Regioselective chemoenzymatic syntheses of ferulate conjugates as chromogenic substrates for feruloyl esterases

Olga Gherbovet, Fernando Ferreira, Apolline Clément, Mélanie Ragon, Julien Durand, Sophie Bozonnet, Michael J. O’Donohue and Régis Fauré*

Abstract
Generally, carbohydrate-active enzymes are studied using chromogenic substrates that provide quick and easy color-based detection of enzyme-mediated hydrolysis. For feruloyl esterases, commercially available chromogenic ferulate derivatives are both costly and limited in terms of their experimental application. In this study, we describe solutions for these two issues, using a chemoenzymatic approach to synthesize different ferulate compounds. The overall synthetic routes towards commercially available 5-bromo-4-chloro-3-indolyl and 4-nitrophenyl 5-O-feruloyl-α-L-arabinofuranosides were significantly shortened (from 7 or 8 to 4–6 steps), and the transesterification yields were enhanced (from 46 to 73% and from 47 to 86%, respectively). This was achieved using enzymatic (immobilized Lipozyme® TL IM from Thermomyces lanuginosus) transesterification of unprotected vinyl ferulate to the primary hydroxy group of α-L-arabinofuranosides. Moreover, a novel feruloylated 4-nitrocatechol-1-yl-substituted butanetriol analog, containing a cleavable hydroxylated linker, was also synthesized in 32% overall yield in 3 steps (convergent synthesis). The latter route combined the regioselective functionalization of 4-nitrocatechol and enzymatic transferuloylation. The use of this strategy to characterize type A feruloyl esterase from Aspergillus niger reveals the advantages of this substrate for the characterizations of feruloyl esterases.

Introduction
The development of “white biotechnology” is underpinned by advances in enzyme discovery and engineering, areas that are being driven by metagenomics and in vitro-directed enzyme evolution. These techniques procure massive discovery or the creation of new enzyme-encoding sequences, filling up databases with a wealth of information. However, while resolving an early step in the discovery pipeline, these techniques progressively create a new bottleneck regarding enzyme characterization. Therefore, there is a pressing need to extend the enzymologist’s toolbox, providing informationally rich high-
throughput screens that can not only attribute an activity to putative enzymes but also procure some qualitative details on enzyme properties. In this respect, the availability of easy-to-use chromogenic substrates that can provide both qualitative and quantitative assays and are compatible with automatized protocols is a crucial issue.

Feruloyl esterases (Faes; EC 3.1.1.73 and the family CE1 of the CAZy classification [1]) are of interest, both because of their role in the deconstruction of complex plant-based materials and also as synthetic tools for the preparation of bioactive compounds with potential antioxidant properties [2-5]. Operating via a two-step serine protease mechanism involving a conserved Ser-His-Asp/Glu catalytic triad [6,7], Faes catalyze the hydrolysis of ester bonds linking hydroxycinnamoyl groups to the glycosyl moieties of plant-based polysaccharides, such as arabinoxylans and arabinans. In this respect, Faes are important components of plant-cell-wall-degrading enzymatic arsenals since the hydrolysis of trans-ferulate-polysaccharide linkages contributes to the breakdown of intermolecular bonds that structure the lignocellulosic matrix. Moreover, Faes are useful tools to obtain commercially relevant ferulic acid, which represents up to 3 wt % of plant the cell wall dry weight [8].

So far, the detection and characterization of Faes has mainly relied on either the use of HPLC or on UV–visible spectrophotometry, using natural or synthetic compounds [9,10]. The latter, which are used in high-throughput screening (HTS) assays, fall into three categories. The simplest are feruloyl esters of chromogenic moieties [11-14], such as p-nitrophenol or short-chain alkyl groups (e.g., methyl ferulate). More elaborate and biologically relevant substrates contain a feruloylated α-L-arabinofuranosyl moiety [12,15-17]. These structurally more complex compounds are obtained using multistep syntheses, considerably limiting the availability. Moreover, they might be specific to certain subcategories of feruloyl esterases [18-20], and their use involves a challenging tandem reaction [21]. Finally, the synthesis of other, more generic esters that can be used to assay esterases, including Faes, and lipases have been reported [22,23].

In this work, we revisit the preparation of simple feruloylated substrates, such as the 5-bromo-4-chloro-3-indolyl and 4-nitrophenyl 5-O-feruloylated α-L-arabinofuranosides 1a and 1b. Although these substrates are commercially available, their synthesis involves 7 or 8 steps [15-17]. This contributes to their rather high retail costs (e.g., as of July 29th, 2020, 2500 and 778 € per 100 mg for 1a and 1b, respectively), which are approximately 19- and 14-fold higher than non-feruloylated precursors. Therefore, our aim was to simplify the synthesis in order to reduce the cost. Furthermore, we describe the short synthesis of the new feruloylated chromogenic substrate 12, a molecule that obviates the need for a glycosyl moiety while containing a cleavable hydroxylated linker that mimics the natural geometry and physicochemical properties of glycosidic linkages.

