

Biochemical and gene expression analysis for the selection of elite oil palm ortets

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Abstract

The rising global demand for palm oil necessitates innovative strategies to accelerate yield improvement and ensure sustainable production. Conventional propagation methods are limited by long regeneration cycles, making tissue culture a more practical alternative that relies on elite ortet identification. This preliminary study aimed to evaluate crude palm oil quality, leaf metabolite profiles, and gene expression as early indicators for selecting elite ortets. Mature fruits were used to assess oil yield, free fatty acid, and β -carotene contents, while leaves were analyzed for chlorophyll and total phenolic content. Based on biochemical screening, three candidate ortets were further evaluated for the expression of acetyl-CoA carboxylase (*ACCase*) and phytoene synthase (*PSY*) genes. Oil yield ranged from 77-83%, FFA 5-9%, and β -carotene 200-450 mg kg⁻¹ CPO across candidates. Candidate 3.7 exhibited the highest oil yield and elevated *ACCase* expression, while candidate 6.12 showed the highest β -carotene content and *PSY* expression. Candidate 5.7 demonstrated relatively low FFA and favorable chlorophyll levels. These results suggest an association between *ACCase* expression and oil yield, as well as between *PSY* expression and β -carotene accumulation. Collectively, these findings support a proof-of-concept for an integrated profiling strategy to select elite ortets for advanced tissue culture propagation.

[Keywords: carotene content, clonal propagation, crude palm oil, *Elaeis guineensis*]

Introduction

Crude palm oil (CPO) is one of the world's largest vegetable oil commodities, serving as a key raw material for the food, textile, cosmetics,

pharmaceutical, and biodiesel industries (Hambali & Muzdalifah, 2020). Global demand continues to rise, with palm oil accounting for 51.14% of global vegetable oil consumption (Mustafa, 2022). In 2022, production exceeded 400 million tons, representing an increase of ~160% compared to 2000 (FAOSTAT, 2023). This sharp growth highlights the need for more efficient propagation systems to generate high-yielding and superior planting materials. Conventional propagation is limited by long growth cycles, whereas tissue culture provides a scalable alternative for producing uniform seedlings (Azahra & Ernayunita, 2023).

Tissue culture involves isolating plant tissues and cultivating them in an artificial medium enriched with nutrients and growth regulators, enabling the mass production of uniform seedlings (Kushairi et al., 2010; Baharuddin et al., 2025). This technique relies on explants derived from carefully selected parent plants, or ortets (Azahra & Ernayunita, 2023). Ortets are typically chosen based on agronomic traits such as oil yield and quality, growth vigor, crown architecture, and resistance to pests and diseases (Pratiwi et al., 2020). Conventional selection is primarily based on vegetative traits and bunch analysis, but such approaches are time-consuming and may not reliably predict oil quality or biochemical potential (Azahra & Ernayunita, 2023).

A promising strategy is to complement phenotypic evaluation with molecular and biochemical indicators. Key biosynthetic pathways strongly influence oil yield and metabolite profiles, including fatty acid synthesis regulated by acetyl-CoA carboxylase (*ACCase*) and carotenoid biosynthesis mediated by phytoene synthase (*PSY*) (Tranbarger et al., 2011; Wan Sulaiman et al., 2011). To describe the relationship between key traits and their underlying molecular physiology, we also

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measured the expression of *ACCase* and *PSY* to evaluate the potential of these oil palm ortets.

Based on this gap, the present study preliminarily investigates the oil and metabolite content of several candidate ortets. This integrative approach provides a scientific basis for identifying elite parent material to support efficient clonal propagation through tissue culture. This research aimed at determining chemical parameters to assess the initial potential of candidate ortets, including oil quality and metabolite content of oil palm leaves. Furthermore, this research also aims at evaluating how *ACCase* and *PSY* gene expression levels relate to oil yield and β -carotene content in the selected ortets, supporting the overall assessment of the ortet's potential. Therefore, this preliminary study aims at applying this integrated approach on a limited set of candidate ortets to identify promising individuals and establish proof-of-concept. The findings from this initial investigation will guide and inform a larger, more comprehensive study currently underway on a broader population.

