

High-Resolution Correlative Microscopy Approach for Nanobio Interface Studies of Nanoparticle-Induced Lung Epithelial Cell Damage

Rok Podlipec,* Luka Pirker, Ana Krišelj, Gregor Hlawacek, Alessandra Gianoncelli, and Primož Pelicon



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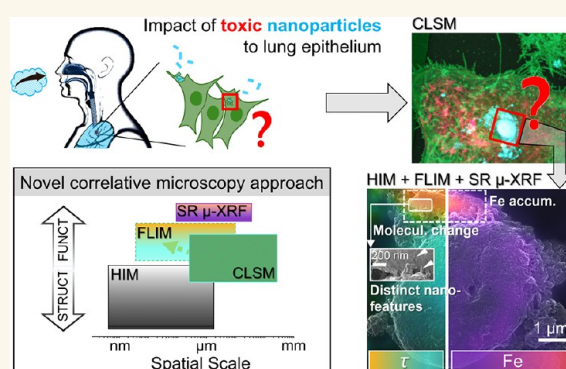
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ABSTRACT: Correlated light and electron microscopy (CLEM) has become essential in life sciences due to advancements in imaging resolution, sensitivity, and sample preservation. In nanotoxicology—specifically, studying the health effects of particulate matter exposure—CLEM can enable molecular-level structural as well as functional analysis of nanoparticle interactions with lung tissue, which is key for the understanding of modes of action. In our study, we implement an integrated high-resolution fluorescence lifetime imaging microscopy (FLIM) and hyperspectral fluorescence imaging (fHSI), scanning electron microscopy (SEM), ultrahigh resolution helium ion microscopy (HIM) and synchrotron micro X-ray fluorescence (SR μ XRF), to characterize the nanobio interface and to better elucidate the modes of action of lung epithelial cells response to known inflammatory titanium dioxide nanotubes (TiO₂ NTs). Morpho-functional assessment uncovered several mechanisms associated with extensive DNA, essential minerals, and iron accumulation, cellular surface immobilization, and the localized formation of fibrous structures, all confirming immunomodulatory responses. These findings advance our understanding of the early cellular processes leading to inflammation development after lung epithelium exposure to these high-aspect-ratio nanoparticles. Our high-resolution experimental approach, exploiting light, ion, and electron sources, provides a robust framework for future research into nanoparticle toxicity and its impact on human health.

KEYWORDS: nanobio interface, TiO₂ nanotubes, lung epithelium inflammation, correlated light and electron microscopy, synchrotron micro X-ray fluorescence, helium ion microscopy



INTRODUCTION

Among the novel microscopy approaches, correlated light and electron microscopy (CLEM) has attracted considerable interest in the life sciences community, particularly following recent developments in the spatial resolution and sensitivity capabilities of individual techniques.^{1,2} Advanced live cell imaging, with its targeted labeling of individual cellular components, can provide molecular specificity and the observation of dynamic changes.^{3,4} In addition, the advanced imaging of fixed cells, when performed in high vacuum, can provide ultrastructural information with nanometer spatial resolution⁵ and chemical information with submicron resolution.^{6,7} When successfully combined in the same region of interest (ROI), a correlative microscopy approach offers several significant advantages. It provides comprehensive functional and structural insights at multiple scales, increased accuracy through

cross-validation of data, and improved contextual understanding through the analysis of dynamic and static processes on live and fixed cells, respectively.

One of the highly relevant scientific areas that can be addressed by correlative microscopy is the field of nanotoxicology, which studies the environmental and, in particular, health effects of the ubiquitous particulate matter in the polluted air.⁸ In particular, ambient ultrafine particles (UFPs) and engineered nanomaterials (NMs) less than 100 nm in size can

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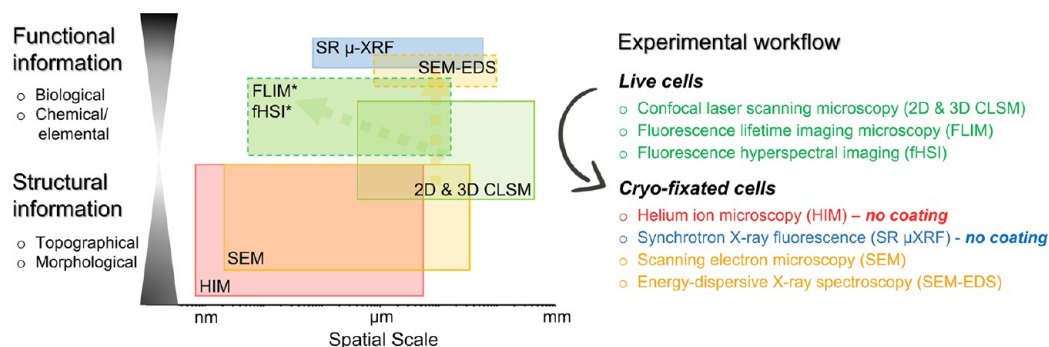


Figure 1. Novel multimodal microspectroscopy-based correlative microscopy enables a comprehensive investigation of the studied nanobio interface across a broad spatial scale, from tens of micrometers down to the nanometer level. This approach provides both functional and structural insights, which in our study focus on the interface between lung epithelial cells and exposed NMs. The colored areas illustrate the typical imaging ranges for each technique, highlighting the type and extent of information obtained. * The spatial resolution extends beyond the optical limit, as FLIM and fHSI can probe molecular interactions at a subdiffraction scale.

