

# ***Apostasia* mitochondrial genome analysis and monocot mitochondria phylogenomics**

Shijie Ke<sup>1</sup>, Ding-Kun Liu<sup>1</sup>, Xiong-De Tu<sup>1</sup>, Xin He<sup>1</sup>, Meng-Meng Zhang<sup>1</sup>, Meng-Jia Zhu<sup>1</sup>, Diyang Zhang<sup>2</sup>, Cuili Zhang<sup>2</sup>, Siren Lan<sup>1,2\*</sup> and Zhong-Jian Liu<sup>2\*</sup>

<sup>1</sup>Fujian Colleges and Universities Engineering Research Institute of Conservation and Utilization of Natural Bioresources, College of Forestry, Fujian Agriculture and Forestry University, Fuzhou, 350002, China

<sup>2</sup>Key Laboratory of Orchid Conservation and Utilization of National Forestry and Grassland Administration at College of Landscape Architecture and Art, Fujian Agriculture and Forestry University, Fuzhou, 350002, China

3210422027@fafu.edu.cn (S.K.); fjliudk@163.com (D.-K.L.); tttxd163@163.com (X.-D.T); 5220422102@fafu.edu.cn (X.H.); 1220428020@fafu.edu.cn (M.-M.Z.); 1200455008@fafu.edu.cn (M.-J.Z.); diyangzhang@126.com (D.Z.); cuilizhang@fafu.edu.cn (C.Z.).

\*Correspondence: zjliu@fafu.edu.cn; lkzx@fafu.edu.cn;

## Supplementary Tables

Table S1 Gene content of *A.shenzhenica* mitogenome.

Group of Genes	Gene Name
Complex I (NADH dehydrogenase)	<i>nad1, nad3, nad4L(2), nad5*, nad6, nad7****, nad9</i>
Complex III (ubiquinol cytochrome c reductase)	<i>cob</i>
Complex IV (cytochrome c oxidase)	<i>cox1, cox3</i>
Complex V (ATP synthase)	<i>atp1, atp4, atp6, atp8, atp9</i>
Cytochrome c biogenesis	<i>ccmB, ccmC, ccmFc*, ccmFn</i>
Ribosomal proteins (SSU)	<i>rps1, rps3*, rps4, rps7, rps10*, rps11, rps12, rps13, rps14, rps19</i>
Ribosomal proteins (LSU)	<i>rpl2*, rpl5, rpl16</i>
Maturases	<i>matR</i>
Transport membrane protein	<i>mttB</i>
Ribosomal RNAs	<i>rrn5, rrn18</i>
Transfer RNAs	<i>trnD-GUC, trnA-Met, trnQ-UUG, trnM-CAU, trnK-UUU, trnQ-UUG, trnP-UGG, trnW-CCA, trnA-Cys, trnF-GAA, trnA-Phe, trnA-Lys, trnA-Gln, trnA-Thr, trnY-GUA, trnC-GCA</i>
Cp transfer proteins	<i>psaJ</i>

\*: represents intron number; Gene(2): Number of copies of multi-copy genes.

Table S2. Ka/Ks analysis of *A.shenzhenica* and four different Asparagales genes;

