

Table S1. Accession table for vouchers used in phylogenetic analyses (bolded text indicates new GenBank accessions).

No.	Species	Voucher	Country	ITS	ETS	<i>KEA1</i> intron 11	<i>KEA1</i> intron 17	<i>TEB</i> exon 17	<i>matK</i>	<i>ndhF</i>
1	<i>Haematostemon guianensis</i> Sandwith	Wurdack 4350 (US) ^a	Guyana	KP794434	MF502427	—	—	MF502846	MF502706	MF502776
2	<i>Plukenetia africana</i> Sond.	Bartsch 1859 (US)	Namibia	MF502513	MF502431	MF502579	MF502651	MF502848	MF502710	MF502780
3	<i>P. africana</i>	Guy 2367 (MO)	Botswana	MF502514	—	—	—	—	—	MF502781
4	<i>P. africana</i>	Palomoti 1086 (MO)	Botswana	MF502515	MF502432	—	—	—	MF502711	MF502782
5	<i>P. africana</i>	Pope et al. 834 (MO)	Botswana	MF502516	MF502433	MF502580	—	—	MF502712	MF502783
6	<i>P. ankaranensis</i> L.J.Gillespie	Gillespie et al. 10697 (CAN) ^a	Madagascar	KP794438	MF502434	MF502581	MF502652	MF502849	MF502713	MF502784
7	<i>P. ankaranensis</i>	Lees s.n. (CAN)	Madagascar	KP794437	MF502435	—	MF502653	MF502850	MF502714	MF502785
8	<i>P. brachybotrya</i> Müll.Arg.	Acevedo-Rodriguez 14416 (NY)	Peru	MF502517	MF502436	MF502582	MF502654	MF502851	MF502715	MF502786
9	<i>P. brachybotrya</i>	Araujo-M. et al. 1722 (MO)	Bolivia	KP794452	MF502437	MF502583	MF502655	MF502852	MF502716	MF502787
10	<i>P. brachybotrya</i>	Fuentes & Torrico 5398 (MO)	Bolivia	MF502518	MF502438	MF502584	—	—	MF502717	MF502788
11	<i>P. brachybotrya</i>	Galiano et al. 6612 (MO)	Peru	MF502519	MF502439	—	—	—	MF502718	MF502789
12	<i>P. brachybotrya</i>	Seidel & Vaquiata 7733 (MO)	Bolivia	MF502520	MF502440	MF502585	MF502656	MF502853	MF502719	MF502790
13	<i>P. aff. brachybotrya</i>	Ledezma et al. 921 (CAN)	Bolivia	MF502511	MF502428	MF502576	MF502649	—	MF502707	MF502777
14	<i>P. aff. brachybotrya</i>	Sperling et al. 5873 (MO)	Brazil	MF502512	MF502429	MF502577	MF502650	MF502847	MF502708	MF502778
15	<i>P. aff. brachybotrya</i>	Sperling et al. 6161 (CAN)	Brazil	—	MF502430	MF502578	—	—	MF502709	MF502779
16	<i>P. carabiasiae</i> J.Jiménez Ram.	Meave et al. 1550 (MO)	Mexico	MF502521	MF502441	MF502586	—	—	MF502720	—
17	<i>P. cf. carolis-vegae</i>	Calatayud et al. 2643 (CAN)	Peru	MF502527	MF502448	MF502592	—	—	—	—
18	<i>P. cf. carolis-vegae</i>	Huamantupa et al. 6445 (CAN)	Peru	MF502528	MF502449	MF502593	MF502660	—	MF502724	—
19	<i>P. cf. carolis-vegae</i>	Monteagudo et al. 14125 (MO)	Peru	MF502529	MF502450	MF502594	MF502661	—	MF502725	MF502793
20	<i>P. cf. carolis-vegae</i>	Monteagudo et al. 15252 (MO)	Peru	MF502530	MF502451	—	—	—	MF502726	MF502794
21	<i>P. cf. carolis-vegae</i>	Rojas et al. 3863 (CAN)	Peru	MF502532	MF502453	MF502595	—	—	—	—

