

PROPOSAL

OPEN INNOVATION

TAHUN 2025





OPEN INNOVATION 2025

• RUANG LINGKUP

Dosen/Peneliti:

1. Peningkatan Oil Content: rekayasa metabolit untuk meningkatkan biosintesis minyak sawit, serangga pollinator & artificial pollination, aplikasi booster atau beneficial untuk meningkatkan oil to dry mesocarp

Maksimal 300 juta per Proposal

Mahasiswa :

1. Ekologi Polination
2. Breeding Elaeidobius K
3. Dinamika Reproduksi Serangga Polinator
4. Artificial Pollination
5. Aplikasi Beneficial Microbes atau booster untuk meningkatkan oil content

Maksimal 10 juta per Proposal

• KRITERIA PENILAIAN SELEKSI

Kriteria penilaian hasil Open Innovation akan meliputi:

1. Impact Terhadap Penyelesaian Masalah Di BGA (50%)
 2. Tingkat Technology Readiness Level (TRL) (25%)
 3. Kelayakan Anggaran dengan Program yang Diusulkan (15%)
 4. Kelayakan Aplikasi dan Scalability (10%)
-
1. Impact Terhadap Penyelesaian Permasalahan di BGA (50%)
 2. Metodologi riset (25%)
 3. Luaran hasil riset (25%)

• RANGKAIAN KEGIATAN

1. Inovator upload proposal ke <https://dashboard.bumitama.co.id/inovasi/>
2. Seleksi proposal administrasi dan presentasi
3. Pengumuman peserta lolos seleksi
4. Kick off Open Innovation
5. Pelaksanaan Riset Project
6. Evaluasi Project
7. Laporan Akhir

**Format proposal maksimal
12 Halaman menggunakan
PPT. Submit proposal
dalam bentuk file pdf.**

1

Justifikasi & Tujuan penelitian

2
halaman

2

Metodologi penelitian

2
halaman

3

Rincian Kebutuhan Biaya

2
halaman

4

Ganchart Detail Kegiatan
Penelitian dan Target Output

2
halaman

5

Analisis Cost & Benefit
(Impact Hasil Penelitian)

2
halaman



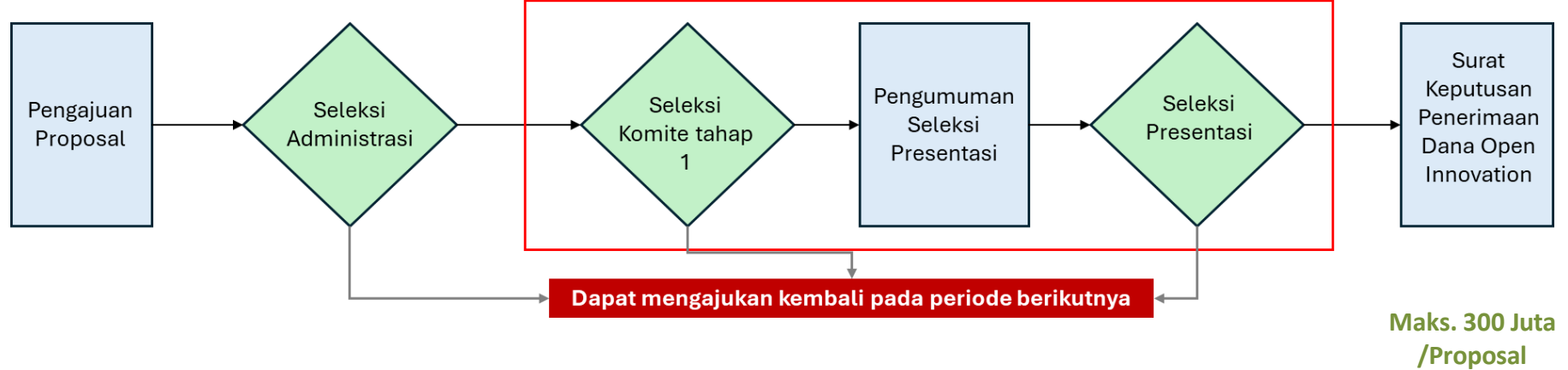
MEKANISME PENGAJUAN PROPOSAL

Skema OI Dosen:

Fokus Open Innovation Dosen/Peneliti:

1. Peningkatan Oil Content: rekayasa metabolit untuk meningkatkan biosintesis minyak sawit, serangga pollinator & artificial pollination, aplikasi booster atau beneficial untuk meningkatkan oil to dry mesocarp

Dosen/Peneliti:

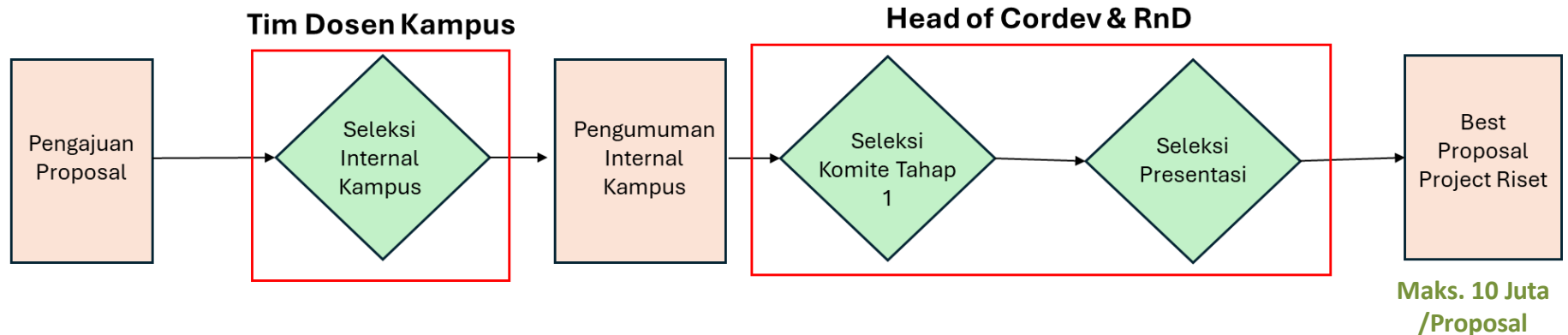


Skema OI Mahasiswa:

Fokus Open Innovation Mahasiswa :

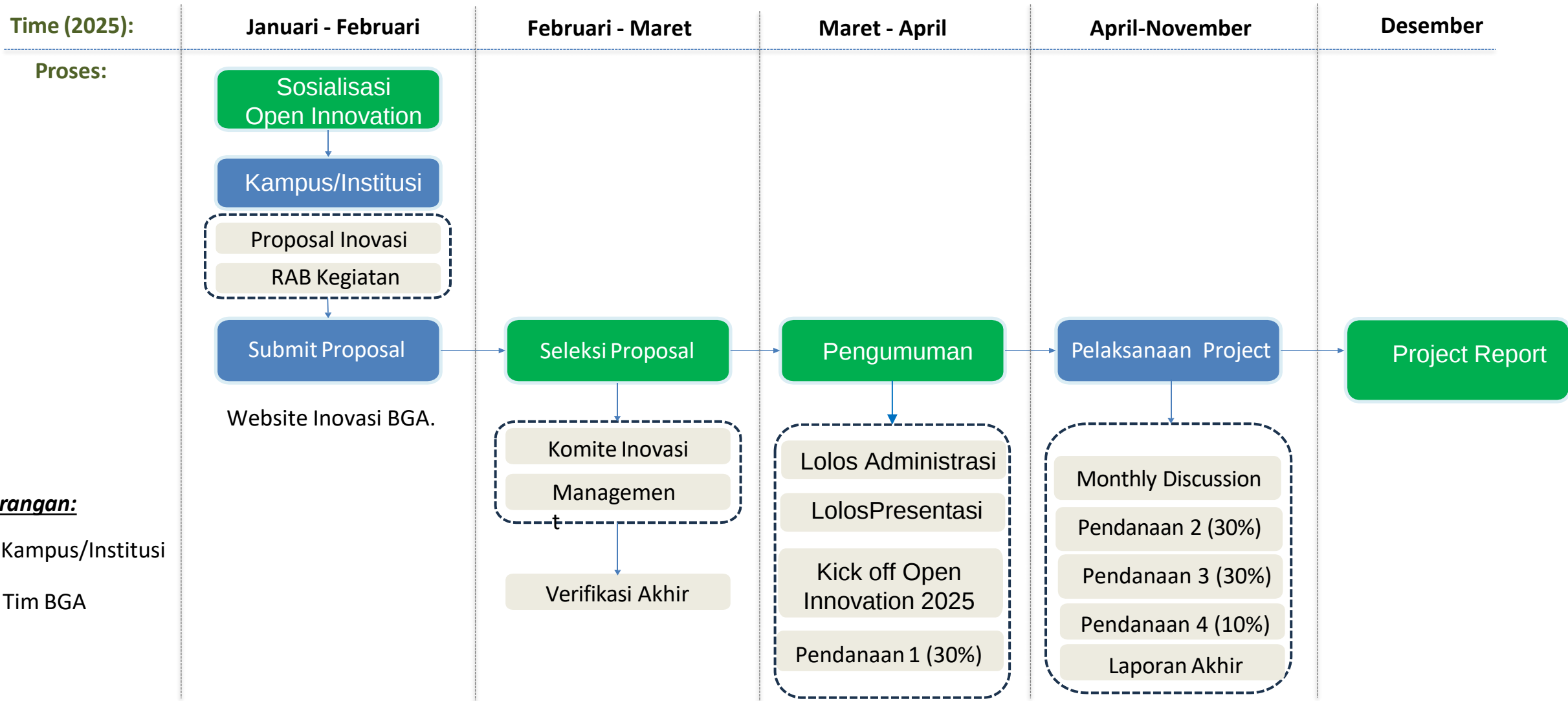
1. Ekologi Polination
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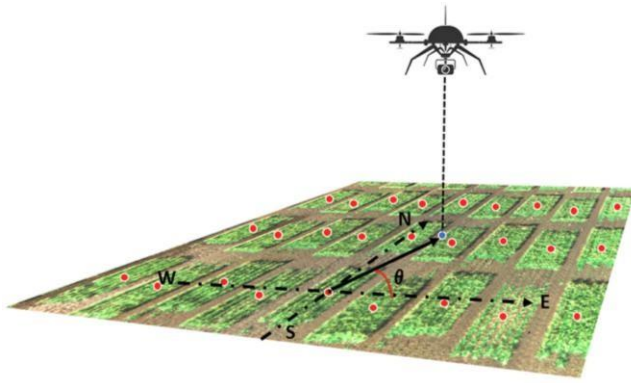
Mahasiswa :





