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Introduction

Rising demand for transportation fuels, diminishing reserves of fossil oil, and the concerns with fossil fuel derived environmental pollution as well as the green-house gas emission derived climate change have resulted in the compelling need for alternative, sustainable new energy sources(1). Algae-based biofuels have been considered one of the promising alternatives to fossil fuels as they can overcome some of these issues (2-4). The current state-of-art of algal biofuel technologies have primarily focused on biodiesel production through prompting high algal lipid yields under the nutrient stress conditions. There are less interests of using algae-based carbohydrate and proteins as carbon sources for the fermentative production of liquid fuel compounds or other high-value bioproducts(5-7). However, under robust algae growth conditions, algal carbohydrate and proteins typically comprise up to ~ 80% of the ash-free dry weight of microalgae biomass (8, 9). Therefore, production of algal biofuel through comprehensive utilization of all algae biochemical components and the addition of high energy density fuel compounds with “fit for purpose” properties or the high value bioproducts will both diminish the process cost and improve the overall process feasibility.

Terpenes are a group of natural products with over 55000 structurally chemical compounds. Compared to biodiesel and other short-, medium-chain alcohols, these cyclic hydrocarbons, containing near zero oxygen content, not only have various biological functionalities(10-14) but also harbor high energy density making them a particularly attractive candidates as “drop-in” fuel candidates for aviation fuels(15-20). In this study, we demonstrated the concept of a "one-pot" bioconversion of algal carbohydrate and protein into terpenes as advanced biofuel compounds and the high value bioproducts through the development of engineered microbial consortium.

Results and Discussion

Caryophyllene and chamigrene, natural bicyclic sesquiterpene (C15) compounds, are common components presenting in the essential oils from various plants (21-24). Recent study suggested that

the blending of hydrogenated sesquiterpanes (in particular carophyllanes, which have a moderate cetane number and only moderately high viscosity, with synthetic branched paraffins to raise cetane and reduce viscosity could produce biosynthetic fuels that meet applicable jet and diesel specifications(25). Therefore, caryophyllene and its isomers have been deemed to be among the top three most promising candidates for jet fuel with the high energy density (26). In our previous study, we firstly discovered and functionally characterized caryophyllene and chamigrene synthases from endophytes(27). Furthermore, we demonstrated the feasibility of bioconversion of algal protein into terpene through terpene biosynthesis reconstruction into mutant *E.coli* strain YH40. Based on the previous studies, hereby, we developed a synthetic microbial consortium and investigated the production of caryophyllene, chamigrene, and other terpene products in one pot fermentation using algal hydrolysate of microalgae monocultures from strain *Reed nannochloropsis sp* as well as natural benthic algal assemblages cultivated from wastewater. To achieve this, the terpene biosynthesis pathway was reconstructed into *E.coli* strain YH40(7), designated for the conversion of algal protein into caryophyllene or chamigrene, and DH1, designated for the conversion of algal carbohydrate into caryophyllene or chamigrene, respectively, as described in previous studies(17). The caryophyllene and chamigrene yields were investigated under three different combinations of inoculum YH40-CI4A-CS/DH1-CI4A-CS at ratios of 2:1, 1:1, 0.5:1 as well as the single strains YH40-CI4A-CS or DH1-CI4A-CS alone. As shown in figure 1 (A), when the co-culture of two strains containing caryophyllene synthases were grown on algal hydrolysate from strain *Reed nannochloropsis sp*, the consortia at an inoculum ratio 1:1 (consortia R1) produced the highest titer of total terpene, up to 507.4 mg/L, including 471 mg/L of sesquiterpene, 36.4 mg/L of monoterpene as well as 124.4 mg/L of caryophyllene. Correspondingly, the consortia R1 consumed the highest amount of algal carbohydrates and proteins, which accounted for 48.2% of total algal carbohydrates and 36% of total algal proteins in the media. Compared to the consortia R1, the consortia R2 and R0.5 consumed less of the total algal biomass, with correspondingly lower concentrations of terpenes. The strain YH40-CI4A-CS alone produced the least amount of total terpene (274.7 mg/L), sesquiterpene (232.1 mg/L) and caryophyllene (14.4 mg/L) while DH1-CI4A-CS yielded 30% higher sesquiterpene and total terpene than strain YH40-CI4A-CS as well as 4 times higher titer of caryophyllene (75.2 mg/L). Compositional analysis of the *Reed nannochloropsis sp*. biomass indicated that the biomass was 20% carbohydrates and 58% protein (data not shown). Based on this data, the highest terpene yield that was achieved corresponded to ~42 mg total terpene/ g algae from consortia R0.5 with 37.4 mg sesquiterpene/ g algae and 6.6 mg caryophyllene/ g algae, as shown in figure 1(D).