22	<i>P. cf. carolis-vegae</i>	Valenzuela et al. 5197 (CAN)	Peru	MF502534	MF502455	MF502597	MF502663	—	—	—
23	<i>P. cf. carolis-vegae</i>	van der Werff et al. 17532 (CAN)	Peru	MF502535	MF502456	MF502598	MF502664	MF502856	MF502729	MF502797
24	<i>P. cf. carolis-vegae</i>	Vasquez et al. 33145 (MO)	Peru	MF502536	MF502457	MF502599	MF502665	MF502857	MF502730	MF502798
25	<i>P. cf. carolis-vegae</i>	Woytkowski 6670 (MO)	Peru	MF502537	MF502458	MF502600	MF502666	MF502858	MF502731	MF502799
26	<i>P. conophora</i> Müll.Arg.	Hart 1621 (MO)	Democratic Republic of the Congo	MF502523	MF502443	MF502588	—	—	—	—
27	<i>P. conophora</i>	Nemba & Thomas 434 (MO)	Cameroon	KP794457	MF502444	MF502589	MF502657	—	MF502722	MF502792
28	<i>P. corniculata</i> Sm.	Huq & Haroon 10780 (GH)	Bangladesh	MF502524	MF502445	MF502590	MF502658	MF502854	—	—
29	<i>P. corniculata</i>	Huq & Haroon 10780 (MO)	Bangladesh	MF502525	MF502446	MF502591	MF502659	MF502855	MF502723	—
30	<i>P. decidua</i> L.J.Gillespie	Rakotomalaza 597 (CAN)	Madagascar	MF502526	MF502447	—	—	—	—	—
31	<i>P. huayllabambana</i> Bussmann, C.Tellez & A.Glenn	Quipusco 381 (MO)	Peru	MF502531	MF502452	—	—	—	MF502727	MF502795
32	<i>P. huayllabambana</i>	Tellez et al. 4 (MO)	Peru	MF502533	MF502454	MF502596	MF502662	—	MF502728	MF502796
33	<i>P. lehmanniana</i> (Pax & K.Hoffm.) Huft & L.J.Gillespie	Acevedo-Rodriguez & Daly 1658 (MO)	Ecuador	KP494443	MF502459	MF502601	MF502667	MF502859	MF502732	MF502800
34	<i>P. lehmanniana</i>	Clark 3953 (MO)	Ecuador	MF502538	MF502460	MF502602	MF502668	MF502860	MF502733	MF502801
35	<i>P. lehmanniana</i>	Silverstone-Sopkin & Giralda-Gesini 8019 (MO)	Colombia	MF502539	MF502461	MF502603	—	MF502861	MF502734	MF502802
36	<i>P. lehmanniana</i>	Zak & Jaramillo 3401 (MO)	Ecuador	MF502540	MF502462	MF502604	MF502669	MF502862	MF502735	MF502803
37	<i>P. lorentensis</i> Ule	Grandez 19608 (AMAZ) ^a	Peru	KP794453	MF502463	MF502605	MF502670	MF502863	MF502736	MF502804
38	<i>P. lorentensis</i>	McDaniel & Rimachi 22451 (MO)	Peru	MF502541	MF502464	MF502606	—	—	MF502737	MF502805
39	<i>P. lorentensis</i>	Rimachi 5122 (MO)	Peru	MF502542	MF502465	MF502607	—	—	MF502738	MF502806
40	<i>P. lorentensis</i>	Solomon 7972 (MO)	Bolivia	MF502543	MF502466	MF502608	—	MF502864	MF502739	MF502807
41	<i>P. lorentensis</i>	Vásquez & Jaramillo 3283 (MO)	Peru	MF502544	MF502467	MF502609	—	—	—	—
42	<i>P. lorentensis</i>	Vasquez et al. 38069 (MO)	Peru	MF502545	MF502468	MF502610	MF502671	MF502865	MF502740	MF502808
43	<i>P. madagascarensis</i> Leandri	Andrianjafy 1648 (CAN)	Madagascar	MF502546	MF502469	MF502611	MF502672	—	MF502741	—

