

Table S5. Location of mt-like nuclear tRNA genes in *G. raimondii* nuclear genome

tRNA	Chr	Identity	Length (bp)	Start	End
trnW-CCA	1	100	74	23,186,930	23,186,857
trnS-GGA	1	100	87	23,199,416	23,199,330
trnS-GCT	1	100	88	23,231,316	23,231,229
trnF-GAA	1	100	74	23,231,737	23,231,664
trnP-UGG	1	98.68	76	23,232,031	23,231,957
trnM-CAU-1	1	100	81	23,342,432	23,342,512
trnM-CAU-cp	1	100	73	23,419,701	23,419,629
trnM-CAU-2	1	100	73	23,450,444	23,450,372
trnW-CCA	1	98.65	74	23,474,675	23,474,748
trnV-GAC	1	98.61	72	23,483,961	23,483,890
trnY-GUA	1	100	83	23,549,199	23,549,117
trnN-GUU	1	100	72	23,550,166	23,550,095
trnC-GCA	1	100	71	23,552,368	23,552,298
trnE-UUC	1	100	72	23,580,669	23,580,598
trnH-GUG	1	100	74	23,582,754	23,582,681
trnS-UGA	1	100	87	23,616,081	23,615,995
trnK-UUU	1	100	73	23,616,989	23,616,917
trnG-GCC	1	100	72	23,645,773	23,645,702
trnM-CAU-2	1	98.63	73	23,651,459	23,651,387
trnI-UAU	1	100	86	23,661,010	23,660,925
trnD-GUC	1	100	74	23,676,388	23,676,315
trnQ-UUG	1	100	72	23,818,761	23,818,690
trnH-GUG	1	93.06	72	42,544,620	42,544,549
trnH-GUG	2	92	75	26,982,032	26,981,959
trnM-CAU-2	2	95.89	73	52,875,794	52,875,722
trnM-CAU-2	3	90.41	73	13,883,827	13,883,756
trnD-GUC	4	98.61	72	3,555,795	3,555,866
trnS-GGA	4	92.86	84	32,001,956	32,001,873
trnS-GGA	5	95.35	86	7,648,967	7,649,052
trnW-CCA	5	94.59	74	49,260,616	49,260,689
trnM-CAU-2	5	94.52	73	60,057,635	60,057,707
trnD-GUC	7	93.24	74	13,216,197	13,216,124
trnK-UUU	7	98.63	73	22,092,280	22,092,208
trnS-GGA	7	91.76	85	24,488,177	24,488,093
trnM-CAU-cp	7	98.59	71	44,344,852	44,344,922
trnS-GGA	8	96.55	87	26,955,720	26,955,634
trnM-CAU-2	9	95.89	73	22,125,081	22,125,153
trnM-CAU-2	10	91.43	70	28,957,917	28,957,848
trnS-GGA	11	94.32	88	24,281,095	24,281,009
trnS-GGA	13	89.66	87	16,190,205	16,190,119
trnQ-UUG	13	94.2	69	23,940,068	23,940,000
trnW-CCA	13	100	74	36,753,933	36,753,860
trnM-CAU-1	13	97.53	81	52,706,781	52,706,701
trnS-GGA	13	97.7	87	52,748,418	52,748,504