penetrate deep into the lungs and induce a variety of chronic diseases, including inflammation, fibrosis, and cancer.^{9–11} For safety assessment and prediction of nanoparticle-induced adverse outcomes, an understanding of the mode of action at the molecular level, starting from the moment the particles interact with the lung tissue, is critical. To this end, several innovative correlative microscopy approaches have been introduced in recent years to elucidate the interaction at the nanobio interface and the subsequent impact of nanomaterials and/or elements on biological matter.^{12–36} A commonly used approach is CLEM¹² which mostly combines highly specific and targeted fluorescence microscopy (FM) with transmission electron microscopy (TEM).^{13–15} More recently, super-resolution fluorescence imaging combined with TEM and scanning electron microscopy (SEM) has been implemented to address the complex biological questions at the nanoscale.¹⁶ While CLEM remains a prominent approach, recent years have seen the development of several complementary techniques. These correlate FM with X-ray tomography,¹⁷ dark-field microscopy (DFM),¹⁸ atomic force microscopy (AFM),^{19,20} and ion-induced luminescence.²¹ Additionally, to enhance and support CLEM, a few ion-based methods have also been introduced.^{22,23} For more detailed and quantitative analysis of the specific molecules and elements present in biological tissues, correlative microscopy approaches often include secondary ion mass spectrometry (SIMS) combined with electron microscopy,^{24–26} and novel high-resolution helium ion microscopy (HIM).^{27–29} In addition to SIMS, proton-induced X-ray emission (PIXE) and X-ray fluorescence (XRF) analytical techniques are also used to quantify chemical elements in biological samples, and are combined with FM³⁰ and EM,^{31,32} respectively. Other correlative approaches include the combination of XRF with AFM³³ or XRF with ion beam techniques,^{34–36} for elemental quantification and complementary information.

Despite the continuous development of new correlative microscopy approaches in nanomaterial toxicity research, the full potential of multimodalities and their combinations has not yet been fully exploited, with many opportunities and challenges.³⁷ In this study, we present a novel correlative microscopy pipeline that can investigate the modes of action and toxic effects of UFPs and NMs, both of which pose the greatest risk to the lungs, as they can penetrate deep into the alveolar region and potentially further into the bloodstream.³⁸ We are particularly focused on the worldwide and one of the most

widely used industrial NMs, titanium dioxide (TiO₂), which in the form of high surface area nanotubes (NTs) induces lung inflammation^{39,40} and is likely to be carcinogenic for humans.⁴¹ We demonstrate the utility of an innovative multimodal and multiscale experimental workflow that integrates several complementary imaging and spectroscopy techniques on live and fixed cells under a high vacuum (Figure 1). In order to preserve the nanoscale morphology and properties of the investigated nanobio interfaces on the surfaces, we use a rapid cryofixation technique without chemical cross-linking, which could otherwise compromise sample size, shape, or cell membranes at the investigated spatial scale.^{42–46} We must acknowledge that the development of sample preparation and preservation for high-vacuum, high-resolution correlative microscopy is one of the major challenges in the field and can play a key role in influencing the physicochemical properties of samples and the subsequent interpretation of results.^{7,33,34,37,47}

The present study aims to cover a wide range of detectable spatial scales (down to the nanometer level) and gain insight into both the functional and structural properties of the samples under investigation. In order to obtain comprehensive knowledge, data gathered from multimodal fluorescence microspectroscopy performed on living cells were integrated with data obtained from high-resolution and high elemental sensitivity X-ray, ion, and electron-based techniques performed in high vacuum. These approaches provided new insights into the physicochemical properties of cell-excreted and cell-deposited TiO₂-biological matter (TiO₂-bio) composites aggregated and immobilized on the surface of the lung epithelium. In addition, we gained deeper morpho-functional insights into the inflammatory and anti-inflammatory mechanisms triggered by the binding and transport of DNA and other molecules to TiO₂ NTs. We also uncovered the possible mechanism of initiation of an acute cellular response via the formation of a fibrous network over the nanoparticles. These findings contribute to a deeper understanding of the current knowledge in the field of UFPs and NMs-induced toxicity on the lung epithelium.

RESULTS AND DISCUSSION

Impact of TiO₂ NTs on DNA Binding and the Transport in Apoptotic Lung Epithelial Cells. To gain new insights into the response of lung epithelial cells to exposed metal oxide NMs, we first performed an integrated multimodal high-resolution fluorescence imaging on live cells exposed to TiO₂ NTs for 2

