

# Testing highly variable chloroplast and nuclear markers in *Caesalpinia s.l.*

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*Caesalpinia pulcherrima*, G.D. Carr  
[http://www.botany.hawaii.edu/FACULTY/CARR/images/poi\\_pul\\_cu.jpg](http://www.botany.hawaii.edu/FACULTY/CARR/images/poi_pul_cu.jpg)

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POSSIBILITIES



***Caesalpinia gilliesii***

Source : M.-A. Archambault



***Caesalpinia decapetala***

Source: Forest & Kim Starr  
<http://www.hear.org/starr/images/full/starr-011205-0121.jpg>



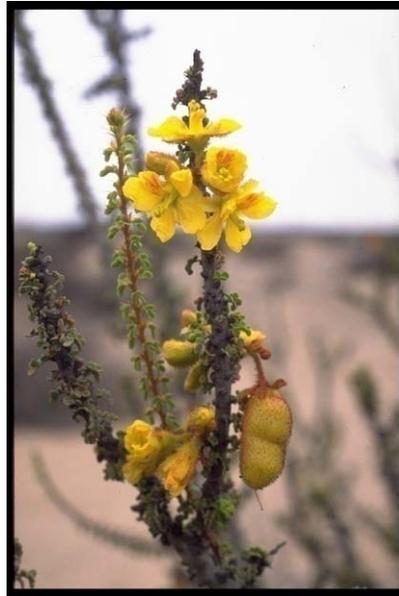
***Caesalpinia echinata***

Source: G. P. Lewis  
<http://www.arkive.org/pau-brasil/caesalpinia-echinata/image-G2285.html>



***Caesalpinia calycina***

Source: G. P. Lewis



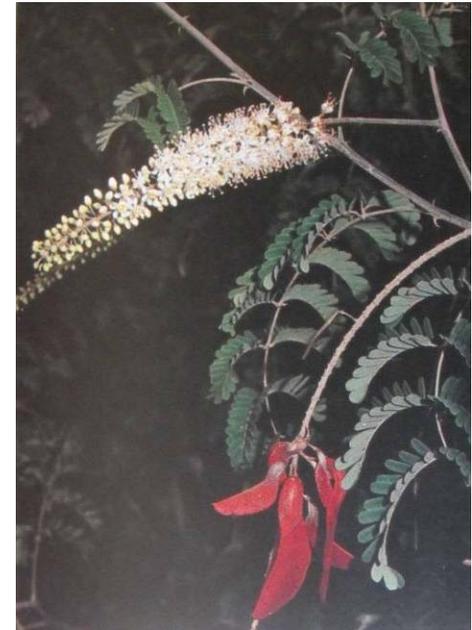
***Balsamocarpon brevifolium***

Source: G. P. Lewis



***Caesalpinia meladenia***

Source: Donovan Bailey



***Pterolobium stellatum***

Source: G. P. Lewis

# Taxonomic classification of *Caesalpinia* s.l.

(Lewis 2005)

## Proposal to reinstate 8 genera in *Caesalpinia* sensu lato

- Does molecular evidence to support this classification?
- What are the generic delimitations?
- Relationships among genera?

<u>Genus</u>	<u>Nb species</u>
<i>Balsamocarpon</i>	1
<i>Cordeauxia</i>	1
<i>Moullava</i>	1
<i>Stahlia</i>	1
<i>Stuhlmannia</i>	1
<i>Zuccaginia</i>	1
<i>Stenodrepanum</i>	1
<i>Cenostigma</i>	2
<i>Haemotaxylon</i>	3
<i>Hoffmannseggia</i>	24
<i>Pomaria</i>	16
<i>Pterolobium</i>	11
<b><i>Caesalpinia</i> (<i>Coulteria</i>)</b>	10
<b><i>Caesalpinia</i> (<i>Erythrostemon</i>)</b>	~ 13
<b><i>Caesalpinia</i> (<i>Guilandina</i>)</b>	7 – 18
<b><i>Caesalpinia</i> (<i>Libidibia</i>)</b>	8
<b><i>Caesalpinia</i> (<i>Mezoneuron</i>)</b>	26
<b><i>Caesalpinia</i> (<i>Poincianella</i>)</b>	~ 36
<b><i>Caesalpinia</i> (<i>Tara</i>)</b>	3
<b><i>Caesalpinia sensus stricto</i></b>	~ 25