In terms of co-culture of these two engineered strains containing chamigrene on the hydrolysate of benthic algal assemblages, the experimental results showed that the terpene yield reached 187 mg/L total terpene at the ratio of 2 (YH40-CI4A-CPS/DH1-CI4A-CPS), including 87 mg/L of monoterpene and 100 mg/L of sesquiterpene, in which the chamigrene was the major product accumulated up to 62 mg/L. The synthetic microbial consortia produced similar total terpene at the ratio of 1 and 0.5 (YH40-CI4A-CPS/DH1-CI4A-CPS), which were ~150 mg/L of total terpene. The microbial consortium at ratio 1 yielded the highest concentration of sesquiterpene (113 mg/L) as well as chamigrene (80 mg/L) among three consortia while the monoterpene yield was the lowest (34.5 mg/L). The strains YH40-TS and DH1-TS alone only produced 26 and 43 mg/L of total terpene, respectively, indicating relatively inefficient bioconversion of algal biomass. Compared to single strain, the synthetic microbial consortia produced 2.5-6.2 times higher total terpene concentration, suggesting that both algal carbohydrate and protein can be more effectively converted in the single-pot process. In terms of algal carbohydrate and amino acid consumption, none of the synthetic consortia were able to completely consume the algal carbohydrates and amino acids. The 2:1 consortium ratio utilized the highest amount of algal biomass, corresponding to 36.8% of total carbohydrates and 31.3% of algal amino acids. The other two consortia ratios consumed similar amount of the total carbohydrates and algal amino acids, which were 10-15% less than the 2:1 consortium. Strain YH40—CI4A-CPS utilized approximately half of the algal amino acids in the medium but algal carbohydrate consumption was minimal (3.8% of total carbohydrate). Strain DH1-TS consumed both algal carbohydrates (37.8 % of total carbohydrate) and amino acids (23.3% of algal amino acids) in the medium. Composition analysis indicated that carbohydrate and protein accounts for 74.2% of the mixed benthic biomass ash free dry weight (HydroMentia, Inc). Based on these data, the 2:1 consortium ratio produced the highest terpene yield at 30.5 mg terpene/ g algae while the 1:1 and 1:2 consortium ratios yielded 27.0 and 28.5 mg terpene/ g algae, respectively. The strain YH40—CI4A-CPS only produced 3.3 mg terpene/ g algae, which was lower than 8.7 mg terpene/ g algae yielded by strain DH1-CI4A-CPS, as shown in figure 2 (B).

Compared to total terpene yield produced from the benthic polyculture biomass in previous study, the consortium employing *Reed nannochloropsis sp.* Monoculture produced more than one fold higher titer of total terpene. In the consortium used for bioconversion of the benthic polyculture biomass, the chamigrene synthase (JGI protein ID 322581) gene was expressed as the last enzyme in the terpene biosynthesis pathway. Compared to the multiple sesquiterpene produced by caryophyllene synthase in this study, chamigrene synthase only produces a single sesquiterpene(chamigrene) with a limited number of monoterpenes(17), which was likely a reason for the higher yield of total terpene from *Reed nannochloropsis sp.* Furthermore, the ash content of the benthic polyculture biomass was more than 50%

of total mass, compared to 5.9% of *Reed nannochloropsis sp.* (data not shown). The higher ash content of the benthic polyculture biomass resulted in higher ion strength in the final algal hydrolysates (fermentation medium), which retarded the cell growth and further compromised the terpene yield.

Additionally, according to the techno-economic analysis of current state-of-art technology of essential oil production, the extraction yield of essential oil were ranged from 0.1% to 1% of plant tissue, corresponding to 1mg-10 mg essential oil/ g plant tissue(28, 29) due to the relative low concentration of essential oil in plant tissue(30). Compared to the extraction yield of essential oil from plant tissue, the engineered strains in this study increased the terpene yield about 3~40 times, which makes it a promising alternative pathway for terpene production.

