

Seq1	Seq2	Tot Score	Tot Energy	Seq1 Position	Seq2 Position	Length
hsa-miR-181a	GAS5-AS1	83	-22.52	3 22	589 607	19

Query: 3' UGAGUGGCUGUCGCAACUUACAA 5'

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Ref: 5' CGTC-CCGCCGGGTGGGTGGG 3'

Seq1	Seq2	Tot Score	Tot Energy	Seq1 Position	Seq2 Position	Length
hsa-miR-181a	LINC01278	81	-20.12	1 24	872 897	25

Query: 3' UGAGUGGCUGUCGCAA--CUUACAA 5'

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Ref: 5' GCGAGCCGGGCGCGTTCAGGGTGGT 3'

Seq1	Seq2	Tot Score	Tot Energy	Seq1 Position	Seq2 Position	Length
hsa-miR-181a	MIAT	73	-20.45	4 23	135 153	19

Query: 3' UGAGUGGCUGUCGCAACUUACAA 5'

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Ref: 5' TGACACCGTCAGC-ATGGCTGTA 3'