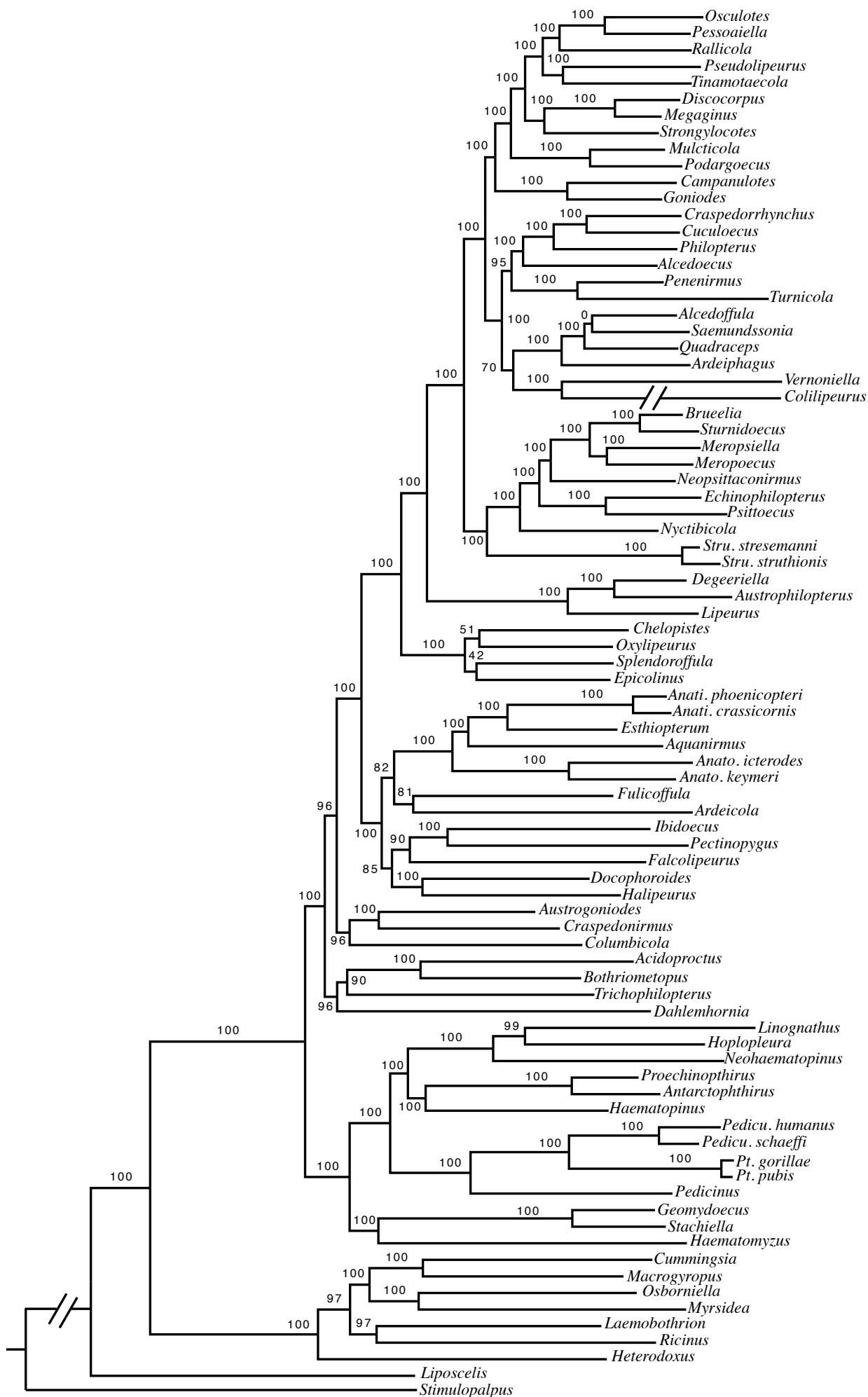