# Why past studies have failed to determine generic delimitations:

**27 generic names associated with *Caesalpinia* s.l.**

**Lack of synapomorphic/diagnostic characters in morphology** (Lewis and Schrire, 1995).

**Not enough variation in selected genetic markers**

(Simpson and Miao, 1997, CpDNA restriction sites; Simpson *et al.* 2003, trnL-F)

**Not enough species to represent *Caesalpinia* s.l.**

(Simpson *et al.*, 2004, 2005, Hoffmannseggia ; Simpson *et al.*, 2006, Pomaria)

(Sotuyo *et al.*, 2007, *Caesalpinia hintonii* complex; Bruneau *et al.*, 2008)

# Taxonomic classification of *Caesalpinia* s.l.

(Lewis 2005)

## Research Objectives

1. Adequate taxonomic sampling
2. Find variable genetic markers
3. Biogeographic analysis to explore evolution of SDTF

<u>Genus</u>	<u>Nb species</u>	<u>Nb species in lab</u>
<i>Balsamocarpon</i>	1	1
<i>Cordeauxia</i>	1	1
<i>Moullava</i>	1	1
<i>Stahlia</i>	1	1
<i>Stuhlmannia</i>	1	1
<i>Zuccaginia</i>	1	1
<i>Stenodrepanum</i>	1	1
<i>Cenostigma</i>	2	2
<i>Haemotaxylon</i>	3	2
<i>Hoffmannseggia</i>	24	5
<i>Pomaria</i>	16	2
<i>Pterolobium</i>	11	1
<b><i>Caesalpinia (Coulteria)</i></b>	<b>10</b>	<b>6</b>
<b><i>Caesalpinia (Erythrostemon)</i></b>	<b>~ 13</b>	<b>3</b>
<b><i>Caesalpinia (Guilandina)</i></b>	<b>7 – 18</b>	<b>5</b>
<b><i>Caesalpinia (Libidibia)</i></b>	<b>8</b>	<b>5</b>
<b><i>Caesalpinia (Mezoneuron)</i></b>	<b>26</b>	<b>11</b>
<b><i>Caesalpinia (Poincianella)</i></b>	<b>~ 36</b>	<b>33</b>
<b><i>Caesalpinia (Tara)</i></b>	<b>3</b>	<b>3</b>
<b><i>Caesalpinia sensus stricto</i></b>	<b>~ 25</b>	<b>8</b>

# Pilot test of highly variable markers

## Chloroplast markers

<u>Marker</u>	<u>Source</u>
1. <i>psbA-trnH</i>	(Sotuyo et al., 2007; Simpson et al., 2004)
2. <i>rps16</i>	(Haston et al., 2005)
3. <i>rpL16</i>	(Marazzi et al., 2006)
4. <i>trnD<sup>GUC</sup>-trnT<sup>GGU</sup></i>	(Simon et al., 2009; Kursar et al., 2009; Shaw et al., 2005)
5. <i>trnS<sup>GCU</sup>-trnG<sup>UUC</sup></i>	(Shaw et al., 2007)
6. <i>rpoB-trnC<sup>GCA</sup></i>	(Shaw et al., 2005)
7. <i>trnC<sup>GCA</sup>-ycf6</i>	(Demesure et al. 1995; Shaw et al., 2005)
8. <i>ycf6-psbM</i>	
9. <i>NdhF-rpL32</i>	(Shaw et al., 2007)
10. <i>rpL32-trnL<sup>UAG</sup></i>	(Shaw et al., 2007)
11. <i>matK</i>	(Bruneau et al., 2008)

## Low-Copy Nuclear markers \*

<u>Marker</u>	<u>Source</u>
1. LEAFY/ Floricaula	(Archambault and Bruneau, 2004)
2. CNGC4	(Choi et al., 2006)
3. ARG10	(Choi et al., 2006)
4. AIGP	(Choi et al., 2006)
5. UDPDG	(Choi et al., 2006)
6. tRALS	(Choi et al., 2006)
7. MMKI	(Choi et al., 2006)
8. Sqdl	(Li et al., 2008)

\* At least one sample was cloned for each marker.



