

Bacterial siderophores: diversity, uptake pathways and applications

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[H1] Introduction

Iron is a vital nutrient for almost all living organisms, spanning from microorganisms to mammals^{1,2}. Its significance in biology stems from its ability to form bonds with oxygen, the redox reactivity of the Fe²⁺/Fe³⁺ couple, its catalytic role and its capacity to form metal complexes. As a result, iron has a crucial role in oxygen transport and functions as an ideal cofactor for enzymes involved in electron transfer reactions, including those in cellular respiration, DNA synthesis and repair, and antioxidant 8 defense mechanisms^{1,2}. For optimal growth, bacteria require iron concentrations in the range of 10⁻⁵ to 10^{-7} M Ref.³. However, these conditions are seldom met in aerobic environments because iron is poorly bioavailable. In the presence of oxygen and at neutral pH, iron primarily exists as ferric iron (Fe(III), which is highly insoluble and precipitates as various insoluble iron oxide compounds (solubility 12 equilibrium constant: K_s (Fe(OH)₃) = 2,8 × 10⁻³⁹ for Fe³⁺ versus K_s (Fe(OH)₂) = 4,9 × 10⁻¹⁷ for Fe²⁺). In the context of host infections, iron is also scarcely bioavailable, as the host actively limits the availability 14 of this metal to pathogens through a strategy known as nutritional immunity $4-10$. This strategy involves proteins such as ferritin (an iron storage protein) that sequester iron. Additionally, of the host produces hormones and proteins that regulate iron availability ; for example, the hormone hepcidin is produced in response to inflammation and reduces iron absorption in the intestine, thereby limiting iron availability for bacteria. Finally, iron-binding proteins such as lactoferrin, transferrin and calprotectin are produced at the infection site, rendering it unavailable. This array of strategies leads to a reduced concentration of free ferric iron in human body fluids and the infection sites.

21 A prevalent strategy bacteria use to access iron relies on siderophores, which are renowned for their 22 exceptionally high affinity for ferric iron^{11,12} (Table 1). Siderophores are secondary metabolites that are synthesized by microorganisms through biosynthetic pathways involving several enzymes, 24 including non-ribosomal peptide synthetases (NRPS) and polyketide synthases (PKS)^{13,14}. Generally, they exhibit molecular weights ranging from 200 to 2000 Da. Once synthesized, siderophores are excreted into the bacterial environment to scavenge iron and transport it back into the bacteria 27 through specialized transporters¹⁵. Afterwards, iron release from siderophores in the bacterial cells 28 occurs via various molecular mechanisms¹⁶. Often, microorganisms can produce 2-3 different siderophores, and they possess the ability to exploit siderophores that have been produced by other 30 microorganisms (xenosiderophores)¹⁷.

 In addition to chelating iron, siderophores can also chelate numerous other metals, including biological metals like Cu, Co, Ni, toxic metals like Al, Ga, Pb, or radionuclides like U, albeit with lower affinity 33 compared to iron^{18,19}. Due to their metal-chelating properties, siderophores have increasingly found

 applications in recent years as biofertilizers, biopesticides, and agents for controlling pathogens in 2 agriculture and aquaculture^{20–23}. In bioremediation, they chelate toxic metals, aiding environmental 3 cleanup²⁴. Siderophores also show promise in iron chelation therapy, antibiotic delivery and medical 4 imaging^{25,26}. In cancer research, they may hinder cancer cell growth^{27,28}.

 In this Review, I highlight and discuss recent advances in our understanding of the diverse molecular mechanisms involved in siderophore–mediated iron acquisition in bacteria. We now recognize that these mechanisms can be exceptionally varied and more complex than those previously described for *Escherichia coli*. This diversity of molecular mechanisms is exemplified by the pathogen *Pseudomonas* 9 aeruginosa, which exhibits numerous iron import pathways that are mediated by siderophores¹². Furthermore, as the range of applications for siderophores continues to expand across various fields, including agriculture and medicine, the technical potential of these bacterial secondary metabolites are also discussed.

[H1] Iron acquisition using siderophores

 In general, siderophore-mediated iron acquisition by bacteria involves a series of steps: sensing iron starvation; synthesizing and secreting siderophores; chelating iron; importing the siderophore–Fe complexes via specific transport systems; and finally releasing iron from these siderophores, inside the bacterial cells.

 The transcriptional repressor Fur (ferric uptake regulator) is the major actor to sense iron starvation. 20 As long as iron levels are sufficient, Fur binds ferrous iron (Fe $^{2+}$) as a cofactor, undergoing a conformational change that enables it to bind to specific DNA sequences called Fur boxes in the 22 promoter regions of target genes encoding proteins like those involved in siderophore biosynthesis 23 and iron acquisition by siderophores²⁹⁻³¹. Under conditions of iron limitation, intracellular iron levels decrease, which leads to the dissociation of iron from Fur. Consequently, Fur undergoes a structural change that reduces its DNA-binding affinity, which results in the de-repression of the target genes 26 transcription³². In addition to Fur regulation, other layers of control exist at the post-transcriptional 27 and translational regulation levels mediated by riboswitches and small RNAs^{33,34}. All these mechanisms ensure that the bacterium can efficiently acquire iron while responding to changes in iron availability. 29 For more details, see the following reviews and articles $31,35-42$.

[H2] The biosynthesis of siderophores and its cellular organization.

 Out of the 500 different siderophores identified, only a handful have had their biosynthetic pathways thoroughly investigated and elucidated. Enterobactin, which is produced by *E. coli*, was the first 3 siderophore for which the biosynthesis pathways has been reported⁴³. Since then, the biosynthesis of 4 pyoverdine I and pyochelin, which are produced by P. *aeruginosa*¹⁴, mycobactins⁴⁴, which is produced 5 by *Mycobacterium tuberculosis*, and yersiniabactin^{13,45,46}, which is produced by *Yersinia pestis*, have also been described. The biosynthesis of siderophores invariably entails a multi-step process that is coordinated by specific enzymes. It commences with the synthesis of precursor molecules, typically amino acids, aromatic rings and small carboxylic acids, which are subsequently assembled to form the 9 final siderophore structure by NRPS or $PKS^{44,47-49}$. Further refinement of the nascent siderophore structure may occurs through acylation and cyclization. The mature siderophore molecule is then released into the extracellular environment. NRPS assemble peptides by linking amino acids together, whereas PKS construct molecules by joining together acetyl, malonyl or methylmalonyl units, typically forming polyketides. Both NRPS and PKS are composed of several modules, with each module being responsible for adding a specific building block, enabling the synthesis of diverse and structurally 15 complex molecules (see the following reviews for more details on NRPS sand PKS⁵⁰⁻⁵²).

