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Calcium Ions Tune the Zinc-Sequestering Properties and Antimicrobial Activity of Human S100A12

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Abstract

Human S100A12 is a host-defense protein expressed and released by neutrophils that contributes to innate immunity. Apo $S100A12$ is a 21-kDa antiparallel homodimer that harbors two $Ca(II)$ binding EF-hand domains per subunit and exhibits two His₃Asp motifs for chelating transition metal ions at the homodimer interface. In this work, we present results from metal-binding studies and microbiology assays designed to ascertain whether $Ca(II)$ ions modulate the $Zn(II)$ -binding properties of S100A12 and further evaluate the antimicrobial properties of this protein. Our metal depletion studies reveal that $Ca(\Pi)$ ions enhance the ability of S100A12 to sequester $Zn(\Pi)$ from microbial growth media. We report that human S100A12 has antifungal activity against *Candida albicans, C. krusei, C. glabrata* and *C. tropicalis*, all of which cause human disease. This antifungal activity is $Ca(II)$ -dependent and requires the $His₃Asp$ metal-binding sites. We expand upon prior studies of the antibacterial activity of S100A12 and report Ca(II)-dependent and strainselective behavior. S100A12 exhibited *in vitro* growth inhibitory activity against *Listeria monocytogenes*. In contrast, S100A12 had negligible effect on the growth of *Escherichia coli* K-12 and *Pseudomonas aeruginosa* PAO1. Loss of functional ZnuABC, a high-affinity Zn(II) import system, increased the susceptibility of *E. coli* and *P. aeruginosa* to S100A12, indicating that S100A12 deprives these mutant strains of $Zn(II)$. To evaluate the $Zn(II)$ -binding sites of S100A12 in solution, we present studies using Co(II) as a spectroscopic probe and chromophoric small-molecule chelators in $Zn(II)$ competition titrations. We confirm that $S100A12$ binds $Zn(II)$ with a 2:1 stoichiometry, and our data indicate sub-nanomolar affinity binding. Taken together, these data support a model whereby $S100A12$ uses $Ca(II)$ ions to tune its $Zn(II)$ -chelating properties and antimicrobial activity.

Graphical abstract

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Electronic Supplementary Information

Complete experimental methods, Tables S1–S14, Figures S1–S10.

Introduction

Neutrophils are an important cellular component of the innate immune system.^{1,2} These white blood cells are recruited to sites of infection and release antimicrobial factors into the extracellular space that include reactive oxygen and nitrogen species, proteases, antimicrobial peptides and metal-chelating proteins. In this work, motivated by the need to further elucidate the interplay between neutrophils, metal homeostasis, and microbial pathogenesis, we combine chemistry and biology to investigate a $Ca(II)$ - and $Zn(II)$ -binding protein named S100A12 that is housed in the neutrophil cytoplasm and deployed at sites of infection.

S100A12 (also termed calgranulin C, P6, CGRP, CO-Ag, CAAF-1, EN-RAGE) was first isolated from porcine granulocytes in 1994 and subsequently identified in specimens from other mammals.^{3–9} In 1995, the serendipitous discovery of $S100A12$ in human neutrophils was reported;⁴ this work revealed that S100A12 constitutes \approx 5% of total cytosolic proteins in resting neutrophils, and further investigations provided its amino acid sequence and gene structure.^{10,11} S100A12 is associated with the inflammatory response and a variety of human pathologies that include disorders of the gastrointestinal tract (e.g. Crohn's disease, ulcerative colitis),¹² type 2 diabetes,¹³ arthritis,¹⁴ Alzheimer's disease,¹⁵ and cystic fibrosis.¹⁶ S100A12 can interact with the receptor for advanced glycation end products $(RAGE)^{17,18}$ and toll-like receptor 4 (TLR4), ¹⁹ and the latter interaction is implicated in S100A12 activation of proinflammatory signaling in human monocytes. Moreover, S100A12 likely contributes to host defense against invading pathogens.^{9,20} Evidence for anti-parasitic activity was first suggested during studies of *Onchocerca volvulus* extracts obtained from tissues of onchocerciasis patients.⁹ These specimens were contaminated with various neutrophil proteins, including S100A12. Subsequent investigations demonstrated that S100A12 possesses *in vitro* anti-parasitic activity against the parasite *Brugia malayi*. 21 Recently, elevated S100A12 levels were observed in gastric biopsy samples from patients colonized with *Helicobacter pylori*. ²⁰ Moreover, *in vitro* studies revealed that S100A12 has antibacterial activity against *H. pylori* and inhibits the activity of its Cag type-IV secretion system, both of which involve $Zn(II)$ chelation by S100A12.²⁰ Taken together, these clinical and biological investigations indicate that S100A12 is an abundant and important player in human biology, and provide multiple avenues for further exploration at the molecular and physiological levels. From the perspective of metals in biology, understanding how S100A12 contributes to metal homeostasis in broad terms, and elucidating whether its metal-bound forms are relevant players in human health and disease, is of particular interest.