# Selected markers for *Caesalpinia s.l.*

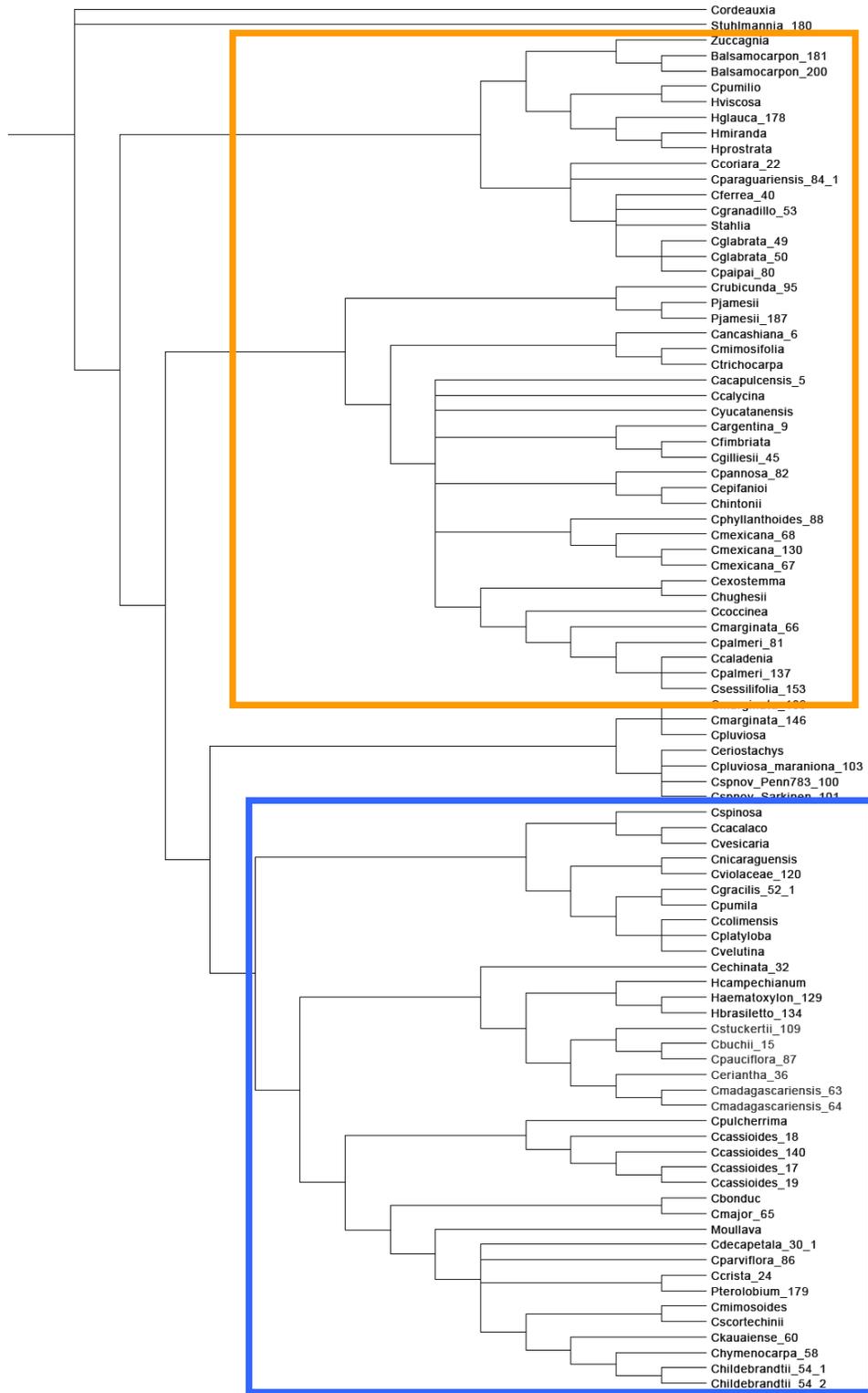
1. ***rps16*** (Peltophorum group, Haston et al., 2005)
2. ***trnD<sup>GUC</sup>-trnT<sup>GGU</sup>*** (Simon et al., 2009; Kursar et al., 2009; Shaw et al., 2005)
3. ***ycf6-psbM*** (Demesure et al. 1995; Shaw et al., 2005)
4. ***trnL*: Bruneau lab + Simpson et al., 2003**
5. ***ITS*: Bruneau lab + Pomaria + Hoffmannseggia**  
(Simpson et al., 2005; 2006)

