



Published in final edited form as:

J Am Chem Soc. 2019 September 11; 141(36): 14052–14056. doi:10.1021/jacs.9b06592.

Enzymatic Intermolecular Hetero-Diels–Alder Reaction in the Biosynthesis of Tropolonic Sesquiterpenes

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Abstract

Diels–Alder reactions are among the most powerful synthetic transformations to construct complex natural products. Despite that increasing of enzymatic intramolecular Diels–Alder reactions have been discovered, natural intermolecular Diels–Alderase are rarely described. Here, we report an intermolecular hetero-Diels–Alder reaction in the biosynthesis of tropolonic sesquiterpenes and functionally characterize EupfF as the first fungal intermolecular hetero-Diels–Alderase. We demonstrate that EupfF catalyzed the dehydration of a hydroxymethyl-containing tropolone (5) to generate a reactive tropolone *o*-quinone methide (6) and might further stereoselectively control the subsequent intermolecular hetero-Diels–Alder reaction with (1*E*,4*E*,8*Z*)-humulenol (8) to produce enantiomerically pure neosetophomone B (1). Our results reveal the biosynthetic pathway of 1 and expand the repertoire of activities of Diels–Alder cyclases.

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Supporting Information

The Supporting Information is available free of charge on the ACS Publications website at DOI: 10.1021/jacs.9b06592.

Experimental details, spectroscopic and computational data (PDF)

The authors declare no competing financial interest.

Diels–Alder (DA) reactions, in which a 1,3-diene and a dienophile react to form an unsaturated cyclohexane regio- and stereoselectively are one of the most powerful synthetic strategies for the synthesis of complex natural products.¹ Both intramolecular and intermolecular Diels–Alder cycloadditions are presumed to introduce structure complexity and biological activity in natural compounds. Although increasing numbers of intramolecular Diels–Alderase or [4 + 2]-cyclases² such as LepI,³ SpnF,⁴ PyrE3,⁵ IccD,⁶ PvhB,⁷ and others,⁸ have been discovered in Nature, direct biochemical evidence of enzyme-catalyzed intermolecular DA reactions have rarely been described except for macrophomate synthase and riboflavin synthase.² Notably, natural products probably biosynthesized via an intermolecular Diels–Alder reaction are frequently reported from bacteria, fungi, and plants⁹ (Figure S1).

Remarkably, several members of the tropolone-sesquiterpene family of meroterpenoids (Figure S2) represented by neosetophomone B (1),¹⁰ epolone B (2),¹¹ eupenifeldin (3),¹² and pycnidione,¹³ isolated as optically pure compounds from a variety of fungi, were proposed to be biosynthesized via one or two tandem hetero Diels–Alder (hDA) reactions (Figure 1).¹⁴ Structurally, these meroterpenoids possess a unique central 11-membered macrocycle flanked by one or two dihydropyrans coupled to tropolones. Biologically, 1 and 3 were found as very potent antitumor agents with activity at the nanomole level in *in vitro* tumor cell lines,^{10,12} while 2 and pycnidione are stromelysin inhibitors and potential antiarthritic drugs.¹³

Studies toward the biomimetic synthesis of tropolone-sesquiterpenes showed the generation of monotropolone adducts via hDA reaction under thermal conditions achieved success only for unnatural model compounds.¹⁵ In fact, the total synthesis of these tropolone-sesquiterpenes has not been completed thus far. It is therefore of interest to identify strategies and enzymes Nature uses to synthesize compounds of this family, especially the structures with observed in 1.

Biosynthetically, hDA reactions of a tropolone *o*-quinone methide with humulene were proposed for the biosynthesis of 1 and 2, which further served as an intermediate for the biosynthesis of eupenifeldin and pycnidione, respectively^{11,16} (Figure 2a). While biosynthesis of the tropolone core in stipitonic acid has been identified (Figure 2a), including a nonreduced polyketide synthase (NR-PKS, TropA), a FAD-dependent monooxygenase (FMO, TropB), and an α -ketoglutarate-dependent dioxygenase (α -KGD, TropC),¹⁷ the enzymatic basis for the DA reaction for 1 biosynthesis has remained unexplored. Here, we report mapping of the cascade of enzymatic steps that furnish the structure of 1, which include a cytochrome P450 monooxygenase (P450), a short-chain dehydrogenase (SDR), and an intermolecular hetero-Diels–Alderase (hDAase) as keys. We demonstrate that the hDAase is responsible for the formation of a highly reactive tropolone *o*-quinone methide and may further stereoselectively control the subsequent intermolecular hDA cycloaddition to produce 1.

