

Supplementary tables

Table S1 Information of Cynipoidea species in this study.

Voucher specimens	Species	Sex	Locality	Date	Collector/Collect type	Identification reference
ZJUH_20220001	<i>Anacharis</i> sp.	Male	Zhangjiakou, Hebei, China	21-AUG-2005	Ying Zhang	[1,2]
ZJUH_20220002	<i>Aegilips</i> sp.	Male	Shangrao, Jiangxi, China	07-AUG-2015	Ning Mao	[1,2]
ZJUH_20220003	<i>Melanips</i> sp.	Male	Linshi, Tibet, China	09-AUG-2019	Xiaofei Li	[1,2]
ZJUH_20220004	<i>Prosaspicera validispina</i>	Female	Linshi, Tibet, China	16-AUG-2019	Zekai Li	[1-3]
ZJUH_20220005	<i>Pujadella villari</i>	Female	Linshi, Tibet, China	16-AUG-2019	Zekai Li	[1,4]
ZJUH_20220006	<i>Endecameris</i> sp.	Female	Tianmu Mountain, Zhejiang, China	29-AUG-2018	Zekai Li	[1,2]
ZJUH_20220007	<i>Ganaspini</i> sp.	Male	Linshi, Tibet, China	04-AUG-2019	Yellow trap	[1,2]
ZJUH_20220008	<i>Trybliographa</i> sp.	Male	Linshi, Tibet, China	16-AUG-2019	Xingzhou Ma	[1,2]
ZJUH_20220009	<i>Figites</i> sp. 1	Male	Foping, Shaanxi, China	28-JUL-2013	Jiangli Tan	[1,2]
ZJUH_20220010	<i>Figites</i> sp. 2	Male	Sonoma, California, USA	07-SEP-2008	Peter H. Kerr	[1,2]
ZJUH_20220011	<i>Ibalia</i> sp.	Male	Shangrao, Jiangxi, China	07-AUG-2015	Shajiangyang Mao	[1,5]
ZJUH_20220012	<i>Paramblynotus</i> sp.	Female	Republic of the Congo	20-OCT-2008	Yves Braet, Michael J. Sharkey	[1,2]
ZJUH_20220013	<i>Oberthuerella sharkeyi</i>	Female	Republic of the Congo	20-OCT-2008	Yves Braet, Michael J. Sharkey	[1,2,6]
ZJUH_20220014	<i>Tessmannella kiplingi</i>	Female	Rikaze, Tiibet, China	01-AUG-2018	Yellow trap	[1,2,6]
ZJUH_20220015	<i>Saphonecrus</i> sp.	Female	Shangrao, Jiangxi, China	07-AUG-2015	Shajiangyang Mao	[1,7,8]
ZJUH_20220016	<i>Parnips nigripes</i>	Female	Madrid, Spain	19-MAR-2001	Jose Luis Nieves-Aldrey	[1,9]

Table S2 The best schemes of partition and substitution models selected in nucleotide dataset.

Optimal partition	Best Model	Partition names
Partition 1	GTR+I+G	c1p2, c3p2, c2p2, a6p2
Partition 2	GTR+I+G	n4lp1, n5p1, n4p1, n3p1
Partition 3	GTR+I+G	cbp1, n1p1, c2p1, a6p1, c3p1
Partition 4	GTR+G	c1p3, c3p3, a6p3, c2p3, n3p3
Partition 5	GTR+I+G	n6p1, n2p1, a8p1
Partition 6	GTR+I+G	n4p2, n4lp2, n5p2, n3p2
Partition 7	GTR+G	n6p3, n1p3, cbp3
Partition 8	GTR+G	n5p3, n4lp3, n4p3
Partition 9	GTR+I+G	n1p2, cbp2
Partition 10	GTR+I+G	a8p2, n6p2, n2p2
Partition 11	GTR+I+G	c1p1
Partition 12	GTR+G	n2p3, a8p3

Table S3 Length of tRNA and rRNA genes in the Cynipoidea mitogenomes.

