

Supplementary materials

Table S1 The primer design for miRNA RT-PCR

prostate related miRNAs	miRNA sequence	primer	reference
Let-7c-5p	TGAGGTAGTAGGTTGTATGGTT	TGAGGTAGTAGGTTGTATGGTT	1
Let-7e-5p	TGAGGTAGGAGGTTGTATAGTT	TGAGGTAGGAGGTTGTATAGTT	1
miR-100-5p	AACCCGTAGATCCGAACCTTGTG	ACCCGTAGATCCGAACCTTGTG	1, 2
miR-101-3p	TACAGTACTGTGATAACTGAA	CGGTACAGTACTGTGATAACTGAA	3
miR-103-3p	AGCAGCATTGTACAGGGCTATGA	GCAGCATTGTACAGGGCTATG	4
miR-106a-5p	CAAAGTGCTAACAGTGCAGGTAG	AAGTGCTAACAGTGCAGGTAG	1,3
miR-106b-5p	TAAAGTGCTGACAGTGCAGAT	TAAAGTGCTGACAGTGCAGAT	3
miR-107-3p	AGCAGCATTGTACAGGGCTATCA	GCAGCATTGTACAGGGCTATC	1, 3, 4
miR-10a-5p	TACCCTGTAGATCCGAATTTGTG	TACCCTGTAGATCCGAATTTGTG	3, 5
miR-125a-5p	TCCCTGAGACCCTTTAACCTGTGA	CCTGAGACCCTTTAACCTGTG	3
miR-125b-5p	TCCCTGAGACCCTAACCTGTGA	TCCCTGAGACCCTAACCTGTG	2, 3
miR-126-3p	TCGTACCGTGAGTAATAATGCG	TCGTACCGTGAGTAATAATGCG	3
miR-126-5p(*)	CATTATTACTTTTGGTACGCG	GGCATTATTACTTTTGGTACGCG	3
miR-128-3p	TCACAGTGAACCGGTCTCTTT	TCACAGTGAACCGGTCTCTT	3
miR-141-3p	TAACACTGTCTGGTAAAGATGG	CGTAACACTGTCTGGTAAAGATGG	1, 2, 3
miR-143-3p	TGAGATGAAGCACTGTAGCTC	TGAGATGAAGCACTGTAGCTC	2, 3
miR-145a-5p	GTCCAGTTTTCCAGGAATCCCT	CCAGTTTTCCAGGAATCCC	1, 3
miR-146a-5p	TGAGAACTGAATTCCATGGGTT	TGAGAACTGAATTCCATGGGTT	3
miR-148a-3p	TCAGTGCACTACAGAACTTTGT	TCAGTGCACTACAGAACTTTGT	3
miR-151-3p	CTAGACTGAGGCTCCTTGAGG	TAGACTGAGGCTCCTTGAGG	6
miR-155-5p	TTAATGCTAATTGTGATAGGGGT	GTTAATGCTAATTGTGATAGGGGT	7
miR-16-5p	TAGCAGCACGTAAATATTGGCG	TAGCAGCACGTAAATATTGGCG	3, 4
miR-181a-5p	AACATTCAACGCTGTCCGGTGAGT	CATTCAACGCTGTCCGGTGAG	3
miR-182-5p	TTTGGCAATGGTAGAACTCACACCG	TGGCAATGGTAGAACTCACAC	3
miR-194-5p	TGTAACAGCAACTCCATGTGGA	TGTAACAGCAACTCCATGTGG	3
miR-195a-5p	TAGCAGCACAGAAATATTGGC	TAGCAGCACAGAAATATTGGC	3
miR-200a-3p	TAACACTGTCTGGTAACGATGT	TAACACTGTCTGGTAACGATGT	3
miR-200b-3p	TAATACTGCCTGGTAATGATGA	CGTAATACTGCCTGGTAATGATGA	3
miR-200c-3p	TAATACTGCCGGGTAATGATGGA	TAATACTGCCGGGTAATGATGG	3
miR-203-3p	GTGAAATGTTTAGGACCACTAG	CGGTGAAATGTTTAGGACCACTAG	3
miR-205-5p	TCCTTCATTCCACCGGAGTCTG	TCCTTCATTCCACCGGAGTC	2, 3
miR-20a-5p	TAAAGTGCTTATAGTGCAGGTAG	TAAAGTGCTTATAGTGCAGGTAG	1, 3, 8
miR-21a-5p	TAGCTTATCAGACTGATGTTGA	CGGTAGCTTATCAGACTGATGTTGA	1, 3
miR-210-3p	CTGTGCGTGTGACAGCGGCTGA	TGCGTGTGACAGCGGCTGA	3
miR-214-3p	ACAGCAGGCACAGACAGGCAGT	AGCAGGCACAGACAGGCAG	3
miR-218-5p	TTGTGCTTGATCTAACCATGT	CGTTGTGCTTGATCTAACCATGT	3

