Supporting Information

Davis et al. 10.1073/pnas.1403157111



Fig. S1. Malpighiales phylogeny inferred from nucleotide sequences of the plastid inverted repeat region. Fifty-percent maximum-likelihood majority-rule consensus phylogeny is shown here. Well-supported relationships between Centroplacaceae, Elatinaceae, and Malpighiaceae are highlighted in gray. Values are maximum-likelihood bootstrap percentages and Bayesian posterior probabilities, respectively.



Fig. S2. Phylogeny and divergence time estimates of Malpighiaceae inferred using BEAST. The three fossil calibration points described in the main text are starred. Old World clades that have lost the oil-bee association are shown in red; New World clades that have lost the oil-bee association are shown in blue. Numbers near nodes are Bayesian posterior probabilities/maximum-likelihood bootstrap percentages; a hyphen indicates that the node is not present in the maximum-likelihood 50% majority-rule consensus tree. Divergence time estimates in million years with associated confidence intervals are shown in blue.

Table S1. Morphological characters scored by Davis and Anderson (1)

- Traits
- 1. Habit
- 2. Stems rooting at nodes
- 3. Vegetative hairs
- 4. Vegetative hairs
- 5. Phyllotaxy
- 6. Stipule position
- 7. Stipule connation
- . . .
- 8. Stipule persistence
- 9. Stipules enclosing buds
- 10. Petiole glands
- 11. Lamina margin
- 12. Lamina glands
- 13. Inflorescence position
- 14. Inflorescence ultimate units

15. Inflorescence decussate

- 16. Cincinni
- 17. Bract and bracteole size
- 18. Bracts and bracteoles scalelike
- 19. Bract persistence
- 20. Peduncle
- 21. Pedicel
- 22. Bracteole glands
- 23. Bracteole persistence
- 24. Bracteoles enclosing buds
- 25. Cleistogamous flowers
- 26. Breeding system
- Loi Diccuing Syste
- 27. Sepal number
- 28. Sepal aestivation
- 29. Sepal margin, glands
- 30. Calyx in anthesis
- 31. Calyx in fruit
- 32. Calyx glands

33. Calyx gland attachment

- 34. Calyx glands long-stalked
- 35. Petal number
- 36. Petals in bud
- 37. Petal color
- **38. Petals clawed** 39. Petal hairs
- ss. retai nans
- 40. Corolla symmetry
- 41. Disk
- 42. Stamen number
- (in bisexual or male flowers)
- 43. Stamens sterile
 - (in bisexual or male flowers)
- 44. Filament sizes (fertile)
- 45. Filament fusion
- 46. Anther hairs
- 47. Anther bristles, apex
- 48. Anther wings
- Score 0: tree or woody shrub; 1: erect suffrutex; 2: trailing suffrutex; 3: vine; 4: herb 0: no; 1: yes 0: unicellular; 1: multicellular 0: 2-branched; 1: basifixed; 2: stellate 0: opposite or whorled; 1: alternate 0: stem between petioles or beside petiole; 1: petiole margin; 2: petiole inner face; 3: not present 0: distinct; 1: connate, same leaf; 2: connate, opposite leaves; 3: connate, opposite and same leaves; 4: not present 0: long-persistent; 1: soon-deciduous; 2: not present 0: no; 1: yes 0: absent; 1: present 0: without true teeth; 1: toothed 0: absent; 1: abaxial surface; 2: margin or v. sl. within; 3: adaxial surface 0: terminal or terminal and axillary; 1: axillary 0: pseudoracene or thyrse; 1: umbel of (2)4(6) flowers; 2: umbel of >six flowers; 3: 1(2) flower(s); 4: verticil of four flowers; 5: spike; 6: consistently two flowers 0: mostly not; 1: all or mostly yes; 2: proximally decussate, distally not; 3: too tight to tell; 4: single flowers 0: one-flowered; 1: two to several-flowered 0: full-sized leaves; 1: much reduced; 2: apparently absent 0: no; 1: yes; 2: absent 0: persistent; 1: deciduous; 2: absent 0: 0-1 mm long; 1: >1 mm long 0: well developed, > bracteoles; 1: absent or very short, < bracteoles 0: eglandular; 1: one or both glandular; 2: bracteoles absent 0: persistent; 1: deciduous; 2: absent 0: no; 1: yes; 2: absent 0: absent; 1: present 0: bisexual; 1: morphologically bisexual, functionally unisexual; 2: morphologically bisexual or male; 3: unisexual, dioecious 0: 5; 1: 2-4 0: imbricate; 1: valvate 0: absent; 1: present 0: erect or appressed; 1: revolute; 2: reflexed 0: hardly enlarged; 1: enlarged, papery 0: none; 1: 10 on five sepals; 2: eight on lateral four sepals; 3: four (fused) on lateral four sepals; 4: up to five glands; 5: six on lateral four sepals 0: completely on free sepals; 1: half or more on free sepals; 2: mostly below free sepals; 3: not present 0: no; 1: yes; 2: not present 0: five; 1: two to four 0: exposed; 1: concealed by sepals 0: yellow; 1: pink (+white); 2: white; 3: lilac 0: yes; 1: no 0: absent; 1: present, abaxial surface; 2: present, adaxial surface; 3: present, margin 0: nearly radial; 1: bilateral, NW type; 2: bilateral, Acridocarpus type 0: absent; 1: present 0: >10; 1: 10; 2: 7-9; 3: six opposite sepals + posterior petal; 4: five opposite sepals; 5: 1-4 0: none; 1: staminodes opposite five sepals; 2: staminodes opposite anterior-lateral sepals; 3: posterior three staminodes; 4: filament opposite posterior petal without anther; 5: staminodes opposite posterior-lateral sepals; 6: staminodes opposite petals, posterior-lateral sepals; 7: staminodes opposite three anterior sepals; 8: staminodes opposite five petals 0: subequal; 1: longer opposite sepals; 2: longer opposite some or all petals; 3: longer opposite sepals, posterior-lateral petals; 4: anterior three shorter; 5: posterior three shorter; 6: shorter opposite posterior-lateral petals; 7: very long opposite anterior sepal; 8: stouter opposite posterior-lateral petals; 9: anterior one shorter 0: distinct; 1: connate at base or higher 0: absent; 1: present 0: absent; 1: present 0: absent; 1: present