Recently, we coisolated 1 and 3 from the wild-type *Penicillium janthinellum*, a producer of alanditrypinone.¹⁸ Analysis of the crude extract by liquid chromatography mass spectrometry (LC–MS) led to the observation of minor 1 and major 3 with characteristic UV

absorption of tropolones and molecular weights of 384 and 548, respectively (Figure S3). Compounds 1 and 3 were fully characterized by NMR and circular dichroism spectra as enantiomerically pure neosetophomone B and eupenifeldin, respectively. To investigate the biosynthetic pathway of 1, the genome of *P. janthinellum* was queried for the presence of the trop-like gene cluster. Bioinformatics analysis led to the identification of only one *trop*-like NR-PKS gene cluster (named *eupf* cluster) in its genome (Figures 2b and S4). In addition to three enzymes (EupfA–C) which are homologues to TropA, TropB, and TropC, respectively, the *eupf* cluster encodes additionally a cytochrome P450 monooxygenase (EupfD), a short-chain dehydrogenase (EupfE), two hypothetic enzymes (EupfF and EupfG), as well as an ABC transporter and regulators (Figure 2b). EupfF and EupfG show high homology to AsR5 and AsR6 (64% and 62%), respectively, which were recently suggested as putative hetero-Diels–Alderase and humulene synthase, respectively, in the biosynthesis of xenovulene A.¹⁹ A biosynthetic gene cluster *eup* in *Phoma* sp. has been recently identified by gene deletion for the biosynthesis of eupenifeldin and EupF was also suggested as putative hDAase.²¹ However, the function of AsR5 and EupF has not been successfully characterized. It was scrutable that EupfABC were responsible for the formation of stipitaldehyde, and EupfG served as a humulene synthase to form a sesquiterpene, while the only oxygenase EupfD could catalyze the hydroxylation at the terpene moiety at an uncertain stage. The putative hDAase EupfF, homologous to AsR5 and EupF, was proposed to be involved in the cyclization between diene and dienophile.

To verify the *eupf* gene cluster and elucidate the biosynthetic pathway of 1, heterologous expression experiments in *Aspergillus nidulans* were conducted. As expected, expression of *eupfABC* in *A. nidulans* led to the production of major stipitaldehyde (4, Figure 3a), which was suggested as an intermediate for biosynthesis of xenovulene A and other tropolone-sesquiterpenes. To form a diene precursor for the DA reaction, the aldehyde moiety in 4 has to be reduced first. As the only potential reductase in the *eupf* cluster, *EupfE* is highly likely to be involved in this transformation. To verify the function of EupfE, intron-free *eupfE* was cloned from cDNA, heterologously expressed in *Escherichia coli*, and purified to homogeneity (Figure S6). Compared to control experiments where missing EupfE did not generate new products, incubation of EupfE with 4 and NADPH led to complete conversion to a new product 5 (Figure 3b) with characteristic UV absorption of tropolone and MW of 182, 2 Da more than that of 4 (Figure S7). Although 5 was not stable enough, careful isolation and quick NMR analysis led to the full identification of 5 as we expected (Figure 3d). Dehydration of 5 either spontaneously or enzymatically was proposed to generate the reactive tropolone *o*-quinone methide (6).

