Supporting Information

A dual-trigger entropy drive circuit based on competitive hybridization for high-specific enzyme-free detection of single nucleotide polymorphisms

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Table S1 The oligonucleotides used in the experiments

Name	Sequence (5'-3')
A	BHQ3-TTAAGATCTCAGGGGTGGCAAGCACTGATCAGTCACACTAG TTAAGCTGTAAGGGG-BHQ1
A1	GCACTGATCAGTCACACTAGTTAAGCTGTAAGGGG-BHQ1
B1	TTGCCACCCCTGA-CY5
B2	FAM-AGCTTAACTAGTG
B3	TGACTGATCAGTGC
MT	ACCCCTTACATCTTAAC
WT	ACCCCTTAGATCTTAAC
F1	TGACTGATCAGTGCTTGCCACCCCTGAGA
F2	ACAGCTTAACTAGTGTGACTGATCAGTGC
Mis-1	ACCCCTTAAATCTTAAC
Mis-2	ACCCCTTATATCTTAAC
Mis-3	ACGCCTTATATCTTAAC
Mis-4	ACCCCTTATATCTTGAC

Table S2 The recovery rate test of the detection limit

Add	1 fM	5 fM	10 fM
Test	1.3 fm	5.9 fm	10.9 fm
Recovery	130.3%	118.5%	109.1%