44	<i>P. madagascarensis</i>	Gillespie 4175 (CAN)	Madagascar	MF502547	MF502470	MF502612	MF502673	MF502866	MF502742	MF502809
45	<i>P. madagascarensis</i>	Villiers et al. 4899 (MO)	Madagascar	MF502548	MF502471	MF502613	MF502674	MF502867	MF502743	MF502810
46	<i>P. penninervia</i> Müll.Arg.	Atha et al. 1001 (MO)	Belize	KP794455	MF502472	MF502614	MF502675	MF502868	MF502744	MF502811
47	<i>P. penninervia</i>	Carnevali & Duno 7594 (MO)	Mexico	MF502549	MF502473	MF502615	MF502676	MF502869	MF502745	MF502812
48	<i>P. penninervia</i>	Martínez 10527 (MO)	Mexico	KP794456	MF502474	MF502616	MF502677	MF502870	MF502746	MF502813
49	<i>P. penninervia</i>	Martinez 17705 (DAV)	Mexico	MF502550	MF502475	MF502617	MF502678	MF502871	MF502747	MF502814
50	<i>P. penninervia</i>	McPherson 8461 (MO)	Panama	KP794454	MF502476	MF502618	MF502679	MF502872	MF502748	MF502815
51	<i>P. penninervia</i>	Wallnöfer & Frisch 5996 (MO)	Guatemala	MF502551	MF502477	MF502619	—	MF502873	MF502749	MF502816
52	<i>P. cf. penninervia</i>	Gentry 47799 (MO)	Colombia	MF502522	MF502442	MF502587	—	—	MF502721	MF502791
53	<i>P. polyadenia</i> Müll.Arg.	Croat 20285 (MO)	Peru	MF502552	—	—	—	—	MF502750	MF502817
54	<i>P. polyadenia</i>	Gillespie 4314 (CAN)	Smithsonian Greenhouse ex French Guiana	MF502553	MF502478	MF502620	MF502680	MF502874	MF502751	MF502818
55	<i>P. polyadenia</i>	Wurdack 5288 (US)	Guyana	MF502554	MF502479	MF502621	MF502681	MF502875	MF502752	MF502819
56	<i>P. serrata</i> (Vell.) L.J.Gillespie	Davidse 10480 (MO)	Brazil	MF502555	MF502480	—	—	—	MF502753	MF502820
57	<i>P. serrata</i>	Forzza et al. 5328 (RB)	Brazil	MF502556	MF502481	MF502622	MF502682	—	MF502754	MF502821
58	<i>P. serrata</i>	Goldenberg et al. 1424 (RB)	Brazil	MF502557	MF502482	MF502623	MF502683	—	MF502755	MF502822
59	<i>P. serrata</i>	Peixoto et al. 4154 (MO)	Brazil	KP794458	MF502483	MF502624	—	—	MF502756	MF502823
60	<i>P. serrata</i>	Sartori & Pardo 11 (RB)	Brazil	MF502558	MF502484	MF502625	MF502684	—	MF502757	MF502824
61	<i>P. serrata</i>	Thomas 10221 (NY)	Brazil	MF502559	MF502485	MF502626	MF502685	—	—	MF502825
62	<i>P. stipellata</i> L.J.Gillespie	Aguilar 8193 (MO)	Costa Rica	KP794448	MF502486	MF502627	MF502686	MF502876	MF502758	MF502826
63	<i>P. stipellata</i>	Cardinal-McTeague 8 (CAN) ^a	Costa Rica	MF502560	MF502487	MF502628	MF502687	MF502877	MF502759	MF502827
64	<i>P. stipellata</i>	Ibarra Manriquez & Sinaca 1115 (MO)	Mexico	MF502561	MF502488	MF502629	MF502688	MF502878	—	MF502828
65	<i>P. stipellata</i>	Liesner 3088 (MO)	Costa Rica	KP794451	MF502489	MF502630	MF502689	MF502879	—	MF502829
66	<i>P. stipellata</i>	Morales & Rojas 5342 (MO)	Costa Rica	KP794450	MF502490	MF502631	MF502690	MF502880	MF502760	MF502830