To identify the dienophile precursor, EupfG, the putative humulene synthase was expressed in *A. nidulans*, which led to the detection of a compound 7 with the same molecular weight (*m/z* 204) but different retention time from that of α -humulene (9) standard indicated by GC–MS analysis (Figure S8), which suggested 7 was an isomer of α -humulene. Isolation of 7 was found to be difficult due to its high volatility. The structure of 7 was suggested by coexpression of *eupfG* and *eupfD* in *A. nidulans*, which led to the production of 8 (Figure 3c, trace (i)) with MW of 220, 16 Da more than that of α -humulene. Compound 8 was fully characterized by MS and NMR spectra as a hydroxylated product of 7 (Figure 3d), which also supported that 7 was (1*E*,4*E*,8*Z*)-humulene. Different from α -humulene in which all

three olefins are in the *E* form, compounds 7 and 8 possess a *Z* form of the olefinic bond at ^{8,9}, which is consistent with the *Z* form of alkene in 1. As a result, EupfG was identified as a new sesquiterpene synthase producing (1*E*,4*E*,8*Z*)-humulene other than α -humulene.

We realized that 5 could be *O*-methylated at C₉-OH gradually and partly in methanol (Figure S9). Methylated 5 was isolated and fully characterized by NMR spectra as a tautomer mixture (2:1). The identification of methylated 5 (6a) also suggested the spontaneous conversion from 5 to 6 occurred before it underwent 1,4- or 1,6-addition by methanol (Figure S9). Due to the incomplete spontaneous conversion, we believed that enzyme was required to generate 6 from 5, and EupfF, the putative hetero-Diels–Alderase, was presumed to be involved in this process. However, repeated attempts to obtain soluble EupfF for biochemical assay failed. Fortunately, an enzyme, EupF, highly homologous to EupfF (81% similarity/67% identity), was found in the genome of *Phoma* sp., which was suggested but not characterized as putative hDAase for the biosynthesis of eupenifeldin.²⁰ Attempts to obtain soluble EupF (42.0 kDa) from *Escherichia coli* BL21(DE3) were successful (Figure S10). To explore if EupF catalyzed the dehydration of 5, we performed *in vitro* EupF assay with 5 as the sole substrate. Since the expected product 6 was extremely instable, we added 1% glycerol in the *in vitro* system to trap 6. Consequently, 72.5% of 5 converted to two main products, 6b and 6c, with molecular weights of 256 and 346, respectively (Figure S11), by EupF after 2 h, while only 21.8% of 5 could be transformed without EupF. On the basis of the MS and UV, 6b and 6c were suggested to be 1,4-nucleophilic addition products of 6 with glycerol and 5, respectively. The results from the *in vitro* assay supported that dehydration of 5 could occur spontaneously, but EupF remarkably accelerated this process. Hence, the diene precursor 6 could be identified as tropolone *o*-quinone methide.

With 5 and 8 in hand, we were able to examine whether EupF was directly involved in the hDA reaction. We performed an *in vitro* reaction containing 30 μ M EupF, 4.0 mM 5, and 1.0 mM 8 in 50 mM bicine buffer (pH 8.0). Whereas the control reaction without EupF did not lead to apparent new products, EupF catalyzed the conversion of 5 and 8 into 1 (Figures 4a and S12). Compound 1's functionality as an enantiopure compound indicated the hDA reaction was enzymatically controlled since racemic products would be expected from the spontaneous reaction. Interestingly, when the natural substrate 8 was switched to unnatural substrate humulenol (10), which was obtained from expression of *eupf D* in *A. nidulans* by feeding humulene (9), in the *in vitro* EupF assay, two products 2 and 11 (1:1 ratio) were detected (Figure 4a, trace v). To obtain enough 2 and 11 for structural elucidation, 9 was fed to *A. nidulans* coexpressing *eupf ABCDEF* (Figure S12). Finally, compounds 2 and 11 were fully characterized by NMR and CD as natural epolone B and its isomer, named isoeopolone B (Figure 4c), respectively. Trace amounts of 2 and 11 were also detectable (Figure 4a, trace vi) in the absence of EupF with low yield compared to the enzymatic reaction. Hence, we suggested (1) the cycloaddition between the reactive tropolone *o*-quinone methide 6 and dienophile (8 or 10) could occur spontaneously, (2) EupF directly controlled the stereoselectivity of hDA reaction in the case of natural substrate 8, and (3) unnatural substrate 10 was not recognized by EupF but underwent spontaneous cycloaddition with 6. Thus, EupF was fully characterized as the first hDAase, which differs from all intramolecular DAases in phylogenetic analysis (Figure S13) and has no recognizable cofactor-binding site. A model of EupfF (EupF) computed with SWISS-MODEL and template 2p4o suggest a predicted

six-bladed propeller (Figure S14) that may provide suitable space for substrates. Although His, Lys, and Arg have been found as potential active sites for dehydration and DA reaction, ²¹ mutagenesis experiments showed that conserved H37, R51, R92, and R323 were not catalytic residues for EupF (Figures S15–16), which suggested EupF was different from other DAases.

