

## Supporting Information

### **A DNA biosensors-based microfluidic platform for attomolar real-time detection of unamplified SARS-CoV-2 virus**

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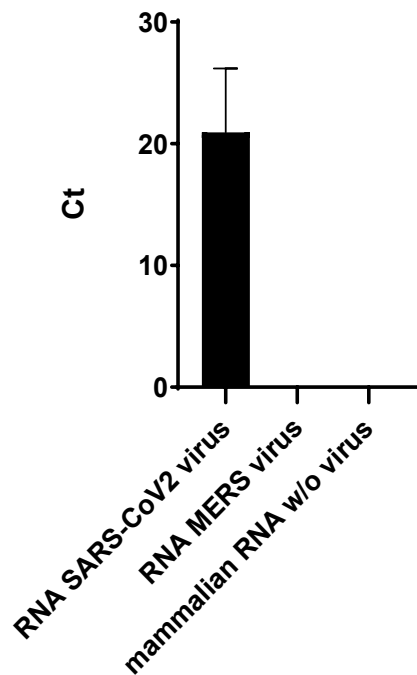
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## Specificity of PP1 for SARS-CoV-2 virus



**Figure S1.** Specific amplification of RNA from SARS-CoV-2 positive sample and not from SARS-CoV-2 negative samples. The primers used to perform the qPCR were: 5' TGAAGCAAGGTGAAATCAAGGA 3' and 5' AACAGCAAGAAGTGCAACGCCAAC 3' (as probe **PP1**).

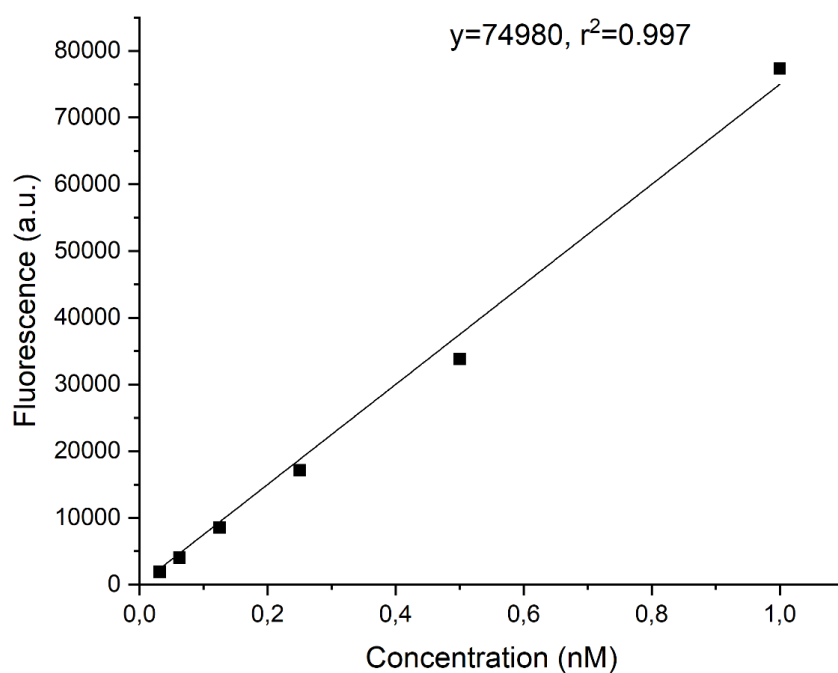
## Quantification of immobilized ssDNA probe on functionalized SiO<sub>2</sub> slides

Slide	S-PP1	S-Suc (negative control)
DNA amount quantified (per face)	2.90 pmol 1.94 pmol 2.46 pmol 3.57 pmol	0.11 pmol 0.21 pmol 0.00 pmol
<b>Average</b>	<b>2.7 ± 0.7 pmol</b>	<b>0.10 ± 0.10 pmol</b>

