cysteine thiyl radical attacks the unsaturated imidazole ring of L-histidine.

To initiate this study, we produced OvoA from *E. tasmaniensis* as previously reported.[13] Typical OvoA reactions contained L-cysteine, L-histidine, 1 μM FeSO₄, 1 mM ascorbate, 50 mM Tris HCl and 50 mM NaCl. The reactions were performed at 26 °C and were monitored by HPLC at 220 nm. Addition of either ascorbate or D-isoascorbate to the reaction mixture constituted a major improvement over our previous protocols[13] because the antioxidants increased OvoA activity by nearly 100-fold (Fig. S1, ESI†). From a 800 mL reaction mixture containing 10 mg of OvoA, 1 mM L-histidine and 1 mM L-cysteine we were able to purify 100 mg of 5-L-histidyl-L-cysteine sulfoxide (**2**, Fig. 1, Fig. S2, ESI†),
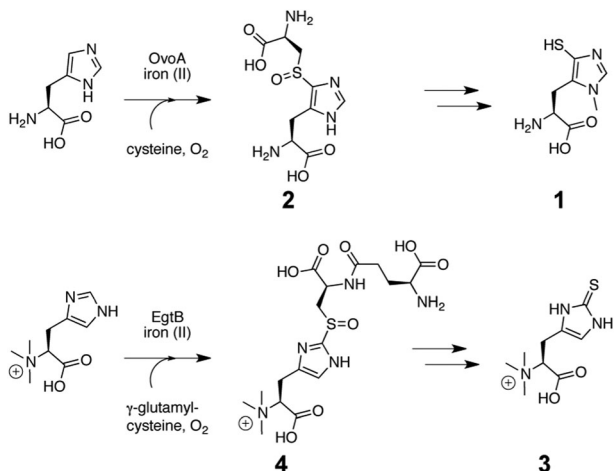
Fig. 1 Biosynthesis of Ovothiol (**1**) in *Erwinia tasmaniensis* via 5-L-histidyl-L-cysteine sulfoxide intermediate (**2**).[13] Bottom: biosynthesis of ergothioneine (**1**) in *Mycobacterium smegmatis* via a 2-*N,N,N*-α-trimethyl-L-histididyl-γ-L-glutamyl-L-cysteine sulfoxide intermediate (**4**).[14] OvoA and EgtB are the first known sulfoxide synthases.

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Table 1 Michaelis–Menten parameters of OvoA catalyzed sulphurization of imidazole derivatives[a]

	Sulfoxide product	k_{cat} [s^{-1}]	K_M [M^{-1}]	k_{cat}/K_M
Sulphur donor				
L-Cysteine	2	3.3×10^0	3.5×10^{-4}	9.4×10^3
Sulphur acceptor				
L-Histidine (5)	2	3.4×10^0	3.4×10^{-4}	1.0×10^4
Histamine (7)	26	3.6×10^{-1}	2.3×10^{-4}	1.6×10^3
L-Histidinamide (6)	25	5.7×10^{-1}	3.1×10^{-4}	1.8×10^3
D-Histidine (9)	28 and 29	7.8×10^{-1}	6.7×10^{-4}	1.2×10^3
4-Methyl imidazole (8)	27	n.a.	n.a.	1.8×10^1

[a] Reactions containing 20 mM Tris HCl pH 8.0, 20 mM NaCl, 1 uM FeSO$_4$, 0.28 uM OvoA, 2 mM tris(2-carboxyethyl)phosphine (TCEP) and 1 mM ascorbate were incubated at 26 °C. Product formation was monitored by HPLC at 220 nm. Products were identified by ESI MS (Table S1, ESI). Stated values are within an error margin of $\pm10\%$ (Fig. S3 and S6, ESI).

demonstrating than OvoA is able to deliver thiolated L-histidine at preparative scales.

OvoA catalysis is characterized by a k_{cat} of 3.3 s^{-1} and a k_{cat}/K_M of 9.4×10^3 M^{-1} s^{-1} and 1.0×10^4 M^{-1} s^{-1} for L-cysteine and L-histidine respectively (Table 1, Fig. S3, ESI†). The pH dependence of $k_{cat}/K_{M,his}$ follows a bell shaped curve with an activity maximum at pH 7.3 flanked by the kinetic pK_as 6.8 and 8.0 (Fig. S4, ESI†). Alkaline pH limits activity due to a declining k_{cat}. At lower pH $K_{M,his}$ becomes limiting. The lower kinetic pK_a coincides with the pK_a of L-histidine, suggesting that OvoA binds this substrate in deprotonated form.

