Supporting Information

One-pot biocatalytic synthesis of *rac*-syringaresinol from a lignin-derived phenol

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Abbreviations: BPA, bisphenol A; EUGO, eugenol oxidase; HRP, horseradish peroxidase, DHSA, dihydrosinapyl alcohol, DHCA, dihydroconiferyl alcohol.

 Rosetta
 V166A, V166T, Y168A, V276T, M282G, M282S, L381I, L381Q, I391A, I391V, G392A, S394A, S394V, A423L, Q425L, I427V

 Rational
 G165S, L381N, A423I, Q425E, I427A, H434D, H434L, L438D, L438E, I427G

	EUGO8X	EUGO10X	
Space group	P1	P 2 ₁ 2 ₁ 2 ₁	
Unit cell axes (Å)	113.56,142.99, 154.60	57.65, 98.05, 177.34	
Unit cell angles (°)	114.87, 97.00, 93.28	90, 90, 90	
Resolution (Å)	2.3	1.65	
PDB code	8BAP	8BAM	
R _{sym} ^{<i>a,b</i>}	0.1592 (0.8421)	0.09249 (1.028)	
CC _{1/2}	0.988 (0.547)	0.998 (0.674)	
Completeness ^b (%)	98.21 (97.67)	99.41 (95.48)	
Unique reflections	379400	120212	
Redundancy	3.5 (3.6)	6.4 (6.3)	
I/σ^b	7.2 (1.6)	13.8 (1.7)	
N° of non-hydrogen atoms protein/FAD ligand water Average B value (Å ²)	68547/16x53 16x15 1287 34.51	9917/2x53 2x15 703 23.43	
R _{crys} ^c (%)	19.3	17.2	
R _{free} ^c (%)	24.9	20.6	
Rms bond length (Å)	0.009	0.007	
Rms bond angles (°)	1.02	0.94	

 Table S2. Data collection and refinement statistics for EUGO8X and EUGO10X.

^{*a*} $R_{sym}=\sum |I_i-<I>|/\sum I_i$, where I_i is the intensity of ith observation and <I> is the mean intensity of the reflection.

^b Values in parentheses are for reflections in the highest resolution shell.

^c $R_{cryst}=\sum |F_{obs}-F_{calc}|/\sum |F_{obs}|$ where F_{obs} and F_{calc} are the observed and calculated structure factor amplitudes, respectively. R_{cryst} and R_{free} were calculated using the working and test sets, respectively.



Figure S1. Mechanistic proposal for the oxidation of dihydrosinapyl alcohol by EUGO. Flavin acts as a hydride acceptor for dihydrosinapyl alcohol, generating a reactive quinone methide intermediate. The intermediate is quenched either by deprotonation of the adjacent carbon to form an alkene (blue arrows), or base-catalysed hydroxylation by a water molecule (pink arrows).



Figure S2. HPLC analysis of relevant compounds. Dihydrosinapyl alcohol, sinapyl alcohol, and syringaresinol are shown in blue, green and red, respectively.



Figure S3. HPLC calibration curves of dihydroconiferyl alcohol, coniferyl alcohol, dihydrosinapyl alcohol and syringaresinol.



Figure S4. Temperature effect on stability of dihydrosinapyl alcohol. Dihydroconiferyl alcohol (5 mM) was incubated in KPi buffer (50 mM, pH 7.5) containing DMSO (10% v/v) at the indicated temperatures, and the mixture was assayed by HPLC.



Figure S5. Optimization of the one-pot conversion of dihydrosinapyl alcohol to syringaresinol by an oxidase-peroxidase cascade reaction. A) Effect of additional hydrogen peroxide on the formation of syringaresinol from dihydrosinapyl alcohol by an oxidase-HRP cascade reaction. Using dihydrosinapyl alcohol (5 mM), oxidase (10 μ M), and HRP (10 μ M), the reactions with a gradient addition of hydrogen peroxide were performed in KPi buffer (50 mM, pH 7.5) at 37 °C, 150 rpm. B) Effects of delayed addition of HRP and hydrogen peroxide on the formation of syringaresinol by an oxidase-HRP cascade reaction. The reactions were initialized with dihydrosinapyl alcohol (5 mM) and oxidase (10 μ M), and then HRP with a gradient concentration and HRP (10 μ M) combined with hydrogen peroxide at 5 mM and 10 mM were added after 3 hours.



Figure S6. Analysis of product mixture isolated from 20 mg conversion of dihydrosinapyl alcohol using EUGO10X. A) ¹H NMR spectrum of product mixture of sinapyl alcohol and dihydrosinapyl alcohol. B) GC-EI-MS mass spectrum of the sinapyl alcohol reaction product. The NMR peaks formed which are distinct from those of dihydrosinapyl alcohol align well with literature values for sinapyl alcohol.¹



Figure S7. Analytical data of syringaresinol product isolated from 20 mg conversion of dihydrosinapyl alcohol using EUGO10X and HRP. A) ¹H NMR spectrum. B) ¹³C NMR spectrum. C) GC-EI-MS mass spectrum of the syringaresinol product.

¹**H NMR** (400 MHz, CDCl₃) δ_{H} 6.51 (s, 4H, aromatic protons), 5.45 (s, 2H, -OH), 4.67 (d, J = 3.9 Hz, 2H, H-1), 4.22 (dd, 2H, J = 7.4, 6.5 Hz, H-4), 3.86 – 3.80 (m, 14H, H-4 & -OMe), 3.08 – 3.00 (m, 2H, H-2).

 $^{13}\textbf{C}$ NMR (101 MHz, CDCl_3) δ_{C} 147.2, 134.3, 132.1, 102.7, 86.1, 71.8, 56.4, 54.3.



