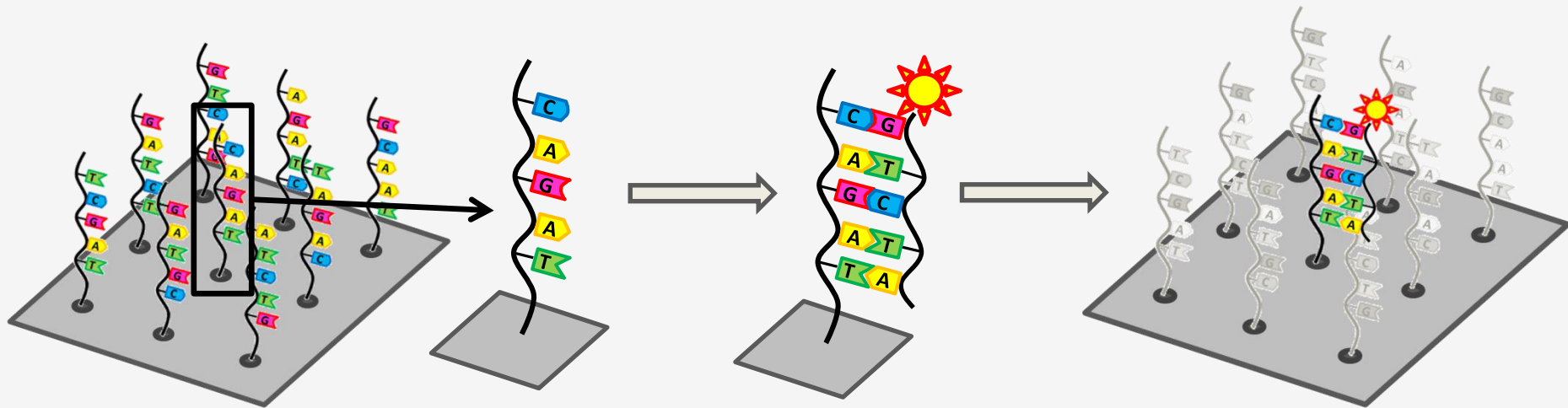
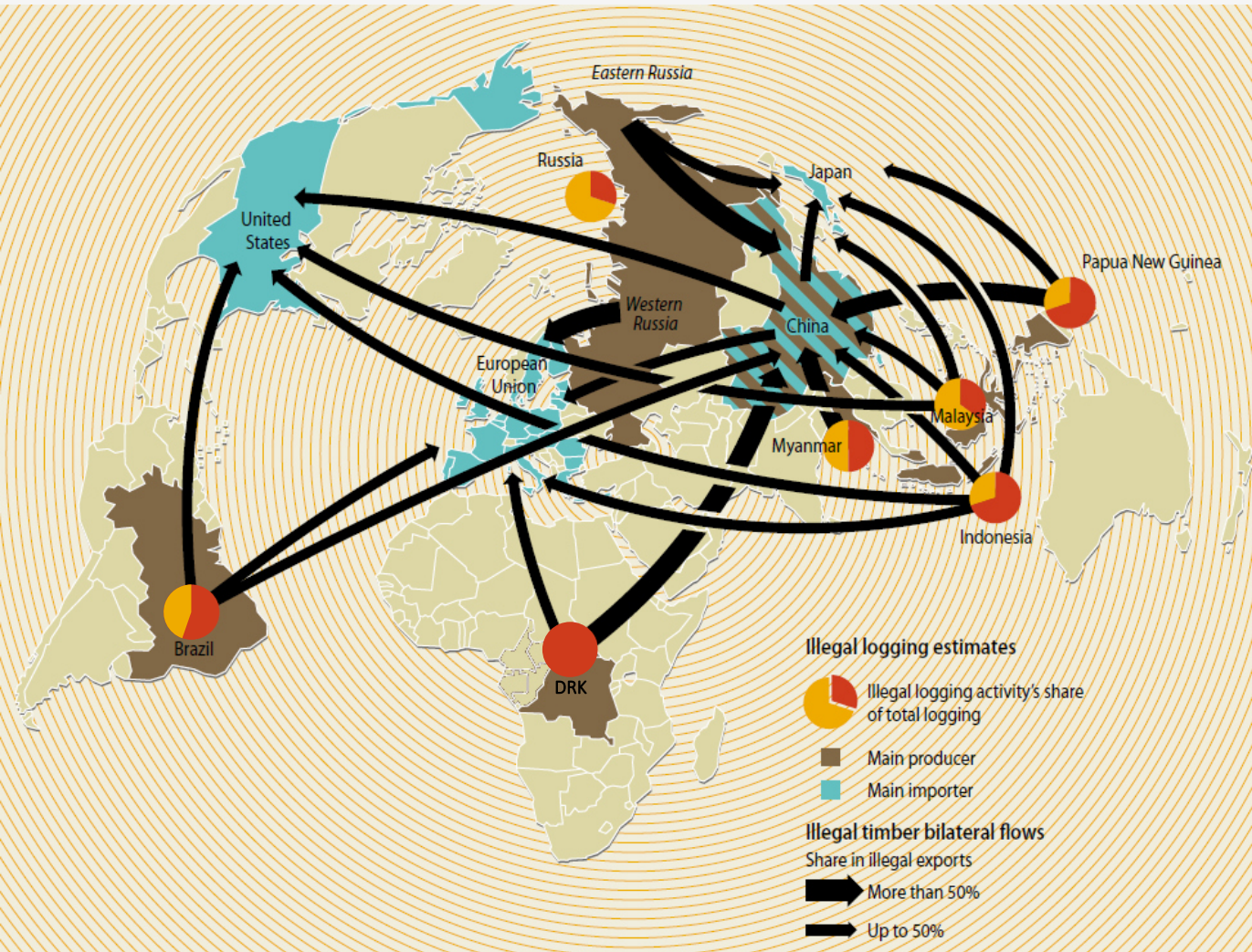


# Development of a DNA macro array for the genetic identification of trade-relevant, CITES wood species and their look alike

Hamburg, 2023



## Illegal logging and trade



- 15 – 30 % of the global logging is illegal
- Sales volume of 51 – 152 bn US\$ <sup>[1]</sup>
- 50 – 90 % of the national harvest



- Loss of biodiversity
- No guarantee for species conservation

[1] Nellemann (2012) INTERPOL Environmental Crime Programme; INTERPOL (2021), factsheet

[2] Hoare (2015) Chatham House Report

## Legal regulations in species protection



- 15 – 30 % of the global logging is illegal
- Sales volume of 51 – 152 bn US\$
- 50 – 90 % of the national harvest



- Loss of biodiversity

Convention on International Trade in Endangered Species of Wild Fauna and Flora

Up to 50%

## Methods for wood species identification



[1] modified

### Problem

There is a lack of available, easy-to-use, low-cost tools to check the legality of a timber product.