**Table S1.** Quantity of surface-immobilized ssDNA probe on **S-PP1** and **S-Suc** slides.

Name	Sequence
PP1	5'AmMC6-AACAGCAAGAAGTGCAACGCCAAC
Cy3-comp. PP1	5'Cy3-GTTGGCGTTGCACTTCTTGCTGTT

**Table S2.** DNA sequences corresponding to PP1 and 5'-Cy3-labelled complementary DNA strand.

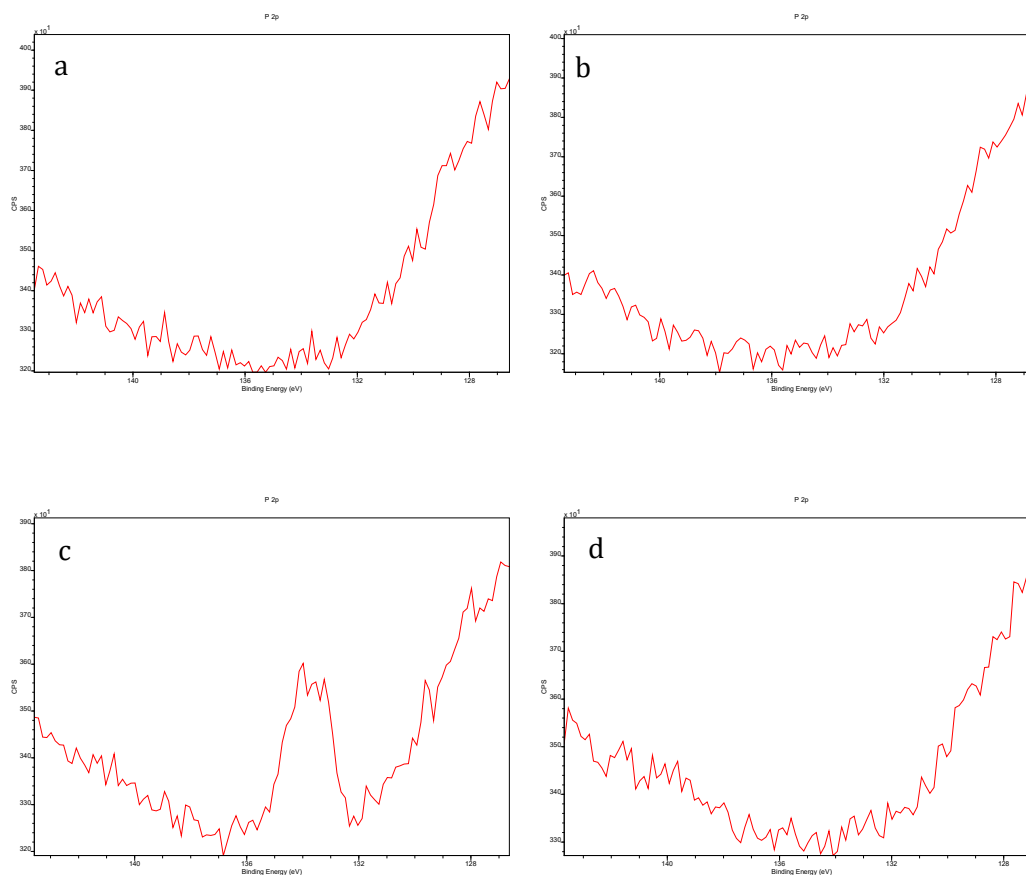


**Figure S2.** Representative standard curve for the quantification of Cy3-complementary **PP1** strand liberated from a surface.