**Results and Discussion**

**Chemoenzymatic synthesis of 5-O-feruloylated α-L-arabinofuranosides**

The synthesis of the chromogenic 5-O-feruloylated α-L-arabinofuranosides 1a and 1b is usually achieved using a multistep pathway that involves trapping the furanose conformation, anomeric activation, glycosidation, regioselective deprotection of the primary hydroxy group, feruloylation, and final deprotection to yield the target molecule [12,15-17]. Additionally, the temporary protection of the functional groups is sometimes used during the synthesis in order to facilitate certain steps. Using an alternative approach, we employed a one-step regioselective transesterification of the unprotected vinyl ferulate 2 (synthesized in 56% in-house yield and in up to 77% previously reported yield [24,25] in one step on a gram scale) using Lipozyme® TL IM (a commercially available immobilized lipase from *T. lanuginosus* that efficiently catalyzes the transesterification of cinnamates) [26-28] and readily available and reasonably cheap 5-bromo-4-chloro-3-indolyl or 4-nitrophenyl α-L-arabinofuranosides. This afforded the corresponding feruloylated derivatives 1a and 1b (Figure 1). The yields (73 and 86% for the indolyl and 4-nitrophenyl derivatives, respectively) characterizing the regioselective enzymatic feruloylation of the primary hydroxy group compare favorably with the previously reported overall yield (46 and 47%, respectively, in three steps) [15-17,29], which relate to the selective enzymatic O-5-deacetylation and esterification of the primary hydroxy group and a final deprotection of the 2,3-O-acetyl groups of the glycoside and the O-acetyl group of the ferulate moiety. The fact that lipase-catalyzed transesterification obviates the need for protection and deprotection is a considerable advantage because the final deprotection in the chemical pathway is complicated by the presence of another ester linkage within the molecules [12-15]. In principle, the method described herein is generic, and thus applicable to other chromogenic α-L-arabinofuranosidic compounds, such as 4-nitrocatechol (4NTC), 2-chloro-4-nitrophenyl, and umbelliferyl derivatives.

To demonstrate the suitability of 1a for qualitative in situ screening of microbial colonies growing on solid agar medium, the compound (300 µg/mL) was incorporated into top agar, along with an α-L-arabinofuranosidase from *Thermobacillus xylanilyticus* (TxAbf). This provided the means to reveal colored microbial colonies expressing the type A Fae from
Aspergillus niger (AnFaeA, Figure 1C). Coloration is the result of successive reactions: (i) release of the free 5-bromo-4-chloro-indoxyl-3-ol by an enzyme cascade, wherein TxAbf-catalyzed cleavage of the glycosidic bond was made possible by the prior release by AnFaeA of the ferulate moiety [21] and (ii) spontaneous oxidation and subsequent dimerization forming the insoluble blue 5-bromo-4-chloro-substituted indigo dye [15]. In an alternative demonstration, 1a was also used for colony level detection of FAE using yeast cells (Yarrowia lipolytica [30]) that actually coexpressed AnFaeA and TxAbf (data not shown).

Investigating the enzymatic transferuloylation reaction of non-glycosidic motifs

While lipase-catalyzed transesterifications of cinnamate or phenolic derivatives onto glycosidic structures have been extensively described [25,27,28,31-33], data related to the regioselectivity of transesterifications of hydroxylated alkyl and/or aryl moieties are sparser [26,34-36]. The extent to which Lipozyme® TL IM catalyzes feruloyl transfer reactions involving substituted benzylic alcohols was thus investigated to establish its usefulness for the preparation of both various polyhydroxylated molecules of interest (e.g., antioxidants) [4,26] and of novel chromogenic feruloylated substrates with various physicochemical features for screening applications. Accordingly, we observed that transesterifications only occurred when using primary benzylic alcohols; no phenol acylation was detected with hydroxynitrobenzylic alcohol (Table 1, no side product of 4 and 9 with transfer on an aromatic secondary alcohol; i.e., phenol), 2-chloro-4-nitrophenol, or 4NTC (data not shown). Additionally, the exact position of the benzylic alcohol affected the transfer, with ortho-substitutions (R1) displaying a hindered electron-withdrawing group. This led to a low target product
yield (i.e., 5, 6, 8, and 9), a lower reactivity, and/or a higher hydrolysis of the vinyl ferulate into ferulic acid.

Synthesis of L-arabinofuranoside-free 4-nitrocatechol-1-yl–linker–ferulate chromogenic substrate 12 and evaluation of 12 as a chromogenic substrate for Fae assays

To synthesize chromogenic (±)-4-O-(2-hydroxy-4-nitrophenyl)-1-O-trans-feruloyl-1,2,4-butanetriol (4NTC–linker–Fe, 12), which contains 4NTC bound via a cleavable linker to a ferulate motif, a multistep route was devised (Scheme 1). First, a shorter, more practical pathway towards racemic (±)-4-O-(2-hydroxy-4-nitrophenyl)-1,2,4-butanetriol (11) was developed. Compared to the previously reported 4-step synthesis [37], two drawbacks were circumvented, notably avoiding (i) the preparation of the volatile (S)-1-ido-3,4-O-isopropylidene-3,4-butanediol intermediate and (ii) the use of a protected version of the chromogenic linker 11, either for the extra hydroxy group of the catecholyl moiety or the secondary hydroxy group of the linker [23,37]. It is noteworthy that a racemic mixture of this linker has previously been used to prepare chromogenic substrates for esterases [23,38].