 The first-described biosynthesis of a siderophore was that of enterobactin, a tris-catecholate siderophore that binds ferric iron through three 2,3-dihydroxybenzoate (DHB) groups. DHB is synthesized from chorismate by three enzymes: EntA, EntB and EntC (with EntC functioning as an isochorismate synthase, EntB as an isochorismatase, and EntA as a 2,3-dihydro-2,3-dihydroxybenzoate 20 dehydrogenase)^{46,53,54} (Figure 1). The subsequent stage involves the assembly of three molecules of DHB and three L-serines to form a molecule of enterobactin. This assembly is facilitated by the NRPS 22 EntF and three additional enzymes: EntE, EntB and EntD⁵⁵. DHB is activated by EntE and transferred to EntB. The NRPS EntF catalyzes the formation of each ester and amide bond during the assembly of the 24 three DHB units. EntD functions as a 4'-phosphopantetheinyl transferase, necessary for activating one of the sub-domains of EntF and EntB.

 Enterobactin biosynthesis involves one NRPS (EntF), but most siderophore biosynthesis pathways involve multiple NRPS or PKS enzymes, as well as complex maturation processes of the synthesized molecule. This complexity is evident in the biosynthesis pathways of pyoverdine I. Pyoverdines constitute a family of chromo-peptide siderophores produced by fluorescent *Pseudomonas* strains, with pyoverdine I, II, and III representing distinct variants within this family, each distinguished by its unique structure56,57 . Pyoverdine I functions as the primary siderophore produced by *P. aeruginosa* PAO1, whereas pyoverdine II and III are alternative variants produced by other strains of *P. aeruginosa*. The core structure of pyoverdine I comprises an eleven-amino-acid peptide that is assembled by four 34 NRPSs (PvdL, PvdI, PvdJ, and PvdD)^{14,58} (Figure 2). Within this peptide backbone, two unusual amino

 acids, L-2,4-diaminobutyrate (L-Dab) and L-N5-formyl-N5-hydroxyornithine (L-fOHOrn), are 2 incorporated, with L-Dab being synthesized by the aminotransferase PvdH⁵⁹ and L-fOHOrn being 3 generated by the L-ornithine-N5-oxygenase PvdA and the hydroxyornithine transformylase PvdF^{60–62}. Additionally, during the initial assembly by NRPSs, a myristic or myristoleic acid is likely to be linked to the first amino acid of the peptide to ensure the association of the pyoverdine precursor with the 6 bacterial inner membrane during synthesis^{14,58}. After its assembly by the NRPSs, the pyoverdine 7 precursor is translocated from the cytoplasm into the periplasm by the inner membrane ATP-binding-8 cassette (ABC) transporter PvdE⁶³. In the periplasm, the myristic or myristoleic acid moiety is removed 9 by the acylase PvdQ $64,65$. Moreover, a portion of the peptide undergoes an auto-cyclization (involving L-Dab and D-Tyr residues) to form the characteristic chromophore of pyoverdines. This cyclization 11 process requires the copper-dependent tyrosinase PvdP^{66,67} and the oxidoreductase PvdO^{68,69}. Finally, the release of siderophores into the bacterial environment involves efflux systems, with PvdRT–OpmQ 13 being responsible for the secretion of pyoverdine I (Refs. $70,71$).

 Proteomic analyses of isolated inner membranes of *P. aeruginosa* have provided evidence that a portion of each of the four NRPSs involved in pyoverdine I biosynthesis (PvdL, PvdI, PvdJ, and PvdD) is 17 associated with the inner membranes and forms protein complexes⁷². This enzymatic complex was initially termed a 'siderosome' in the literature. However, since the term siderosome also refers to a membrane-bound organelle found in spleen macrophages, containing aggregates of hemosiderin, we will avoid using the term siderosome in this review to prevent any confusion. Instead, we will use the term multi-enzymatic complex.

22 More recently, multicolor single-molecule localization microscopy approaches showed colocalization 23 between PvdL and the other three NRPS⁷³, which further suggests the existence of a multi-enzymatic complex and that PvdL may have a central role in the assembly of this complex at the level of the inner leaflet of the cytoplasmic membrane. Concerning the other cytoplasmic enzymes involved in pyoverdine I biosynthesis, pull-down assays using PvdA, the enzyme responsible for producing LfOHOrn in *P. aeruginosa*, have revealed its interaction with NRPS PvdJ and PvdL⁷² . Co-purification assays have also demonstrated that PvdA binds to PvdF, with both enzymes catalyzing fOHOrn 29 synthesis⁷⁴. Furthermore, in vivo single-molecule tracking and Förster resonance energy transfer, as measured by fluorescence lifetime imaging, demonstrated that PvdA predominantly diffuses while bound to large complexes in the cytoplasm of *P. aeruginosa*, with a small, exchangeable trapped 32 fraction^{75,76}. PvdA was also found to interact with different stoichiometries with the four NRPS (PvdL, PvdI, PvdJ, and PvdD): several PvdA molecules interacted with PvdI, whereas PvdA formed one-to-one complexes with PvdJ⁷⁶. All of these data provide further confirmation of the existence of multi-

 enzymatic complexes associated with the inner membrane and composed of all four NRPSs, PvdA and PvdF (concerning PvdH no data are currently available) (Figure 2C).

 Currently, it remains unknown how the association of these multi-enzymatic complexes involved in pyoverdine I biosynthesis occurs with the inner membrane. PvdA possesses a hydrophobic, inner- membrane-anchoring domain at its N-terminus, facilitating its association with the inner 6 membrane^{72,77}. PvdA could help PvdL association with the inner membrane. Furthermore, the presence of a myristic acid chain attached to the first amino acid of the pyoverdine I backbone has also led to the suggestion that pyoverdine I biosynthesis takes place at the inner leaflet of the cytoplasmic 9 membrane⁶⁵. The association of PvdL and possibly all multi-enzymatic complexes involved in pyoverdine I biosynthesis to the cytoplasmic membrane may occur via PvdA, the myristic acid chain of the pyoverdine I precursor, or through some other unknown mechanism.

 This discovery of multi-enzymatic complexes for the biosynthesis of pyoverdine I offers a completely 13 novel perspective on the organization of siderophore biosynthesis in bacterial cells^{72,75,76,78,79}. Such organizations probably also exist for the synthesis of other siderophores in other bacterial species. A similar spatial organization may also exist for the NRPS involved in the biosynthesis of pyochelin, with 16 the NRPS PchE co-localizing with the enzyme PchA. Moreover, the fatty acid coenzyme-A ligase FadD1 has been demonstrated to be necessary for optimal pyochelin production in *P. aeruginosa* 18 cells⁸⁰, which suggests that in the case of pyochelin biosynthesis a siderophore precursor also exists with a fatty acid chain. For mycobactin in *Mycobacterium tuberculosis*, the fatty acid ligase FadD33 20 (also known as MbtM) has been reported to transfer a lipophilic aliphatic chain to the ε -amino group 21 of the lysine mycobatin core, thus having a role in the biosynthesis of this siderophore⁸¹. Moreover, EntD, one of the enzymes in *E. coli* involved in enterobactin biosynthesis, has been identified as a membrane protein: membrane fractionation techniques and protease accessibility studies have indicated that certain domains of EntD are associated with the inner leaflet of the cytoplasmic 25 membrane⁸². These various observations have led to the hypothesis that siderophore biosynthesis may occur in proximity to the cytoplasmic membranes.