Materials and Methods

Plant materials

Plant materials consisted of fruits and leaves collected from Tenera oil palms planted in 2010 at the Indonesian Oil Palm Research Institute (IOPRI), Bogor Unit, Ciomas, Bogor, Indonesia. Sampling was conducted under uniform field conditions. Mature fruits were harvested at a comparable stage of ripeness. Leaves were collected from the same position on the frond.

In this study, the candidate ortet is identified by a unique code (e.g., 3.7, 4.3, 5.5, etc). All fruits and leaves analyzed were collected from the same coded candidates. Mature fruits harvested from each candidate were used for CPO quality assessment, including oil yield, FFA content, and β -carotene concentration. Leaves (17th fronds) from the corresponding candidates were analyzed for total chlorophyll and total phenolic content. Based on the combined biochemical screening results, three candidate ortets with comparatively favorable parameters were selected for further molecular analysis. Young fruits harvested from these candidates were subsequently used to evaluate the expression levels of *ACCase* and *PSY* genes.

Determination of metabolite contents of CPO

Metabolite contents of CPO analyzed in this study include oil yield, FFA content, and β -carotene. CPO yield was measured using the modified method of Priyatama et al. (2023), with modifications based on the amount of sample and solvents used. The mesocarp (fruit flesh) was separated from the seeds

and cut into small pieces. The dried samples were ground after overnight drying (± 16 hours) in an oven. A 5 g sample was wrapped in filter paper and placed in a Soxhlet tube. The Soxhlet apparatus was assembled with a condenser and a fat flask containing 100 mL of petroleum benzene. Extraction was performed at 100°C and terminated when the solvent in the Soxhlet tube appeared clear. The fat flask was removed and dried in an oven at 100°C for 1 hour. The flask was reweighed to determine oil yield. Free fatty acid content in CPO was determined by the titrimetric method based on SNI 01-2901-2006 (AOCS, 2004). Two grams of CPO was dissolved in 50 mL of ethanol and heated to 40°C. Phenolphthalein (PP) was added as an indicator and titrated with 0.1 N NaOH. β -carotene was extracted from 0.1 g of CPO with 25 mL of *n*-hexane. β -carotene absorbance was measured at λ 446 nm (Sarip et al., 2023).

Determination of chlorophyll contents

Chlorophyll content was measured using the method of Singh et al. (2019). Chlorophyll was extracted from 0.1 g of palm leaf with 10 mL of 80% acetone. Chlorophyll absorbance was measured at λ 645 and 663 nm. The colorimetric assay for phenolic compounds was based on the Folin-Ciocalteu (FC) method (Leng et al., 2017) that involved phenolic extraction, pure phenolic compound (gallic acid) calibration, and absorbance measurement after color development. Phenolics were extracted from 0.2 g of dried palm leaves with 20 mL of methanol. A 1000 ppm gallic acid standard solution was diluted to 50, 75, 100, 150, and 200 ppm. Sample filtrate/standard solution (0.1 mL) was mixed with 0.5 mL FC reagent and 5 mL distilled water, vortexed, and left for 6 min. The mixture was reacted with 1.5 mL 20% Na_2CO_3 and adjusted to 10 mL with distilled water. Each reaction mixture was incubated for 2 hours before measuring absorbance at λ 765 nm.

cDNA preparation

Total RNA was isolated using the GeneJET RNA Purification Kit (Thermo Scientific, 2025). The concentration and purity of the isolated RNA were measured using a NanoDrop™ 2000 Spectrophotometer. For cDNA synthesis, 1 μg of total RNA was reverse-transcribed using ReverTra Ace™ qPCR RT Master Mix with gDNA Remover (Toyobo). PCR amplification was performed using MyTaq HS Red Mix (Bioline) in a 10 μL reaction containing 5 μL 2 $\times$ master mix, 0.4 μM of each primer listed in Table 1, and 1 μL cDNA template. The amplification was conducted under the following conditions: initial denaturation at 95°C for 1 min; 35 cycles of 95°C for 15 s, 55–60°C for 15 s,

and 72°C for 30 s; and a final extension at 72°C for 5 min. PCR products were resolved on 1% agarose gel.