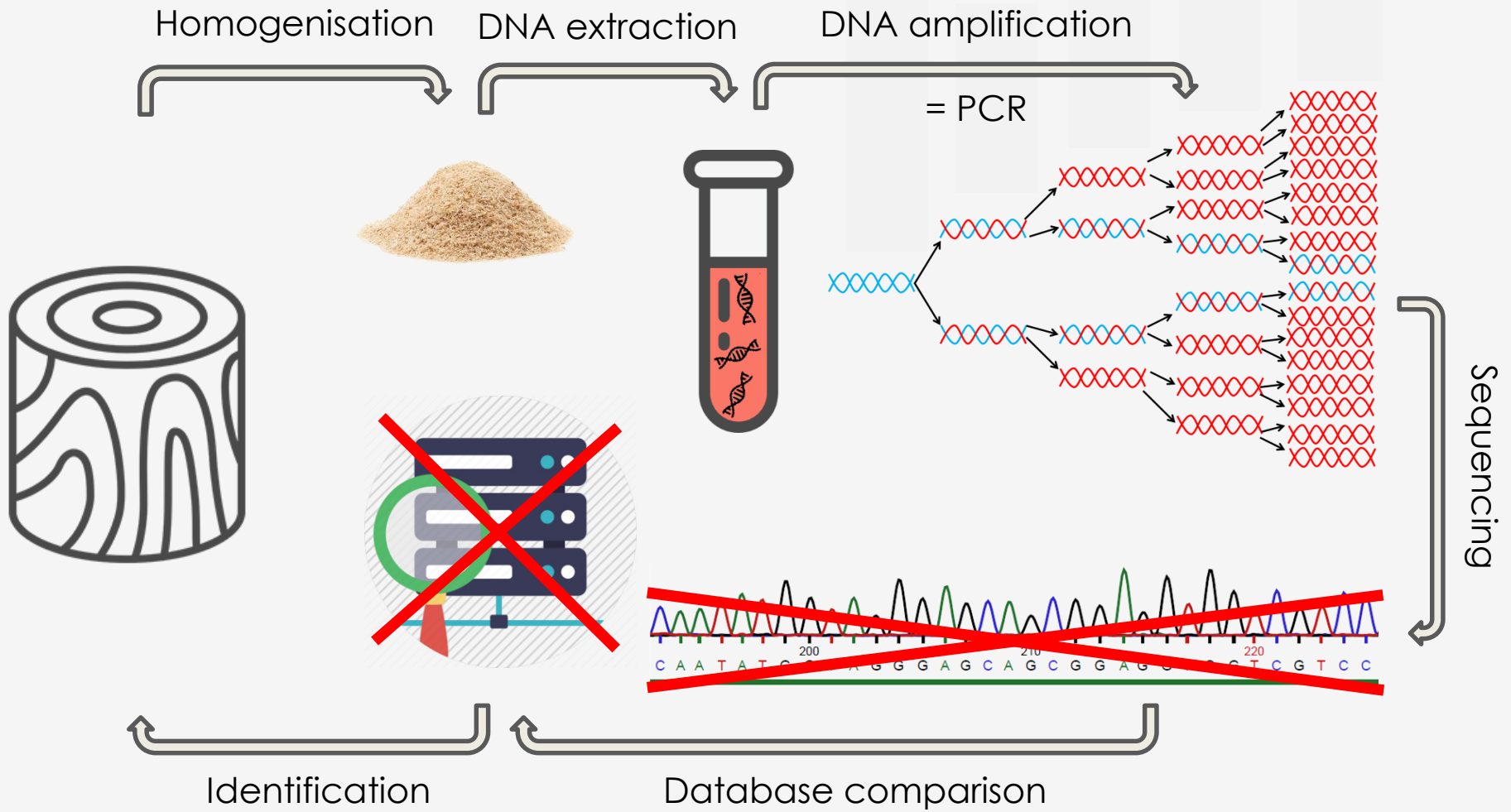
[2]

### Goal

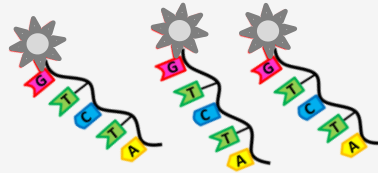
Development of a fast, cheap, genetic, low-tech tool for identifying CITES timber species without anatomical knowledge.

