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Selective Inhibition of Collagen Prolyl 4-Hydroxylase in Human Cells

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Abstract

Collagen is the most abundant protein in animals. Its overproduction is associated with fibrosis and cancer metastasis. The stability of collagen relies on post-translational modifications, the most prevalent being the hydroxylation of collagen strands by collagen prolyl 4-hydroxylases (CP4Hs). Catalysis by CP4Hs enlists an iron cofactor to convert proline residues to 4-hydroxyproline residues, which are essential for the conformational stability of mature collagen. Ethyl 3,4-dihydroxybenzoate (EDHB) is commonly used as a “P4H” inhibitor in cells, but suffers from low potency, poor selectivity, and off-target effects that cause iron deficiency. Dicarboxylates of 2,2'-bipyridine are among the most potent known CP4H inhibitors but suffer from a high affinity for free iron. A screen of biheteroaryl compounds revealed that replacing one pyridyl group with a thiazole moiety retains potency and enhances selectivity. A diester of 2-(5-carboxythiazol-2-yl)pyridine-5-carboxylic acid is bioavailable to human cells and inhibits collagen biosynthesis at concentrations that neither cause general toxicity nor disrupt iron homeostasis. These data anoint a potent and selective probe for CP4H and a potential lead for the development of a new class of antifibrotic and antimetastatic agents.

Collagen is the principal component of bone, connective tissues, and the extracellular matrix in animals.¹ The overproduction of collagen is associated with a variety of diseases, including fibrotic diseases² and cancers.^{3–7} The stability of collagen relies on

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ASSOCIATED CONTENT

Supporting Information

Experimental procedures, compound characterization, Table S1, and Figures S1–S9. This material is available free of charge via the Internet at <http://pubs.acs.org>

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posttranslational modifications that occur throughout the secretory pathway.⁸ By far the most prevalent of these modifications is the hydroxylation of collagen strands by collagen prolyl 4-hydroxylases (CP4Hs), which are Fe(II)- and α -ketoglutarate (AKG)-dependent dioxygenases (FAKGDs) located in the lumen of the endoplasmic reticulum.⁹ Catalysis by CP4Hs converts (2*S*)-proline (Pro) residues in procollagen strands into (2*S*,4*R*)-4-hydroxyproline (Hyp) residues (Figure 1A), which are essential for the conformational stability of mature collagen triple helices.¹⁰ Importantly, CP4Hs are validated targets for treating both fibrotic diseases¹¹ and metastatic breast cancer.⁶

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Like all enzymes of the FAKGD superfamily, catalysis by CP4Hs requires Fe(II) and the cosubstrates AKG and dioxygen.^{12–14} The Fe(II) is bound by a conserved His-X-Asp/Glu...X_n...His motif, and AKG chelates to enzyme-bound Fe(II) using its C-1 carboxylate and C-2 keto groups, while the C-5 carboxylate group forms hydrogen bonds and engages in Coulombic interactions with a basic residue (typically, arginine or lysine). All FAKGDs are believed to effect catalysis through a similar two-stage mechanism in which AKG is first decarboxylated oxidatively to generate a highly reactive Fe(IV)=O species (ferryl ion), which effects the hydroxylation of a hydrocarbon via a radical rebound process.^{12–14}

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In vertebrates, CP4Hs exist as $\alpha_2\beta_2$ tetramers. In these tetramers, the α -subunit contains the catalytic and substrate-binding domains, and the β -subunit is protein disulfide isomerase, which is a multifunctional protein that is responsible for maintaining the α -subunit in a soluble and active conformation.⁹ Three isoforms of the α -subunit, α (I), α (II), and α (III), have been identified in humans.⁹ All α -subunit isoforms form tetramers with the β -subunit, which we refer to herein as the CP4H1, CP4H2, and CP4H3 holoenzymes. CP4H1 is the most prevalent of the isoforms, and has been characterized most extensively. Whereas the structure of the tetrameric complex is unknown, those of the individual domains of the α (I)-subunit have provided insight into the interaction of CP4Hs with the procollagen substrate, as well as the dimerization of the α (I)-subunits to facilitate formation of the tetramer.^{15–18}