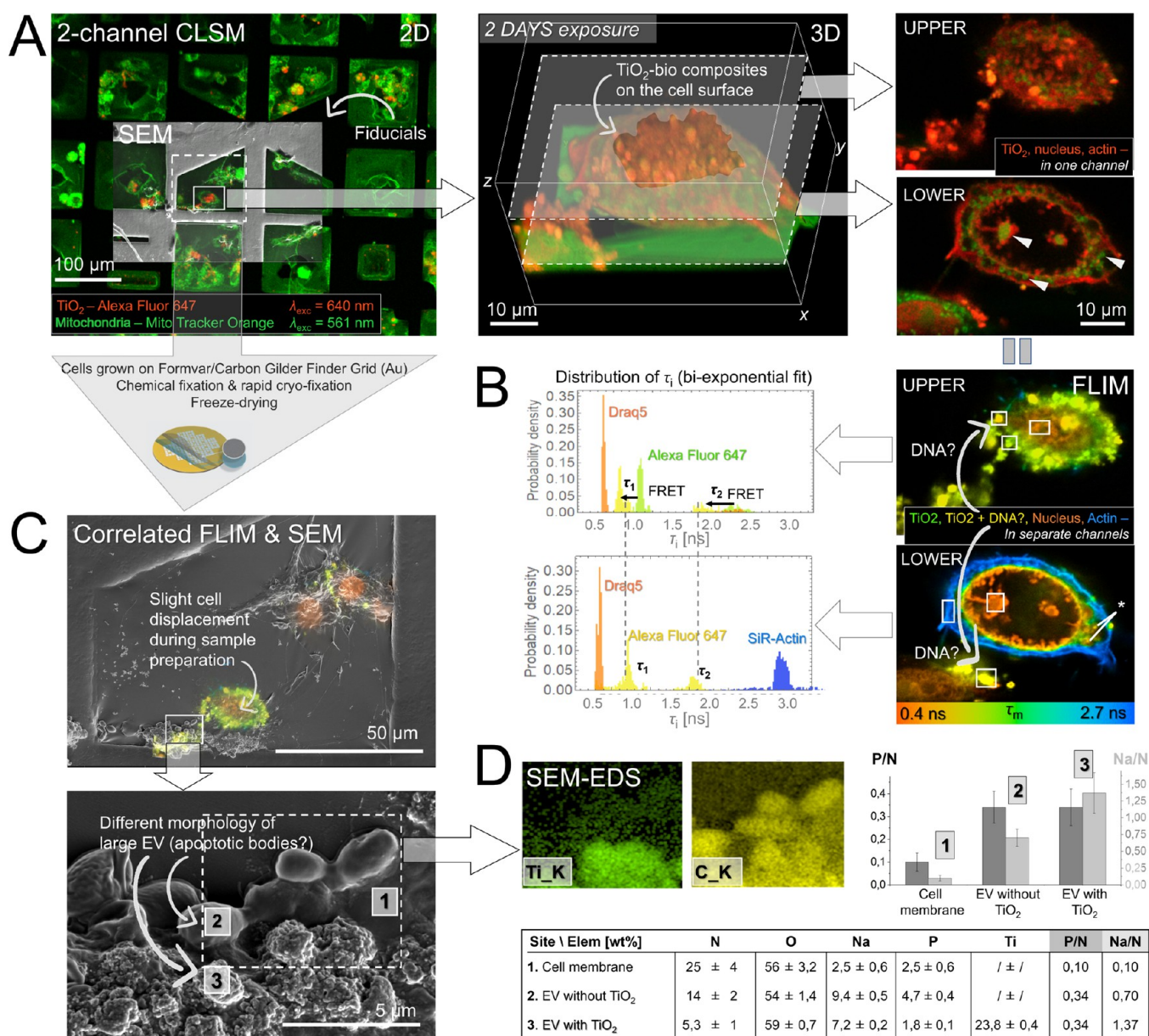


Figure 2. Correlative multimodal fluorescence microscopy and SEM-EDS (multimodal CLEM) applied on lung epithelial LA4 cells exposed to TiO₂ NTs for 2 days reveal physicochemical properties of the nanobio interface and the insights of DNA binding and transport during plausible cell apoptosis. (A) Low to high magnification 2D/3D CLSM showing TiO₂-bio composites on the cell surface and the morphology of the cell nuclei, actin and TiO₂ (in red) and mitochondria (in green). (B) The same site with correlated FLIM capable of precisely distinguishing between individual labeled structures (color-coded from red to blue), using a single laser source ($\lambda = 640$ nm). Shifted distribution of τ_1 and τ_2 from biexponential fit (green to yellow in the chart on the left) indicate on local FRET between TiO₂ and DNA dyes, revealing plausible binding of DNA to TiO₂ surface. (C) Correlated FLIM and SEM after a successful registration and overlap of fiducials shown in (A). High magnification images reveal large EVs with different morphologies, denoted with 2 and 3. (D) Chemical characterization of the same structures performed by SEM-EDS confirms not only high accumulation of biological matter (C) on nanoparticles' surface (Ti) but also a few times higher content of organic phosphate normalized to biological matter (P/N) (right chart and table), indicating a high but mechanistically different DNA load inside large EVs.

days with a meaningful surface dose ($3 \mu\text{g}/\text{cm}^2$), as discussed in [Supplementary Comment #1](#) (Figure 2A,B). Cells were grown on special 10 nm thin Finder grid Formvar/Carbon substrates, with fiducials used for correlative microscopy (Figure 2A, arrow). Prior to measurement, the observed cell mitochondria, actin network, and nuclei were stained to characterize any potential structural and/or functional changes that may have occurred as a result of nanoparticle exposure. To track and determine potential colocalization between stained cellular structures and nanoparticles, TiO₂ NTs were fluorescently

labeled prior to administration according to the well-established protocol.⁴⁸ To be able to successfully register and overlap the correlated images by identifying the focal plane, we first applied low-magnification imaging (10 $\times$) using two-channel confocal laser scanning microscopy (CLSM). In addition to clearly visualizing the grid, used as a fiducial (Figure 1A, in black), it was possible to distinguish between TiO₂ (in red) and mitochondria (in green). To delve deeper into the interaction between TiO₂ and cells, we performed high-resolution imaging in three dimensions (3D) at 60 $\times$ magnification (Figure 1A, second

image). The 3D image shows the localization and aggregation of TiO₂ on the cell surface (white arrow), a phenomenon observed and described in our recent study.³⁹ Here, we further unravel the high heterogeneity of the composites in morphology and size (upper image plane (UPPER)). However, the similar excitation and emission profiles of the fluorescent dyes prevented a clear differentiation between the TiO₂, nucleus, and actin structures beneath the cell membrane (lower image plane (LOWER)), using a 2-channel CLSM.