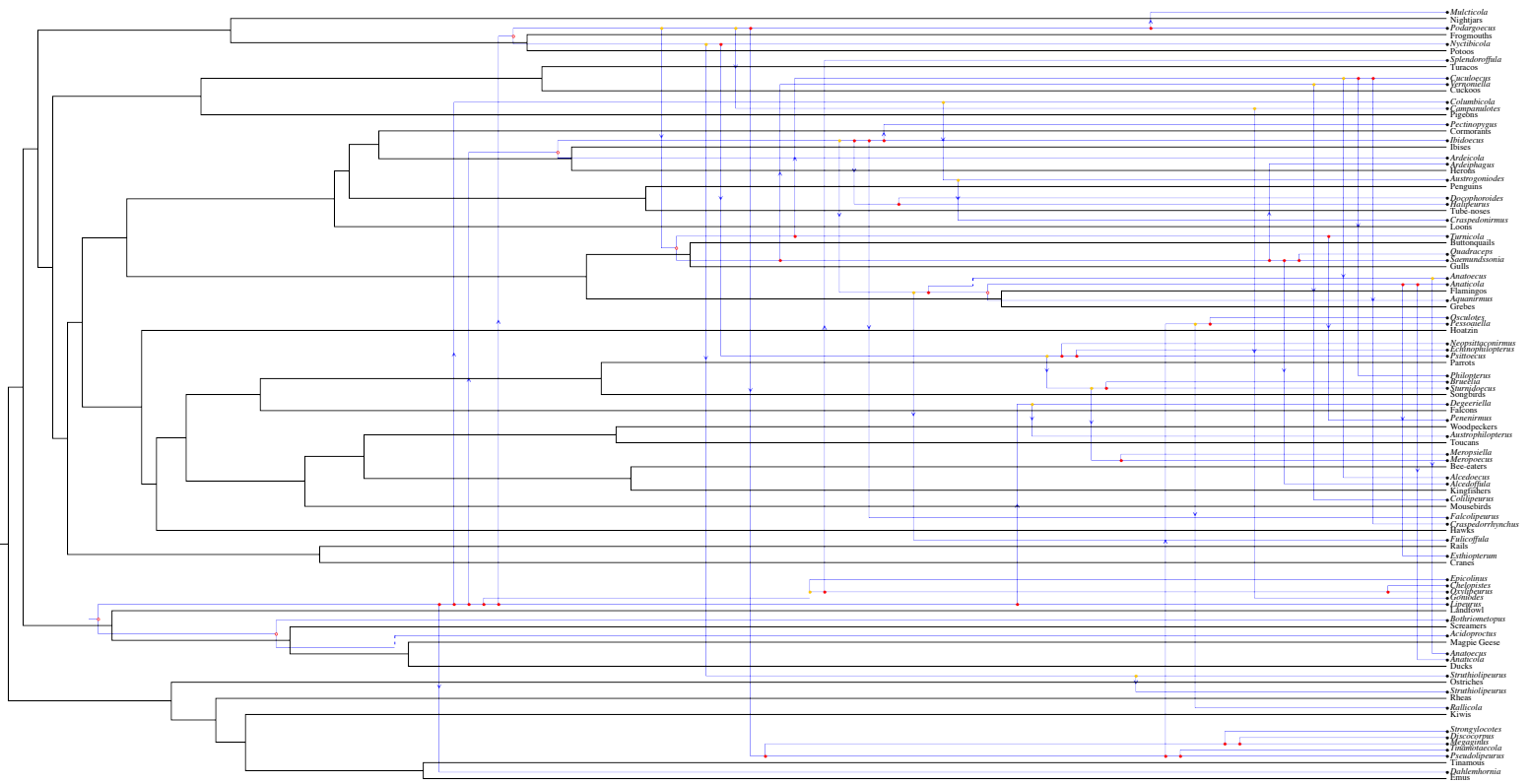


