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# Adsorption and Inactivation of SARS-CoV-2 on the surface of anatase TiO<sub>2</sub>(101)

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## Abstract

We investigated the adsorption of severe acute respiratory syndrome corona virus 2 (SARS-CoV-2), the virus responsible for the current pandemic, on the surface of the model catalyst  $\text{TiO}_2(101)$  using atomic force microscopy, transmission electron microscopy, fluorescence microscopy and X-ray photoelectron spectroscopy, accompanied by density functional theory calculations. Three different methods were employed to inactivate the virus after it was loaded on the surface of  $\text{TiO}_2(101)$ : i) ethanol, ii) thermal and iii) UV treatments. Microscopic studies demonstrate that the denatured spike proteins and other proteins in the virus structure readsorb on the surface of  $\text{TiO}_2$  under thermal and UV treatments. The interaction of the virus with the surface of  $\text{TiO}_2$  was different for the thermally and UV treated samples compared to the sample inactivated via ethanol treatment. AFM and TEM results on the UV-treated sample suggested that the adsorbed viral particles undergo damage and photocatalytic oxidation at the surface of  $\text{TiO}_2(101)$  which can affect the structural proteins of SARS-CoV-2 and denature the spike proteins in 30 minutes. The role of Pd nanoparticles (NPs) was investigated in the interaction between SARS-CoV-2 and  $\text{TiO}_2(101)$ . The presence of Pd NPs enhanced the adsorption of the virus due to the possible interaction of the spike protein with the NPs. This study is the first investigation of the interaction of SARS-CoV-2 with the surface of single crystalline  $\text{TiO}_2(101)$  as a potential candidate for virus deactivation applications. Clarification of the interaction of the virus with the surface of semiconductor oxides will aid in obtaining a deeper understanding of the chemical processes involved in photo-inactivation of microorganisms, which is important for the design of effective photocatalysts for air purification and self-cleaning materials.

## Introduction

In late 2019, the Corona virus disease (COVID-19) caused by the severe acute respiratory syndrome corona virus 2 (SARS-CoV-2) led to a global pandemic.<sup>1</sup> SARS-CoV-2, with a spherical shape and a size of around 80-120 nm,<sup>2</sup> has four important structural proteins

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3 namely spike (S), membrane (M), envelope (E), and nucleocapsid (N) proteins and some  
4 non-structural proteins that can adjust the cellular responses<sup>3</sup>(Figure 1). S-protein, with a  
5 club-like shape of 20 nm in length, is the outermost structure in SARS-CoV-2 and is respon-  
6 sible for the entrance of the virus into the cell.<sup>4-6</sup> SARS-CoV-2 can be transmitted through  
7 droplets produced by the infected person, by direct contact, via contaminated surfaces and  
8 by the inhalation of small respiratory droplets and aerosol particles.<sup>7</sup> It has been reported  
9 that SARS-CoV-2 can survive more than three hours in aerosols<sup>8</sup> and is stable on stainless-  
10 steel surfaces and plastic with a half-life of approximately 5.6 and 6.8 hours, respectively.<sup>9</sup>  
11 Moreover, Ong et al. reported that the RNA of SARS-CoV-2 was observed in air.<sup>10</sup> Owing to  
12 the role of air and surface contamination in SARS-CoV-2 transmission, the development of  
13 self-decontaminating materials with low cytotoxicity should be considered. Recently, various  
14 treatment methods have been reported for the inactivation of viruses such as chemical dis-  
15 infection with alcohol,<sup>11</sup> formaldehyde,<sup>12</sup> peroxide,<sup>13</sup> hypochlorous acids<sup>14</sup> and mechanical  
16 sterilization using ultraviolet (UV) light irradiation<sup>15</sup> and heating.<sup>16</sup> Disinfection methods  
17 can affect the lipid bilayer and denature the spike proteins during the early stages of virus  
18 deactivation.<sup>17,18</sup> Due to the high surface stability of SARS-CoV-2 that enhances the ef-  
19 ficiency of virus transmission,<sup>17</sup> cost effective and efficient methods are required for virus  
20 inactivation on surfaces.

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39 Solid state antiviral materials are preferable to solvent-based ones due to their improved  
40 robustness and long term stability.<sup>19</sup> Oxide nanomaterials are promising candidates for the  
41 removal and inactivation of viral/bacterial infections as well as for preventing pathogen dis-  
42 semination.<sup>20,21</sup> Titanium dioxide (TiO<sub>2</sub>), with unique properties such as low toxicity, high  
43 photoactivity and high commercial availability compared to other semiconductors,<sup>22,23</sup> has  
44 received considerable attention for environmental applications such as solid-state antiviral  
45 materials, self-cleaning surfaces and air and water purification systems.<sup>24,25</sup> Multiple com-  
46 peting reaction pathways can be responsible for virus inactivation over nanomaterials, such  
47 as catalytic oxidation, metal ion release, photo-thermal effects, and formation of reactive  
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oxygen species (ROS).<sup>26</sup> Among different advanced oxidation processes, which are used for the elimination of organic and inorganic pollutants and inactivation of microorganisms, photocatalysis is the most desirable method owing to the high ability in inactivation of different types of contaminants.<sup>27</sup> By passing the polluted air over a photocatalyst, pollutants can adsorb on the photocatalyst surface and can be converted to harmless and low toxic compounds via the formation of holes ( $h^+$ ) with high oxidation potential<sup>19</sup> and ROS such as superoxide radical ( $\bullet O_2^-$ ), hydroxyl radical ( $\bullet OH$ ) and  $H_2O_2$  under light irradiation.<sup>28-30</sup> Consequently, photocatalyst surfaces can facilitate virus disinfection, mineralize the organic compounds in the virus structure<sup>31</sup> and inhibit the spread of the virus via the contaminated surfaces.<sup>32,33</sup> Different methods such as the combination of noble metal and metal oxides have been reported to improve the photocatalytic activity of  $TiO_2$  for the inactivation of SARS-CoV-2 in solution.<sup>19,34</sup> Nanoparticles (NPs) with a small size and high surface to volume ratio show high adsorption capacity for amino acids and peptides.<sup>35</sup> It was shown that gold (Au) and silver (Ag) NPs interact with peptide and human immunodeficiency virus 1 (HIV-1), respectively.<sup>36,37</sup> NPs can bind to the viral genome and modulate the viral transcription.<sup>38</sup> Recently, it was reported that SARS-CoV-2 can adsorb on metal surfaces such as Au NPs,<sup>39</sup> oxygen-containing substrates (e.g. glass, paper, wood),<sup>40</sup> and surface hydroxyl groups.<sup>41</sup> Depending on the surface functional groups, active sites and surface humidity, the adsorption capacity of SARS-CoV-2 can be influenced.<sup>41</sup> The functional groups of amino acids in S-proteins of virus, play a vital role in the adhesion process. Recently, the photoactivity of Ag- $TiO_2$  single atom nanozyme,<sup>42</sup> Ag NPs@ $TiO_2$ ,<sup>34</sup>  $Cu_xO/TiO_2$ <sup>43</sup> and  $TiO_2/Ti-O$  on  $Al_2O_3$  balls<sup>44</sup> have been investigated for the removal and inactivation of SARS-CoV-2. The studies were performed in aqueous solution and only the reaction solution was tested to identify the virus inactivation applying catalysts. In fact, most of the data surrounding microorganism inactivation, especially virus inactivation, is reported in the liquid phase<sup>34,42,43</sup> and there is limited knowledge about the adsorption of SARS-CoV-2 on the surface of catalysts or at the interface between the virus and the catalyst. Therefore, obtaining more detailed information

