

Supporting information

Highly sensitive and multiplexed miRNAs analysis based on digitally encoded silica microparticles coupled with RCA-based cascade amplification

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Table S1. Sequences of oligonucleotides and miRNAs

Name	Sequence (5' to 3')
Padlock probe- Let-7a	p- CTACT ACCTC A GGAG ATGAG TGACT GACGA GCTGA GGCCT ATGTG CATTC GCTGA GG A AC TAT AC AAC
Capture probe- Let-7a	NH ₂ -AAAAAAAAAAAAAGGCCTATGTG <u>CATTCGCTGA</u> TGCGACCT
Padlock probe- MiRNA-21	p- CTG ATAAGCTA GGAGATGAGT GACTGACGA G CTGAGG G CAT TCTCCATGT G CTGAGG TCAA C ATCAGT
Capture probe- miR-21	NH ₂ -AAAAAAAAAAAAA <u>GGGCATTCTC</u> <u>CATGTGCTGATGCGACCT</u>
RCA primer UT	TCG TCA GTCACT CA TCTCC Cy3-AGGTCGCA
Designed sequence c for let-7a	TCAGC GAATG CACAT AGGCC
Let-7a	U GAGGU AGUAG GUUGUAUAGUU
Let-7b	UGAGGUAGUAGGUUGUGUGUU
Let-7c	UGAGGUAGUAGGUUGUAUGUU
MiRNA-21	UAGCUUAUCAGACUGAUGUUGA'

Table S2. The Comparison of Existing Amplification Methods for miRNA analysis

Detection method	LOD	Dynamic range	References
RCA	10 pM	-	1
Branched RCA	18 fM	3	2
Dumbbell probe-mediated RCA (D-RCA)	1 fM	8	3
RCA-assisted NESA	100 fM	4	4
P-ERCA	0.2 aM	3	5
Loop mediated isothermal amplification (LAMP)	1 pM	7	6
Exponential amplification reaction (EXPAR)	0.1 aM	10	7
Duplex specific nuclease signal amplification	100 fM	5	8
RCA based cascade amplification	0.5 fM	5	This work

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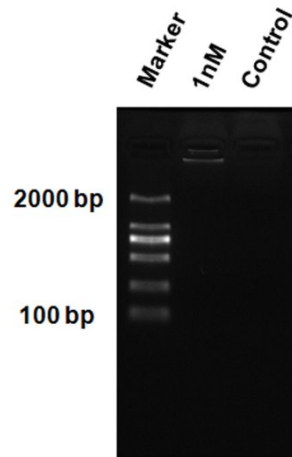
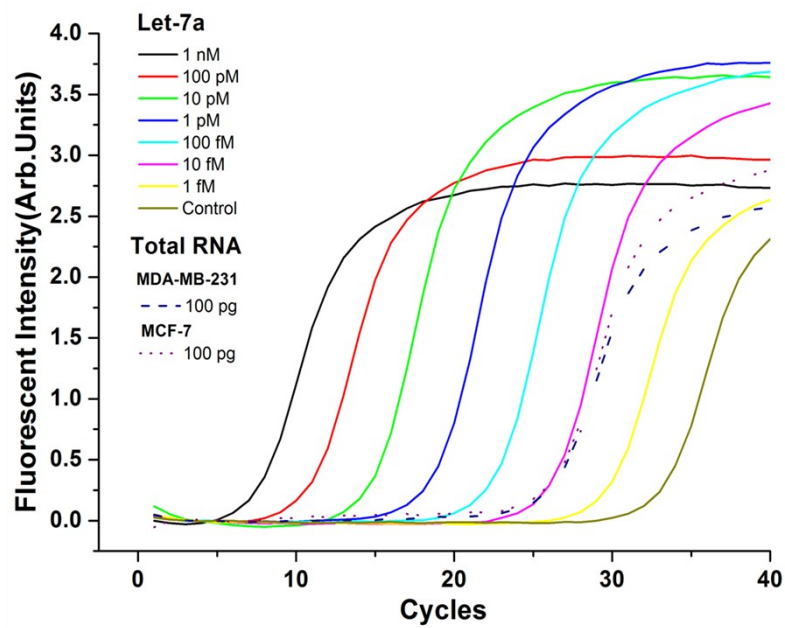


Fig. S1 Agarose gel electrophoresis image of the RCA product, with let-7a of 1 nM and DNA marker of 100–2000 bp.

Fig. S2

A



B

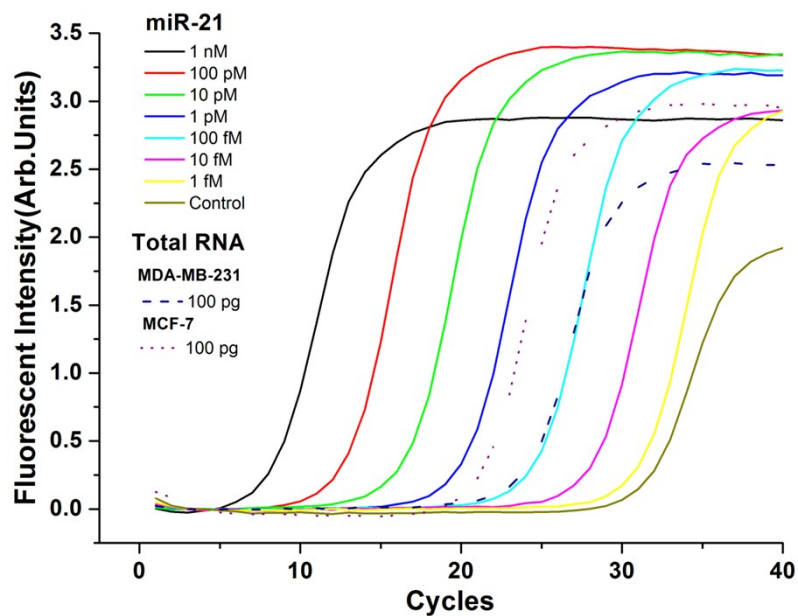


Fig. S2 (A) qRT-PCR analysis of let-7a with synthetic standards of concentrations ranging from 1 fM to 1 nM, and 100 pg total RNA from MDA-MB-231 and MCF-7 cell lines. (B) qRT-PCR analysis of miR-21 with synthetic standards of concentrations ranging from 1 fM to 1 nM, and 100 pg total RNA from MDA-MB-231 and MCF-7 cell lines.