- Parsimony analysis for each marker
- Parsimony for combined matrix of five markers

# Parsimony results

- Same phylogenetic groups in the strict consensus trees in the independent and combined analysis
- Relationships amongst these groups is unclear : weak backbone of the phylogeny

	<b>trnL</b>	<b>ITS</b>	<b>rps16</b>	<b>trnD-trnT</b>	<b>ycf6-psbM</b>
Nb Taxon	122	104	103	90	48
Nb Characters	605	725	953	1663	1513
Parsimony-Informative Characters	119 (17,6%)	471 (55,6 %)	160 (16,8%)	249 (15,0%)	191 (12,6%)
Nb Trees	+10,000	664	+10,000	+10,000	8
Length Trees	343	3227	436	732	469
Consistency Index	0.7289	0.3042	0.7385	0.7077	0.7655
Retention Index	0.9018	0.6958	0.8874	0.8481	0.8634



Outgroup: *Cordeauxia*  
**Parsimony Analysis of Combined Matrix**  
 Strict consensus of 10,000 trees

**Caesalpinia group 1**  
**Neotropics**

- Balsamocarpon*      *Zuccagnia*
- Hoffmannseggia*      *Libidibia*
- Stahlia*      *Poincianella*
- Erythrostemon*      *Pomaria*

90 taxa (~75 species)  
 5369 characters

**Pluviosa clade**

Fast Bootstrap:

**Caesalpinia group 2** 5000 replicates  
**Old tropics + Neotropics** no branch swapping.

- Tara*      *Coulteria*
- Haematoxylon*      *Caesalpinia* ss
- Moullava*      *Guilandina*
- Mezoneuron*      *Pterolobium*