To identify these individual structures, we proceeded with fluorescence lifetime imaging microscopy (FLIM), integrated into the same experimental setup. FLIM, a powerful imaging technique for measuring the lifetime of fluorescent molecules independent of signal intensity,³ also provides quantitative insights into the properties and dynamic changes of the surrounding molecular environment⁴⁹—an advantage we later utilized. With FLIM, we were able to identify the individual labeled structures (Figure 2B), which was not possible with the 2-channel CLSM. The fluorescence lifetime, as measured in each image pixel, is color coded with the mean lifetime (τ_m), which is the weighted average of the individual lifetime components (τ_i) from the biexponential fit (see eq 1). The FLIM analysis, performed after using a single excitation laser ($\lambda = 640$ nm), allowed to distinguish between actin structures (shown in blue), a nucleus undergoing a distinct phase separation due to specific biophysical and/or biochemical processes⁵⁰ (shown in orange), and TiO₂ with a wide τ_m distribution (shown in green to yellow). The latter indicates a high degree of structural and molecular variability and most likely local attachment of DNA molecules and/or their fragments to the surface of nanoparticle composites (see the regions indicated by the arrow). The proximity of the Draq5 dye, intercalated in the DNA structure at a distance of less than 10 nm, to the Alexa Fluor 647 dye attached to TiO₂ plausibly induces a fluorescence resonance energy transfer (FRET) transition, which is known to shorten the fluorescence decay time.⁵¹ This is evidenced by the blue shift in both the τ_1 and τ_2 distributions (Figure 2B, top left image). We also observed the Draq5 signal locally concentrated in the cell cytoplasm (see the asterisk). With the additional experiments taking advantage of the combined FLIM and fluorescence hyperspectral imaging (fHSI), we further confirmed the local and widespread presence of DNA in the cytoplasm, apparently colocalized with the TiO₂ NTs (Figure S1), suggesting a possible downstream inflammation.⁵² These intracellularly formed nanobio composites may represent one of the key early states in the cell clearance mechanism that evolves into the observed surface-immobilized TiO₂-bio composites.

To gain further insight into the structural and functional properties of the formed composites in relation to DNA binding, we then prepared the samples for a correlative SEM (Figure 2C). Due to the chemical fixation treatment of 4% PFA and 2% GA and the fragile thin Formvar/carbon support films, we observed a slight displacement of the investigated cell and its structures, as indicated by the white arrow. However, this was not a limitation in this study, as the identification of the same extracellular vesicles (EVs) in fixed cells was sufficient for subsequent morphological and chemical characterization (Figure 2C,D). The SEM measurements revealed the presence of distinct morphological features of large EVs that resembled apoptotic bodies (ABs).⁵³ Additional SEM-EDS chemical mapping (Figure 2D) with typical spectra (Figure S2) revealed that rough surface topography was colocalized with the presence of TiO₂ (Ti map in green), while all large EVs exhibited a similar

amount of organic matter (C map in yellow). Given that the electron penetration depth and interaction volume for the detected characteristic X-rays are on the order of a few microns at $E_0 = 15$ keV,⁵⁴ and considering the much thinner measured layer of organic matter in EVs with a high nanoparticle content (Ti 23.8 wt %), we confirmed a concentrated accumulation and extensive binding of organic matter (lipids and proteins) to the TiO₂ surface—consistent with findings from our recent study.⁵⁵ Through detailed chemical analysis of the distinct regions labeled 1–3 (Figure 2C), we confirmed previous FLIM-based observations, suggesting the very likely presence of DNA in EVs (Figure 2D). The weight percentage (wt %) of the elements shown in the table and in the corresponding bar charts shows a clear, more than 3-fold, elevation of phosphate (P) and sodium (Na) inside EVs (sites 2 and 3) compared to the surrounding cell membrane (site 1). The values are normalized to the concentration of organic matter, in our case organic nitrogen (P/N and Na/N), within each measured region to remove the potential effects of different densities and interaction volumes, all of which affect EDS detection. It is noteworthy that there was no discernible difference in P/N between EVs with high and negligible TiO₂ content, indicating a similar concentration of DNA or its fragments in both cell-excreted structures. However, the different morphologies of these large EVs, observed by both high magnification SEM and FM on live cells, suggest completely different functions and modes of action for DNA loading, packaging, and transport, all of which are considered essential for cellular homeostasis and immune response modulation.⁵⁶

The most likely cause of the high P accumulation in flat-surfaced EVs (site 2) is the high concentration of DNA with associated binding proteins in the so-called plasma membrane blebbing process, followed by its release from the cells in the form of apoptotic bodies, which are the largest class of EVs. This has the biological implication of impending apoptosis of the measured cell, with the shape indicating an early stage of this process. However, the initiation of cell apoptosis may well be indicated by the observed chromatin condensation, which is known to be one of the sequences of nuclear changes culminating in cell apoptosis,⁵⁷ and the observed round mitochondria, which are known to occur under conditions that compromise mitochondrial function, such as cell apoptosis⁵⁸ (Figure 2A, see arrows in lower right image). On the other hand, the EVs with a rough surface and a significant accumulation of TiO₂ appear significantly different from the previously observed EVs, casting doubt on whether these structures are truly apoptotic bodies. The most likely cause of the high concentration of organic phosphates (P), further supported by the highest accumulation of mineral Na⁺ among the measured sites, is a physical rather than a biological mechanism, as previously reported.^{59,60} Namely, the DNA phosphate backbone has been found to play the major role in TiO₂ adsorption,⁵⁹ while Na⁺ has been found to preferentially adsorb to the TiO₂ surface as inner-sphere complexes,⁶⁰ both under high ionic strength conditions,⁵⁹ such as our cellular environment. Preferential DNA binding to the TiO₂ surface, being found in both cytoplasmic (Figure S1) and cell surface regions (Figure 2), may have important biological implications for disturbing cellular homeostasis. Interference with transcription or even replication processes, as observed in our recent study,⁴¹ may lead to genotoxic effects.⁶¹ On the other hand, Na⁺ binding, if carried out intracellularly, could disrupt the ion homeostasis, leading to disruption of membrane potential and