Table S1. Cont.	
Traits	Score
49. Anther dehiscence	0: longitudinal slits; 1: apical or subapical pores
50. Connective extended,	0: no; 1: yes
fleshy (fertile anthers)	
51. Anther size (fertile)	0: subequal; 1: larger opposite sepals; 2: larger opposite petals;
	3: larger opposite posterior-lateral petals; 4: posterior three smaller;
	5: smaller opposite anterior-lateral petals; 6: anterior three smaller;
	7: anterior one larger; 8: larger opposite three anterior sepals;
	9: anterior three larger
52. Androecium symmetry	0: nearly radial; 1: strongly bilateral
53. Pollen symmetry	0: radial; 1: global
54. Pollen ectoapertures	0: present; 1: absent
55. Endoapertures, number	0: 3; 1: 4–12
56. Pollen diameter (μm)	0: ≤ 22; 1: >22
57. Receptacle hairs between	0: absent or nearly so; 1: present, abundant
filaments and gynoecium	
58. Carpel number (chasmogamous flowers)	0: three; 1: two; 2: five
59. Ovules per locule	0: one (or none); 1: two or more
60. Carpels sterile	0: none; 1: anterior; 2: one posterior; 3: anterior +one posterior
61. Carpel fusion in ovary	0: distinct or connate at base only; 1: connate whole ventral face or axis
62. Style number (chasmogamous flowers)	0: as many as carpels; 1: one on anterior carpel; 2: two on posterior carpels;
	3: two on anterior carpel + one posterior carpel
63. Style fusion	0: distinct; 1: connate in styles; 2: connate in stigmas
64. Gynoecium symmetry	0: nearly radial; 1: strongly bilateral
65. Style thickness	0: subulate, slender; 1: uniform, thick; 2: uniform, slender
66. Stigma	0: terminal or nearly so, tiny; 1: internal, short, large; 2: internal, long-decurrent, large;
	3: internal, tiny; 4: terminal, large, capitate or truncate; 5: terminal, large,
	reniform; 6: terminal, large, elongated
67. Style dorsal extension	0: none; 1: angle to hook; 2: foliole
68. Styles distally	0: entire; 1: bifid
69. Fruit texture	0: dry; 1: fleshy
70. Fruit dehiscence	0: schizocarpic, not releasing seed; 1: indehiscent; 2: loculicidally
	dehiscent, releasing seeds; 3: septicidally dehiscent, releasing seeds
71. Fruit wall	0: smooth; 1: setiferous; 2: dorsal wing dominant; 3: lateral wing(s)
	dominant; 4: winglets, ruffles; 5: aculeate; 6: dorsal or dorsal+lateral crest(s)
72. Carpophore	0: absent; 1: present
73. Eumascagnioid disk	U: absent; 1: present
74. Arii 75. Fadaaaa	U: absent; I: present
75. Endosperm	U: absent; 1: present

Characters with significant difference between oil-bee pollinated clades are highlighted in bold. An acceleration in trait diversification in oil-bee clades is indicated in blue; a deceleration in oil-bee free clades are in bold and italics.

1. Davis CC, Anderson WR (2010) A complete generic phylogeny of Malpighiaceae inferred from nucleotide sequence data and morphology. Am J Bot 97(12):2031–2048.

PNAS PNAS