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3 regarding the adsorption/inactivation behavior of SARS-CoV-2 at the surface and interface  
4 of  $\text{TiO}_2$  as well as the role of NPs can lead to the development of efficient antiviral coatings  
5 and aid in the design of advanced materials for the photocatalytic inactivation of viruses.  
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7 In this study, we investigated the adsorption of SARS-CoV-2 on single crystalline  $\text{TiO}_2(101)$ ,  
8 the thermodynamically most stable facet of anatase,<sup>45</sup> and on  $\text{TiO}_2(101)$  supported Pd NPs.  
9 Thermal and ethanol treatments were employed to inactivate the SARS-CoV-2 particles ad-  
10 sorbed on the surface of  $\text{TiO}_2(101)$ . Moreover, the UV induced photo-inactivation of SARS-  
11 CoV-2 adsorbed on  $\text{TiO}_2(101)$  was investigated. These methods were employed before the  
12 samples were transferred out of the biosafety level (BSL)-3 laboratory. We characterized the  
13 samples using atomic force microscopy (AFM), transmission electron microscopy (TEM),  
14 scanning tunneling microscopy (STM), fluorescence microscopy (FM) and X-ray photoelec-  
15 tron spectroscopy (XPS), combined with density functional theory (DFT) calculations. The  
16 main objective of our present work is to identify the adsorption and inactivation behavior of  
17 SARS-CoV-2 on the surface of the model photocatalyst  $\text{TiO}_2(101)$ .  
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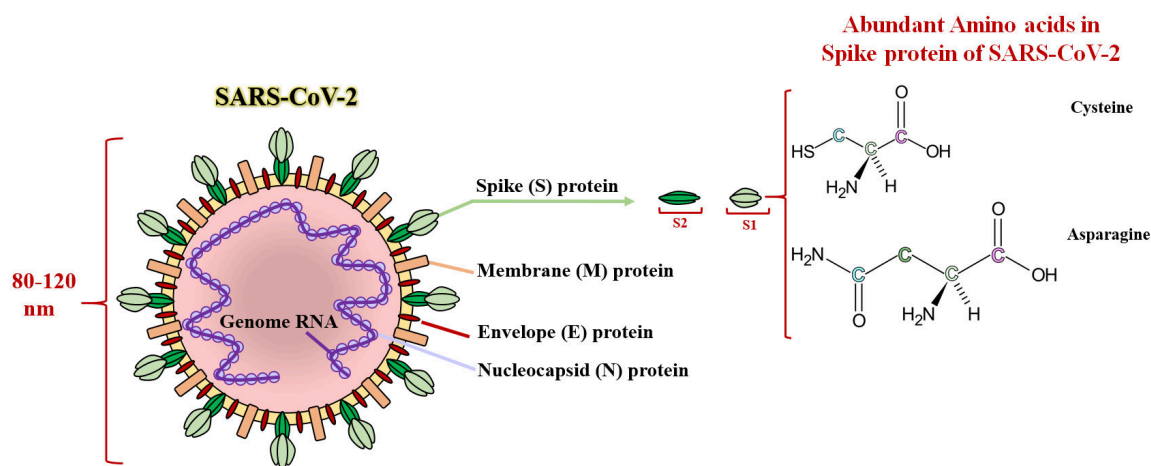


Figure 1: Schematic structure of SARS-CoV-2.

## Results and discussion

### Characterization of Cell Culture Media (CCM) on $\text{TiO}_2(101)$

As described in the methods section, SARS-CoV-2 suspension contains CCM. To obtain more information about the CCM structure, as well as to distinguish it from the virus particles, atomic force microscopy was employed to study the CCM on the surface of  $\text{TiO}_2(101)$ . According to the AFM images (Figure 2 a and b), no particle structure was observed. CCM forms a fiber-like structure on the surface of  $\text{TiO}_2(101)$  with a thickness of around 14-45 nm (Figure 2 c).

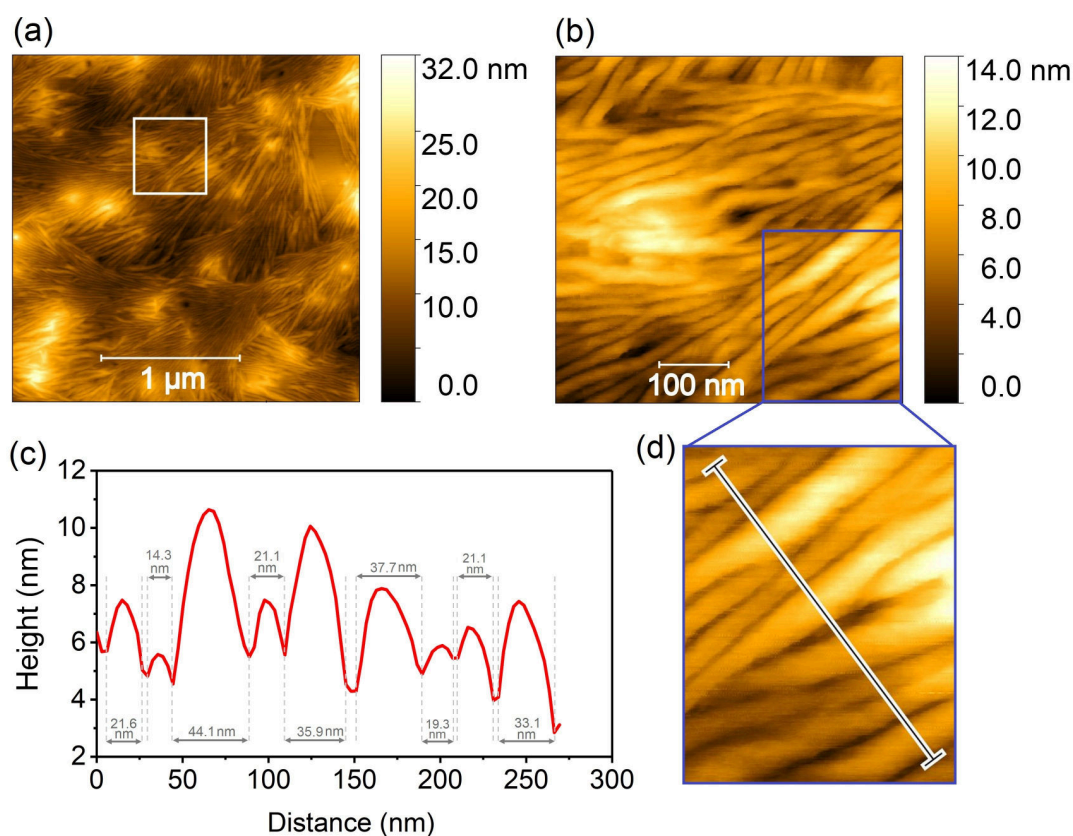


Figure 2: (a) AFM ( $2.5 \times 2.5 \mu\text{m}^2$ ) image of CCM adsorbed on the surface of  $\text{TiO}_2(101)$  after thermal treatment at  $70^\circ\text{C}$ . (b) AFM image ( $0.5 \times 0.5 \mu\text{m}^2$ ) of marked area with white rectangle in image (a). (c) Line scan profile from top to down in zoomed area (d) of image (b).

### Adsorption of SARS-CoV-2 on the surface of TiO<sub>2</sub>(101)

To further confirm the adsorption of SARS-CoV-2 on TiO<sub>2</sub>(101), fluorescence microscopy was applied. Anatase TiO<sub>2</sub>(101) coated with CCM was measured as a reference sample. Rabbit anti-SARS-CoV-2 nucleocapsid primary antibodies and anti-rabbit secondary antibodies conjugated to Alexa568 were used to fluorescently label the virus particles in the samples by indirect immunofluorescence. Mean fluorescence intensities for SARS-CoV-2 and CCM on TiO<sub>2</sub>(101) were obtained to be 6160 and 2902, respectively. The comparatively high fluorescence intensity of the virus sample confirms the adsorption of SARS-CoV-2 on the surface of TiO<sub>2</sub>. Compared to the virus sample, only faint green signal was detected for CCM on TiO<sub>2</sub>(101) (see Figure S1 in the supporting information) attributed to autofluorescent proteins and other components in CCM.

### Comparing the various inactivation methods of SARS-CoV-2 adsorbed on the surface of TiO<sub>2</sub>(101)

Inactivation of the Corona virus was necessary for the transfer of materials from the biosafety level (BSL)-3 laboratory to other laboratories allowing for accelerated studies against SARS-CoV-2 during the pandemic.<sup>13</sup> Recently, Lyonais et al. reported a new inactivation method in solution media that kept the virus particles intact but inactivated.<sup>12</sup> They found that under heating treatment (58 °C for 30 min) in solution media, the virus molecules lost their spherical structure and became damaged, while upon formaldehyde (FA) treatment SARS-CoV-2 was inactivated but the structure remained intact.<sup>12</sup> In this study, thermal inactivation was done by incubating the sample at 70 °C for 30 min after loading the SARS-CoV-2 on the surface of TiO<sub>2</sub>(101) (see Methods section) and then the sample was analyzed by AFM and TEM. Although this inactivation method denatures the proteins, it still allows one to obtain information about the adsorption behavior of SARS-CoV-2 at the surface of TiO<sub>2</sub>(101). Based on the AFM images shown in Figure 3 a and b (i), several small and large

