

Supporting Information

Colorimetric Detection of Norovirus in Oyster Samples through DNAzyme as a Signaling

Probe

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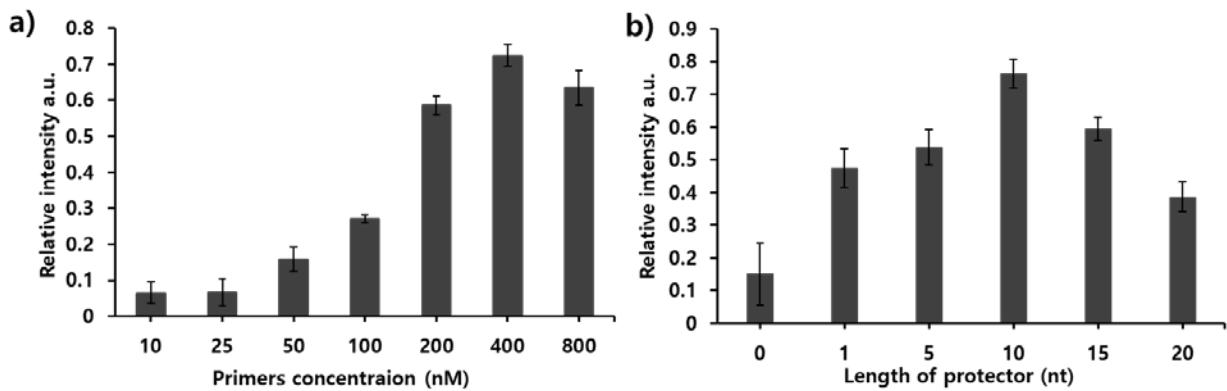


Figure S1. Optimization of colorimetric detection of noroviruses. (a) Primer concentration; (b) Protector sequence length.

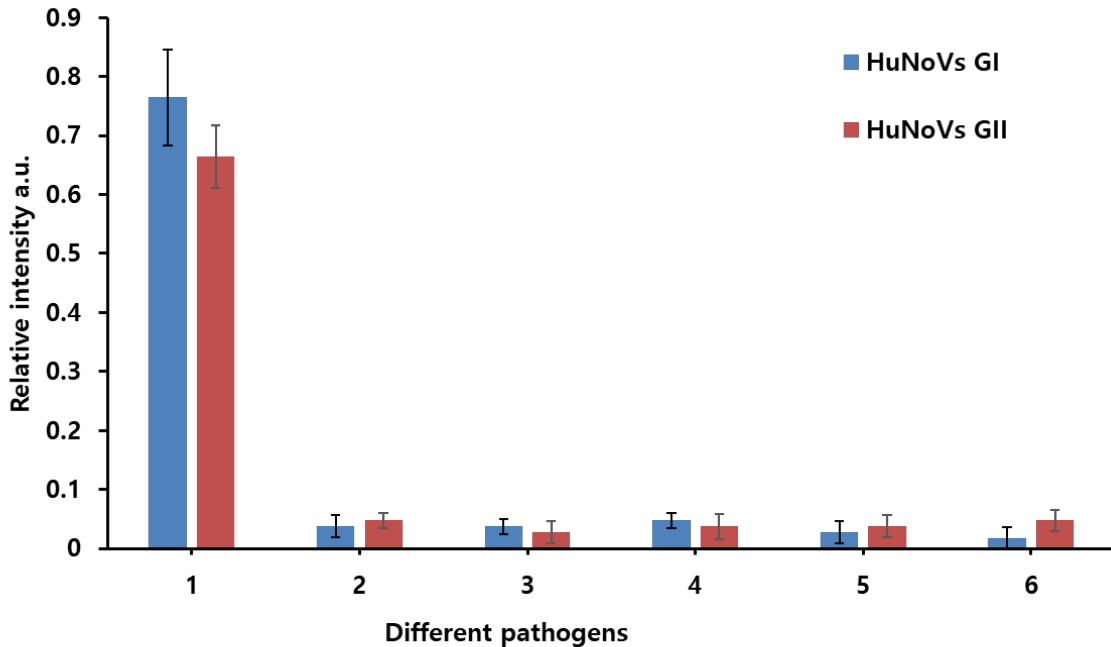


Figure S2. Specificity of proposed strategy towards HuNoVs GI and GII. 1, HuNoVs; 2, *E. coli* O157:H7; 3, *Listeria monocytogenes*; 4, *Vibrio parahaemolyticus*; 5, *Salmonella* Typhimurium; 6, *Bacillus cereus*. The concentration of HuNoVs GI, HuNoVs GII, and all bacteria were 10^3 , 10^3 , and 10^4 cfu mL $^{-1}$, respectively. Error bars are represent the standard deviations from three representative experiments (n = 3).

Table S1. The comparison of proposed strategy with other reported methods for the detection of HuNoV

Detection methods	Detection limit	Sample tested	Detection time (min)	Reference
Turbidimeter detection (RT-LAMP)	10^2 and 10^3 copies/tube, HuNoV GI and GII	Buffer	90	1
Colorimetric detection (RT-LAMP) hydroxynaphthol blue dye (HNB)	10^3 copies per reaction, HuNoV GII	Buffer	60	2
Fluorescence detection (RT-LAMP)	10 copies/reaction, Murine norovirus GI	Stool	45	3
Electrochemical detection (gold-immobilized synthetic peptide)	7.8 copies/mL, HuNoV GI	Buffer	120	4
Electrochemical detection (graphene-gold nano-composite aptasensor)	100 pM, HuNoV GII	Biological fluid	50	5
Fluorescence detection (FAM-labeled aptamer)	3.3 ng/mL, HuNoV GII	Mussel tissue	50	6
Colorimetric detection (split G-quadruplex)	4 nM, HuNoV GII	Buffer	160	7
Colorimetric (HRPzyme-integrated PCR)	Single copy/tube, HuNoV GI and GII	Oyster	60	This study

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