[1] Schmitz et al. (2019) „The Timber Tracking Tool Infogram.“

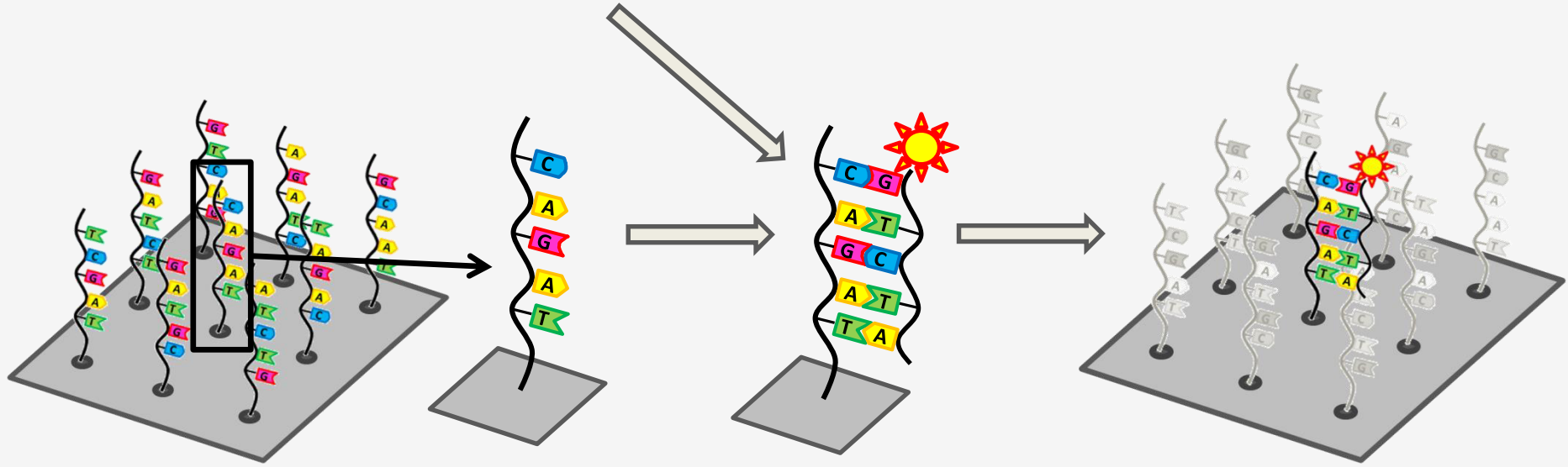
[2] Dormontt et al. (2015); Lowe et al. (2016); Wiedenhoeff et al. (2016); Gasson et al. (2021); Low et al. (2022)



DIG-labelled  
sample DNA



Detection  
reaction



Macro array  
with oligos

Specific  
oligo

Hybridisation of  
oligo and sample

Macro array with  
bonded sample



- No sequencing required
  - No species knowledge required
  - Parallel analysis of several species within one product
  - Low-tech approach  
= cost efficient
- Application in producing countries
- Commercially applicable as DNA chip

- Dependent on accessible DNA
- No application for processed products such as paper or coal

*Carapa guianensis*, *C. procera*  
*Entandrophragma* spp.  
*Khaya* spp.  
*Leplaea cedrata*, *L. thompsonii*  
*Sandoricum koetjape*



True Mahogany  
*Swietenia* spp.

*Guibourtia arnoldiana*, *G. ehie*  
*Copaifera salikounda*  
*Hymenaea courbaril*



Bubinga  
*Guibourtia demeusei*

*Alstonia scholaris*  
*Dyera costulata*  
*Antiaris toxicaria*  
*Brosimum alicastrum*



*Donella pruniformis*  
*Endospermum moluccanum*  
*Jacaranda copaia*  
*Neolamarckia cadamba*  
*Pterygota bequaertii*, *P. macrocarpa*  
*Terminalia ivorensis*,  
*T. superba*

Ramin  
*Gonystylus* spp.



Cedro  
*Cedrela* spp.



Red sanders  
*Pterocarpus santalinus*

*Pterocarpus erinaceus*, *P. indicus*,  
*P. macrocarpus*, *P. soyauxii*



Guaiacum wood  
*Guaiacum* spp.

*Plectrocarpa arborea*  
*Chlorocardium rodiei*  
*Handroanthus heptaphyllus*  
*H. serratifolius*

[1]



Database construction

Design of the oligonucleotids

Optimization macro array

Macro array tests

Sampling

Comparison of sequences

Lievens et al. 2012

Pre-testing in groups

DNA extraction

Searching for species-specific sections



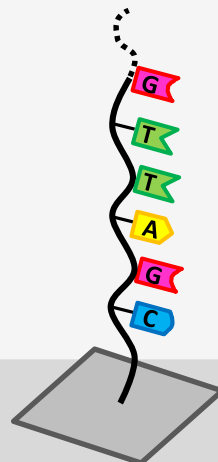
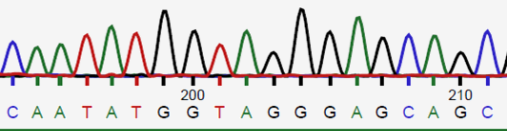
Cross-testing

Amplification of the Barcode

Oligo manufacture



Sequencing



6,7h to 3,2h  
time reduction

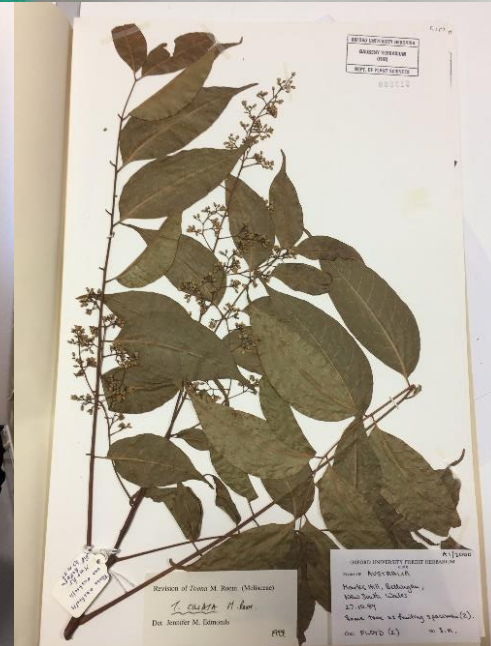
Holzart Sonde	Probe A	Probe B	Probe C	Probe D	Probe E	...
1	x					
2		x			x	
3			x			
4				x		
5				x	x	
6						
7		x				
8						
9	x					
10						
...						