 The spatial co-localization of enzymes within a multi-enzymatic complex located by the inner 28 membrane represents a significant advantage. This organization optimizes the efficient transfer of siderophore precursors among different enzymes and prevents the dispersion of these precursors throughout the cytoplasm, where they could potentially chelate intracellular metals. Such organization may also be important for the secretion of the synthesized siderophores out of the bacterial cells. Many questions remain regarding the organization of enzymes in such a multi-enzymatic complex. How do NRPSs interact with each other within the complex? What impact does this organizational

1 structure have on the enzymatic activity of each individual enzyme? Are the enzymes within the 2 complex continually active in the production of siderophores?

3

4 **[H2] Iron chelation by siderophores and high diversity in their chemical** 5 **structures.**

6 Siderophores possess a remarkable ability to scavenge and capture even trace amounts of iron from 7 their surroundings due to their high affinity for iron ions, especially ferric iron (Fe $3+$) (Table 1). This 8 heightened affinity can be attributed to their chemical structure and the presence of bidentate 9 constituents such as catecholates, hydroxamates, α -hydroxy-carboxylates and phenolate groups^{12,83} 10 (Figure 3). Siderophores often consist of three bidentate groups, which in an octahedral geometry 11 around the coordinate with Fe^{3+} ions^{12,83,84}. However, it is not uncommon to find siderophores with 12 fewer than three bidentate groups, in which case, the coordination stoichiometry may be 2:1 or 3:1 13 for siderophore: $Fe³⁺$.

 Currently, approximately 500 siderophores with diverse chemical structures have been identified, which highlights the vast variability in their chemical compositions. As mentioned above, whereas some siderophores exclusively contain catecholate, hydroxamate or α-hydroxy-carboxylate chelating 17 groups, many others consist of mixtures of different ligands^{12,83,85}. This structural heterogeneity reflects the adaptive strategies used by microorganisms in diverse ecological niches to thrive under varying iron availability conditions. These intricate variations in their structures arise from the diverse enzymatic pathways involved in their biosynthesis across different microbial species.

21 Although siderophores are primarily known for their high affinity for iron, they also coordinate many 22 other metal ions, albeit with lower affinities compared to iron^{18,19} (see below). For example, the 23 formation constants for the hydroxamate siderophore desferrioxamine B with Ga³⁺, Al³⁺, and In³⁺ range 24 between 10²⁰ and 10²⁸ M⁻¹, whereas that with Fe³⁺ is 10³⁰ M⁻¹ (Refs. ^{18,86}). Similarly, formation constants 25 for pyoverdine with Zn²⁺, Cu²⁺, and Mn²⁺ fall between 10¹⁷ and 10²² M⁻¹, whereas that with Fe³⁺ is 10³² 26 M⁻¹ (Ref. ⁸⁷). Screening with sixteen different metals (Ag⁺, Al³⁺, Cd²⁺, Co²⁺, Cu²⁺, Cu²⁺, Ga³⁺, Hg²⁺, 27 Mn²⁺, Ni²⁺, Pb²⁺, Sn²⁺, Tb³⁺, Tl⁺ and Zn²⁺) revealed that pyoverdine and pyochelin, the two siderophores 28 produced by *P. aeruginosa*, are capable of chelating all these metals^{88,89}. Another example is with 29 vanadium, which can be chelated by the siderophores pyochelin, putrebactin, pyoverdine, 30 enterobactin, and desferrioxamine B^{90-93} .

[H2] Uptake and delivery of iron by siderophores.

 In Gram-negative bacteria, all reported cases have demonstrated that, regardless of the type of siderophore or bacterial species, iron uptake through siderophores begins with the uptake of 4 siderophore–Fe complexes across the outer membrane by TonB-dependent transporters (TBDTs) ¹⁵. These transporters are named after the inner membrane protein TonB, which forms a complex with two other inner membrane proteins, ExbB and ExbD. The TonB–ExbB–ExbD complex functions as a molecular 'motor', that transmits energy from the inner membrane proton motive force to the TBDT in the outer membrane, facilitating the translocation of the siderophore-iron complex into the 9 periplasm^{94–96}. These TBDTs demonstrate remarkable specificity in binding to siderophores¹⁵. Consequently, when a bacterial species can use different siderophores to access iron, it will express 11 distinct TBDTs, each tailored to a specific siderophore or closely related ones^{15,17,97,98} (Figure 4a). For instance, *P. aeruginosa* can express at least 14 different TBDTs involved in siderophore-mediated iron 13 acquisition via siderophores or xenosiderophores¹⁷ (Supplementary Table 1). Furthermore, *E. coli* strains use up to nine TBDTs, for iron uptake by siderophores or xenosiderophores, and *Salmonella* 15 species use up to six TBDTs, all dedicated to the utilization of siderophores or xenosiderophores⁹⁷⁻⁹⁹ (Supplementary Table 1). It is important to note that even among strains of a species, there can be strain-to-strain variability in the number of siderophores produced, xenosiderophores used, and genes encoding TBDTs present in the genomes. Moreover, the presence of various iron uptake pathways in bacterial genomes underscores the significance of iron for bacterial survival and indicates the existence 20 of intense competition among microorganisms for this essential resource.

 In Gram-positive bacteria, no outer membrane is present, and the bacterial cells feature specific receptors anchored in the cytoplasmic membrane that recognize and bind siderophore–iron complexes. Upon binding, these proteins transfer the siderophore–iron complex to an ABC transporter 25 embedded in the cytoplasmic membrane¹⁰⁰. Like TBDTs in Gram-negative bacteria, these ABC transporters, along with their corresponding membrane-bound receptors, exhibit high siderophore specificity. For example, *Staphylococcus aureus* can express multiple systems at its cell surface, each 28 dedicated to importing iron via a specific type of siderophore or xenosiderophore^{100,101} (Figure 4c).

 In both, Gram-negative and Gram positive bacteria, the subsequent critical step in iron acquisition involves releasing iron from the siderophore–iron complex. In all the mechanisms reported to date, iron reduction occurs. In Gram-negative bacteria, this step can occur either in the periplasm or 32 cytoplasm, depending on the siderophore pathway or bacterial species¹⁶ (Figure 4B). In cases of cytoplasmic dissociation, siderophore–Fe complexes must traverse the inner membrane, where ABC

 transporters primarily participate in this step, though some protonmotive-dependent transporters 2 have also been documented¹⁶. Dissociating ferric iron from high-affinity siderophores necessitates specific mechanisms, which can vary among siderophores and bacterial species. These strategies invariably entail iron reduction, often accompanied by enzymatic modification or cleavage of the 5 siderophore¹⁶. Additionally, changes in pH can influence the stability of the siderophore–iron complex, with lower intracellular pH weakening the siderophore–iron binding. A vivid illustration of this molecular diversity can be found in the case of enterobactin (Figure 5). In *E. coli*, enterobactin undergoes enzymatic hydrolysis by the esterase Fes in the bacterial cytoplasm, which leads to the 9 release of the iron for intracellular utilization^{85,102} (Figure 5a). By contrast, in *P. aeruginosa*, which uses enterobactin without producing it, the esterase PfeE hydrolyzes enterobactin in the bacterial 11 periplasm¹⁰³ (Figure 5a). In both cases, iron reduction is also necessary to achieve efficient iron release 12 from enterobactin^{103,104,104}. Another example highlighting this diversity is the iron uptake by pyoverdine I (Figure 5b), the primary siderophore used by *P. aeruginosa*. Here, iron is released from the chelator in the bacterial periplasm through a mechanism involving iron reduction, with no chemical 15 modification of the siderophore^{105–107}. Once iron is liberated from the siderophore in the periplasm, it must traverse the inner membrane to access the cytoplasm for utilization. Gram-negative bacteria use 17 specialized ABC transporters to facilitate this crucial step¹⁰⁸.