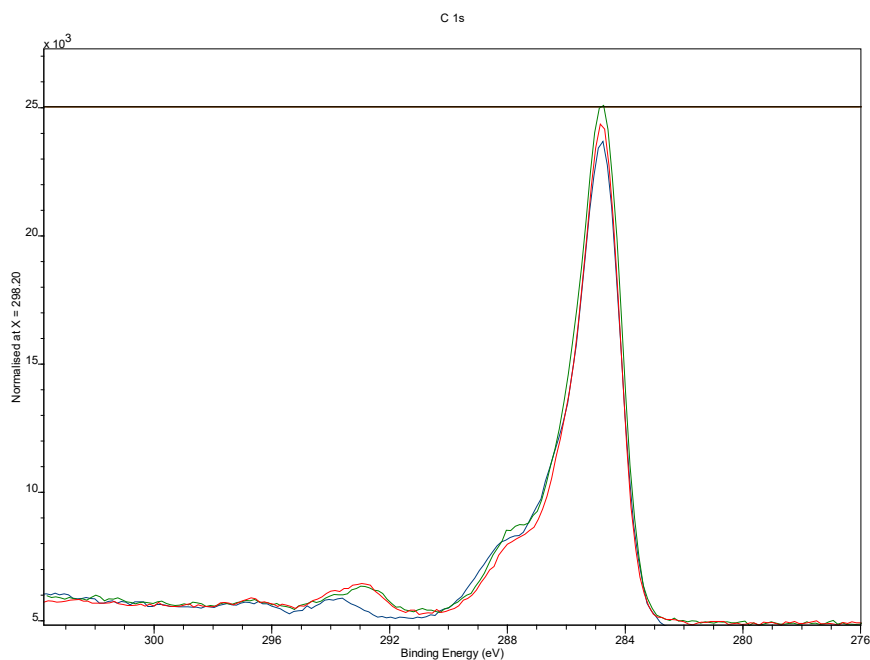
## Characterization of functionalized SiO<sub>2</sub> slides by XPS

Slides	Si at%	C at%	N at%	P at%
S-NH <sub>2</sub>	42.48	49.15	8.37	0.00
S-Suc	39.32	52.62	8.06	0.00
S-PP1	38.77	54.08	6.86	0.30

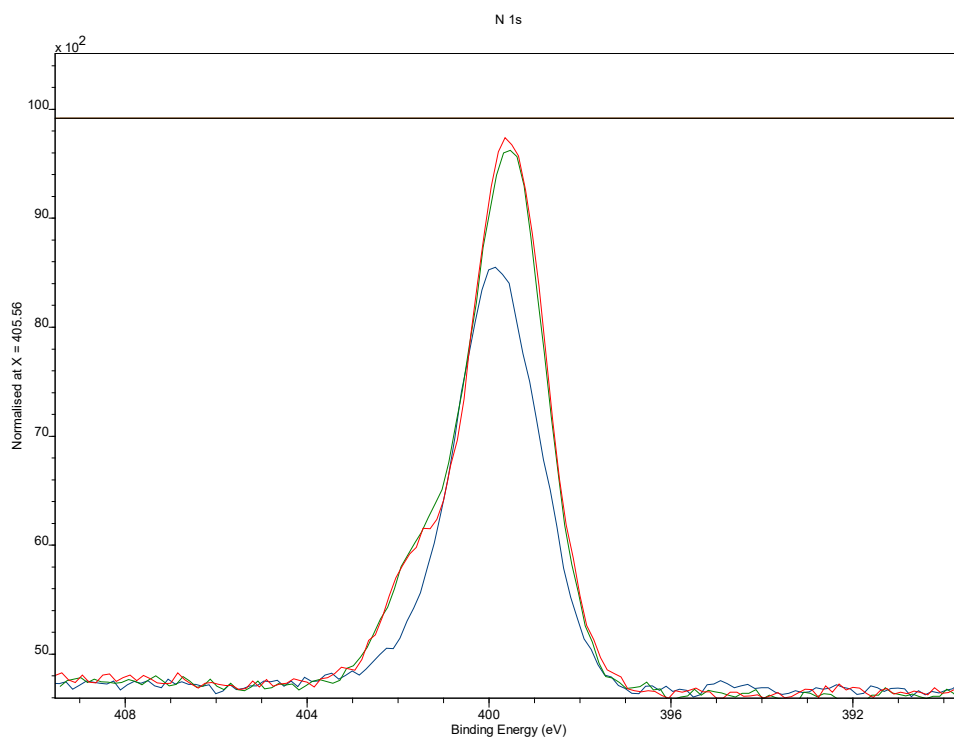
**Table S3.** Surface relative atomic concentration of C, N, O and S detected via XPS for amino-modified slides **S-NH<sub>2</sub>**, succinic-modified slides **S-Suc** and DNA-functionalized slides **S-PP1**.