The alkylation of 4NTC with homoallylic bromide under basic conditions gave a mixture of mono- and dialkylated 4NTC derivatives, with 10 (38%) predominating because of the preferential formation of the phenolate at the para-position (relative to the nitro group) [39]. Osmium tetroxide-mediated dihydroxylation in the presence of N-methylmorpholine N-oxide (NMNO)
afforded 11 in 90% yield. Finally, the regioselective transferuloylation of the primary hydroxy group of the triol derivative 11 with Lipozyme® TL IM was performed, and the expected chromogenic substrate 12 was isolated in a high yield of 94%. Accordingly, the synthesis of the chromogenic ferulate 12 was achieved in 32% overall yield in 3 steps from commercial reactants (convergent synthesis using a slight excess of the synthesized vinyl ferulate 2) and without the requirement to perform a final deprotection.

As expected, the investigation of the stability of the chromogenic substrate 12 using UV–visible spectrophotometry revealed that, unlike 4-nitrocatechol-1-yl ferulate (4NTC–Fe), which undergoes spontaneous hydrolysis even at a neutral pH value and 40 °C [13], the presence of the alkyl linker procures a higher stability over a wide pH value range (up to pH 9.0), irrespective of the temperature. This is because in compound 12, the ferulate moiety is not directly linked to the good leaving group 4NTC (pKₐ = 6.61 [40]). Instead, it is bonded to the linker with a pKₐ value that can be compared either to that of glycerol (pKₐ = 13.61) or Л-arabinofuranosyl (pKₐ = 11.31) [41], meaning that it is a poor leaving group. Moreover, our observations regarding the linker stability are consistent with the known stability of the ester linkages under basic conditions.

The usefulness of 4NTC–linker–Fe (12) for the characterization of Faes was investigated (Figure 2), measuring 4NTC release by AnFaeA [42] at 40 °C. The enzyme-catalyzed reaction leads to the cleavage of the ester bond linking the ferulate to the linker–4NTC moiety, and thus to the accumulation of linker–4NTC. Therefore, working in a discontinuous mode, 4NTC is quantified by submitting samples removed from the reaction to the oxidative action of sodium periodate at 0 °C and registering the absorbance at 530 nm (under alkaline conditions) [23,37,38,43]. Importantly, it is vital to include a stoichiometric amount of ethylene glycol to avoid further oxidation of free 4NTC by sodium periodate (Figure 2B).

The specific activity (SA) of AnFaeA on 4NTC–linker–Fe (12) was determined to be 3 IU/mg of the protein (with IU corresponding to the international unit of the Fae hydrolytic activity), a value comparable to that measured on destarched wheat bran (3 IU/mg) [44] containing 5-O-feruloylated α-L-arabinofuranosyl moieties, but lower than that (40 IU/mg, unpublished data) measured using the more labile 4NTC–Fe. Therefore, although 4NTC–Fe is a practical synthetic probe for both the high-throughput screening and the preliminary characterization of the Fae activity [13], 4NTC–linker–Fe (12) is almost certainly a better analogue of ferulate linkages found in plant-based structures.

**Conclusion**

The use of immobilized Lipozyme® TL IM provides the means to perform the regioselective transesterification of the vinyl ferulate 2 to the primary hydroxy group of benzylc alcohols and polyhydroxylated compounds. Three compounds suitable for the detection and/or characterization of Fae activity were synthesized in a straightforward protocol that holds the potential to greatly reduce the cost of the substrates 1a and 1b. Moreover, the enzyme-driven convergent synthesis of 12 affords a
novel substrate that is highly suitable for the characterization of feruloyl esterases.

**Experimental**

**Materials and general methods**

4-Nitrophenyl α-L-arabinofuranosides and 5-bromo-4-chloro-3-indolyl α-L-arabinofuranosides were purchased from Carbosynth and Lipozyme® TL IM (immobilized lipase from *T. lanuginosus*, 250 IUN/g. IUN = interesterification unit) was supplied by Novozymes. The reaction evolution was monitored by analytical thin-layer chromatography using silica gel 60 F 254 precoated plates (Merck). Spots were visualized using UV light of 254 nm wavelength, followed by soaking in a 0.1% (w/v) orcinol solution containing a sulfuric acid/ethanol/water 3:72.5:22.5, v/v/v mixture, followed by charring. Purification by column chromatography was performed using a Reveleris® flash chromatography automated system (Büchi) equipped with prepacked irregular silica gel 40–63 μm cartridges (FlashPure EcoFlex, Büchi). NMR spectra were recorded on a Bruker Avance II 500 spectrometer at 298 K. Chemical shifts (δ) are given in ppm, with the signals of the residual solvents as the internal references [45]. The coupling constants (J) are reported in Hertz (Hz), with singlet (s), doublet (d), triplet (t), doublet of doublets (dd), doublet of triplets of doublets (ddd), broad (br), and quadruplet of triplets (qt) as abbreviations. Analyses and assignments were made using 1D (1H, 13C, and J-modulated spin–echo (*J*mod)) and 2D NMR experiments (COSY and HSQC). High-resolution mass spectrometry (HRMS) analyses were performed at PCN-ICMG (Grenoble, France). Optical rotations were measured using a JASCO P-2000 polarimeter at 20 °C.

