

**S11 Table. Statistics of the noncoding RNA in the Yap hadal snailfish genome.**

Type	Number	Average length (bp)	Total length (bp)	% of genome
miRNA	1,651	107.67	177,759	0.022429
tRNA	9,143	82.48	754,147	0.10306
rRNA	3,572	118.95	424,874	0.058083
	18S	2,707	253,565	0.034652
	28S	340	118,145	0.016146
	5.8S	42	6,309	0.000862
	5S	483	46,855	0.006403
snRNA	1,323	136.87	181,085	0.024747
	CD-box	262	27,718	0.003788
	HACA-box	152	23,938	0.003271
	splicing	869	123,127	0.016826
Total	-	-	2,137,522	0.292111