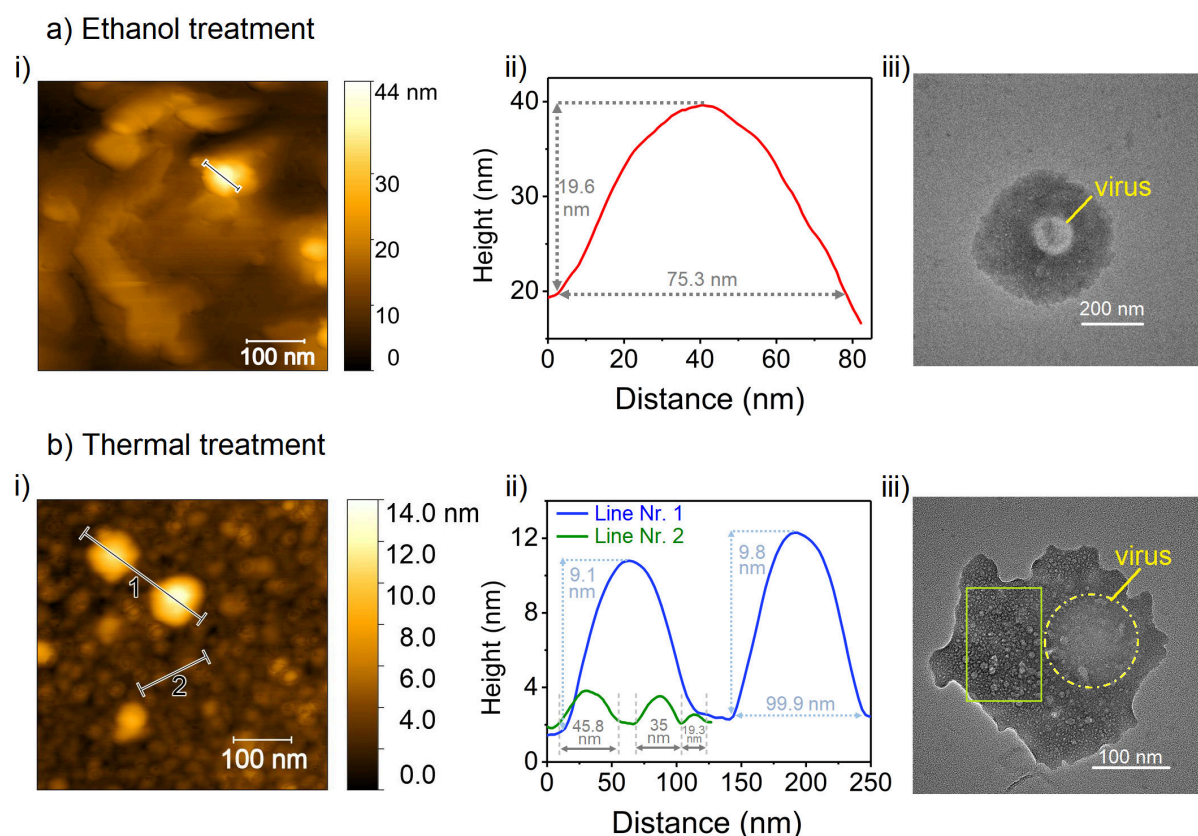


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3 particles were observed for SARS-CoV-2 adsorbed on TiO<sub>2</sub>(101) following thermal inactiva-  
4 tion, while no particles were detected in CCM on the surface of TiO<sub>2</sub> (Figure 2). Therefore,  
5 these particles come from the virus structure. According to the AFM images for ethanol and  
6 thermal treatments (Figure 3), the observed spherical particles with the particle diameter in  
7 the range of 75-100 nm correspond to the virus structure. However the height of the particles  
8 in these images is smaller than the expected virus diameter (80-120 nm). Due to drying of the  
9 sample, viruses become flatter on the surface of TiO<sub>2</sub>(101). Based on the line scan profiles  
10 (Figure 3a and b (ii)), the heights of adsorbed SARS-CoV-2 on TiO<sub>2</sub> are 19.6 nm and 9-10  
11 nm under ethanol and thermal treatments, respectively. It is known that temperature can  
12 affect the structure of biomacromolecules such as protein, lipids and nucleic acids.<sup>46,47</sup> Since  
13 the thermal treatment was performed before the AFM measurement, the virus proteins and  
14 membrane could have denatured and adsorbed on the surface of TiO<sub>2</sub>. Contrary to ethanol  
15 treatment, the area surrounding the virus is covered by the cell by-products layer, which have  
16 a smaller size than SARS-CoV-2 (Figure 3 b(ii)). In fact, after ethanol treatment most of  
17 the adsorbed virus particles were removed from the catalyst surface via the surface washing.  
18 Lipid bilayer envelope in SARS-CoV-2 contains the S, E, and M proteins.<sup>40</sup> The virus can be  
19 inactivated by disruption of the lipid envelope. Under ethanol treatment, ethanol dissolves  
20 this layer that leads to the inactivation of the virus.<sup>48</sup>

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23 Under ethanol and thermal treatments, AFM images could not show the S-protein on the  
24 virus structure. Due to the virus movements on the surface of TiO<sub>2</sub>(101) during the ad-  
25 sorption, and the subsequent inactivation methods, the S-proteins may break. We are not  
26 able to directly monitor the mechanism of the interaction of the virus with the surface of  
27 the catalyst and their subsequent self-assembly, as they could not be delivered out of the  
28 BSL-3 laboratory. However, for the first time we have clear evidence that the virus interacts  
29 with the surface and that the conditions at the interface play an instrumental role in the  
30 mechanism of adsorption and denaturation of the virus.

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33 Recently, Rath and Kumar reported that the spike protein conformation depends on tem-

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3 perature.<sup>47</sup> At temperatures exceeding 50 °C, the receptor binding motif (RBM) in the spike  
4 protein structure is completely closed, resulting in the inactivation of SARS-CoV-2. Clearly,  
5 ethanol washes the virus away, and therefore there are less virus particles on the surface of  
6 TiO<sub>2</sub>(101), while through thermal treatment the adsorbed viruses are dried on the surface  
7 of TiO<sub>2</sub> and therefore more proteins are dissociated from the virus structure after thermal  
8 treatment. Denatured S-proteins and other virus proteins (from the virus envelope and  
9 membrane) dissociate and adsorb at the surface of TiO<sub>2</sub>(101). Figure 3 a and b (iii) show  
10 the TEM images of SARS-CoV-2 adsorbed on TiO<sub>2</sub>(101) that was initially inactivated by  
11 ethanol and thermal treatments, respectively. Spherical particles with a size of around 110  
12 nm correspond to SARS-CoV-2. As we expected, compared to ethanol treatment, small par-  
13 ticles in the diameter range of 10-45 nm were observed in the TEM image of the thermally  
14 treated sample (see the labeled region with a yellow rectangle in Figure 3 b (iii)). These  
15 results are in good agreement with the AFM images shown in Figure 3 a and b, confirming  
16 the effect of heat on the virus structure and its inactivation.  
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31 Figure 3: AFM ( $0.5 \times 0.5 \mu\text{m}^2$ ) images of SARS-CoV-2 adsorbed on  $\text{TiO}_2(101)$  (i), line scan  
 32 profile measured along the particles in AFM images (black lines) (ii) and TEM images (iii)  
 33 of SARS-CoV-2 adsorbed on  $\text{TiO}_2(101)$  after ethanol (a) and thermal (b) treatments.  
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### 35 X-ray photoelectron spectroscopy (XPS) analysis of adsorbed cell culture media 36 (CCM) and inactivated SARS-CoV-2 on the surface of $\text{TiO}_2(101)$ 37

38 To obtain more information on the chemical composition of the system and to further prove  
 39 the adsorption and interaction of the virus with the surface of  $\text{TiO}_2(101)$  as well as the  
 40 role of different functional groups on virus adsorption, we performed XPS experiments. To  
 41 better distinguish the adsorbed virus from CCM using the XPS results, the adsorbed CCM  
 42 on  $\text{TiO}_2(101)$  before and after thermal treatment were also measured. We also studied the  
 43 interaction of the denaturated peptide chain of the spike protein theoretically through sim-  
 44 plified models involving single amino acids, i.e. cysteine (cys) or asparagine (asn), onto the  
 45 anatase  $\text{TiO}_2(101)$  surface, whose optimized structures and adsorption energies are reported  
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3 in Figure 4. Then, for each of these models, we calculated the C 1s binding energies.

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5 Our aim is not to determine the most stable adsorption configuration of the amino acids,  
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7 but it is to conceive configurations that resemble cysteine or asparagine as a part of a long  
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9 peptide chain and understanding their interactions with the surface. This goal requires that  
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11 either both or at least one of the amino acid groups involved in the peptide bonds of the  
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13 actual primary chain (amino on one side and carboxylic on the other) are only weakly inter-  
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15 acting with the surface. In the case of cysteine, we considered the carboxyl, the amino and  
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17 the thiol as adsorbing groups: our results suggest that covalent, electrostatic and hydrogen  
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19 bond interactions take place. The first model (Figure 4 b) is an exception with respect to  
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21 this requirement, because both the amino and the carboxylic groups are interacting strongly  
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23 with the surface, whereas the side chain of the amino acid containing the thiol (-SH) group  
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25 is not. This configuration has been computed for comparison with a similar adsorption  
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27 model, which has been previously reported in literature as the most favored configuration  
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29 for *cys*/TiO<sub>2</sub>(101).<sup>49</sup>

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31 In this work, we find that it is not as stable as another configuration (Figure 4 d), where  
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33 the thiol and the amino groups are coordinated to two surface Ti atoms and the OH of  
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35 the carboxylic group establishes a hydrogen bond with a surface O atom. We should note  
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37 here that the de-protonation of the thiol group is found to be unfavorable and has not been  
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39 reported. Also, the zwitterionic form of the amino acid on the surface was only found in the  
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41 model in Figure 4 b, where, the proton is transferred from the amino group to a surface O  
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43 atom. We expect that the presence of water in the simulations would further stabilize the  
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45 zwitterionic form, as suggested by several studies.<sup>49-51</sup> In the case of asparagine, we limited  
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47 the study to one configuration, where only the side chain is involved in the interaction with  
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49 the surface, except for a H-bond of the carboxylic OH, again with the aim of representing  
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51 asparagine as part of a virtual peptide chain.