OPEN INNOVATION 2025



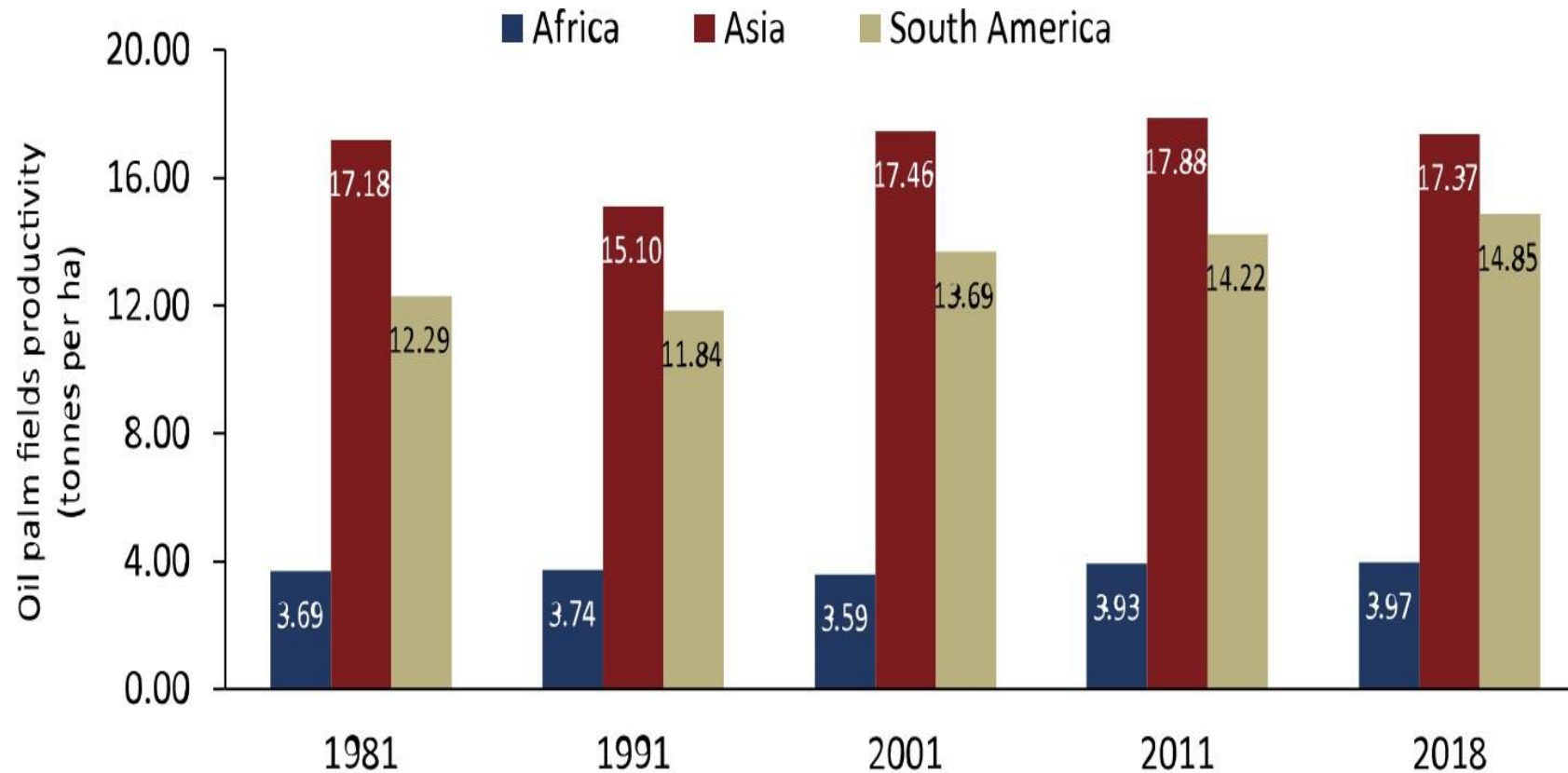


INOVASI TERINTEGRASI UNTUK MEMAJUKAN INDUSTRI SAWIT INDONESIA

PT. BUMITAMA GUNAJAYA AGRO (BGA)