**Figure S3.** XPS spectra of P 2p region of (a) amino-modified slides **S-NH<sub>2</sub>**; (b) **S-Suc** slides; (c) DNA-functionalized slides **S-PP1**; (d) **S-Suc** slides incubated with **PP1** probes without coupling agent.



**Figure S4.** XPS spectra of C 1s region of amino-modified slides **S-NH<sub>2</sub>** (red), succinic-modified slides **S-Suc** (green) and DNA-functionalized slides **S-PP1** (blue). Normalized at 298.20eV.



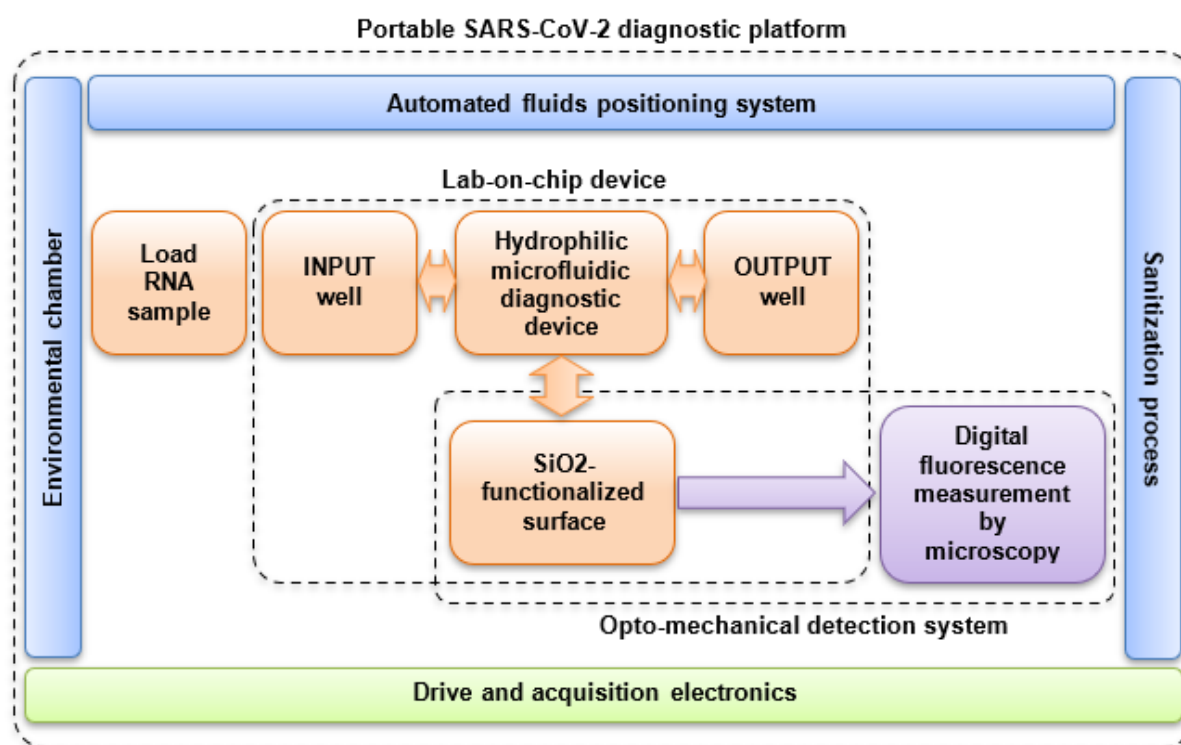
**Figure S5.** XPS spectra of N 1s region of amino-modified slides **S-NH<sub>2</sub>** (red), succinic-modified slides **S-Suc** (green) and DNA-functionalized slides **S-PP1** (blue). Normalized at 405.56eV