Real-time PCR (RT-PCR) setup

Quantitative real-time PCR (qRT-PCR) was performed using SensiFAST™ SYBR® Hi-ROX Kit (Bioline) in [x system]. Each 10 µL reaction contained 5 µL 2× master mix, 0.4 µM of each primer described previously, 1 µL cDNA template, and nuclease-free water. The amplification conditions were: 95°C for 2 min, followed by 40 cycles of 95 °C for 10 s, 60°C for 10 s, and 72°C for 15 s. Melting curve analysis was subsequently performed to confirm primer specificity, with the following steps: 95°C for 15 s, 60°C for 1 min, and 95°C for 15 s. Each sample was analyzed in three technical replicates and three biological replicates. Relative gene expression levels were calculated using the $2^{-\Delta\Delta C_t}$ method (Livak & Schmittgen, 2001), where C_t values of target genes were normalized against the housekeeping gene (actin) to obtain ΔC_t . The $\Delta\Delta C_t$ values were calculated relative to the mean ΔC_t of all samples.

Amplification efficiency was assumed to be comparable between target and reference genes.

Results and Discussion

CPO quality: oil yield and FFA content

The oil yield of ten candidate ortets ranged from 77% to 83% (w/w) based on mesocarp dry weight (Figure 1). Candidate 3.7 recorded the highest yield at 83.56%, approximately 2–6% higher than the other candidates. Similarly, candidate 5.7 exhibited a comparatively high yield of 81.64%. These elevated CPO yields highlight both candidates for further evaluation. The yields reported here are higher than those obtained by Sarip et al. (2023), who recorded ~75% using Soxhlet extraction, yet align with Hasibuan (2020), who reported 80–84% yields from boiled and mechanically pressed fresh fruit. This consistency suggests that the extraction protocol applied in the present study is robust, with minor differences likely reflecting intrinsic variation in fruit physiology rather than methodological artifacts.

Table 1. Primer pairs used in RT-qPCR

Gene	Primer sequence	Gene function
<i>ACCase</i>	5'-GAAGCACCWTYCCTGCMYT	Fatty acid biosynthesis
	5'-CKHGTGGRGCCAYACRAT	
<i>PSY</i>	5'-TCCGACTGTGGCTTTCTTCT	β -Carotene biosynthesis
	5'-AGTCACACCAGCCTCTGCTT	
<i>Actin</i>	5'-TGCTGATCGTATGAGCAAGGAAA	Housekeeping gene
	5'-GAAATCCACATCTGCTGGAAGGT	

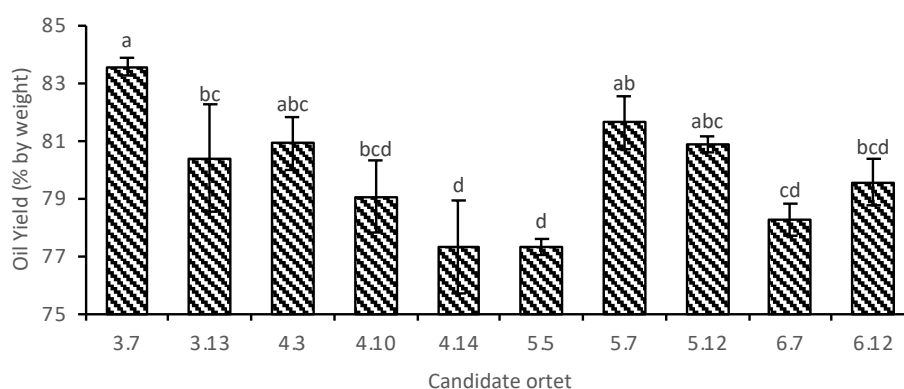


Figure 1. CPO yield from extraction. Data are presented as mean $\pm$ standard error (SE) (n = 3). ANOVA showed a $p < 0.05$. Different letters indicate significant differences.

