



Wood Identification Tool Development and Application: *Examples from Indonesia*

Presented by

Iskandar Z Siregar

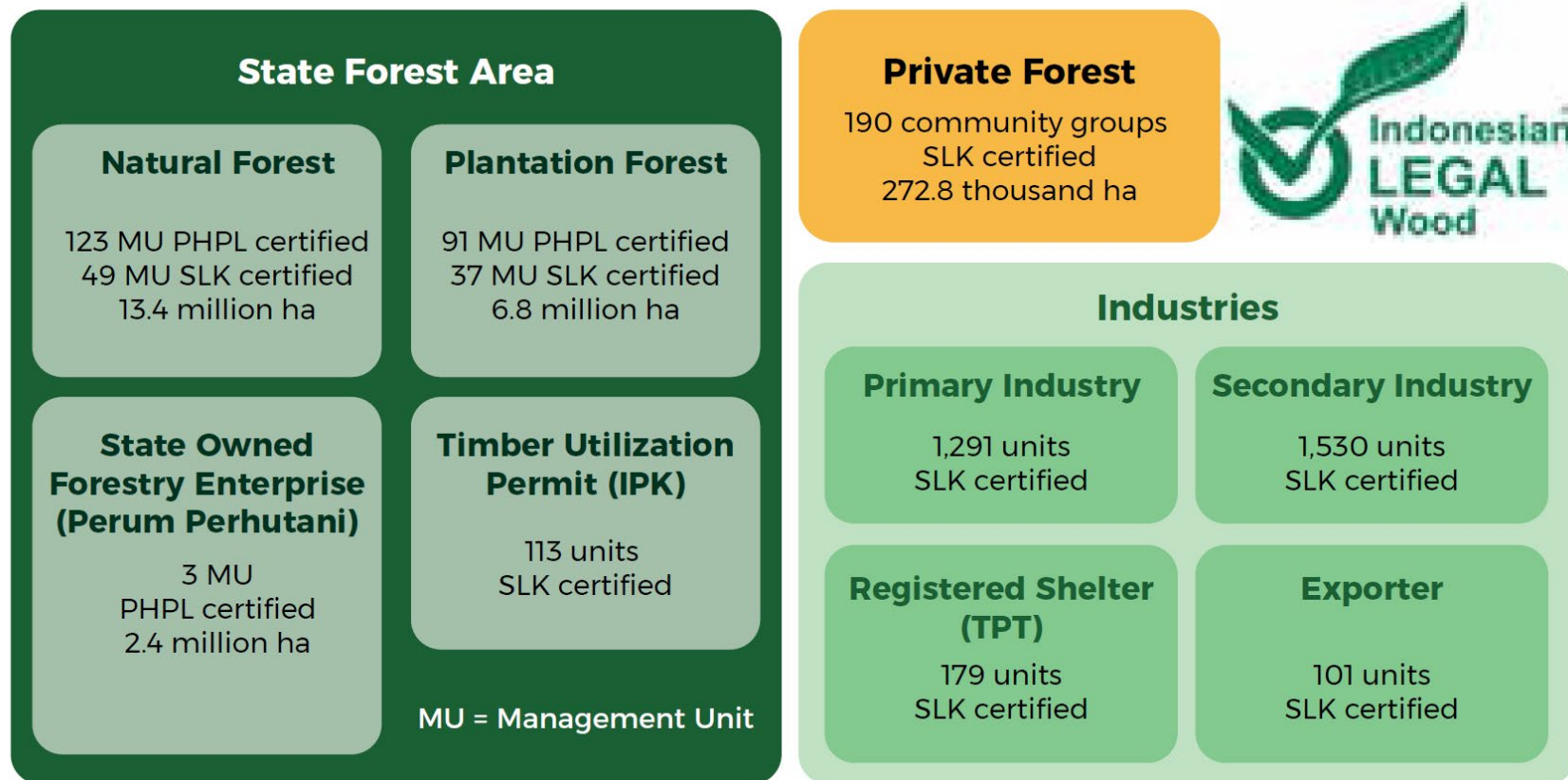
Department of Silviculture, Faculty of Forestry

IPB University, Bogor, Indonesia

Email: siregar@apps.ipb.ac.id

Introduction

Progress – SFM Certification



Progress of PHPL (Management of Sustainable Production Forest) and SLK (Certification of Timber Legality) certification, as of December 2017 (DJPHPL-MoEF, 2018)

Introduction

Timber Legality System - SVLK

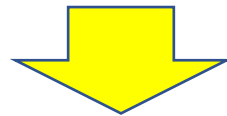


■ Introduction

Tree Genetics and Wood DNA Studies in Indonesia

- Red Meranti
- Ironwood
- Ebony

- Population genetics
- DNA Barcodes



Technology for determining origin and species of timber??



Red Meranti & Ironwood Research

Population genetics, DNA Barcodes & Wood DNA

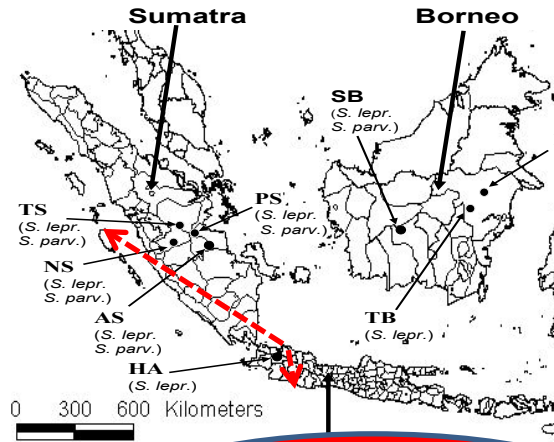


Results

Population Genetic Studies#1

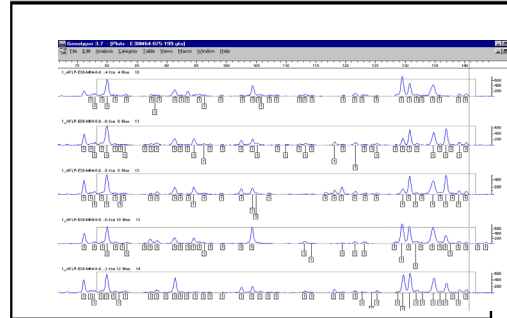
Red Meranti (*Shorea leprosula*)

AFLP Markers



Transfer of materials (seeds): Siberut, Sukabumi

FOERDAI-Haurbentes Experimental Forest (He=0.186)



