## **Supplementary Information**

## SARS-CoV-2 detection using a nanobody-functionalized voltammetric device

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**Figure S1**. (a) Affinity and size exclusion chromatography: (b) SDS-PAGE, acrylamide 15 %, Coomassie blue staining: 1: Total. 2: Pellet. 3: Soluble fraction. 4: Nickel / Flow-through. 5: Gel filtration / Elution



**Figure S2.** SPR binding curve of RBD to VHH-72 modified gold SPR chip (recorded on Biacore T200). The SPR binding corves were recorded on a Biocore T200 (Cytiva Life Science, using as running buffer HBS-P+  $1\times$  (containing 0.1 M HEPES, 1.5 M NaCl and 0.5% v/v Surfactant P20. A CM5 SPR chip was employed and after activation using NHS/EDC (15 mM in HBS-P+  $1\times$ ) for 30 sec (flow rate= 5µL min<sup>-1</sup>, VHH-72 (100 µg mL<sup>-1</sup> in HBS-P+  $1\times$ ) was flown over surface for 30 sec at 5µL min<sup>-1</sup> to reach a modification response of 200 RU (equal to 0.2 ng mm<sup>-2</sup>). The interaction with RBD (in HBS-P+) was at a flow of 30µL min<sup>-1</sup>.



**Figure S3. High resolution of the C1s region of interfaces E1-E5 (Table 1)**. The gold surface was modified with 3-mercaptopropinic acid before (**E1**) and after (**E2**) VHH-72 immobilization. (E3) Direct linkage of VHH-72. Gold electrode modified with 3-mercaptopropinic acid, followed by NH<sub>2</sub>-PEG-maleimide linkage before (**E4**) and after (**E5**) VHH-72-13C as immobilization.

|           |   | Atomic percentage (%) |            |         |         |
|-----------|---|-----------------------|------------|---------|---------|
| Electrode | Modification strategy                           | C1s                   | <b>O1s</b> | N1s     | S2p     |
| <b>E1</b> | 3-Mercaptopropionic acid                        | 75.3±0.3              | 20.6±0.2   | -       | 4.1±0.5 |
|           | (MPA)   |                       |            |         |         |
| E2        | 1. MPA  | 69.8±0.5              | 18.9±0.4   | 8.9±0.2 | 2.4±0.5 |
|           | 2. VHH-72                                       |                       |            |         |         |
| E3        | 1. VHH-72-13C                                   | 69.9±0.4              | 20.2±0.2   | 8.8±0.4 | 1.1±0.2 |
| <b>E4</b> | 1. MPA  | 69.5±0.4              | 20.1±0.8   | 6.2±0.5 | 4.2±0.5 |
|           | 2. NH <sub>2</sub> -PEG <sub>6</sub> -maleimide |                       |            |         |         |
| E5        | 1. 3-Mercaptopropionic acid                     | 65.3±0.5              | 21.4±0.6   | 9.2±0.2 | 4.1±0.7 |
|           | 2. $NH_2$ -PEG <sub>6</sub> -maleimide          |                       |            |         |         |
|           | 3. VHH-72-13C (ou                               |                       |            |         |         |
|           | VHH11D4-13C)                                    |                       |            |         |         |

Table S1. Element percentage obtained from XPS analysis for electrodes E1-E5

The success of the integration of VHH-72 was validated by XPS analysis. The C1s high resolution spectra of the 5 interfaces are seen in **Figure S3**. The C1s spectrum of **E1** can be deconvoluted into bands at 285.0 eV (C-C/C-H), 286.7 eV (C-S/C-O) and 288.2 (O-C=O) eV in accordance with the chemical composition of the electrode surface. The C1s high resolution spectrum of the surface **E2**, prepared by direct linking of VHH-72 *via* amide bond formation,

can be curve-fitted with several bands at 285.0 eV (C-C/C-H), 286.7 eV (C-N/C-S/C-O) and 288.6 eV (O-C=O/ N-C=O).

The C1s spectrum of interface **E3**, obtained through the direct linkage of VHH-C13 to gold surface, shows comparable features as E2, with bands at 285.0 eV (C-C/C-H), 286.7 eV (C-S/C-O/C-N) and 288.6 eV (O-C=O/N-C=O).

The core level XPS spectrum of the C1s of the pegylated gold interface functionalised with maleimide terminal group (**E4**) can be fitted with several components at 285.0 eV (C-C/C-H), 286.3 eV (C-O) and 288.7 eV (C=O and N-C=O), in full agreement with the chemical composition of the surface. Incorporation of VHH-72-C13 or VHH-11D4-13C *via* thiol-maleimide bond formation (**E5**) induces slight changes in the C1s band with contributions at 285.0 eV (C-C/C-H), 286.3 eV (C-O, C-N, C-S) and 288.2 eV (C=O). The increase of the band at 286.3 eV is due to the introduction of additional C-N and C-S bonds brought by the VHH-72-C13 or VHH-11D4-13C.



Figure S4. Differential pulse voltammograms of different electrodes: (a) Gold electrode before (black) and after (blue) VHH-72 immobilization (surface **E2**) in ferrocenemethanol (1 mM in 0.1 M PBS, pH 7.4). (b) Gold electrode before (black) and after (blue) VHH-72 (surface **E2**) immobilization in  $Fe(CN)_6^{4-/3-}$  (1 mM in 0.1 M PBS, pH 7.4). (c) Gold electrode modified with VHH-72-13C (surface **E3**) and via maleimide-modified PEG linker (surface **E5**) immobilization in ferrocenemethanol (1 mM in 0.1 M PBS, pH 7.4).



Figure S5: Correlation of Ct values with SARS-CoV-2 infectivity: Vero E6 cells ( $2.5 \times 10^5$  cells/well) were infected with 10-fold dilutions of a SARS-CoV-2 isolate. The plates were incubated for 6 days in 5% CO<sub>2</sub> atmosphere at 37 °C and examined daily using an inverted microscope (ZEISS Primovert) to evaluate the extent of the virus-induced cytopathic effect in cell culture. Calculation of estimated virus concentration was carried out by the Spearman and Karber method<sup>1,2</sup> and expressed as TCID<sub>50</sub>/mL (50% tissue culture infectious dose). TCID<sub>50</sub>/mL values were transformed to PFU mL<sup>-1</sup> by using the formula PFU mL<sup>-1</sup> = TCID<sub>50</sub>/mL × 0.7 RNA extraction.<sup>3</sup> Negative RT-PCR results were set to Ct= 50. The results are expressed as the mean ± SEM of at least 4 independent samples for each group.



Figure S6: Calibration curves for RBD on VHH-72-13C and VHH11D4-13C modified interfaces. Change in current density responses towards RBD standard addition on gold electrodes modified with (a) VHH-72-C13 *via* direct linkage (surface E3) with a PEG linker (surface E5), and (b) CHH11D4-13C using ferrocenemethanol (1 mM in 0.1 M PBS, pH 7.4) as redox probe. The results are expressed as the mean  $\pm$  SEM of at least 3 independent samples for each group.

## **Supplementary References**

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