Human S100A12 (92-residues, 10.5 kDa monomer) is a member of the S100 family of Ca(II)-binding peptides that harbors two EF-hand domains and forms homooligomers. Five crystal structures of human S100A12 are available and provide insight into its oligomerization properties and coordination chemistry.^{22–25} The first crystal structure revealed a homodimer with four $Ca(II)$ ions bound at the EF-hand domains.²² This structure also revealed two His₃Asp motifs formed at the homodimer interface, which are comprised of His15 and Asp25 of one subunit and His85 and His89 of the other subunit. The crystal structures of Cu(II)- and Ca(II)-bound S100A12 and Zn(II)-bound S100A12 established a 2:1 M(II):S100A12 homodimer stoichiometry ($M = Cu$, Zn) with each M(II) ion coordinated

by a His₃Asp motif (Figure 1).²⁴ Comparison of the available apo and holo structures shows that metal binding modulates the oligomerization properties of $\text{S100A12}.^{22-25}$ For instance, apo S100A12 crystallized as a dimer whereas the Zn(II)-bound form crystallized as a tetramer.²⁵ Thereafter, solution studies showed that $Zn(II)$ and $Ca(II)$ binding to S100A12 provide varying oligomeric states that include dimers, tetramers and hexamers.²⁶ These studies exemplify the complex speciation of S100A12, and suggest a model whereby different forms afford functional diversity.²⁷

Beyond crystallographic characterization of the Cu(II)- and Zn(II)-bound forms, only limited information about how S100A12 coordinates Zn(II) and other first-row transition metal ions in solution is available, and some of the data appear contradictory. A seminal investigation of porcine S100A12 revealed stoichiometric Zn(II) binding (K_d < 10 nM), and found that Zn(II) coordination enhanced the affinity of this protein for Ca(II) by several orders of magnitude.³ More recently, a study of $Zn(II)$ binding by human S100A12 reported K_d values for Zn(II) in the micromolar range ($K_{d1} = 16 \mu M$, $K_{d2} = 83 \mu M$).²⁶ Given that the pig and human forms share 70% sequence identity and have conserved His3Asp motifs, we had difficulty reconciling these data. Moreover, the study of human S100A12 in *H. pylori* infection supported a role for this protein in $Zn(II)$ sequestration,²⁰ a function that requires formation of a high-affinity Zn(II) complex.

These observations indicate that re-evaluation of the $Zn(\Pi)$ -binding properties of human S100A12 is necessary. Moreover, S100A12 chelates both Ca(II) and Zn(II) at different sites (Figure 1), and whether Ca(II) ions influence the Zn(II)-binding properties of S100A12 is unknown. In prior work, we discovered that human calprotectin (S100A8/S100A9 oligomer) uses Ca(II) to modulate its M(II)-binding ($M = Mn$, Fe, Zn) properties and exhibits enhanced M(II) affinities in the presence of high Ca(II) concentrations.^{28–30} We hypothesized that S100A12 may exhibit similar behavior given its accepted function as an extracellular $Zn(II)$ -chelating protein.^{20,26}

Many outstanding questions regarding the antimicrobial activity and host-defense function of S100A12 also remain. To the best of our knowledge, no extensive studies detailing the *in vitro* antimicrobial activity of S100A12 are available in the literature. This situation contrasts the wealth of data reported for other factors produced by human neutrophils, including calprotectin and the α-defensins.31,32 The limited *in vitro* data available for *Staphylococcus aureus* and *H. pylori* indicate that human S100A12 has strain-selective antibacterial activity and support its contribution to the host metal-withholding response.20,32

In this work, we address these issues and examine the metal-binding properties and antimicrobial activity of human S100A12 under conditions of low and high Ca(II). In the presence of excess Ca(II), S100A12 readily sequesters Zn(II) from standard microbial growth media, exhibits growth inhibitory activity against a variety of pathogenic fungi, and provides antibacterial activity against the human gastrointestinal pathogen *Listeria monocytogenes*. Our data support a model where S100A12 responds to the local Ca(II) concentration to modulate its Zn(II)-sequestering ability and antimicrobial activity, and effectively turn on these functions in the Ca(II)-rich environment of the extracellular space.

Results and Discussion

Human S100A12 Depletes Zn(II) from Microbial Growth Medium

To probe which metal ions S100A12 sequesters from microbial growth medium, we conducted a series of metal depletion experiments (Figure 2, Tables S4–S13, Supplementary Information). We treated standard growth media employed for *in vitro* antifungal (YPDbased medium) or antibacterial (TSB-based medium) activity assays with varying concentrations of S100A12 (0 – 250 µg/mL, t = 20 h, T = 30 °C). Following treatment of the medium with S100A12, we employed spin filtration to separate the protein from the treated medium. We quantified the amount of unbound metal in the treated medium by inductively coupled plasma-mass spectrometry (ICP-MS). Because the protein and treated medium were separated by spin filtration, this procedure does not afford an equilibrium measurement because the metal concentrations in the filtered sample are not precisely the "free" metal concentrations in a solution at equilibrium. Nevertheless, it provides an assessment of which metal ions can be sequestered by S100A12 in complex samples. To determine whether the presence of Ca(II) influences metal sequestration, we prepared and analyzed samples with and without a \approx 2-mM Ca(II) supplement.