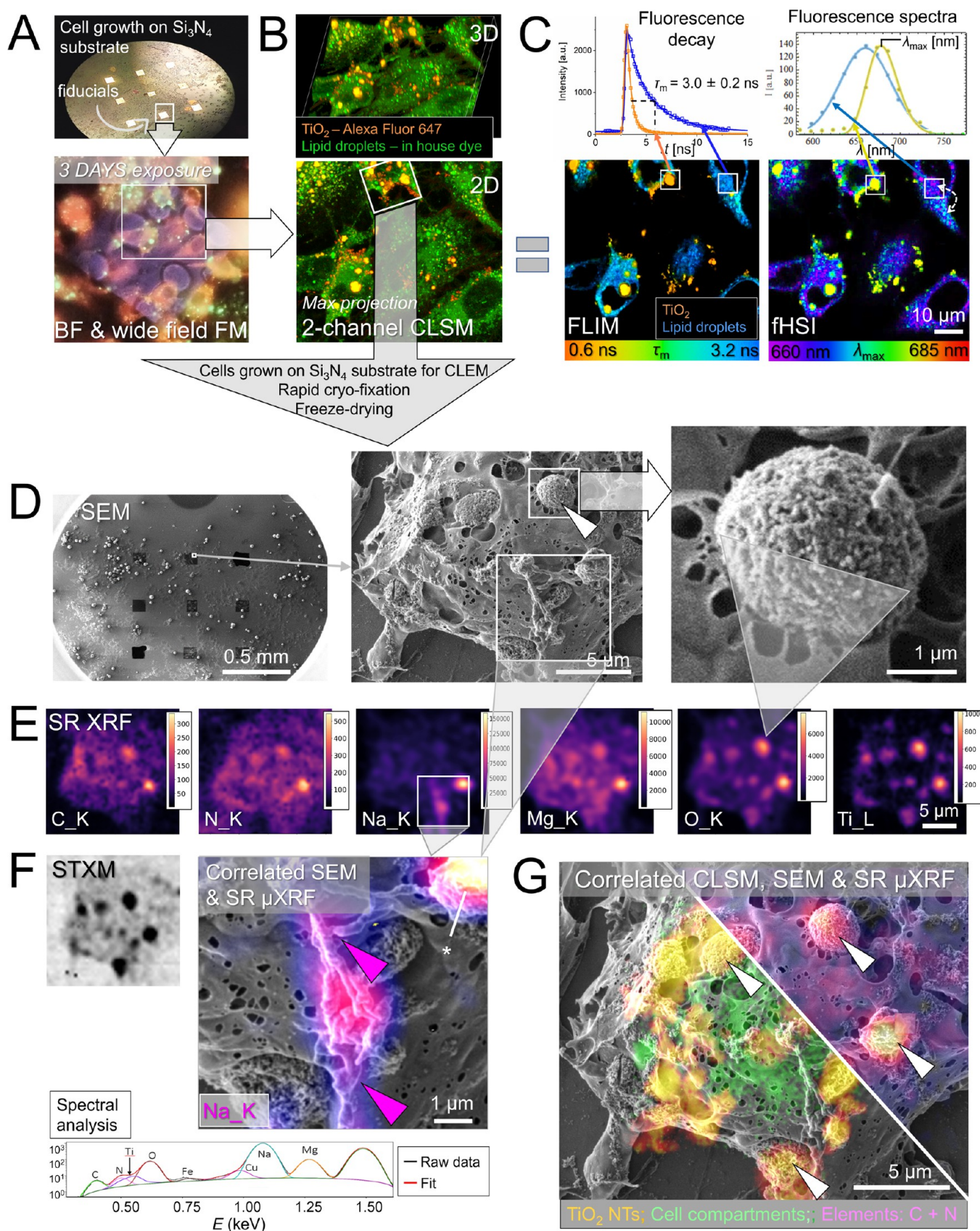


Figure 3. Novel multimodal CLEM combined with synchrotron μXRF (CLEM- μXRF) revealing in-depth physicochemical and morpho-functional properties of inflammatory TiO_2 NPs-rich cell-excreted composites immobilized on the surface of lung epithelial LA4 cells. (A) Combined BF and wide field FM employed to visualize cell growth on 100 nm thick transparent silicon nitride (Si_3N_4) films of the 100×100 μm dimension used as fiducials and optimal substrate for the correlative microscopy both reflection and transmission mode (μXRF). (B) 2D and 3D CLSM performed on the same site to show labeled cells (in green) and cell-excreted TiO_2 composites (in orange) with an improved resolution.

Figure 3. continued

(C) Same site was characterized with additional FLIM and fHSI, capable of distinguishing between individual labeled structures and of inspecting local microenvironment differences, such as molecular conformations or polarity. The upper charts illustrate FLIM decay curves and fluorescence spectra for the two distinct sites. (D) Low to high magnification SEM images of the same site performed after rapid cryo-fixation of the sample to preserve distinct structural features of the TiO₂-bio composites on the cell surface (bottom image). (E) SR μ XRF elemental maps from the same site showing the extensive binding of intrinsic biomolecules (C and N) and essential minerals (Na⁺ and Mg²⁺) on the TiO₂-rich local sites. (F) Left supporting atomic number (Z) and density-sensitive STXM map with the typical XRF spectra collected from the TiO₂-bio composite site (denoted with an asterisk) and its color-coded fits for individual elements. Right – high-magnification correlated Na⁺ map and SEM further highlighting very likely functional relationship between composition and filamentous structure with potential inflammatory implications. (G) Correlated live cell CLSM and high-vacuum SEM and μ XRF revealing morpho-functional properties on the distinct sites (white arrows).

the related processes. Moreover, the immobilization of TiO₂-rich, irregularly shaped EVs, on the cell surface by the intertwined actin fibers from the inside (Figure S3, super-resolution STED microscopy on live cells) and fibrin-like fibers from the outside reported in Figure 5, could affect the normal clearance of apoptotic cells and potentially introduce pathological conditions for inflammation or autoimmunity.