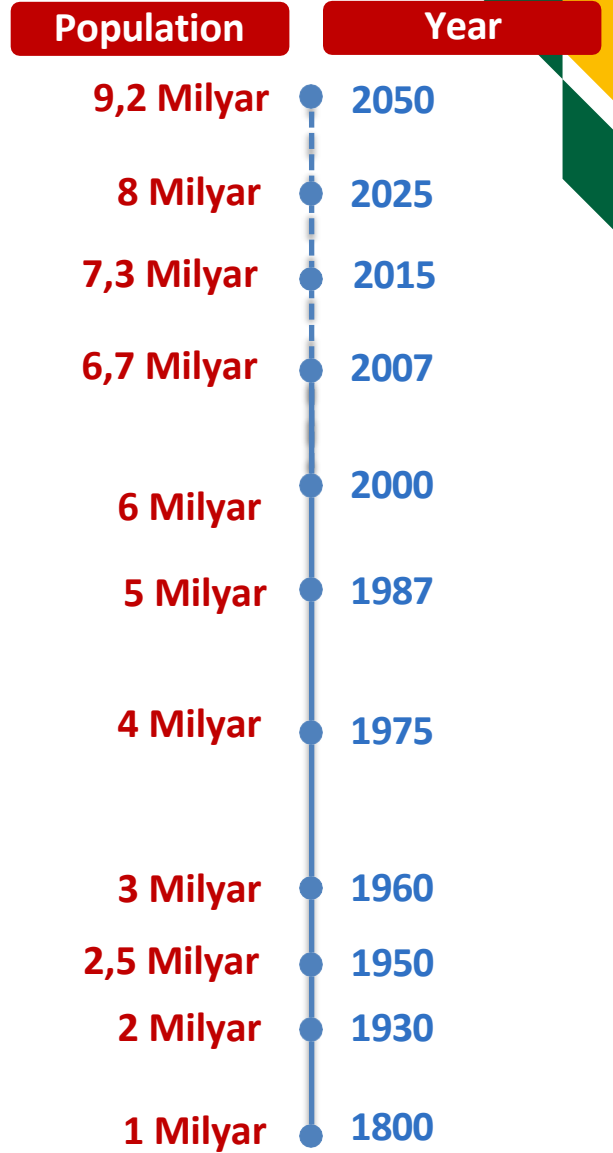


KEUNGGULAN KOMPARATIF DAN KOMPETITIF INDONESIA & TANAMAN SAWIT



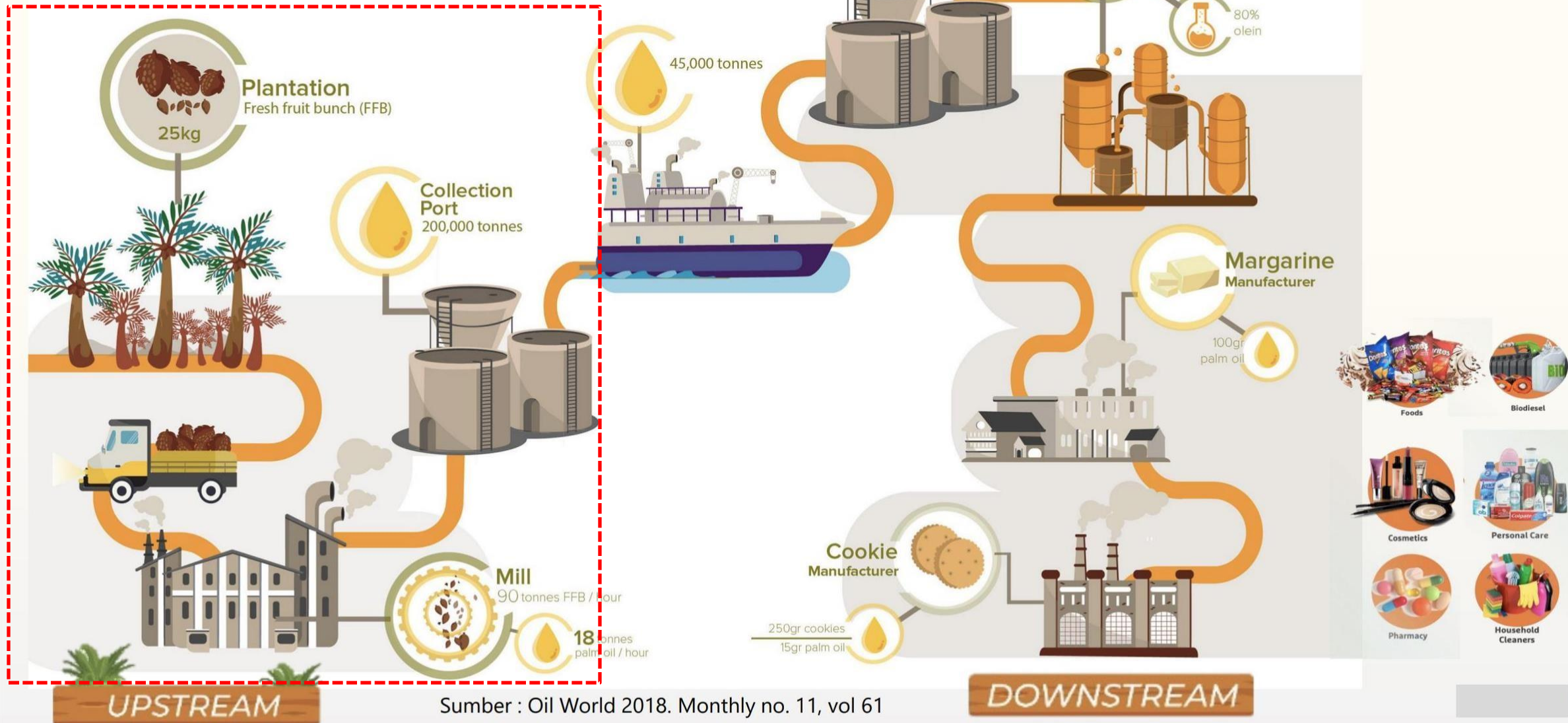
A comparative look at how Africa's oil palm productivity is in relation to other continents

WORLD POPULATION GROWTH



Palm Oil Supply Chain

NOTE : Volumes in this infographic are averages.
There is a lot of variation in processing, transport capacity and ingredient use



BUMITAMA GUNAJAYA AGRO (BUMITAMA AGRI)



Planted Area : 186,628 Ha
Nucleus : 131,099 Ha (70%)
Plasma : 55,529 Ha (30%)



Estates : 68 Estates
Mills : 17 Unit
Total Capacity : 1,065 Tph



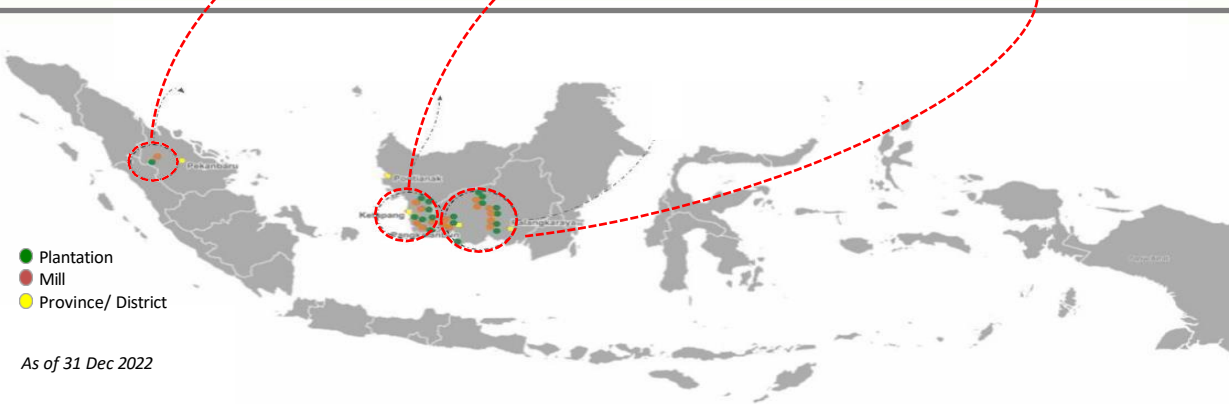
Staff : 1,500
Worker (eff) : 26,000



Riau:
 • 2K ha Planted Area
 • 1 Mill (90 tph capa)

West Kalimantan:
 • 81K ha Planted Area
 • 5 Mills (300 tph capa)

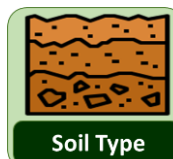
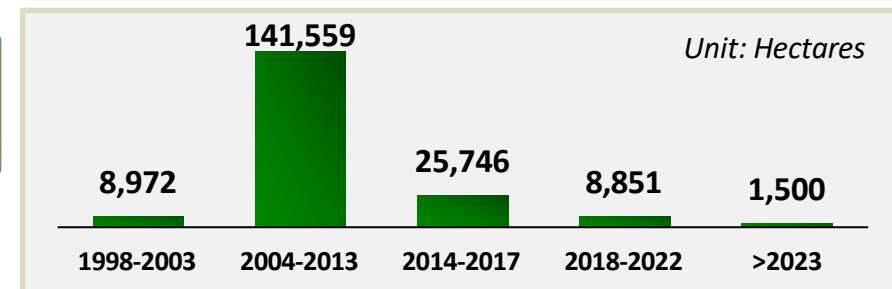
Central Kalimantan:
 • 105K ha Planted Area
 • 9 Mills (675 tph capa)



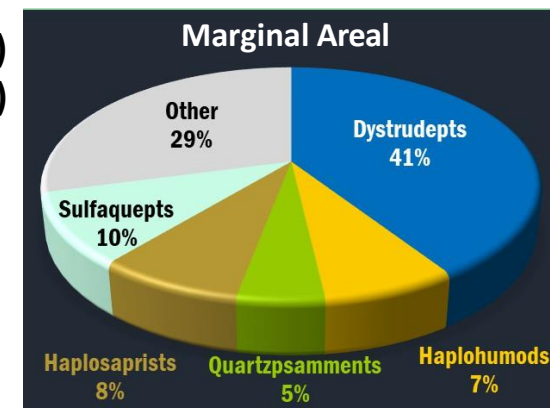
No. of Cooperatives : 56 Units
No. of Families : 35,599



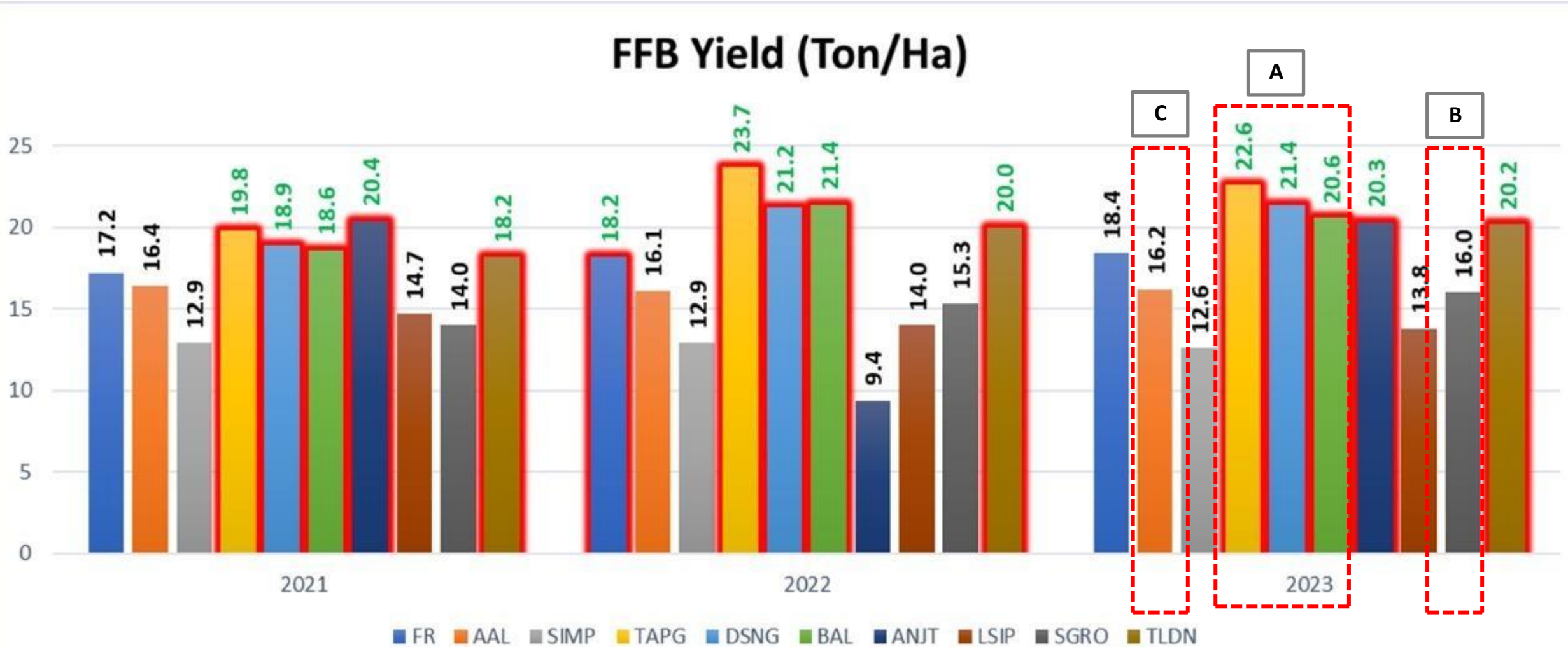
- **Bumitama Corporate University (BCU)** - : 2 Unit Learning & Development
- **Central Workshop** : 2 unit
- **Research Center & Innovation** : 4 Unit



