

Table S4. Tandem repeats in the *D. tangerinum* mitogenome

Indices	Period Size	Copy Number	Consensus Size	Percent Matches	Percent Indels	Score	A	C	G	T	Entropy (0-2)
3650--3680	15	2.0	16	93	6	55	64	0	19	16	1.29
27206--27238	17	1.9	17	100	0	66	12	12	6	69	1.35
39364--39401	14	2.5	16	91	8	62	36	13	5	44	1.66
78010--78042	17	1.9	17	100	0	66	69	6	12	12	1.35
136471--136509	21	1.9	19	90	10	60	53	7	25	12	1.65
192471--192577	45	2.4	45	98	0	205	10	23	19	46	1.80
193993--194032	20	2.0	20	90	0	62	65	12	0	22	1.26
229852--229879	13	2.2	13	100	0	56	46	0	21	32	1.52
235017--235058	17	2.5	17	100	0	84	40	0	23	35	1.55
327784--327816	17	1.9	17	100	0	66	12	12	6	69	1.35
343947--343972	10	2.6	10	100	0	52	61	0	7	30	1.24
376187--376219	17	1.9	17	100	0	66	12	12	6	69	1.35
405417--405449	17	1.9	17	100	0	66	12	12	6	69	1.35
459258--459284	14	1.9	14	100	0	54	48	14	7	29	1.71
512110--512146	15	2.4	16	90	4	58	35	5	8	51	1.55
512794--512832	17	2.3	17	100	0	78	25	10	0	64	1.25
512872--512920	17	2.9	17	90	3	73	44	4	2	48	1.33

Parameters: Total length (Period size * Copy number) >= 30 and Percent Matches >= 90%