Conclusion

Algae-based biofuels production has primarily focused on biodiesel production through transesterification of algal lipids. Under robust algal biomass accumulation conditions, carbohydrate and proteins typically comprise up to ~80% of the ash-free dry weight of algae biomass. Therefore, a comprehensive process for bioconversion of algal carbohydrates and proteins to high energy density fuels and value-added bioproducts should significantly improve the algal fuel process feasibility. In this study, we demonstrated simultaneous bioconversion of algal carbohydrates and proteins to terpenes which are attractive candidates for high energy density aviation fuels and other intermediate to high value bio-based chemicals applications. Using an engineered microbial consortium, greater than 30% of the carbohydrates and proteins from both a wastewater-based mixed algal feedstock and monoculture of strain *Reed nannochloropsis sp* were converted to terpenes, including both monoterpenes and sesquiterpenes. This microbial consortium concept for comprehensive utilization of algal biomass offers a versatile path forward for the production of fuels and active bioproducts from algae.

Material and Methods

Strains and Plasmids

The *E.coli* strain DH1 was obtained from Joint BioEnergy Institute (JBEI). The mutant *E.coli* strain YH40 (BW25113/F' [traD36, proAB+, lacIqZΔM15]ΔglnA, ΔgdhAΔluxSΔlsrA) was generously provided by Professor James C Liao from University of California, Los Angeles (UCLA). The plasmid pBbE1a-TSP_{up} containing the terpene biosynthesis pathway and the plasmid pBbE7k-TS were

constructed in our previous study. The two plasmids were co-transformed into strains DH1 and YH40, respectively.

Terpene production from a microbial consortium on algal hydrolysates

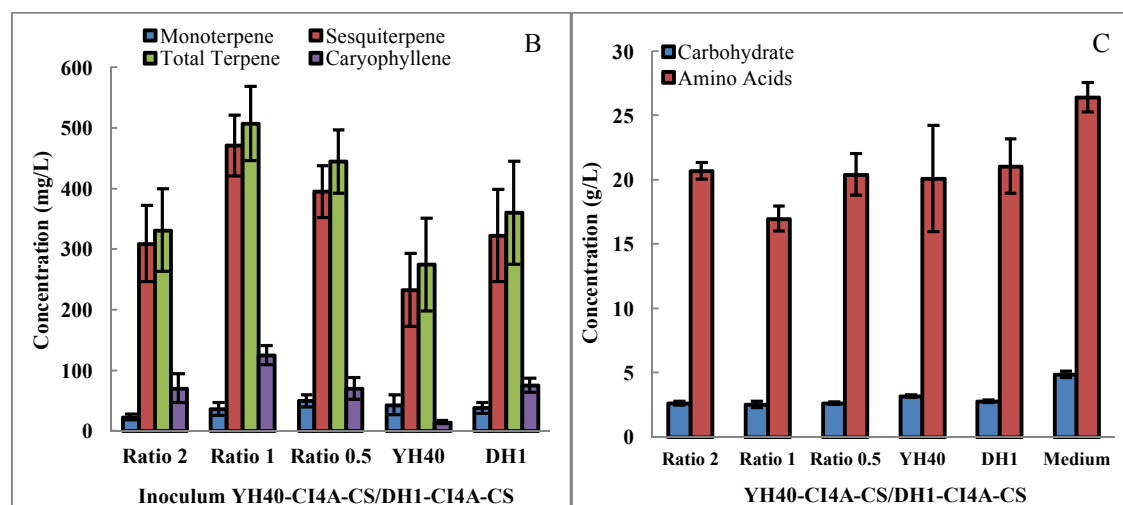
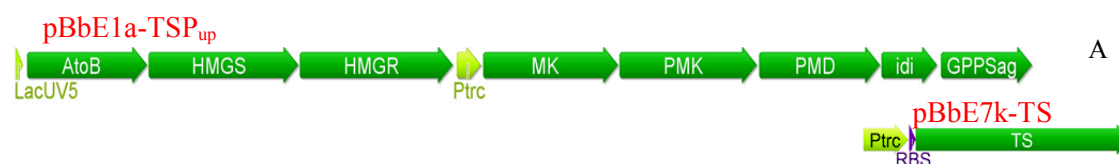
Algal biomass samples from both sources were pretreated according to protocols from the National Renewable Energy Laboratories and hydrolyzed with 2 mg/mL Pronase (promega, CA) following the manufacturer's protocol. The pretreated and hydrolyzed algal biomass was sterilized through filtration. *E. coli* strains DH1 and YH40 each containing the terpene biosynthesis pathway were cultured into 15ml of LB medium as described in the previous study. The overnight cultures were centrifuged and the cell pellets were re-suspended into 4 ml of pretreated algal hydrolysate. Various ratios (2:1, 1:1, 1:2) of engineered YH40 to DH1 were inoculated into the algal hydrolysate at a final concentration of 10% v/v. The culture were incubated at 37°C, 220 rpm and induced with 1 mM IPTG once the OD reached 0.8. The flasks were cap-sealed and cultured for another 72 hours at 25°C, 180 rpm for terpene production. Analytical samples were taken at the initial and end point of fermentation. The concentrations of total carbohydrate and amino acids were determined according to the established colorimetric protocols. The terpene profile and concentration was determined as described in the previous study(17, 27).