Species	trnW	trnC	trnY	trnL2	trnK	trnD	trnG	trnA	trnR	trnN	trnS1	trnE	trnF	trnH	trnT	trnP	trnL1	trnS2	trnV	trnI	trnQ	trnM	rrnS	rrnL
<i>Anacharis</i> sp.	72	63	71	71	72	70	70	69	69	70	62	69	66	74	67	72	71	69	73	71	67	69	824	1406
<i>Aegilips</i> sp.	67	68	67	65	69	67	66	57	65	65	61	72	64	65	64	72	67	69	66	68	68	68	792	1360
<i>Melanips</i> sp.	65	68	72	69	77	68	73	72	70	67	63	67	68	73	67	68	71	67	70	68	71	72	877	1418
<i>Prospicera validispina</i>	68	69	68	67	70	66	54	66	63	65	63	68	69	69	63	64	65	67	67	69	70	67	848	1360
<i>Pujadella villari</i>	67	69	72	69	69	70	68	68	71	67	64	68	65	53	67	66	68	68	70	66	-	68	858	1464
<i>Endecameris</i> sp.	72	65	65	65	72	74	69	66	69	65	60	68	66	71	65	77	67	67	66	65	71	63	879	1361
<i>Ganaspini</i> sp.	68	64	68	68	67	73	68	67	70	73	63	69	64	65	61	69	70	67	68	70	70	67	822	1231
<i>Trybliographa</i> sp.	67	66	65	66	69	69	66	66	66	67	60	65	-	68	65	67	68	68	66	69	67	67	812	1375
<i>Figites</i> sp. 1	-	72	69	68	68	68	68	64	67	-	65	66	65	64	65	69	-	-	-	65	70	67	811	1384
<i>Figites</i> sp. 2	66	67	70	69	69	68	66	66	68	69	63	66	67	66	67	66	-	68	70	66	69	70	813	1355
<i>Ibalia</i> sp.	69	64	66	67	72	69	67	65	67	68	64	68	63	65	65	69	74	68	59	67	67	75	811	1357
<i>Paramblynotus</i> sp.	66	63	64	66	64	64	67	66	70	67	60	69	64	66	63	65	69	66	69	65	69	68	824	1287
<i>Oberthuerella sharkeyi</i>	52	67	67	67	75	68	66	64	72	69	62	66	64	67	63	58	69	68	69	68	71	68	815	1362
<i>Tessmannella kiplingi</i>	69	63	68	67	69	65	69	69	67	66	62	58	62	53	62	55	69	64	71	70	70	67	793	1362
<i>Saphonecrus</i> sp.	68	69	65	70	69	66	68	70	69	69	64	67	67	69	64	66	72	68	68	69	70	68	844	1409
<i>Parnips nigripes</i>	70	67	67	67	70	68	66	65	65	66	64	67	65	69	64	66	54	68	66	69	70	70	821	1377

Table S4 Relative synonymous codon usage in 16 Cynipoidea mitogenomes.