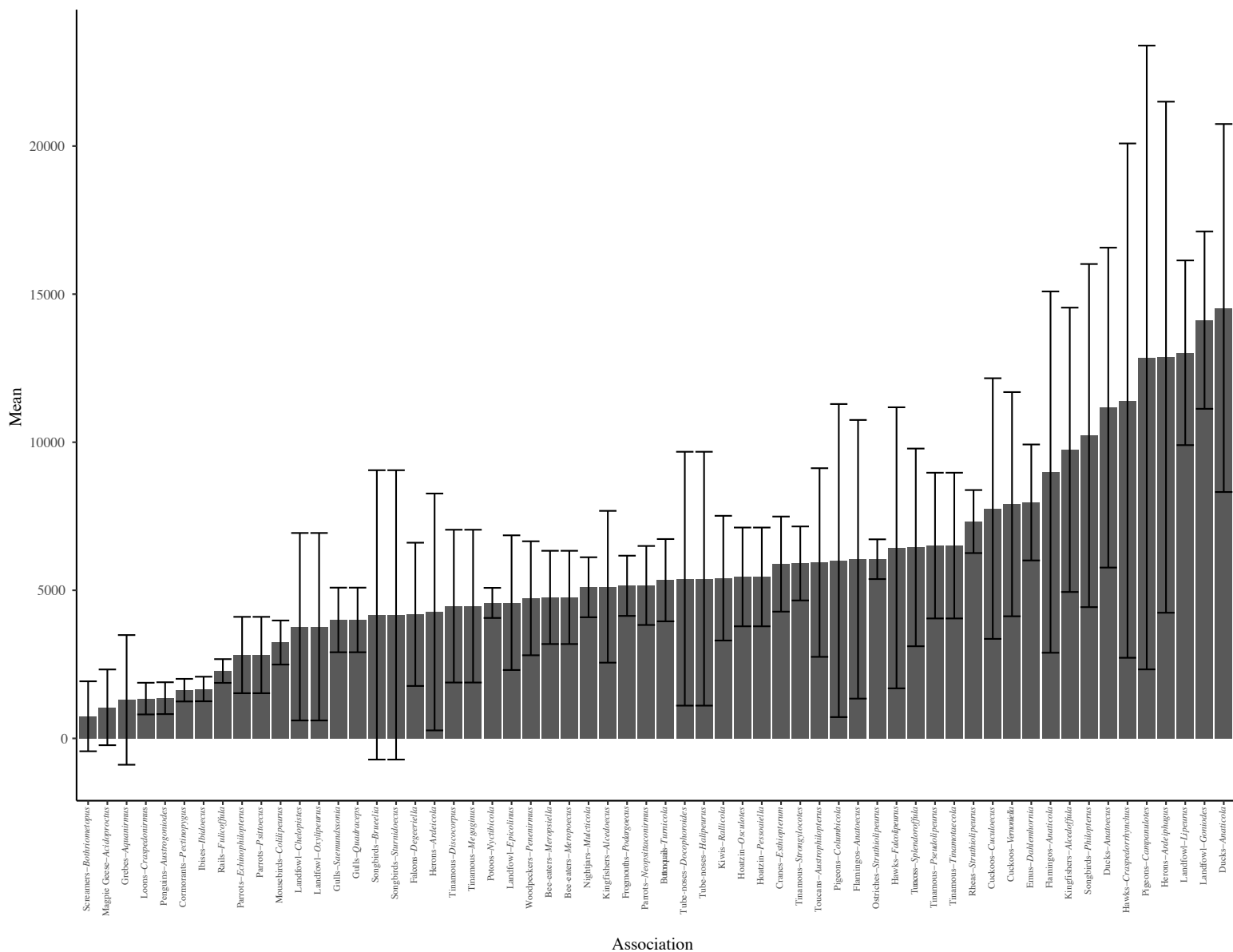
Supplementary Figures



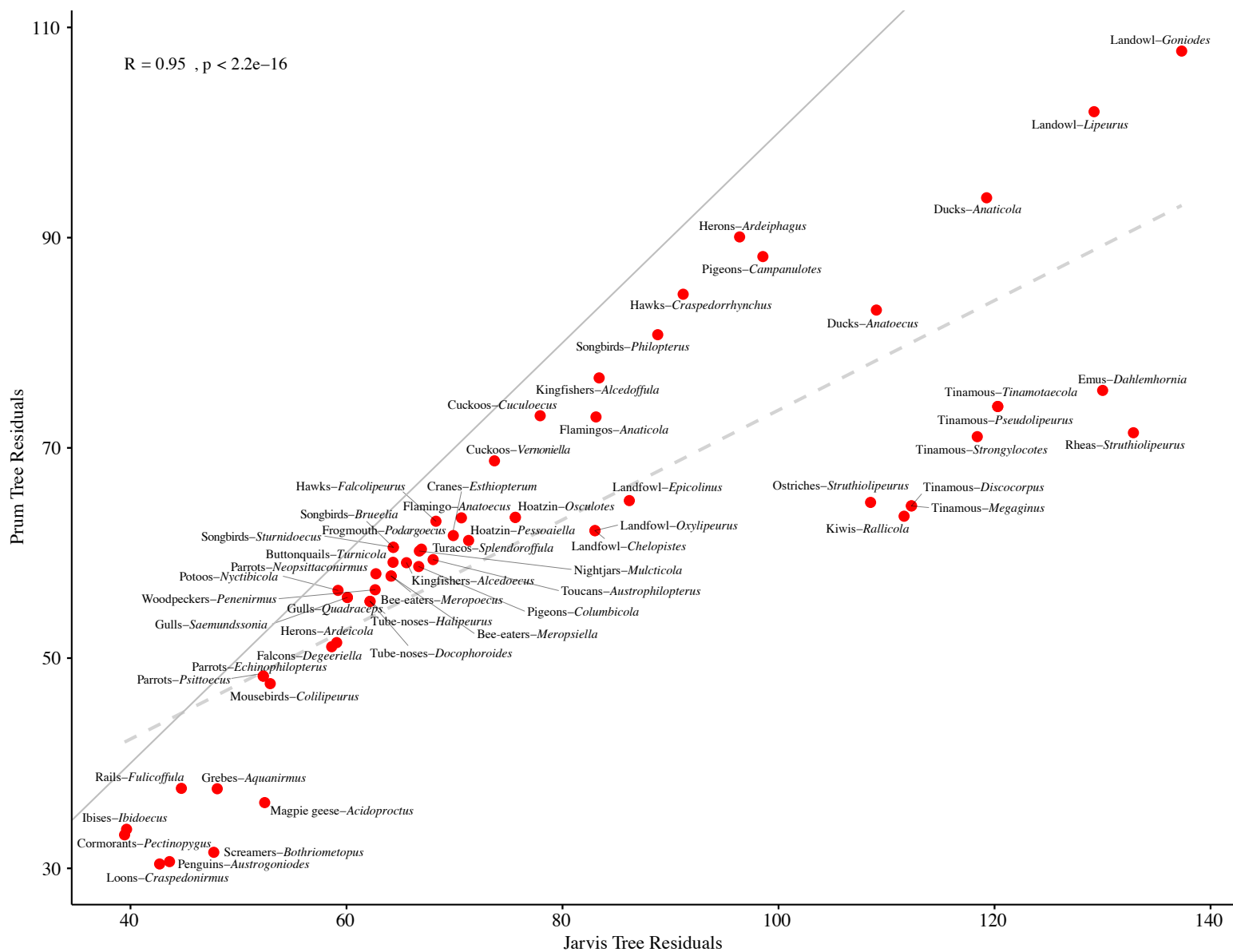
Supplementary Figure 1: Phylogeny based on nonpartitioned maximum likelihood analysis of all codon positions. Nodes labelled with bootstrap support.



Supplementary Figure 2: One possible reconstruction from the Jane cophylogenetic analysis of the Prum *et al.*² topology (black) and feather louse topology (blue) derived from the partitioned maximum likelihood analysis with 3rd codon positions removed. Blue arrows indicate directionality of a host-switch. Hollow red circles identify cospeciation events. Filled colored (red and yellow) circles identify duplication events. Dashed lines represent a loss (sorting event).



Supplementary Figure 3: Bar graph of the mean residual link scores between feather lice and their avian hosts as suggested by Paco analysis of the Prum *et al.*² topology and the feather louse maximum likelihood topology with 3rd positions removed. Whiskers and fences display the 95% confidence intervals produced that contains the true mean residual link score for each bird louse association. Statistics are derived from $n = 9,999$ permutations of the two topologies for respective bird louse associations tested.



Supplementary Figure 4: Scatter plot of residuals from Paco analysis using the Prum *et al.*² versus Jarvis *et al.*¹ avian tree topologies when analyzed with the feather louse maximum likelihood tree topology with 3rd codon positions removed. Statistics are derived from n = 9,999 permutations of the two topologies for respective bird louse associations tested.