67	<i>P. stipellata</i>	Refugio Cedillo Trigas 3510 (MO)	Mexico	MF502562	MF502491	MF502632	MF502691	MF502881	MF502761	MF502831
68	<i>P. stipellata</i>	Urbina 1155 (MO)	Nicaragua	KP794449	MF502492	MF502633	MF502692	MF502882	MF502762	MF502832
69	<i>P. supraglandulosa</i> L.J.Gillespie	Acevedo-Rodriguez 6022 (US)	Suriname	MF502563	MF502493	MF502634	MF502693	MF502883	MF502763	MF502833
70	<i>P. verrucosa</i> Sm.	Barrabe & Crozier 145 (US)	French Guiana	MF502564	MF502494	MF502635	MF502694	MF502884	MF502764	MF502834
71	<i>P. verrucosa</i>	Herrera & Kaemar 10073 (CAN)	Suriname	MF502565	MF502495	—	—	—	—	—
72	<i>P. verrucosa</i>	Hoffman 5917 (US)	Suriname	MF502566	MF502496	MF502636	MF502695	MF502885	MF502765	MF502835
73	<i>P. volubilis</i> L.	Bell 93-546 (US)	Peru	KP794447	MF502497	MF502637	MF502696	MF502886	MF502766	MF502836
74	<i>P. volubilis</i>	Burnham & Krings 1640 (MO)	Ecuador	KP794446	MF502498	MF502638	MF502697	MF502887	MF502767	MF502837
75	<i>P. volubilis</i>	Carrillo & Reyes 448 (MO)	Ecuador	MF502567	MF502499	MF502639	—	—	—	—
76	<i>P. volubilis</i>	Huamantupa 3500 (CAN)	Peru	MF502568	MF502500	MF502640	MF502698	—	—	—
77	<i>P. volubilis</i>	Jaramillo et al. 1237 (MO)	Bolivia	MF502569	MF502501	MF502641	MF502699	—	MF502768	MF502838
78	<i>P. volubilis</i>	Nee 55162 (MO)	Bolivia	KP794445	MF502502	MF502642	MF502700	MF502888	MF502769	MF502839
79	<i>P. volubilis</i>	Parada et al. 206 (CAN)	Bolivia	KP794444	MF502503	MF502643	MF502701	MF502889	MF502770	MF502840
80	<i>P. volubilis</i>	Teran et al. 2502 (MO)	Bolivia	MF502570	MF502504	MF502644	MF502702	—	MF502771	MF502841
81	<i>P. volubilis</i>	Valenzuela 8531 (MO)	Peru	MF502571	MF502505	—	—	—	—	—
82	<i>P. volubilis</i>	Wurdack s.n. (US) ^a	Smithsonian Greenhouse ex Peru	MF502572	MF502506	MF502645	MF502703	MF502890	MF502772	MF502842
83	<i>Romanoa tamnoides</i> (A.Juss.) Radcl.-Sm.	Fuentes 1848 (MO)	Bolivia	MF502573	MF502507	—	—	—	—	—
84	<i>R. tamnoides</i>	Raes & Terceros 177 (MO)	Bolivia	KP794435	MF502508	MF502646	MF502704	—	MF502773	MF502843
85	<i>R. tamnoides</i>	Raes et al. 211 (MO)	Bolivia	MF502574	MF502509	MF502647	MF502705	—	MF502774	MF502844
86	<i>R. tamnoides</i>	Zardini & Chaparro 50824 (MO)	Paraguay	MF502575	MF502510	MF502648	—	—	MF502775	MF502845

^a = silica gel preserved voucher; all others from herbarium sheets

Table S2. Accession table of genomes/transcriptomes used in data-mining analyses.

