

Table S2. The size variation between *V. sinensis* CN/TL (CN/TL), *V. sinensis* QY (QY) and *V. sinensis* (VS) (HM067837).

tRNA/IGN	CN/TL	QY	VS (HM067837)	tRNA/IGN	CN/TL	QY	VS (HM067837)
tRNA ^{Ile}	64	65	64	tRNA ^{Asn}	64	64	64
IGN	(+11	(+14	(+7	IGN	(-3	(-3	(-4
tRNA ^{Gln}	69	69	69	tRNA ^{Ser1}	66	66	68
IGN	(+18	(+16	(+19	IGN	0	0	(-1
tRNA ^{Met}	64	64	64	tRNA ^{Glu}	63	63	63
IGN	0	0	0	IGN	(-2	(+25	(-2
ND2	1023	1023	1023	tRNA ^{Phe}	63	64	63
IGN	(-2	(-2	(-2	IGN	0	0	0
tRNA ^{Trp}	69	69	69	ND5	1729	1729	1729
IGN	(-8	(-8	(-8	IGN	0	0	0
tRNA ^{Cys}	61	61	61	tRNA ^{His}	62	62	63
IGN	0	0	0	IGN	0	0	0
tRNA ^{Tyr}	70	70	70	ND4	1347	1347	1347
IGN	(-41	(-41	(-41	IGN	(-7	(-7	(-7
COI	1578	1578	1578	ND4L	297	297	294
IGN	(-5	(-5	(-5	IGN	(+2	(+2	(+2
tRNA ^{Leu2}	64	64	64	tRNA ^{Thr}	62	62	62
IGN	0	0	0	IGN	0	0	0
COII	688	688	688	tRNA ^{Pro}	64	64	64
IGN	0	0	0	IGN	(+5	(+5	(+5
tRNA ^{Lys}	69	69	70	ND6	516	516	516
IGN	0	0	(-1	IGN	(-1	(-1	(-1
tRNA ^{Asp}	67	67	67	Cyt b	1135	1135	1135
IGN	(+9	(+9	(+9	IGN	0	0	0
ATP8	156	156	156	tRNA ^{Ser2}	67	67	67
IGN	(-4	(-4	(-4	IGN	(+18	(+30	(-20
ATP6	675	675	675	ND1	939	939	939
IGN	(-1	(-1	(-1	IGN	(+1	(+1	(+1
COIII	789	789	789	tRNA ^{Leu1}	64	65	64
IGN	(+4	(+4	(+4	IGN	0	0	0
tRNA ^{Gly}	62	62	62	16S rRNA	1223	1221	1251
IGN	(-3	(-3	(-3	IGN	0	0	(-28
ND3	357	357	357	tRNA ^{Val}	66	66	66
IGN	(-2	(-2	(-2	IGN	0	0	0
tRNA ^{Ala}	65	66	67	12S rRNA	792	790	793
IGN	0	0	(-1	IGN	0	0	0
tRNA ^{Arg}	64	64	63	CR	1015	911	1105
IGN	(-3	(-3	(-3				