	<i>Anacharis</i> sp.		<i>Aegilips</i> sp.		<i>Melanips</i> sp.		<i>Prosaspicera</i> <i>validispina</i>		<i>Pujadella</i> <i>villari</i>		<i>Endecameris</i> sp.		Ganaspini sp.		<i>Trybliographa</i> sp.		<i>Figites</i> sp. 1		<i>Figites</i> sp. 2		<i>Ibalia</i> sp.		<i>Paramblynotus</i> sp.		<i>Oberthuerella</i> <i>sharkeyi</i>		<i>Tessmannella</i> <i>kiplingi</i>		<i>Saphonecrus</i> sp.		<i>Parnips</i> <i>nigripes</i>	
Codon	Count	RSCU	Count	RSCU	Count	RSCU	Count	RSCU	Count	RSCU	Count	RSCU	Count	RSCU	Count	RSCU	Count	RSCU	Count	RSCU	Count	RSCU	Count	RSCU	Count	RSCU	Count	RSCU	Count	RSCU	Count	RSCU
UUU(F)	285	1.67	309	1.84	343	1.88	346	1.94	307	1.77	329	1.81	320	1.78	287	1.68	309	1.9	297	1.79	374	1.96	387	1.95	345	1.78	320	1.77	348	1.89	343	1.93
UUC(F)	57	0.33	26	0.16	22	0.12	10	0.06	39	0.23	34	0.19	39	0.22	55	0.32	17	0.1	35	0.21	8	0.04	10	0.05	42	0.22	42	0.23	20	0.11	12	0.07
UUA(L)	368	4.36	450	4.93	500	5.42	469	5.15	381	4.24	479	5.16	446	4.8	406	4.27	502	5.47	473	5.12	553	5.78	591	5.68	482	4.98	435	4.68	503	5.18	553	5.6
UUG(L)	30	0.36	20	0.22	11	0.12	21	0.23	55	0.61	11	0.12	19	0.2	31	0.33	16	0.17	16	0.17	7	0.07	3	0.03	36	0.37	30	0.32	20	0.21	12	0.12
CUU(L)	38	0.45	40	0.44	21	0.23	32	0.35	37	0.41	38	0.41	40	0.43	43	0.45	12	0.13	30	0.32	11	0.11	20	0.19	24	0.25	24	0.26	24	0.25	15	0.15
CUC(L)	5	0.06	9	0.1	3	0.03	3	0.03	3	0.03	3	0.03	5	0.05	10	0.11	1	0.01	3	0.03	0	0	0	0	0	0	3	0.03	3	0.03	1	0.01
CUA(L)	66	0.78	29	0.32	18	0.2	20	0.22	62	0.69	26	0.28	46	0.5	79	0.83	20	0.22	32	0.35	3	0.03	10	0.1	34	0.35	66	0.71	33	0.34	10	0.1
CUG(L)	0	0	0	0	0	0	1	0.01	1	0.01	0	0	1	0.01	1	0.01	0	0	0	0	0	0	0	0	5	0.05	0	0	0	0	2	0.02
AUU(I)	376	1.66	442	1.86	482	1.97	446	1.89	388	1.83	515	1.93	457	1.81	387	1.65	449	1.92	456	1.88	505	1.98	498	1.96	477	1.87	452	1.83	463	1.9	459	1.94
AUC(I)	76	0.34	33	0.14	7	0.03	25	0.11	36	0.17	19	0.07	49	0.19	82	0.35	18	0.08	29	0.12	6	0.02	9	0.04	32	0.13	41	0.17	24	0.1	15	0.06
AUA(M)	418	1.86	377	1.91	394	1.92	399	1.89	349	1.79	386	1.94	389	1.92	335	1.84	382	1.92	376	1.87	394	1.95	363	1.97	340	1.85	370	1.88	381	1.85	374	1.93
AUG(M)	31	0.14	18	0.09	16	0.08	23	0.11	42	0.21	11	0.06	17	0.08	29	0.16	16	0.08	26	0.13	10	0.05	6	0.03	28	0.15	24	0.12	32	0.15	14	0.07
GUU(V)	53	1.51	59	2.05	54	1.85	68	2.05	77	1.82	33	1.43	61	2.05	67	1.85	66	1.96	69	2.08	28	1.38	37	1.92	48	1.94	67	2.39	58	2.37	44	1.56
GUC(V)	8	0.23	3	0.1	1	0.03	5	0.15	5	0.12	1	0.04	2	0.07	7	0.19	1	0.03	1	0.03	1	0.05	3	0.16	6	0.24	2	0.07	4	0.16	6	0.21
GUA(V)	71	2.03	48	1.67	59	2.02	59	1.77	62	1.47	56	2.43	52	1.75	57	1.57	68	2.01	56	1.68	50	2.47	36	1.87	44	1.78	36	1.29	33	1.35	60	2.12