Seq1	Seq2	Gene	Ks	Ka	Ka/Ks
<i>Apostasia shenzhenica</i>	<i>Allium cepa</i>	<i>atp1</i>	0.1098	0.0232	0.2113
<i>Apostasia shenzhenica</i>	<i>Gastrodia elata</i>	<i>atp1</i>	0.0497	0.0187	0.376257545
<i>Apostasia shenzhenica</i>	<i>Hemerocallis citrina</i>	<i>atp1</i>	0.0762	0.0088	0.115485564
<i>Apostasia shenzhenica</i>	<i>Asparagus officinalis</i>	<i>atp1</i>	0.0646	0.0241	0.373065015
<i>Apostasia shenzhenica</i>	<i>Allium cepa</i>	<i>atp4</i>	0.0415	0.0519	1.25060241
<i>Apostasia shenzhenica</i>	<i>Gastrodia elata</i>	<i>atp4</i>	0.1157	0.0592	0.511668107
<i>Apostasia shenzhenica</i>	<i>Hemerocallis citrina</i>	<i>atp4</i>	0.0232	0.0264	1.137931034
<i>Apostasia shenzhenica</i>	<i>Asparagus officinalis</i>	<i>atp4</i>	0.0351	0.023	0.655270655
<i>Apostasia shenzhenica</i>	<i>Allium cepa</i>	<i>atp6</i>	0.0808	0.0415	0.513613861
<i>Apostasia shenzhenica</i>	<i>Gastrodia elata</i>	<i>atp6</i>	0.1093	0.0483	0.441903019
<i>Apostasia shenzhenica</i>	<i>Hemerocallis citrina</i>	<i>atp6</i>	0.0613	0.0279	0.455138662
<i>Apostasia shenzhenica</i>	<i>Asparagus officinalis</i>	<i>atp6</i>	0.0613	0.0356	0.580750408
<i>Apostasia shenzhenica</i>	<i>Allium cepa</i>	<i>atp8</i>	0.5999	0.2566	0.427737956
<i>Apostasia shenzhenica</i>	<i>Gastrodia elata</i>	<i>atp8</i>	0.1729	0.1048	0.606130711
<i>Apostasia shenzhenica</i>	<i>Hemerocallis citrina</i>	<i>atp8</i>	0.2326	0.1286	0.552880482
<i>Apostasia shenzhenica</i>	<i>Asparagus officinalis</i>	<i>atp8</i>	0.314	0.1342	0.427388535
<i>Apostasia shenzhenica</i>	<i>Allium cepa</i>	<i>atp9</i>	0.0634	0.1055	1.664037855
<i>Apostasia shenzhenica</i>	<i>Gastrodia elata</i>	<i>atp9</i>	0.1105	0.0726	0.657013575
<i>Apostasia shenzhenica</i>	<i>Hemerocallis citrina</i>	<i>atp9</i>	0.0866	0.1053	1.215935335
<i>Apostasia shenzhenica</i>	<i>Asparagus officinalis</i>	<i>atp9</i>	0.086	0.0858	0.997674419
<i>Apostasia shenzhenica</i>	<i>Allium cepa</i>	<i>ccmb</i>	0.1074	0.0408	0.379888268
<i>Apostasia shenzhenica</i>	<i>Gastrodia elata</i>	<i>ccmb</i>	0.0754	0.0578	0.766578249
<i>Apostasia shenzhenica</i>	<i>Hemerocallis citrina</i>	<i>ccmb</i>	0.0646	0.0362	0.560371517
<i>Apostasia shenzhenica</i>	<i>Asparagus officinalis</i>	<i>ccmb</i>	0.0504	0.0362	0.718253968
<i>Apostasia shenzhenica</i>	<i>Allium cepa</i>	<i>ccmc</i>	0.0768	0.0447	0.58203125
<i>Apostasia shenzhenica</i>	<i>Gastrodia elata</i>	<i>ccmc</i>	0.0892	0.0368	0.412556054
<i>Apostasia shenzhenica</i>	<i>Hemerocallis citrina</i>	<i>ccmc</i>	0.0641	0.023	0.358814353
<i>Apostasia shenzhenica</i>	<i>Asparagus officinalis</i>	<i>ccmc</i>	0.0704	0.0269	0.382102273
<i>Apostasia shenzhenica</i>	<i>Allium cepa</i>	<i>ccmfc</i>	0.1076	0.0496	0.460966543
<i>Apostasia shenzhenica</i>	<i>Gastrodia elata</i>	<i>ccmfc</i>	0.0813	0.0827	1.017220172
<i>Apostasia shenzhenica</i>	<i>Hemerocallis citrina</i>	<i>ccmfc</i>	0.0641	0.0502	0.783151326
<i>Apostasia shenzhenica</i>	<i>Asparagus officinalis</i>	<i>ccmfc</i>	0.0731	0.0475	0.649794802