The YPD/Tris medium employed for antifungal activity assays (*vide infra*) contains relatively high concentrations of Fe (\approx 3.5 µM) and Zn (\approx 7 µM), and lower concentrations of Mn (\approx 100 nM), Co (\approx 130 nM), Ni (\approx 100 nM) and Cu (\approx 150 nM) (Figure 2A, Table S4). With the exception of Zn, S100A12 treatment had negligible effect on the concentrations of the first-row transition metals present in the YPD/Tris medium. Zinc depletion occurred, and the extent of Zn depletion depended on the concentration of S100A12 and whether the medium was supplemented with $Ca(\Pi)$. The $Ca(\Pi)$ effect was most obvious at 62.5 and 125 µg/mL S100A12. For instance, at 125 µg/mL S100A12, the Zn concentration in the treated medium decreased by a factor of \approx 3 (−Ca) or \approx 10 (+Ca).

The TSB/Tris medium employed for antibacterial activity assays (*vide infra*) also contains relatively high concentrations of Fe (\approx 3 µM) and Zn (\approx 5 µM), and the other first-row transition metals are markedly less abundant (Figure 2B, Table S9). S100A12 treatment resulted in Ca(II)-dependent depletion of Zn, but not the other transition metals, from this medium. The Ca(II) effect was most pronounced at 62.5 µg/mL S100A12 where negligible Zn(II) depletion occurs in the absence of the Ca(II) supplement and \approx 4-fold reduction in total Zn occurs in the presence of Ca(II).

To confirm that the His₃Asp sites are essential for $Zn(II)$ depletion, we treated YPD/Tris and TSB/Tris with S100A12 His₃Asp (250 μ g/mL), a variant that harbors the point mutations H15A, D25A, H85A, and H89A. This variant cannot bind metal ions at the $His₃Asp$ sites because the four metal-chelating residues are replaced by non-coordinating alanine moieties (Supplementary Information). No metal depletion was observed for the $His₃Asp$ variant in either YPD/Tris or TSB/Tris (Figure 2). These results demonstrate that the His $_3$ Asp sites are essential for Zn(II) depletion from microbial growth media, as expected based on the Zn(II)- S100A12 crystal structure.²⁵

These metal depletion results illuminate three key points about the metal-sequestering properties of S100A12: (i) S100A12 binds Zn(II) with sufficient affinity to deplete it from growth medium; (ii) S100A12 appears to select for Zn(II) over other first-row transition metal ions, at least under these experimental conditions; and (iii) S100A12 uses Ca(II) ions to modulate its $Zn(\Pi)$ -binding properties, and the presence of $Ca(\Pi)$ results in more effective Zn depletion from the medium. Moreover, the TSB/Tris metal-depletion profile of S100A12 is markedly different from that of calprotectin, a metal-sequestering S100 family member that harbors two different sites for transition metal ions (His₃Asp and His₄/His₆) and reduces the levels of Mn, Fe, Ni, Cu, and Zn from this medium.³⁰ The apparent $Zn(II)$ selectivity of S100A12 is consistent with prior solution studies of this protein, which provided no evidence for the formation high-affinity Mn(II)-S100A12 and Fe(II)-S100A12 complexes in the absence or presence of $Ca(II)$.^{30,33} The selective $Zn(II)$ depletion is also reminiscent of studies of a His₄ variant of calprotectin that only harbors an interfacial His3Asp site for chelating transition metals. This variant only depleted $Zn(II)$ from TSB/Tris mixtures.³⁰

Human S100A12 Exhibits Calcium-Dependent Antifungal Activity

Fungi have a significant $Zn(II)$ requirement, and $Zn(II)$ acquisition by fungal pathogens is important for virulence.^{34–40} We therefore questioned whether $S100A12$ exhibits antifungal activity that results from its ability to bind Zn(II). To the best of our knowledge, an antifungal activity for S100A12 has not been reported. We evaluated the growth inhibitory activity of S100A12 against four *Candida* strains that cause human disease. *C. albicans* is a commensal organism of the human gastrointestinal tract and opportunistic human pathogen that can invade the mucosa and cause bloodstream infections.41,42 *C. krusei, C. glabrata*, and *C. tropicalis* are emerging human health threats, form biofilms, and exhibit resistance to antifungals such as the fluconazoles. $43-45$ When cultured in the YPD/Tris medium employed in the metal-depletion studies (Figure 2), each fungal strain grew to an OD_{600} value of ≈ 0.6 in the absence and presence of a \approx 2-mM Ca(II) supplement (Figure 3, Figure S5). Addition of S100A12 (0 – 500 µg/mL) to *C. albicans* resulted in concentration-dependent growth inhibition, and this effect was enhanced when $Ca(\Pi)$ was present in the medium (Figure 3A). The Ca(II) enhancement was most striking at 125 μ g/mL S100A12 where the OD₆₀₀ value decreased to ≈ 0.3 in the absence of Ca(II), and negligible *C. albicans* growth was observed when Ca(II) was included in the medium.