richtig positive Reaktion  
 falsch positive Reaktion

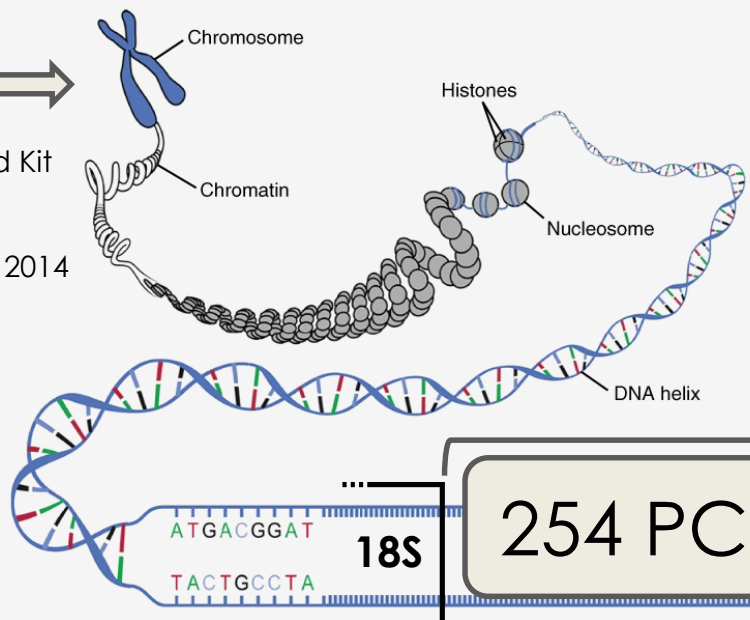
# Sampling



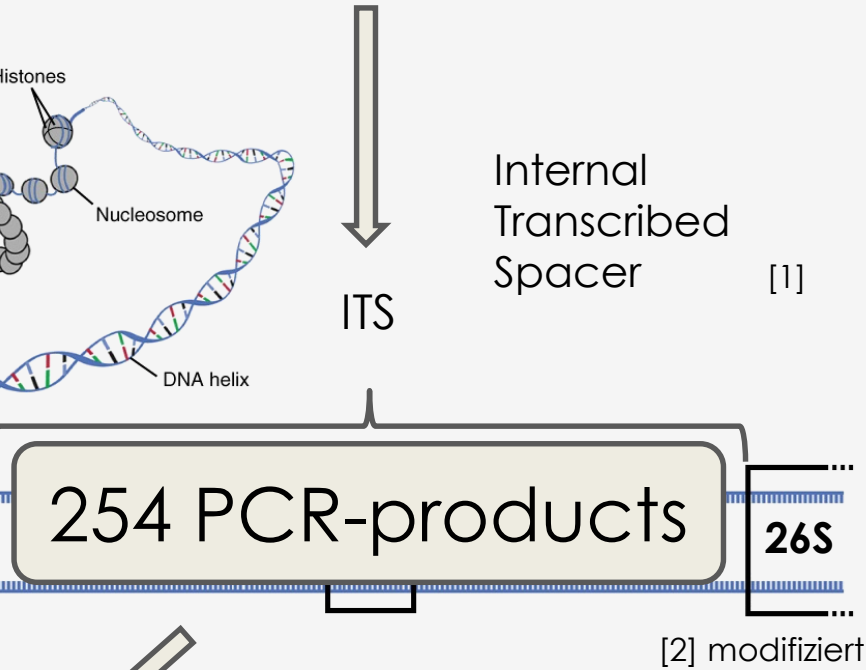
264 samples



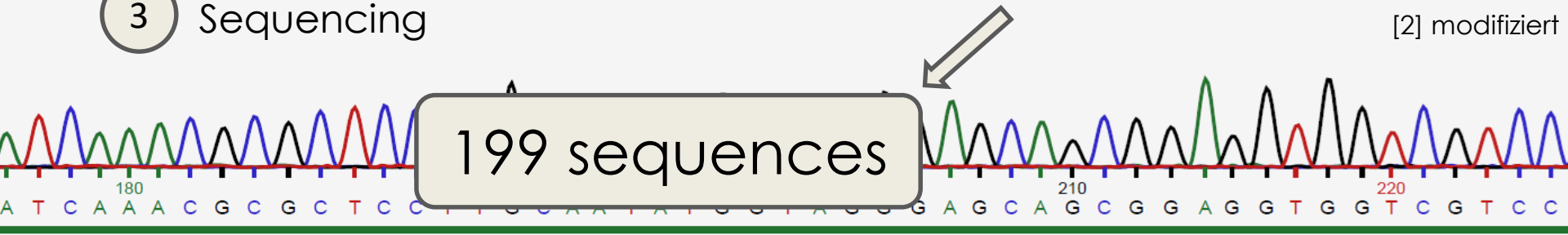
# 1 DNA extraction



# 2 PCR of the barcode



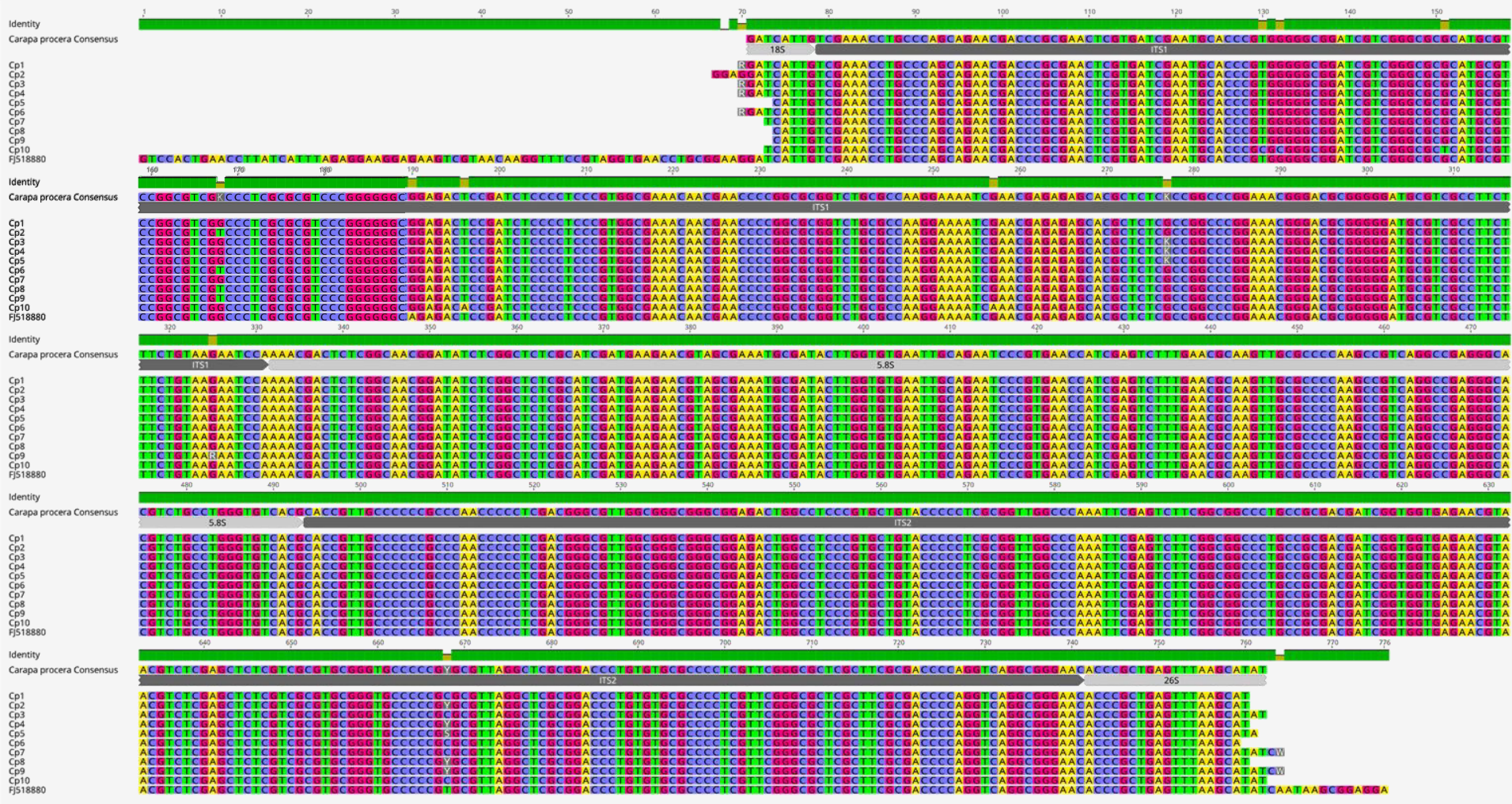
# 3 Sequencing



[1] Baldwin et al. (1995)

[2] nach CFCF 2016 – Carl Fredrik online: [https://upload.wikimedia.org/wikipedia/commons/b/b4/0321\\_DNA\\_Macrostructure.jpg](https://upload.wikimedia.org/wikipedia/commons/b/b4/0321_DNA_Macrostructure.jpg)