<i>Apostasia shenzhenica</i>	<i>Allium cepa</i>	<i>cob</i>	0.051	0.0277	0.543137255
<i>Apostasia shenzhenica</i>	<i>Gastrodia elata</i>	<i>cob</i>	0.0506	0.023	0.454545455
<i>Apostasia shenzhenica</i>	<i>Hemerocallis citrina</i>	<i>cob</i>	0.0512	0.0172	0.3359375
<i>Apostasia shenzhenica</i>	<i>Asparagus officinalis</i>	<i>cob</i>	0.0511	0.0195	0.381604697
<i>Apostasia shenzhenica</i>	<i>Allium cepa</i>	<i>cox1</i>	0.1136	0.0326	0.286971831
<i>Apostasia shenzhenica</i>	<i>Gastrodia elata</i>	<i>cox1</i>	0.0605	0.0189	0.312396694
<i>Apostasia shenzhenica</i>	<i>Hemerocallis citrina</i>	<i>cox1</i>	0.0596	0.0089	0.149328859
<i>Apostasia shenzhenica</i>	<i>Asparagus officinalis</i>	<i>cox1</i>	0.0631	0.026	0.412044374
<i>Apostasia shenzhenica</i>	<i>Allium cepa</i>	<i>cox3</i>	0.1029	0.0205	0.199222546
<i>Apostasia shenzhenica</i>	<i>Gastrodia elata</i>	<i>cox3</i>	0.0632	0.0309	0.488924051
<i>Apostasia shenzhenica</i>	<i>Hemerocallis citrina</i>	<i>cox3</i>	0.0589	0.0184	0.312393888
<i>Apostasia shenzhenica</i>	<i>Asparagus officinalis</i>	<i>cox3</i>	0.0753	0.015	0.199203187
<i>Apostasia shenzhenica</i>	<i>Allium cepa</i>	<i>mttb</i>	0.0373	0.0292	0.782841823
<i>Apostasia shenzhenica</i>	<i>Gastrodia elata</i>	<i>mttb</i>	0.0709	0.0354	0.499294781
<i>Apostasia shenzhenica</i>	<i>Hemerocallis citrina</i>	<i>mttb</i>	0.0278	0.0197	0.708633094
<i>Apostasia shenzhenica</i>	<i>Asparagus officinalis</i>	<i>mttb</i>	0.0342	0.0197	0.576023392
<i>Apostasia shenzhenica</i>	<i>Allium cepa</i>	<i>nad3</i>	0.0594	0.0505	0.85016835
<i>Apostasia shenzhenica</i>	<i>Gastrodia elata</i>	<i>nad3</i>	0.0844	0.0425	0.503554502
<i>Apostasia shenzhenica</i>	<i>Hemerocallis citrina</i>	<i>nad3</i>	0.0597	0.0544	0.911222781
<i>Apostasia shenzhenica</i>	<i>Asparagus officinalis</i>	<i>nad3</i>	0.0471	0.0465	0.987261146
<i>Apostasia shenzhenica</i>	<i>Allium cepa</i>	<i>nad4L</i>	0.165	0.0206	0.12484848
<i>Apostasia shenzhenica</i>	<i>Gastrodia elata</i>	<i>nad4L</i>	0.1366	0.0723	0.292825768
<i>Apostasia shenzhenica</i>	<i>Hemerocallis citrina</i>	<i>nad4L</i>	0.165	0.0206	0.12484848
<i>Apostasia shenzhenica</i>	<i>Asparagus officinalis</i>	<i>nad4L</i>	0.165	0.0206	0.12484848
<i>Apostasia shenzhenica</i>	<i>Allium cepa</i>	<i>nad6</i>	0.0708	0.0587	0.829096045
<i>Apostasia shenzhenica</i>	<i>Gastrodia elata</i>	<i>nad6</i>	0.0531	0.0599	1.128060264
<i>Apostasia shenzhenica</i>	<i>Hemerocallis citrina</i>	<i>nad6</i>	0.1069	0.0564	0.527595884
<i>Apostasia shenzhenica</i>	<i>Asparagus officinalis</i>	<i>nad6</i>	0.0852	0.0564	0.661971831
<i>Apostasia shenzhenica</i>	<i>Allium cepa</i>	<i>nad7</i>	0.0264	0.0107	0.40530303
<i>Apostasia shenzhenica</i>	<i>Gastrodia elata</i>	<i>nad7</i>	0.0389	0.0113	0.290488432
<i>Apostasia shenzhenica</i>	<i>Hemerocallis citrina</i>	<i>nad7</i>	0.0157	0.0096	0.611464968
<i>Apostasia shenzhenica</i>	<i>Asparagus officinalis</i>	<i>nad7</i>	0.0192	0.0084	0.4375
<i>Apostasia shenzhenica</i>	<i>Allium cepa</i>	<i>nad9</i>	0.075	0.0285	0.38
<i>Apostasia shenzhenica</i>	<i>Gastrodia elata</i>	<i>nad9</i>	0.0611	0.0299	0.489361702
<i>Apostasia shenzhenica</i>	<i>Hemerocallis citrina</i>	<i>nad9</i>	0.0704	0.0161	0.228693182