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Caption of Figures and Tables

Figure 1: Comprehensive conversion of algal carbohydrate and protein into caryophyllene and others terpenes using synthetic microbial consortium on algal hydrolysate of strain *Reed nannochloropsis sp.* A: Construct of terpene biosynthesis pathway, B: caryophyllene and other terpene concentration, C: algal carbohydrate and protein consumption of microbial consortium, D: caryophyllene and other terpene yields based on the substrate consumption.

Figure 2: Comprehensive conversion of algal carbohydrate and protein into caryophyllene and others terpenes using synthetic microbial consortium on algal hydrolysate of benthic polyculture biomass. A: chamigrene and other terpene concentration, B: algal carbohydrate and protein consumption and total terpene yields based on the substrate consumption.



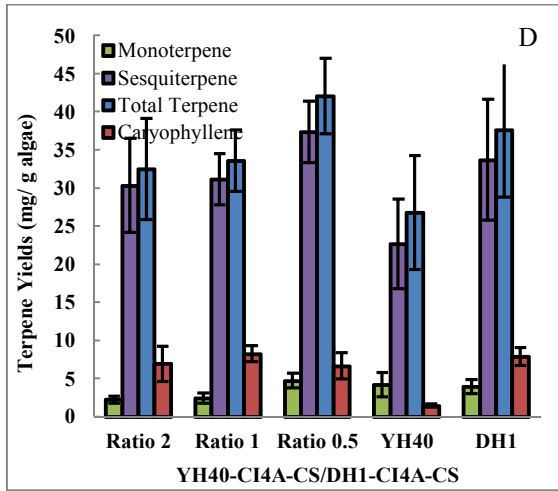


Figure 1.