### Assessment of the stability of S-PP1 slides

Storage conditions	SSC 4X	MiliQ	Tris.HCl 1M	PBS 1X	Dried without solvent	Dried with solvents
DNA density (pmol/cm <sup>2</sup> ) After 10 days	3.07 ± 0.3	2.25 ± 0.78	3.89 ± 0.43	3.00 ± 0.53	2.52 ± 0.13	1.50 ± 0.2
DNA density (pmol/cm <sup>2</sup> ) After 15 days	/	2.55 ± 0.71	/	/	/	/
DNA density (pmol/cm <sup>2</sup> ) After 20 days	/	2.63 ± 0.62	/	/	/	/

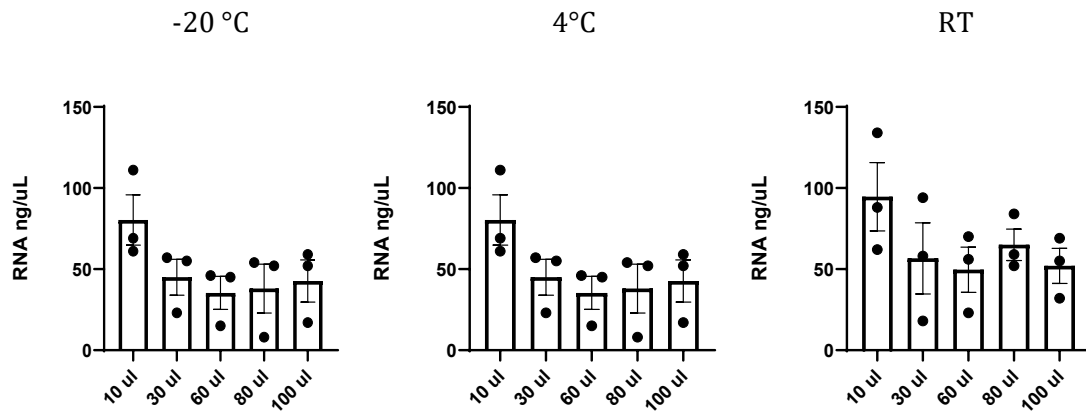
**Table S4.** Quantification of surface immobilized **PP1** DNA probe on **S-PP1** slides after 10 days of storage at 4°C in different media. Quantification of surface immobilized **PP1** DNA probe on **S-PP1** slides after 15 and 20 days of storage at 4°C in MilliQ. Values are expressed as mean ± standard deviation (triplicate).

### Conception diagram of the automated SARS-CoV-2 diagnostics platform



**Figure S6.** Main block diagram of the automatic device for viral screening via DNA-based biosensors.

## Comparison of RNA extraction protocols



**Figure S7.** RNA extraction performed using different concentration of human saliva mixed with Lucigen Quick Extract DNA kit (1:1). No significant difference was observed keeping the kit at -20°C (as manufacturer's instruction), 4°C or RT.