S100A12 also displayed Ca(II)-dependent growth inhibitory activity against *C. krusei, C. glabrata*, and *C. tropicalis* (Figure 3B–D). As observed for *C. albicans*, the presence of the Ca(II) supplement enhanced the antifungal activity of S100A12 against these three strains. Moreover, strain-to-strain variability in S100A12-mediated growth inhibition occurred in both the absence and presence of the Ca(II) supplement. In the absence of a Ca(II) supplement, *C. albicans* and *C. glabrata* were more susceptible to S100A12 than *C. krusei* and *C. tropicalis*. Indeed, neither *C. krusei* nor *C. tropicalis* exhibited growth inhibition in the presence of 500 µg/mL S100A12 when the Ca(II) supplement was omitted, whereas *C. albicans* and *C. glabrata* growth was inhibited to varying degrees at this S100A12 concentration. Taken together, these data establish that (i) S100A12 exhibits *in vitro* antifungal activity against a number of *Candida* strains, (ii) enhanced fungal growth

inhibition occurs under conditions of high Ca(II), and (iii) the susceptibility of *Candida* spp. to S100A12 is strain-dependent.

The His3Asp Motifs Are Essential for Antifungal Activity

We reasoned that the Ca(II)-dependent antifungal activity of $S100A12$ results from $Zn(II)$ sequestration. We performed two additional assays to test this hypothesis and evaluate the requirement of the His3Asp sites for the antifungal activity against *C. albicans*. First, to determine whether the His3Asp sites are essential, we compared the antifungal activity of S100A12 and the His3Asp variant (Figure 4A). This assay revealed that S100A12 His₃Asp (1 mg/mL, +Ca) has no effect on *C. albicans* growth relative to the untreated control. Next, to confirm that the antifungal activity originates from apo His₃Asp sites of S100A12 and not a transition-metal-bound form, we pre-incubated S100A12 with two equivalents of $Zn(II)$ to afford $Zn(II)$ -S100A12. Similar to the His₃Asp variant, $Zn(II)$ -S100A12 (125 µg/mL, +Ca) did not inhibit *C. albicans* growth (Figure 4B).

A comparison of the S100A12 concentrations (125 µg/mL, ≈5 µM S100A12 or ≈10 µM His₃Asp sites) required for maximum antifungal activity with the Zn(II) content (\approx 7 µM) in the YPD/Tris medium is informative. This comparison indicates that an excess of $Zn(I)$ binding sites relative to the $Zn(\text{II})$ content in the growth medium is required for S100A12 to exert its full growth inhibitory activity against *Candida* spp. under these assay conditions. The metal depletion studies indicated that 125 µg/mL S100A12 reduces the Zn level in YPD/Tris to baseline (\approx 620 nM) when Ca(II) is present in the medium. Taken together, these observations suggest that the *Candida* strains evaluated in this work compete with S100A12 for Zn(II) available in the growth medium.

Human S100A12 Exhibits Strain-Specific Antibacterial Activity

To further evaluate the *in vitro* antibacterial activity of human S100A12, we examined its growth inhibitory activity against a panel of five bacterial strains that included *E. coli* K-12, *Pseudomonas aeruginosa* PAO1, *S. aureus* ATCC 25923, *Listeria monocytogenes* ATCC 19115, and *Lactobacillus plantarum* WCSF1. Because a prior investigation reported that S100A12 lacked antibacterial activity against *S. aureus* Neuman,³² we screened for bacterial growth inhibition using a high concentration (1000 µg/mL) of S100A12 in TSB/Tris or MRS/Tris (for *L. plantarum* only) medium with or without a \approx 2-mM Ca(II) supplement (Figure 5). In agreement with prior work,³² we observed that S100A12 (\pm Ca) had no effect on *S. aureus* ATCC 25923 growth. Likewise, S100A12 afforded negligible growth inhibition for *E. coli* K-12 or *P. aeruginosa* PAO1. In contrast, cultures of *L. monocytogenes* and *L. plantarum* treated with S100A12 exhibited Ca(II)-dependent growth inhibition. Negligible growth of *L. monocytogenes* occurred, and the growth of *L. plantarum* was markedly attenuated, when the medium contained the Ca(II) supplement. On the basis of the metal depletion studies, we reason that the strains susceptible to S100A12 are more sensitive to Zn(II) deprivation. Future studies are required to address this notion on a strain-by-strain basis. We note that the inhibited growth of *L. plantarum* by S100A12 is in general agreement with our prior work, which found that the His₃Asp site of calprotectin accounts for the full antibacterial activity against this strain *in vitro*. ³⁰ Moreover, the negligible activity of S100A12 against *S. aureus* and *E. coli* is reminiscent of the calprotectin His₄

variant, which only has a His₃Asp site at the S100A8/S100A9 interface and exhibits attenuated antibacterial activity against *S. aureus* and *E. coli* in mixed TSB/Tris medium.28,32