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The development of CP4H inhibitors has been of interest since the mid-1970s. Like many FAKGDs, human CP4Hs are inhibited by AKG mimics,¹⁹ such as *N*-oxalyl glycine (NOG), pyridine-2,4 dicarboxylic acid (24PDC), pyridine-2,5-dicarboxylic acid (25PDC), and 3,4-dihydroxybenzoic acid (DHB),²⁰ as well as by simple metal chelators, such as 2,2'-bipyridine (bipy) (Figure 1B). These compounds suffer from low potency in cellular assays, insufficient selectivity for CP4H, and intolerable cytotoxicity.^{21,22} Still, the ethyl ester of DHB (that is, EDHB) is often used as a cellular “P4H” inhibitor,^{23,24} even though DHB is not selective for CP4H compared to other FAKGDs, requires high dosing, and leads to an iron-deficient phenotype.²⁴

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The most potent inhibitors of human CP4Hs identified to date are bipyridinedicarboxylic acids (bipyDCs) (Figure 1B). Two bipyDCs have high potency: 2,2'-bipyridine-4,5'-dicarboxylic acid (bipy45'DC)²⁵ and 2,2'-bipyridine-5,5'-dicarboxylic acid (bipy55'DC).²⁶ Both of these bipyDCs inhibit human CP4H competitively with respect to AKG and bind selectively to human CP4H1 compared to PHD2, another human P4H.²⁵

An intrinsic property of bipyDCs limits their utility in a biological context. Like bipy itself, bipyDCs form tight complexes with free iron,²⁵ which is the dominant metal in life.²⁷ We hypothesized that replacing the bipy core with another biheteroaryl scaffold could lead to compounds that retain the potency and selectivity of bipy45'DC and bipy55'DC, but not their high affinity for free iron. Accordingly, we synthesized bipy analogs that retain one pyridyl group but replace the second with six distinct 5- or 6-membered nitrogen-containing heterocycles: pyrazine, pyridazine, oxazole, pyrazole, imidazole, pyrrole, and thiazole (Figure 2). We find that an asymmetric scaffold containing a thiazolyl group has especially desirable attributes as an inhibitor of human CP4H.

RESULTS AND DISCUSSION

Biheteroaryl Compounds

To begin our analysis, we prepared a library of six biheteroaryl compounds (Figure 2) and evaluated them as iron chelators *in vitro*. The library was assembled either from commercial vendors (pypyr and pyim) or by synthesis using palladium-catalyzed cross-coupling reactions (pypyraz, pypyrid, pyox, and pythi). To evaluate the iron affinity of these compounds *in vitro*, we performed titration experiments to determine the half-maximal concentration required to form a complex with 20 μM Fe(II) at pH 7.0. The ensuing $\text{Fe}_{20}\text{-EC}_{50}$ parameter provides a metric for iron affinity.²⁵ With the exception of pyox, all members of the library were observed to form distinct colored $\text{Fe}(\text{ligand})_3^{2+}$ complexes under these conditions (see: Supporting Information).

The $\text{Fe}_{20}\text{-EC}_{50}$ values of the biheteroaryl compounds varied by more than two orders of magnitude (Figure S1; Table 1). All, however, were significantly greater than that of bipy or bipyDCs.²⁵ The $\text{Fe}_{20}\text{-EC}_{50}$ value appeared to rely on both the ring size and $\text{p}K_{\text{a}}$ of the conjugate acid of the alternative heterocycle, as scaffolds with 5-membered rings were substantially weaker chelators than were those with 6 membered rings and $\text{p}K_{\text{a}}$ value correlated positively with iron affinity.

