

**Colorimetric and photothermal dual-modal analysis of
cancer-related gene based on G-quadruplex-hemin cascade
signal amplification**

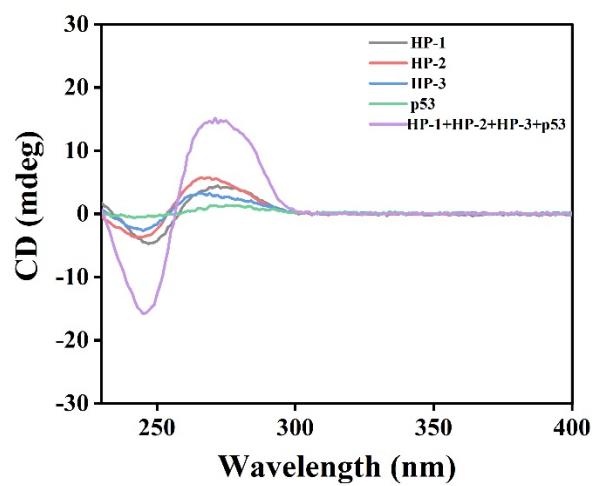


Fig S1. Circular dichroism of different DNA strands.

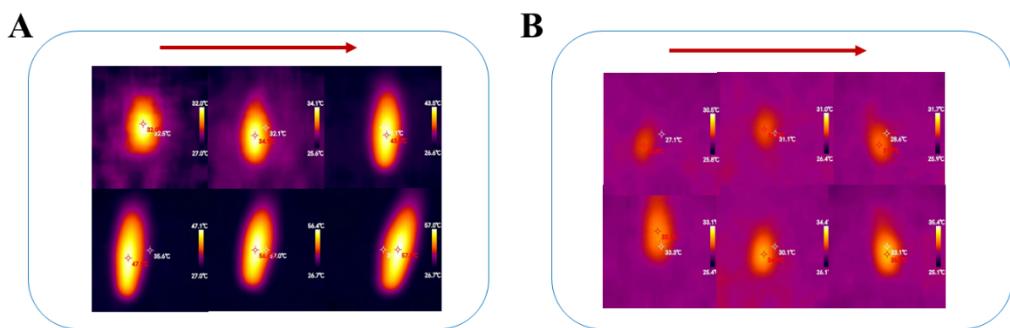


Fig S2. (A) Thermograms of different mutant p53 gene concentrations in the presence of I⁻; Thermograms of different mutant p53 gene concentrations in the absence of I⁻.

Table S1. DNA sequences used in this experiment

Primer name	DNA sequence (from 5'-3')	Number of bases
HP-1	AAA AAA ACA AATTAA ACA CCT TGA GTC TTC CAG TGT GAT GAA GAA GAA GGT GTT TAA GTA	60
HP-2	AGG GCG GGT GGG TGT TTA AGT TGG AGA ATT GTA CTT AAA CAC CTT CTT CTT GGG T	55
HP-3	TGG GTC AAT TCT CCA ACT TAA ACT AGA AGA AGG TGT TTA AGT TGG GTA GGG CGG G	55
mutant p53	TCA TCA CAC TGG AAG ACT C	19
A	TCG TCA CAC TGG AAG ACT C	19
B	TCA TCA CAC CGG AAG ACT C	19
C	TCA TCA CAC TGG AAG GCT C	19
D	TCG TCA CAC CGG AAG ACT C	19
E	TCA TCA CAC CGG AAG GCT C	19
F	TCG TCA CAC CGG AAG GCT C	19

Table S2 Recovery assay of mutant p53 gene in human serum samples

Sample	Add (U L⁻¹)	Found (U L⁻¹)	RSD (%)	Recovery (%)
1	0.000	No Found		
2	0.100	0.100		100.8
		0.096	0.048	95.99
		0.100		100.7
3	10.00	9.991		99.91
		10.02	0.219	100.2
		9.977		99.77
4	100.0	100.57		100.6
		100.38	1.910	100.4
		100.76		100.8