AFLP Fragments

S. lepr= *Shorea leprosula*
S. parv= *Shorea parvifolia*

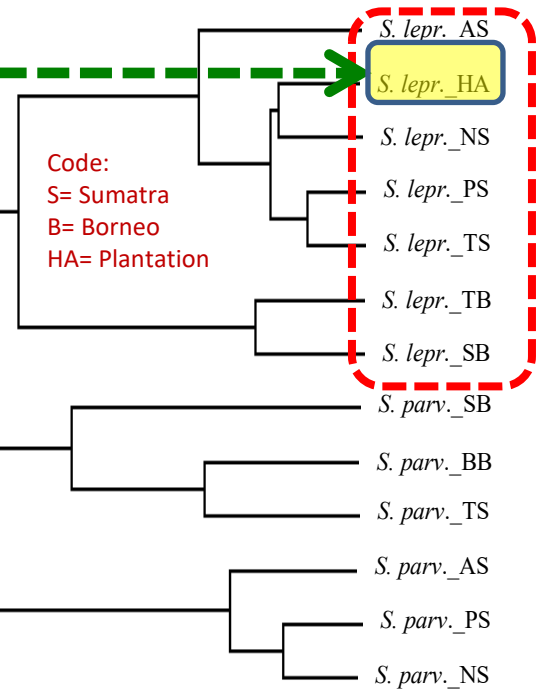


Table 2 Genetic diversity within populations of *S. leprosula* and *S. parvifolia*

Population	Sample size	Polymorphic loci	PPL (%)	n_a	n_e	H_e	H'_e	I
<i>S. lepr</i> _HA	34	34	60.71	1.607	1.318	0.186	0.204	0.280
<i>S. lepr</i> _TB	23	26	40.43	1.404	1.184	0.113	0.143	0.182
<i>S. lepr</i> _AS	9	37	66.07	1.661	1.355	0.208	0.258	0.316
<i>S. lepr</i> _PS	1	20	51.20	1.512	1.380	0.163	0.200	0.248
<i>S. lepr</i> _TS	22	28	50.00	1.500	1.281	0.160	0.177	0.238
<i>S. lepr</i> _NS	16	28	50.00	1.500	1.250	0.151	0.184	0.232
<i>S. lepr</i> _SB	18	27	48.21	1.482	1.249	0.145	0.169	0.221
Mean	19	30	53.32	1.533	1.274	0.161	0.191	0.245
Total	133	52	92.86	1.929	1.347	0.211	0.212	0.330
SD				0.260	0.352	0.184	0.251	
<i>S. parv</i> _BB	16	19	33.93	1.339	1.201	0.115	0.154	0.171
<i>S. parv</i> _AS	33	43	76.79	1.768	1.371	0.222	0.238	0.337
<i>S. parv</i> _PS	23	33	58.93	1.589	1.228	0.143	0.167	0.228
<i>S. parv</i> _TS	14	21	37.50	1.375	1.159	0.097	0.137	0.152
<i>S. parv</i> _NS	28	28	50.00	1.500	1.189	0.119	0.143	0.188
<i>S. parv</i> _SB	21	30	53.57	1.536	1.211	0.135	0.161	0.214
Mean	22.5	29	51.79	1.518	1.227	0.138	0.167	0.215
Total	135	48	85.71	1.857	1.336	0.205	0.204	0.319
SD				0.353	0.353	0.185		0.256

PPL Percentage of phenotypically polymorphic loci, n_a observed number of alleles per locus, n_e effective number of alleles per locus, H_e Nei's (1973) gene diversity, I Shannon's information index (Lewontin 1972), SD standard deviation of total values H'_e Nei's gene diversity estimated with the computer program AFLP-SURV 1.0 (Vekemans 2002)

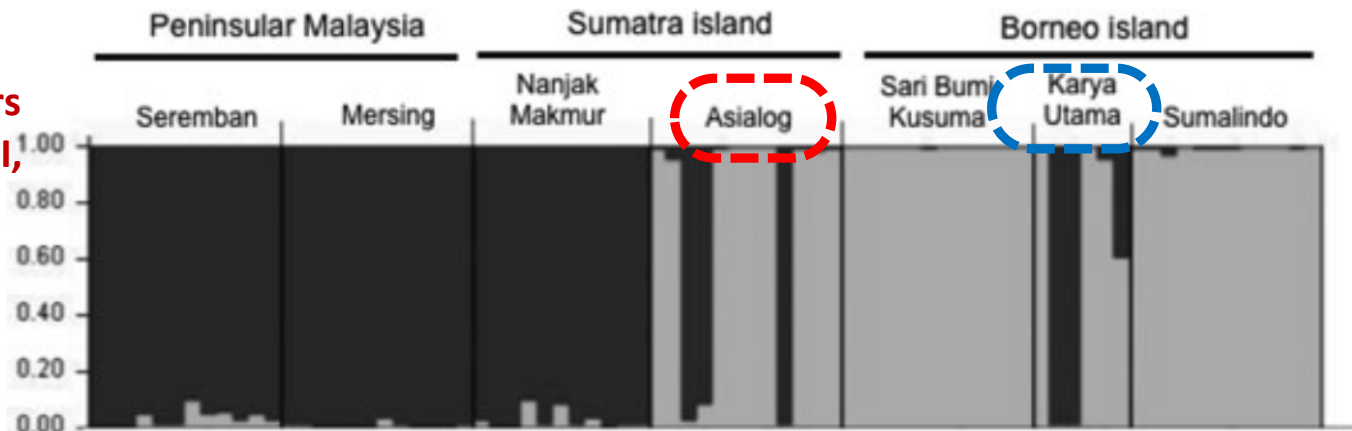
Center of genetic diversity for *S. leprosula* in ex Asia Log Concession (PT. REKI), Sumatra