*Erythrostemon calycina*

G. P. Lewis



*Caesalpinia poiteana*

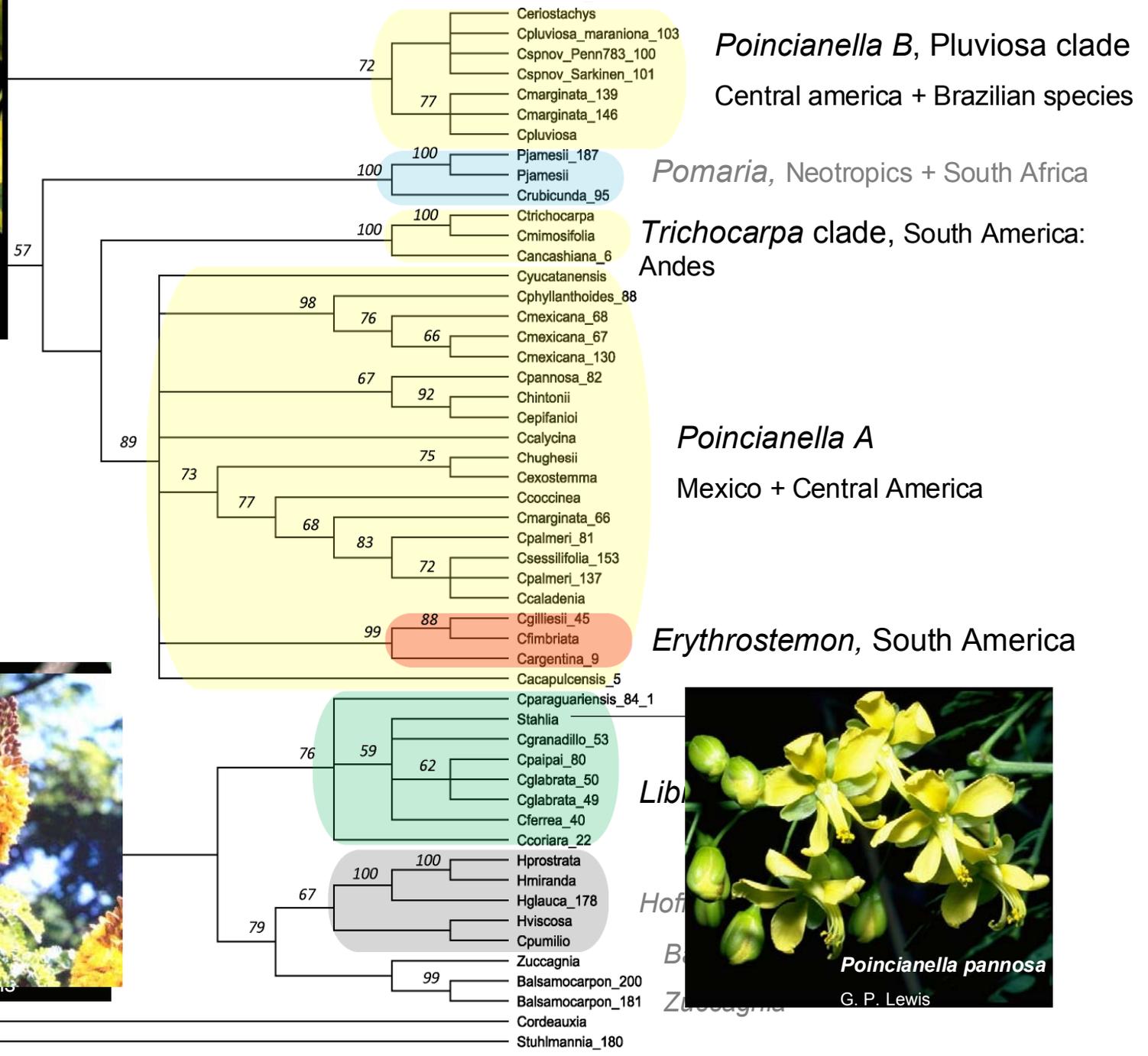


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**Caesalpinia group 1**