Biheteroaryl Dicarboxylates *in Vitro*

Inspired by the low affinity of the biheteroaryl compounds for free iron, we prepared a small library of biheteroaryl dicarboxylates (Figure 2) and interrogated those compounds as both iron chelators and inhibitors of human CP4H. With the exceptions of pyimDC (which was synthesized via a classical route to substituted imidazoles²⁸) and pypyrDC (which was synthesized by a 1,3-dipolar cycloaddition of a 2 ethynylpyridine with ethyl diazoacetate²⁹), all of the biheteroaryl dicarboxylates were synthesized by either palladium-catalyzed direct arylation of an appropriate heterocycle (pypyrazDC, pypyridDC, pyoxDC, pyoxDC*, pythiDC or pythiDC*) with a functionalized 2 bromopyridine³⁰⁻³⁴ or palladium-catalyzed oxidative cross-coupling (pypyrroleDC) between a functionalized pyridine *N*-oxide and an *N* protected pyrrole.³⁵ Typically, direct arylation using methyl- or ethyl-protected carboxylate esters allowed synthesis of the target compounds in 2–4 steps with an acceptable yield. For pyoxDC and pythiDC, cross-coupling yields using the typical inner-sphere base pivalic acid (PivOH) were prohibitively low (<5%, data not shown). We found that the addition of 1 adamantanecarboxylic acid rather than PivOH improved yields markedly (see: Supporting

Information) and encourage the continued investigation of 1 adamantanecarboxylic acid as an inner-sphere base in palladium-catalyzed direct arylation reactions.

We investigated iron chelation by the biheteroaryl dicarboxylates in a manner similar to that for the parent scaffolds. To our surprise, we were not able to detect complex formation by spectrophotometry for any of the biheteroaryl dicarboxylates at concentrations up to 1 mM, suggesting that the affinity of these compounds for free iron would be negligible in a biological context. Previously, we reported that various bipyDCs have $Fe_{20}\text{-}EC_{50}$ values that are similar to that of bipy itself,²⁵ so our discovery that biheteroaryl dicarboxylates investigated herein have an $Fe_{20}\text{-}EC_{50}$ value >1 mM represents an improvement of at least an order of magnitude.

Next, we assessed the biheteroaryl dicarboxylates as inhibitors of human CP4H1. To separate any inhibitory effect that derives from iron sequestration rather than enzymic binding, we employed previously described assay conditions (10 μM compound and 50 μM $FeSO_4$) in which potent chelators like bipy do not cause inhibition.²⁵ In this initial screen (Figure S2), we found that some biheteroaryl dicarboxylates showed little or no inhibition of human CP4H1, consistent with the inability of their heteroatoms to participate in an enzymic interaction. (Both pypyridDC and pypyrDC even showed modest activation under these conditions by a mechanism that is unclear.) Notably, we found that pyimDC, pyoxDC, and pythiDC were inhibitors of human CP4H1, with pyimDC and pythiDC demonstrating potency only a bit weaker than that of the bipyDCs. Importantly, the regioisomers pythiDC* and pyoxDC* did not show significant inhibition, suggesting that proper regiochemistry is essential for inhibition.

Unlike oxazole or thiazole, imidazole exists as two tautomers, one with a proton on N1 (as in the depiction of pyimDC in Figure 2) and another with a proton on N3. Although we did not observe the formation of a complex between pyimDC and free iron by spectrophotometry, we examined this issue more thoroughly. We found that pyimDC was able to deter the formation of the $Fe(\text{bipy})_3^{2+}$ complex in a dose-dependent manner (Figure S3). Moreover, competition required a free carboxylate on the imidazole ring. These data are consistent with the formation of a $Fe(\text{pyimDC})_2$ complex with N1 bound to iron. To eliminate this mode of binding, we synthesized NMe-pyimDC (Figure 2), which is an analogue of pyimDC that is methylated on N1. We found that NMe-pyimDC was able to deter the formation of the $Fe(\text{bipy})_3^{2+}$ complex, but only at high concentrations (Figure S3). We also found that NMe pyimDC is an inhibitor of human CP4H *in vitro*, but that its potency is less than that of pyimDC (Figure S2).

In subsequent dose–response experiments, we found the inhibition curves for pyimDC, pythiDC, and pyoxDC to be sigmoidal, yielding IC_{50} values in the low-micromolar range (Figure 3A). The potencies of pyimDC and pythiDC were approximately 10-fold greater than that of pyoxDC. A Lineweaver–Burke analysis of inhibition of CP4H by pythiDC demonstrated competitive inhibition with respect to the AKG cosubstrate (Figure 3B).