To support the proposed pathway, we performed density functional theory (DFT) studies on the Diels–Alder reaction at the PBE0-D3(BJ)/def2-TZVP level of theory.^{22–27} These calculations indicate that this cycloaddition is very facile (Figures 4b and S17). Transition states TS-1 and TS-2 lead to product 1 and are highly asynchronous with the two forming bonds differing by 0.49 and 0.45 Å, respectively. There is a 0.25 electron charge transfer in the transition state, in TS-1 the heterodiene has a –0.25 charge, and dienophile has a +0.25 charge. The charges are similar in TS-2. It is probable that the rate-determining step is generation of the *o*-quinone methide 6. After formation of 6, the reaction barriers of TS-1 and TS-2 are purely entropic (TS-1 $H^\ddagger = -1.2$, $G^\ddagger = +13.8$ kcal·mol⁻¹, TS-2 $H^\ddagger = -0.6$, $G^\ddagger = +15.1$ kcal·mol⁻¹). These reactions are enthalpically barrierless because the heterodiene is highly electron-deficient and a highly stable aromatic product is formed.

Unexpectedly, neither *in vitro* assay with 5 and 8 as substrates nor feeding experiments (Figure S18) led to production of 3 (only 1 was detected), which highly indicated that other or additional enzymes were required for the right side DA reaction in the biosynthesis of 3. Further investigation on the biosynthesis of bistropolone-sesquiterpenes is in progress.

In conclusion, we biochemically characterized EupfF (EupF) as the first fungal intermolecular hetero-Diels–Alderase and demonstrated that EupfF catalyzed the generation of a reactive tropolone *o*-quinone methide and further stereoselectively controlled the subsequent intermolecular hetero-Diels–Alder reaction to produce 1. Our discovery expands the collection of increasing DAases from fungi and opens a new strategy for mining and combinational biosynthesis of tropolone-sesquiterpenes.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

ACKNOWLEDGMENTS

This work was supported financially by the CAMS Innovation Fund for Medical Sciences (CIFMS, No. 2017-I2M-4-004 to Y.H., and 2018-I2M-3-005 to Y.C.) and the Drug Innovation Major Project (2018ZX09711001-006 to Y.H.) as well as R01GM 124480 and NSF (CHE-1806581) to K.N.H. The UCLA Institute of Digital Research and Education (IDRE) provided computational resources for all calculations. We are grateful to Dr. Minghua Chen for ECD calculations.