GUG(V)	8	0.23	5	0.17	3	0.1	1	0.03	25	0.59	2	0.09	4	0.13	14	0.39	0	0	7	0.21	2	0.1	1	0.05	1	0.04	7	0.25	3	0.12	3	0.11
UCU(S)	86	1.99	71	1.85	100	2.52	83	2	86	2.1	71	1.62	78	1.79	83	1.95	74	1.9	74	1.76	127	3.32	63	1.69	63	1.57	63	1.6	90	2.29	100	2.61
UCC(S)	10	0.23	17	0.44	5	0.13	4	0.1	24	0.59	20	0.46	24	0.55	32	0.75	6	0.15	12	0.29	4	0.1	5	0.13	10	0.25	13	0.33	9	0.23	4	0.1
UCA(S)	122	2.83	117	3.05	105	2.64	126	3.04	72	1.76	143	3.26	119	2.73	95	2.24	124	3.18	123	2.93	62	1.62	127	3.4	129	3.2	115	2.92	106	2.7	85	2.22
UCG(S)	0	0	0	0	0	0	4	0.1	12	0.29	0	0	0	0	1	0.02	0	0	2	0.05	0	0	1	0.03	4	0.1	5	0.13	1	0.03	0	0
CCU(P)	66	1.89	54	2.08	67	2.58	61	2.44	55	1.93	40	1.58	49	1.9	44	1.41	64	2.42	62	2.34	80	3.23	57	2.3	48	1.83	51	2	56	2.15	69	2.65
CCC(P)	23	0.66	9	0.35	5	0.19	2	0.08	16	0.56	5	0.2	15	0.58	37	1.18	2	0.08	4	0.15	2	0.08	3	0.12	9	0.34	11	0.43	9	0.35	1	0.04
CCA(P)	48	1.37	40	1.54	32	1.23	35	1.4	35	1.23	55	2.18	38	1.48	42	1.34	39	1.47	38	1.43	16	0.65	39	1.58	46	1.75	36	1.41	39	1.5	33	1.27
CCG(P)	3	0.09	1	0.04	0	0	2	0.08	8	0.28	1	0.04	1	0.04	2	0.06	1	0.04	2	0.08	1	0.04	0	0	2	0.08	4	0.16	0	0	1	0.04
ACU(T)	59	1.33	54	1.33	52	1.65	60	1.94	76	2.11	47	1.59	61	1.72	75	1.55	61	1.92	64	1.88	57	2.07	35	1.32	50	1.67	51	1.7	54	1.7	65	2.1
ACC(T)	25	0.56	16	0.4	2	0.06	5	0.16	17	0.47	8	0.27	22	0.62	50	1.04	3	0.09	8	0.24	1	0.04	2	0.08	6	0.2	15	0.5	9	0.28	4	0.13
ACA(T)	91	2.06	91	2.25	71	2.25	58	1.87	46	1.28	63	2.14	59	1.66	66	1.37	63	1.98	61	1.79	51	1.85	67	2.53	62	2.07	53	1.77	64	2.02	52	1.68
ACG(T)	2	0.05	1	0.02	1	0.03	1	0.03	5	0.14	0	0	0	0	2	0.04	0	0	3	0.09	1	0.04	2	0.08	2	0.07	1	0.03	0	0	3	0.1
GCU(A)	30	1.71	31	1.8	44	2.48	39	2.11	43	2.02	35	2.09	24	1.5	22	1.38	33	2	36	2.03	39	2.48	34	2.34	29	1.97	31	1.97	26	1.93	54	2.73
GCC(A)	9	0.51	10	0.58	6	0.34	2	0.11	16	0.75	4	0.24	10	0.63	15	0.94	2	0.12	7	0.39	2	0.13	0	0	3	0.2	5	0.32	2	0.15	2	0.1
GCA(A)	31	1.77	27	1.57	20	1.13	31	1.68	22	1.04	28	1.67	28	1.75	25	1.56	30	1.82	26	1.46	22	1.4	24	1.66	25	1.69	24	1.52	26	1.93	23	1.16
GCG(A)	0	0	1	0.06	1	0.06	2	0.11	4	0.19	0	0	2	0.13	2	0.13	1	0.06	2	0.11	0	0	0	0	2	0.14	3	0.19	0	0	0	0
UAU(Y)	162	1.62	170	1.63	206	1.89	191	1.82	178	1.66	190	1.83	148	1.71	118	1.37	187	1.79	179	1.76	210	1.94	208	1.93	197	1.82	187	1.72	199	1.82	218	1.95
UAC(Y)	38	0.38	38	0.37	12	0.11	19	0.18	37	0.34	18	0.17	25	0.29	54	0.63	22	0.21	24	0.24	7	0.06	7	0.07	20	0.18	31	0.28	20	0.18	6	0.05
CAU(H)	48	1.45	49	1.63	63	1.97	52	1.73	49	1.53	50	1.72	49	1.61	48	1.5	60	1.9	61	1.94	54	1.93	60	1.94	51	1.67	59	1.87	48	1.55	58	1.9
CAC(H)	18	0.55	11	0.37	1	0.03	8	0.27	15	0.47	8	0.28	12	0.39	16	0.5	3	0.1	2	0.06	2	0.07	2	0.06	10	0.33	4	0.13	14	0.45	3	0.1
CAA(Q)	47																															