Like bipy45'DC and bipy55'DC,^{26,25} pyimDC, pyoxDC, and pythiDC could bind in the AKG binding pocket and use their second carboxyl group to form additional interactions

with residues in the active site of CP4H. If so, then the biheteroaryl dicarboxylates should exhibit structure–activity relationships similar to those of bipy45′DC and bipy55′DC for the inhibition of PHD2, another P4H enzyme that has been characterized extensively. We find that pyimDC, pythiDC, and pyoxDC inhibit human PHD2 only weakly (Figure S4), with pythiDC displaying especially low potency.

Biheteroaryl Dicarboxylates *in Cellulo*

Encouraged by the results *in vitro*, we next sought to determine if pyimDC and pythiDC could be suitable for use *in cellulo*. First, we determined the effect of these compounds on iron metabolism in human cells. Because human CP4Hs are validated therapeutic targets for breast cancer, we chose the MDA-MB-231 human breast cancer cell line as a primary model system.⁶ We performed additional analyses in human embryonic kidney (HEK) cells. We assessed iron metabolism with immunoblots for ferritin, the transferrin receptor (TfR), and the transcription factor HIF-1 α , all of which are known to give distinct phenotypes depending on the status of iron in a human cell.²⁴ More specifically, levels of ferritin and TfR are regulated by iron regulatory proteins 1 and 2 (IRPs), which have iron-dependent RNA-binding activity that modulates the expression of target genes at the level of translation.^{36,37} The stability of HIF-1 α is dependent on the prolyl 4-hydroxylase activity of PHD2, which is an activity that is inherently sensitive to the iron status of the cell. Thus, iron-deficient cells exhibit ferritin levels that are lower and TfR and HIF-1 α levels that are higher than those of untreated cells.

Ethyl dihydroxybenzoate (EDHB) an AKG mimic that is commonly used as a “P4H” inhibitor in cell culture models, served as our benchmark.^{23,24} Calculations suggested that the diethyl esters of pyimDC and pythiDC have log P values conducive to cellular uptake (Table S1), and we synthesized those two esters. Moreover, the iron affinity of these diethyl esters remained sufficiently low (see: Section XXII in the Supporting Information and Figure S1), encouraging their use in cellular assays. Cultured MDA-MB-231 cells are known to secrete large amounts of collagen.⁶ Due to the importance of CP4H-dependent hydroxylation for collagen stability, the level of collagen secreted by MDA-MB-231 cells is dependent on the activity of this enzyme. Thus, MDA-MB-231 cells are ideal for investigations of both iron deficiency and CP4H activity.

Toward this end, we treated MDA-MB-231 cells with biheteroaryl dicarboxylates, and assayed for cytotoxicity and indicators of iron deficiency. None of the esterified biheteroaryl dicarboxylates exhibited cytotoxic activity at high micromolar concentrations (see: Section XI in the Supporting Information). Cells treated with EDHB demonstrated a strong iron-deficient phenotype, as expected (Figures 4A and 4B). In contrast, cells treated with diethyl pythiDC appeared to be normal at concentrations as high as 500 μ M. Interestingly, diethyl pyimDC showed an intermediate phenotype characterized by a significant decrease in ferritin levels but without an associated accumulation of TfR or HIF-1 α . This phenotype was, however, not observable at diethyl pyimDC concentrations \leq 56 μ M. The results in MDA-MB-231 cells were replicated in HEK cells. Again, treatment with diethyl pythiDC and low levels of diethyl pyimDC (as well as simply pyim and pythi; Figure S5) did not affect the level of TfR, HIF-1 α , or ferritin (Figure S6).

Next, we assessed the function of IRPs in cells treated with the esterified biheteroaryl compounds. Using a radiolabeled RNA ligand, we performed electrophoretic mobility shift assays on IRPs from treated MDA-MB-231 cells. Whereas DFO, EDHB, and bipy all caused significant increases in RNA-binding by IRPs, diethyl pyimDC and diethyl pythiDC did not (Figure S7). This result is again consistent with these compounds having little effect on cellular iron levels.

Lastly, we examined the effect of treating MDA-MB-231 cells with the esterified biheteroaryl compounds on the cellular secretion of type I collagen (Figure 4C), which relies on CP4H activity.⁶ We found that treatment with either diethyl pyimDC or diethyl pythiDC reduced the level of secreted collagen significantly (Figure 4C). Moreover, the efficacy of diethyl pyimDC or diethyl pythiDC was indistinguishable from that of diethyl bipy55'DC. No inhibition of catalysis by CP4H1 or PHD2 was observed *in vitro* with these diethyl esters at a concentration of 100 μ M (data not shown), confirming that the inhibitory potential of these compounds is masked by esterification. Lastly, treatment with diethyl pyimDC and diethyl pythiDC did not appear to affect the levels of human CP4H1 itself (Figure S8), consistent with the observed reduction in collagen secretion arising from inhibition of CP4H.