# Consensus sequences

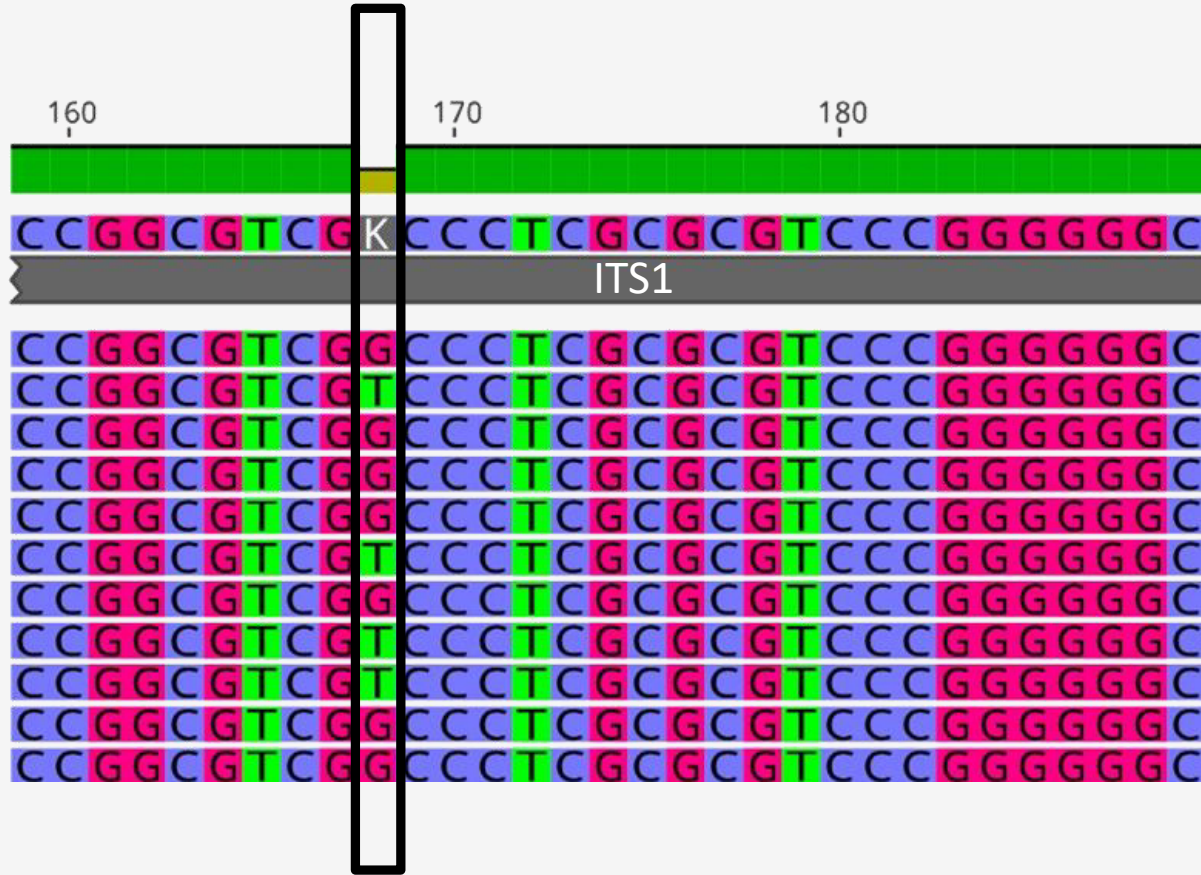


Consensus sequences

Identity

*Carapa procera* Consensus

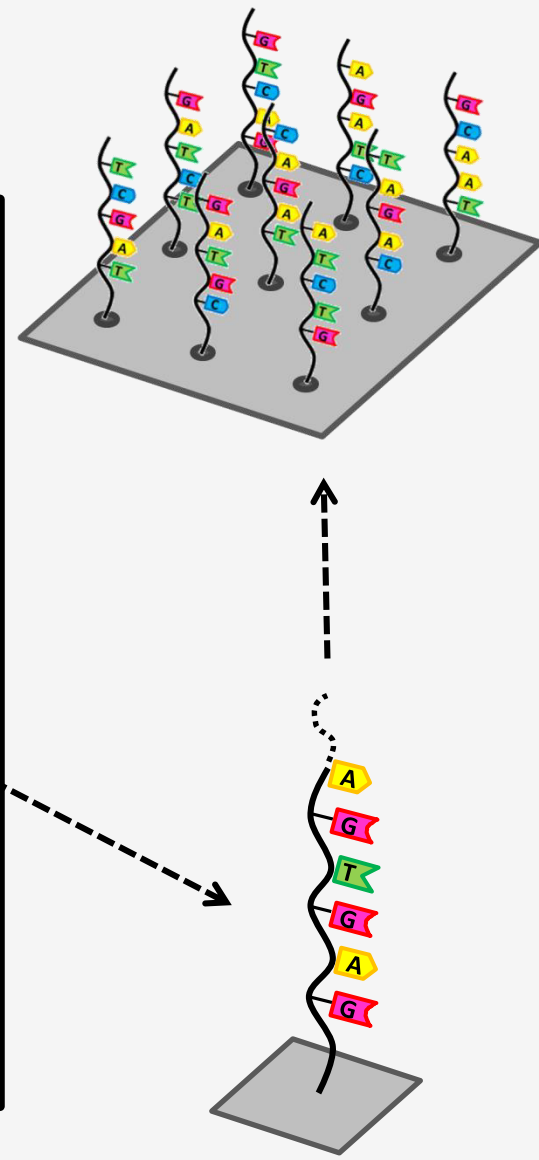
- Cp1
- Cp2
- Cp3
- Cp4
- Cp5
- Cp6
- Cp7
- Cp8
- Cp9
- Cp10
- FI518880