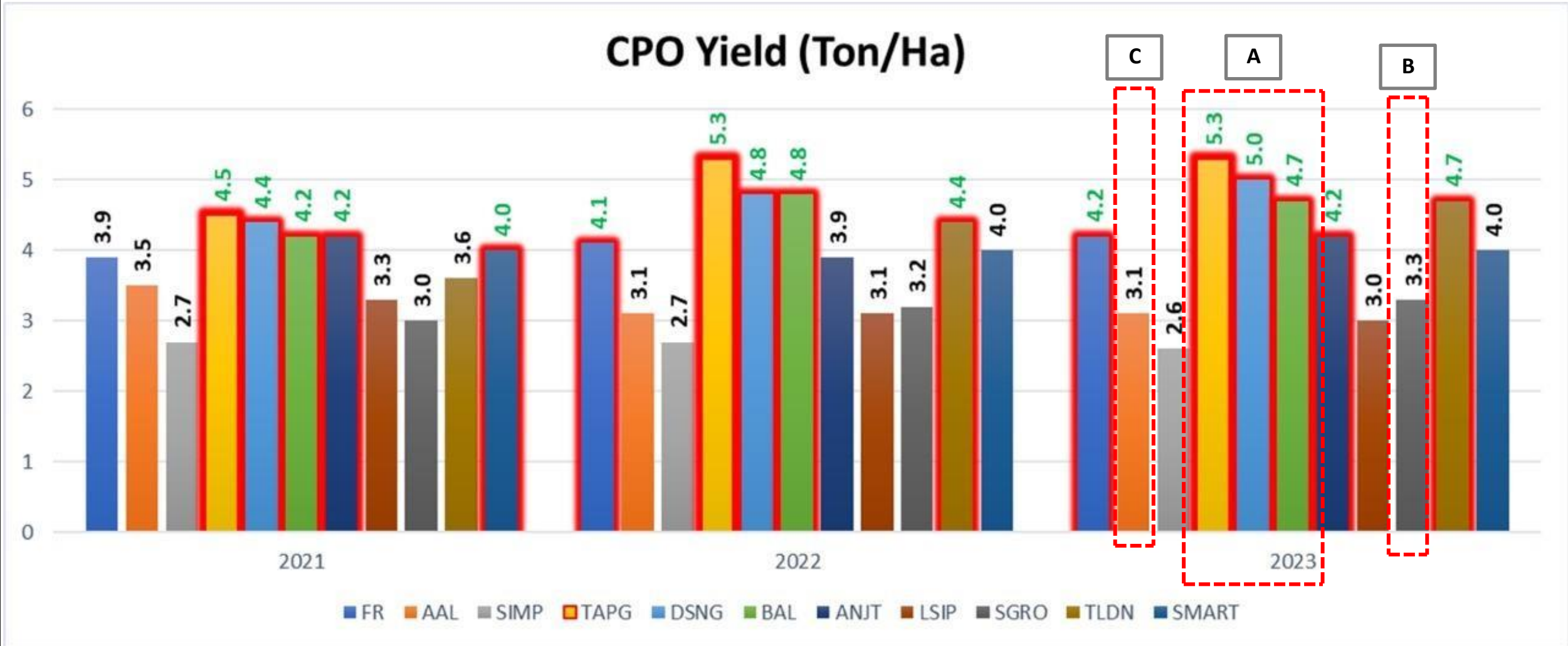
Mineral : 55,880 Ha (30%)
Marginal: 130.748 Ha (70%)



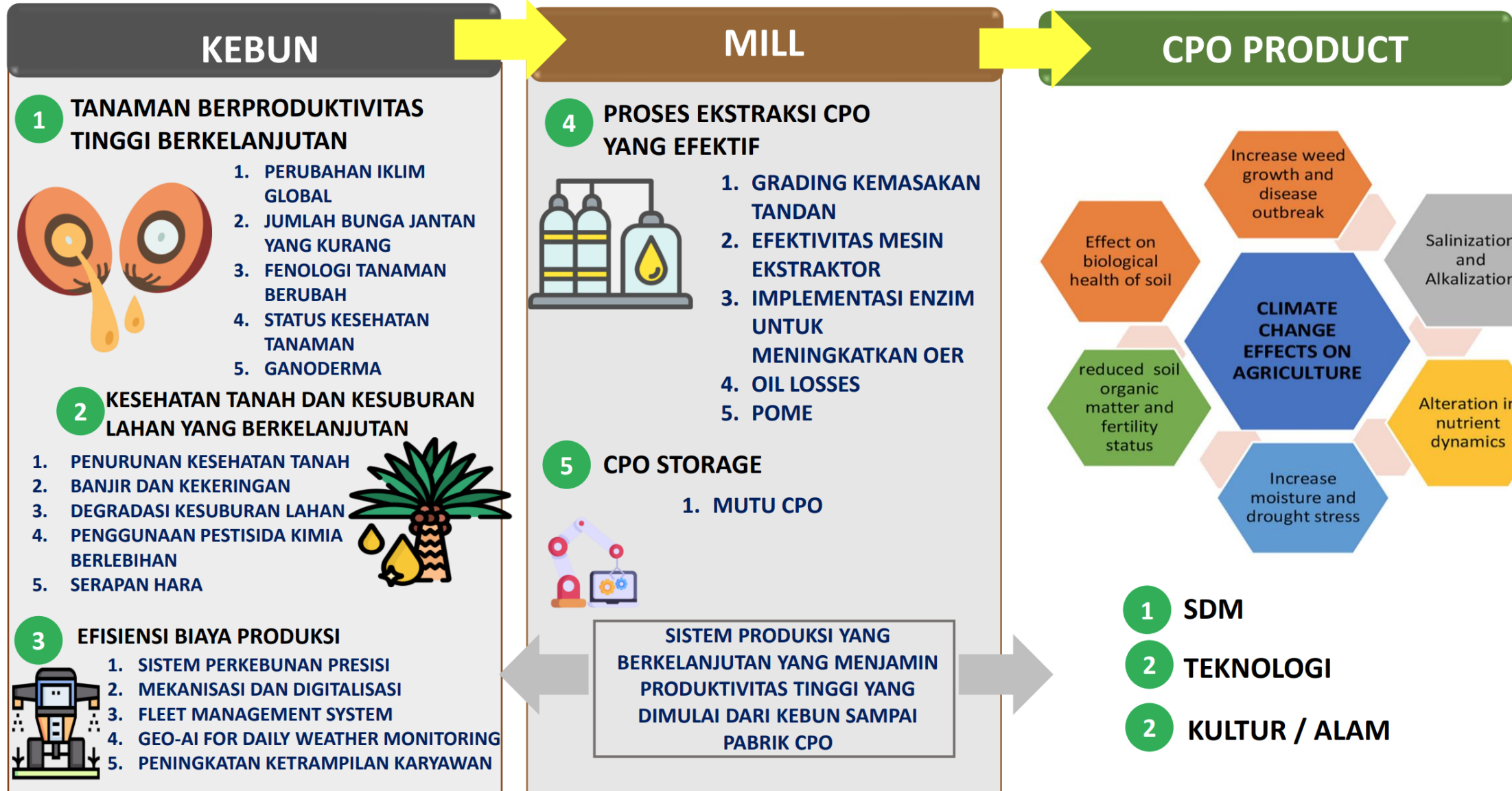
PRODUKTIVITAS BEBERAPA KEBUN DI INDONESIA