No.	Taxon	Type	Assembly file	Reference
1	<i>Croton tiglium</i> L.	Transcriptome	VVPY_Croton_tiglium_-SOAPdenovo-Trans-assembly.fa	1KP Project
2	<i>Euphorbia mesembryanthemifolia</i> Jacq.	Transcriptome	LSLA_Euphorbia_mesembryanthemifolia_juv_-SOAPdenovo-Trans-assembly.fa	1KP Project
3	<i>E. mesembryanthemifolia</i>	Transcriptome	VPDX_Euphorbia_mesembryanthemifolia_mat_-SOAPdenovo-Trans-assembly.fa	1KP Project
4	<i>Jatropha curcas</i> L.	Draft genome	GCF_000696525.1_JatCur_1.0_rna.fna	Sato et al. 2011
5	<i>Manihot grahamii</i> Hook.	Transcriptome	XNLP_Manihot_grahamii_-SOAPdenovo-Trans-assembly.fa	1KP Project
6	<i>Plukenetia volubilis</i> L.	Transcriptome	PlutentiaAssemGADC01000001-GADC01066310.fasta	Wang et al. 2012
7	<i>Ricinus communis</i> L.	Draft genome	TIGR_castorWGS_release_0.1.cds.fsa	Chan et al. 2010
8	<i>R. communis</i>	Draft genome	TIGR_castorWGS_release_0.1.gene.fsa	Chan et al. 2010
9	<i>R. communis</i>	Transcriptome	PAZJ_Ricinus_communis_1kp_-SOAPdenovo-Trans-assembly.fa	1KP Project

Table S3. List of molecular markers, primer sequence, and amplification/sequencing protocols.

Name	Primer (5' to 3')	Reference	Recommended Protocol
Nuclear DNA (nDNA)			
External Transcribed Spacer (ETS)			
ETS F1	GTT ATG TTT GGY GTT TCG GSA ATG CT	Designed here	Amplification and sequencing: ETS Pluk F2 + 18S 5' R Annealing temperature: 52°C
ETS F2	ATG ATC GTT TGS TTT GGC AGG CTC	Designed here	
18S 5' R	GAA TTA GTT CAT ACT TAC ACA TGC ATG	Designed here	
Internal Transcribed Spacer (ITS)			
ITS 5a [F]	CCT TAT CAT TTA GAG GAA GGA G	Stanford et al. 2000	Amplification and sequencing were done for the Full region (ITS 5a + u4) or in two overlapping fragments: ITS1 (ITS 5a + u2) and ITS2 (ITS p3 + u4). Sometimes ITS p2 and 4 were used as slightly internal sequencing primers. Annealing temperature: 55°C
ITS u2 [R]	GCG TTC AAA GAY TCG ATG RTT C	Cheng et al. 2016	
ITS p2 [R]	GCC RAG ATA TCC GTT GCC GAG	Cheng et al. 2016	
ITS p3 [F]	YGA CTC TCG GCA ACG GAT A	Cheng et al. 2016	
ITS u4 [R]	RGT TTC TTT TCC TCC GCT TA	Cheng et al. 2016	
ITS 4 [R]	TCC TCC GCT TAT TGA TAT GC	White et al. 1990	
KEA1 intron 11			
KEA1-i11 F	GAA TCA GAT ATT GCK CCA TAT CGT G	Designed here	Amplification and sequencing: KEA1-i11 F + R Annealing temperature: 55°C
KEA1-i11 R	AGA AGC TTT GGA TCA ATA GAC ATT C	Designed here	
KEA1 intron 17			
KEA1-i17 F	AGG TCC TTC ATA AAG TTG GTG CWG A	Designed here	Amplification and sequencing: KEA1-i17 F + R Annealing temperature: 65°C
KEA1-i17 R	GTG CAA GAA CAG CAG CTG CCA ACT G	Designed here	
TEB exon 17			
TEB F	GCR AAG AAR TCA AGA ATG GTG CWC G	Designed here	Amplification and sequencing: TEB F + R2 Annealing temperature: 55°C
TEB R1	CTC TCY TCA TCA GGC CAM GAA TCC A	Designed here	
TEB R2	ATT SAT AGG ACC CTT CTC ACA GGC A	Designed here	
Plastid DNA (cpDNA)			
matK (full coding sequence)			
matK F1	AAT CCT CTA TCC TTG CTT CAA	Designed here	Amplification and sequencing were designed to capture the full coding region in three overlapping fragments, an effective protocol for poor quality DNA. matK Pluk F1 + R1, F2 + R2, and F3 + R3. Annealing temperature: 55°C
matK R1	CTC ATG AAG AAA GAG TYG TAA TAA A	Designed here	
matK F2	AAT ACC TTA CCC CAT CCA TCT A	Designed here	
matK R2	ATT GGA ACT ATT GTA TCG AGT TTC	Designed here	
matK F3	GTA CGG AGT CAA ATG YTA GAA	Designed here	
matK R3	CCT TAC TCG TAT ACT GTA TGA GC	Designed here	
ndhF 3' (partial coding sequence)			
ndhF 1318F	GGA TTA ACY GCA TTT TAT ATG TTT CG	Olmstead and Sweere 1994	Amplification and sequencing: ndhF 1318F + 2110Ri (or 1703R, to obtain a partial sequence for poorer quality samples) Annealing temperature: 49°C
ndhF 1703R	GGC TCC AAT AAA YAA AGT	Beilstein et al. 2006	
ndhF 2110Ri	TCA ATT ATT CGT TTA TCA A	Steinmann and Porter 2002	