The concentration of FFA is a critical indicator of CPO degradation and overall oil quality. Across the ten candidate ortets, FFA content ranged from 5% to 9% (w/w) (Figure 2). Candidate 4.14 exhibited the highest FFA level, whereas candidates 5.12 and 5.7 recorded the lowest. These values exceeded the maximum threshold established by SNI 2901-2021 ($\leq 5\%$). Nonetheless, the measured range is consistent with previous studies. Igile et al. (2023) reported FFA contents between 2.44% and 12.76%, while Japir et al. (2017) documented values of 8.7% and 3.8% across two oil palm fruit types. Similarly, Ekwenye (2010) observed varietal differences, with FFA levels ranging from 4.46% to 7.33% in Dura and from 6.02% to 14.76% Tenera, indicating a genetic predisposition toward higher FFA accumulation in the latter.

Free fatty acids are produced through lipase-catalyzed hydrolysis of triacylglycerols (Mahesar et al., 2016; Shidiq et al., 2022). Its accumulation is influenced by multiple factors, including storage duration, processing temperature, and fruit moisture content. Compared to triacylglycerols, FFAs are more prone to oxidation, which promotes rancidity and reduces oil quality (Fokou et al., 2009). Hudori (2016) further demonstrated that mesocarp storage conditions substantially influence FFA levels, with prolonged storage leading to marked increases. In extreme cases, FFA concentrations can reach 20% (Ekwenye, 2010), underscoring the susceptibility of

CPO to post-harvest degradation. Similarly, Japir et al. (2017) reported that high temperatures accelerate triacylglycerol breakdown, while fermentation processes involving lipolytic enzymes also enhance triacylglyceride hydrolysis (Fokou et al., 2009).

Metabolite content: β -carotene, chlorophyll, and phenolic compounds

Secondary metabolites in plants are broadly classified into terpenoids, phenolic compounds, and nitrogen-containing compounds (Anulika et al., 2016). In this study, we focused on β -carotene, chlorophyll, and phenolic compounds as key indicators of oil palm physiological status and CPO quality.

β -carotene ($C_{40}H_{56}$) was detected in concentrations ranging from 200 to 450 mg kg⁻¹ CPO (Figure 3). Candidate 6.12 contained the highest level (440 mg kg⁻¹) of β -carotene, consistent with reported ranges of 200–500 ppm (Tai & Brunner, 2019) and 305–506 ppm (Hasibuan, 2020). Freshly processed fruit often contains higher levels (542–868 ppm) (Silalahi, 2022). Beyond its role as a natural pigment, β -carotene functions as an antioxidant and is enzymatically converted to retinol (vitamin A) by β -carotene dioxygenase (Sari & Mayasari, 2018), linking its accumulation to both nutritional and industrial value.

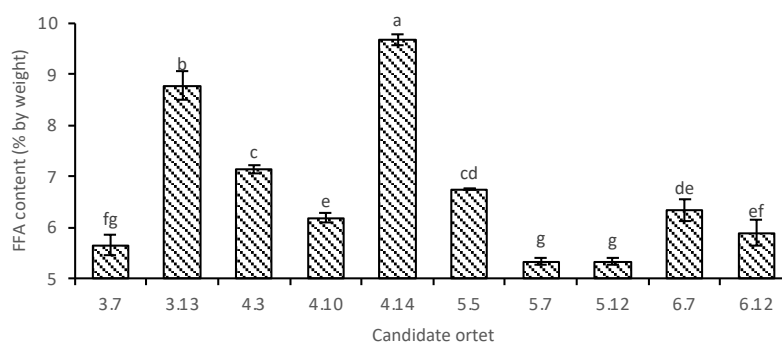


Figure 2. FFA content of CPO. Data are presented as mean $\pm$ standard error (SE) (n = 3). ANOVA showed a $p < 0.05$. Different letters indicate significant differences.