To elaborate on this hypothesis, the observed dense structure of immobilized EVs on the apoptotic cell surface could partially block the secretion pathways of the typical anti-inflammatory signals, thereby disrupting the typical anti-inflammatory environment. Moreover, if apoptotic cells are not cleared efficiently and in a timely manner by macrophages due to their overload in high particulate matter exposure and plausibly reduced motility,⁶² they may progress to secondary necrosis and lose their membrane integrity.⁶³ By releasing immunostimulatory content or so-called damage-associated molecular patterns (DAMPs), the cellular dynamics then promote inflammation and recruitment of immune cells. DAMPs are released through the lesions in the plasma membrane orchestrated by the plasma membrane protein Ninjurin-1 (NINJ1), which is structured into supramolecular filamentous assemblies as recently shown by super-resolution microscopy.⁶⁴ A meaningful observation is that the polymerized filaments range in size from a few tens of nanometers to a few hundred nanometers—closely matching the dimensions of the lesions or holes we found on the cell surface close to the anchored TiO₂-rich excreted structures (Figure 3D). Thus, the holes may not necessarily be entirely a consequence of the incomplete sample preservation during cryofixation and freeze-drying, but may actually represent the plasma membrane lesions. However, further studies with better statistics are needed to confirm this.

Taken together, this provides an additional possible mode of action in the initiation of chronic lung injury, complementing the well-established mechanisms of PM- and TiO₂-induced macrophage phagocytosis dysfunction,^{65–67} and our recently discovered nanomaterial cycling between the lung alveolar surface and macrophages,³⁹ both of which lead to chronic inflammation.

In-Depth Morpho-Functional Characterization of Cell-Excreted and Surface-Immobilized TiO₂-Bio Composites.

To further investigate the morphological and structural properties of TiO₂-bio composites immobilized on the cell surface, as well as to gain further insight into their chemical properties at the submicron scale, we implemented an extended approach, a multimodal CLEM combined with synchrotron μ -XRF (CLEM- μ XRF), where we joined 2-channel CLSM, FLIM and fHSI on living cells with SEM and μ -XRF on cryo-fixed cells (Figure 3). To improve the quality of the imaging capability and sample preservation in our novel correlative microscopy

pipeline, we used special 100 nm thick silicon nitride (Si₃N₄) substrates/windows (Figure 3A) that are transparent to soft X-rays and avoided chemical fixation to preserve the nanoscale features of the investigated nanobio structures (Figure 3D), respectively. In order to improve the resolution and sensitivity of the analysis of particularly light elements, we integrated spatially resolved synchrotron radiation SR μ XRF (Figure 3E,F), with its ability to achieve up to an order of magnitude better, submicron spatial resolution and up to several orders of magnitude better elemental sensitivity⁶⁸ than SEM-EDS, making it a valuable tool for investigating intricate biological processes such as ours.

Lung epithelial cells were grown on a surface-modified Si₃N₄ substrate with fiducials in the form of multiple 100 × 100 μ m transparent windows and exposed to fluorescently labeled TiO₂ NPs for a period of 3 days. We observed a large number of TiO₂-rich structures, measuring up to a few microns in size, as shown in green and orange using a wide-field FM (Figure 3A) and enhanced-resolution CLSM (Figure 3B), respectively. For visualization, the cells were stained with the in-house-synthesized lipid droplets (LDs) dye. Due to partitioning of the dye to other lipid-rich environments during the incubation period, the contrast of the LDs was reduced, as shown by the green color (Figure 3B). With CLSM, in comparison to wide field FM, we were able to better distinguish each of the numerous TiO₂ NT-rich composites, which varied in size and shape.

To gain further insight into the local properties and interactions in the near-molecular environment of the observed TiO₂ NTs-rich structures and LDs, we performed FLIM and fHSI imaging/mapping at the same site (Figure 3C). The typical and easily distinguishable FLIM and fHSI spectra from the two distinct cell compartments within the white rectangles and corresponding fits using a biexponential function (eq 1) and an empirical intensity-normalized log-normal function with nanometer peak position resolution,⁶⁹ respectively, are shown in the top. The fHSI of LDs and lipid-rich regions showed spectral shifts of up to 10 nm, ranging from $\lambda_{\text{max}} = 660$ nm (in purple) to $\lambda_{\text{max}} = 670$ nm (in blue), indicating local microenvironmental differences, such as molecular conformation, polarity, etc.⁷⁰ This was supported by FLIM data, with LDs showing a reduced mean fluorescence lifetime (τ_m) (lighter blue) compared to those of other lipid-rich regions (darker blue), indicating an increased packing density of the dye within LDs. The analysis of TiO₂ NPs-rich structures showed less heterogeneity, but still with a considerable range in τ_m , ranging from 0.6 to 1.0 ns, also indicating differences in the packing density. An increased τ_m may also indicate a greater distance between fluorophores, possibly due to lipid interspacing, as shown previously.³⁹

To probe deeper into the properties and possible consequent effects of the TiO₂ NTs-bio interaction, we introduced to our