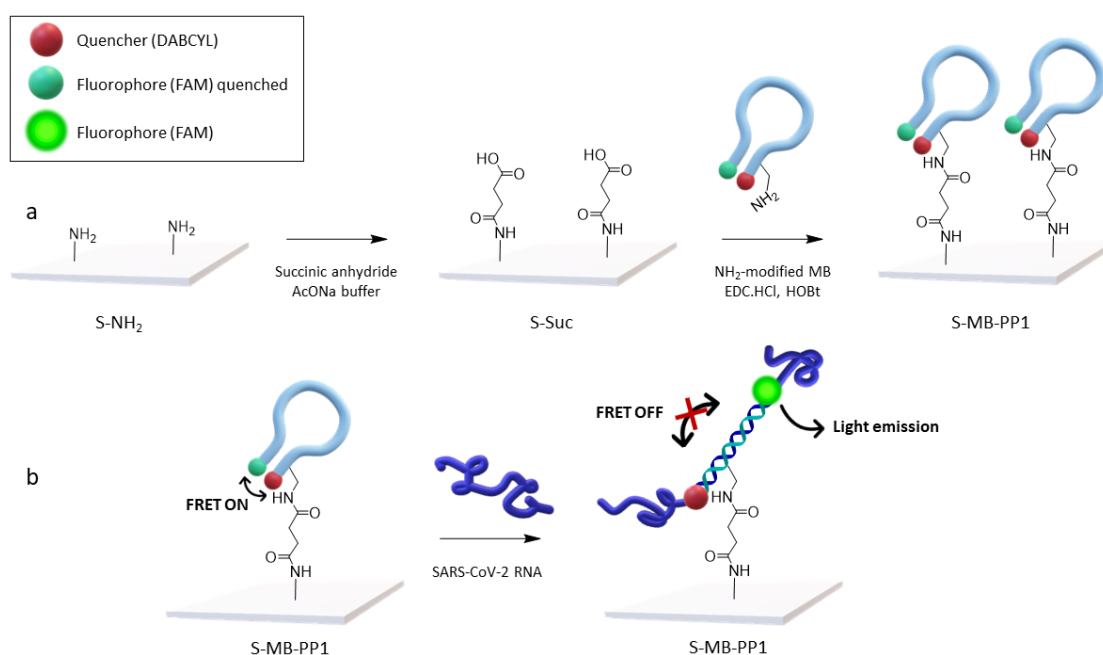
## Evaluation of sensing slides functionalized with the molecular beacon MB-PP1 probe

Name	Sequence
MB-PP1	5'AmMC6-TT TTT TTT TTT TT-FAM- TGA CGG AAC AGC AAG AAG TGC AAC GCC AAC CCG TCA-DABCYL-3'

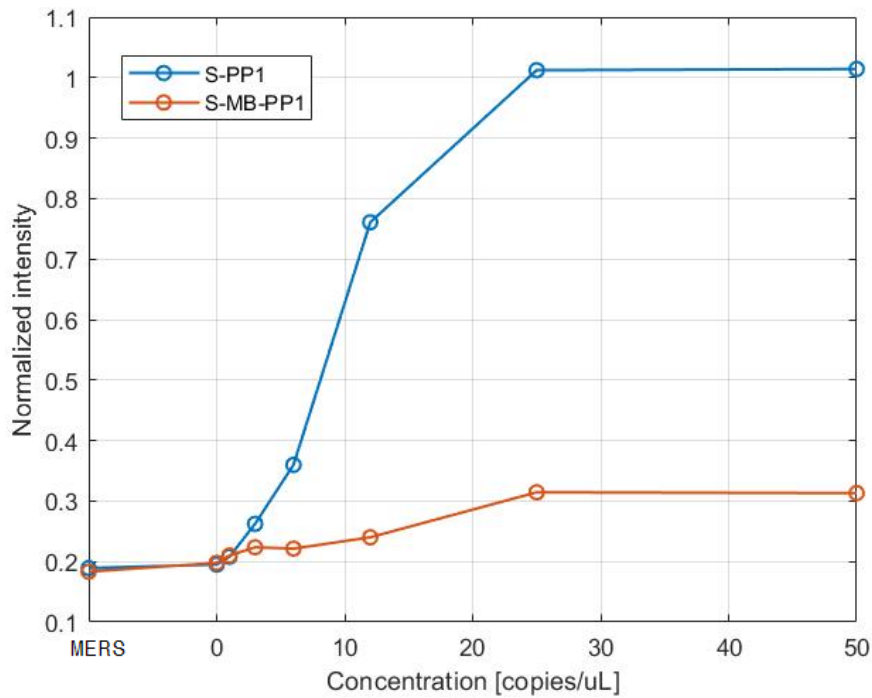
**Table S5.** DNA sequence corresponding to **MB-PP1**. DABCYL: N-[4-(4-dimethylamino)phenylazo]benzoic acid; FAM: Carboxyfluorescein. The FAM fluorophore was internally integrated to the probe through click chemistry.