Bacteria with Defective Zinc Acquisition Machinery Are Susceptible to S100A12

To evaluate whether bacterial strains with defective $Zn(II)$ uptake machinery exhibit enhanced susceptibility to S100A12, we performed antibacterial activity assays with Δ*znuA* mutants of *E. coli* K-12 and *P. aeruginosa* PAO1 obtained from the Keio Collection⁴⁶ and Seattle *Pseudomonas aeruginosa* PAO1 Transposon Mutant Library, ⁴⁷ respectively. ZnuABC is a high-affinity $Zn(II)$ uptake system employed by Gram-negative bacteria.^{48–51} ZnuA is the periplasmic $Zn(II)$ -binding protein that captures $Zn(II)$ and delivers the ion to ZnuBC, which are located in the inner membrane.⁵² Bacterial strains with defective ZnuABC transport machinery are more susceptible to Zn(II) deprivation than their native counterparts.^{49, 53} Indeed, S100A12 completely attenuated the growth of *E. coli znuA* at 125 µg/mL in both the absence and presence of a Ca(II) supplement (Figure 6). *P. aeruginosa znuA* also exhibited enhanced sensitivity to S100A12; however, the growth inhibition was only partial and required the presence of $Ca(\Pi)$ in the medium (Figure S6). A similar observation was reported in studies examining the *in vitro* antimicrobial activity of calprotectin against *P. aeruginosa* PA14.51 In this prior work, calprotectin did not provide complete growth inhibition of the wild-type strain or *znuA* mutant, but the growth of the *znuA* strain was impaired relative to the wild-type strain.⁵¹ These results afforded a new

hypothesis where *P. aeruginosa* PA14 has additional and as-yet undiscovered Zn(II) acquisition machinery that allows for $Zn(\Pi)$ uptake in the absence of $ZnuA$.⁵¹ Subsequent investigations of *P. aeruginosa* PAO1 and a *znuA* strain cultured under conditions of low Zn(II) revealed the expression of genes associated with additional import pathways associated with $Zn(\Pi)$ acquisition as well as up-regulation of genes that encode $Zn(\Pi)$ independent proteins, which prokaroytes can employ in place of Zn(II)-containing proteins when $Zn(II)$ is limited (e.g., $Zn(II)$ -independent ribosomal proteins).⁵⁴ The current results with S100A12 are also consistent with these studies.

Human S100A12 Is a High-Affinity Zinc Chelator

The metal-depletion and antimicrobial activity assays presented in this work indicate that $S100A12$ coordinates $Zn(II)$ with sufficient affinity to sequester the ion and that $Ca(II)$ ions accentuate this function. Prior studies have examined the Zn(II)-binding properties of S100A12 in solution. In seminal characterization of porcine S100A12, metal-binding titrations that monitored the intrinsic fluorescence of S100A12 revealed stoichiometric $Zn(II)$ binding $(K_{d,Zn} \quad 10 \text{ nM})$.³ More recently, intrinsic fluorescence was employed to study $Zn(II)$ chelation by human S100A12.²⁶ In this work, titration of human S100A12 with Zn(II) resulted in a gradual change in protein emission, and this titration curve was fit to afford K_d _{Zn} values of 16 and 83 µM for the two His₃Asp sites.²⁶ It was difficult for us to reconcile how the Zn(II) dissociation constants for these two orthologues could vary by several orders of magnitude, and our conclusions from the metal-depletion studies and antifungal activity assays are contrary to the prior study of the human orthologue. As a consequence, we decided to further evaluate the metal-binding properties of human

S100A12 in solution using techniques we employed to characterize the Zn(II)-binding properties of calprotectin.²⁸

We first employed $Co(II)$ as a probe for the His₃Asp sites because it exhibits rich spectroscopic properties as a result of its $3d^7$ electronic configuration.⁵⁵ Addition of Co(II) to S100A12 caused the solution to change from colorless to pink (75 mM HEPES, 100 mM NaCl, pH 7.0), and the optical absorption spectrum of Co(II)-S100A12 exhibited ligand field transitions centered at 563 nm ($\varepsilon = 820 \text{ M}^{-1} \text{cm}^{-1}$) (Figure 7A). This spectroscopic signature is reminiscent of four- or five-coordinate $Co(II)$ species,⁵⁵ including $Co(II)$ -calprotectin where the optical features in the $d-d$ range are dominated by Co(II) bound at the His₃Asp site.28 Moreover, Co(II)-binding titrations revealed that the intensity of this signal increased until two equivalents of $Co(II)$ per S100A12 homodimer were added. The sharp inflection point indicates a 2:1 Co(II)/S100A12 stoichiometry (Figure 7B), in agreement with two His3Asp sites per homodimer. The Co(II)-binding studies were performed in the absence and presence of excess Ca(II) and comparable spectra were obtained under both sets of conditions (Figure S7).