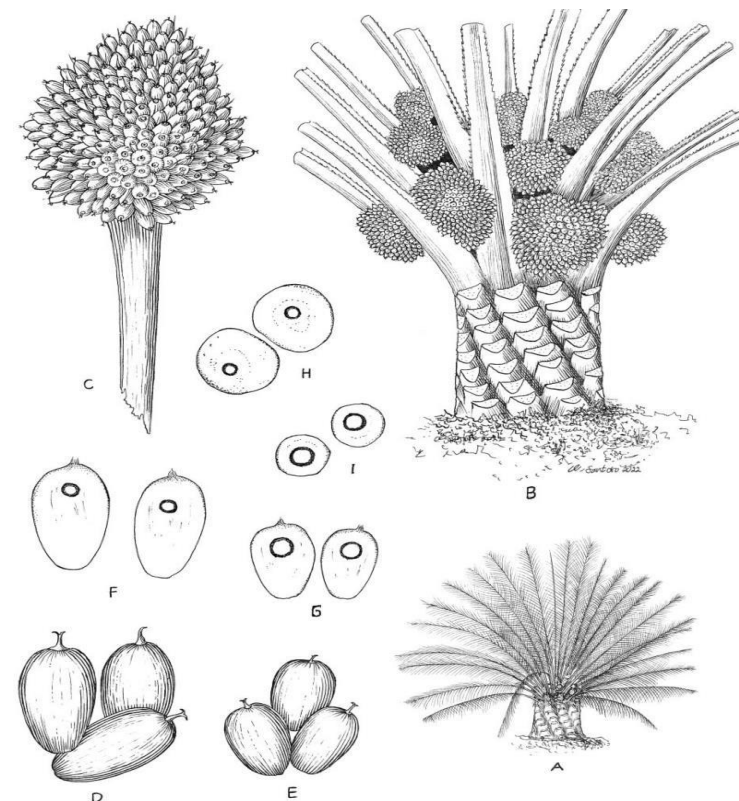
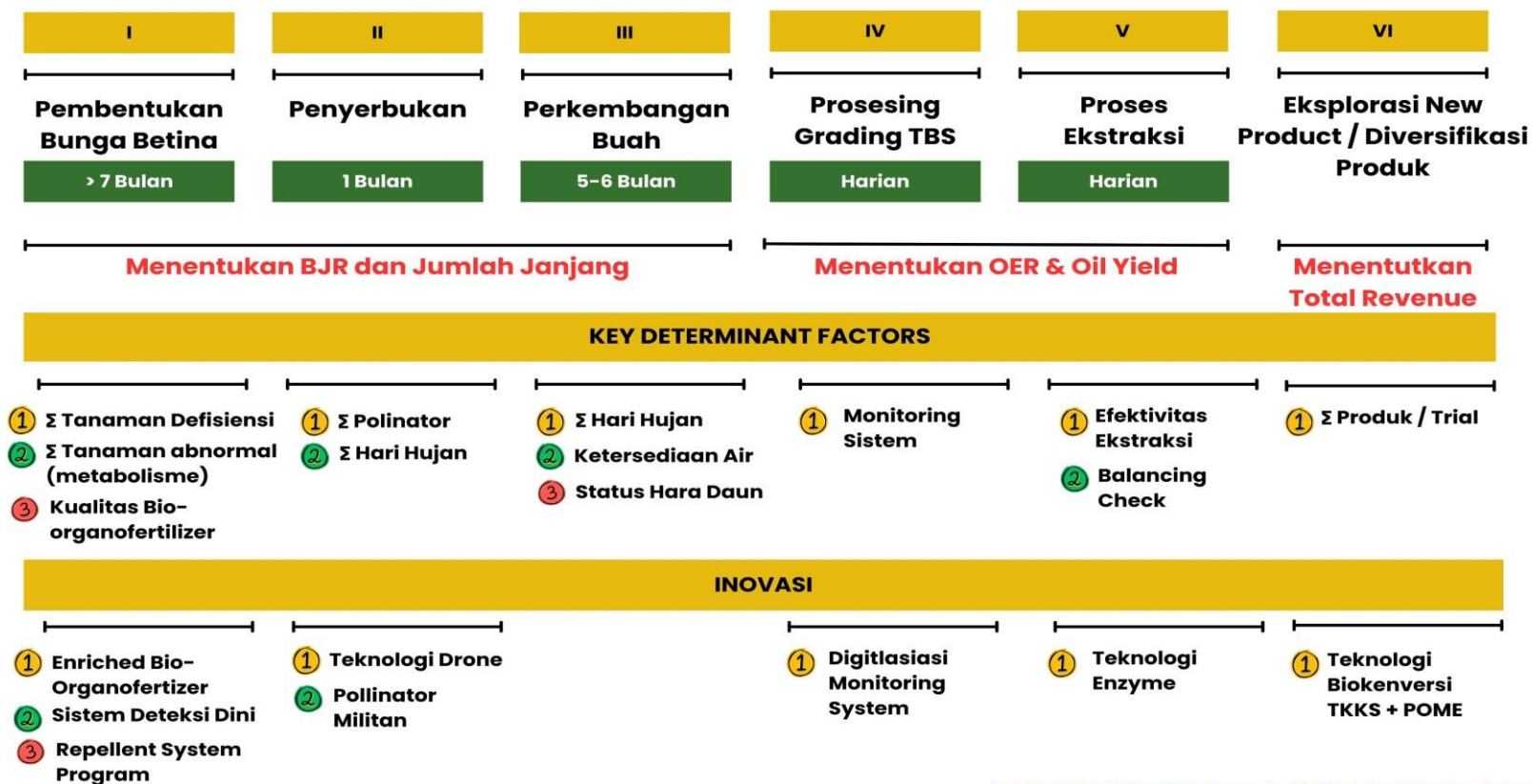
KOMPARASI CPO YIELD (Ton/Ha)



TANTANGAN INDUSTRI KELAPA SAWIT



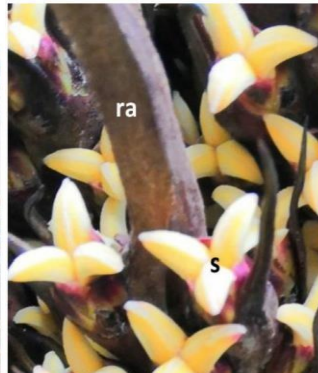
PENINGKATAN TBS & CPO MERUPAKAN URGENSI SAAT INI



7 HARI



7 HARI



2 HARI

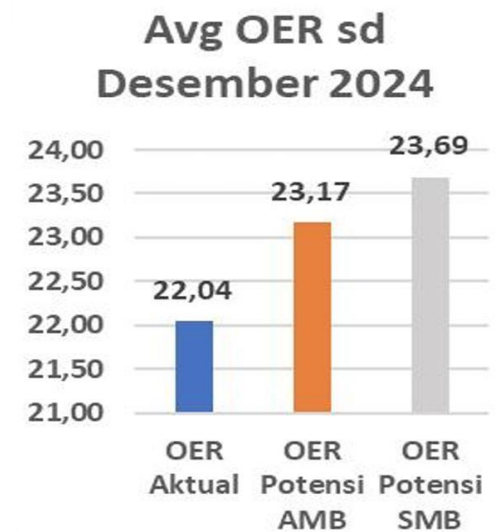
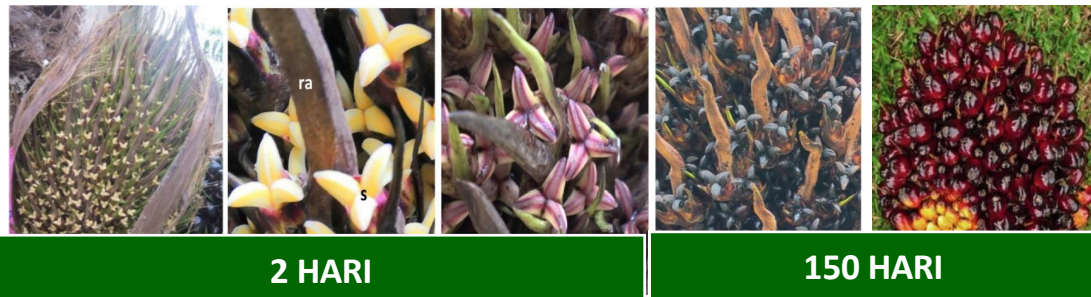


150 HARI

PENINGKATAN TBS & CPO MERUPAKAN URGENSI SAAT INI

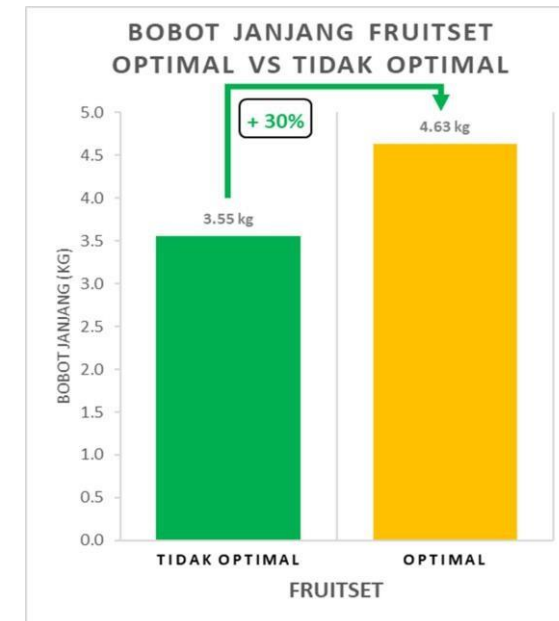
FOKUS INOVASI : BAGAIMANA MENINGKATKAN OIL CONTENT?