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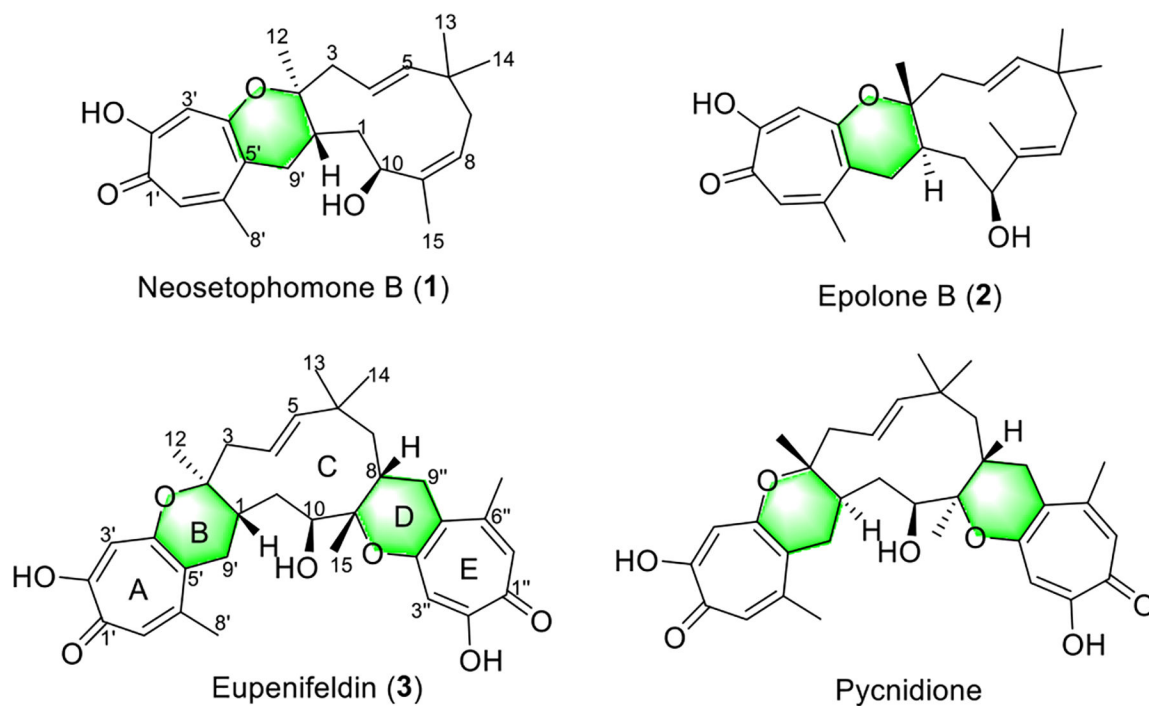


Figure 1.
Representative tropolone-sesquiterpenes probably generated from intermolecular Diels-Alder reactions.