**General procedure for enzymic transesterifications**

The enzymatic transesterification steps were performed according to a published protocol [32]. Briefly, Lipozyme® TL IM (1 g, reusable) was added to a solution of alcohol (0.30 mmol, 1 equiv) and the vinyl ferulate 2 (100 mg, 0.45 mmol, 1.5 equiv) in acetonitrile (6 mL). The reaction mixture was stirred overnight at 37 °C, then filtered, the filter cake was washed with acetone, and the filtrate was evaporated to dryness. The residue was recovered in ethyl acetate, and washed three times with saturated aqueous sodium hydrogen carbonate. The combined organic phases were dried over anhydrous sodium sulfate, filtered, and concentrated under reduced pressure. Flash chromatography (gradient of ethyl acetate in petroleum ether from 0 to 50%) afforded the pure ferulates 1a, 1b, 3–9, and 12.

5-Bromo-4-chloro-3-indolyl 5-O-trans-feruloyl-α-L-arabinofuranoside (1a, 107 mg, 0.19 mmol, 73%). Green-yellow foam. The NMR data (CD3OD) were consistent with those previously reported [15].

4-Nitrophenyl 5-O-trans-feruloyl-α-L-arabinofuranoside (1b). The application of the general procedure for enzymatic transesterifications was used with commercial 4-nitrophenyl α-L-arabinofuranoside (50 mg, 0.18, mmol, 1 equiv) and the vinyl ferulate 2 (61 mg, 0.28 mmol, 1.5 equiv) to give 1b (71 mg, 0.16 mmol, 86%) as a yellowish powder. The NMR data (CD3OD) were consistent with those previously reported [16,17].

3-Nitrobenzyl trans-ferulate (3, 78 mg, 0.24 mmol, 79%), white powder. 1H NMR (500 MHz, (CD3)2CO, δ) 8.32 (t, *J* = 1.7, 1H, CH of Bn), 8.21 (ddd, *J* = 8.3, 2.2, 1.1, 1H, CH of Bn), 7.90 (ddd, *J* = 7.6, 1.7, 1.0, 1H, CH of Bn), 7.71 (t, *J* = 7.6, 1.6, CH of Bn), 7.68 (d, *J* = 15.8, 1H, CH=CHCO2), 7.37 (d, *J* = 2.0, 1H, CH of Fe), 7.17 (ddd, *J* = 8.3, 2.0, 1H, CH of Fe), 6.87 (d, *J* = 8.3, 1H, CH of Fe), 6.50 (d, *J* = 15.8, 1H, CH=CHCO2), 5.38 (s, 2H, CH2 of Bn), 3.92 (s, 3H, OMe); 13C NMR (125 MHz, (CD3)2CO, δ) 167.2 (C=O), 150.3 (Cq), 149.4 (Cq), 148.8 (Cq) 146.7 (CH=CHCO2), 140.2 (Cq), 135.1 (CH of Bn), 130.8 (CH of Bn), 127.3 (Cq), 124.2 (CH of Fe), 123.6 (CH of Bn), 123.4 (CH of Bn), 116.1 (CH of Bn), 115.1 (CH=CHCO2), 111.4 (CH of Fe), 65.1 (CH2 of Bn), 56.4 (OMe); HRESIMS (m/z): [M – H]− calcd for C12H12NO6, 328.0821; found, 328.0833.

2-Hydroxy-5-nitrobenzyl trans-ferulate (4, 79 mg, 0.23 mmol, 76%), white powder. 1H NMR (500 MHz, (CD3)2CO, δ) 8.29 (d, *J* = 2.8, 1H, CH of Bn), 8.14 (dd, *J* = 8.9, 2.8, 1H, CH of Bn), 7.68 (d, *J* = 15.8, 1H, CH=CHCO2), 7.38 (d, *J* = 1.8, 1H, CH of Fe), 7.17 (dd, *J* = 8.0, 1.8, 1H, CH of Fe), 7.12 (d, *J* = 8.9, 1H, CH of Bn), 6.88 (dd, *J* = 8.0, 1H, CH of Fe), 6.52 (d, *J* = 15.8, 1H, CH=CHCO2), 5.33 (s, 2H, CH2 of Bn), 3.92 (s, 3H, OMe); 13C NMR (125 MHz, (CD3)2CO, δ) 167.5 (C=O), 162.0 (Cq) 150.3 (Cq), 148.8 (Cq), 148.7 (Cq), 146.6 (CH=CHCO2), 141.6 (Cq), 127.4 (Cq), 126.3 (2CH of Bn), 124.3 (CH of Fe), 116.5 (CH of Bn), 116.1 (CH of Fe), 115.2 (CH=CHCO2), 111.3 (CH of Fe), 61.1 (CH2 of Bn), 56.4 (OMe); HRESIMS (m/z): [M – H]− calcd for C12H12NO7, 344.0770; found, 344.0778.