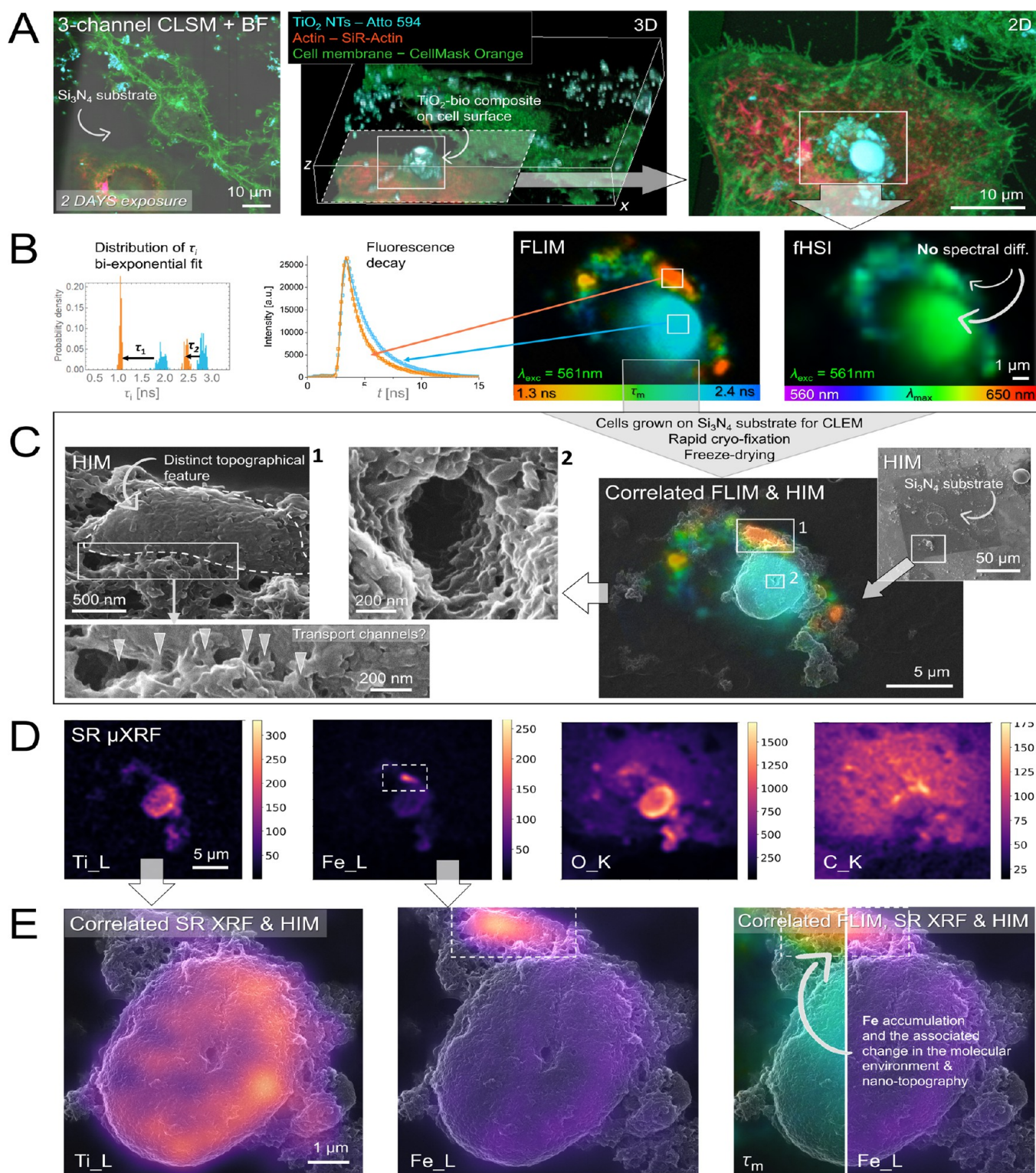


Figure 4. Novel correlated high-resolution FLIM, SR μ XRF and HIM providing in-depth morpho-functional assessment of inflammatory TiO_2 -bio cell-excreted composites immobilized on the surface of lung epithelial LA4 cells. (A) Cells grown on thin Si_3N_4 transparent substrate (in gray, BF image) with labeled cell membranes (in green), actin (in red) and nanoparticles (in light blue) acquired with 3-channel CLSM using 60 $\times$ water immersion objective. 3D CLSM revealing size, shape and topography of a large, few micron-sized TiO_2 -bio composite formed on the cell surface (marked with an arrow). (B) fHSI and FLIM imaging and analysis of the investigated structure uncovering no particular spectral but significant fluorescence lifetime contrasts, respectively (see the distributions and decays on the left). (C) Correlated FLIM and HIM exhibiting colocalization of distinct fluorescence lifetime map (in orange) and nanoscale topographical features shown in the images on the left. Further ultrahigh magnification HIM reveals intriguing morpho-functional features, from the apparent multiple transport channels into the local site with an accumulated Fe (1) (marked with arrows), to a void or tunnel formed in the center of the TiO_2 -bio composite (2), which could facilitate exchange between insight and outside of the cell. (D) SR μ XRF elemental maps of the investigated region showing metals and biomolecules

Figure 4. continued

distributed locally and more evenly, respectively. (E) Changes in the physical properties of the molecular environment at the nanobio interface plausibly caused by the high accumulation of Fe, as shown by the correlated FLIM, SR μ XRF, and HIM.