 Whether in Gram-negative or Gram-positive bacteria, various reductases involved in the process of 19 dissociating iron from siderophores have been described (for a recent review, see Ref. 109). Some of these reductases are soluble, whereas others are membrane-anchored. Among them are siderophore- interacting proteins (SIPs), which are flavoproteins. One notable example is the YqjH protein from *E.* 22 *coli*, which reduces iron in enterobactin–iron and dicitrate–iron complexes⁸⁵. Additionally, a few reductases have been identified that utilize a [2Fe-2S] cluster, known as the ferric siderophore 24 reductase (FSR) family¹¹⁰. These reductases are found in Gram-positive bacteria like in *Bacillus halodurans* and in Gram-negative bacteria, exemplified by the FhuF protein, implicated in iron release 26 mechanisms from hydroxamate-type siderophores in *E. coli*^{110–112}. Furthermore, a recent discovery 27 introduces a new class of siderophore–Fe membrane reductases, including FpvG and FoxB, involved in 28 iron acquisition by pyoverdine and desferrioxamine G in *P. aeruginosa*^{107,113}. The X-ray structure of FpvG reveals a fold with two tightly bound heme molecules that likely have a role in the iron reduction 30 . process¹¹³. These mechanisms of iron dissociation from siderophores remain poorly understood and have received much less scientific attention compared to other aspects of iron import by siderophores.

 In conclusion, siderophores exhibit remarkable diversity in their chemical structures (Figure 2), reflecting probably their adaptation to diverse ecological niches and iron-scavenging strategies. One can imagine that this structural diversity has arisen through evolutionary processes, with bacteria

 developing siderophores featuring specific chemical attributes optimized for efficient iron scavenging in varying environments, contributing to their ecological success and survival. The evolution of siderophore chemical structures has likely occurred alongside a diversification of mechanisms for dissociating siderophore-Fe complexes in bacteria, often displaying distinct and complex mechanisms. These mechanisms of iron release from siderophores can also vary between species for the same siderophore as shown with the example of iron acquisition by enterobactin in *E. coli* and *P. aeruginosa*.

[H2] Uptake and delivery of other metals than iron by siderophores.

9 Siderophores can chelate many metals besides iron $89,114$, but whether these metals are imported into bacteria by siderophores remains a frequently asked question. Since an iron reduction step is most often essential for releasing iron from the siderophore and delivering it to bacterial cells, one could assume that no metal other than iron would be delivered to bacteria by siderophores. This iron reduction step, necessary for releasing iron from siderophores, may be a critical control point for the metal selectivity of these iron uptake pathways, helping to avoid the release of metals other than iron from siderophores. In *P. aeruginosa*, it has been shown that pyoverdine–metal complexes other than pyoverdine–Fe are re-excreted by efflux systems because probably the metal cannot be removed from 17 the siderophore.

 However, data also show the opposite — that siderophores can import other biologically relevant metals besides iron. It was shown that yersiniabactin can bind several divalent transition metals, including Ni and Cu, and that *E. coli* can use yersiniabactin to acquire Cu and Ni during metal limitation¹¹⁶ . Moreover, it has been shown that the probiotic bacterium *E. coli* Nissle 1917 (or 'Nissle') 22 utilizes yersiniabactin to import Zn, enabling Nissle to grow in zinc-limited media¹¹⁷. Additionally, yersiniabactin in *Yersinia pestis* has been shown to overcome zinc limitation during infection of both 24 mammalian and insect hosts¹¹⁸. All these data suggest that yersiniabactin behaves more like a metallophore than a siderophore. However, this behavior may be specific to yersiniabactin or siderophores with chemical structures similar to yersiniabactin. Pyochelin, like yersiniabactin, contains 27 a phenolate–thiazoline core in its structure (Figure 3) and is capable of transporting metals other than 28 iron into *P. aeruginosa* cells⁸⁸. However, it has not been confirmed whether these uptakes have 29 biological significance. Currently, the mechanism of iron or metal release from yersiniabactin or pyochelin is unknown, and it may involve siderophore degradation rather than iron reduction. It would be interesting to investigate whether siderophores with chemical structures similar to yersiniabactin can also facilitate the uptake of other biologically relevant metals into bacteria, potentially having a crucial role in bacterial physiology and virulence.

[H1] Siderophores in competition and infection

 Within bacterial communities, a fierce competition unfolds among microorganisms for resources, and siderophores have a pivotal role in this competition for iron. Studies have demonstrated that reduced iron availability leads to the proliferation of siderophore-producing bacterial families and decreases in the abundance of non-siderophore iron-dependent taxa within the intestinal microbiota, underscoring 6 the significance of siderophores^{119,120}. Furthermore, microbial species within a microbiome that produce siderophores generate a variety of these molecules with distinct chemical structures, which results in varying affinities for iron. Additionally, bacteria often use their own siderophores to access iron and can also exploit siderophores produced by other bacteria (xenosiderophores) by expressing the necessary transporters (TBDT in Gram-negative bacteria or ABC transporters in Gram-positive bacteria, as depicted in Figure 4). As a result, the diverse microorganisms within a microbiome exhibit differential capabilities to acquire iron, which substantially affects their growth and survival within the 13 microbiota¹²¹. Some bacteria may outcompete others by producing or utilizing siderophores with higher iron-binding affinities, thereby gaining a competitive edge in iron-limited environments. For example, bacteria with the ability to use a wide range of siderophores, such as *P. aeruginosa* (Supplementary table 1), are more competitive than *Salmonella enterica* subsp*. enterica* serovar Typhimurium within bacterial communities. This competition for iron mediated by siderophores can 18 influence the composition and dynamics of the entire microbial community¹²¹. Moreover, siderophore 19 production by beneficial bacteria may contribute to pathogen exclusion in the microbiota¹²¹.

20 Although the full extent of the importance of siderophores in microbial communities and iron 21 competition remains incompletely understood, key factors in this competition undoubtedly include 22 the affinity of siderophores for iron and the ensuing competition for the chelation of this metal; and 23 the ability of the bacteria to use an extensive array of siderophores, particularly those with high iron affinity.