Slide	S-MB-PP1	S-Suc (negative control)
DNA amount quantified (per face)	5.42 pmol 3.67 pmol	0.73 pmol 0.00 pmol
Average	4.5 ± 0.9 pmol	0.37 ± 0.52 pmol

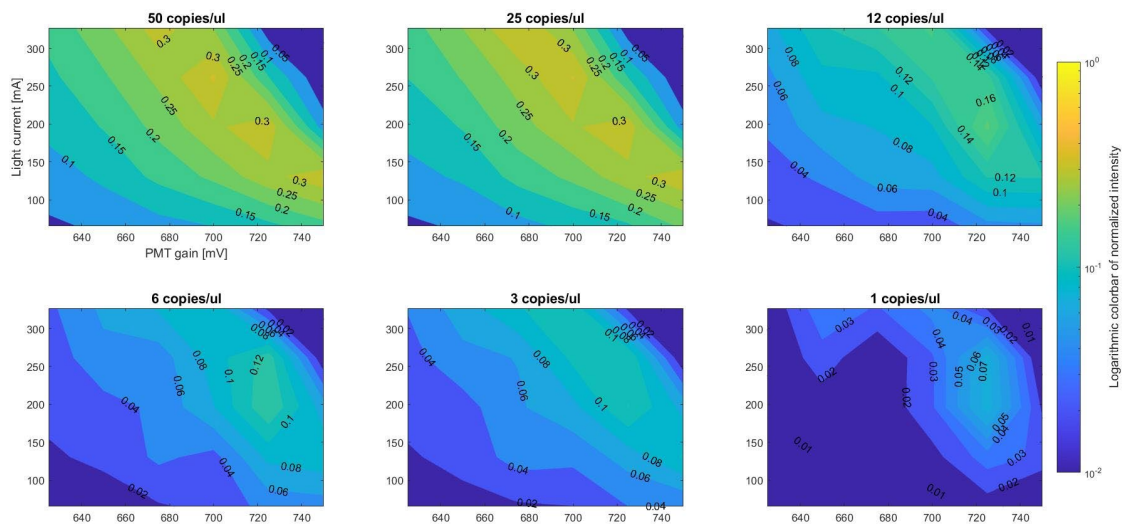
**Table S6.** Quantity of surface-immobilized **MB-PP1** probes on **S-MB-PP1** and **S-Suc** slides



**Scheme S1.** a: Conjugation of **MB-PP1** probes to the surface of sensing slides. b: Principle of SARS-CoV-2 RNA detection with **S-MB-PP1** slides. DABCYL: N-[4-(4-dimethylamino)phenylazo]benzoic acid; FAM: Carboxyfluorescein; FRET: Fluorescence Resonance Energy Transfer.



**Figure S8.** Comparison of the sensing properties of **S-PP1** and **S-MB-PP1** slides. Light intensities emitted by SYBR Green I (**S-PP1** samples) and carboxyfluorescein (**S-MB-PP1** samples) were converted by the PMT and normalized by subtraction of baseline values (**S-PP1** slides with SYBR Green I only or **S-MB-PP1** slides). **S-PP1** and **S-MB-PP1** were incubated with saliva samples containing SARS-CoV-2 virus at concentrations of 50, 25, 12, 6, 3, 1 copies/μL or MERS virus at a concentration of 50 copies/μL.



**Figure S9.** Difference between **S-MB-PP1** samples and baseline at decreasing SARS-CoV-2 viral loads (50, 25, 12, 6, 3, 1 copies/μL) represented as logarithmic colorbars of normalized intensities over LED current intensity (mA) and PMT gain (mV).