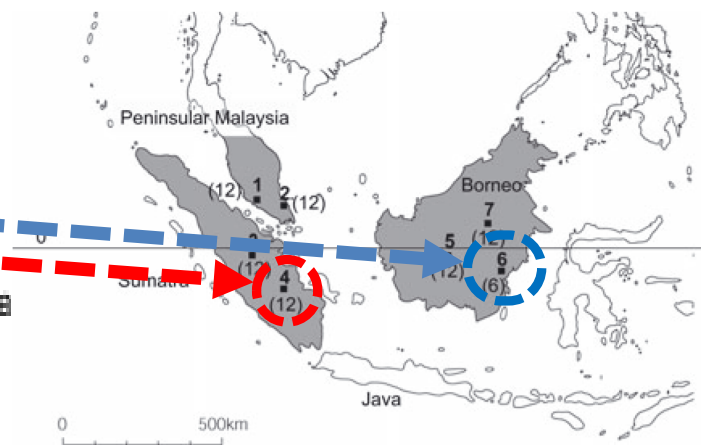
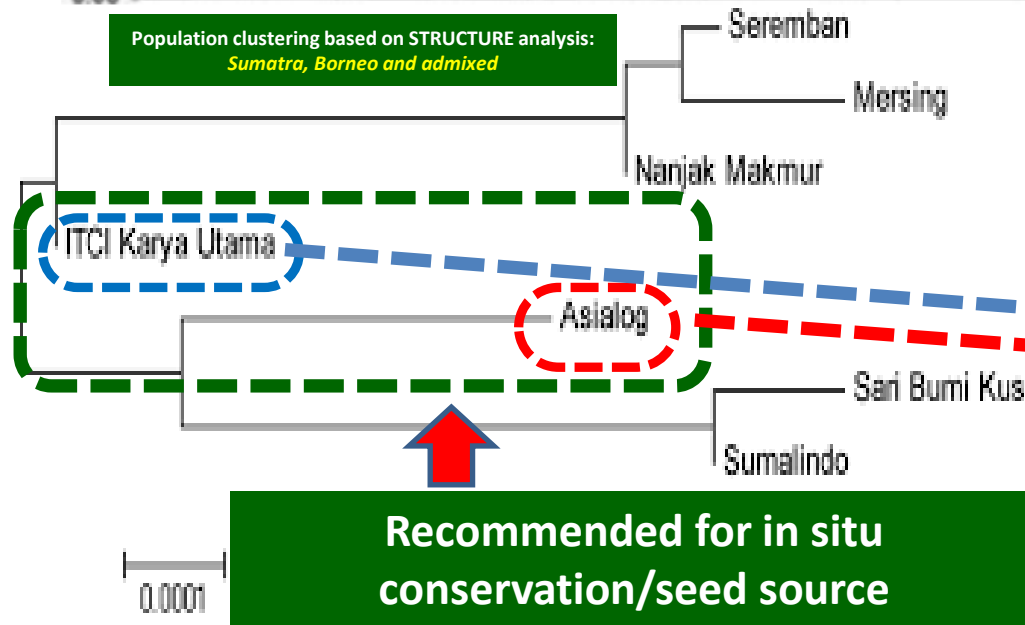
Shorea parvifolia

Results

Five nuclear gene markers (GapC, GBSSI, PgiC, SBE2 and SODH)



Population Genetic Studies#2

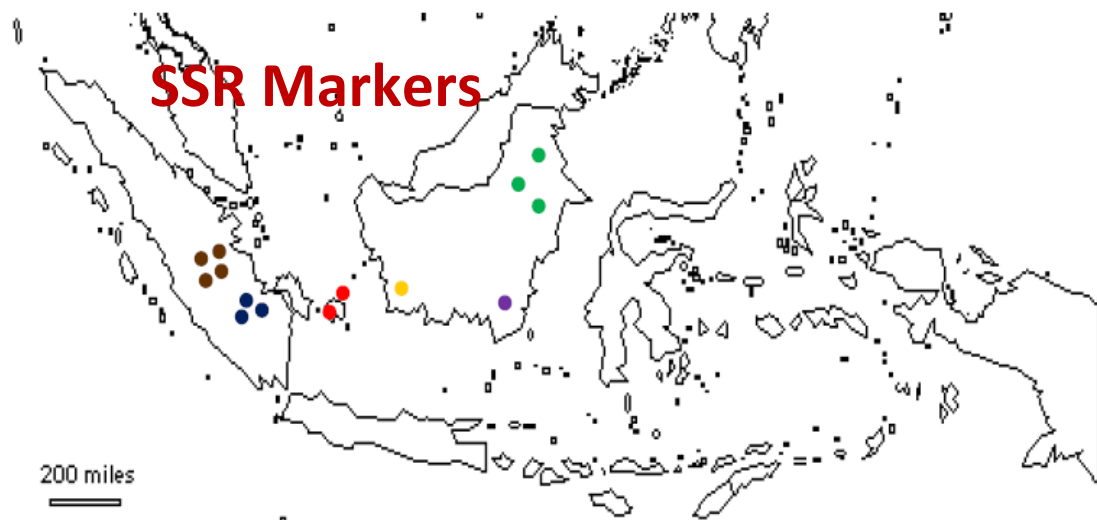


Iwanaga et al. (2012)

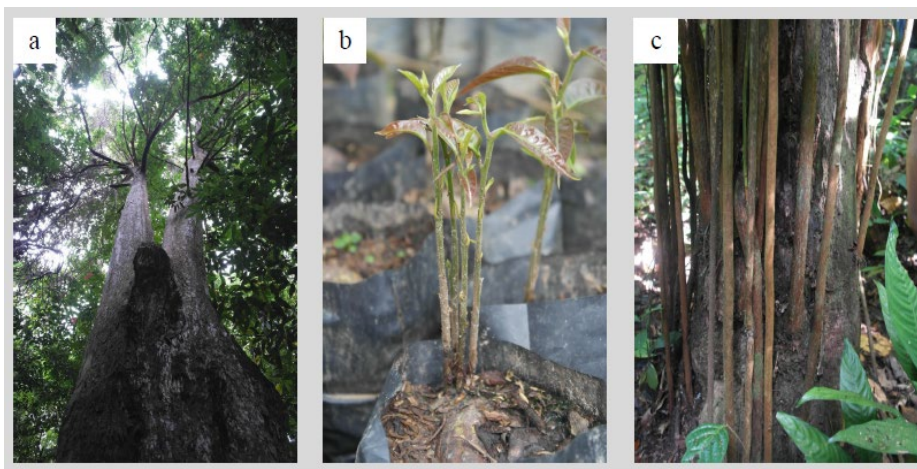
Results

Population Genetic Studies#3

Ironwood (*Eusideroxylon zwageri*)



Symbol	Provinces	No.population
●	Jambi	4 populations
●	South sumatra	3 populations
●	Belitung	2 populations
●	East Kalimantan	3 populations
●	West Kalimantan	1 population
●	South Kalimantan	1 population



Purba et al. (unpublished)

- Collection of leaves from mature trees and seedlings for DNA extraction throughout the natural distribution area in Indonesia (n= 1048 samples)
- Ten suitable genetic markers (SSR) have been developed
- Population structure of ironwood (*E. zwageri*) were determined by analyzing the level of genetic differentiation within and among populations

Results

The objective is to generate DNA sequences of vascular plant species in Sumatra and combine it with morphological analysis in order to provide a reliable species identification tool for vascular plants in tropical forest