 During infection, a battle for iron occurs between the host and pathogens. As mentioned in the introduction, the host uses a defense strategy known as nutritional immunity, whereby essential 27 nutrients such as iron are restricted from pathogens $4-9$. To overcome this iron restriction, bacteria 28 produce and release siderophores, which scavenge iron from the host, including from proteins like 29 transferrin and ferritin^{122,123}. By acquiring iron, bacteria can enhance their growth and survival within 30 the host, thereby promoting the establishment and progression of infection¹²⁴. Additionally, pathogens can exploit xenosiderophores produced by other microorganisms, including commensal or competing ones, further complicating the interplay between pathogens, the host immune system and the 33 microbiota^{10,125}. Due to their crucial role in infection, siderophores are often regarded as critical virulence factors that contribute to the pathogenesis of bacterial infections by facilitating iron

 acquisition and promoting bacterial survival and proliferation within the host environment. 2 Understanding the role of siderophores in infection can offer insights into novel therapeutic strategies targeting bacterial iron acquisition pathways.

[H1] Versatile tools for innovative applications

 Due to their high affinity for iron and their ability to chelate metals other than iron with strong affinities, siderophores have emerged as innovative, potent, and versatile tools in various fields, including agriculture, bioremediation, metal recovery and medical applications (Table 2). Below, I have briefly outlined these various applications to provide an overview of the potential uses for these molecules.

[H2] Application in agriculture.

 Siderophores hold promise for enhancing agricultural practices, particularly as biofertilizers and biopesticides, offering benefits such as improved plant growth, nutrient uptake and overall crop 15 $health¹²⁶$.

 As previously discussed for bacteria, iron is also an essential nutrient for plant growth. However, iron often exists in insoluble forms in soils. Even though plants produce their own siderophores to access iron (phytosiderophores), the additional presence of siderophore-producing bacteria can aid in solubilizing iron by solubilizing it from various insoluble iron oxide compounds (Supplementary Figure 20 1a), thus making it more accessible for plant absorption, thus preventing iron deficiency chlorosis (for 21 a review see Ref. 127). Additionally, as siderophores can also form complexes with other biological metal 22 ions, such as Zn, Cu and Mn, the presence of siderophore-producing bacteria in the soil may enhance 23 the solubility and availability of these nutrients to plants¹²⁷. This leads to improved nutrient uptake and enhanced plant growth.

 Furthermore, siderophores can have a role in protecting plants against certain pathogens by disrupting their iron acquisition processes (Supplementary Figure 1b). This contribution leads to improved plant health and reduced disease incidence. For instance, *Bacillus* spp. inhibits the growth of *Fusarium oxysporum* by producing siderophores that scavenge iron from the environment^{128,129}. These findings 29 underscore how the use of siderophores or siderophore-producing microorganisms can contribute to more sustainable and efficient agricultural practices, ultimately resulting in increased crop productivity and improved soil health (reducing the need for synthetic chemical fertilizers).

 Siderophores have also demonstrated potential in the biocontrol of fish pathogens in aquaculture 2 settings^{130–133}. Aquatic farming, including fish, shellfish and aquatic plants, often faces challenges related to disease outbreaks caused by various pathogens. Beneficial microorganisms that produce siderophores can outcompete fish pathogens for available iron in the environment. This competition reduces the growth and survival of the pathogens, aiding in the control of disease outbreaks. The use of siderophores as biocontrol agents can help reduce the reliance on chemical treatments for managing fish diseases, thereby minimizing the risk of antibiotic resistance.

8 It is essential to note that although the potential benefits of using siderophores in agriculture and aquaculture are promising, practical implementation necessitates careful consideration of factors such as specific pathogens, the microbial community within agricultural or aquaculture environments, and potential effects on non-target organisms.

[H2] Application in bioremediation.

 Siderophores have also displayed potential for applications in bioremediation, a process that uses living organisms or their products to detoxify or remove pollutants from contaminated environments 16 (for a review see Ref.). As already mentioned above, siderophores possess the ability to chelate a wide range of metals, including toxic ones such as heavy metals (Ar, Cd, Pb, Hg, Cr, Ni, Cu, etc.), metalloids (Te or Po) and radionuclides (U), although their affinities for these metals are generally 19 Iower compared to their affinity for Fe³⁺ (Ref. 134). Consequently, siderophores can have a substantial 20 role in remediating environments that are contaminated with metal pollutants. When released into contaminated environments, siderophores can bind to these toxic metal ions, forming stable 22 complexes that increase the solubility and availability of metals for uptake by microorganisms through iron uptake pathways. For instance, a study identified siderophore production by *Pseudomonas brenneri* MF957286 as a tool for removing Cr(VI) from contaminated soils¹³⁵. Marine bacteria- produced siderophores have been found to chelate polonium (Po) and protactinium (Pa), whereas 26 pyoverdine produced by *P. fluorescence* has demonstrated its ability to chelate uranium (Ur)^{136,137}. Another research group increased siderophore-mediated bioaccumulation of Cd(II) by *Bacillus subtilis*¹³⁸ . Additionally, it was shown that *P. azotoformans* produces a mixed-type catecholate and hydroxamate siderophore when grown on arsenic-contaminated soil, which was more efficient than 30 citric acid or EDTA in removing arsenic from the soil¹³⁹. However, it is important to note that complexation with siderophores does not guarantee the entry and accumulation of toxic metals or metalloïds within bacterial cells. When bioaccumulation of siderophore–metal complexes occurs, it can often be toxic to bacteria, affecting their growth. In response, bacteria may develop various

1 resistance mechanisms, such as increasing the expression of efflux systems¹⁴⁰. To overcome such 2 challenges, physical methods like precipitation or biosorption can be used to remove siderophore-3 metal complexes from contaminated soils, liquids or waste¹⁴¹.

 Siderophores can also be integrated with other bioremediation techniques, including microbial consortia and phytoremediation (Supplementary Figure 1c), to create synergistic effects and enhance 6 overall remediation outcomes¹⁴². In phytoremediation, a bioremediation strategy that use plants to extract pollutants from soil, siderophores can improve metal solubilization and subsequent metal uptake by plants (Supplementary Figure 1c). A study using siderophore-producing *P. aeruginosa*, *P. fluorescens* and *Ralstonia metallidurans* showed that rhizobial inoculation with these species in maize 10 plants increased the phytoremediation of chromium (Cr) and lead (Pb) 143 .

 Finally, studies have also indicated that siderophores can mitigate the toxicity of asbestos, silicate minerals present in soil that have been used in construction for many years. Asbestos is highly toxic due to its substantial iron content, which generates free radicals and reactive oxygen species (ROS) through the Fenton reaction, which can lead to DNA damage. Inhalation of asbestos fibers can result in severe health issues, including lung fibrogenesis, mesothelioma, pleural calcification and various cancers. Iron can be extracted from asbestos fibers using siderophores or siderophore-producing 17 bacteria (for a review see Ref.).

 In summary, siderophores hold great promise as tools for enhancing bioremediation efforts within a range of bioremediation strategies.

[H2] Recovering rare earth elements.

