

**Supplementary Table 13. Statistics on the annotation of non-coding RNA of the *C. goeringii* genome.**

Type	Copy	Average length (bp)	Total length (bp)	% of genome	
miRNA	147	120.8503	17,765	0.000437	
tRNA	493	73.8438	36,405	0.000896	
rRNA	rRNA	1,544	286.0699	441,692	0.010872
	18S	371	773.5445	286,985	0.007064
	28S	264	178.9091	47,232	0.001163
	5.8S	134	145.9179	19,553	0.000481
	5S	775	113.4477	87,922	0.002164
	snRNA	528	118.375	62,502	0.001538
snRNA	CD-box	307	101.4788	31,154	0.000767
	HACA-box	75	153.6133	11,521	0.000284
	splicing	146	135.8014	19,827	0.000488