Parsimony Analysis

Strict consensus of 10,000 trees

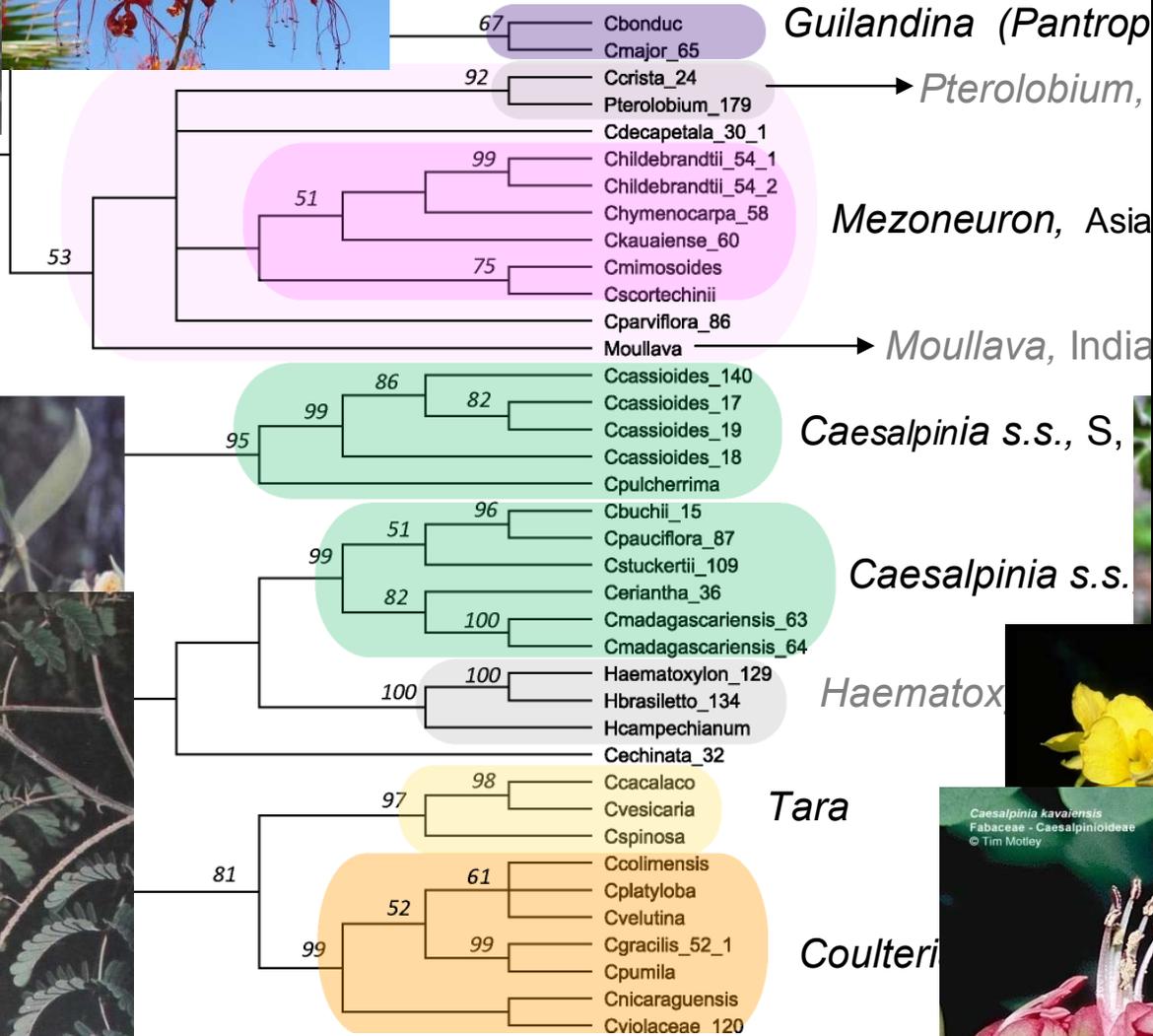
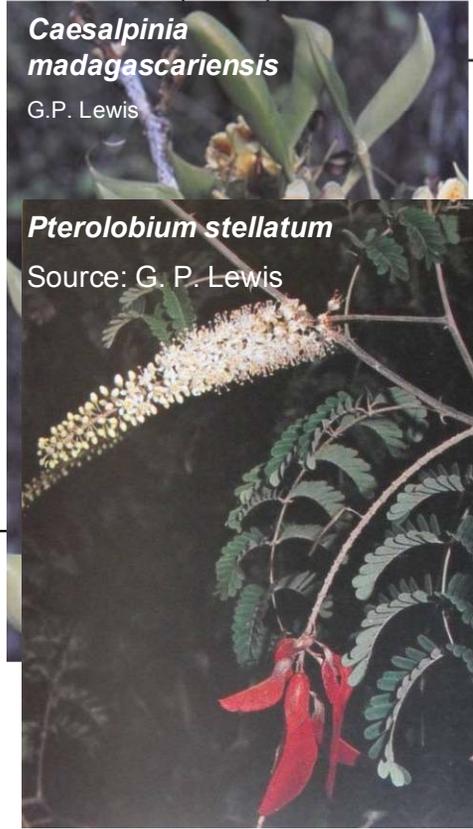