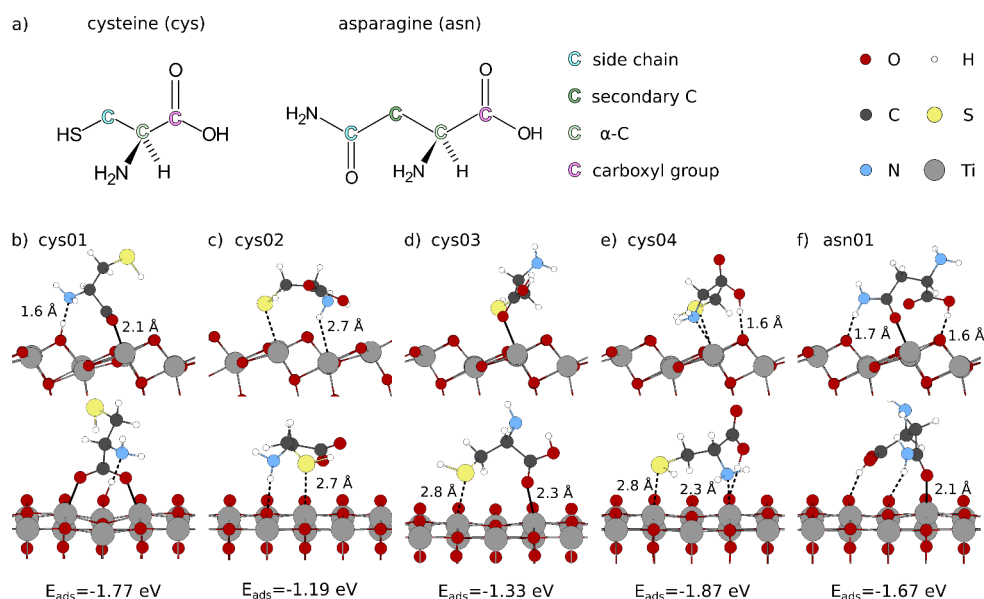


Figure 4: (a) Chemical structure of cysteine and asparagine in the neutral form, along with the chemical labeling of C atoms. (b)-(f) Left (upper row) and front (lower row) views of the optimized models of cys or asn adsorbed on anatase  $\text{TiO}_2(101)$  surface. In each panel, the adsorption energy ( $E_{\text{ads}} = E_{\text{cys,asn}/\text{TiO}_2} - E_{\text{cys,asn}} - E_{\text{TiO}_2}$ ) and the most relevant distances are reported. The color code of the atom is reported in the top corner of the image. The solid/dashed black lines mark covalent bonds/electrostatic or H-bond interactions.

Experimentally, we focused on the core-level spectra for the Ti 2p, C 1s, O 1s, and N 1s for the as-prepared surface of  $\text{TiO}_2$  (i), the same surface following the deposition of CCM (ii) and subsequent thermal treatment (iii), and inactivated SARS-CoV-2 on  $\text{TiO}_2$  after heating to 70 °C (iv) (Figure 5). Regarding the experimental Ti 2p core level spectra shown in Figure 5 a, two dominant peaks were observed at binding energies (BEs) 458.3 and 464.1 eV that correspond to the Ti 2p<sub>3/2</sub> and Ti 2p<sub>1/2</sub> spin-orbit split components  $\text{Ti}^{4+}$  in  $\text{TiO}_2(101)$ .<sup>52</sup> The small peaks located at 457.2 and 462.8 eV belong to the Ti 2p<sub>3/2</sub> and Ti 2p<sub>1/2</sub> components, attributed to small amounts of  $\text{Ti}^{3+}$ .<sup>53</sup> The C 1s core level for CCM can be fitted with five components at BEs of 283.6 eV (C=C),<sup>54</sup> 284.95 eV (C-H),<sup>55</sup> 286.5 eV (C-N and C=N),<sup>56</sup> C-OH,<sup>57</sup> C-SH<sup>58</sup>, 287.9 eV (C=O and O=C-N), and 289.4 eV (O=C-O).<sup>54</sup> Compared to CCM, the ratio of observed peaks at 284.95 eV and 286.5 eV was greater for the virus containing sample. As it was expected, this is due to the hydrocarbon groups present as side chains in the protein structure and this is greater in the virus adsorbed sample due to

a higher number of proteins and lipids. Figure 6a shows the difference spectrum of C 1s obtained by subtracting the C 1s spectrum of the virus from CCM. Based on this result, the contribution of virus in the C 1s spectrum was obtained. The dominant contribution was related to the C–H and C=O in the virus protein structure.

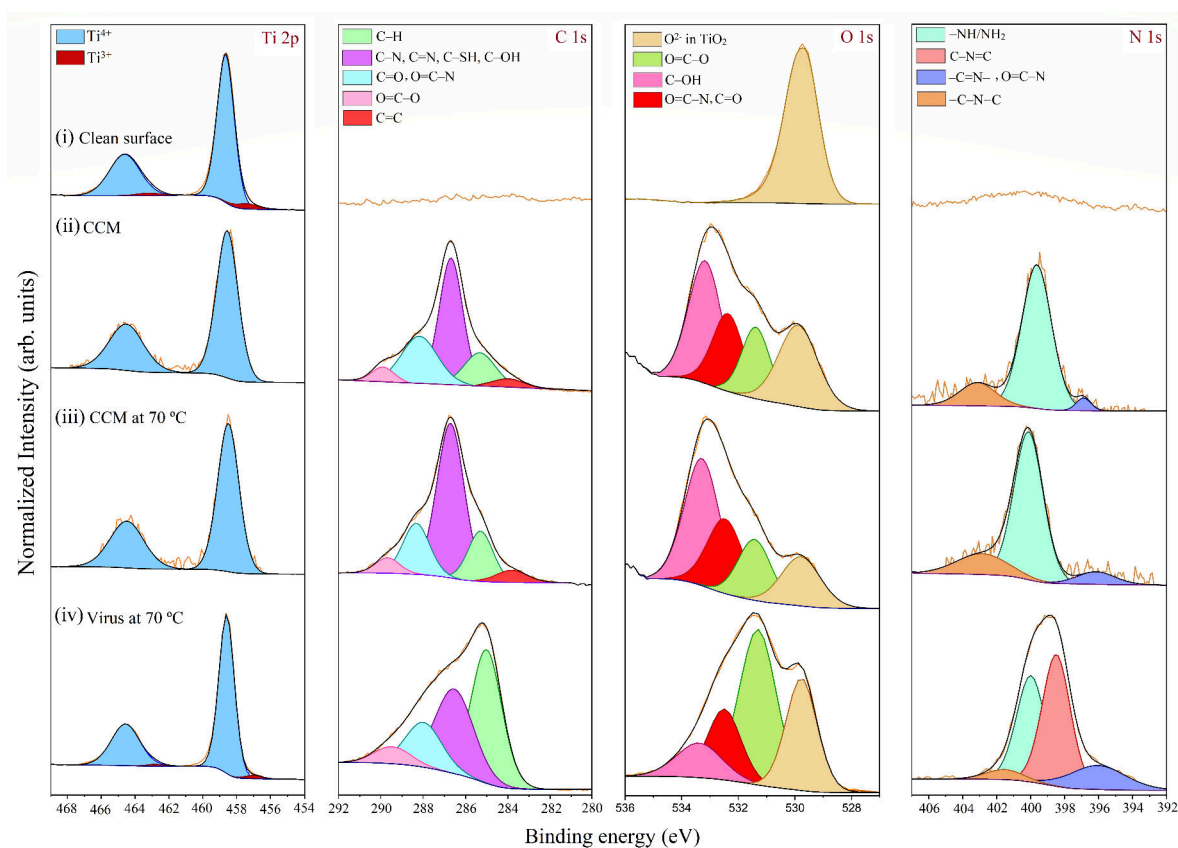


Figure 5: Core-level photoelectron spectra of Ti 2p (a), C 1s (b), O 1s (c), and N 1s (d) for clean surface of TiO<sub>2</sub>(101) (i), adsorbed CCM before (ii) and after (iii) thermal treatment, and adsorbed SARS-CoV-2 after heating at 70 °C (iv). SARS-CoV-2 adsorbed on TiO<sub>2</sub>(101) was measured at P22 beamline with a photon energy of 3.4 keV, while measurements for the adsorbed CCM before and after thermal treatment were performed at a photon energy 1.4 keV.

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The actual interaction of the virus with the surface can be confirmed by the C 1s simulated results, shown in Figure 6 b-e, where three broad components can be resolved, matching the experimental ones (represented by green, magenta and cyano sticks, respectively). The two features at higher binding energy (BE) are shifted by  $\sim 1.5$  and  $2.8$  eV with respect to the secondary C (Figure 4 a) in the side chain of asn (that is computed to be the least bound), against an experimental shift of 1.55 and 2.95 eV. The peak at the lowest BE, besides the secondary C atoms (C-H), includes also the C $\alpha$  of both cys (for all configurations considered) and asn. This would provide a rationalization for the high intensity of such peak, as observed in the experimental XPS (see Table 1 in the supporting information (S2)). At intermediate BEs, our models suggest the contribution of the C atoms in the carboxylic group, which is however strongly dependent on the adsorption configuration. This agrees with the broad experimental feature at 286.5 eV. Finally, at the highest BEs we register the contribution of the C atoms in the side chains of the amino acids, either the C bound to the thiol for cys or the C in the amide group for asn. The position of this third feature is in worse agreement with experiments, which is probably due to the contribution of other amino acids in the spike protein not considered in the computational model.