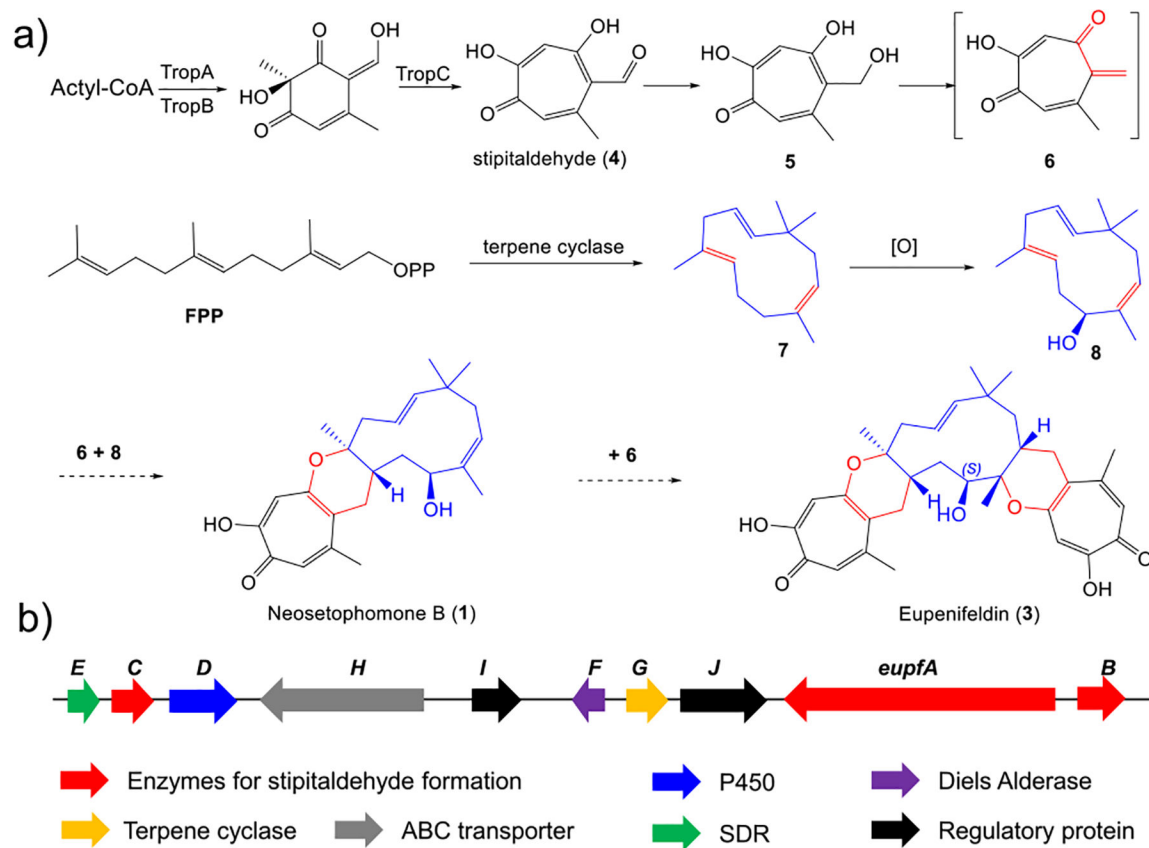


Figure 2.
Proposed biosynthetic pathways of 1 and 3 (a) and the *eupf* gene cluster in *P. janthinellum* (b).

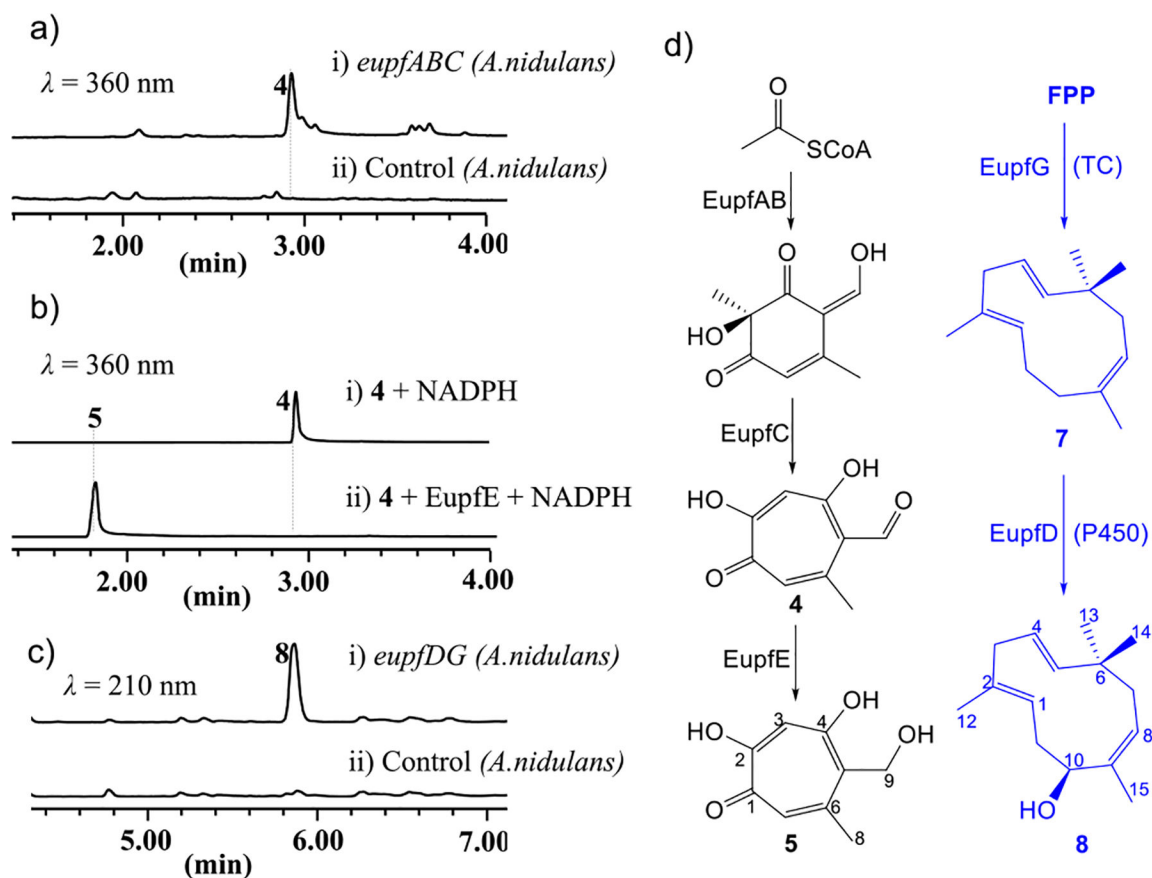


Figure 3. Generation of potential precursors in the biosynthesis of 1. (a–c) LCMS traces from heterologous expression of *eupfABC/DG* in *A. nidulans* or from *in vitro* assay of EupfE. (d) Generation of 5 and 8.