Supplementary Tables

Table 1:

Summary of Jane cophylogenetic analyses with Prum topology:

All nucleotide sites:										
Solution ID	Cospeciations	Duplications	Duplications &		Failures to		Cost	Ancestral Host	Host-switches to	
			Host Switches	Losses	Diverge	Palaeognaths			Ancestral host prior to Palaeognath host-switch	
1	6	15	38	2	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
2	6	15	38	2	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
3	6	15	38	2	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
4	7	15	37	4	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
5	7	15	37	4	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
6	7	15	37	4	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
7	6	14	39	1	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
8	6	14	39	1	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
9	6	14	39	1	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
10	6	14	39	1	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
11	6	14	39	1	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
12	6	14	39	1	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
13	7	14	38	3	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
14	7	14	38	3	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
15	7	14	38	3	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
16	7	14	38	3	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
17	7	14	38	3	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
18	7	14	38	3	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
19	7	14	38	3	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
20	7	14	38	3	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
21	7	14	38	3	0	93	Galloanserae	4	Anseriformes, Nyctibius, Podargus, Opisthocomus	
3rd codon positions removed:										
Solution ID	Cospeciations	Duplications	Duplications &		Failures to		Cost	Ancestral Host	Host-switches to	
			Host Switches	Losses	Diverge	Palaeognaths			Ancestral host prior to Palaeognath host-switch	
22	6	15	38	2	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
23	6	15	38	2	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
24	6	15	38	2	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
25	7	15	37	4	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
26	7	15	37	4	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
27	7	15	37	4	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
28	6	14	39	1	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
29	6	14	39	1	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
30	6	14	39	1	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
31	6	14	39	1	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
32	6	14	39	1	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
33	6	14	39	1	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
34	6	14	39	1	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
35	6	14	39	1	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
36	6	14	39	1	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
37	6	14	39	1	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
38	6	14	39	1	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
39	6	14	39	1	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
40	7	14	38	3	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
41	7	14	38	3	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
42	7	14	38	3	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
43	7	14	38	3	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
44	7	14	38	3	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
45	7	14	38	3	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
46	7	14	38	3	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
47	7	14	38	3	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
48	7	14	38	3	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
49	7	14	38	3	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
50	7	14	38	3	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	
51	7	14	38	3	0	93	Galloanserae	4	Galliformes, Nyctibius, Podargus, Opisthocomus	

Supplementary Table 1: Summary of Jane solutions derived from cophylogenetic analysis of the Prun avian tree topology and the respective feather louse topology analyzed (3rd codon positions removed and all nucleotide sites).

Summary of Jane cophylogenetic analyses with Jarvis topology:

All nucleotide sites:

Solution ID	Cospeciations	Duplications &		Failures to		Cost	Host-switches to		Ancestral host prior to Palaeognath host-switch
		Duplications	Host Switches	Losses	Diverge		Palaeognaths		
52	6	14	39	1	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
53	6	14	39	1	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
54	6	14	39	1	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
55	6	14	39	1	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
56	6	14	39	1	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
57	6	14	39	1	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
58	6	14	39	1	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
59	6	14	39	1	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
60	6	14	39	1	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
61	6	14	39	1	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
62	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
63	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
64	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
65	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
66	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
67	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
68	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
69	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
70	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
71	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
72	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
73	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
74	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
75	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
76	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
77	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
78	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
79	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
80	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
81	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
82	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
83	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
84	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius
85	7	14	38	3	0	93	Galloanserae	3	Anseriformes, Podargus, Nyctibius

3rd codon positions removed:

Solution ID	Cospeciations	Duplications &		Failures to		Cost	Host-switches to		Ancestral host prior to Palaeognath host-switch
		Duplications	Host Switches	Losses	Diverge		Palaeognaths		
86	6	14	39	1	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
87	6	14	39	1	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
88	6	14	39	1	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
89	6	14	39	1	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
90	6	14	39	1	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
91	6	14	39	1	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
92	6	14	39	1	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
93	6	14	39	1	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
94	6	14	39	1	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
95	6	14	39	1	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
96	6	14	39	1	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
97	6	14	39	1	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
98	6	14	39	1	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
99	6	14	39	1	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
100	6	14	39	1	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
101	6	14	39	1	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
102	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
103	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
104	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
105	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
106	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
107	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
108	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
109	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
110	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
111	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
112	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
113	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
114	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
115	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
116	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
117	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
118	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
119	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
120	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
121	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
122	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
123	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
124	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius
125	7	14	38	3	0	93	Galloanserae	3	Galliformes, Podargus, Nyctibius