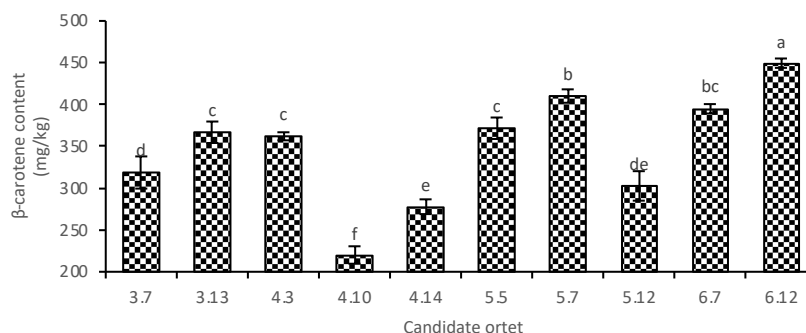


Figure 3. β -carotene content of CPO. Data are presented as mean $\pm$ standard error (SE) (n = 3). ANOVA showed a $p < 0.05$. Different letters indicate significant differences.

Chlorophyll, the primary pigment in photosynthesis, exists in two forms in higher plants: chlorophyll a (dark green) and chlorophyll b (light green) (Holidi et al., 2016; Mustafa et al., 2015). Chlorophyll a serves as the main electron donor in reaction centers, while chlorophyll b expands light-harvesting efficiency (Mulero et al., 2022). Total chlorophyll in fresh oil palm leaves ranged from 1.4 to 1.7 mg g⁻¹ (Figure 4), consistent with previous reports of 1–1.5 mg g⁻¹ (Manoh et al., 2021). The chlorophyll a/b ratio varied between 2.5 and 4.5, within the normal physiological range (Sonobe et al., 2020). Notably, candidate 6.7 exhibited a high a/b ratio, suggesting growth under high light intensity, whereas candidate 3.7 displayed the lowest, consistent with shaded conditions (Zhang et al., 2020). Variations in chlorophyll concentration and a/b ratios highlight how environmental factors, including temperature and rainfall, modulate photosynthetic efficiency and, ultimately, productivity (Djs et al., 2020).

Phenolic compounds, characterized by hydroxyl groups attached to aromatic rings, are another major class of secondary metabolites with key roles in plant defense (Mahardani & Yuanita, 2021). Here, total phenolic content in dried leaves ranged from 200 to 450 mg kg⁻¹ (Figure 5), aligning with previous findings of 50–300 ppm (Safa et al., 2024)

and 200–500 ppm (Hasan & Manan, 2020). Phenolics are synthesized in response to environmental stress (Hanin & Pratiwi, 2017) and function as antioxidants by donating hydrogen atoms to neutralize reactive oxygen species (Andrés et al., 2023). They also protect against UV-B radiation and DNA damage (Kuljarusnont et al., 2024), suggesting that elevated levels reflect both genetic predisposition and environmental stress adaptation.

Within our limited population, integrating metabolite and oil quality profiles reveals three promising candidate ortets. Candidate 3.7 combined high oil yield with elevated phenolic content, indicating superior stress tolerance and stability. Candidate 6.12 contained the highest β -carotene concentration, conferring added nutritional and antioxidant value. Meanwhile, candidate 5.7 exhibited low FFA levels and favorable chlorophyll profiles, traits associated with improved oil quality and photosynthetic capacity. However, the synergistic roles of phenolics (stress defense), chlorophyll (photosynthetic efficiency), and β -carotene (nutritional and oxidative protection) suggest a multifaceted contribution to CPO productivity and stability (Djs et al., 2020; Fokou et al., 2009; Hanin & Pratiwi, 2017).