DNA Barcodes

DNA barcode markers (matK and rbcL); n= 5100 samples; > 1100 species

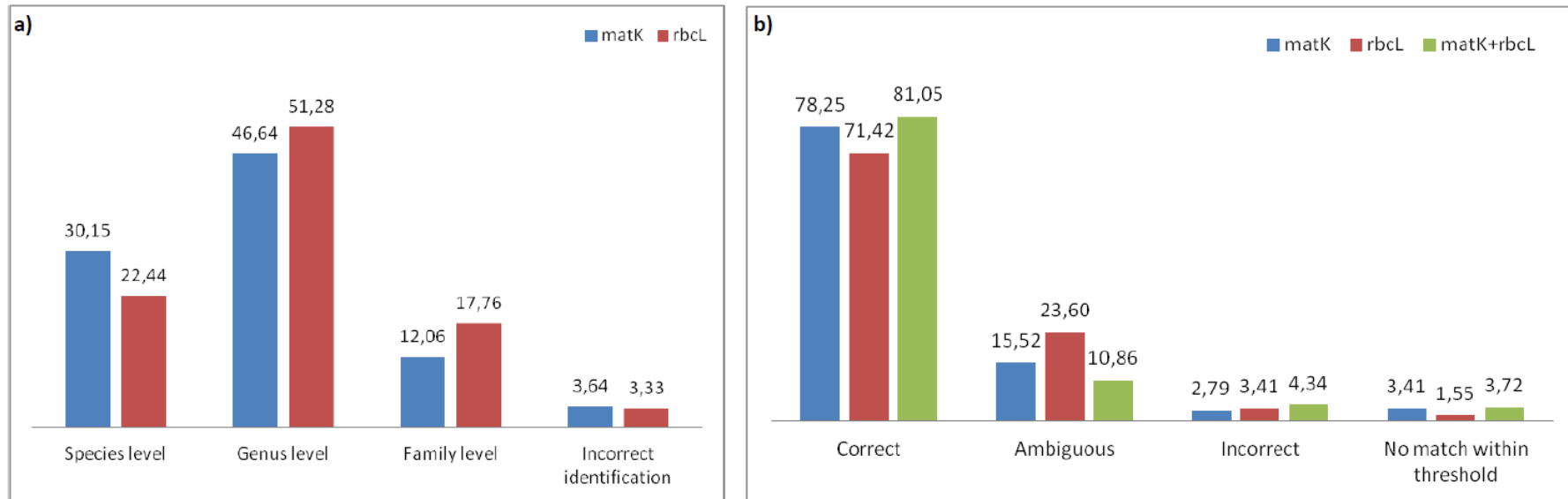


Fig 1. a) Level of match comparison between morphological identification and molecular identification (in percentage) with matK and rbcL as markers, b) Identification success of DNA barcodes (in percentage) using TaxonDNA with matK, rbcL, and combination of both markers

Results

Shorea spp

DNA Isolation from Wood#1

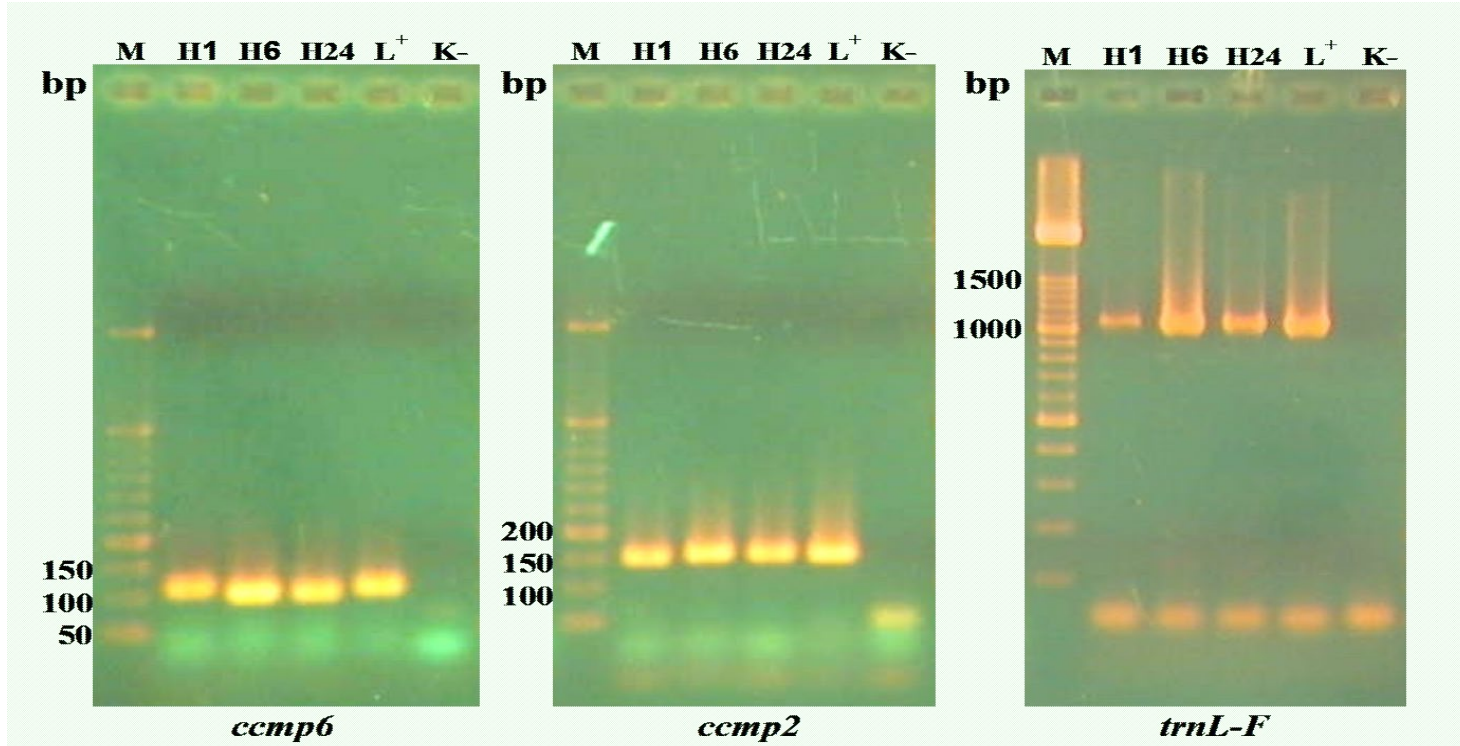


Figure 1. PCR profiles of the DNA samples amplified with primers *trnL-F* (Taberlet et al., 1991), *ccmp2* and *ccmp6* (Weising and Gardner, 1999).

Length of cpDNA fragment amplified by *ccmp6*, *ccmp2* and *trnL-F* was about 0.1, 0.15 and 1.1 kb, respectively. Samples on gel lanes are: M = Size standard; H1, H6 and H24 = wood DNA of Meranti (botanical name unknown), *Shorea leprosula* and *Shorea ovalis*, respectively; L⁺ = Positive control (leaf DNA); K⁻ = Negative control (water).