CAG(Q)	4	0.16	1	0.04	2	0.09	6	0.27	12	0.51	5	0.19	1	0.04	5	0.18	3	0.12	8	0.33	1	0.04	0	0	4	0.15	13	0.46	5	0.21	1	0.04
AAU(N)	222	1.71	275	1.82	265	1.96	263	1.93	203	1.59	242	1.85	216	1.72	181	1.58	258	1.89	232	1.8	277	1.99	299	1.97	244	1.83	259	1.77	262	1.83	265	1.91
AAC(N)	37	0.29	27	0.18	6	0.04	10	0.07	52	0.41	20	0.15	35	0.28	48	0.42	15	0.11	26	0.2	2	0.01	5	0.03	23	0.17	34	0.23	24	0.17	12	0.09
AAA(K)	137	1.81	133	1.89	139	1.93	136	1.96	119	1.72	128	1.91	147	1.87	128	1.83	137	1.89	140	1.85	148	1.97	143	1.97	148	1.85	136	1.79	154	1.93	138	1.97
AAG(K)	14	0.19	8	0.11	5	0.07	3	0.04	19	0.28	6	0.09	10	0.13	12	0.17	8	0.11	11	0.15	2	0.03	2	0.03	12	0.15	16	0.21	6	0.08	2	0.03
GAU(D)	40	1.4	55	1.75	57	2	54	1.89	51	1.67	49	1.75	41	1.55	39	1.42	52	1.93	50	1.82	49	2	55	1.96	52	1.82	49	1.85	53	1.83	51	1.89
GAC(D)	17	0.6	8	0.25	0	0	3	0.11	10	0.33	7	0.25	12	0.45	16	0.58	2	0.07	5	0.18	0	0	1	0.04	5	0.18	4	0.15	5	0.17	3	0.11
GAA(E)	68	1.89	53	1.8	61	1.82	64	1.91	37	1.17	70	1.94	67	1.81	54	1.66	64	1.78	53	1.61	68	2	61	1.97	61	1.82	47	1.52	55	1.59	65	1.91
GAG(E)	4	0.11	6	0.2	6	0.18	3	0.09	26	0.83	2	0.06	7	0.19	11	0.34	8	0.22	13	0.39	0	0	1	0.03	6	0.18	15	0.48	14	0.41	3	0.09
UGU(C)	23	1.53	28	1.93	23	1.92	21	1.83	23	1.64	19	2	24	2	22	1.76	23	1.92	25	2	25	1.92	25	1.85	24	1.71	23	1.84	29	1.87	30	2
UGC(C)	7	0.47	1	0.07	1	0.08	2	0.17	5	0.36	0	0	0	0	3	0.24	1	0.08	0	0	1	0.08	2	0.15	4	0.29	2	0.16	2	0.13	0	0
UGA(W)	70	1.65	73	1.85	70	1.77	73	1.87	54	1.32	79	1.95	70	1.79	68	1.74	71	1.87	67	1.72	75	1.97	70	1.92	69	1.89	63	1.7	68	1.84	76	1.95
UGG(W)	15	0.35	6	0.15	9	0.23	5	0.13	28	0.68	2	0.05	8	0.21	10	0.26	5	0.13	11	0.28	1	0.03	3	0.08	4	0.11	11	0.3	6	0.16	2	0.05
CGU(R)	10	1	11	1.13	18	1.71	19	1.85	5	0.48	8	0.8	14	1.37	15	1.54	18	1.85	13	1.27	12	1.2	10	0.98	14	1.3	18	1.71	12	1.23	11	1.07
CGC(R)	3	0.3	1	0.1	1	0.1	0	0	1	0.1	3	0.3	2	0.2	2	0.21	0	0	1	0.1	0	0	0	0	0	0	0	0	0	0	0	0