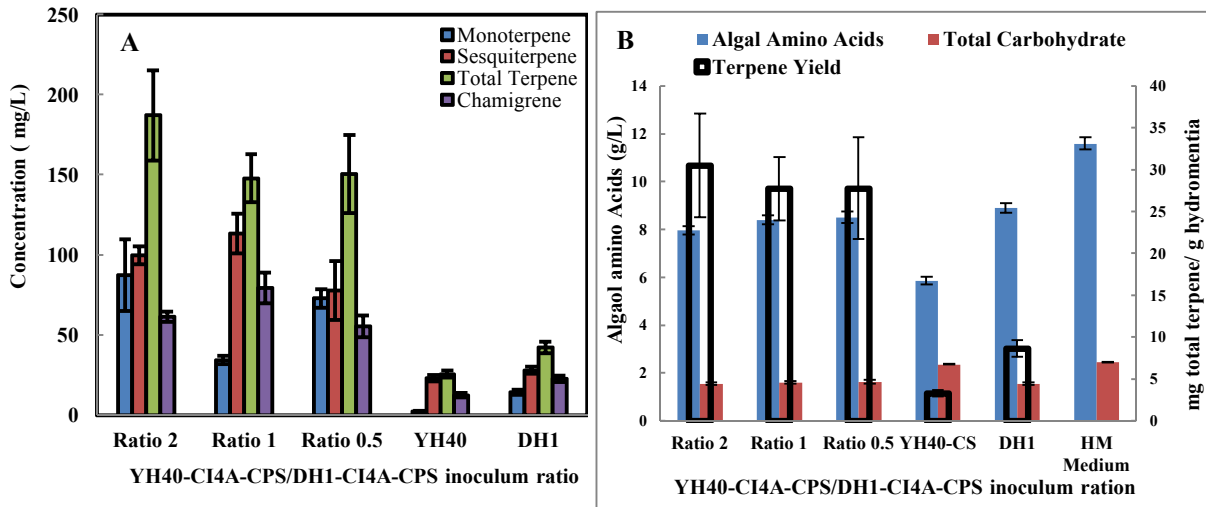


Figure 2

Reference

1. Lynd LR, van Zyl WH, McBride JE, Laser M. Consolidated bioprocessing of cellulosic biomass: an update. *Current Opinion in Biotechnology*. 2005;16(5):577-83.
2. Moody JW, McGinty CM, Quinn JC. Global evaluation of biofuel potential from microalgae. *Proceedings of the National Academy of Sciences of the United States of America*. 2014;111(23):8691-6.
3. Razeghifard R. Algal biofuels. *Photosynthesis Research*. 2013;117(1-3):207-19.
4. Luque R. Algal biofuels: the eternal promise? *Energy and Environmental Science*. 2010;3:254-7.
5. El-Mashad HM. Biomethane and ethanol production potential of *Spirulina platensis* algae and enzymatically saccharified switchgrass. *Biochemical Engineering Journal*. 2015;93:119-27.
6. Babujanarthanam R, Kavitha P. Simultaneous Saccharification and Fermentation of Dilute Acid Pretreated Red Algae (*Gelidiella acerosa*) for Bioethanol Production. *Energy Sources Part a-Recovery Utilization and Environmental Effects*. 2014;36(12):1305-14.
7. Huo Y-X, Cho KM, Rivera JGL, Monte E, Shen CR, Yan Y, et al. Conversion of proteins into biofuels by engineering nitrogen flux. *Nature Biotechnology*. 2011;29(4):346-U160.
8. Wang H, Ji B, Wang J, Guo F, Zhou W, Gao L, et al. Growth and biochemical composition of filamentous microalgae *Tribonema* sp as potential biofuel feedstock. *Bioprocess and Biosystems Engineering*. 2014;37(12):2607-13.
9. Chen C-Y, Zhao X-Q, Yen H-W, Ho S-H, Cheng C-L, Lee D-J, et al. Microalgae-based carbohydrates for biofuel production. *Biochemical Engineering Journal*. 2013;78:1-10.
10. Zhang Z, Guo S, Liu X, Gao X. Synergistic Antitumor Effect of alpha-pinene and beta-pinene with Paclitaxel against Non-small-cell Lung Carcinoma (NSCLC). *Drug research*. 2015;65(4):214-8.
11. Rufino AT, Ribeiro M, Sousa C, Judas F, Salgueiro L, Cavaleiro C, et al. Evaluation of the anti-inflammatory, anti-catabolic and pro-anabolic effects of E-caryophyllene, myrcene and limonene in a cell model of osteoarthritis. *European journal of pharmacology*. 2015;750C:141-50.
12. Kovac J, Simunovic K, Wu Z, Klanecnik A, Bucar F, Zhang Q, et al. Antibiotic Resistance Modulation and Modes of Action of (-)-alpha-Pinene in *Campylobacter jejuni*. *Plos One*. 2015;10(4).
13. Han L, Yang L, Liu B, Cheng X. Trans-caryophyllene suppresses tumor necrosis factor (TNF alpha)-induced inflammation in human chondrocytes. *European Food Research and Technology*. 2014;239(6):1061-6.
14. Guo K, Mou X, Huang J, Xiong N, Li H. Trans-Caryophyllene Suppresses Hypoxia-Induced Neuroinflammatory Responses by Inhibiting NF-kappa B Activation in Microglia. *Journal of Molecular Neuroscience*. 2014;54(1):41-8.
15. Strobel G. The story of mycodiesel. *Current Opinion in Microbiology*. 2014;19:52-8.
16. Riyaz-Ul-Hassan S, Strobel G, Geary B, Sears J. An Endophytic *Nodulisporium* sp from Central America Producing Volatile Organic Compounds with Both Biological and Fuel Potential. *Journal of Microbiology and Biotechnology*. 2013;23(1):29-35.
17. AM Gladden CAT, Connie Gao, Greg O'Bryan, Amy J Powell, Adam M Scheer, Kevin Turner, Weihua Wu, Eizadora T Yu. Tailoring Next-Generation Biofuels and Their Combustion in Next-Generation Engines. Sandia Report. 2013:2013-10094.
18. Strobel G, Singh SK, Riyaz-Ul-Hassan S, Mitchell AM, Geary B, Sears J. An endophytic/pathogenic *Phoma* sp from creosote bush producing biologically active volatile compounds having fuel potential. *Fems Microbiology Letters*. 2011;320(2):87-94.
19. Strobel GA, Knighton B, Kluck K, Ren Y, Livinghouse T, Griffin M, et al. The production of myco-diesel hydrocarbons and their derivatives by the endophytic fungus *Gliocladium roseum* (NRRL 50072) (vol 154, pg 3319, 2008). *Microbiology-Sgm*. 2010;156:3830-3.
20. Griffin MA, Spakowicz DJ, Gianoulis TA, Strobel SA. Volatile organic compound production by organisms in the genus *Ascocoryne* and a re-evaluation of myco-diesel production by NRRL 50072. *Microbiology-Sgm*. 2010;156:3814-29.
21. Malingré T HH, Batterman S, Bos R, Visser J. The essential oil of *cannabis sativa*. *Planta Medica*. 1975;28(1):56-61.
22. Kpadonou Kpoviessi BG1 LE, Kpoviessi DS, Gbaguidi F, Yehouenou B, Quetin-Leclercq J, Figueredo G, Moudachirou M, Accrombessi GC. Chemical variation of essential oil constituents of *Ocimum gratissimum*

