An all-in-one homogeneous DNA walking nanomachine and its application for intracellular analysis of miRNA

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Figure S1 FRET induced by the hybridization of HP-1 (green) and HP-2 (red). (A) Fluorescence excitation and emission spectra of FITC. (B) Fluorescence excitation and emission spectra of TAMRA. (C) The occurrence of FRET of HP-1/HP-2 after annealing.



Figure S2 Standard linear calibration curves of (A) FITC-labeled HP-1 (Ex: 494 nm; Em: 517 nm) and (B) TAMRA-labeled HP-2 (Ex: 558 nm; Em: 583 nm). The intersection points of the dashed lines show the relative fluorescence intensities of HP-1 and HP-2 released from the AuNPs using DTT (10 mM) treatment.



Figure S3 Fluorescence intensity ratio (F_T/F_F) of the nanomachine in the presence of PBS, DMEM with 10% FBS, RPMI-1640 with 10% FBS, DNase I, cell lysate and miR-21.



Figure S4 (A) Scheme showing the nanomachine treated by target miR-21 or nontarget miRNAs (miR-16, miR-26a, and miR-214). (B) Fluorescence intensity ratio (F_T/F_F) of the nanomachine in the presence of PBS, miR-16, miR-26a, miR-214, and miR-21.



Figure S5 Cell viability determined by MTT cytotoxicity assays. MCF-7 cells were incubated with nanomachine (6 nM) for 0, 3, 6, 12, 18 and 24 h.



Figure S6 Confocal images of MCF-7 cells incubated with the nanomachine (6 nM) at different time points. Scale bar: $20 \ \mu m$.



Figure S7 Confocal images of MCF-7 cells treated with LysoTracker blue and nanomachine for 1 h and 3 h, respectively. Scale bar: 20 µm.



Figure S8 The relative expression levels of miR-21 in MCF-7, HeLa and L02 by qRT-PCR.



Figure S9 Confocal images of miR-21 in HeLa cells using the DNA walking nanomachine after transfecting the cells with NC (negative control), miR-21 mimic or inhibitor. Scale bar: 20 µm.

Table S1. Sequences	of oligonucleotides.
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Name	Sequence (5' to 3')	
LS-1(4 nt)	CAGACTGATGTTGATTT-SH	
LS-1(6 nt)	AGACTGATGTTGATTT-SH	
LS-1(8 nt)	GACTGATGTTGATTT-SH	
LS-2	CTTTGGGGTAGCTTTTT-SH	
HP-1(4 nt)	TCAACATCAGTCTG <u>ATAA</u> GCTACCCCTTTGGGGGTA GC-FITC	
HP-1 (6 nt)	TCAACATCAGTCT <u>GATAAG</u> CTACCCCTTTGGGGGTA G-FITC	
HP-1 (8 nt)	TCAACATCAGTC <u>TGATAAGC</u> TACCCCTTTGGGGGTA -FITC	
HP-2	TAMRA-GCTACCCCAAAGGGGGTAGCTTATCAGACT GATAGTCTGATAA	
Molecular beacon	FITC-TCAACATCAGTCTGATAAGCTAAAAAAAAAA GATGTTGA-BHQ1	
miR-21	UAGCUUAUCAGACUGAUGUUGA	
miR-21 mimic	UAGCUUAUCAGACUGAUGUUGA	
miR-21 inhibitor	-21 inhibitor TCA ACATCAGTCTGATAAGCTA	
miR-21 forward	GTGCAGGGTCCGAGGT	
miR-21 reverse	GCCGCTAGCTTATCAGACTGATGT	
U6 forward	CTCGCTTCGGCAGCACA	
U6 reverse	AACGCTTCACGAATTTGCGT	
miR-16	UAGCAGCACGUAAAUAUUGGCG	
miR-26a	UUCAAGUAAUCCAGGAUAGGCU	
miR-214	ACAGCAGGCACAGACAGGCAGU	

Table S2. Comparison of the assay performance of the nanomachine with previously reported methods for intracellular miRNA detection.

	Method	LOD	Readout	Ref.		
1	Au Nanoflare Probe	0.68 nM	Fluorescence	1		
2	Isothermal Circular Strand Displacement	129.4 pM	Fluorescence	2		
	Polymerization (B-ICSDP)					
3	Hairpin-Fuelled Catalytic Nanobeacons	67 pM	Fluorescence	3		
4	Single-Layer Perfluorinated Tungsten	0.75 nM	Fluorescence	4		
	Diselenide Nanoplatform					
5	DNAzyme Amplification Strategy	44 pM	Fluorescence	5		
6	HCR System	680 pM	Fluorescence	6		
7	DNA Walking Nanomachine	26 pM	Fluorescence	This work		

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