1. Meningkatkan jumlah bunga betina yang terbentuk
2. Meningkatkan kuantitas pollen pada inflorescens bunga jantan
3. Meningkatkan populasi serangga pollinator dan agresivitasnya untuk berkunjung ke bunga betina
4. Memastikan penyerbukan terjadi
5. Meningkatkan Fruit to Bunch (Fruitset & Modifikasi bentuk buah)
6. Meningkatkan Oil to Dry Mesocarp
7. Meningkatkan Rasio Mesocarp to Fruit
8. Meningkatkan ekspresi gen-gen yang terkait dalam biosintesis minyak

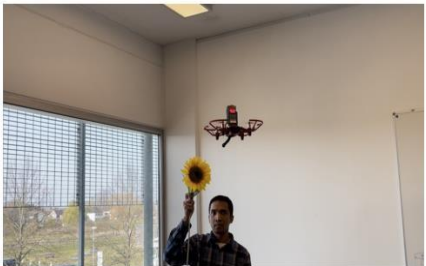
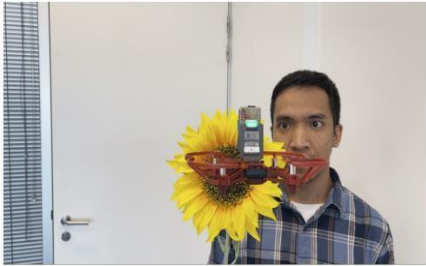
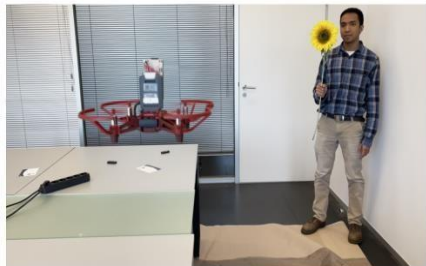
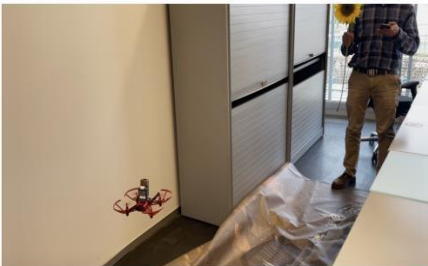


Luas dan Populasi Kelapa sawit BGA

Region	Luas	Populasi
Mentaya	30,566	3,828,131
Pundu	44,608	5,680,816
Kotawaringin	29,810	4,005,409
Kendawangan	20,930	2,835,198
Sei Rasau	24,127	3,308,801
Nanga Tayap	23,906	3,021,393
Rokan Hulu	2,299	314,098
Sei Melayu	11,174	1,471,007
TOTAL	187,420	24,464,853



Autonomus Pollination Drone for the future Agriculture



Differential Metabolite Profiles during Fruit Development in High-Yielding Oil Palm Mesocarp

Huey Fang Teh^{1*}, Bee Keat Neoh¹, May Ping Li Hong¹, Jaime Yoke Sum Low¹, Theresa Lee Mei Ng¹, Nalisha Ithnin¹, Yin Mee Thang¹, Mohaimi Mohamed², Fook Tim Chew³, Hirzun Mohd. Yusof¹, Harikrishna Kulaveerasingam¹, David R. Appleton¹

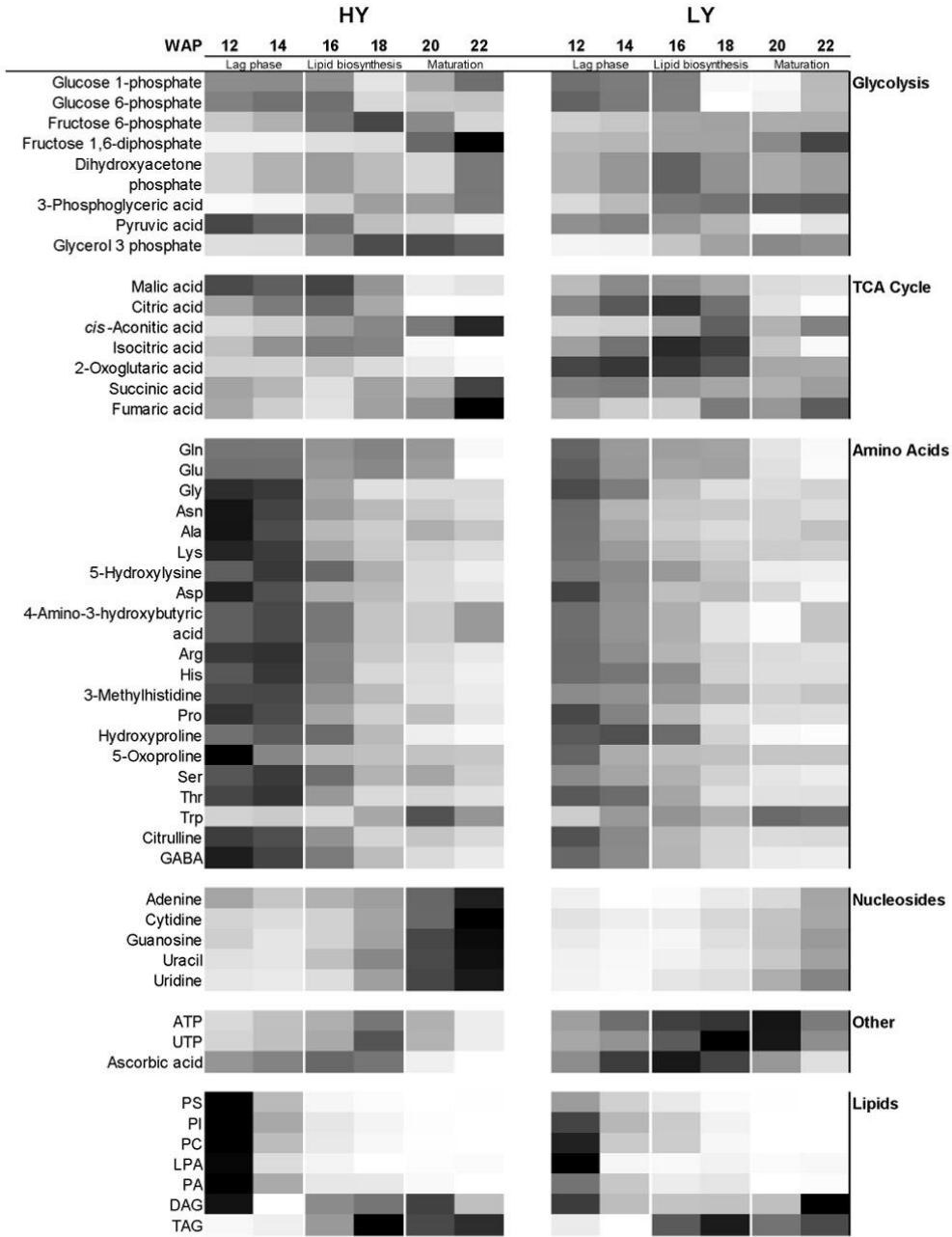
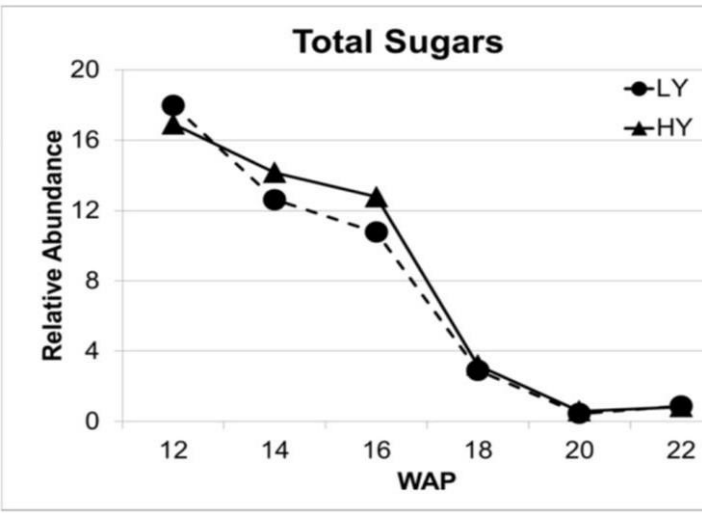
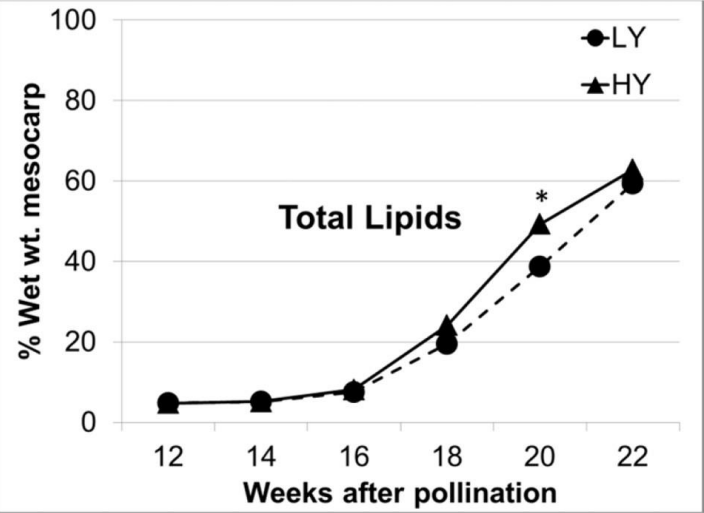
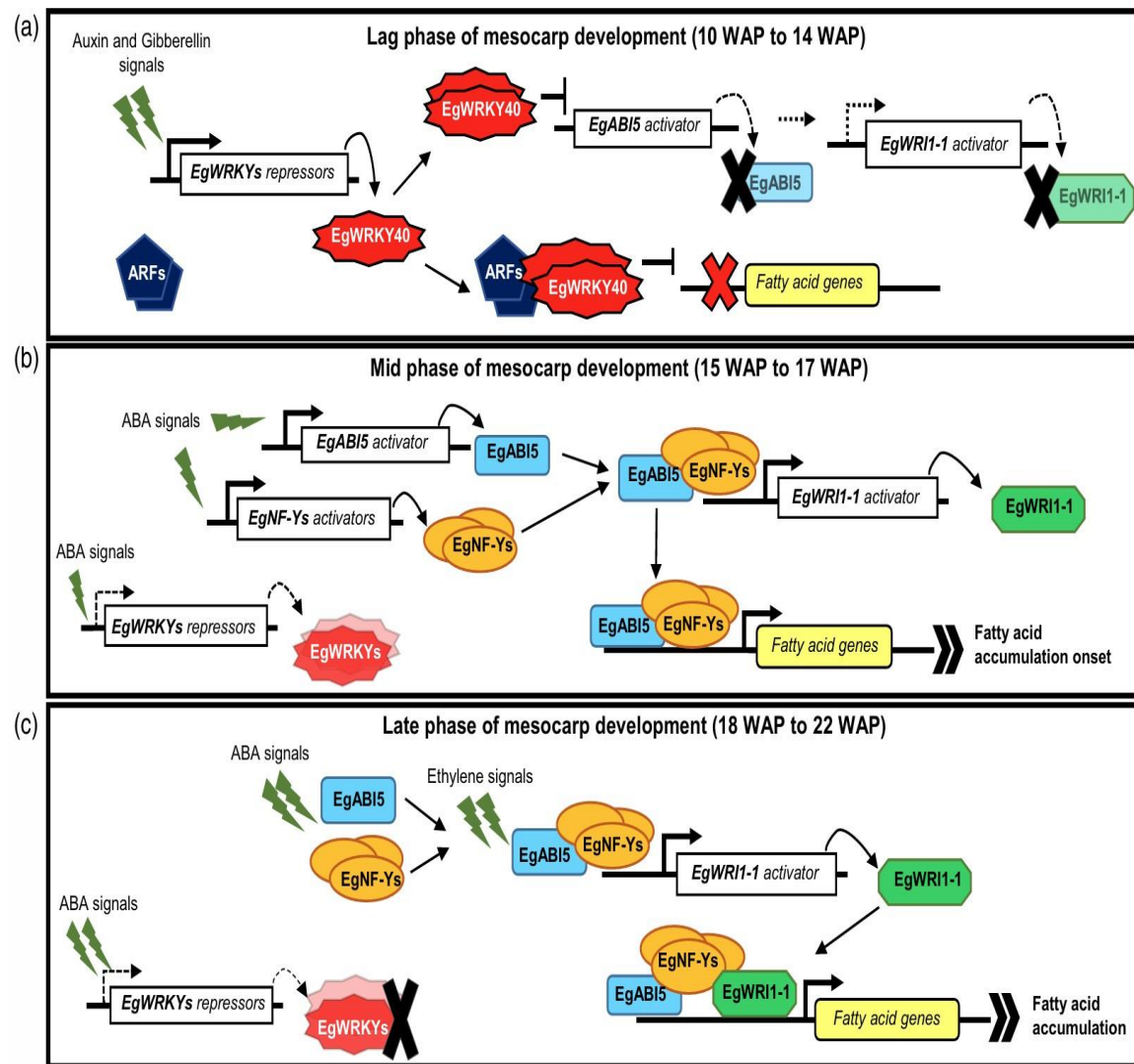


