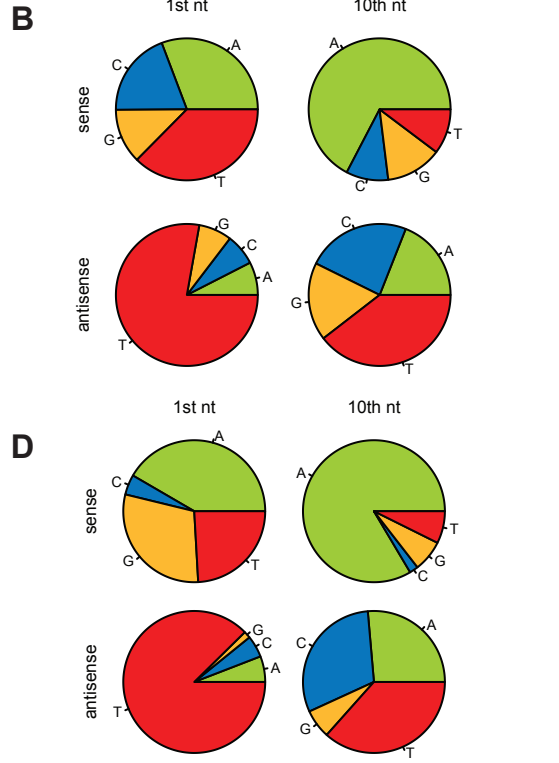
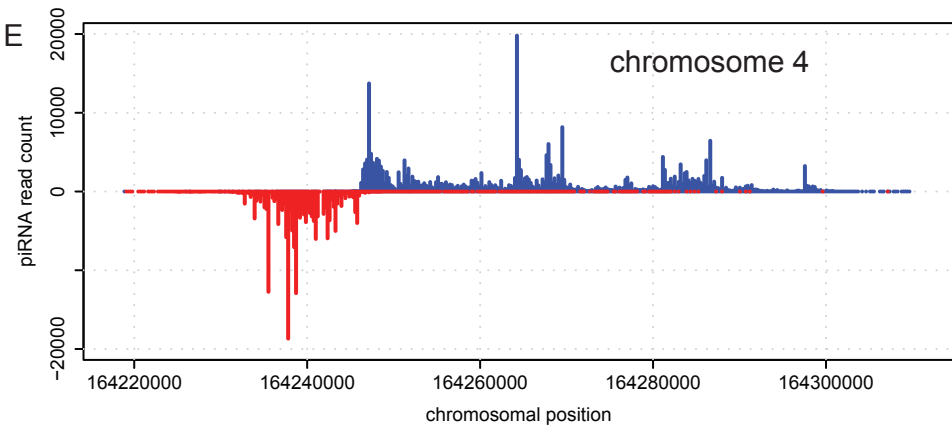
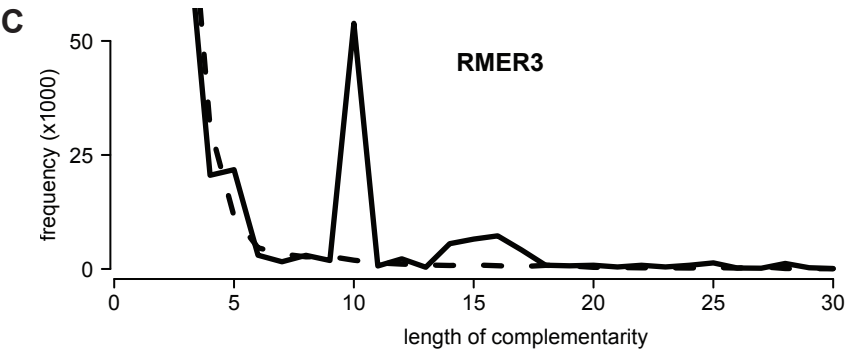
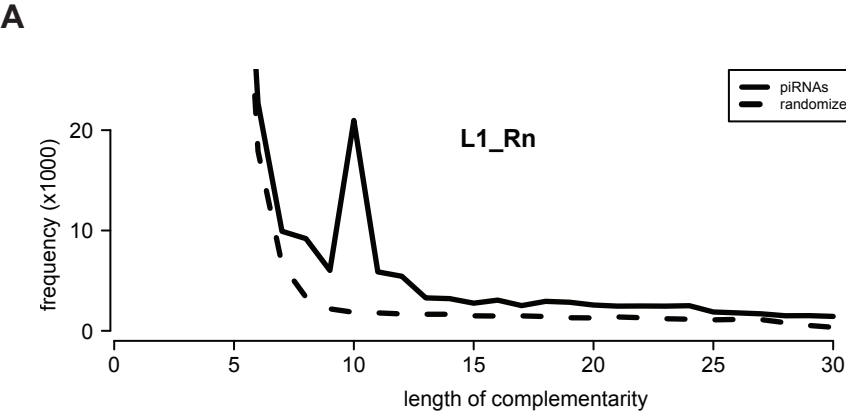




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characteristic	piRNA clusters	outside clusters
Gene count (genes/Mb)	19	4.8
Intergenic regions > 100 kb	63%	29%
Repeat sequence content	30.5%	38.4%
CG content	46.2%	41.9%
sequence conserved > 50%	3.90%	5.37%