# Comparison of sequences

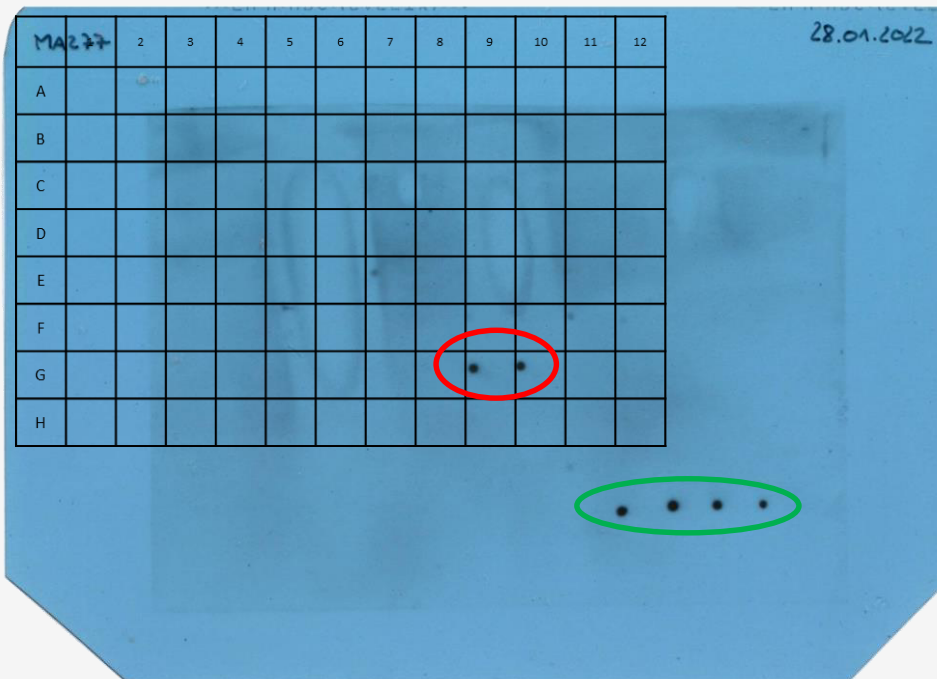


148 oligos



## Evaluation of the X-ray films

Scanned X-ray film



Probe grid

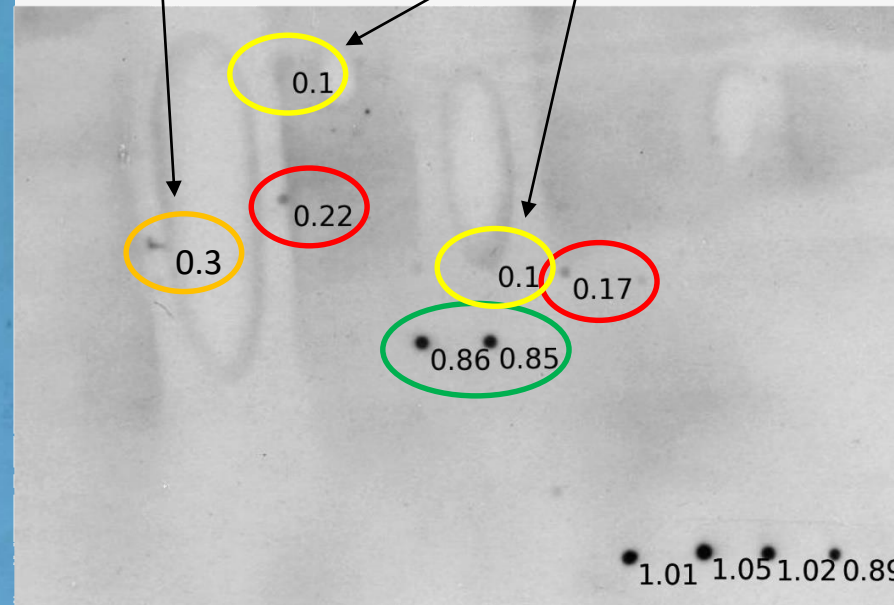
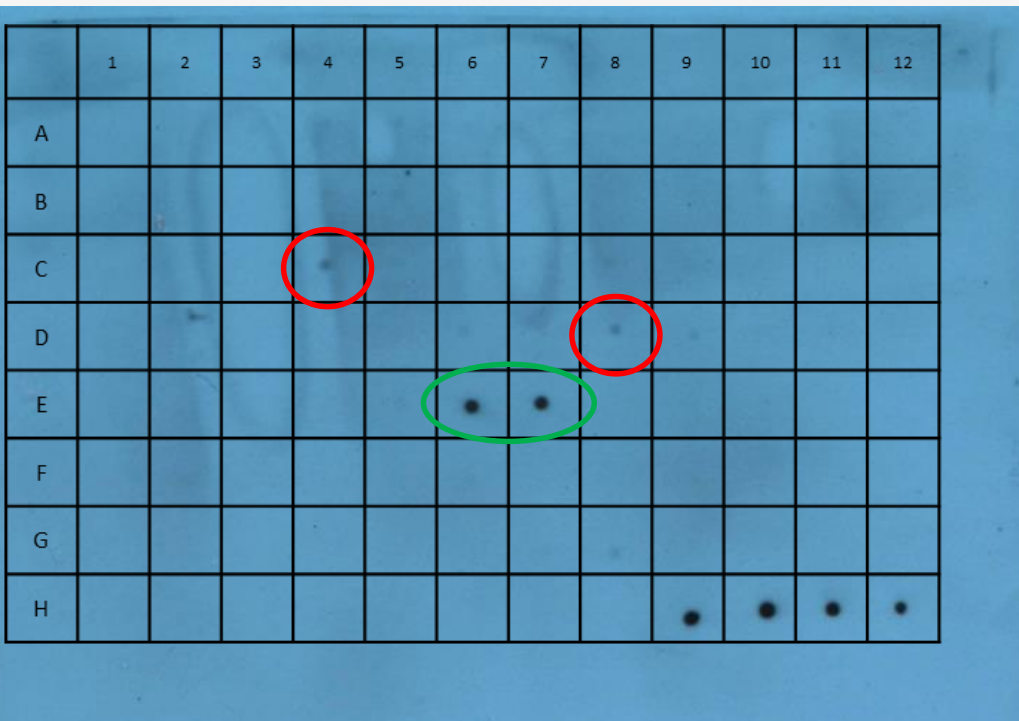
	1	2	3	4	5	6	7	8	9	10	11	12
A	Smahagoni181	Mahagoni for4	Smahagoni649	Smacrophylla112	Macrophylla rev1	Carapa for3	Carapa for2	Carapa rev1	Cprocera 50	Entango for2	Entcyl for2	Ecylicridum264
B	Eutile 86	Khaya for2	Khaya rev1	Khaya243	Kivorensis 106	Lcedrata 35	Lcedrata 459	Lthompsonii35	Lthompsonii457	Cedfissilis for1	Cedodorata for1	Cedrela 29
C	Toona70	Gpelle 221	Gtess 225	Gtess 579	Garnoldiana79	Garnoldiana219	Garnoldiana629	Gehie 44	Csalikounda41	Csalikounda93	Hcourb for1	Hcourb for2
D	Perinaceus615	Psantalinus487	Psantalinus682	Pindicus 487	Pindicus 597	Pmacrocarpus71	Pindimacro32	Psouauxii 501	Gonystylus48	Gonystylus137	Ramin for1	Gonystylus422
E	Ramin rev1	Alstonia 481	Aboonei 408	Atoxicaria 149	Atoxicaria 553	1Balicastrum483	1Balicastrum502	Lprunifomis33	Dprunifomis512	Jcopaia 28	Jcopaia 510	Jcopaia 589
F	Koto for1	Pbequaertii175	Pmacrocarpa26	Limba for1	Limba rev1	Tivorensis 111	Tivorensis 224	Tsuperba 219	Guaiaicum for1	Gcoulteri 172	Gofficiale172	Guaoffici for1
G	Gua sanct for1	Gua sanct rev1	Bulnesia for1	Bulnesia rev1	Parborea 34	Chlorocardium111	lpe for1	Handroanthus211	lpe rev1			
H									Holz for1	Holz for2	Holz rev1	Holz rev2

