

Supplementary Table

Table S1. **Node support (ML bootstrap percentage (BP) and Bayesian posterior probability (PP)) for phylogenetic analyses.** Node numbers refer to the phylogeny in Figures 3 and S1.

Dashes refer to nodes with <50 BS or 0.5 PP support.

Node#	ML		Bayes	
	Constrained (C)	Unconstrained	Constrained (C)	Unconstrained
1	-	-	-	-
3	-	-	-	-
5	-	-	0.78	0.8
6	-	-	-	-
7	65	67	0.97	0.98
10	83	79	1.0	0.99
11	92	89	1.0	1.0
14	87	87	1.0	1.0
17	-	-	-	-
18	-	-	-	-
19	-	-	0.57	0.66
21	C, 97	90	C, 1.0	1.0
23	-	53	-	0.56
26	-	-	0.54	0.5
28	C, 74	-	C, 1.0	0.97
30	-	-	0.51	0.53
32	68	56	0.94	0.93
35	-	-	0.74	0.77
36	-	-	-	-
37	-	-	0.59	0.62
39	C, 94	92	C, 1.0	1.0
41	75	75	0.94	0.95
44	-	-	-	-
45	-	-	-	-
47	-	-	0.65	0.59
49	-	-	-	-
51	55	52	0.91	0.9
54	-	-	0.67	0.67
56	-	-	-	0.5
58	-	-	-	-
59	82	82	1.0	1.0
61	83	80	1.0	1.0
63	93	93	1.0	1.0
65	56	-	-	-
68	-	-	-	-
69	70	72	0.93	0.93

72	-	-		0.94	0.95
74	59	54		0.97	0.97
76	98	97		1.0	1.0
79	-	-		0.98	0.98
80	-	-		0.84	0.61
82	100	100		1.0	1.0
85	-	-		0.61	0.5
86	C, -	-		C, 1.0	-
87	82	69		1.0	0.99
89	67	61		0.81	0.82
91	98	98		1.0	1.0
94	-	-		0.94	0.71
95	63	-		0.99	0.94
97	95	94		1.0	1.0
100	82	77		0.96	0.96
101	100	100		1.0	1.0
104	99	99		1.0	1.0
107	-	-		-	-
108	C, 55	-		C, 1.0	-
110	-	-		0.69	-
113	C, 91	-		C, 1.0	0.99
115	100	100		1.0	1.0