2-Chloro-5-nitrobenzyl trans-ferulate (5, 48 mg, 0.13 mmol, 44%), white powder. 1H NMR (500 MHz, (CD3)2CO, δ) 8.42 (br d, *J* = 3.0, 1H, CH of Bn), 8.26 (dd, *J* = 9.0, 3.0, 1H, CH of Bn), 7.81 (d, *J* = 9.0, 1.0, CH of Bn), 7.71 (d, *J* = 16.0, 1H, CH=CHCO2), 7.40 (d, *J* = 1.8, 1H, CH of Fe), 7.19 (dd, *J* = 8.0, 1.8, 1H, CH of Fe), 6.88 (d, *J* = 8.0, 1H, CH of Fe), 6.56 (d, *J* = 16.0, 1H, CH=CHCO2), 5.43 (s, 2H, CH2 of Bn), 3.93 (s, 3H, OMe); 13C NMR (125 MHz, (CD3)2CO, δ) 167.1 (C=O), 164.6 (Cq), 150.4 (Cq), 148.8 (Cq), 147.1 (CH=CHCO2), 140.6 (Cq), 137.4 (Cq), 131.7 (CH of Bn), 127.3 (Cq), 125.2 (2CH of Bn), 124.4 (CH of Fe), 116.1 (CH of Fe), 114.7 (CH=CHCO2),...
111.4 (CH of Fe), 63.0 (CH$_2$ of Bn), 56.4 (OMe); HRESIMS (m/z): [M − H]$^-$ calcd for C$_{17}$H$_{13}$ClNO$_6$, 362.0431; found, 362.0431.

2-Fluoro-5-nitrobenzyl trans-ferulate (6, 48 mg, 0.14 mmol, 46%), white powder. $^1$H NMR (500 MHz, CD$_3$CO$_2$H, δ) 8.43 (dd, $J = 3.2, 2.9, 1$H, CH of Bn), 8.33 (ddd, $J = 9.0, 4.3, 3.0$ Hz, 1H, CH of Bn), 7.67 (d, $J = 16.0, 1$H, CH=CHCO$_2$), 7.49 (t, $J = 9.0, 1$H, CH of Fe), 7.36 (d, $J = 1.8, 1$H, CH of Fe), 7.16 (dd, $J = 8.0, 1.8, 1$H, CH of Fe), 6.87 (d, $J = 8.0, 1$H, CH of Fe), 6.49 (d, $J = 16.0, 1$H, CH=CHCO$_2$), 5.39 (s, 2H, CH$_2$ of Bn), 3.91 (s, 3H, OMe); $^{13}$C NMR (125 MHz, CD$_3$CO$_2$H, δ) 167.1 (C=O), 161.5 (d, $J = 257.4, C_q$), 150.4 (C$_q$), 148.8 (C$_q$), 146.9 (CH=CHCO$_2$), 127.2 (C$_q$), 127.1 (d, $J = 6.1, 1$H, CH of Bn), 126.9 (d, $J = 10.5, 1$H, CH of Bn), 126.7 (d, $J = 17.1, C_q$), 124.3 (CH of Fe), 117.7 (d, $J = 24.3, 1$H, CH of Fe), 116.1 (1H, CH of Fe), 114.7 (CH=CHCO$_2$), 111.4 (CH of Fe), 59.6 (d, $J = 3.9, 1$H, CH of Fe), 56.3 (OMe); HRESIMS (m/z): [M − H]$^-$ calcd for C$_{17}$H$_{13}$FONO$_6$, 346.0727; found, 346.0731.

4-Nitrobenzyl trans-ferulate (7, 96 mg, 0.29 mmol, 97%), white powder. $^1$H NMR (500 MHz, CD$_3$OD, δ) 8.27 (m, 2H, CH of Bn), 7.72 (m, 2H, CH of Bn), 7.69 (d, $J = 16.0, 1$H, CH=CHCO$_2$), 7.35 (d, $J = 2.0, 1$H, CH of Fe), 7.17 (dd, $J = 8.2, 2.0, 1$H, CH of Fe), 6.88 (d, $J = 8.2, 1$H, CH of Fe), 6.51 (d, $J = 16.0, 1$H, CH=CHCO$_2$), 5.38 (s, 2H, CH$_2$ of Bn), 3.92 (s, 3H, OMe); $^{13}$C NMR (125 MHz, CD$_3$OD, δ) 167.1 (C=O), 150.3 (C$_q$), 148.8 (C$_q$), 148.6 (C$_q$) 146.7 (CH=CHCO$_2$), 145.4 (C$_q$), 129.4 (CH of Fe), 127.3 (C$_q$), 124.4 (CH of Fe), 124.2 (CH of Bn), 116.0 (CH of Fe), 115.0 (CH=CHCO$_2$), 111.4 (CH of Fe), 65.1 (CH$_2$ of Bn), 56.4 (OMe); HRESIMS (m/z): [M − H]$^-$ calcd for C$_{17}$H$_{14}$NO$_6$, 328.0821; found, 328.0825.