Using the same kinetic assay we then profiled the substrate specificity of OvoA. We previously determined that OvoA is highly specific for L-cysteine as a sulphur donor and does not accept other thiols such as D-cysteine, N-acetyl-L-cysteine, γ-L-glutamyl-L-cysteine, glutathione, or thiophosphate.[13] In comparison, the specificity for the sulphur acceptor proved to be significantly broader. With the first compound series (6–14, Fig. 2) we probed the importance of the amino acid moiety in the sulphur acceptor, and using the second series (15–22, Fig. 2) examined whether OvoA accepts L-amino acids with alternative side chains.

L-Histidinamide (6) and histamine (7) are converted to products 25 and 26 only ten fold less efficiently than L-histidine. 4-Methyl imidazole is also a substrate demonstrating that the amino acid moiety of the sulphur acceptor is not essential for catalysis. On the other hand, we found no measurable turnover ($k_{cat}/K_M < 1$ M^{-1} s^{-1}) of 4-bromo imidazole (10), 4(5)-(hydroxymethyl)imidazole (11), urocanic acid (12), N-α-acetyl-L-histidine (13) and carnosine (14).

Amino acid modifying enzymes are usually characterized by high enantioselectivity. Therefore we were surprised that D-histidine is fairly well tolerated as a substrate by OvoA (Table 1). However, detailed HPLC and ^{1}H NMR analyses (Fig. S7, ESI†) identified the reaction product as a mixture of 63% 5-D-histidyl-L-cysteine sulfoxide (28) and 37% 2-D-histidyl-L-cysteine sulfoxide (29). We believe that the two regio-isomers result from two different binding modes of D-histidine to the OvoA active site in which either the C$_2$ or the C$_5$ position is presented to the reactive center. A preference for either binding mode may also
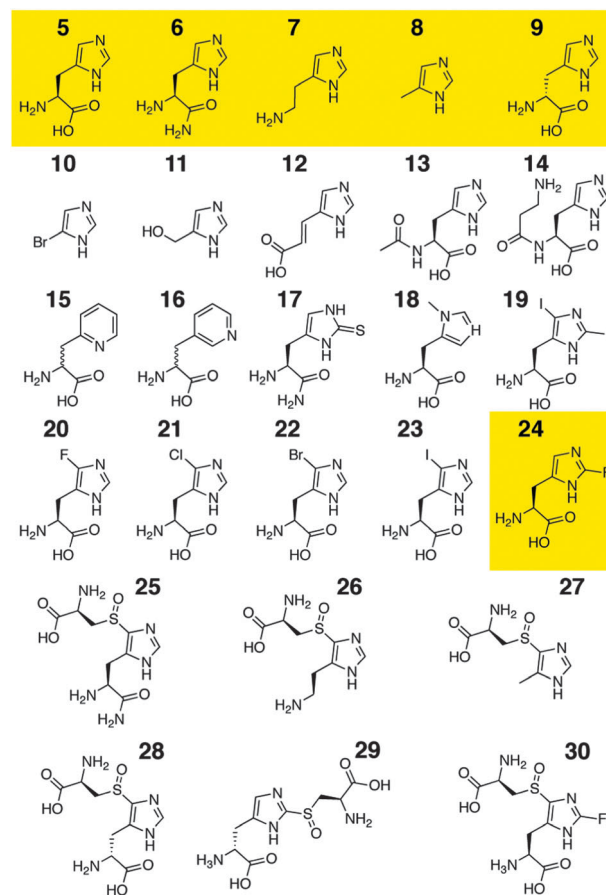

Fig. 2 L-Histidine (5) and histidine analogs (6–24) tested for OvoA catalyzed turnover. Yellow: OvoA substrates. Sulfoxides 25–30 are ESI-MS identified OvoA products (Table S1, ESI†).

explain the different regiospecificity for imidazole sulphurization in EgtB and OvoA (Fig. 1).