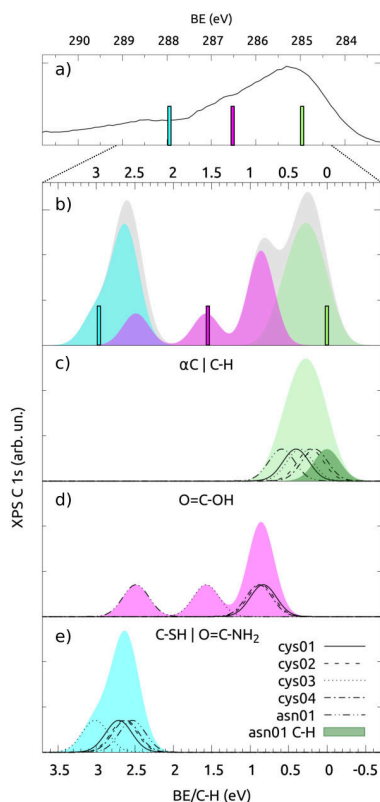


Figure 6: (a) Experimental C 1s XPS spectrum of the virus only, after CCM subtraction. The vertical sticks report the position of the fitted components shown in Figure 5 b, using the same color code. (b) Simulated C 1s XPS spectrum obtained as the overlap of all C species in all models. The simulated spectrum is aligned with the experimental one by matching the position of the component with the lowest BE to that of the fitted C-H feature and the dotted lines highlight the zoom in the energy range. (b), (c) and (d) report the separated contributions along with the individual components originated by inequivalent C atoms in various models considered. The BEs are reported with respect to the calculated energy of the secondary C in asn (C-H).

The O 1s core level peak of CCM was de-convoluted into four peaks (Figure 5 c). These peaks were observed at BEs 529.8 eV, 531.4 eV, 532.3, and 533.1 eV, which are assigned to the lattice oxygen in TiO<sub>2</sub>, O=C-O, C=O/O=C-N (peptide or amide bond) and C-OH species, respectively.<sup>57,59</sup> All these peaks were also observed in the O 1s core level spectrum of the virus and CCM samples after thermal treatment. The changes observed in the peaks ratio in O 1s spectrum of the virus sample after thermal treatment are related to the presence of proteins and lipid structures in the virus. The N 1s core level spectrum after adsorption of CCM (Figure 5 d) can be fit with three components at binding energies of 396.8 eV, 399.6



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3 eV, and 402.7 eV that correspond to the imine group ( $-C=N-$  and  $O=C-N$ ),<sup>60</sup>  $-NH_2/-NH$   
4 and  $-C-N-C$ ,<sup>61</sup> respectively. Due to the presence of glutamine and non-essential amino  
5 acids in CCM, the amine and imine components were observed in the N 1s spectrum of  
6 CCM. In addition, dulbecco's modified eagle's medium (DMEM) in CCM contains four  
7 fold concentrations of amino acids and vitamins. Some of these vitamins contains imine  
8 group for which the corresponding peak was observed in the N 1s spectra. After thermal  
9 treatment, the  $-C=N-$  peak was slightly changed and broadened which could be related  
10 to the reaction between amino acids and carbonyl groups resulting in formation of imines  
11 through elimination of  $H_2O$  under thermal treatment. Similar behavior was detected for  
12 the  $-C=N-$  peak in the virus sample. In fact under thermal treatment, denatured proteins  
13 lose their 3D structure and can show conformational changes. Moreover, through inter-chain  
14 cross linking between  $-NH_2$  and  $-COOH$  groups in the virus proteins new compounds can  
15 be formed, which can change the C 1s, O 1s and N 1s peak ratios and shape of virus sample  
16 compared to the CCM.  
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### Photocatalytic inactivation of SARS-CoV-2 adsorbed on the surface of TiO<sub>2</sub>(101)

UV (200-280 nm) irradiation is a potentially effective disinfection method that is widely used for disinfection of surfaces and is able to deactivate different types of viruses including SARS-CoV-2.<sup>62</sup> Under UV irradiation, RNA absorbs the UV photon which blocks the transcription via the formation of pyrimidine dimers, while absorption of UV radiation by amino acids disturbs the activity and structural function of the proteins.<sup>63</sup> However, the disinfection mechanism of SARS-CoV-2 under UV irradiation is not clear and requires further investigation. Recently, Lo et al.<sup>64</sup> reported the effect of UV-C (253.7 nm) irradiation on inactivation of SARS-CoV-2 in the liquid phase. TEM and protein damage testing results revealed that UV-C can deactivate the virus particles while their viral morphology remains intact.<sup>64</sup> Immunoblotting results of viral S- and N-proteins confirmed that protein degradation might not be the main reason for the UV-induced inactivation of the virus.<sup>64</sup> UV light with wavelengths ranging from 220-280 nm can damage the viral genome since this wavelength is close to the absorption wavelength of the virus nucleic acid.<sup>65</sup>

To study the effect of UV light on the virus adsorbed on the surface of TiO<sub>2</sub>(101), the sample was exposed to UV light for 30 min (with a fixed distance between the sample and UV lamp of 10 cm) and then characterized by AFM and TEM out of the BSL-3 laboratory.<sup>66</sup> A Plaque assay test was performed in order to investigate the number of infectious viruses present after UV treatment. Contrary to the dark condition, where the virus particles remained active, complete virus inactivation was achieved after 30 min of UV irradiation.

The AFM images of the UV-treated samples (Figure 7 a) shows spherical particles with a size of around 80-120 nm (shown in the line profile in Figure 7 b) corresponding to the adsorbed SARS-CoV-2 particles on the surface of TiO<sub>2</sub>(101). Some particles did not change in size and are intact with spike, confirming that one of the inactivation mechanisms under UV irradiation could be RNA damage without an obvious effect on viral structural proteins. Other particles with smaller size (20-30 nm) are related to the viral proteins membrane which dissociated from the virus structure under UV light and readsorbed on the surface of

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TiO<sub>2</sub>(101) (Figure 7 a, b). Figure 7 c displays the TEM images of SARS-CoV-2 adsorbed on the surface of TiO<sub>2</sub>(101) after UV treatment. These results reveal that the virus morphology changed after UV treatment and the virus diameter was enlarged in line with the AFM results (see the labeled region with a yellow rectangle in Figure 7 a). In addition, surface S-proteins were not detected in the TEM results which confirmed that virus particles lost S-proteins following adsorption on the surface of TiO<sub>2</sub> and subsequent UV inactivation. Our results suggest that, the TiO<sub>2</sub>(101)/UV photocatalytic system inactivated SARS-CoV-2 by viral genome damage and viral proteins degradation.

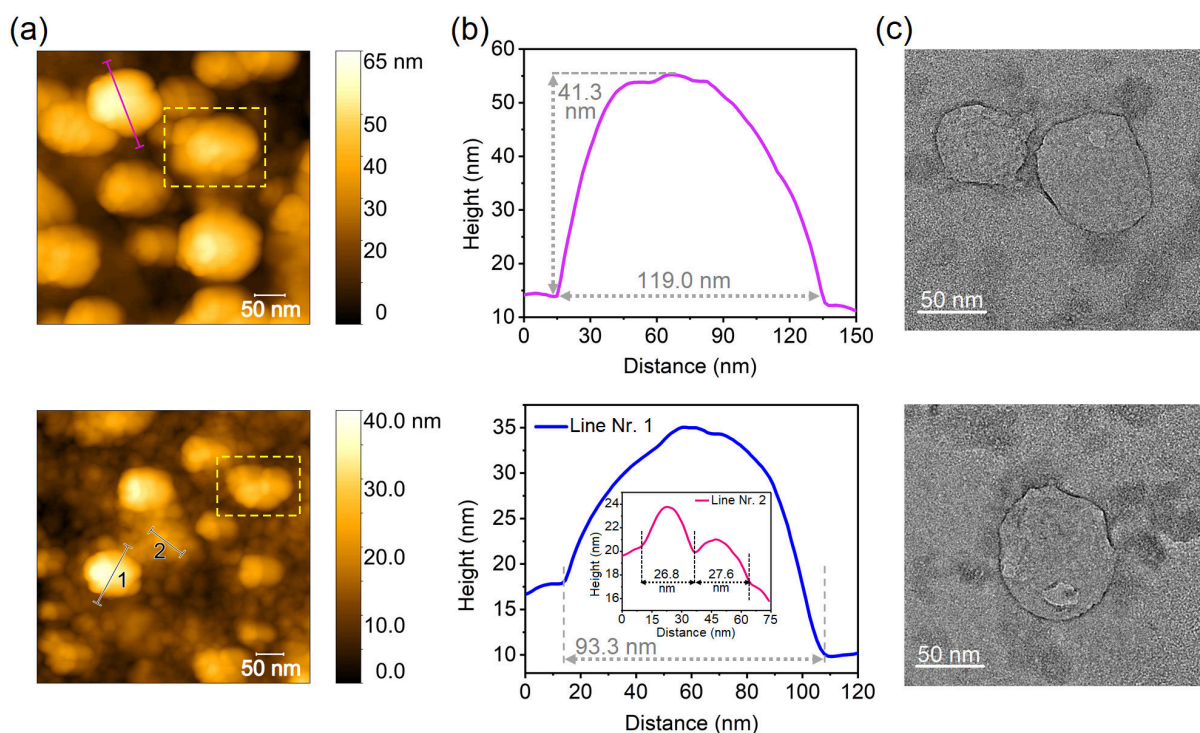


Figure 7: (a) AFM ( $0.5 \times 0.5 \mu\text{m}^2$ ) images of SARS-CoV-2 adsorbed on TiO<sub>2</sub>(101) after UV treatment and (b) line scan profile measured along the particles in the AFM images. (c) TEM images of SARS-CoV-2 adsorbed on TiO<sub>2</sub>(101) after UV treatment.