Conclusions

We have identified compounds that can replace EDHB in experiments with cultured cells. Unlike EDHB, we found that diethyl pythiDC, diethyl pyimDC, and even diethyl bipy55'DC inhibit CP4H activity in cultured cells at concentrations that do not cause iron deficiency. Given the subtle ferritin phenotype caused by diethyl pyimDC and the measurable iron affinity of bipy55'DC and related bipyDCs *in vitro*,²⁵ we put forth diethyl pythiDC as a preferred replacement for EDHB. Moreover, an esterified pythiDC could serve as a lead for clinical development.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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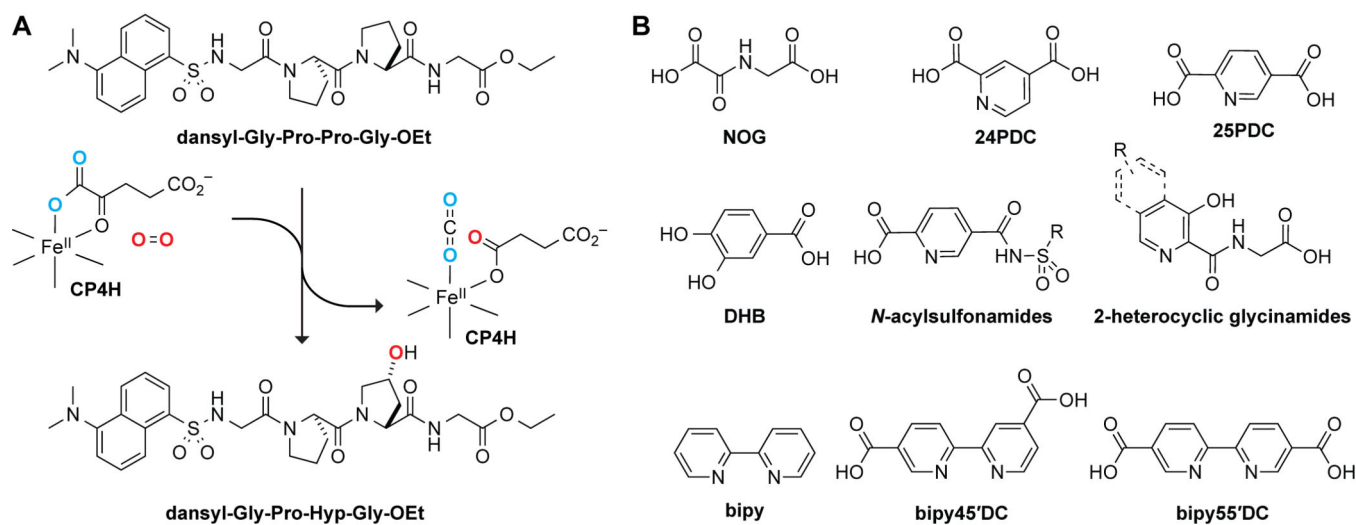


Figure 1. Reaction catalyzed by collagen prolyl 4-hydroxylase (CP4H) and its inhibition. (A) In an Fe(II)- and AKG-dependent reaction, CP4Hs catalyze the hydroxylation of Pro residues in collagenous peptides to form Hyp residues. (B) Examples of known human CP4H inhibitors.