→ *Brosimum alicastrum*

Wood species: *Brosimum alicastrum*

Measurement results from artefacts

Measurement is caused by unstable background

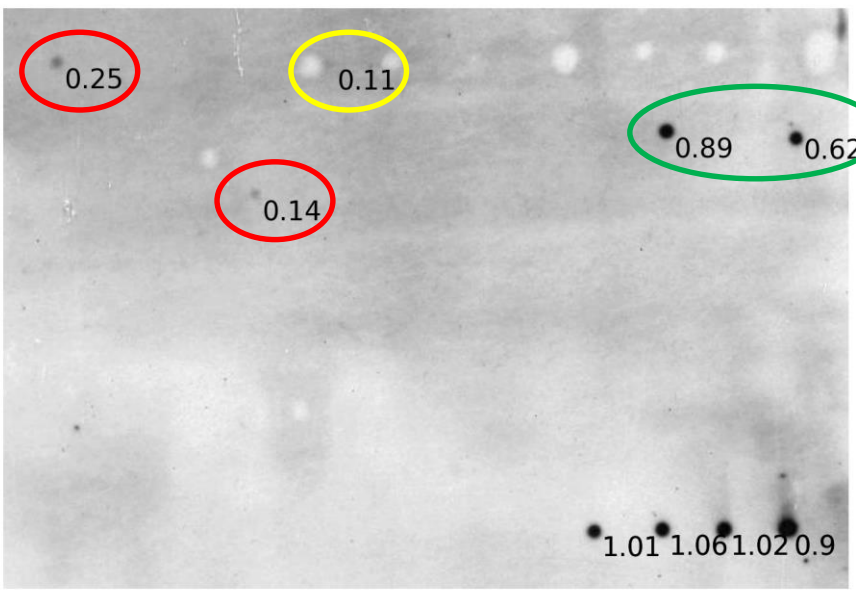


Threshold value 0.1

→ automated signal evaluation with manual check







	<i>K. grandifoliola</i>	<i>K. ivorensis</i>	<i>K. nyasica</i>	<i>K. senegalensis</i>	<i>L. cedrata</i>	<i>L. thompsonii</i>	<i>C. angustifolia</i>	<i>C. fissilis</i>	<i>C. odorata</i>	<i>T. ciliata</i>	<i>T. sureni</i>
<i>K. grandifoliola</i>	0,00	0,00	0,00	0,00	0,00	0,00	0,25	0,00	0,00	0,00	0,00
<i>K. ivorensis</i>	0,00	0,00	0,00	0,00	0,20	0,00	0,00	0,00	0,00	0,00	0,00
<i>K. nyasica</i>	0,00	0,00	0,00	0,00	0,30	0,00	0,00	0,00	0,00	0,00	0,00
<i>K. senegalensis</i>	0,00	0,00	0,01	0,02	0,01	0,00	0,12	0,00	0,00	0,00	0,00
<i>L. cedrata</i>	0,00	0,04	0,00	0,00	0,00	0,00	0,00	0,00	0,02	0,11	0,00
<i>L. thompsonii</i>	0,00	0,01	0,00	0,02	0,00	0,03	0,00	0,00	0,00	0,00	0,00
<i>C. angustifolia</i>	0,00	0,05	0,03	0,04	0,05	0,03	0,00	0,00	0,00	0,01	0,04
<i>C. fissilis</i>	0,00	0,04	0,01	0,01	0,05	0,02	0,17	0,00	0,05	0,00	0,00
<i>C. odorata</i>	0,00	0,00	0,02	0,00	0,00	0,00	0,00	0,02	0,00	0,00	0,00
<i>T. ciliata</i>	0,00	0,02	0,01	0,00	0,00	0,00	0,00	0,00	0,00	0,00	0,07
<i>T. sureni</i>	0,00	0,00	0,00	0,00	0,00	0,00	0,00	0,00	0,00	0,00	0,00
Eucylindricum264	0,00	0,02	0,01	0,00	0,00	0,00	0,00	0,00	0,00	0,00	0,00
Eutile86	0,00	0,00	0,00	0,15	0,00	0,00	0,00	0,75	0,00	0,03	0,00
Khaya for2	0,00	0,07	0,00	0,00	0,00	0,00	0,00	0,76	0,80	0,89	0,92
Khaya rev1	0,00	0,05	0,00	0,00	0,01	0,00	0,00	0,92	0,97	0,96	0,98
Khaya243	0,02	0,00	0,01	0,00	0,00	0,00	0,00	0,45	0,56	0,84	0,75
Kivorensis106	0,00	0,00	0,00	0,00	0,00	0,00	0,00	0,02	0,03	0,14	0,00
Lcedrata35	0,02	0,00	0,00	0,00	0,00	0,00	0,00	0,00	0,00	0,04	0,00
Lcedrata459	0,01	0,00	0,09	0,00	0,03	0,00	0,00	0,00	0,00	0,00	0,00
Lthompsonii35	0,00	0,00	0,00	0,00	0,00	0,00	0,00	0,00	0,00	0,00	0,00
Lthompsonii457	0,01	0,01	0,00	0,19	0,00	0,00	0,00	0,01	0,00	0,01	0,00
Ced fissilisC6for1	0,00	0,00	0,00	0,00	0,00	0,00	0,00	0,00	0,00	0,00	0,00
Ced odorataC6for1	0,00	0,00	0,01	0,05	0,03	0,00	0,02	0,00	0,03	0,00	0,00
Cedrela29	0,00	0,00	0,02	0,04	0,02	0,00	0,00	0,01	0,00	0,10	0,00

- Eucylindricum264
- Eutile86
- Khaya for2
- Khaya rev1
- Khaya243
- Kivorensis106
- Lcedrata35
- Lcedrata459
- Lthompsonii35
- Lthompsonii457
- Ced fissilisC6for1
- Ced odorataC6for1
- Cedrela29

## Oligos

148 tested → 81 selected → 66 appropriate

## Wood species

32 of 53 species identifiable

15 species grouped in 2 to 3 species



Holzart \ Sonde	<i>G. demeuseii</i>	<i>G. pellegriniana</i>	<i>G. tessmannii</i>	<i>G. arnoldiana</i>	<i>G. ehie</i>
<i>G. pellegriniana</i> 221	•	•			
<i>G. tessmannii</i> 225	•	•	•		
<i>G. arnoldiana</i> 79				•	
<i>G. ehie</i> 44					•

Interpretation of the oligo required



Holzart \ Sonde	<i>C. guianensis</i>	<i>C. procera</i>
Carapa for3	•	•
Carapa for2	•	•
Carapa rev1	•	•
<i>C. procera</i> 50		•

2 species no DNA

(*Alstonia scholaris*, *Chlorocardium rodiei*)

4 species no appropriate oligos

(*S. koetjape*, *D. costulata*, *E. moluccanum*, *N. cadamba*)

## Goal

Development of a rapid, low-cost, genetic tool to identify CITES timber species without anatomical knowledge

## Method

Micro array → time reduction of 6,7h to 3,2h

ITS as barcode region → Establishment of an internal database

Oligo design: sequences comparison → 148 developed → 66 appropriate  
32 + 15 wood species identifiable or definable

## Result

Identification is possible within one day

Method with low investment and running costs

No anatomical knowledge required

## Outlook

Extensive validation with samples from trade necessary  
Application in countries of origin

Practicability → new control method

→ Combating the illegal timber trade + Compliance with species protection

# Thank you