22 The ability of siderophores to efficiently chelate metals beyond iron¹³⁴ make them valuable candidates 23 in processes for the recovery of rare earth elements (for a review see Ref.). These elements encompass lanthanum (La), cerium (Ce), praseodymium (Pr), neodymium (Nd), promethium (Pm), samarium (Sm), europium (Eu), gadolinium (Gd), terbium (Tb), dysprosium (Dy), holmium (Ho), erbium (Er), thulium (Tm), ytterbium (Yb), lutetium (Lu), scandium (Sc) and yttrium (Y). Rare earth elements 27 have pivotal roles in various modern technologies, including electronics, renewable energy and 28 advanced materials. The recovery of these elements from diverse sources holds significant interest due to their strategic importance and limited global availability. Siderophores could be instrumental in developing methods for selectively recovering specific rare earth elements from complex mixtures. They are capable of forming stable complexes with rare earth elements, rendering them more soluble 32 in aqueous solutions^{145,146}. This complexation process facilitates the extraction and separation of rare

 earth elements from other metals in various solutions, presenting a more environmentally friendly alternative to traditional extraction methods that involve harsh chemicals.

[H2] Metal sensors.

 The high affinity of siderophores for metal ions has proven valuable in the development of metal 6 sensors or methods for metal detection (for a review see Ref. $26,147$). Siderophore-based metal sensors find application in monitoring metal contamination in environmental samples, including water, soil and sediment, offering insights into metal pollution levels in ecosystems. Furthermore, siderophores can be used to detect metal ions in biological samples. In the field of medical diagnostics, siderophores have been explored as tools for detecting abnormal metal ion concentrations in bodily fluids, which 11 could indicate certain health conditions $148,149$.

 In certain cases, the spectral properties of siderophores have been directly used as visual indicators of the presence of metal ions. Some siderophores undergo spectral changes, including variations in 14 fluorescence and/or absorbance, upon binding to specific metal ions^{89,150,151}. These alterations in fluorescence or absorbance can be detected and quantified, which enables the sensitive detection of metal ions. Siderophores can be modified and covalently linked to fluorophores; when such conjugates 17 bind to a metal ion, the fluorescence properties of the probe change^{148,149}. Siderophore-modified 18 electrodes, known as electrochemical sensors, can be used¹⁵². The binding of metal ions to the siderophore on the electrode surface can induce changes in electrical properties, which can be measured and correlated with the concentration of the metal ion. Siderophores can also be incorporated into nanoparticle structures to create sensitive metal sensors, referred to as 22 nanoparticle-based sensors^{153,154}. Furthermore, siderophores can be combined with genetically 23 engineered bacteria or other microorganisms to develop bioreporters¹⁵⁵. These bioreporters emit fluorescence or other signals when they encounter specific metal ions, providing a live, biological 25 indicator of metal presence. These sensors offer advantages such as specificity, sensitivity, and potential adaptability for on-site or real-time detection.

[H2] Imaging agent.

 Siderophores can be modified to carry imaging agents, such as radioactive isotopes or contrast agents used in various imaging modalities (for example, positron emission tomography, single-photon 31 emission computed tomography, magnetic resonance imaging (MRI))¹⁵⁶. For example, siderophores can enhance the contrast in MRI images by binding to certain metal ions, which affects the relaxation

 properties of nearby water molecules. This can improve the visibility of specific tissues or structures 2 during MRI scans^{157,158}. These modified siderophores can be designed to selectively accumulate in specific tissues or cells of interest (Supplementary Figure 1e). Such imaging agents can detect bacterial 4 infections in small animal imaging experiments¹⁵⁹. Modified siderophores can be engineered to target cancer cells that overexpress specific receptors. For example, pyoverdine was conjugated to synthesized superparamagnetic iron oxide nanoparticles. These superparamagnetic iron oxide 7 nanoparticles-pyoverdine conjugates were covalently bound to a MUC1 aptamer¹⁶⁰. An aptamer is a short, single-stranded oligonucleotide that can bind to a specific target molecule with high affinity and specificity. The conjugate could be used as a diagnostic agent, since a type of MUC1 is aberrantly overexpressed on the majority of cancer cell surfaces, such as in human colon cancer. The superparamagnetic iron oxide nanoparticles-pyoverdine-MUC1 complex was also loaded with doxorubicin, an anti-cancer agent, to create a targeted drug 13 delivery system¹⁶⁰. This compound was detected at the tumor site in mice using ex vivo fluorescence imaging. In vivo analysis revealed significant tumor inhibition and increased survival rates in the mice receiving this compound compared to untreated mice. Inflammation is often associated with increased iron metabolism, and siderophores labeled with imaging agents can 17 be used to detect and monitor inflammatory processes in various conditions¹⁶¹. The use of 18 siderophores in imaging is an area of ongoing research and innovation $162-165$.

[H2] Iron chelation therapy.

 Iron chelation therapy is a medical treatment aimed at reducing excess iron levels in the body, which 22 can occur due to conditions such as hereditary hemochromatosis, thalassemia and certain types of 23 anemias¹⁶⁶. In this approach, synthetic siderophores or siderophore analogs can be used to bind excess 24 iron in the body, and the resulting iron complexes are excreted through urine and feces^{28,167}. The remarkably affinity of siderophores for iron ions enables them to efficiently and selectively bind to excess iron in the bloodstream and tissues, making this affinity crucial for the success of iron chelation 27 therapy. Desferrioxamine (Desferal), deferasirox (Exjade), and deferiprone (Ferriprox) (Supplementary 28 Figure 1f) have been the most important US FDA-approved iron chelators over the past years^{28,167}. DFO is a natural siderophore, whereas the other two compounds are synthetic molecules.

[H2] Cancer treatment.

 Siderophores have also garnered attention in the context of cancer research. Cancer cells often require higher levels of iron to fuel their rapid growth and proliferation. In fact, a link exists between iron 3 overload and the progression of cancer (for a review see Ref. ¹⁶⁸). Consequently, cancer cells frequently manifest alterations in iron metabolism and heightened expression of iron-regulating proteins. Researchers have explored the idea of using siderophores or siderophore-like compounds to specifically address the heightened iron requirements of cancer cells. The fundamental concept behind 7 this approach is to hinder the growth of cancer cells by depriving them of the iron essential for their survival. For example, ferrichrome has demonstrated a robust capacity to suppress colon cancer cells 9 without impeding the growth of non-cancerous intestinal cells¹⁶⁹. Similarly, desferrioxamine has 10 undergone testing against metastatic hepatocellular carcinoma¹⁷⁰. However, such an approach is complex and necessitates thorough consideration of potential side effects and interactions with normal cells. To address this challenge, numerous artificial iron chelators, including pyridoxal isonicotinoyl hydrazone, thiosemicarbazones and O-tresox, have been developed for cancer therapy 14 (for a review see Ref. $28,171$). However, currently no compound has received approval for clinical use.

[H2] Pathogen sensor.