Supplementary Table 2: Summary of Jane solutions derived from cophylogenetic analysis of the Jarvis avian tree topology and the respective feather louse topology analyzed (3rd codon positions removed and all nucleotide sites).

Louse species	Family	Host	Total Reads	SRR Accession
<i>Trichophilopterus babakotophilus</i>	Philopteridae	<i>Propithecus verreauxi</i>	38,735,400	SRR5308144
<i>Dahlemhorna asymmetrica</i>	Philopteridae	<i>Dromaius novaehollandiae</i>	65,382,868	SRR5308359
<i>Struthiolipeurus stresemanni</i>	Philopteridae	<i>Rhea americana</i>	106,684,732	SRR5308383
<i>Struthiolipeurus struthionis</i>	Philopteridae	<i>Struthio camelus</i>	66,838,096	SRR5308365
<i>Rallicola</i> sp.	Philopteridae	<i>Apteryx</i> sp.	54,822,042	SRR5308364
<i>Strongylocotus lipogonus</i>	Philopteridae	<i>Rhynchotus rufescens</i>	68,077,110	SRR5308142
<i>Megaginus tataupensis</i>	Philopteridae	<i>Crypturellus tataupa</i>	58,725,252	SRR5308131
<i>Discocorpus mexicanus</i>	Philopteridae	<i>Crypturellus cinnamomeus</i>	102,833,150	SRR5308387
<i>Pseudolipeurus plumbeus</i>	Philopteridae	<i>Crypturellus tataupa</i>	45,289,078	SRR5308356
<i>Tinamotaecola elegans</i>	Philopteridae	<i>Eudromia elegans</i>	91,157,284	SRR5308366
<i>Bothriometopus macrocnemis</i>	Philopteridae	<i>Chauna torquata</i>	125,502,194	SRR5088466
<i>Acidoproctus rostratus</i>	Philopteridae	<i>Dendrocynna viduata</i>	72,086,096	SRR5308389
<i>Oxylpeurus chiniri</i>	Philopteridae	<i>Ortalis vetula</i>	71,982,724	SRR5308134
<i>Chelopistes texanus</i>	Philopteridae	<i>Ortalis vetula</i>	61,406,098	SRR5308114
<i>Epicolinus clavatus</i>	Philopteridae	<i>Colinus virginianus</i>	73,404,240	SRR5308393
<i>Goniodes ortygis</i>	Philopteridae	<i>Colinus virginianus</i>	91,028,250	SRR5308120
<i>Lipeurus caponis</i>	Philopteridae	<i>Gallus gallus</i>	56,279,502	SRR5308373
<i>Anatoecus icterodes</i>	Philopteridae	<i>Anas cyanoptera</i>	63,288,740	SRR5308111
<i>Anaticola crassicornis</i>	Philopteridae	<i>Anas clypeata</i>	108,620,518	SRR5308340
<i>Multicola bacurau</i>	Philopteridae	<i>Nyctidromus albigollis</i>	63,680,908	SRR5308374
<i>Podargoecus papuensis</i>	Philopteridae	<i>Podargus papuensis</i>	38,463,532	SRR5308376
<i>Nyctibicola longirostris</i>	Philopteridae	<i>Nyctibius jamaicensis</i>	62,948,748	SRR5308388
<i>Columbicola columbae</i>	Philopteridae	<i>Columba livia</i>	56,608,482	SRR5308115
<i>Campanulotes compar</i>	Philopteridae	<i>Columba livia</i>	117,145,090	SRR5308113
<i>Austrogoniodes waterstoni</i>	Philopteridae	<i>Eudiptula minor</i>	84,049,742	SRR5308390
<i>Craspedonirmus immer</i>	Philopteridae	<i>Gavia immer</i>	54,000,574	SRR5308116
<i>Pectinopygus varius</i>	Philopteridae	<i>Phalacrocorax varius</i>	108,163,928	SRR5308135