### The role of Pd nanoparticles on SARS-CoV-2 adsorption on Pd/TiO<sub>2</sub>(101)

To study the role of TiO<sub>2</sub> supported NPs on the adsorption behavior of SARS-CoV-2, Pd NPs were grown on the surface of TiO<sub>2</sub>(101) using evaporation of Pd in UHV via physical

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3 vapor deposition (see Methods section). Scanning tunneling microscopy (STM) (Figure 8 a)  
4 showed that the width of the Pd NPs is 3-4 nm (Figure 8 b). SARS-CoV-2 virus was loaded  
5 on the surface of the Pd/TiO<sub>2</sub>(101) and then inactivated by heating at 70 °C. AFM image of  
6 SARS-CoV-2 adsorbed on Pd/TiO<sub>2</sub>(101) exhibits a high number of particles with an average  
7 size around 80-120 nm that are related to the virus (Figure 8 c). Furthermore, larger particles  
8 were observed in the AFM image (Figure 8 d), which come from the interaction of the virus  
9 with Pd NPs resulting in the formation of complex structures. Similar behavior was reported  
10 for the Ag/TiO<sub>2</sub> catalyst, in which the interaction of the spike protein with Ag atoms and  
11 the subsequent formation of a SARS-CoV-2/Ag/TiO<sub>2</sub> complex enhanced the inactivation of  
12 the virus.<sup>42</sup> Wang et al. found that increasing the amount of Ag NPs on TiO<sub>2</sub> enhances the  
13 adsorption of SARS-CoV-2 in the liquid phase, although they did not study the surface of the  
14 solid catalyst and corresponding adsorption behavior of the virus.<sup>42</sup> Based on our observation  
15 and literature reports, we propose that the cause of higher adsorption and interaction of the  
16 virus with the Pd/TiO<sub>2</sub>(101) surface could be due to the binding of spike proteins with Pd  
17 NPs which enhances the adsorption efficiency compared to the bare TiO<sub>2</sub>. Virus proteins  
18 contain carboxylic acid (-COOH), hydroxyl (-OH), amine (-NH<sub>2</sub>), and carbonyl (C=O)  
19 functional groups.<sup>41</sup> Based on the surface charge of the substrate these functional groups  
20 can interact with the surface and enhance the adsorption ability of samples.<sup>67-69</sup> Cysteine  
21 (C<sub>3</sub>H<sub>7</sub>NO<sub>2</sub>S) and asparagine (C<sub>4</sub>H<sub>8</sub>N<sub>2</sub>O<sub>3</sub>) are two abundant amino acids in the S-protein  
22 structure<sup>42</sup> with terminating carboxylic groups which are responsible for the connection of  
23 the virus to the surface of TiO<sub>2</sub>. Moreover, the thiol group in the amino acid of the S-protein  
24 can interact with Pd NPs and generate complex structures (palladium cysteine thiolate). The  
25 interaction between the thiol group of the protein with the surface of Ag<sup>70</sup> and Au<sup>71</sup> metal  
26 NPs were observed earlier. So, the role of different functional groups in S-protein amino  
27 acids is important to understand the adsorption mechanism of virus on the surface of TiO<sub>2</sub>.  
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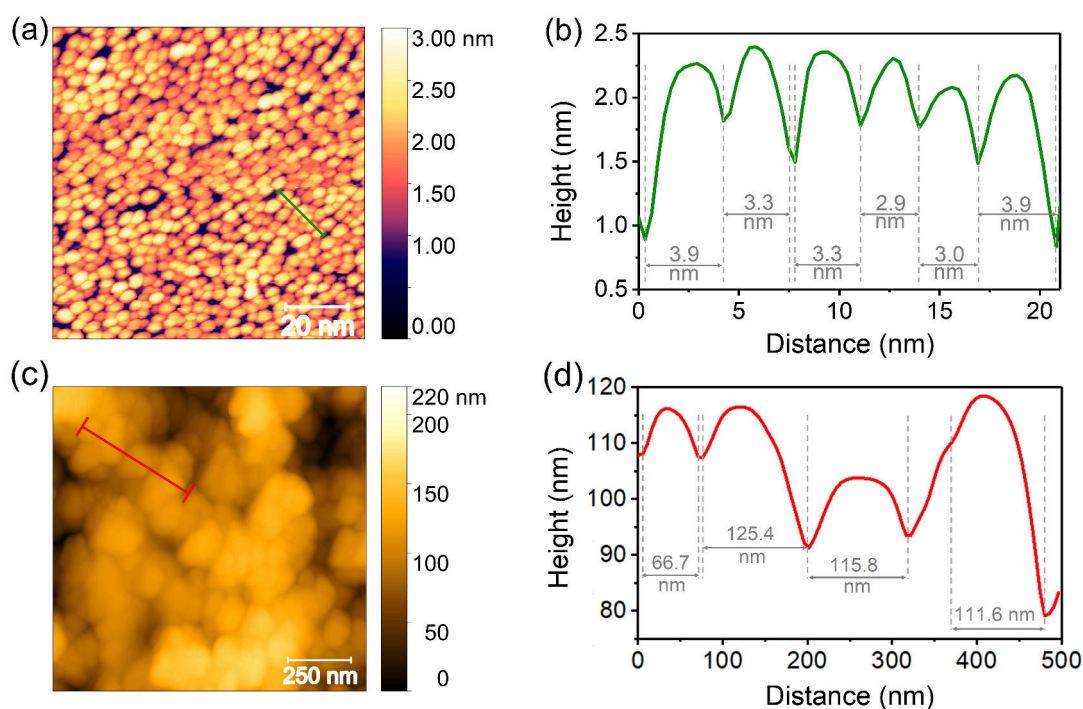


Figure 8: (a) STM image ( $0.1 \times 0.1 \mu\text{m}^2$ ) of Pd/TiO<sub>2</sub>(101) at room temperature (Tunneling parameters: 1.5 V, 0.2 nA). (b) Line scan profile of marked particles in the STM image. (c) AFM ( $1.2 \times 1.2 \mu\text{m}^2$ ) image of SARS-CoV-2 adsorbed on the surface of Pd/TiO<sub>2</sub>(101) catalyst after thermal treatment, and (d) line scan profile along the red line in the AFM image.

### Proposed mechanism of adsorption and inactivation of SARS-CoV-2 on TiO<sub>2</sub>(101)

The main step of virus adsorption on the surface of TiO<sub>2</sub> is related to the interaction of S-protein with the oxide surface, due to the high content of amino acids. Nitrogen and sulfur in cysteine and asparagine, with their paired electrons, can interact with Ti atoms which act as Lewis acid sites. Moreover, the surface oxygen atoms are Brønsted basic sites, and act as hydrogen (–H) bond acceptor.<sup>49</sup> So, functional groups such as –NH<sub>2</sub>, –COOH, –SH, and –OH in the S-protein structure groups can interact with surface Ti and oxygen sites and enhance the adsorption of the virus particles. The charge of viral particles is dependent on the solution pH. The isoelectric point (IEP) of the virus is lower than 7.0.<sup>39</sup> Owing to the variety of proteins in virus structure and the large size of the virus, both negative and positive charges of surface functional groups exist in the virus structure between the range of pH=5.0-8.0 that