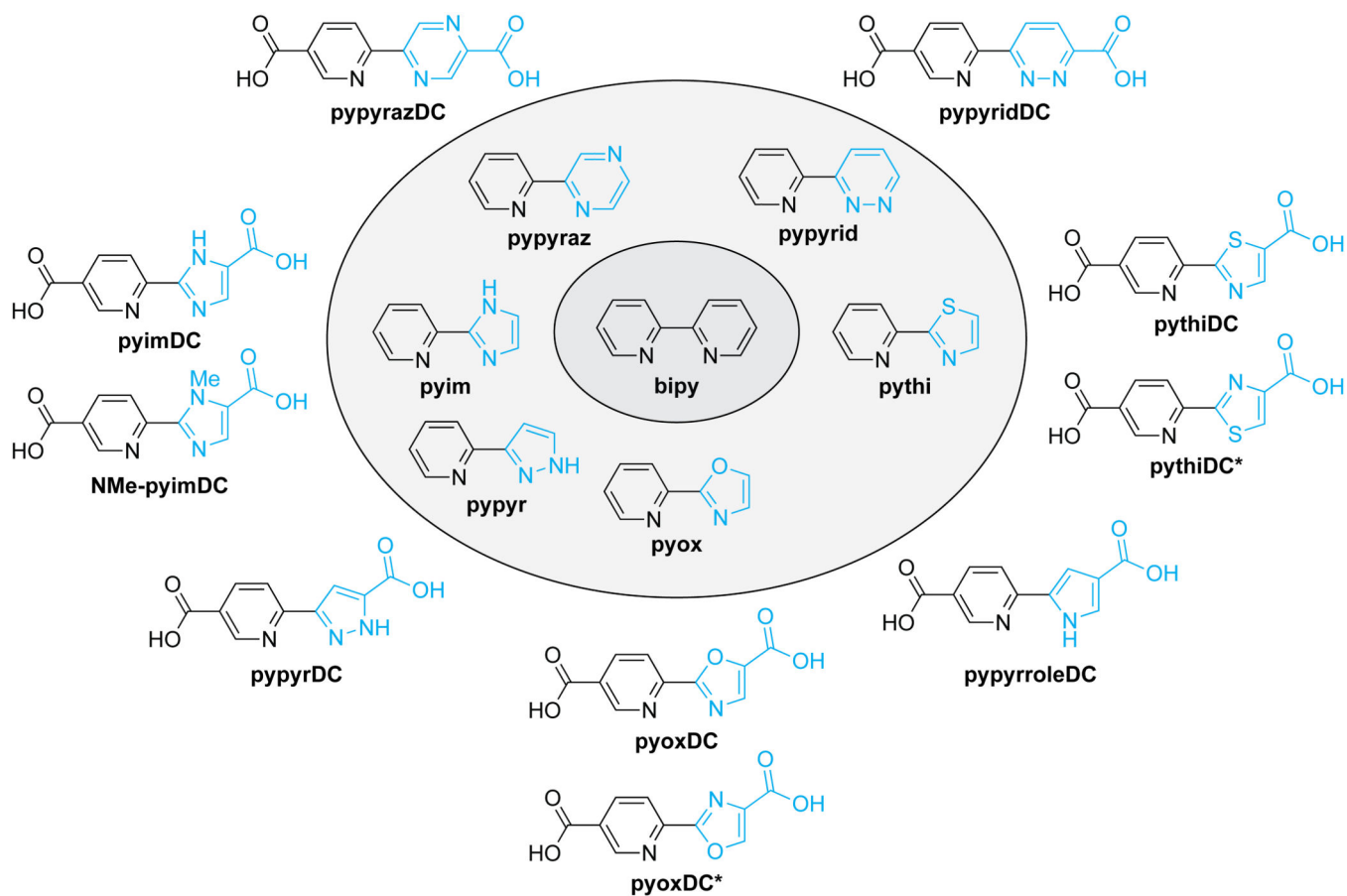