CGA(R)	24	2.4	24	2.46	21	2	19	1.85	23	2.19	26	2.6	23	2.24	20	2.05	21	2.15	23	2.24	28	2.8	31	3.02	29	2.7	21	2	23	2.36	28	2.73
CGG(R)	3	0.3	3	0.31	2	0.19	3	0.29	13	1.24	3	0.3	2	0.2	2	0.21	0	0	4	0.39	0	0	0	0	0	0	3	0.29	4	0.41	2	0.2
AGU(S)	26	0.6	22	0.57	24	0.6	25	0.6	25	0.61	18	0.41	25	0.57	24	0.56	19	0.49	32	0.76	16	0.42	19	0.51	23	0.57	23	0.58	28	0.71	19	0.5
AGC(S)	1	0.02	0	0	0	0	2	0.05	9	0.22	1	0.02	2	0.05	7	0.16	2	0.05	1	0.02	1	0.03	2	0.05	4	0.1	6	0.15	3	0.08	1	0.03
AGA(S)	84	1.95	68	1.77	81	2.04	80	1.93	68	1.66	93	2.12	81	1.86	75	1.76	75	1.92	72	1.71	93	2.43	79	2.11	82	2.04	71	1.8	65	1.66	94	2.46
AGG(S)	16	0.37	12	0.31	3	0.08	8	0.19	32	0.78	5	0.11	20	0.46	23	0.54	12	0.31	20	0.48	3	0.08	3	0.08	7	0.17	19	0.48	12	0.31	3	0.08
GGU(G)	27	0.73	29	0.84	31	0.83	36	0.97	27	0.64	38	1.13	34	0.93	30	0.75	45	1.15	40	1.01	15	0.43	24	0.69	39	1.08	37	0.99	33	0.87	24	0.64
GGC(G)	4	0.11	1	0.03	2	0.05	4	0.11	15	0.36	6	0.18	3	0.08	11	0.28	4	0.1	4	0.1	1	0.03	1	0.03	3	0.08	6	0.16	0	0	4	0.11
GGA(G)	86	2.32	83	2.41	98	2.61	90	2.43	31	0.74	86	2.55	66	1.81	72	1.8	77	1.97	78	1.97	115	3.33	95	2.73	86	2.39	55	1.48	87	2.3	97	2.6
GGG(G)	31	0.84	25	0.72	19	0.51	18	0.49	95	2.26	5	0.15	43	1.18	47	1.18	30	0.77	36	0.91	7	0.2	19	0.55	16	0.44	51	1.37	31	0.82	24	0.64

Table S5 Synonymous and nonsynonymous substitutional analysis of 13 proteinencoding genes.

Genes	Ka	Ks	Ka/Ks
<i>atp6</i>	0.2160	0.4137	0.5221
<i>atp8</i>	0.3506	0.3096	1.1322
<i>cox1</i>	0.1177	0.4494	0.2619
<i>cox2</i>	0.2184	0.4107	0.5318
<i>cox3</i>	0.2036	0.4434	0.4592
<i>cytb</i>	0.1859	0.3787	0.4909
<i>nad1</i>	0.1972	0.3913	0.5039
<i>nad2</i>	0.3319	0.3241	1.0243
<i>nad3</i>	0.2038	0.4126	0.4940
<i>nad4</i>	0.2362	0.3696	0.6391
<i>nad4L</i>	0.2648	0.3498	0.7568
<i>nad5</i>	0.2309	0.3909	0.5908
<i>nad6</i>	0.3041	0.2566	1.1850

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