Results

Comparison of Two DNA Isolation Methods on 49 Wood Species

DNA Isolation from Wood#2

Nama Sampel	Sampel		Nama Sampel	Sampel	
	K	S		K	S
<i>Acacia mangium</i> Willd.	+	+	<i>Vitex sp.</i>	+	+
<i>Miconia eminei</i> Engler	+	+	<i>Miconia eminei</i> Engler	+	+
<i>Ochroma saramensis</i> Miq.	+	+	<i>Ochroma saramensis</i> Miq.	+	+
<i>Calophyllum macrocarpum</i> Hook.f.	+	+	<i>Calophyllum macrocarpum</i> Hook.f.	+	+
<i>Ochroma grandifolia</i> Kuntze	+	+	<i>Ochroma grandifolia</i> Kuntze	+	+
<i>Parosperum elongatum</i>	+	+	<i>Parosperum elongatum</i>	+	+
<i>Hopsea laevigata</i> (Poir.) Endert	+	+	<i>Hopsea laevigata</i> (Poir.) Endert	+	+
<i>Canthium coriandrinum</i> (Blanc.) R.A. Howard	+	+	<i>Canthium coriandrinum</i> (Blanc.) R.A. Howard	+	+
<i>Elaeagnus ovalis</i> (Miq.) Dandy	+	+	<i>Elaeagnus ovalis</i> (Miq.) Dandy	+	+
<i>Hopsea sampel</i> Koehrh.	+	+	<i>Hopsea sampel</i> Koehrh.	+	+
<i>Duroia ilicifolia</i> Muiray	+	+	<i>Duroia ilicifolia</i> Muiray	+	+
<i>Juglans sp.</i>	+	+	<i>Juglans sp.</i>	+	+
<i>Diospyros pilosulobata</i> Blanco	+	+	<i>Diospyros pilosulobata</i> Blanco	+	+
<i>Omeiopsis arborea</i> Koehrh.	+	+	<i>Omeiopsis arborea</i> Koehrh.	+	+
<i>Veronica arborea</i> Back. Han.	+	+	<i>Veronica arborea</i> Back. Han.	+	+
<i>Tecoma grandis</i> L.f.	+	+	<i>Tecoma grandis</i> L.f.	+	+
<i>Cassia siamensis</i> Lank.	+	+	<i>Cassia siamensis</i> Lank.	+	+
<i>Koompassia malaccensis</i> ex Benth.	+	+	<i>Koompassia malaccensis</i> ex Benth.	+	+
<i>Sandoricum lasiagae</i> (Horn.) Merr.	+	+	<i>Sandoricum lasiagae</i> (Horn.) Merr.	+	+
<i>Dryobalanops aromatica</i> C.F. Gaertn.	+	+	<i>Dryobalanops aromatica</i> C.F. Gaertn.	+	+
<i>Alseodaphne moluccana</i> (L.) W.Mill.	+	+	<i>Alseodaphne moluccana</i> (L.) W.Mill.	+	+
<i>Alseodaphne moluccana</i> (L.) W.Mill.	+	+	<i>Alseodaphne moluccana</i> (L.) W.Mill.	+	+
<i>Diospyros conferta</i> Skoott	+	+	<i>Diospyros conferta</i> Skoott	+	+
<i>Terminalia balata</i> (Roxb.) Steud.	+	+	<i>Terminalia balata</i> (Roxb.) Steud.	+	+
<i>Falcataria moluccana</i> Blume	+	+	<i>Falcataria moluccana</i> Blume	+	+



(Ramdhani & Siregar, 2017)

49 wood species

Methods	Wood sample types	DNA markers (barcodes)	Total samples	Amplification success (%)
CTAB	Dry	ITS	49	21 (43%)
		rbcl		10 (20%)
	Fresh	ITS	20	20 (100%)
		rbcl		20 (100%)
QDPMK (Qiagen)	Dry	ITS	49	23 (47%)
		rbcl		13 (27%)
	Fresh	ITS	20	19 (95%)
		rbcl		19 (95%)

BIODIVERSITAS
Volume 20, Number 6, June 2019
Pages: 1748-1752

ISSN: 1412-033X
E-ISSN: 2085-4722
DOI: 10.13057/biodiv.d2006055

Short Communication: DNA extraction from stored wood of *Falcataria moluccana* suitable for barcoding analysis

HASYATI SHABRINA^{1*}, ULFAH J. SIREGAR², DEDED D. MATRA³, KOICHI KAMIYA⁴,
ISKANDAR Z. SIREGAR^{5*}

¹Tropical Silviculture Program, Department of Silviculture, Faculty of Forestry, Institut Pertanian Bogor, Jl. Ulin, Dramaga Campus, Bogor 16680, West Java, Indonesia. Tel/fax +62 251 8626806. *email: hasyati_shabrina@apps.ipb.ac.id
²Department of Silviculture, Faculty of Forestry, Institut Pertanian Bogor, Jl. Ulin, Dramaga Campus, Bogor 16680, West Java, Indonesia. Tel/fax +62 251 8626806. *email: siregar@apps.ipb.ac.id, iszaregar@yahoo.com
³Department of Agronomy and Horticulture, Faculty of Agriculture, Institut Pertanian Bogor, Jl. Meranti, Dramaga Campus, Bogor, West Java, Indonesia
⁴Forest Resource Department, Faculty of Agriculture, Ehime University, 3-5-7 Tarumi, Matsuyama, Ehime Prefecture, Japan

Manuscript received: 24 February 2019. Revision accepted: 29 May 2019.

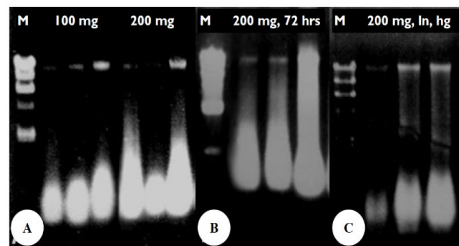


Figure 1. DNA extracted from A) 0.1 (lane 2-4) and 0.2 (lane 5-7) g tissue, frozen 24 hours, B) 0.1 g tissue, frozen 72 hours, and C) 0.2 g tissue with liquid nitrogen and grind using mortar. M: A HindIII

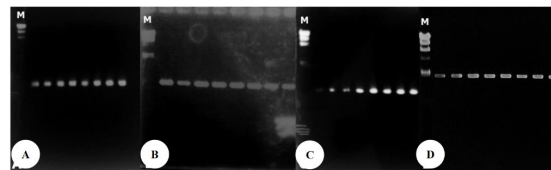


Figure 2. The amplification results using *psbA-trnH* intergenic spacer primer with DNA from previous extraction methods as template: A) 100 mg frozen for 24 hours in 30 °C; B) 200 mg frozen for 24 hours in 30 °C; C) 100 mg frozen for 72 hours in 30 °C; D) 200 mg frozen with liquid nitrogen; M: 1 kb DNA ladder.