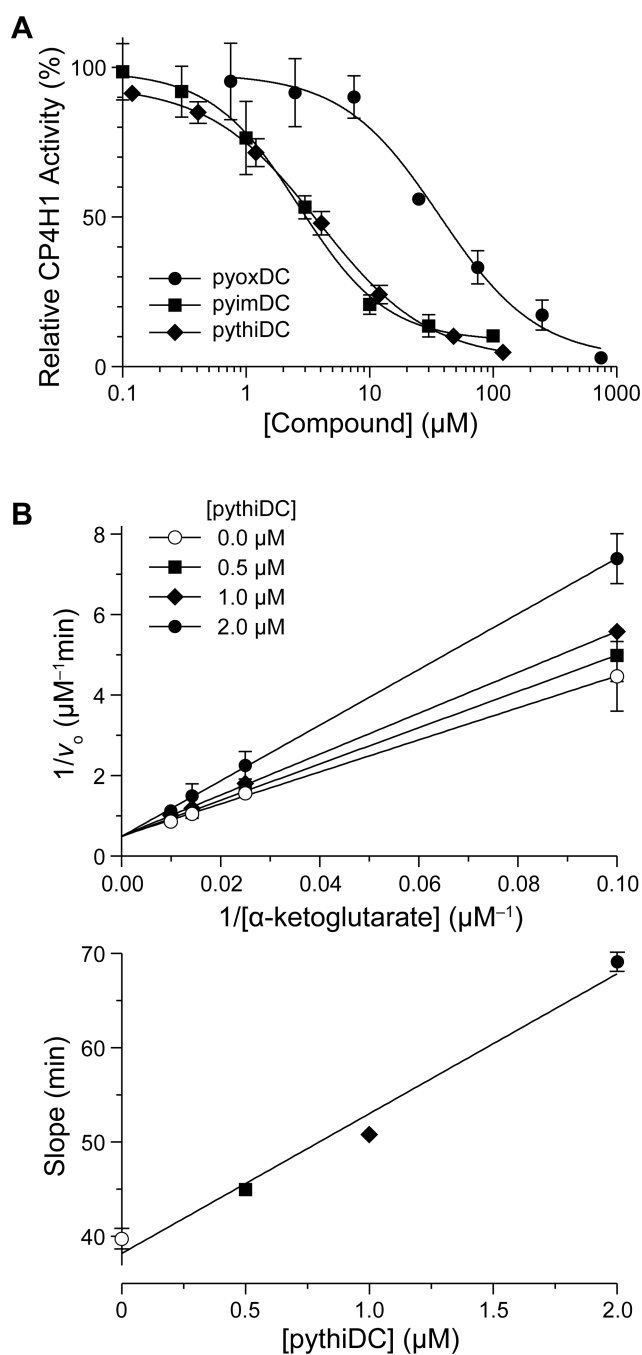