When we assayed OvoA with L-amino acids with imidazole derivatives as side chains we found much less substrate promiscuity. OvoA does not accept compounds 15–23 as substrates (Table 1). The remarkable exception in this series is 2-fluoro-L-histidine (24). Although an excellent L-histidine isostere,[16] 2-fluoro-L-histidine presents a much more acidic (p$K_a \approx 1$) and electron poor side chain.[16] The redox potential of 2-fluoro-L-histidine is not known, but studies on indoles and phenols suggest that fluorination can increase the redox potential of aromatic moieties by 60–200 mV.[17,18] Despite these changes OvoA converts 2-fluoro-L-histidine to 2-fluoro-5-L-histidyl-L-cysteine sulfoxide (30, HRMS: calcd 309.0669, found 309.0663) with almost the same efficiency as it converts L-histidine to 2 (Fig. S8, ESI†).

The apparent insensitivity of OvoA to the redox potential of the sulphur acceptor calls our earlier attempts at explaining the underlying catalytic mechanism into question.[13] These proposals (1–3, Fig. 3) predicted sulfoxidation of the substrate L-cysteine as a requisite step to generate the enzyme bound oxo iron(IV) species (a) which then mediates C–S bond formation (b, c or d).[13] This second step could proceed via homolytic cleavage of the imidazole C$_5$–H bond (b, mechanism 1). Because the resulting sp^2
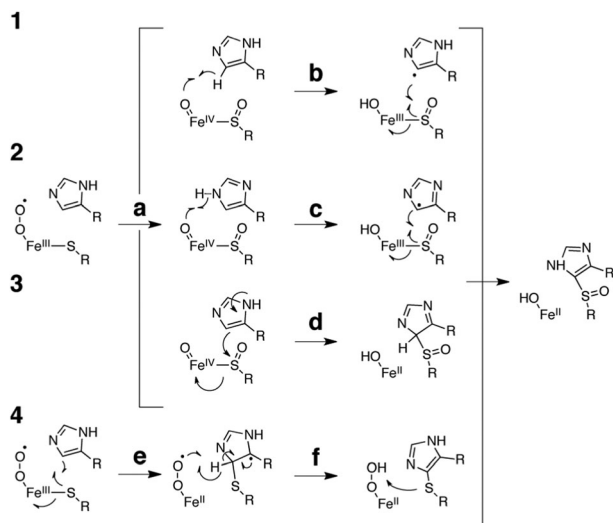
Fig. 3 Proposed catalytic mechanism for OvoA catalyzed oxidative sulphur insertion into the imidazole C_5–H bond of L-histidine. The current data are most consistent with mechanism 4.
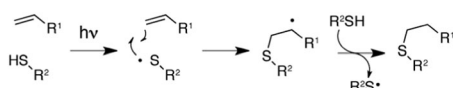
Fig. 4 Proposed mechanism of the thiol-ene reaction.[15]

radical is very unstable we would expect this to be the rate limiting step.[19] Yet, when OvoA was assayed with deuterated D/L-histidine, we could not detect any substrate kinetic isotope effect ($KIE_{substrate}$).[13]

A second mechanism proposed one-electron oxidation of the imidazole ring coupled with deprotonation of the resulting imidazyl radical cation (**c**, mechanism 2). This reaction is still a difficult and likely rate limiting task for the oxo iron(IV) species.[19] The absence of a significant solvent kinetic isotope effect ($KIE_{solvent} = 1.2 \pm 0.1$, Fig. S5, ESI†) and the observation that the electron poor 2-fluoro-L-histidine is an efficient OvoA substrate suggest that this step is either not rate limiting or does not occur. The third mechanism which implicates the imidazole ring as a nucleophile (**d**, mechanism 3) is similarly inconsistent with efficient turnover of 2-fluoro-L-histidine, and with the absence of a $KIE_{solvent}$.