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3 can enhance the adsorption efficiency of viruses on various surfaces.<sup>41,72</sup> Our results suggest  
4 that interaction of different functional groups in the virus proteins structure are responsible  
5 for adsorption of SARS-CoV-2 on the surface of TiO<sub>2</sub>. During the thermal treatment virus  
6 proteins separate from the virus structure and adsorb on the surface of TiO<sub>2</sub>(101). This  
7 changes of the virus structure under thermal treatment leads to the inactivation of virus on  
8 the surface. Three mechanisms can be responsible for SARS-CoV-2 inactivation under UV  
9 light irradiation on the surface of TiO<sub>2</sub>(101): i) viral genome damage via absorption of UV  
10 irradiation by nucleic acid, ii) direct denaturation of viral proteins under UV light, and iii)  
11 photocatalytic oxidation of the viral particles. Utilizing TiO<sub>2</sub> as a photocatalyst enhances the  
12 disinfection efficiency of the surface owing to the oxidation/reduction of organic compounds  
13 in the viral proteins. Figure 9 shows the proposed mechanism of adsorption and inactivation  
14 of SARS-CoV-2 at the surface of TiO<sub>2</sub>(101).  
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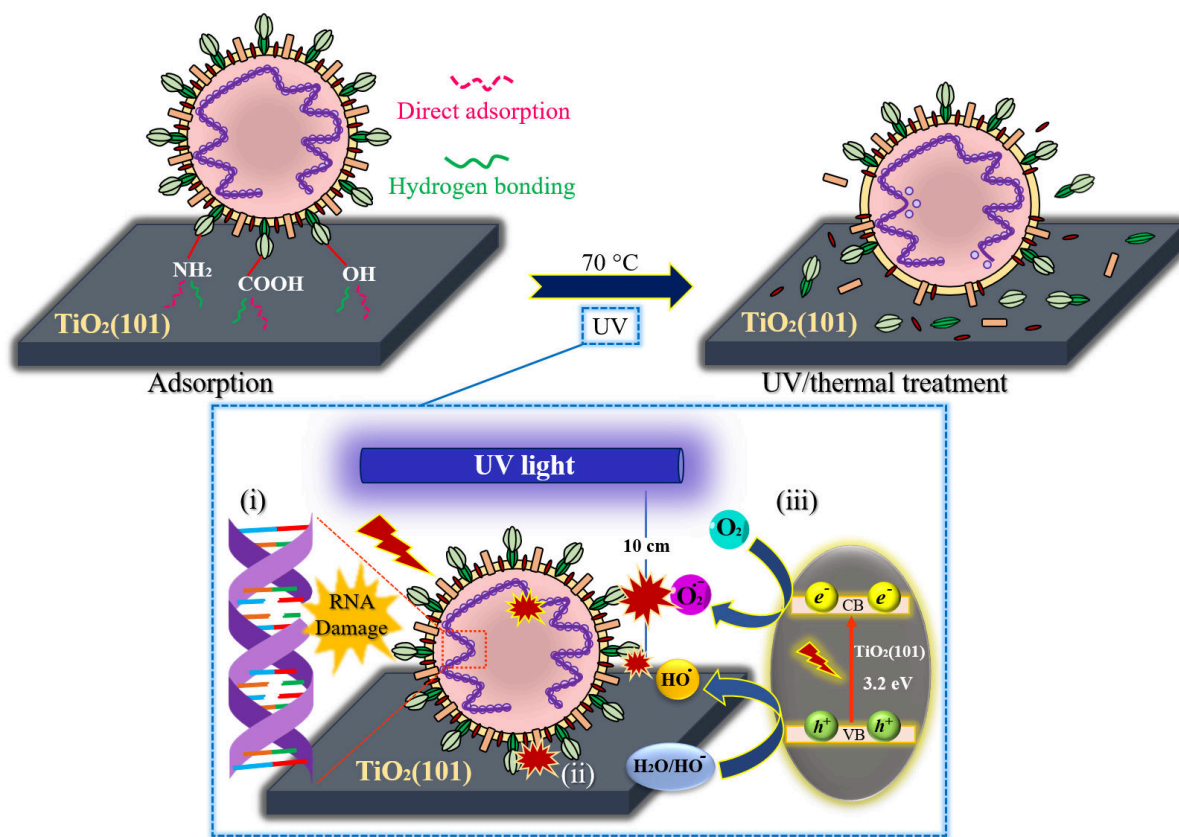


Figure 9: Proposed mechanism of SARS-CoV-2 adsorption, and the effect of thermal and UV treatments at the surface of  $\text{TiO}_2(101)$ . (i), (ii) and (iii) refer to the RNA damage, direct denaturation of S-protein and photocatalytic oxidation mechanism, respectively.

## Conclusion

SARS-CoV-2 was adsorbed on the surface of the single crystalline anatase  $\text{TiO}_2(101)$  and was studied after inactivation under thermal (at  $70\text{ }^\circ\text{C}$ ), UV and ethanol treatments. Different microscopy techniques such as AFM, TEM and Fluorescence microscopy were used to investigate the adsorption, interaction and morphology of the virus/ $\text{TiO}_2(101)$ . Based on the AFM and TEM results, SARS-CoV-2 showed different interaction mechanisms under thermal, UV and ethanol treatments on the surface of titanium dioxide. In fact, following heat treatment of SARS-CoV-2 adsorbed on single crystal anatase surface, the virus proteins denature and dissociate from the virus structure, and subsequently re-adsorb on the catalyst surface. Thermal treatment changes the virus structure via the dissociation of proteins, while

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3 ethanol dissolves the lipid bilayer resulting in the inactivation of the virus. Additionally, we  
4 performed XPS at the Ti 2p, C 1s, N 1s and O 1s core levels, which we combined with the  
5 theoretical XPS characterization of different models at the DFT level. The virus proteins  
6 lost their activity via close contact with the surface of TiO<sub>2</sub> and interaction of –NH<sub>2</sub> and  
7 –COOH groups at the surface of TiO<sub>2</sub>, which was observed in the XP spectra. Moreover, the  
8 adsorbed virus showed aggregation behavior on the surface of the Pd/TiO<sub>2</sub>(101) catalyst.  
9 Interaction of functional groups of amino acids of the spike protein played vital role in the  
10 adhesion process, which highlighted the role of Pd NPs on virus adsorption. Plaque assay  
11 results demonstrated that SARS-CoV-2 particles adsorbed on the surface of TiO<sub>2</sub>(101) were  
12 efficiently inactivated by UV light within 30 min. The AFM and TEM results confirmed  
13 that UV irradiation has an effect on the adsorbed viral particles on the surface of TiO<sub>2</sub>(101)  
14 and changes the virus morphology, due to photocatalytic oxidation at the surface of TiO<sub>2</sub>.  
15 The information obtained in this study will be vital for the development of air and surface  
16 cleaning materials from oxide catalysts/photocatalysts that are urgently required during the  
17 ongoing pandemic.  
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## 35 Experimental

### 36 Materials

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39 The SARS-CoV-2 stock was supplied from Technical University of Munich (TUM). Sodium  
40 Pyruvate and fetal bovine serum (FBS) were supplied from Thermo Fischer Scientific GmbH  
41 (Dreieich, Germany). Glutamine, bovine serum albumin, Dulbecco's phosphate buffered  
42 saline (PBS), Bovine serum Albumin (BSA), and saponin were purchased from Sigma Aldrich.  
43 Non-essential amino acids (NEAA) and Dulbecco's Modified Eagle's Medium (DMEM) were  
44 purchased from Lonza (Lonza, Walkersville, MD, USA) and from Gibco (New York, USA).  
45 Penicillin/Streptomycin was obtained from Gibco (New York, USA).  
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## Preparation of SARS-CoV-2

The SARS-CoV-2 isolate hCoV-19/Germany/BAV-PL-virotum-nacq/2020 (GISAID accession ID: EPI\_ISL\_582134) was derived from a nasopharyngeal swap of a 2 years old boy. The SARS-CoV-2 stock was propagated in the biosafety level 3 (BSL3) laboratory at the institute of Virology, TUM. VeroE6 cells at a confluency of about 60% were inoculated with 1.0 mL of the isolated virus and 9.0 mL of DMEM containing 5.0% FBS, L-Glutamine (2.0 mM), Penicillin /Streptomycin (50 U mL<sup>-1</sup>), 1.0% non-essential amino acids, and sodium pyruvate (1.0 mM), at 37.0 °C. 24- and 28-hours post-inoculation, the supernatant was harvested and centrifuged for 10 min at 300xg and stored at -80 °C.

## Preparation of Pd/TiO<sub>2</sub>(101) catalyst

The anatase TiO<sub>2</sub>(101) single crystal (5 mm x 5 mm x 2 mm, Surface net Ltd.) was prepared by repeated 1 keV Ar<sup>+</sup> ion bombardment and 850 K anneal cycles in a backpressure of 1 × 10<sup>-6</sup> mbar O<sub>2</sub>, until XPS showed the surface to be free of contamination. The Pd nanoparticles were deposited at room temperature on the surface of vacuum prepared titanium dioxide (as outlined above) by means of Pd metal vapor deposition in an UHV chamber, with a base pressure of 1 × 10<sup>-10</sup> mbar.