 Due to the strong, specific, and selective interaction between siderophore–Fe complexes and their 18 transporters, such as TBDTs in the case of Gram-negative bacteria¹⁵, siderophores have been considered in the development of pathogen sensors. These sensors utilize the interaction between siderophores and pathogens to create detection methods that are capable of identifying the presence 21 of harmful microorganisms (for reviews see Refs. 26,147). Siderophores are modified with a fluorescent 22 or colorimetric tag. When the siderophore–Fe complex binds to a specific pathogen (binding to the TBDT at the bacterial surface), fluorescence or color changes occur, indicating the presence of the pathogen (Supplementary Figure 1e). Another strategy involves using siderophores in conjunction with bioreporter systems, where the presence of the pathogen induces a specific response in a genetically engineered organism. This response can be easily detectable, providing a visible indicator of pathogen presence. For instance, researchers have developed a sensor system based on siderophores and gold- plated chips, combined with microscopy analysis, that is capable of detecting and distinguishing *Yersinia enterocolitica*, *S. aureus*, *Mycobacterium smegmatis* and *P. aeruginosa*¹⁷² . Others immobilized pyoverdine on planar aluminum chip substrates and used Raman microspectroscopy to isolate and 31 identify *P. aeruginosa* and *P. fluorescence* from tap water¹⁷³. These different sensors offer rapid detection, as the interaction between the siderophore and pathogen can occur relatively quickly in

 various environments, including food, water, clinical samples and clinical diagnostics, producing a visible or measurable signal.

[H2] Vectorization of antibiotics.

 One of the bottlenecks in the development of new antibiotics is the ability of these drugs to enter bacteria and especially to cross the outer membrane of Gram-negative bacteria. Iron-uptake systems that use siderophores can function as pathways for drug import into bacteria. Indeed, by covalently attaching antibiotics to siderophores, these compounds can exploit the bacterial iron uptake pathway 9 to deliver antibiotics directly into bacterial cells¹⁷⁴ (Supplementary Figure 1d). This strategy was not invented by scientists, but rather, bacteria have evolved such compounds to outcompete and kill other bacteria for resources. Natural siderophore-antibiotic conjugates, such as albomycins, ferrimycins, 12 danomycins or salymycins, are produced by microorganisms¹⁷⁵. As each bacterial species uses its own siderophores, and not necessarily those of other bacterial species, it is possible to design siderophore- conjugated antibiotics to target specific bacterial species or strains. Many different siderophore– 15 antibiotics have been designed and tested^{176–178}. The results have shown promising in vitro activity against various clinically relevant multidrug-resistant Gram-negative bacteria. However, these 17 compounds have not yet made it to the market, except for Cefiderocol¹⁷⁹⁻¹⁸¹. Cefiderocol was developed by Shionogi and approved by the USA Food and Drug Administration in 2019 and European Medicines Agency in 2020, respectively, and was released on the market of the United States of America and Europe. Cefiderocol contains a cephalosporin moiety linked to a catechol functional 21 group, which acts as an iron chelator (for a review see Ref.). The uptake of this compound occurs 22 through the iron-uptake pathways and in the periplasm, it inhibits the bacterial cell wall synthesis, ultimately leading to cell death. As with any antibiotic, the development of resistance is inevitable. Regarding the resistance mechanisms observed so far against cefiderocol, the most frequently reported mechanisms are those observed against β-lactams, as well as reduced expression or mutation 26 of genes encoding the TBDTs involved in cefiderocol-iron uptake (for a review, see Ref.). The example of cefiderocol demonstrates that iron import pathways can be promising import pathways for antibiotics when vectorized by siderophores. However, resistance phenomena cannot be avoided, and the development of this type of strategy requires a thorough understanding of the biology related to the expression and functioning of these iron import pathways in bacteria, particularly in pathogenic bacteria.

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[H1] Conclusion

 In conclusion, the diversity of siderophore chemical structures and the array of molecular mechanisms involved in iron acquisition by siderophores reflect the intricate interplay between microorganisms 4 and their environments, and underscores the remarkable adaptability of microorganisms to acquire this essential nutrient in various environments. These molecular mechanisms, which encompass siderophore biosynthesis, iron chelation, and iron import and delivery into bacterial cells, enable microorganisms to thrive in iron-limited ecosystems and contribute to the dynamics of microbial communities. Understanding the complexity of siderophore-mediated iron acquisition is crucial for elucidating microbial physiology, ecology and evolution, as well as for developing strategies to use siderophores and the siderophore-dependent microbial iron uptake pathways for environmental, biotechnological and biomedical applications. Indeed, siderophores, with their diverse chemical structures and exceptional metal-chelating properties, represent fascinating and versatile molecules offering opportunities for innovative applications across multiple disciplines, and they may unlock innovative solutions for diverse challenges.

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1 **Table 1: Stability constants of siderophores for iron.**

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1 **Table 2: Applications of siderophore properties in various fields, ranging from agriculture to** 2 **bioremediation and medical science.**

 Figure 1. Biosynthesis pathway of enterobactin in *Escherichia coli***.** The biosynthesis of enterobactin involves the synthesis of 2,3-dihydroxybenzoate (DHB) from chorismate by three enzymes: EntA, EntB 4 and EntC. Enterobactin is then formed by assembling three molecules of DHB and three L-serines, facilitated by the non-ribosomal peptide synthetases (NRPS) EntF together with EntE and EntD enzymes. EntF catalyzes the formation of ester and amide bonds, whereas EntD functions as a 4'- phosphopantetheinyl transferase that is essential for activating sub-domains of EntF and EntB. The stereochemistry of the chemical structures of the siderophore presented is not specified.

 Figure 2. Biosynthesis pathway of pyoverdine I in *Pseudomonas aeruginosa***. a)** The precursor of the siderophore pyoverdine I is an 11-amino acid peptide that is assembled in the cytoplasm of *Pseudomonas aeruginosa* cells by four non-ribosomal peptide synthetases (NRPS), namely PvdL, PvdI, PvdJ and PvdD. Additionally, three enzymes (PvdH, PvdA and PvdF) are involved in synthesizing atypical amino acids utilized by the NRPS. PvdL is the NRPS involved in the assembling of the tripeptide L-Glu– L-Tyr–D-Dab (in blue like PvdL). Next, L-Ser, L-Arg, L-Ser and L-OHOrn (all in green) are incorporated successively by PvdI, L-Lys and L-OHOrn (in pink) by PvdJ and the two L-Thr (in yellow) by PvdD. From the initial step of peptide assembly, a myristate or myristoleate chain is incorporated on the first residue (L-Glu) by PvdL. **b)** The cytoplasmic precursor is exported to the periplasm, where it undergoes maturation involving other enzymes (PvdQ, PvdP and PvdO), which leads to the removal of the fatty acid chain and cyclization of the chromophore. **c) The relative size of the enzymes is shown.** PvdL, PvdI, PvdD and PvdD are NRPS, with very different molecular weights (480, 570, 240 and 274 kDa for PvdL, PvdI, PvdJ and PvdD respectively). The molecular weights of three enzymes, PvdH, PvdA and PvdF, involved in synthesizing atypical amino acids are 50, 50 and 31 kDa. **d) Schematic view of the multi-enzymatic complex formed between the different enzymes involved in pyoverdine 1 biosynthesis in** *P. aeruginosa***.** The gene organization is illustrated, along with a schematic view of the multi-enzymatic complex formed by PvdL–PvdI–PvdJ–PvdD–PvdH–PvdF–PvdA (colours as in part c). 27 Parts c and d is adapted with permission from Gasser et al., 2020 (Ref.).