Addition of Zn(II) to Co(II)-S100A12 resulted in loss of the absorption feature (Figure S8), which indicates that S100A12 binds $Zn(II)$ with higher affinity than Co(II). This behavior is expected based on the Irving-Williams series.⁵⁶ To confirm the Zn(II)-S100A12 stoichiometry and further investigate the Zn(II) affinity, we successfully competed S100A12 with employed three different chromophoric Zn(II) chelators. Zincon is a colorimetric Zn(II) indicator that with a dissociation constant in the micromolar range ($K_{d,Zn} \approx 10 \mu M$).^{57,58} In both the absence and presence of excess Ca(II), no change in Zincon absorbance occurred until after two equivalents of Zn(II) per S100A12 homodimer were added, demonstrating that S100A12 outcompeted Zincon for Zn(II) (Figure 7C). These titrations confirmed the expected 2:1 Zn(II):S100A12 homodimer stoichiometry and indicated that S100A12 coordinates Zn(II) with sub-micromolar affinity. Next, we examined the competition of S100A12 with the fluorescent Zn(II) sensors MagFura-2 ($K_{d, Zn} \approx 20$ nM)^{59,60} and FluoZin-3 ($K_{d,Zn}$ = 9 nM).⁶¹ These experiments were performed only in the absence of Ca(II) because both MagFura-2 and FluoZin-3 exhibit a fluorescence response to this divalent cation. S100A12 outcompeted MagFura-2 (Figure 7D) as well as FluoZin-3 (Figure S10), suggesting that S100A12 coordinates two equivalents of $Zn(II)$ with low- or subnanomolar affinity in the absence of Ca(II). Taken together, the outcomes of these $Zn(II)$ competition titrations are in general agreement with prior characterization of porcine S100A12 where stoichiometric Zn(II) binding ($K_{d,Zn}$ < 10 nM) was observed,³ as well as the metal-depletion studies presented in this work.

The metal depletion studies and antimicrobial activity assays indicate that the $Zn(II)$ -binding properties of $S100A12$ are modulated by Ca(II) ions. In prior studies of the Zn(II)-binding properties of human calprotectin, we employed a Ca(II)-insensitive Zn(II) sensor named Zinpyr-4 (ZP4, $K_{d,Zn}$ = 650 pM),⁶² and discovered that CP exhibits enhanced Zn(II) affinity in the presence of $Ca(II).^{28}$ We sought to employ the same strategy to further elucidate how Ca(II) modulates Zn(II) chelation by S100A12; however, our preliminary S100A12/ZP4 experiments were plagued by artifacts that precluded using ZP4 in a quantitative or reliable manner (data not shown). In particular, mixtures of ZP4, S100A12, and Zn(II) exhibited

unstable emission. The origin of this confounding behavior is unclear, and various issues can arise when employing competitors in metal-binding titrations.⁶³ We note that formation of the Zn(II)-S100A12 complex contributed to the problem because stable emission was observed for ZP4/S100A12/Zn(II) mixtures that contained the His3Asp variant. As a result, it is possible that (i) formation of ternary complexes involving Zn(II)-bound S100A12 and ZP4 occurred, and/or (ii) Zn(II)-dependent oligomerization of S100A12 causes ZP4 to associate with the protein. Regarding the latter scenario, the self-association properties of S100A12 are complex. Zinc binding is reported to cause the homodimer to self-associate and form tetramers and hexamers, 26 and the Zn(II)-S100A12 crystal structure revealed exposure of a hydrophobic patch with $Zn(II)$ binding.²⁵ On the basis of the metal-depletion studies and antimicrobial activity assays, we expect that the presence of Ca(II) ions increases the binding affinity of S100A12 for Zn(II), as we discovered for calprotectin,²⁸ and as-yet unidentified competitors or alternative methods are required to probe this notion further.

Human S100A12, Metal Homeostasis and Innate Immunity

This work informs current understanding of how S100A12 contributes to metal homeostasis and innate immunity. We present the discovery that S100A12 uses Ca(II) ions to modulate its Zn(II)-sequestering properties and antimicrobial activity. We also report the discovery that S100A12 has *in vitro* antifungal activity. We hypothesize this activity results from the fungi succumbing to $Zn(I)$ deprivation. These studies provide a guide for future investigations of human S100A12 in the contexts of the host/microbe interaction, metal homeostasis, and innate immunity.