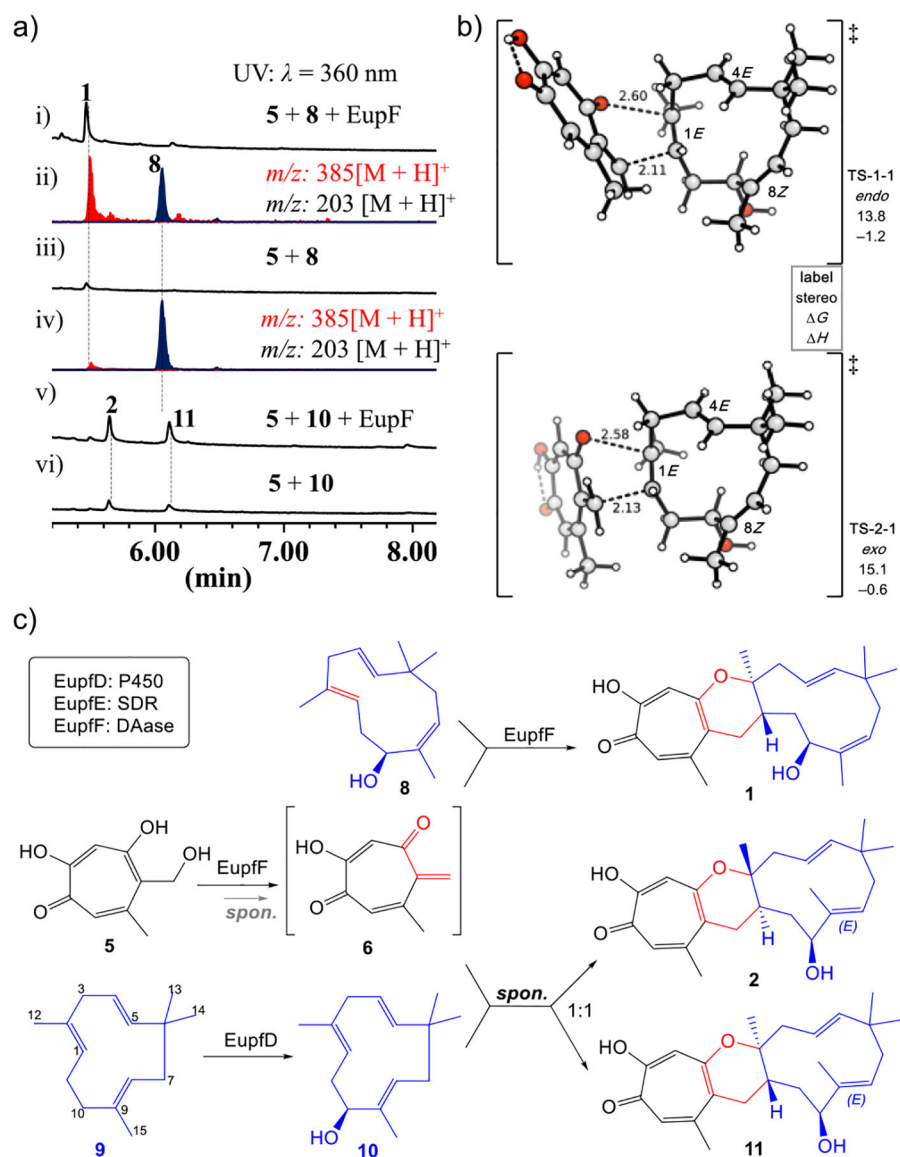


Figure 4. Biosynthesis of 1 and 2. (a) LCMS traces from *in vitro* assay of EupF. (b) DFT-computed free energies for hDA reaction between 8 and 6. (c) Biosynthetic pathway of 1.