All PCR amplifications used BIOLASE DNA Polymerase (Bioline) in 15uL reactions. Recipes included: 9.2uL H₂O; 1.5uL 10x NH₄ Reaction Buffer; 0.6uL 50mM MgCl₂ Solution [2.0mM]; 0.3uL 10mM dNTP Mix [0.2mM]; 0.375uL 10uM Primer (each) [0.25uM]; 1.5uL 1mg/mL Bovine Serum Albumin (BSA; Sigma Aldrich) [0.1 mg/mL]; 0.15uL 5U/uL BIOLASE DNA Polymerase [0.05U/uL]; 1uL Genomic DNA. Nuclear markers included 3uL 5M Betaine (Sigma Aldrich) [1M], in which case H₂O was reduced to 6.2uL. Thermocycler protocols followed: 3min 94°C Initiation; 35 cycles of (i) 30s 94°C Denaturing, (ii) 1m (see table) Annealing, (iii) 30s 72°C Extension; 10min 72°C Final extension.

Table S4. Biogeographical distribution matrix (A = Mexico to NW South America, B = Amazon/Guiana, C = Atlantic Forest, D = Tropical Africa, E = Madagascar, F = SE Asia).

Taxon	A	B	C	D	E	F
<i>Haematostemon guianensis</i>	0	1	0	0	0	0
<i>Plukenetia africana</i>	0	0	0	1	0	0
<i>Plukenetia ankaranensis</i>	0	0	0	0	1	0
<i>Plukenetia brachybotrya</i>	0	1	0	0	0	0
<i>Plukenetia aff. brachybotrya</i>	0	1	0	0	0	0
<i>Plukenetia carabiasiae</i>	1	0	0	0	0	0
<i>Plukenetia cf. carolis-vegae</i>	0	1	0	0	0	0
<i>Plukenetia conophora</i>	0	0	0	1	0	0
<i>Plukenetia corniculata</i>	0	0	0	0	0	1
<i>Plukenetia decidua</i>	0	0	0	0	1	0
<i>Plukenetia lehmanniana</i>	1	0	0	0	0	0
<i>Plukenetia loretensis</i>	0	1	0	0	0	0
<i>Plukenetia madagascariensis</i>	0	0	0	0	1	0
<i>Plukenetia penninervia</i>	1	0	0	0	0	0
<i>Plukenetia cf. penninervia</i>	1	0	0	0	0	0
<i>Plukenetia polyadenia</i>	0	1	0	0	0	0
<i>Plukenetia serrata</i>	0	0	1	0	0	0
<i>Plukenetia stipellata</i>	1	0	0	0	0	0
<i>Plukenetia supraglandulosa</i>	0	1	0	0	0	0
<i>Plukenetia verrucosa</i>	0	1	0	0	0	0
<i>Plukenetia volubilis</i>	0	1	0	0	0	0
<i>Romanoa tamnoides</i>	0	0	1	0	0	0