<i>Apostasia shenzhenica</i>	<i>Asparagus officinalis</i>	<i>nad9</i>	0.0706	0.0188	0.266288952
<i>Apostasia shenzhenica</i>	<i>Allium cepa</i>	<i>rps7</i>	1.1904	0.6126	0.514616935
<i>Apostasia shenzhenica</i>	<i>Gastrodia elata</i>	<i>rps7</i>	0.0287	0.0659	2.296167247
<i>Apostasia shenzhenica</i>	<i>Hemerocallis citrina</i>	<i>rps7</i>	0.032	0.0652	2.0375
<i>Apostasia shenzhenica</i>	<i>Asparagus officinalis</i>	<i>rps7</i>	0.032	0.0652	2.0375
<i>Apostasia shenzhenica</i>	<i>Allium cepa</i>	<i>rps12</i>	0.1699	0.0432	0.254267216
<i>Apostasia shenzhenica</i>	<i>Gastrodia elata</i>	<i>rps12</i>	0.0685	0.0321	0.468613139
<i>Apostasia shenzhenica</i>	<i>Hemerocallis citrina</i>	<i>rps12</i>	0.1685	0.04	0.237388724
<i>Apostasia shenzhenica</i>	<i>Asparagus officinalis</i>	<i>rps12</i>	0.1618	0.0804	0.496909765

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Table S3. Prediction of RNA editing sites in *A.shenzhenica* mitogenome

Type	RNA -editing	Number	Percentage
hydrophobic	CTT (L) => TTT (F)	8	28.85%
	CCG (P) => CTG (L)	17	
	CCA (P) => CTA (L)	32	
	CCC (P) => TTC (F)	2	
	CCT (P) => CTT (L)	25	
	GCG (A) => GTG (V)	2	
	CCC (P) => CTC (L)	11	
	CTC (L) => TTC (F)	7	
	CCT (P) => TTT (F)	12	
	GCT (A) => GTT (V)	3	
	GCC (A) => GTC (V)	1	
hydrophilic	CGT (R) => TGT (C)	28	14.42%
	CGC (R) => TGC (C)	9	
	CAT (H) => TAT (Y)	15	
	CAC (H) => TAC (Y)	8	
hydrophobic-hydrophilic	CCC (P) => TCC (S)	13	10.82%
	CCA (P) => TCA (S)	8	
	CCT (P) => TCT (S)	16	
	CCG (P) => TCG (S)	8	
hydrophilic-hydrophobic	TCG (S) => TTG (L)	34	45.43%
	TCA (S) => TTA (L)	52	
	TCT (S) => TTT (F)	38	
	ACT (T) => ATT (I)	3	
	TCC (S) => TTC (F)	29	
	ACA (T) => ATA (I)	3	
	CGG (R) => TGG (W)	27	
	ACG (T) => ATG (M)	2	
	ACC (T) => ATC (I)	1	
hydrophilic-stop	CAA (Q) => TAA (X)	1	0.48%
	CGA (R) => TGA (X)	1	

Table S4 Distribution of penta and hexa SSRs in the *A.shenzhenica* mitogenome.

ID	SSR type	SSR	Start	End	Location
1	penta	(TCCGA)3	20000	20014	IGS(nad4L, nad5)
2	penta	(AAAAG)3	30671	30685	IGS(nad4L, nad5)
3	penta	(TCTTA)3	126755	126769	IGS(nad5, trnC-GCA)
4	penta	(TAAGG)3	191619	191633	IGS(nad9, tRNA-Thr)
5	hexa	(CCCCGG)3	225860	225877	IGS(trnF-GAA, rps7)
6	penta	(TAAAG)3	313314	313328	IGS(PSA, rpl16)
7	hexa	(GGCTTG)3	324866	324883	IGS(rpl2, rps11)
8	penta	(TACAA)3	353917	353931	IGS(rps13, trnK-UUU)
9	penta	(TCATT)3	444355	444369	IGS(rps4, trnQ-UUG)
10	penta	(AAATG)3	543309	543323	IGS(rn18, nad6)
11	penta	(TTTAG)3	579039	579053	IGS(ccmB, nad7)
12	penta	(GAAAA)3	582124	582138	IGS(ccmB, nad7)