Synthesis of the feruloylated 4-nitrocatechol-1-yl-substituted butanoylantranol 4NTC–linker–Fe (12)

2-O-(But-3-enyloxy)-5-nitrophenol (10). To a solution of 4-nitrocathecol (4NTC, 1.00 g, 6.44 mmol, 1 equiv) in dry DME (8 mL) were added potassium carbonate (1.06 g, 7.67 mmol, 1.2 equiv) and homoaalicylic bromide (670 µL, 6.51 mmol, 1 equiv) at 40 °C. After overnight stirring at 40 °C, the reaction mixture was concentrated under reduced pressure. The residue was redissolved in ethyl acetate, washed with a saturated aqueous sodium hydrogen carbonate solution, and brine. The combined organic phases were dried over anhydrous sodium sulfate, filtered, and concentrated under reduced pressure. Flash chromatography (gradient of ethyl acetate in petroleum ether from 0 to 50%) afforded 10 (516 mg, 2.47 mmol, 38%) as a solid. $^1$H NMR (500 MHz, CDCl$_3$, δ) 7.82 (dd, $J = 9.0, 3.0, 1$H, CH of 4NTC), 7.80 (d, $J = 3.0, 1$H, CH of 4NTC), 6.90 (d, $J = 9.0, 1$H, CH of 4NTC), 5.91–5.83 (m, 1H, =CH$_2$), 5.23–5.16 (m, 2H, CH$_2$=), 4.21 (t, $J = 6.6, 2$H, CH$_2$O), 2.63 (br

 qt, $J = 6.6, 1.6, 2$H, CH$_2$); $^{13}$C NMR (500 MHz, CDCl$_3$, δ) 151.3 (C$_q$), 145.8 (C$_q$), 142.1(C$_q$), 133.2 (C=CH), 118.1 (CH$_2$=), 116.8 (CH of 4NTC), 110.4 (CH of 4NTC), 110.1 (CH of 4NTC), 68.5 (CH$_2$O), 33.2 (CH$_2$); HRESIMS (m/z): [M − H]$^-$ calcd for C$_{10}$H$_{10}$NO$_4$, 208.0610; found, 208.0615.

(+)−4-O-(2-Hydroxy-4-nitrophenyl)-1,2,4-butanoliter (11). A solution of 10 (157 mg, 0.75 mmol, 1 equiv) in acetonitrile/water 2.5−1 v/v (3.5 mL) was treated at 25 °C under stirring with NaNMO (106 mg, 0.90 mmol, 1.2 equiv) and osmium tetroxide (2.5 wt % solution in tert-butanol, 38 µL) and stirred at room temperature for 18 h. 10% (w/v) aqueous sodium sulfite (0.5 mL) was added, and stirring was continued for 30 min. The product was extracted with ethyl acetate (three times with 10 mL) and washed with brine. The combined organic phases were dried over anhydrous sodium sulfate, filtered, and concentrated under reduced pressure, which afforded the expected compound 11 (165 mg, 0.68 mmol, 90%) as a solid. [tr$_D^{20}$] = 0 (c 2.0, CH$_3$OH). The NMR data (CD$_3$OD) of the racemate 11 were consistent with those previously reported for the pure (S)-enantiomer [37] ([tr$_D^{20}$] = −52 (c 2.0, CH$_3$OH)).

Screening of Fae+ microorganisms in solid medium using X−α-L-Araf–Fe (1a)

The Y. lipolytica AnFaeA(+) strain was used to inoculate solid YNB medium (1.7 g/L YNB without casamino acid, 5 g/L ammonium chloride, 20 mL/L oleic acid, 10 g/L n-glucose, 2 g/L casamino acid, and 15 g/L bacto agar in 100 mM citrate–phosphate buffer at pH 5). The petri dishes were incubated for 48 h at 30 °C and then overlayed with a prepara-
tion of 1% (w/v) molten top agar containing the chogenic substrate 1a (300 µg/mL and 0.5% DMSO) and TsAbf (2 IU/mL). Once the top agar was solid, incubation at 37 °C for 1 h allowed the color to develop. Y. lipolytica strains that contained no AnFaeA gene were also checked to remain colorless after the addition of the chogenic substrate and the auxiliary enzyme.

Liquid-medium-based colorimetric assays using 4NTC–linker–Fe (12)

In a typical experiment, discontinuous enzyme assays were performed in triplicates under buffered conditions (100 mM sodium phosphate at pH 6.0) in the presence of 1.8 mM 12 and 3.6% DMSO, final concentrations. For the assay, this solution was preincubated at 40 °C before AnFaeA addition. Aliquots (25 µL) were stopped by cooling (at 0 °C) every 6 min over a 24 min period and mixed with 45 µL of a cooled 10 mM NaIO4 solution (pH 2.0). After being kept for 5 min at 0 °C throughout, 45 µL of ethylene glycol were added, followed by 135 µL of 2 M Na2CO3 after 5 min. The OD values at 530 nm were recorded on an Infinite M200 PRO (TECAN) microplate reader. One international unit (IU) of Fae specific activity (SA, expressed in µmol/min/mg or IU/mg) corresponds to the amount of released 4NTC (in µmol) per minute per milligram of protein. Negative controls containing all of the reactants except the enzyme were always included in order to monitor and correct for spontaneous hydrolysis of the substrate. Control reactions containing 12, 11 and 4NTC without enzyme, and 4NTC without enzyme and ethylene glycol were also prepared.

