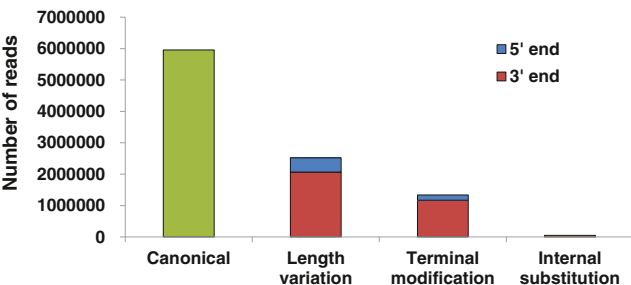
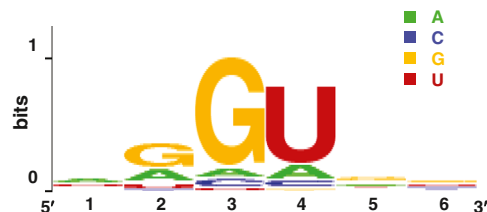


A**C****B**

miR-124	Sequence variants	Reads	Percent	
Length variations	UACAAUUAAGGCACGCGGUGAAUGCCAAGAGCG			
	UUAAGGCACGCGGUGAAUGC	2075	1.97%	
	UUAAGGCACGCGGUGAAUGCC	5393	5.13%	
	UUAAGGCACGCGGUGAAUGCCA	8328	7.92%	
	UAAGGCACGCGGUGAAUGC	6335	6.03%	
	UAAGGCACGCGGUGAAUGCC	14215	13.53%	
	UAAGGCACGCGGUGAAUGCCA	50082	47.66%	
	UAAGGCACGCGGUGAAUGCCAA	8418	8.01%	
	UAAGGCACGCGGUGAAUGCCAAG	278	0.26%	
	AAGGCACGCGGUGAAUGCC	431	0.41%	
	AAGGCACGCGGUGAAUGCCA	1727	1.64%	
	AGGCACGCGGUGAAUGCCA	88	0.08%	
Terminal modification (5')	CAAGGCACGCGGUGAAUGCCA	236	0.22%	
Terminal modification (5')	AAAGGCACGCGGUGAAUGCCA	191	0.18%	
Terminal modification (3')	UAAGGCACGCGGUGAAUGCCA U	4928	4.69%	
Terminal modification (3')	UAAGGCACGCGGUGAAUGCCA A	2251	2.14%	

0% 20% 40% 60%