miR-22-3p	AAGCTGCCAGTTGAAGAAGCTGT	AGCTGCCAGTTGAAGAAGCTG	3
miR-221-3p	AGCTACATTGTCTGCTGGGTTTC	CTACATTGTCTGCTGGGTTTC	1, 3
miR-222-3p	AGCTACATCTGGCTACTGGGT	AGCTACATCTGGCTACTGGG	3
miR-223-3p	TGTCAGTTTGTCAAATACCCCA	TGTCAGTTTGTCAAATACCCCA	1, 3
miR-224-5p	TAAGTCACTAGTGGTTCCGTT	TAAGTCACTAGTGGTTCCGTT	3
miR-23a-3p	ATCACATTGCCAGGGATTTC	ATCACATTGCCAGGGATTTC	3
miR-23b-3p	ATCACATTGCCAGGGATTACC	ATCACATTGCCAGGGATTACC	3
miR-24-3p	TGGCTCAGTTCAGCAGGAACAG	GCTCAGTTCAGCAGGAACAG	1, 3
miR-25-3p	CATTGCACTTGTCTCGGTCTGA	CATTGCACTTGTCTCGGTCTG	3
miR-26a-5p	TTCAAGTAATCCAGGATAGGCT	TTCAAGTAATCCAGGATAGGCT	1, 3
miR-26b-5p	TTCAAGTAATTCAGGATAGGT	CGGTTCAAGTAATTCAGGATAGGT	1
miR-27b-3p	TTCACAGTGGCTAAGTTCTGC	TTCACAGTGGCTAAGTTCTGC	3
miR-296-5p	AGGGCCCCCCTCAATCCTGT	GCCCCCCTCAATCCTG	2, 3
miR-298-5p	GGCAGAGGAGGGCTGTTCTTCC	CAGAGGAGGGCTGTTCTTC	1
miR-29a-3p	TAGCACCATCTGAAATCGGTTA	TAGCACCATCTGAAATCGGTTA	3
miR-301a-3p	CAGTGCAATAGTATTGTCAAAGC	CAGTGCAATAGTATTGTCAAAGC	3
miR-30b-5p	TGTAAACATCCTACACTCAGCT	TGTAAACATCCTACACTCAGCT	3
miR-30c-5p	TGTAAACATCCTACACTCTCAGC	TGTAAACATCCTACACTCTCAGC	1, 3
miR-31-5p	AGGCAAGATGCTGGCATAGCTG	GCAAGATGCTGGCATAGCTG	3
miR-320-3p	AAAAGCTGGGTTGAGAGGGCGA	AAAAGCTGGGTTGAGAGGGC	3
miR-328-3p	CTGGCCCTCTCTGCCCTTCCGT	CCCTCTCTGCCCTTCCG	4
miR-330-5p	TCTCTGGGCCTGTGTCTTAGGC	TCTCTGGGCCTGTGTCTTAG	3
miR-331-3p	GCCCCTGGGCCTATCCTAGAA	CCCTGGGCCTATCCTAGA	3
miR-345-5p	GCTGACCCCTAGTCCAGTGCTT	CTGACCCCTAGTCCAGTGC	3
miR-346-5p	TGTCTGCCCGAGTGCCTGCCTCT	TGCCCGAGTGCCTGCCTCT	1
miR-34a-5p	TGGCAGTGTCTTAGCTGGTTGT	GGCAGTGTCTTAGCTGGTTG	3, 9
miR-34b-5p	AGGCAGTGTAAATTAGCTGATTGT	AGGCAGTGTAAATTAGCTGATTGT	3, 4
miR-34c-5p	AGGCAGTGTAGTTAGCTGATTGC	GGCAGTGTAGTTAGCTGATTGC	3, 9
miR-375-3p	TTGTTCGTTCCGGCTCGCGTGA	TTGTTCGTTCCGGCTCGCGT	1, 3
miR-449a-5p	TGGCAGTGTATTGTTAGCTGGT	TGGCAGTGTATTGTTAGCTGG	3, 9
miR-451-5p	AAACCGTTACCATTACTGAGTT	CGGAAACCGTTACCATTACTGAGTT	1
miR-485-3p	AGTCATACACGGCTCTCCTCTC	AGTCATACACGGCTCTCCTC	4
miR-486-5p	TCCTGTACTGAGCTGCCCCGAG	CTGTACTGAGCTGCCCCGA	4
miR-574-3p	CACGCTCATGCACACACCCACA	GCTCATGCACACACCCACA	1, 4
miR-7a-5p	TGGAAGACTAGTGATTTTGTGT	CGTGGAAGACTAGTGATTTTGTGT	3
miR-7a-2-3p	CAACAAGTCCCAGTCTGCCACA	ACAAGTCCCAGTCTGCCACA	7
miR-92a-3p	TATTGCACTTGTCCCAGCCTG	TATTGCACTTGTCCCAGCCTG	3, 4
miR-92b-3p	TATTGCACTCGTCCCAGCCTC	TATTGCACTCGTCCCAGCCTC	4
miR-93-5p	CAAAGTGCTGTTTCGTGCAGGTAG	AAGTGCTGTTTCGTGCAGGTAG	1
miR-96-5p	TTTGGCACTAGCACATTTTGTCT	TTTGGCACTAGCACATTTTGTCT	3
miR-99a-5p	AACCCGTAGATCCGATCTTGTG	AACCCGTAGATCCGATCTTGTG	3
U6		CGCAAATTCGTGAAGCGTTCC	

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Table S2 Primer design for Drosha and Dicer1 RT-PCR

primer name	sequence
Q-mDrosha-S	GAGCCTAGAGGAAGCCAAACA
Q-mDrosha-AS	GCCGGACGTGAGTGAAGAT
Q-dicer1-S3	GCAAGGAATGGACTCTGAGC

Q-dicer1-AS3	GGGGACTTCGATATCCTCTTC
Q-mActb-S	CTTTCCAGCCTTCCTTCTTG
Q-mActb-AS	CAGCACTGTGTTGGCATAGAG