Acknowledgements

The NMR work carried out in this work at TBI (Toulouse, France) was performed with the equipment of Meta-Toul (Metabolomics & Fluxomics Facilities, Toulouse, France, http://www.metatoul.fr). MetaToul is part of the national infrastructure MetaboHUB (The French National Infrastructure for Metabolomics and Fluxomics, http://www.metahub.fr) and is supported by grants from the Région Midi-Pyrénées, the European Regional Development Fund, SICOVAL, IBiSA-France, CNRS and INRAE. We thank the ICEO facility dedicated to enzyme screening and discovery and part of the Integrated Screening Platform of Toulouse (PICT, IBiSA) for providing access to their equipment. The authors wish to acknowledge the support from the ICMG Chemistry Nanobio Platform (Grenoble, France) for HRMS analyses. Moreover, the authors acknowledge that this work is based on a manuscript that is included in the PhD thesis by J. D. titled “Approches multiples d’ingénierie pour l’utilisation d’enzymes hydrolytiques comme outils de synthèse”. INSA de Toulouse, France, 2017 (https://tel.archives-ouvertes.fr/tel-02011218).

Funding

This work was supported by the European Union’s Seventh Programme for Research, Technological Development and Demonstration under Grant Agreement No 613868, the OPTIBIOCAT project (to O. G. and J. D.) by the French National Research Agency, grant ANR-05-PNRB-002, the project SPPECABBE (to M. R.), and the Région Midi-Pyrénées grant DAER-Recherche 07009817 (to F. F.).

ORCID® iDs

Mélanie Ragon - https://orcid.org/0000-0002-1367-4764
Julien Durand - https://orcid.org/0000-0002-5631-6210
Sophie Bozonnet - https://orcid.org/0000-0001-5091-2209
Michael J. O’Donohue - https://orcid.org/0000-0003-4246-3938
Régis Fauré - https://orcid.org/0000-0002-5107-9009

Preprint

A non-peer-reviewed version of this article has been previously published as a preprint: https://doi.org/10.3782/bxiv.2020.107.v1

References

1. Lombard, V.; Golaconda Ramulu, H.; Drula, E.; Coutinho, P. M.; Henrissat, B. Nucleic Acids Res. 2014, 42, D490–D495. doi:10.1093/nar/gkt1178
2. Faulds, C. B. Phytochem. Rev. 2010, 9, 121–132. doi:10.1007/s11101-009-9156-2
3. Dumon, C.; Song, L.; Bozonnets, S.; Fauré, R.; O’Donohue, M. J. Process Biochem. (Oxford, U. K.) 2012, 47, 346–357. doi:10.1016/j.procbio.2011.06.017
4. Razzagh-Asli, N.; Garrido, J.; Khazraei, H.; Borges, F.; Fruizi, O. Curr. Med. Chem. 2013, 20, 4436–4450. doi:10.2174/0929867313209990141
5. Oliveira, D. M.; Mota, T. R.; Oliva, B.; Segato, F.; Marchiosi, R.; Ferrarese-Filho, O.; Faulds, C. B.; dos Santos, W. D. Bioprocess. Technol. 2019, 278, 408–423. doi:10.1016/j.biortech.2019.01.064
6. Prates, J. A. M.; Tarbouriech, N.; Charnock, S. J.; Fontes, C. M. G. A.; Ferreira, L. M. A.; Davies, G. J. Structure 2001, 9, 1183–1190. doi:10.1016/s0969-2126(01)00684-0
7. Wong, D. W. S. Appl. Biochem. Biotechnol. 2006, 133, 87–112. doi:10.1385/abab:133:2:87
8. Dilokpimol, A.; Mäkelä, M. R.; Aguilar-Pontes, M. V.; Benoit-Gelber, I.; Hildén, K. S.; de Vries, R. P. Biotechnol. Biofuels 2016, 9, 231. doi:10.1186/s13068-016-0651-6
9. Ramos-de-la-Peña, A. M.; Contreras-Esquível, J. C. J. Mol. Catal. B: Enzym. 2016, 130, 74–87. doi:10.1016/j.molcatb.2016.05.004
10. Ramírez-Velasco, L.; Armentdín-Ruíz, M.; Rodríguez-González, J. A.; Müller-Santos, M.; Asaff-Torres, A.; Mateos-Díaz, J. C. Comb. Chem. High Throughput Screening 2016, 19, 616–626. doi:10.2174/1388207196661110154722
11. Hegde, S.; Srinivas, P.; Muralikrishna, G. Anal. Biochem. 2009, 387, 128–129. doi:10.1016/j.ab.2009.01.001
12. Zhang, S.-B.; Ma, X.-F.; Pei, X.-Q.; Liu, J.-Y.; Shao, H.-W.; Wu, Z.-L. J. Mol. Catal. B: Enzym. 2012, 74, 36–40. doi:10.1016/j.molcatb.2011.08.011
13. Gherbovet, O.; Fauré, R.; Ferreira, F.; Durand, J.; Ragon, M.; Hostyn, G.; Record, E.; Bozonnet, S.; O’Donohue, M. J. Mol. Catal. B: Enzym. 2016, 126, 24–31. doi:10.1016/j.molcatb.2016.01.012

14. Mastihubová, M.; Mastihuba, V.; Kremnicky, L.; Willet, J. L.; Côté, G. L. Synlett 2001, 1559–1600. doi:10.1055/s-2001-17450