 Figure 3. Structures of siderophores. Shown are structures of different siderophores. Highlighted are in blue the catechol chelating groups; in green the hydroxamates; in pink the carboxylates; and in orange the phenolates. The stereochemistry of the chemical structures of the siderophores presented is not specified.

 Figure 4. Iron uptake in Gram negative and Gram positive bacteria. a) Siderophore-Fe transport through the outer membrane in Gram-negative bacteria. Iron uptake begins with the translocation of s siderophore-Fe complexes across the outer membrane by TonB-dependent transporters (TBDTs)¹⁵. Each siderophore or closely related siderophore has a specific TBDT that facilitates its uptake by the bacterial cells. The energy required for this transport is provided by the TonB–ExbB–ExbD complex. This complex functions as a molecular 'motor', that transmits energy from the inner membrane proton motive force to the TBDT in the outer membrane, facilitating the translocation of the siderophore-iron 8 complex into the periplasm⁹⁴⁻⁹⁶. More TBDTs are expressed than TonB-ExbB-ExbD complexes, indicating that a single TonB–ExbB–ExbD complex interacts with several TBDTs. I modified the text to better explain the message of Figure 4a.**] b)** Iron release from siderophores In Gram-negative bacteria. After the uptake of siderophore-Fe complexes across the outer membrane, the release of iron from the siderophore can occur either in the periplasm or the cytoplasm, depending on the 13 siderophore and bacterial species¹⁶. In the first case, after its dissociation from the chelator, iron associates with periplasmic binding proteins (PBP) to avoid any precipitation and is then transported across the inner membrane by a specific ABC transporter. In the second case, the siderophore-Fe binds in the periplasm to a PBP associated with an ABC transporter, which transports the siderophore-Fe complex further across the inner membrane. Iron release from siderophores always involves an iron reduction step, which can be associated or not with a chemical modification or hydrolysis of the siderophore. **c) Iron acquisition by siderophores in Gram-positive bacteria, exemplified by iron uptake in** *Staphylococcus aureus*100,101 **.** *S. aureus* produces two hydroxy acid-type siderophores, staphyloferrin A and staphyloferrin B, along with a hydroxamate-type siderophore containing citric acid residues called staphylobactin. It can also utilize hydroxamate (such as ferrichrome, desferrioxamine B, aerobactin and coprogen) and catecholate-type xenosiderophores. Another siderophore possibly produced by *S. aureus* is aureochelin, mentioned in a single article from 1997 25 (Ref. 189), with no recent literature to support the finding or identification of the transporter involved. Hence, aureochelin is not included in the figure. Although its structure remains unknown, it might resemble a phenolate catecholate-type siderophore. Iron uptake by staphyloferrin A and B, 28 staphylobactin, and xenosiderophores involves specific ABC transporters associated with receptors anchored in the inner membrane. Upon entry into the cytoplasm, iron is released from the siderophores through mechanisms involving iron reduction and pathway-specific mechanisms. Although the Fhu system has been extensively characterized in *S. aureus*, homologs have been identified in *L. monocytogenes*190, the *B. cereus* group191, and *S. pyogenes*¹⁹² . Like *S. aureus*, species within the *Staphylococcus*, *Listeria*, and *Bacillus* genera can utilize exogenous catecholate siderophores via the Sst uptake system, and these bacterial species can also import iron through this uptake pathway 35 using catecholamine stress hormones such as epinephrine, norepinephrine and dopamine^{191,193-198}.

 Figure 5. Iron acquisition by enterobactin and pyoverdine I. a) Iron acquisition by enterobatin in *P. aeruginosa* **and** *E. coli***.** Iron acquisition by enterobactin (Ent) in *P. aeruginosa* and *E. coli* is a process characterized by distinct mechanisms in each bacterium. *P. aeruginosa* does not produce the enterobactin siderophore but can utilize it. Upon the formation of the Ent–Fe complex in the environment of *P. aeruginosa*, recognition is facilitated by the TonB-dependent transporters (TBDTs) PfeA and PirA, with PfeA acting as the primary TBDT for this siderophore–Fe complex and PirA as the 8 secondary TBDT¹⁹⁹⁻²⁰¹. Following transport across the outer membrane by TBDTs, the iron complex dissociates from Ent in the periplasm. This step involves iron reduction and hydrolysis of Ent into three 10 molecules of 2,3-dihydroxybenzoic acid (DHBS) catalyzed by the esterase PfeE¹⁰³. In contrast, in *E. coli*, 11 a single TBDT, FepA, is involved in the import of the Ent–Fe complex across the outer membrane²⁰²⁻²⁰⁴. Within the periplasm, the Ent–Fe complex binds to the periplasmic protein FepB, which is associated 13 with an ATP-binding cassette transporter, FepBC, facilitating import across the inner membrane^{205–207}. The dissociation of iron from Ent in the cytoplasm involves a reductase, Ydjh, and an esterase, Fes, similar to the process observed in the periplasm of *P. aeruginosa*, resulting in the hydrolysis of Ent into three DHBS molecules104,104 . **b) Iron acquisition by pyoverdine I in** *P. aeruginosa***.** Pyoverdine I (Pvd) **[Au: pyoverdine I throughout? Yes]** is the primary siderophore synthesized and excreted by *P. aeruginosa*. When Pvd has chelated iron in the bacterial environment, the complex is recognized and 19 transported across the outer membrane by two TBDTs, namely FpvA and FpvB^{208–210}. FpvA is the major 20 transporter, whereas FpvB functions as the secondary transporter^{211,212}. Within the periplasm, the Pvd-21 Fe complex associates with two periplasmic proteins, FpvF and FpvC¹⁰⁸. FpvF belongs to the periplasmic 22 siderophore binding protein, and FpvC to the periplasmic iron binding proteins^{108,213}. The FpvF-Pvd-Fe- FpvC complex further associates with the inner membrane complex FpvGHJK, where FpvG functions 24 as a reductase, and the functions of the other three proteins (FpvH, FpvJ and FpvK) remain unknown 25 but are essential for optimal reductase activity of FpvG^{106,107,214}. Iron is reduced within the FpvCFGHJK 26 complex and dissociated from Pvd^{106,107}. The ferrous iron is then specifically chelated by FpvC²¹³, which 27 transports it to the ABC transporter FpvDE for translocation into the cytoplasm¹⁰⁸. Pvd remains associated with FpvF, which guides it to the PvdRT-OpmQ efflux pump, facilitating its export into the 29 extracellular milieu^{214–216}. As Pvd undergoes no chemical modification or degradation during the iron 30 release process, it can readily chelate another ferric ion, initiating the cycle anew¹⁰⁵.

Table of content:

 In this Review, Schalk explores the molecular mechanisms involved in siderophore-mediated iron acquisition in bacteria. In addition, the possible applications for siderophores in the environment, agriculture and medicine are also discussed.