*Poincianella pannosa*

G. P. Lewis

# Caesalpinia group 2

Parsimony Analysis  
 Strict consensus of 10,000 trees



*Guilandina* (Pantrop)

→ *Pterolobium*,

*Mezoneuron*, Asia

→ *Moullava*, India

*Caesalpinia s.s.*, S,

*Caesalpinia s.s.*

*Haematox*

*Tara*

*Coulteri*



[http://www.botany.hawaii.edu/FACULTY/CARR/images/cae\\_kav.jpg](http://www.botany.hawaii.edu/FACULTY/CARR/images/cae_kav.jpg)

**Some of the classification of *Caesalpinia* s.l. by Lewis (2005) seems to be supported by our results, but more work is needed!**

**Developing and scoring a matrix of morphological characters : defining diagnostic characters**

- Collaboration with G. P. Lewis
- Main challenge: *Poincianella-Erythrostemon* (+ *Pomaria*) group

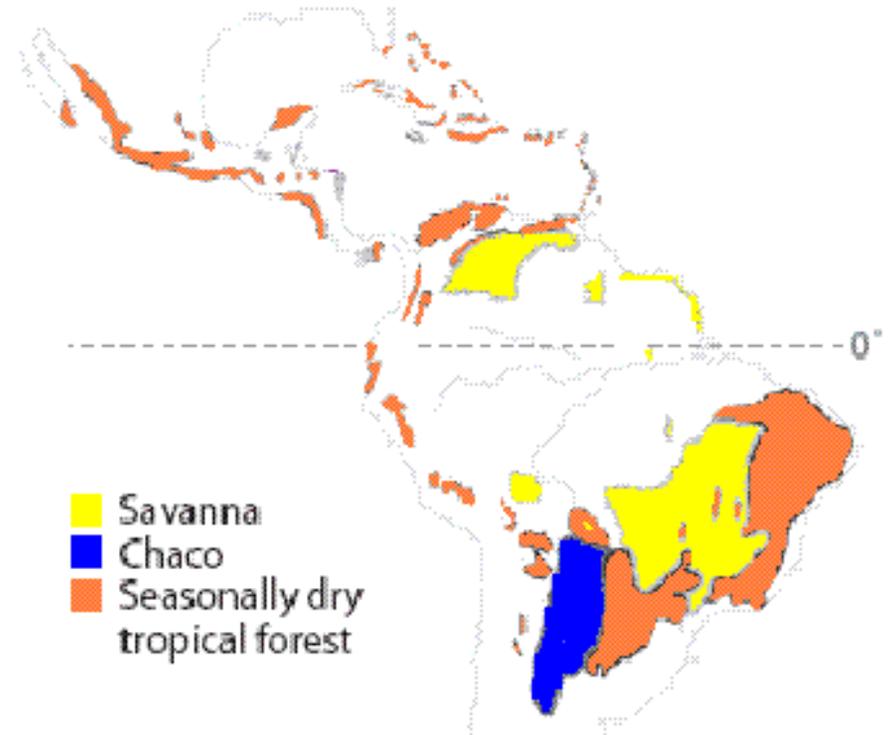
**Biogeographical analyses of Seasonally Dry Tropical Forests**

**Developing *Poincianella-Erythrostemon* group as a biogeographic model to investigate speciation of SDTF:**

- Strong geographic structuring of the phylogeny of *Caesalpinia* s.l.
- Molecular dating analysis with fossil data
  - Mezoneuron (45 mya)
  - *Caesalpinia* sensu stricto (45 mya)

# Seasonally Dry Tropical Forests

- The succulent biome: 5-6 months <100mm of rain, continuous canopy, more fertile soils, low levels of disturbance.
- >50% SDTF in Neotropics: Cactaceae, Bigoniaceae, Leguminosae;
- 97% of remaining SDTF under threat (Miles et al. 2006);
- Disjunct SDTF Arc:
  - High level of beta diversity; endemics.
  - *C. celendinensis*, *H. aphylla*, *C. pluviosa* var. *maraniona*
  - Vicariance or dispersal?



Modified from Pennington et al., 2009

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