<i>Docaphoroides brevis</i>	Philopteridae	<i>Diomedea exulans</i>	54,250,984	SRR5308117
<i>Halipeurus diversus</i>	Philopteridae	<i>Puffinus tenuirostris</i>	52,500,588	SRR5308124
<i>Quadriceps punctatus</i>	Philopteridae	<i>Larus argentatus</i>	68,896,180	SRR5308139
<i>Saemundsonia lari</i>	Philopteridae	<i>Larus novaehollandiae</i>	38,719,424	SRR5308141
<i>Turnicola</i> sp.	Philopteridae	<i>Turnix pyrrhorax</i>	50,940,150	SRR5308379
<i>Ibidoecus bisignatus</i>	Philopteridae	<i>Plegadis chihi</i>	42,590,904	SRR5308126
<i>Esthiopterum brevcephalum</i>	Philopteridae	<i>Grus canadensis</i>	113,004,908	SRR5308385
<i>Ardeiphagus cochlearius</i>	Philopteridae	<i>Cochlearius cochlearius</i>	79,452,682	SRR5308384
<i>Ardeicola expallidus</i>	Philopteridae	<i>Bubulcus ibis</i>	75,255,050	SRR5308391
<i>Fulicoffula longipila</i>	Philopteridae	<i>Fulica americana</i>	52,765,362	SRR5308119
<i>Aquanirmus occidentalis</i>	Philopteridae	<i>Aechmophorus occidentalis</i>	92,913,330	SRR5308392
<i>Anatoecus keymeri</i>	Philopteridae	<i>Phoenicopterus chilensis</i>	98,579,854	SRR5308381
<i>Anaticola phoenicopteri</i>	Philopteridae	<i>Phoenicopterus chilensis</i>	88,679,252	SRR5308382
<i>Alcedoecus</i> sp.	Philopteridae	<i>Halcyon badia</i>	77,411,704	SRR5308110
<i>Alcedoffula alcyonae</i>	Philopteridae	<i>Ceryle alcyon</i>	41,932,010	SRR5308368
<i>Osculotes curta</i>	Philopteridae	<i>Opisthocomus hoazin</i>	81,287,340	SRR5308133
<i>Pessoaiella obsita</i>	Philopteridae	<i>Opisthocomus hoazin</i>	81,580,082	SRR5308145
<i>Splendoroffula</i> sp.	Philopteridae	<i>Tauraco porphyrelopha</i>	51,425,552	SRR5308378
<i>Austrophilopterus cancellatus</i>	Philopteridae	<i>Ramphastos sulfuratus</i>	41,088,016	SRR5308369
<i>Penenirmus auritus</i>	Philopteridae	<i>Sphyrapicus varius</i>	58,531,158	SRR5308137
<i>Cuculoecus africanus</i>	Philopteridae	<i>Chrysococcyx cupreus</i>	62,937,116	SRR5308372
<i>Vernoniella guimaraesi</i>	Philopteridae	<i>Crotophaga ani</i>	46,644,520	SRR5308380
<i>Colilipeurus obscurior</i>	Philopteridae	<i>Colius colius</i>	19,267,276	SRR5308370
<i>Falcolipeurus marginalis</i>	Philopteridae	<i>Cathartes aura</i>	46,795,498	SRR5308118
<i>Craspedorrhynchus subhaematopus</i>	Philopteridae	<i>Accipiter cooperii</i>	95,915,920	SRR5308371
<i>Meropoecus</i> sp.	Philopteridae	<i>Merops ornatus</i>	65,254,844	SRR5308363
<i>Meropsiella</i> sp.	Philopteridae	<i>Merops ornatus</i>	37,474,984	SRR5308362
<i>Degeeriella rufa</i>	Philopteridae	<i>Falco berigora</i>	55,640,798	SRR5308223
<i>Neopsittaconirmus bushae</i>	Philopteridae	<i>Northiella haematogaster</i>	75,033,324	SRR5308361
<i>Echinophilopterus claytoni</i>	Philopteridae	<i>Northiella haematogaster</i>	61,015,492	SRR5308360
<i>Psittococcus</i> sp.	Philopteridae	<i>Cacatua galerita</i>	46,060,548	SRR5308377