Figure 3. Heatmap of important primary metabolite concentrations during HY and LY palm mesocarp development. Heatmap of averaged and normalized mesocarp metabolite concentrations in key biosynthetic and cellular pathways of HY and LY oil palm groups at 12-22 WAP (n=8). Scale from white (lowest concentration) to black (highest concentration) within each metabolite class. doi:10.1371/journal.pone.0061344.g003



WRI1-1, ABI5, NF-YA3 and NF-YC2 increase oil biosynthesis in coordination with hormonal signaling during fruit development in oil palm

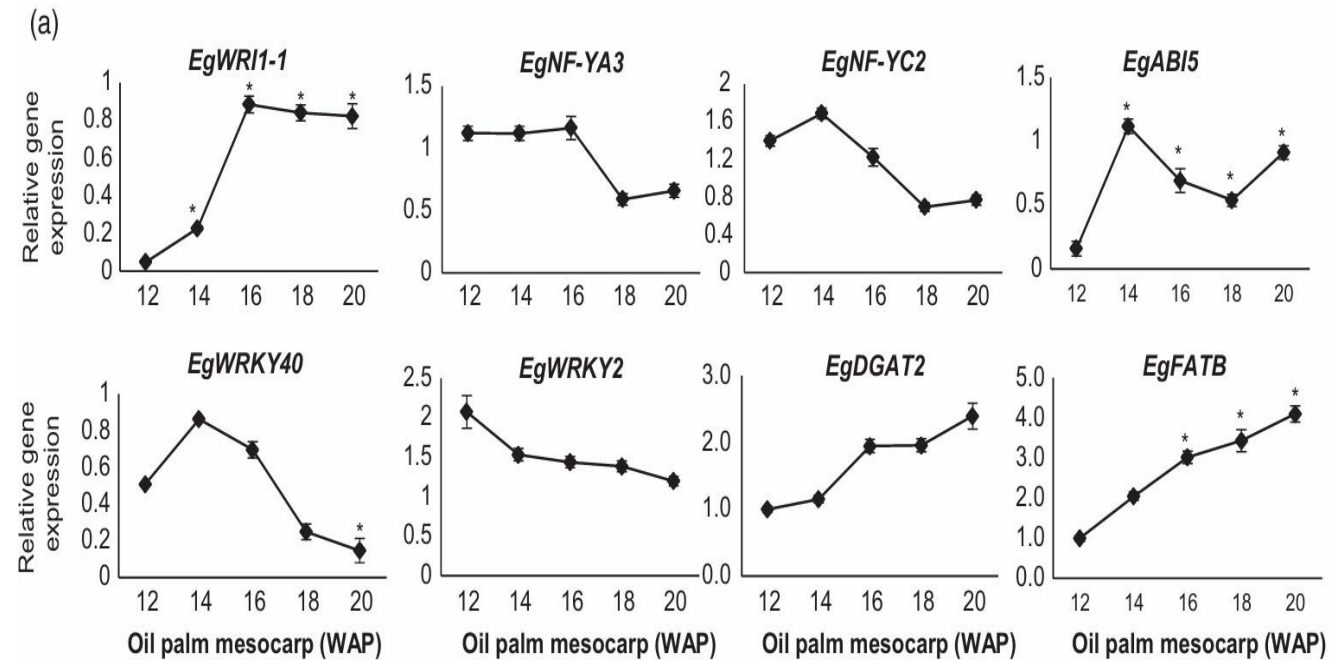


Figure 7. Regulatory model for oil biosynthesis in oil palm mesocarp.

(a) *EgWRKY2* and *EgWRKY40* interact with *EgABI5* to repress its DNA-binding activity to *EgWRI1-1* at the lag phase of mesocarp development. Auxin signals trigger *EgWRKYs* expression that interacts with ARFs to repress transcription of fatty acid biosynthesis genes.

(b) Elevation of ABA signals during mid-phase of mesocarp development inhibits transcription of *EgWRKYs* repressors and triggers transcription activation of *EgABI5* and *EgNF-Ys* activators. The activators form a complex to activate transcription of *EgWRI1-1* and their downstream oil biosynthesis genes.

(c) High ABA and ethylene signaling levels trigger *EgWRI1-1* transcription. The *EgWRI1-1* is recruited into activation complexes containing *EgABI5* and *EgNF-Ys*, to activate the expression of oil biosynthesis genes, which consequently promotes oil accumulation in oil palm mesocarp. [Colour figure can be viewed at wileyonlinelibrary.com].