S100A12 is co-packaged with its fellow metal-sequestering S100 protein calprotectin in the cytoplasm of human neutrophils, which indicates that the proteins are co-released at sites of infection and inflammation. Our current work illuminates similarities and differences between the metal-chelating properties of these two human S100 proteins. Both employ Ca(II) ions to tune the affinities for transition metal ions, which are bound at sites that form at the interface between two subunits. The heterooligomeric nature of calprotectin affords two different sites at the $\text{S}100\text{A}8/\text{S}100\text{A}9$ interface.⁶⁴ One site is a His₃Asp motif similar to the sites of S100A12, whereas the other is an unusual histidine-rich motif. Like S100A12, the His₃Asp site of calprotectin binds $Zn(II)$ with high affinity and has markedly lower affinity for Mn(II) and Fe(II).^{28–30} The histidine-rich motif was first identified as a His₄ motif in the crystal structure of Ca(II)-bound calprotectin and later determined to provide a hexahistidine coordination environment for Mn(II) and Fe(II).^{30,32,33,65} Whether this site coordinates Zn(II) using the hexahistidine motif is currently unclear. Regardless, the histidine-rich site coordinates $Mn(\Pi)$, Fe(II) and $Zn(\Pi)$ with high affinity and the metal selectivity of this site $(Mn(II) < Fe(II) < Zn(II))$ is in agreement with the Irving-Williams series.⁵⁶ We recently proposed that kinetics also contribute to metal sequestration at this site, enabling calprotectin to retain bound $Mn(II)$ and Fe(II).³⁰ On the basis of the current studies, we now question whether $Zn(II)$ chelation by S100A12 reduces the $Zn(II)$ available for capture by calprotectin in extracellular milieu and thereby boosts the ability of calprotectin to capture other nutrient metals like $Mn(II)$ and $Fe(II)$ at the $His₆ site$.

S100A12 is often described as a $Cu(II)$ - and $Zn(II)$ -chelating protein, as evidenced by crystallographic characterization (Figure 1). The YPD/Tris and TSB/Tris growth medium employed in the metal depletion studies contains markedly greater levels of $\text{Zn} \approx 5 \mu\text{M}$) relative to Cu (\approx 100 nM), and we observed negligible depletion of Cu for either medium as a result of S100A12 treatment. This latter result may stem from the relative metal concentrations in the growth medium, and additional investigations are required to clarify the origin of this observation. In particular, investigations of $Cu(II)$ -S100A12 are limited and the Cu(II) affinity of S100A12 is unknown, making further solution studies of copper binding to S100A12 an important avenue for future work. We note here that a proposal in which Cu-S100A12 generates reactive oxygen species and thereby affords anti-parasitic activity has been put forth.^{24, 27} To the best of our knowledge, the reported experimental support for this proposal is insufficient and limited to the fact that S100A12 can bind Cu(II).

From the perspective of human S100A12 structure and function, the results presented in this work augment prior studies that evaluated the biophysical properties of this protein. The reported solution studies and crystallographic characterization of apo and metal-bound S100A12 delineate that the protein has the capacity to exist in many different states.²²⁻²⁷ This complex speciation may give S100A12 functional diversity that is important for cell signaling and the immune response.^{18,26,66} That data presented in this work that reveal S100A12 uses Ca(II) binding to its EF-hand domains to tune its Zn(II)-chelating properties at the His3Asp sites and add another dimension to this behavior. This observation is reminiscent of studies of porcine S100A12, which showed that the presence of Zn(II) increased the Ca(II) affinity by several orders of magnitude.³

Conditions of high $Ca(II)$ enhance the $Zn(II)$ -sequestering ability and antimicrobial activities of S100A12. These observations support a model whereby S100A12 responds to local Ca(II) concentrations to tune its Zn(II)-binding properties and are consistent with its function as an extracellular Zn(II)-chelating protein. The extracellular space contains Ca(II) ions at concentrations that are orders of magnitude greater than those found in the cytoplasm of resting cells.⁶⁷ We reason that the low Ca(II) levels in in resting neutrophils allow S100A12 to maintain an apo state in the cytoplasm, and the high Ca(II) levels in the extracellular space allow S100A12 to sequester $Zn(\Pi)$ once it is deployed at a site of infection or inflammation. In prior work, we put forth the same model for the behavior of calprotectin. Whether this model is general for all S100 proteins that chelate transition metals at interfacial sites is a topic for future investigation.

Experimental Section

Complete experimental methods are provided as Supplementary Information.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Figure 1.

