

Supplementary Table 9. Summary of predicted ncRNA genes in the watercress genome

Type		Number	Average length (bp)	Total length (bp)	Percentage of Genome (%)
rRNA	rRNA	2,043	348.63	712,255	0.21
	18S	67	1,923.66	128,885	0.04
	28S	80	4,534.89	362,791	0.11
	5.8S	72	156.1	11,239	0.00
	5S	1,824	114.77	209,340	0.06
snRNA	snRNA	558	108.58	60,586	0.02
	CD-box	396	97.39	38,565	0.01
	HACA-box	69	129.28	8,920	0.00
	splicing	93	140.87	13,101	0.00
miRNA		320	123.75	39,600	0.01
tRNA		884	75.28	66,551	0.02