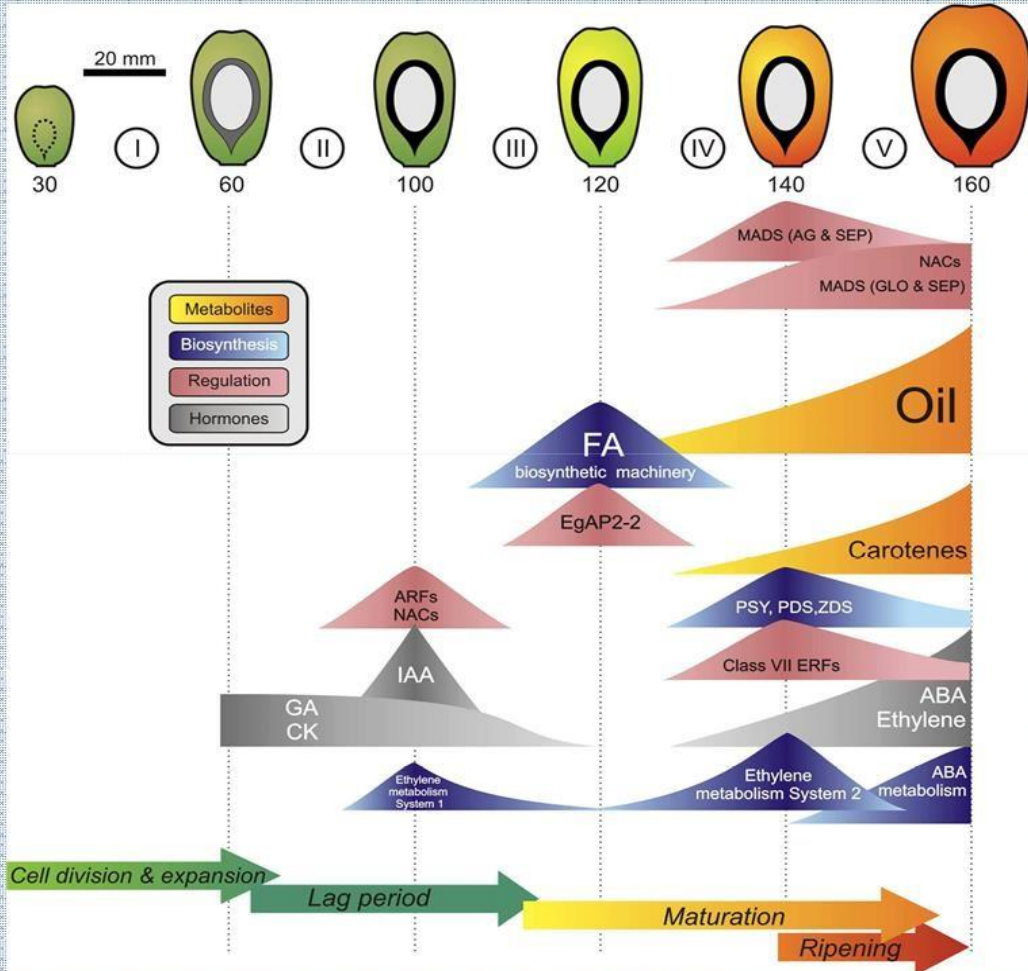


Figure 1. Scheme representing the major events that occur during the phases of mesocarp development.

Fruit development

- Morphological, histological, and biochemical (hormones, lipids) analyses were used to define phases of oil palm fruit development, maturation, and ripening (Fig. 1)
- Phase IV, between 120 and 140 DAP, is characterized by the onset of lipid and carotenoid accumulation
- Very low amounts of all hormones examined, including auxin, GA and cytokinin, ABA, and ethylene, were detected during phase IV
- Oil accumulates within subcellular spherical organelles (10–15 μ m in diameter, six to 12 per mesocarp cell) that occupy the whole volume of the cells (Fig. 2)
- Mesocarp cells also contain distinct regions, presumably chromoplasts, with high carotenoid concentrations (Fig. 2)
- This detailed morphological, histological and biochemical description of fruit development enabled to chose the stages for transcriptome 454 pyrosequencing

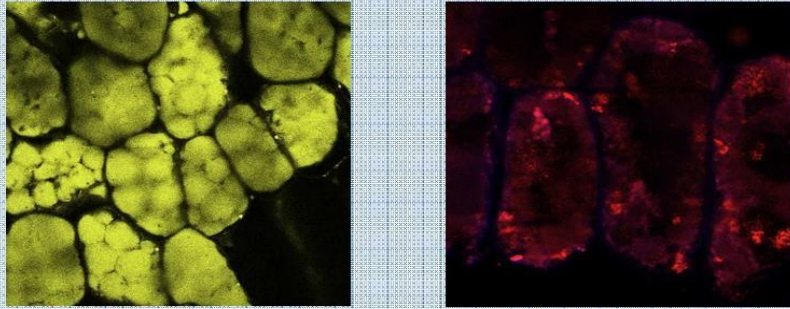


Figure 2. Confocal microscopy of mesocarp oil (left) and β -carotene (right)

Oil biosynthesis

- The core FA synthetic machinery of the oil palm mesocarp is remarkably coordinated at the transcriptional level (Fig. 3)
- The transcription peak of FA biosynthetic genes coincided with the onset of oil accumulation
- A transcript encoding a transcription factor similar to WRINKLED1 is highly expressed in the mesocarp during FA biosynthesis
- Canonical AW elements were found in several transcripts encoding enzymes involved in glycolysis and FA synthesis
- FA synthesis in the plastid and TAG assembly in the ER seem to be governed by two different transcriptional programs
- The last acylating step of TAG assembly may involve several complementary routes

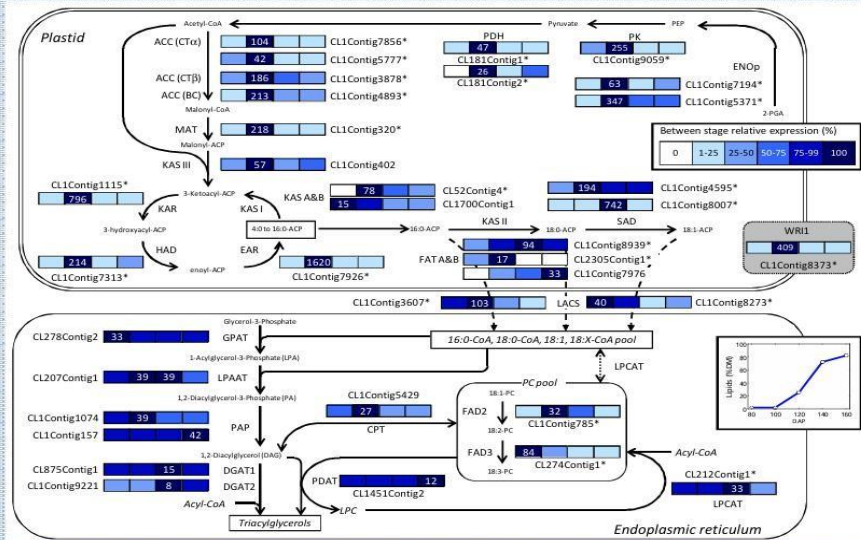


Figure 3. FA synthesis and TAG assembly in the oil palm mesocarp as reconstructed from the 454 pyrosequencing transcriptome. Gene expression levels at 100, 120, 140, 160 DAP are indicated with colored bars. * = significant differential expression

Contact Person



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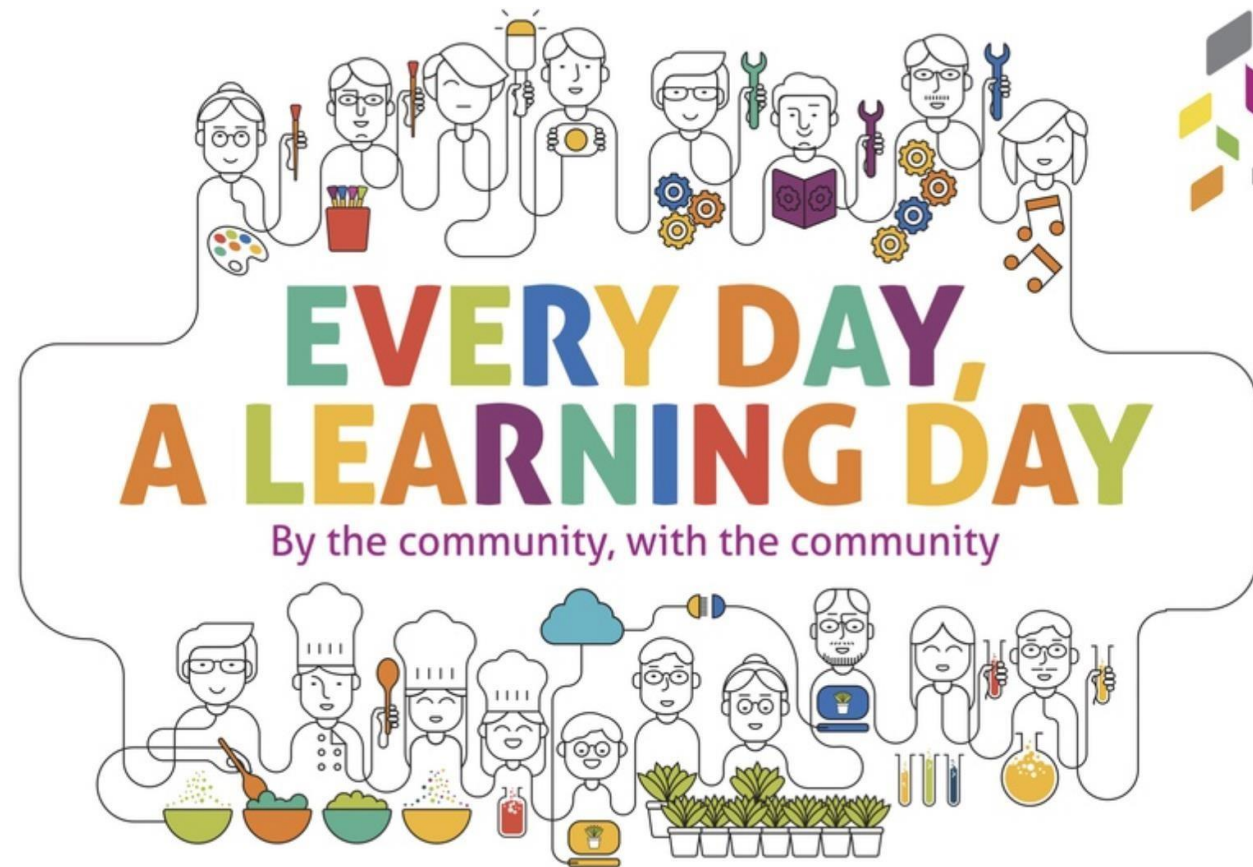
Abid Zulfikar

Corporate Planning &
Continuous Improvement



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THANK YOU



TOGETHER WE CAN