15. Marmuse, L.; Asther, M.; Fabre, E.; Navarro, D.; Lesage-Meesen, L.; Asther, M.; O’Donohue, M.; Fort, S.; Driguez, H. Org. Biomol. Chem. 2008, 6, 1208–1214. doi:10.1039/b717742a

16. Mastihubová, M.; Biely, P. Carbohydr. Res. 2010, 345, 1094–1098. doi:10.1016/j.carres.2010.03.034

17. Mastihubová, M.; Szemesová, J.; Biely, P. Tetrahedron Lett. 2003, 44, 1671–1673. doi:10.1016/s0040-4039(03)00338-8

18. Crepin, V. F.; Faulds, C. B.; Connerton, I. F. Appl. Microbiol. Biotechnol. 2004, 63, 647–652. doi:10.1007/s00253-003-0476-3

19. Underlin, E. N.; Frommhagen, M.; Dilopkimol, A.; van Erven, G.; de Vries, R. P.; Kabel, M. A. Front. Bioeng. Biotechnol. 2020, 8, 332. doi:10.3389/fbioe.2020.00332

20. Hunt, C. J.; Antonopolou, I.; Tankaslæ, A.; Rova, U.; Christakopoulos, P.; Haritos, V. S. Sci. Rep. 2017, 7, 17315. doi:10.1038/s41598-017-17260-x

21. Biely, P.; Mastihubová, M.; van Zijl, W. H.; Prior, B. A. Anal. Biochem. 2002, 311, 68–75. doi:10.1006/abio.2001.4001-3

22. Schmidt, M.; Bornscheuer, U. T. Biometal. Eng. 2005, 22, 51–56. doi:10.1016/j.bioeng.2004.09.004

23. Marmuse, L.; Asther, M.; Navarro, D.; Lesage-Meesen, L.; Asther, M.; Fort, S.; Driguez, H. Carbohydr. Res. 2007, 342, 2316–2321. doi:10.1016/j.carres.2007.06.004

24. Armesto, N.; Ferrero, M.; Fernández, S.; Gotor, V. J. Org. Chem. 2003, 68, 5784–5787. doi:10.1021/jo034387a

25. Mastihubová, M.; Mastihuba, V. Biorg. Med. Chem. Lett. 2013, 23, 5389–5392. doi:10.1016/j.bmcl.2013.07.051

26. Wang, Y.; Zhang, D.-H.; Chen, N.; Zhi, G.-Y. Helv. Chim. Acta 2014, 97, 1656–1670. doi:10.1002/hlca.201300690

27. Chyba, A.; Mastihubová, V.; Mastihuba, M. Biorg. Med. Chem. Lett. 2016, 26, 1567–1570. doi:10.1016/j.biortech.2016.02.010

28. Chyba, A.; Mastihubová, M.; Mastihuba, V. Monatsh. Chem. 2016, 147, 67–79. doi:10.1007/s00706-016-1698-6

29. Mastihubová, M.; Szemesová, J.; Biely, P. Biorg. Med. Chem. Lett. 2006, 16, 1805–1810. doi:10.1016/j.bmcl.2005.10.023

30. Bordes, F.; Fudalé, F.; Dossat, V.; Nicaud, J.-M.; Marty, A. J. Microbiol. Methods 2007, 70, 493–502. doi:10.1016/j.mimet.2007.06.008

31. Otto, R. T.; Scheib, H.; Bornscheuer, U. T.; Pleiss, J.; Syldatk, C.; Schmid, R. D. J. Mol. Catal. B: Enzym. 2000, 8, 201–211. doi:10.1016/s1381-1177(99)00589-2

32. Mastihubová, M.; Mastihuba, V.; Bilaničová, D.; Borekiová, M. J. Mol. Catal. B: Enzym. 2006, 38, 54–57. doi:10.1016/j.molcatb.2005.11.003

33. Zeuner, B.; Kontogeorgis, G. M.; Rissager, A.; Meyer, A. S. New Biotechnol. 2012, 29, 255–270. doi:10.1016/j.nbt.2011.11.011

34. Parmar, V. S.; Prasad, A. K.; Pati, H. N.; Kumar, R.; Azim, A.; Roy, S.; Errington, W. Biocatal. Bioengineering 2019, 2009, 119–134. doi:10.1002/bioe.20191117

35. Malhotra, S.; Calderón, M.; Prasad, A. K.; Parmar, V. S.; Haag, R. Org. Biomol. Chem. 2010, 8, 2228–2237. doi:10.1039/b927021c

36. Wang, J.; Gu, S.-S.; Cui, H.-S.; Wu, X.-Y.; Wu, F.-A. Biocatal. Bioengineering 2014, 158, 39–47. doi:10.1039/b1071177v

License and Terms

This is an Open Access article under the terms of the Creative Commons Attribution License (https://creativecommons.org/licenses/by/4.0/). Please note that the reuse, redistribution and reproduction in particular requires that the author(s) and source are credited and that individual graphics may be subject to special legal provisions.

The license is subject to the Beilstein Journal of Organic Chemistry terms and conditions: (https://www.beilstein-journals.org/bjoc/terms)

The definitive version of this article is the electronic one which can be found at: https://doi.org/10.3762/bjoc.17.30