

**Predicted results using PITA (target score≤-20)**

Gene	microRNA	Sites	Score
Seq1	hsa-miR-671-5p	4	-31
Seq1	hsa-miR-608	14	-30.31
Seq1	hsa-miR-330-5p	6	-30.3
Seq1	hsa-miR-661	8	-27.41
Seq1	hsa-miR-324-5p	4	-27.2
Seq1	hsa-miR-326	6	-26.8
Seq1	hsa-miR-296-3p	11	-26.63
Seq1	hsa-miR-1182	12	-26.45
Seq1	hsa-miR-502-3p	8	-24.8
Seq1	hsa-miR-657	5	-24.62
Seq1	hsa-miR-939	7	-24.62
Seq1	hsa-miR-1254	4	-24.6
Seq1	hsa-miR-638	3	-24.1
Seq1	hsa-miR-1237	8	-23.61
Seq1	hsa-miR-367	2	-23.3
Seq1	hsa-miR-612	5	-23.2
Seq1	hsa-miR-1226	4	-22.7
Seq1	hsa-miR-1268	7	-22.66
Seq1	hsa-miR-760	11	-22.64
Seq1	hsa-miR-766	7	-22.23
Seq1	hsa-miR-373	9	-21.99
Seq1	hsa-miR-520e	9	-21.8
Seq1	hsa-miR-198	8	-21.6
Seq1	hsa-miR-323-5p	5	-21.5
Seq1	hsa-miR-188-5p	5	-21.47
Seq1	hsa-miR-423-5p	9	-21.45
Seq1	hsa-miR-422a	4	-21.33
Seq1	hsa-miR-1231	13	-21.22
Seq1	hsa-miR-26a	4	-21.2
Seq1	hsa-miR-139-5p	5	-21.15
Seq1	hsa-miR-342-5p	10	-21.1
Seq1	hsa-miR-601	6	-20.92
Seq1	hsa-miR-1234	5	-20.9
Seq1	hsa-miR-675	9	-20.84
Seq1	hsa-miR-302c	9	-20.81
Seq1	hsa-miR-361-3p	5	-20.8
Seq1	hsa-miR-582-5p	6	-20.8
Seq1	hsa-miR-510	3	-20.7
Seq1	hsa-miR-520f	10	-20.64
Seq1	hsa-miR-520d-3p	9	-20.56
Seq1	hsa-miR-127-5p	3	-20.5
Seq1	hsa-miR-1266	13	-20.45
Seq1	hsa-miR-544	10	-20.2
Seq1	hsa-miR-1294	11	-20.17
Seq1	hsa-miR-658	13	-20.17

Seq1	hsa-miR-135b	7	-20.14
Seq1	hsa-miR-145	13	-20.11
Seq1	hsa-miR-1260	4	-20.05
Seq1	hsa-miR-663	5	-20.05
Seq1	hsa-miR-765	15	-20.03
Seq1	hsa-miR-1308	10	-20
Seq1	hsa-miR-302e	9	-19.9
Seq1	hsa-miR-92a	2	-19.9
Seq1	hsa-miR-1293	9	-19.78
Seq1	hsa-miR-942	9	-19.74
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Seq1	hsa-miR-223	7	-19.59
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Seq1	hsa-miR-1274a	5	-19.38
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Seq1	hsa-miR-520a-5p	8	-18.4

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Seq1	hsa-miR-129-3p	4	-12
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Seq1	hsa-miR-593	9	-10.89
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Seq1	hsa-miR-520c-5p	4	-9.03
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Seq1	hsa-miR-595	3	-8.75
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