- L. from Benin, and impact on antimicrobial properties and toxicity against *Artemia salina* leach. *Chemistry & biodiversity*. 2012;9(1):139-50.
23. Rodrigues FF1 OL, Rodrigues FF, Saraiva ME, Almeida SC, Cabral ME, Campos AR, Costa JG. Chemical composition, antibacterial and antifungal activities of essential oil from *Cordia verbenacea* DC leaves. *Pharmacognosy research*. 2012;4(3):161-5.
24. Meccia G RL, Velasco J, Díaz T, Usubillaga A, Arzola JC, Ramos S. Chemical composition and antibacterial activity of the essential oil of *Cordia verbenacea* from the Venezuelan Andes. *Natural product communications*. 2009;4(8):1119-22.
25. Benjamin G. Harvey WWM, and Thomas A. Koontz. High-Density Renewable Diesel and Jet Fuels Prepared from Multicyclic Sesquiterpanes and a 1-Hexene-Derived Synthetic Paraffinic Kerosene. *Energy Fuels*. 2015;29(4):2430-6.
26. Nakano C, Kim H-K, Ohnishi Y. Identification of the First Bacterial Monoterpene Cyclase, a 1,8-Cineole Synthase, that Catalyzes the Direct Conversion of Geranyl Diphosphate. *Chembiochem*. 2011;12(13):1988-91.
27. Weihua Wu WT, Craig A. Taatjes, Jorge Alonso-Gutierrez, Taek Soon Lee, John M. Gladden. Rapid Discovery and Functional Characterization of Terpene Synthases from Four Endophytic Xylariaceae. *PLoS One*. 2016.
28. Jonathan Moncada JAT, Carlos A. Cardona. Techno-economic and environmental assessment of essential oil extraction from Oregano (*Origanum vulgare*) and Rosemary (*Rosmarinus officinalis*) in Colombia. *Journal of Cleaner Production*. 2015:1-10.
29. HY. Gong WL, GY. LV, Xiaoying Zhou,*. Analysis of essential oils of *Origanum vulgare* from six production areas of China and Pakistan. *Revista Brasileira De Farmacognosia-Brazilian Journal of Pharmacognosy*. 2014;24:25-32.
30. Iijima Y D-RR, Fridman E, Gang DR, Bar E, Lewinsohn E, Pichersky E. The biochemical and molecular basis for the divergent patterns in the biosynthesis of terpenes and phenylpropenes in the peltate glands of three cultivars of basil. *Plant Physiology*. 2004;136(3):3724-36.