A fourth mechanism could explain the present observations more consistently: in this scheme the OvoA based iron(III)-superoxide complex generates a L-cysteine thiyl radical which attacks the imidazole ring (**e**), followed by rearomatisation (**f**). Subsequent sulfoxidation of the thioether restores the ferrous state of the enzyme and concludes the catalytic cycle. Because the imidazole ring serves as an electrophilic target for the nucleophilic thiyl radical it would not be surprising that 2-fluoro-L-histidine is a well tolerated substrate. According to this mechanism C–S bond formation depends only on the presence of an unsaturated carbon on the sulphur acceptor. This scheme is reminiscent of the thiol-ene reaction which

relies on the ability of photo-generated thiyl radicals to attack olefins as a first step to thioether bond formation (Fig. 4).[15] Given the broad scope of this reaction it seems possible that engineered or evolved sulfoxide synthases can be found which can sulphurize a broad range of unsaturated hydrocarbons.

The reported data present OvoA from *E. tasmaniensis* as an efficient catalyst allowing *in vitro* preparation of the sulphurized product (**2**, Fig. 1) on a 100 mg scale. The substrate specificity profile suggests that OvoA does not require an amino acid moiety on the sulphur acceptor. OvoA converts D-histidine into a mixture of C_2 and C_5 modified products indicating that product specificity is purely a function of substrate positioning in the active site. Finally, our observation that OvoA accepts 2-fluoro-L-histidine as an efficient substrate, coupled with the absence of $KIE_{solvent}$ or $KIE_{substrate}$, point towards a catalytic mechanism which views the sulphur acceptor as the passive target of an iron(III)-superoxide generated thiyl radical. In combination, these properties present OvoA as a promising scaffold for the engineering of tailor made sulphur transferases.

Notes and references

1 A. Palumbo, M. Dischia, G. Misuraca and G. Prota, *Tetrahedron Lett.*, 1982, **23**, 3207–3208.
2 E. Turner, R. Klevit, P. B. Hopkins and B. M. Shapiro, *J. Biol. Chem.*, 1986, **261**, 3056–3063.
3 E. Turner, L. J. Hager and B. M. Shapiro, *Science*, 1988, **242**, 939–941.
4 D. J. Steenkamp and H. S. C. Spies, *Eur. J. Biochem.*, 1994, **223**, 43–50.
5 D. J. Steenkamp, D. Weldrick and H. S. C. Spies, *Eur. J. Biochem.*, 1996, **242**, 557–566.
6 R. N. Vogt, H. S. C. Spies and D. J. Steenkamp, *Eur. J. Biochem.*, 2001, **268**, 5229–5241.
7 M. R. Ariyanayagam and A. H. Fairlamb, *Mol. Biochem. Parasitol.*, 2001, **115**, 189–198.
8 K. H. Weaver and D. L. Rabenstein, *J. Org. Chem.*, 1995, **60**, 1904–1907.
9 A. H. Fairlamb and A. Cerami, *Annu. Rev. Microbiol.*, 1992, **46**, 695–729.
10 R. L. Krauth-Siegel, H. Bauer and H. Schirmer, *Angew. Chem., Int. Ed.*, 2005, **44**, 690–715.
11 C. Jacob, *Nat. Prod. Rep.*, 2006, **23**, 851–863.
12 R. L. Krauth-Siegel and A. E. Leroux, *Antioxid. Redox Signaling*, 2012, **17**, 583–607.
13 A. Braunshausen and F. P. Seebeck, *J. Am. Chem. Soc.*, 2011, **133**, 1757–1759.
14 F. P. Seebeck, *J. Am. Chem. Soc.*, 2010, **132**, 6632–6633.
15 C. E. Hoyle and C. N. Bowman, *Angew. Chem., Int. Ed.*, 2010, **49**, 1540–1573.
16 D. S. Wimalasena, J. C. Cramer, B. E. Janowiak, S. J. Juris, R. A. Meinyk, D. E. Anderson, K. L. Kirk, R. J. Collier and J. G. Bann, *Biochemistry*, 2007, **46**, 14928–14936.
17 T. Liu, P. R. Callis, B. H. Hesp, M. de Groot, W. J. Buma and J. Broos, *J. Am. Chem. Soc.*, 2005, **127**, 4104–4113.
18 M. R. Seyedsayamdost, S. Y. Reece, D. G. Nocera and J. Stubbe, *J. Am. Chem. Soc.*, 2006, **128**, 1569–1579.
19 E. A. Bushnell, G. B. Fortowsky and J. W. Gauld, *Inorg. Chem.*, 2013, **51**, 13351–13356.