Table S5. Manual dispersal matrix between biogeographical areas (adjacent = 1.0, short-distance dispersal = 0.5, long-distance dispersal = 0.1).

	Mex. to NW South Am.	Amazon/ Guiana	Atlantic Forest	Trop. Africa	Madagascar	SE Asia
0 to 35 Mya	A	B	C	D	E	F
Mex. to NW South Am.	1	-	-	-	-	-
Amazon/ Guiana	1.0	1	-	-	-	-
Atlantic Forest	0.1	0.5	1	-	-	-
Trop. Africa	0.1	0.1	0.1	1	-	-
Madagascar	0.1	0.1	0.1	0.5	1	-
SE Asia	0.1	0.1	0.1	0.1	0.1	1

Table S6. Trait matrix used in phylogenetic regression analysis with seed size. Plants size (0 = slender stems; 1 = robust to thick stems), seedling ecology (0 = light gap or forest edge; 1 = shade avoidance, canopy liana strategy), fruit type/ hypothesized dispersal mechanism (0 = dry, dehiscent/ scatter hoarding rodents; 1 = fleshy, indehiscent/ mammal predation), biome (0 = wet dominant; 1 = dry component), fire tolerance (0 = non-fire adapted; 1 = fire adapted).

Taxon	Plant size	Seedling ecology	Fruit type/ dispersal	Biome	Fire tolerance
<i>Haematostemon guianensis</i>	1	0	0	0	0
<i>Romanoa tamnoides</i>	0	0	0	0	0
<i>Plukenetia africana</i>	0	0	0	1	1
<i>Plukenetia ankaranensis</i>	1	0	0	1	0
<i>Plukenetia brachybotrya</i>	0	0	0	0	0
<i>Plukenetia</i> aff. <i>brachybotrya</i>	0	0	0	0	0
<i>Plukenetia carabiasiae</i>	1	1	0	0	0
<i>Plukenetia</i> cf. <i>carolis-vegae</i>	1	0	0	0	0
<i>Plukenetia conophora</i>	1	1	1	0	0
<i>Plukenetia corniculata</i>	0	0	0	0	0
<i>Plukenetia decidua</i>	1	0	0	1	0
<i>Plukenetia huayllabambana</i>	1	0	0	0	0
<i>Plukenetia lehmanniana</i>	1	0	1	0	0
<i>Plukenetia loretensis</i>	0	0	0	0	0
<i>Plukenetia penninervia</i>	0	0	0	0	0
<i>Plukenetia</i> cf. <i>penninervia</i>	0	0	0	0	0
<i>Plukenetia polyadenia</i>	1	1	1	0	0
<i>Plukenetia serrata</i>	1	0	1	0	0
<i>Plukenetia stipellata</i>	0	0	0	0	0
<i>Plukenetia supraglandulosa</i>	0	0	0	0	0
<i>Plukenetia verrucosa</i>	0	0	0	0	0
<i>Plukenetia volubilis</i>	0	0	0	0	0

Table S7. The best evolutionary model for seed size based on ΔAIC and Akaike weights (w_i), using the modified *Plukenetia* BEAST chronogram, \log_{10} (estimated seed volume), and the fitContinuous function in GEIGER.

Model	AIC _c	ΔAIC	w_i
Brownian motion	50.03442	0.0	0.6584417
Ornstein-Uhlenbeck	52.73257	2.69815	0.1708530
Early Burst	52.73430	2.69988	0.1707053