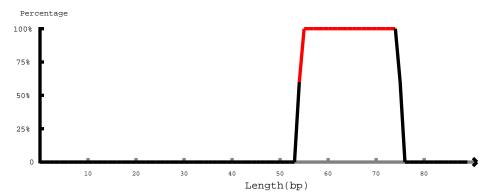
miRge 3.0 Novel miRNA

Name: Sample name: Probability: Chr(GRCh38): Start position of mature miRNA: End position of mature miRNA: Strand: Total read count: Mature miRNA read count:	31128309 + 10 10
Passenger miRNA read count:	0



5'-	CAGCCGGGAUAACUUCCUCAGGAGUUUGGGCUAAAGCUCUCAAAGGGCAAGGGC <mark>CUUAACCUGAAUUUCUGAGC</mark> UAUAUCUCUGAAUUU	-3'
	$\dots (((((((((((((((((((((((((((((((((((($	reads
	CCUUAACCUGAAUUUCUGAGC	4
	CCUUAACCUGAAUUUCUGAGCU	2
	CUUAACCUGAAUUUCUGAGCU	4