Table S5 Distribution of perfect tandem repeats in *A.shenzhenica* mitogenome

No	Repeat sequence	Size	Copy	Percent	start-end	Location
			Number	Matches		
1	TATACATAGAAGCAA	15	2.7	100	173401--173441	IGS(nad9, tRNA-Thr)
2	ATATGTATTAG	11	2.4	100	174049--174074	IGS(nad9, tRNA-Thr)
3	ATAGATATAGAATAAGAATA TATA	24	2	100	174101--174148	IGS(nad9, tRNA-Thr)
4	GAGGAGAAAGGAGGAATGCT CGACCGGAAGGGAACGAGTT ACATACCTGGAACGAGTAGG TT	62	2.3	98	185579--185720	IGS(nad9, tRNA-Thr)
5	ATTTGACTATTTCA	14	1.9	100	203511--203537	IGS(nad9, tRNA-Thr)
6	TGTTACATTTTGT	13	1.9	100	204102--204126	IGS(nad9, tRNA-Thr)
7	ATAGTATATATAAA	14	2.1	100	268146--268174	IGS(matR, tRNA-Cys)
8	TCTTTCTGTCAA	12	2.1	100	305377--305401	IGS(atp4, trnW-CCA)
9	AATAGTTTTAATTATTATGTTG TTTC	26	2	100	418720--418771	IGS(trnM-CAU, rps4)
10	ATAATAATATTATTATTTA	20	2.2	100	431112--431154	IGS(trnM-CAU, rps4)
11	TATAACTTTCTTATCCC	17	2.3	95	558125--558164	IGS(nad6, mttB)
12	CTACTAAAGAAGG	13	2.1	100	561059--561085	IGS(nad6, mttB)
13	TATATTGAATA	11	2.3	100	603499--603523	IGS(trnD-GUC, atp8)

Table S6 The NCBI accession numbers of mitogenomes used in this study

Famliy	Organism	Accession
Asparagus	<i>Allium cepa</i>	AP018390.1
	<i>Hemerocallis citrina</i>	MZ726801-3
	<i>Gastrodia elata</i>	MF070084-102
	<i>Chlorophytum comosum</i>	MW411187.1
	<i>Asparagus officinalis</i>	MT483944.1
Alismatales	<i>Zostera marina</i>	KX808392.1
	<i>Stratiotes aloides</i>	KX808393.1
	<i>Spirodela polyrhiza</i>	NC017840.1
	<i>Butomus umbellatus</i>	KC208619.1
Arecales	<i>Phoenix dactylifera</i>	MH176159.1
	<i>Cocos nucifera</i>	KX028885.1
	<i>Zea perennis</i>	DQ645538.1
	<i>Zea luxurians</i>	DQ645537.1
Poales	<i>Tripsacum dactyloides</i>	NC008362.1
	<i>Zea mays</i>	DQ645536.1
	<i>Chrysopogon zizanioides</i>	MN635785.1
	<i>Coix lacryma-jobi var. puellarum</i>	MT471098.1
	<i>Sorghum bicolor</i>	DQ984518.1
	<i>Sporobolus alterniflorus</i>	MT471321.1
	<i>Eleusine indica</i>	MF616338.1
	<i>Oryza sativa</i>	JN861112.1
	<i>Oryza rufipogon</i>	AP011076.1
	<i>Oryza coarctata</i>	MG429050.1
	<i>Aegilops speltoides</i>	AP013107.1
	<i>Triticum aestivum</i>	GU985444.1
	<i>Hordeum vulgare</i>	MN127974.1
	<i>Lolium perenne</i>	JX999996.1
	<i>Bambusa oldhamii</i>	EU365401.1
Outgroup	<i>Medinilla magnifica</i>	MT043351.1
	<i>Eucalyptus grandis</i>	NC040010.1
	<i>Pyrus betulifolia</i>	MW080658.1
	<i>Nelumbo nucifera</i>	NC030753.1