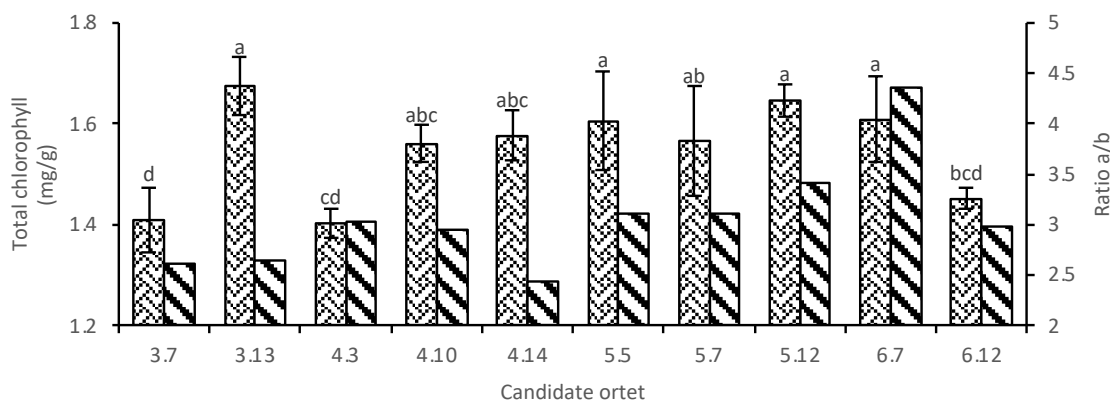


Figure 4. Total chlorophyll content and chlorophyll a/b ratio of oil palm leaves. Data are presented as mean $\pm$ standard error (SE) (n = 3). The ANOVA test yielded a $p < 0.05$. Different letters indicate a significant difference.

Table 2. Expression levels of actin target genes

Gene	Candidate Ortet	Expression level
<i>ACCase</i>	3.7	Down-regulated
	5.7	Down-regulated
	6.12	Down-regulated
<i>PSY</i>	3.7	Down-regulated
	5.7	Down-regulated
	6.12	Down-regulated

Expression of the metabolite-related genes in fresh fruit bunches

The relative expression levels of the *acetyl-CoA carboxylase* (*ACCase*) and *phytoene synthase* (*PSY*) genes, normalized to actin, ranged from 0.02–0.3 and 0.3–0.9, respectively (Figure 6). The comparatively low expression of *ACCase* indicates strong transcriptional repression in the mesocarp tissue analyzed (Table 2). Although *PSY* expression was also relatively low, its higher magnitude compared to *ACCase* suggests preferential activation of carotenoid biosynthesis over fatty acid biosynthesis in this tissue at the time of sampling.

Among the candidate ortets, the expression patterns of *ACCase* and *PSY* aligned with the observed metabolite accumulation. Candidate 3.7 exhibited the highest *ACCase* expression, approximately twofold above the mean level, and correspondingly produced the highest CPO yield. Similarly, candidate 6.12 showed the strongest *PSY* expression, ~1.2-fold higher than average, which was consistent with it containing the greatest β -carotene concentration. The physiological traits of candidate 5.7 were not strongly aligned with the expression of either target gene, which may reflect technical variation in RNA quality or pipetting inconsistencies, as noted in standard qPCR reliability studies (Kozera & Rapacz, 2013).

ACCase catalyzes the committed step of fatty acid biosynthesis, the ATP-dependent carboxylation of acetyl-CoA to malonyl-CoA (Wu & Huang, 2020). In oil palm mesocarp, the enzyme exists in a prokaryotic-type multienzyme complex, a complex comprising biotin carboxylase (BC), biotin carboxyl carrier protein (BCCP), and carboxyltransferase (CT) (Shen et al., 2024). The reaction proceeds in two sequential steps: (i) carboxylation of biotin bound to the BCCP domain, and (ii) transfer of the carboxyl group to acetyl-CoA by the CT domain to yield malonyl-CoA (Milke & Marienhagen, 2020). Thus, the observed variation in *ACCase* expression among ortets provides a plausible biochemical explanation for the differences in oil accumulation.