## Adsorption and inactivation of SARS-CoV-2 on the surface of TiO<sub>2</sub>(101)

To study the adsorption behavior of SARS-CoV-2, 10.0 μL SARS-CoV-2 stock containing 10.000 plaque forming units (PFU) (10<sup>6</sup> PFU/mL) was loaded on the center of catalyst crystal structure so that the whole catalyst surface was covered. To inactivate the adsorbed virus on the surface of TiO<sub>2</sub>(101) and Pd/TiO<sub>2</sub>(101) by thermal inactivation method, the samples were placed in an incubator and heated to 70 °C for 30 min. A similar method was applied for coating the CCM (without SARS-CoV-2) on the surface of TiO<sub>2</sub>(101). To compare the effect of different inactivation methods on adsorbed virus structure, ethanol treatment was

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3 also investigated on adsorbed SARS-CoV-2 particles. Additionally, UV irradiation (Wave-  
4 length: 265 nm; 30 min irradiation time; 10 cm distance to the light source) was investigated  
5 as an inactivation method. For the control experiment, the system was kept in the dark for  
6 30 min. All experiments were performed in a biosafety level (BSL)-3 laboratory. Plaque  
7 assay was performed as described in Stukalov, A. et al. 2021.<sup>73</sup> to determine remaining viral  
8 titers in irradiated samples and check for successful inactivation of infectious virus particles.  
9 After UV treatment, no infectious virus was detected contrary to the dark condition. By  
10 using these methods, we were able to transfer the sample to lower safety laboratories for the  
11 characterization.  
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## 23 **Characterization methods**

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25 **Fluorescence microscopy** Fluorescence microscopy was used to prove binding of  
26 SARS-CoV viral particles to TiO<sub>2</sub> crystals via detection with rabbit anti-SARS-CoV-2 nu-  
27 cleocapsid primary antibodies and anti-rabbit secondary antibodies conjugated to the flu-  
28 orescent dye Alexa568. Since the SARS-CoV-2 adsorbed on TiO<sub>2</sub>(101) was heated at 70  
29 °C as heating treatment, the CCM on TiO<sub>2</sub>(101) was prepared as reference sample under  
30 the same condition. The surface of TiO<sub>2</sub>(101) was covered by CCM and the sample was  
31 dried in an incubator at 70 °C. To reduce non-specific antibody binding, the sample was  
32 covered with blocking solution (PBS supplemented with Saponin (0.25%) and Bovine serum  
33 Albumin (BSA, 3.0%). After 30 min, the sample was washed three times with PBS. Then,  
34 20 µL of blocking solution containing 1:200 diluted primary antibody (anti - SARS-CoV-2  
35 nucleocapsid antibody produced in rabbit, Thermo Fisher Scientific), was added on the top  
36 of the sample. After incubation time (1 h), the sample was washed with PBS to remove  
37 unbound antibody and then blocking solution containing 1:500 fluorescent-dye conjugated  
38 secondary antibody (anti-rabbit antibody conjugated with Alexa568, Thermo Fisher Sci-  
39 entific) was added. After an incubation (30 min) and washing (three times by PBS), the  
40 samples were examined with a fluorescence microscope (Leica DMi8 equipped with a 20x  
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3 NA 0.4 air objective, a Leica DFC9000 GT sCMOS camera, and a Lumencore Sola SE FISH  
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5 365 LED light source).  
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9 **Atomic force microscopy (AFM)** AFM was used for investigating the interaction  
10 of SARS-CoV-2 with the surface of the samples. AFM imaging was performed using a CP-II  
11 instrument from Digital Instruments at DESY Nanolab.<sup>66</sup> The AFM images were recorded  
12 under intermittent (tapping) mode, with different scan sizes ( $2.5 \times 2.5 \mu\text{m}^2$  and  $5.0 \times 5.0$   
13  $\mu\text{m}^2$ ) and 0.996 Hz scan rate.  
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20 **Transmission electron microscopy (TEM)** To investigate the morphology of ad-  
21 sorbed virus on  $\text{TiO}_2$  by TEM, some coating of three substrates that inactivated by ethanol,  
22 thermal and UV treatments was scrapped with a blade. Then, the blades were rinsed with  
23 ethanol and were loaded on the carbon grids. The TEM images of these samples were  
24 recorded on a FEI Talos F200X (typo Thermo Fisher Scientific, USA) at 200 kV.  
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32 **X-ray photoelectron spectroscopy** HAXPES data were collected at beamline P22  
33 at PETRA III at the Deutsches Electron Synchrotron DESY<sup>74</sup> in Hamburg, Germany. A  
34 photon energy of 3.4 keV was used for all experiments, with the energy selected using a Si  
35 (111) double crystal monochromator and a Si (220) post-monochromator. All measurements  
36 were conducted in grazing incidence geometry ( $9^\circ$ ). A Phoibos 225HV analyzer (SPECS,  
37 Berlin, Germany) was used with the small area lens mode and a slit size of 3 mm. Spectra  
38 were collected using a pass energy of 50 eV, and a total energy resolution of about 300 meV.  
39 The XPS measurements for adsorbed CCM on  $\text{TiO}_2(101)$  were carried out using XPS system  
40 in the DESY Nanolab,<sup>66</sup> at the Centre for X-ray and Nano Science, DESY. The X-ray source  
41 employed was Al  $K\alpha$  at 1.4 keV and a Phoibos 150 hemispherical energy analyzer with a  
42 base pressure of  $1.2 \times 10^{-10}$  mbar. The core-level spectra for the CCM adsorbed on  $\text{TiO}_2$   
43 are normalized to the intensity of Ti  $2p_{3/2}$  peaks of clean surface of  $\text{TiO}_2$  while the core-level  
44 spectra for the SARS-CoV-2 adsorbed on  $\text{TiO}_2$  are normalized to the corresponding Ti  $2p_{3/2}$   
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3 peaks. All XPS spectra were fitted by CasaXPS software,<sup>75</sup> while the Shirley background  
4 and Gaussian curves were utilized.  
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9 **Scanning tunneling microscopy (STM)** The STM measurements were performed  
10 in a Scienta Omicron VT SPM with a base pressure of  $4 \times 10^{-11}$  mbar in constant current  
11 mode using a tungsten tip at DESY Nanolab.<sup>66</sup> STM and AFM images processing was done  
12 using the Gwyddion software package.<sup>76</sup>  
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## 16 17 18 **Computational methods** 19

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21 All the calculations have been performed within the density functional theory framework,  
22 through the QUANTUM Espresso suite,<sup>77-79</sup> including the PBE exchange-correlation func-  
23 tional with a plane-waves basis set size of 52 and 575 Ry for the ground state wave function  
24 and charge density, respectively. The TiO<sub>2</sub> anatase (101) surface has been modeled through  
25 a slab made of three triatomic-layers with a 4×1 supercell including at least 11 Å along the  
26 non-periodic *z* direction. The lowermost layer has been kept fixed to the positions optimized  
27 in the bulk structure (with lattice parameters  $a=3.790$  Å and  $c=10.325$  Å) while the others  
28 have been left free to relax. The wave function has been expanded onto a shifted 2×2×1 grid  
29 of points in reciprocal space, generated by the Monkhorst-Pack algorithm. Dispersion inter-  
30 actions have been accounted for by the inclusion of an additional charge-dependent term to  
31 the forces through the Grimme-D3 correction.<sup>80</sup> The XP spectra have been simulated at the  
32 C K-edge through the ΔSCF method,<sup>81</sup> for which an additional pseudopotential, generated  
33 with a missing 1s electron in the core states, is placed at every inequivalent core-excited  
34 atom in the supercell. Such a pseudopotential requires an enlarged plane-waves basis set,  
35 up to 74 and 575 Ry. In addition, since we relied on pseudopotentials, we couldn't directly  
36 calculate the absolute BE but rather its relative change energy with respect to a selected  
37 reference, i.e. the core level shift (CLS). To compare the CLS calculated for inequivalent  
38 atoms in different models, we included in each supercell a common reference, namely an  
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3 ethylene ( $C_2H_4$ ) molecule, in each supercell at more than 15 Å from the surface, minimizing  
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5 its interaction with the slab. A similar approach has been successfully applied in previous  
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7 works.<sup>82–85</sup> Finally, the simulated XP spectra have been constructed through the convolution  
8  
9 of the calculated CLS with Gaussian profiles, with broadening  $\sigma=0.17$  eV.  
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## 11 12 13 14 **Acknowledgement**

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16  
17 This study was supported by the initiative and networking fund of the Helmholtz Associa-  
18  
19 tion of German Research Centers under the CORAERO Project (KA1-Co-06). The authors  
20  
21 acknowledge DESY (Hamburg, Germany), a member of the Helmholtz Association HGF, for  
22  
23 the provision of experimental facilities. Parts of this research were carried out at PETRA  
24  
25 III using beamline P22. Funding for the HAXPES instrument by the Federal Ministry of  
26  
27 Education and Research (BMBF) under framework program ErUM is gratefully acknowl-  
28  
29 edged. MK and HN thank Wolfgang Hammerschmidt for helpful discussion and information.  
30  
31 CDV acknowledges support from the European Research Council (ERC) under the Euro-  
32  
33 pean Union’s HORIZON2020 research and innovation programme (ERC Grant Agreement  
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35 No [647020]).  
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## 41 42 **Supporting Information Available**

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44 The following files are available free of charge.

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47 • S1: Fluorescence images of CCM (a), and SARS-CoV-2 (b) adsorbed on  $TiO_2(101)$   
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49 after thermal treatment.  
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- 51  
52 • S2: X-ray absorption at C 1s edge binding energies (BE): comparison between experi-  
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54 mental and calculated values. The latter are reported as differences from an arbitrary  
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56 reference, which in this case has been set to the component with the lowest BE.  
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