Figure S8. HPLC analysis of the conversion of dihydrosinapyl alcohol by EUGO variants at different stages of engineering. In all chromatograms, peaks 1, 2, 3 and 4 hydroxylated DHSA (Rt = 2.7 min), DHSA ketone (Rt = 3.2 min), DHSA (Rt = 4.0 min) and sinapyl alcohol (Rt = 4.3 min), respectively. The first chromatogram shows the elution of the reference substrate dihydrosinapyl alcohol. The chromatograms of the conversions show depletion of dihydrosinapyl alcohol to form sinapyl alcohol and by-products.





Protein Sequences

<u>EUGO</u>

MTRTLPPGVSDERFDAALQRFRDVVGDKWVLSTADELEAFRDPYPVGAAEANLPSAVVSP ESTEQVQDIVRIANEYGIPLSPVSTGKNNGYGGAAPRLSGSVIVKTGERMNRILEVNEKY GYALLEPGVTYFDLYEYLQSHDSGLMLDCPDLGWGSVVGNTLDRGVGYTPYGDHFMWQTG LEVVLPQGEVMRTGMGALPGSDAWQLFPYGFGPFPDGMFTQSNLGIVTKMGIALMQRPPA SQSFLITFDKEEDLEQIVDIMLPLRINMAPLQNVPVLRNIFMDAAAVSKRTEWFDGDGPM PAEAIERMKKDLDLGFWNFYGTLYGPPPLIEMYYGMIKEAFGKIPGARFFTHEERDDRGG HVLQDRHKINNGIPSLDELQLLDWVPNGGHIGFSPVSAPDGREAMKQFEMVRNRANEYNK DYAAQFIIGLREMHHVCLFIYDTAIPEAREEILQMTKVLVREAAEAGYGEYRTHNALMDD VMATFNWGDGALLKFHEKIKDALDPNGIIAPGKSGIWSQRFRGQNL

<u>EUGO5X</u>

MTRTLPPGVSDERFDAALQRFRDVVGDKWVLSTADELEAFRDPYPVGAAEANLPSAVVSP ESTEQVQDIVRIANEYGIPLHPVSTGKNNGYGGAAPRLSGSVIVKTGERMNRILEVNEKY GYALLEPGVTYFDLYEYLQSHDSGLMLDCPDLGWGSVVGNTLDRGVGYTPYGDHFMWQTG LEVVLPQGEVMRTGMGALPGSDAWQLFPYGFGPFPDGMFTQSNLGIVTKMGIALMQRPPA SQSFLITFDKEEDLEQIVDIMLPLRINMAPLQNVPVLRNIFMDAAAVSKRTEWFDGDGPM PAEAIERMKKDLDLGFWNFYGTLYGPPPLIEMYYGMIKEAFGKIPGARFFTHEERDDRGG HVLQDRHKINNGIPSLDELQLLDWVPNGGHIGFSPVSAPDGREAMKQFEMVRNRANEYNK DYMAQFIIGLREMYHVCLFIYDTADPEAREEILQMTKVLVREAAEAGYGEYRTHNALMDD VMATFNWGDGALLKFHEKIKDALDPNGIIAPGKSGIWPQRFRGQNL

EUGO8X

MTRTLPPGVSDERFDAALQRFRDVVGDKWVLSTADELEAFRDPYPVGAAEANLPSAVVSP ESTEQVQDIVRIANEYGIPLHPVSTGKNNGYGGAAPRLSGSVIVKTGERMNRILEVNEKY GYALLEPGVTYFDLYEYLQSHDSGLMLDCPDLGWGSVVGNTLDRGVGYTPYGDHFMWQTG LEVVLPQGEVMRTGMGALPGSDAWQLFPYGFGPFPDGMFTQSNLGIVTKMGIALMQRPPA SQSFLITFDKEEDLEQIVDIMLPLRINMAPLQNVPVLRNIFMDAAAVSKRTEWFDGDGPM PAEAIERMKKDLDLGFWNFYGTLYGPPPLIEMYYGMIKEAFGKIPGARFFTHEERDDRGG HVLQDRHKINNGIPSLDELQQLDWVPNGGHIGFVPVSAPDGREAMKQFEMVRNRANEYNK DYMAQFVIGLREMYHVCLFIYDTADPEAREEILQMTKVLVREAAEAGYGEYRTHNALMDD VMATFNWGDGALLKFHEKIKDALDPNGIIAPGKSGIWPQRFRGQNL

<u>EUGO10X</u>

MTRTLPPGVSDERFDAALQRFRDVVGDKWVLSTADELEAFRDPYPVGAAEANLPSAVVSP ESTEQVQDIVRIANEYGIPLHPVSTGKNNGYGGAAPRLSGSVIVKTGERMNRILEVNEKY GYALLEPGVTYFDLYEYLQSHDSGLMLDCPELGWGSVVGNTLDRGVGYTPYGDHFMWQTG LEVVLPQGEVMRTGMGALPGSDAWQLFPYGFGPFPDGMFTQSNLGIVTKMGIALMQRPPA SQSFLITFDKEEDLEQIVDIMLPLRINMAPLQNVPVLRNIFMDAAAVSKRTEWFDGDGPM PAEAIERMKKDLDLGFWNFYGTLYGPPPLIEMYYGMIKEAFGKIPGARFFTHEERDDRGG HVLQDRHKINNGIPSLDELQQLDWVPNGGHIGFVPVSAPDGREAMKQFEMVRNRANEYNK DYMASFVIGLREMYHVCLFIYDTADPEAREEILQMTKVLVREAAEAGYGEYRTHNALMDD VMATFNWGDGALLKFHEKIKDALDPNGIIAPGKSGIWPQRFRGQNL

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