(Shabrina et al., 2019)

Falcataria moluccana

Table 1 The nanophotometer result of DNA extracted from all methods used

Starting material	Freezing method	Disrupting method	Maximum DNA yield (ng/μL)	A ₂₆₀ /A ₂₈₀ ratio
100 mg	24 hrs; -30°C	TissueLyser	257.80	1.87-2.25
200 mg	24 hrs; -30°C	TissueLyser	140.45	1.89-2.86
100 mg	72 hrs; -30°C	TissueLyser	214.35	1.91-4.38
200 mg	Liquid nitrogen	Hand grinding	174.10	1.48-2.84

■ Summary

- Focus on populations of widespread species for **baseline genetic information** only (e.g. genetic structure, diversity) → genetic database development
- Application mostly for identifying “**center of genetic diversity**” → FGR conservation strategy
- Wood samples for DNA isolation is **not standard** (methods, size etc) → only for genetic analysis purpose not for others



Ebony Research in Indonesia



Ebony Research

Progress and Preliminary Results

Iskandar Z. Siregar, Fifi Gus Dwiyantri, Essy Harnelly, Muhammad Majiidu, Lina Karlinasari, Ratih Damayanti, M Rafi, Meaghan Parker-Forney



WORLD
RESOURCES
INSTITUTE



Introduction

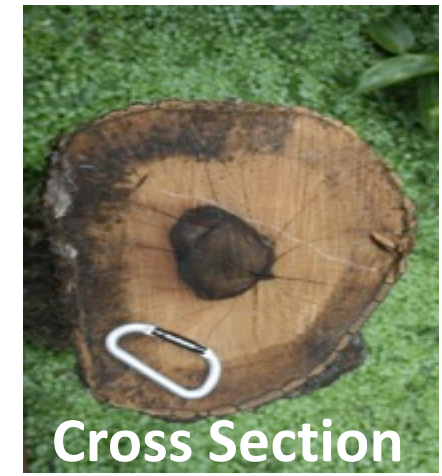
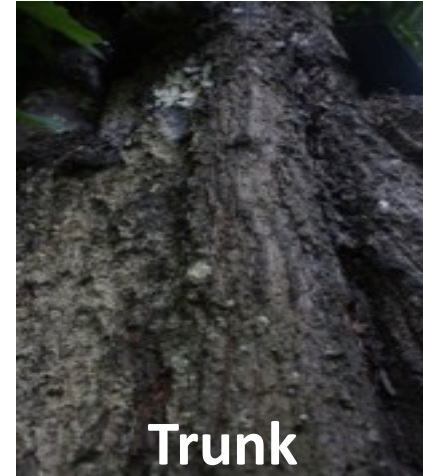


Diospyros celebica
(Ebony)

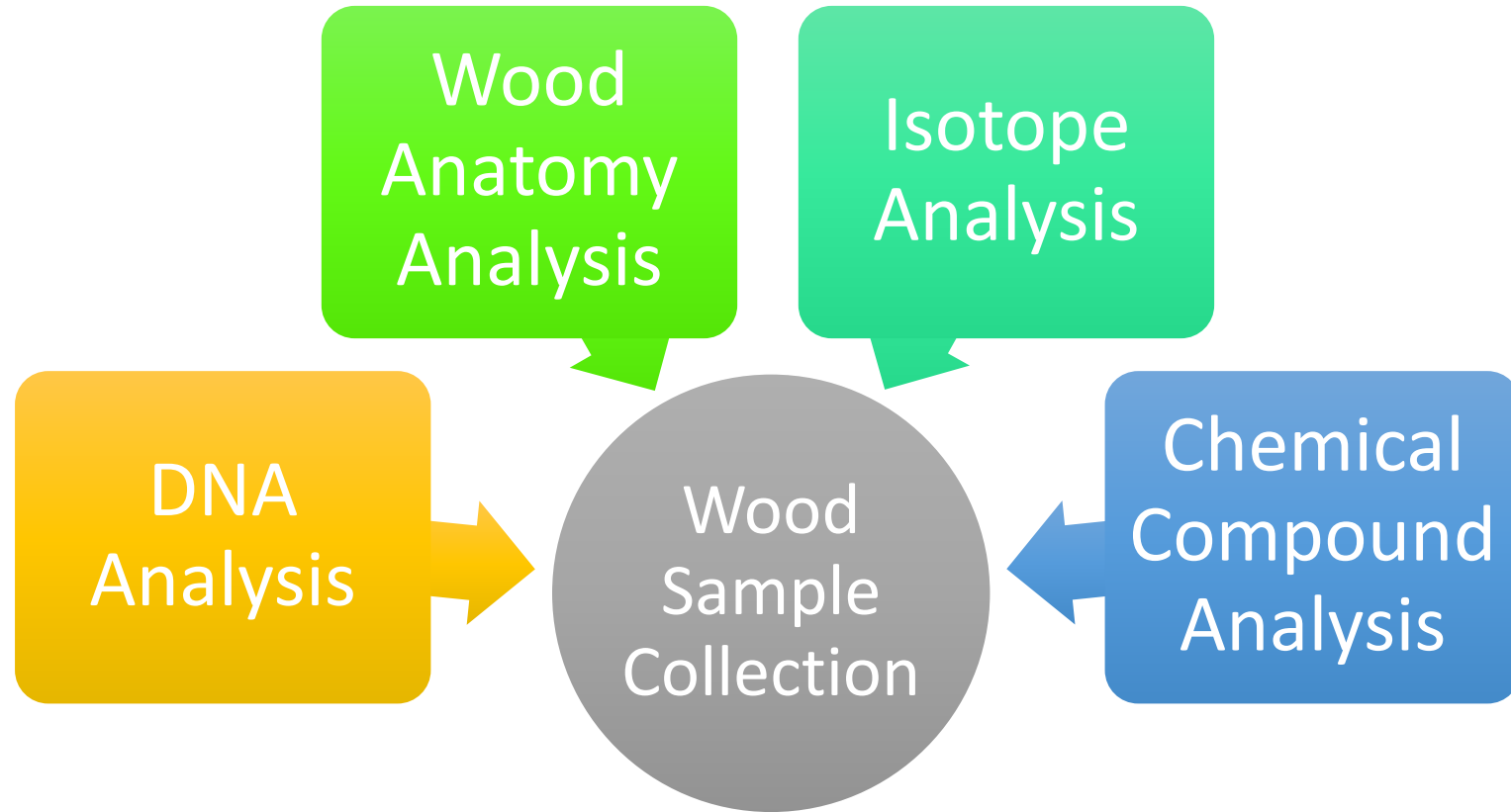
Djarwanto *et al.* 2017

■ Introduction

- Endemic to Sulawesi Island (South Sulawesi, West Sulawesi, Central Sulawesi and North Sulawesi)
- Slow growing species
- Vulnerable species (IUCN 2019)
- Timber for carving, inlay, furniture and musical instruments
- Problem: **Illegal logging** → Wood identificatin tools?