<i>Philopterus</i> sp.	Philopteridae	<i>Tyrannus melancholicus</i>	53,357,072	SRR5308375
<i>Brueelia antiqwa</i>	Philopteridae	<i>Catharus ustulatus</i>	140,698,976	SRR5308112
<i>Sturnidoecus</i> sp.	Philopteridae	<i>Lamprolornis purpureus</i>	59,263,926	SRR5308357
<i>Geomydoecus aurei</i>	Trichodectidae	<i>Thomomys bottae</i>	116,680,800	SRR5308121
<i>Stachiella larseni</i>	Trichodectidae	<i>Mustela vison</i>	38,023,782	SRR5308143
<i>Haematomyzus elephantis</i>	Haematomyzidae	<i>Elephas maximus</i>	94,800,010	SRR5308122
<i>Haematopinus euryternus</i>	Haematopinidae	<i>Bos</i> sp.	64,236,000	SRR5308123
<i>Linognathus spicatus</i>	Linognathidae	<i>Connocchaetes taurinus</i>	42,855,056	SRR5308129
<i>Proechinophthirus fluctus</i>	Echinophthiriidae	<i>Callorhinus ursinus</i>	42,855,056	SRR5308138
<i>Antarctophthirus microchir</i>	Echinophthiriidae	<i>Otaria flavescens</i>	104,403,830	SRR5088465
<i>Hoplopleura arboricola</i>	Hoplopleuridae	<i>Tamias amoenus</i>	174,420,882	SRR5088468
<i>Neohaematopinus pacificus</i>	Polyplacidae	<i>Tamias minimus</i>	154,952,566	SRR5088469
<i>Pedicinus badius</i>	Pedicinidae	<i>Procolobus rufoimtratus</i>	49,099,090	SRR5308136
<i>Pthirus pubis</i>	Pthiridae	<i>Homo sapiens</i>	91,120,738	SRR5088475
<i>Pthirus gorillae</i>	Pthiridae	<i>Gorilla beringei</i>	120,850,588	SRR5088474
<i>Pediculus schaeffi</i>	Pediculidae	<i>Pan troglodytes</i>	128,404,204	SRR1182279
<i>Pediculus humanus</i>	Pediculidae	<i>Homo sapiens</i>	147,704,686	SRR5088472
<i>Ricinus</i> sp.	Ricinidae	<i>Myiothlypis luteoviridis</i>	29,736,768	SRR5308140
<i>Laemobothrion tinnunculi</i>	Laemobothriidae	<i>Falco longipennis</i>	86,487,406	SRR5308127
<i>Myrsidea</i> sp.	Menoponidae	<i>Myiothlypis luteoviridis</i>	35,830,404	SRR5308132
<i>Osborniella crotophagae</i>	Menoponidae	<i>Crotophaga ani</i>	85,359,404	SRR5088470
<i>Macrogryopus costalimai</i>	Gyropidae	<i>Cuniculus paca</i>	65,940,856	SRR5308130
<i>Cummingsia maculata</i>	Trimenoponidae	<i>Lestoros inca</i>	66,405,676	SRR5308146
<i>Heterodoxus spiniger</i>	Boopidae	<i>Canis lupus</i>	84,240,242	SRR5308125
<i>Liposcelis brunnea</i>	Liposcelidae	Non-parasitic	74,000,000	SRR5308128
<i>Stimulopalpus japonicus</i>	Amphientomidae	Non-parasitic	148,541,936	SRR5088476

Supplementary Table 3: Summary of taxonomic sampling and host associations of individual feather lice collected. Number of reads produced from Illumina sequencing and SRR accession numbers are summarized.

References

1. Jarvis, E. D. *et al.* Whole-genome analyses resolve early branches in the tree of life of modern birds. *Science* **346**, 1320–1331 (2014).
2. Prum, R. O. *et al.* A comprehensive phylogeny of birds (Aves) using targeted next-generation DNA sequencing. *Nature* **526**, 569–573 (2015).