By contrast, *PSY* functions as the first committed enzyme in carotenoid biosynthesis (Zhou et al., 2022). The higher *PSY* expression in candidate ortet 6.12 offers a mechanistic insight into the enhanced β -carotene accumulation observed. Together, these findings describe a direct molecular relationship between the expression of these specific biosynthetic genes and the accumulation of their end products in these selected candidates. This correlation provides a physiological context for the observed phenotypic variation within these candidate ortets and underscores the value of an integrated approach for the preliminary characterization of ortet potential.

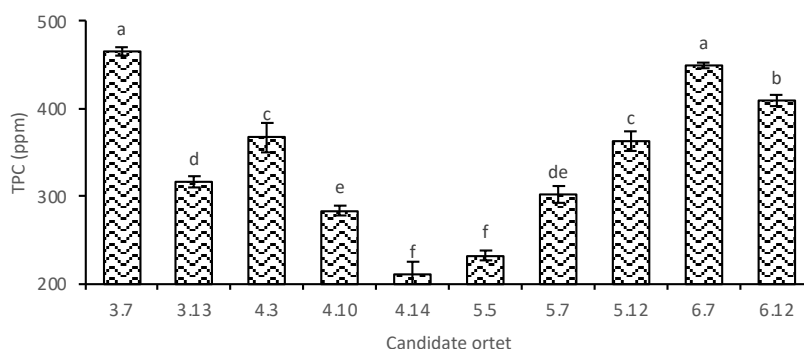


Figure 5. Total phenolic content (TPC) of oil palm leaves. Data are presented as mean $\pm$ standard error (SE) (n = 3). The ANOVA test yielded a p-value < 0.05. Different letters indicate a significant difference.

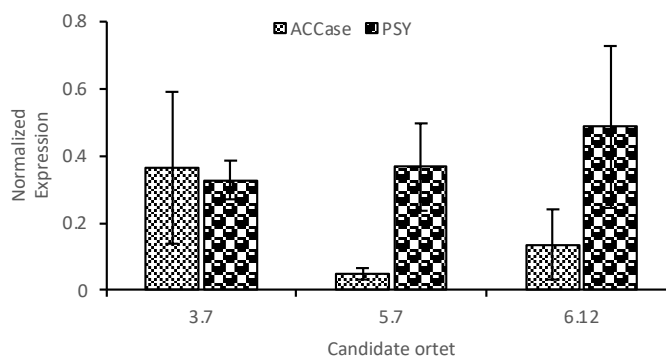


Figure 6. Expression of *ACCase* and *PSY* genes in oil palm fruit

Together, these findings describe a concurrent relationship between the expression of these specific biosynthetic genes and the accumulation of their end products in these selected ortets. The alignment of high *ACCase* expression with high oil yield in candidate ortet 3.7, and high *PSY* expression with high β -carotene in candidate 6.12, provides a physiological context for the observed phenotypic variation and underscores the value of an integrated approach for characterizing ortet potential. However, given the limited sample size of this initial investigation, the identified candidates are promising preliminary rather than definitive elite selections. A more comprehensive study on a larger population is currently ongoing to further investigate these relationships and identify robust selection markers.

Conclusion

This study establishes a foundational framework for selecting elite oil palm ortets by integrating phenotypic screening with molecular profiling. Within our tested population, candidate ortets 3.7, 5.7, and 6.12 emerged as superior candidates. The mechanistic consistency between *ACCase* expression and oil yield in candidate 3.7, and between *PSY* expression and β -carotene content in candidate 6.12, underscores the utility of this approach. While these results are promising, they represent a critical first step; future work must expand to a larger population and translate this proof-of-concept into validated, robust selection criteria for tissue culture applications.

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