Methods



Objectives

- **Main objective:** To setting up a reference data building pipeline for DNA of commercial timber species, *Diospyros celebica* Bakh. (Ebony)
- **Specific objectives:** To collect physical timber reference material and extract its associated DNA for the species *Diospyros celebica* Bakh. (Ebony).





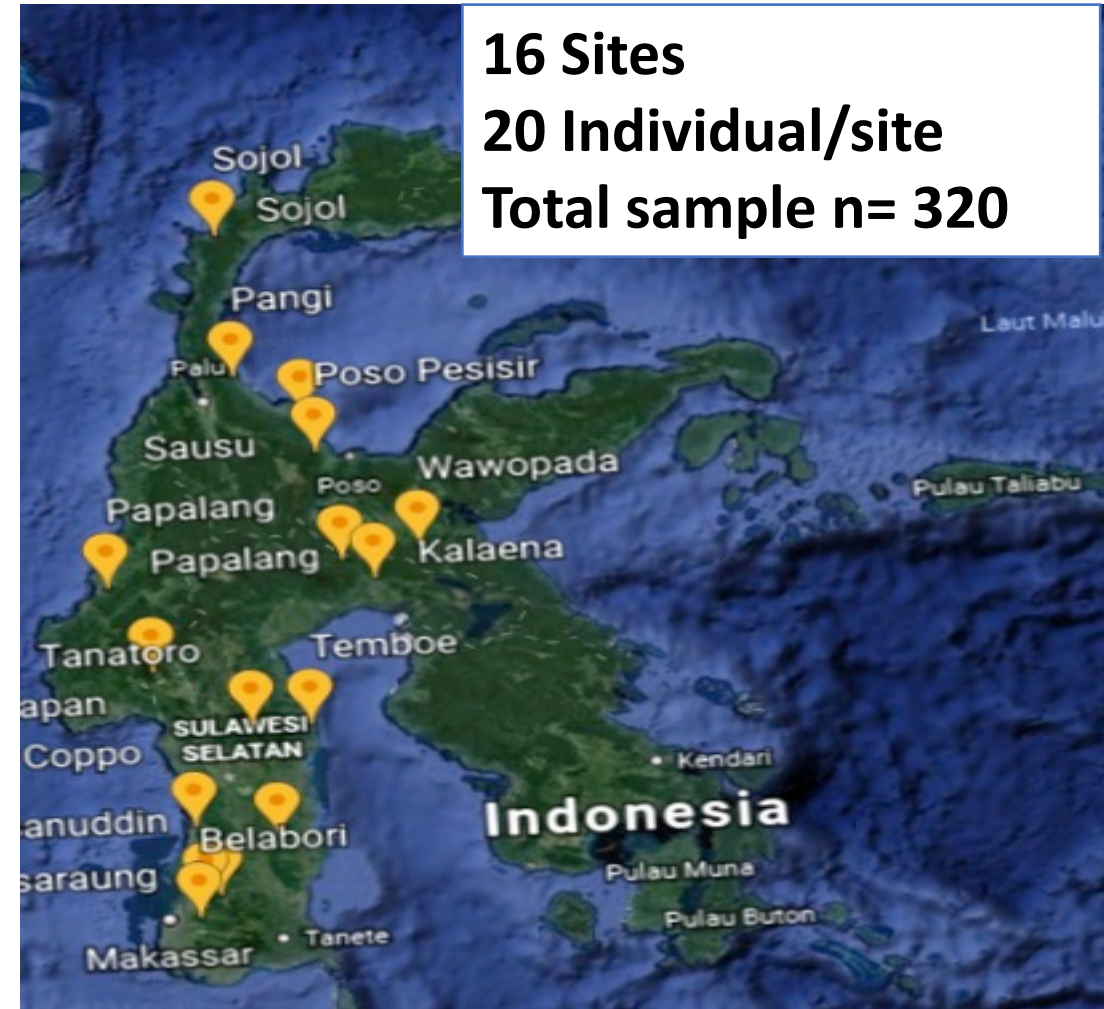
Activity#1:
Wood Collection (Concentrated Species:
***Diospyros celebica*) in Sulawesi**

Objective

To update the current distribution of *Diospyros celebica* in Sulawesi including wood sample collection

D. celebica Wood Sampling Sites

No	Province	Regency	Sub-district	Sampling Sites
1	South Sulawesi	Maros		Batimurung Bulusaraung National Park 
				Hasanuddin University Teaching Forest
		Barru	Barru	Coppo Village
		Sidenreng Rappang (Sidrap)	Pituriase	Tana Toro Protection Forest
		Gowa	Parangleo	Bellabori
		Luwu Timur	Mangkutana	Ponda-Ponda Nature Reserve 
			Mangkutana	Kalaena Nature Reserve
			Mangkutana	Mango Lembo Village
			Mangkutana	Pegunungan Faruhumpeni Nature Reserve
		Luwu	Larompong	Temboe Education Forest and Tourism
		Bone	Ulaweng	Cani Sirenreng Nature Park
2	West Sulawesi	Mamuju	Papalang	Batu Papang Village
			Papalang	Palado
3	Central Sulawesi	Marowali	Lembo	Wawopada Village
		Parigi Moutong	Sausu	Sausu Village
				Pangi Binangga Nature Reserve
		Poso	Poso Pesisir	Peawa Oti Montane Forest
		Donggala		Gunung Sojol Nature Reserve