Figure 2. Biheteroaryl compounds used in this study. Novel moieties are rendered in cyan, and similar compounds are grouped together.

**Figure 3.**

Biheteroaryl dicarboxylates as inhibitors of human CP4H1. (A) Individual points represent the mean (\pm SD) of three independent experiments. Data were fitted to a dose–response equation to determine IC_{50} values: pyimDC, (2.6 ± 0.1) μM ; pyoxDC, (33 ± 8) μM ; pythiDC, (4.0 ± 0.2) μM . (B) Lineweaver–Burke analysis of inhibition by pythiDC. The rate of the reaction catalyzed by CP4H1 with increasing α -ketoglutarate concentration (10–100 μM) was determined in the presence of a fixed concentration of pythiDC (0.0, 0.5, 1.0, or 2.0 μM). Individual points represent the mean (\pm SE) of two independent experiments. Data

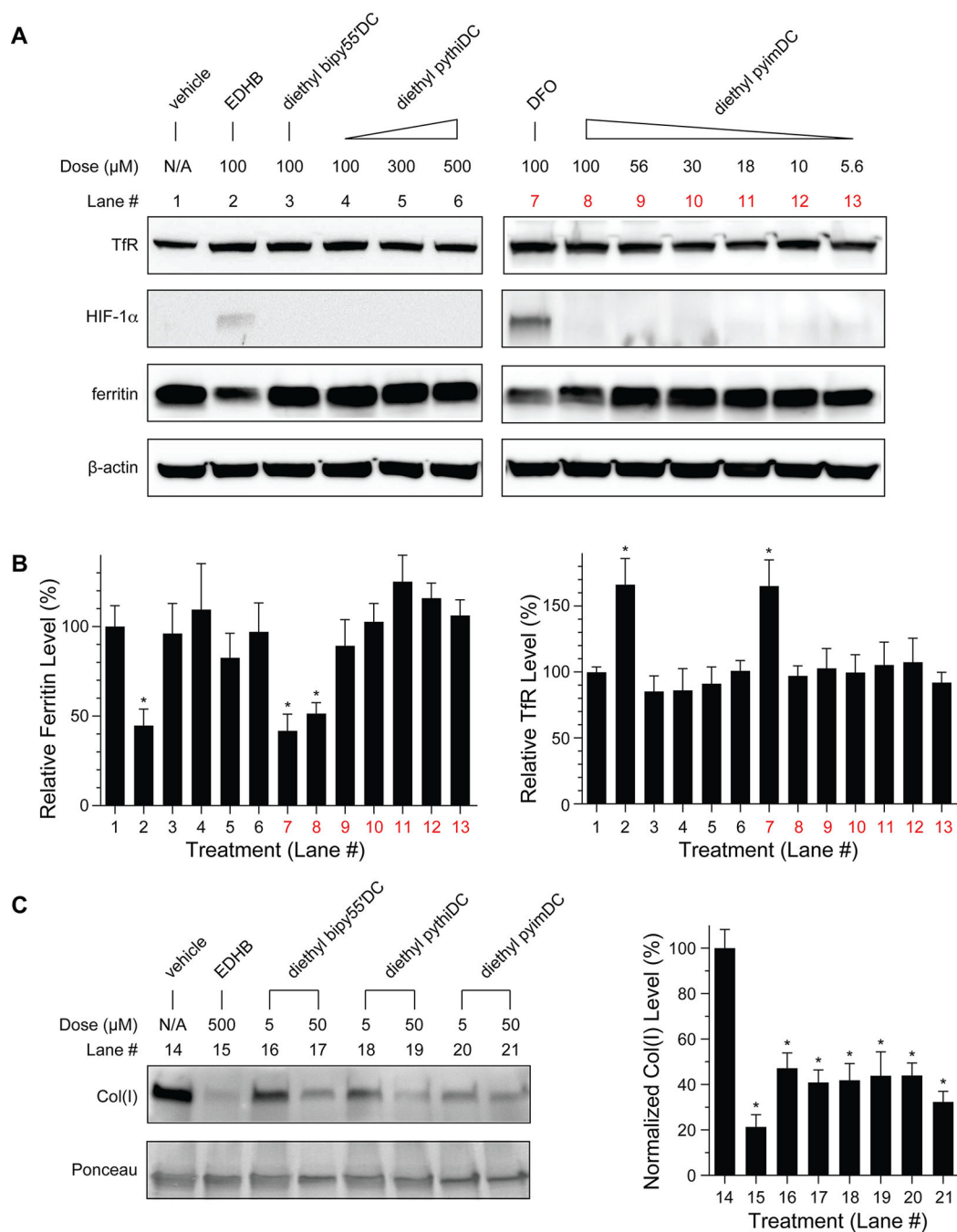
were fitted by linear regression to determine a K_i value of (0.39 ± 0.04) μM for competitive inhibition.

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biheteroaryl compounds on collagen secretion into conditioned media. Blots are representative of at least 5 replicates and quantitations (right) are normalized to total protein using the Ponceau S stained blot.

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Table 1

Iron binding by bipy and biheteroaryl compounds.

Compound	Heteroaryl pK_a^a (ref.)	pK_a^b	Fe(II) Complex λ_{max}^c	Stoichiometry (Ligand:Fe) ^d	Fe ₂₀ -EC ₅₀ ^e
bipy	5.2 (38)	4.3	523	3:1	43 ± 2
pyrid	2.3 (39)	<3	522	3:1	124 ± 1
pypraz	0.8 (40)	<3	523	3:1	400 ± 7
pyim	7.1 (41)	5.4	484	3:1	900 ± 30
pypr	2.5 (42)	4.1	441	ND ^f	2250 ± 100
pythi	2.5 (43)	<3	524	3:1	5100 ± 100
pyox	0.8 (44)	<3	NO ^g	NO ^g	>18,000

^a pK_a value of the conjugate acid of pyridine (bipy) or the non-pyridyl ring.

^b pK_a value of the conjugate acid as estimated by titration of the entire compound; bipy, ref. 45.

^c Determined using spectrophotometry (see: Supporting Information).

^d Determined by Job's method.^{46,47}

^e Determined by titration of 20 μ M FeSO₄; bipy, ref. 25. Values are the mean (\pm SE) of 3 replicates.

^f Not determined due to prohibitive competing absorbance from iron.

^g Not observed.