

Table S10. RNA alignment Rfam results statistics

RNA_Types	Uniq_tags	Uniq_ratio	Total_tags	Total_ratio
rRNA	7,869	84.09%	250,865	91.46%
tRNA	64	0.68%	198	0.07%
snoRNA	857	9.16%	20,185	7.36%
snRNA	0	0.00%	0	0.00%
Other	568	6.07%	3,033	1.11%
Total	9,358	100.00%	274,281	100.00%