■ Field Work: GPS, Tree Measurement



Field Works: Herbarium Collection



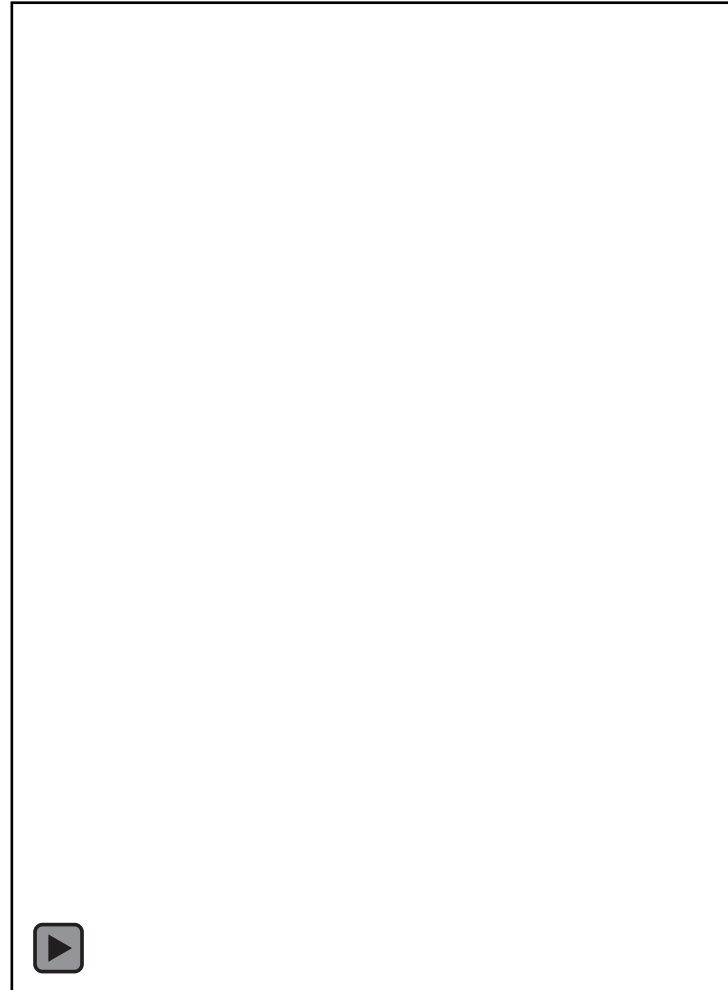
Field Works: Leaf Collection



Field Works: Core Wood Collection



■ Field Works: Core Wood Collection



Activity#2:

DNA Extraction from Ebony (*Diospyros celebica* Bakh.) Dry Wood Samples Collected Using Pickering Punch

Objective

To develop the most efficient method of obtaining DNA from tissues of dry ebony wood

Materials & Methods

1. Core Wood Extraction from Tree Stand Using Pickering Punch (Agroisolab, UK)

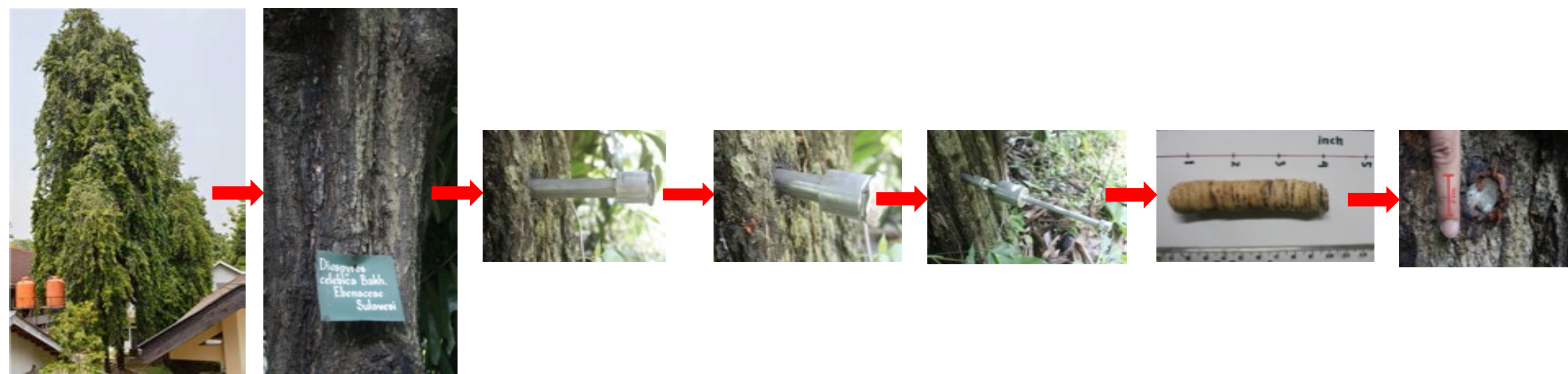
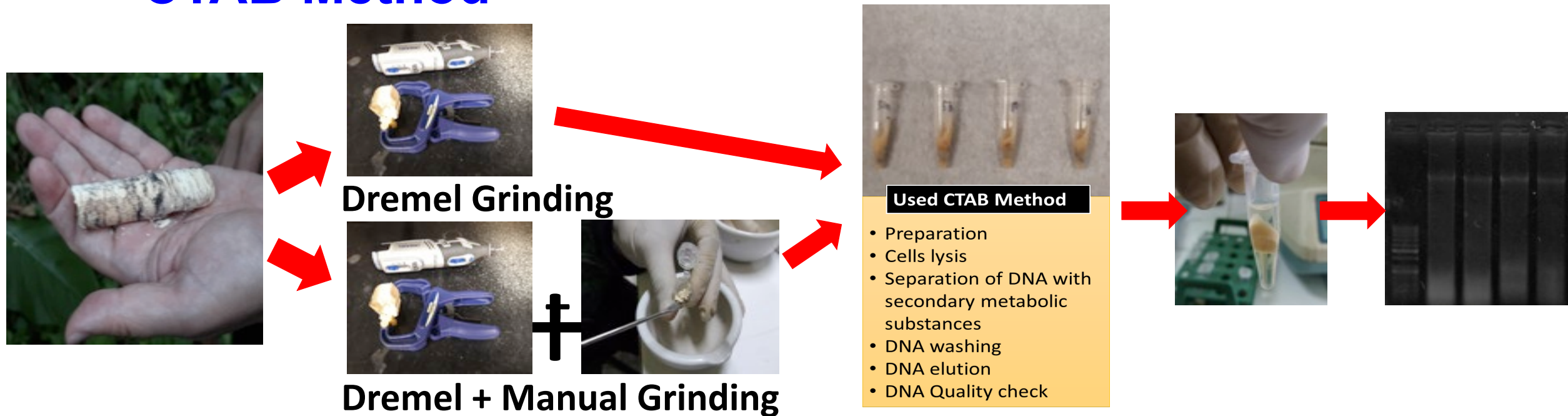


Figure 1. The workflow of ebony wood core extraction using Pickering Punch.

Materials & Methods

2. Genomics DNA Extraction from Dry Wood Using CTAB Method



■ Results & Discussion

Table 1 Presents concentrations and quality of the obtained DNA using two different homogenization methods.

Tree No.	Homogenization Method	DNA Conc. (ng/ μ l)	Purity ($A_{260/280}$)
1	Dremel	215.25	2.532
	Dremel+manual grinding	335.60	2.011
2	Dremel	166.50	2.595
	Dremel+manual grinding	242.75	2.104



Testing the use of Oxford Nanopore's MinION Sequencing Device (in collaboration with Prof Brook Milligan, NMSU)



Closing Remarks

- The **first collection** of samples has been carried out in Sulawesi comprising herbarium, leaves and core wood → Testing the pickering punch.
- Modification of combined the use of “**dremel + manual grinding**” can be recommended for extracting DNA from dry wood → Testing the use of MinION.
- **The core wood** can be used for various purposes → Testing other analyses (isotope, NIR/chemical compound and anatomy).



IPB University
— Bogor Indonesia —

Thank you