Crystal structures of metal-bound human S100A12 and amino acid sequence alignment of S100A12 orthologues. (A) Structure of the Ca(II)- and Cu(II)-bound S100A12 homodimer. $Cu(II)$ ions are shown as teal spheres, and $Ca(II)$ ions as yellow spheres (PDB: 1ODB).²⁴ (B) View of the His₃Asp motif of S100A12 taken from the structure of a $Zn(II)$ -bound form (PDB 2WCB).²⁵ The Zn(II) ion is shown as a brown sphere. (C) Structure of the Zn(II)bound S100A12 homodimer. Zn(II) ions are shown as brown spheres (PDB: $2WCB$).²⁵ (D) Sequence alignment of human, porcine, and bovine S100A12. The secondary structural elements presented above the alignment are for the human form. The transition-metal binding residues are presented in orange.

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Figure 2.

S100A12 depletes Zn(II) from microbial growth media. (A) Metal analysis of YPD/Tris medium treated with 0–250 μg/mL S100A12 or 250 μg/mL S100A12 His3Asp. (B) Metal analysis of TSB/Tris medium treated with $0-250 \mu\text{g/mL}$ S100A12 or 250 $\mu\text{g/mL}$ S100A12

 $His₃Asp.$ The experiments were conducted in the absence (light gray bars) and presence (dark gray bars) of a \approx 2-mM Ca(II) supplement (mean \pm SEM, n α).

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Figure 3.

Human S100A12 exhibits Ca(II)-dependent antifungal activity in mixed YPD/Tris medium. (A) *C. albicans* SC 5314, (B) *C. glabrata* ATCC 200918, (C) *krusei* ATCC 200917, and (D) *C. tropicalis* ATCC MYA-3404. Black, *Candida* spp. treated with S100A12 in the absence of a Ca(II) supplement. Red, *Candida* spp. treated with S100A12 in the presence of a \approx 2mM Ca(II) supplement. The OD₆₀₀ values were recorded at $t = 30$ h (mean \pm SEM, $n = 3$).

Figure 4.

The His3Asp motifs of S100A12 are essential for antifungal activity against *C. albicans*. (A) Antifungal activity of 1000 μ g/mL S100A12 and S10012 His₃Asp (t = 20 h, mean \pm SEM, $n = 3$). (B) Antifungal activity of 125 µg/mL S100A12 and Zn(II)-S100A12 (t = 30 h, mean \pm SEM, $n = 3$). The Zn(II)-S100A12 sample was prepared by pre-incubating S100A12 with 2 equiv. of $Zn(II)$ (11.9 µM) prior to the assay. The bar labeled " $Zn(II)$ " indicates the growth of a culture where 11.9 μ M Zn(II) was added to the YPD/Tris medium. For each assay, the YPD/Tris medium contained a \approx 2-mM Ca(II) supplement.

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Figure 5.

Antibacterial activity of human S100A12 (1000 µg/mL) against *S. aureus* ATCC 25923 (*Sa*), *E. coli* K-12 *(Ec)*, *P. aeruginosa* PAO1 (*Pa*), *L. monocytogenes* ATCC 19115 (*Lm*), and *L. plantarum* WSCF1 (*Lp*) in the absence and presence of a \approx 2-mM Ca(II) supplement. The assays were performed in mixed TSB/Tris medium ($t = 20$ h, $T = 30$ °C).

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Figure 6.

S100A12 exhibits antibacterial activity against *E. coli* with defective Zn(II) uptake machinery in TSB/Tris medium. (A) Antibacterial activity of S100A12 against *Escherichia coli* K-12. (B) Antibacterial activity of S100A12 against *E. coli* K-12 Δ*znuA*. Black, *E. coli* treated with S100A12 in the absence of a Ca(II) supplement. Red, *E. coli* treated with S100A12 in the presence of \approx 2 mM Ca(II). The OD₆₀₀ values were recorded at t = 20 h $(\text{mean} \pm \text{SEM}, n = 3).$

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Figure 7.

S100A12 metal-binding titrations. (A) Optical absorption spectra of S100A12 (200 µM) titrated with 0–5 equiv. of Co(II) at pH 7.0 (75 mM HEPES, 100 mM NaCl) and 25 °C. (B) Plot of absorbance at 550 nm versus equivalents of Co(II) added for titration of S100A12 or His₃Asp (200 μ M). (C) Response of Zincon (25 μ M) to Zn(II) in the presence of S100A12 or His₃Asp (10 μ M) at pH 7.0 (75 mM HEPES, 100 mM NaCl) and 25 °C (mean \pm SEM, *n*=3). The Zn(II)-Zincon complex absorbs at 621 nm. (D) Response of MF2 (10 μ M) to $Zn(II)$ in the presence of S100A12 or His₃Asp (10 μ M) at pH 7.0 (75 mM HEPES, 100 mM NaCl) and 25 °C (mean ±SEM, *n*=3). The Zn(II)-MF2 complex absorbs at 325 nm. Plots for Co(II) titrations and Zincon competitions performed in the presence of Ca(II) are presented in Figures S7 and S9.