pipeline an additional correlated SEM and SR μ XRF (Figure 3D–G). The combination of these advanced techniques provided new insights into the morphological and structural features well below the micron scale. The use of a 150 mM ammonium acetate wash solution to remove buffer salt crystals, which otherwise interfere with structural and chemical analysis,⁷¹ followed by rapid cryofixation by plunge freezing and final lyophilization, allowed the preservation of the majority of cell surface structures, including the studied TiO₂ NPs-rich excreted composites studied (Figure 3D). The high magnification SEM image shows an almost perfect spherical structure (indicated by the white arrow), likely resulting from the most efficient packing of nanoparticles with the lowest surface energy-maximizing stability.⁷² To confirm that the structure is composed of nanoparticles, we observed strong colocalization with the fluorescence signal of a TiO₂ dye, as measured by CLSM in live cells (Figure 3B, outlined in a rectangle). This is further illustrated by the overlap of all correlated results from CLEM- μ XRF in Figure 3G (indicated by the arrows). To obtain further biological information on these intriguing structures formed on the cell surface, the same sample was transferred to a synchrotron facility for the elemental mapping using SR μ XRF (Figure 3E), supported by scanning transmission X-ray microscopy (STXM) (Figure 3F) for simultaneous mapping of atomic number by absorption contrast sensing.⁶⁸ The spectral maps of primarily light elements have been generated from the spectral data and their fits, where we show an example from the site marked with an asterisk. The color-coded theoretical spectra for each of the measured elements fit perfectly to the raw data (in black) with a high signal-to-noise ratio (S/N) due to the high detection sensitivity of the technique. The results confirm not only a high TiO₂ content within highly spherical composites, as evidenced by locally elevated concentrations of O (K-shell emission spectra) and Ti (much weaker L-shell emission spectra), but also a higher content of C, N, Mg and, to some extent, Na (all K-shell emission spectra) at the same locations. This again confirms the extensive physical affinity of biomolecules, such as lipids or cholesterol, shown previously,³⁹ and positive ions to the surface of the TiO₂ NTs, as elaborated in the first results section. It further validates and signifies our recent findings performed on the same nanobio system using a combination of omics, in vitro measurements, and full-atom in silico simulations, which reported strong interaction of amine and phosphate groups of the disordered lipids with the TiO₂ NTs surface.³⁹ Extensive biomolecule binding, forming the so-called lipoprotein corona as clearly demonstrated by the ultrahigh resolution imaging (Figure S4), can significantly disrupt cellular homeostasis by altering protein function, affecting cell signaling and trafficking machinery,⁷³ all of which can potentially lead to immune activation and long-term cellular stress. The local increase in Na⁺ adsorption may be due to the high surface/volume characteristic of anatase TiO₂ NTs, allowing Na⁺ to intercalate into the surface and subsurface layer.⁷⁴ The most interesting observation is the deviation of the distribution of plausibly extracellular Na⁺ from the rest of the elemental maps (Figure 3E), indicating at least partially a different binding mechanism. Using correlated μ XRF and SEM, we observe complete colocalization of the highly elevated Na⁺

sites with distinct filamentous structures formed by cells bridging the TiO₂-bio composites (Figure 3F, right image, purple arrows). Good correlation of the Na⁺ map with distinct physical properties not only highlights a fundamental link between composition and structure but also suggests a potential functional relationship. The preferential binding is likely due to the high concentration of negatively charged biomolecules within the filamentous structure. Among these, heparan sulfate (HS) chains stand out, as they possess the highest negative charge density of any known biological macromolecule and are commonly found on cell surfaces.⁷⁵ These molecules also play a central role as mediators in inflammatory processes,⁷⁶ suggesting local inflammatory activity on TiO₂-bio composites bridged with these filamentous structures. In addition, the sequestration of both intrinsic and/or extracellular Na⁺ and Mg²⁺ may result in alteration of ion homeostasis, disruption of normal associated cellular functions, and induction of oxidative stress.

To further elucidate the intricate morpho-functional properties of the studied nanobio interface at the nanoscale, we correlated multimodal CLSM with SR μ XRF and ultrahigh resolution helium ion microscopy (HIM), performed for the first time (Figure 4).

By targeted fluorescence labeling prior to live cell imaging of lung epithelial LA4 cells, which had been exposed to TiO₂ NTs for a period of 2 days, we were able to perform a detailed characterization of highly specific nanobio structures formed on the cell surface (Figure 4A,B). Again, a large, condensed nanobio composite, up to a few microns in size, with an approximately spherical shape, was observed and imaged in detail in the multichannel 2D and 3D (Figure 4A). Moreover, we performed additional FLIM and fHSI imaging (Figure 4B), which allowed us to investigate the potential heterogeneity of the molecular environment in the immediate vicinity of the reporting fluorophores attached to the surfaces of TiO₂ NTs. While fHSI showed no significant spectral variation across the composites, indicating an absence of any particular local molecular and physical changes in the immediate nanometer vicinity, FLIM, revealed significant deviations in the fluorescence lifetimes (τ_m) at the same locations, as indicated by the rectangles and corresponding fluorescence decay curves and distributions of τ_i (Figure 4B, left). This may be indicative of different degrees of dye aggregation/self-quenching⁷⁷ due to the different surrounding molecular environment. However, a more in-depth investigation using an expanded set of complementary techniques is required to better understand the cause.

Therefore, the sample grown on Si₃N₄ substrate was cryofixed immediately after fHSI/FLIM imaging and transferred to successive correlated high-vacuum SR μ XRF and ultrahigh-resolution HIM for the most accurate chemical and morphological analysis, respectively. A detailed examination of the surface and chemical speciation provided new insights that now better explain the differences observed in the local molecular environment in living cells (Figure 4C–E). Using extremely sensitive, submicrometer resolution SR μ XRF (Figure 4D), we revealed the presence of localized, highly concentrated iron (Fe) (indicated by the dashed rectangle), as opposed to more widely distributed titanium (Ti), and, in particular, intrinsic carbon (C)

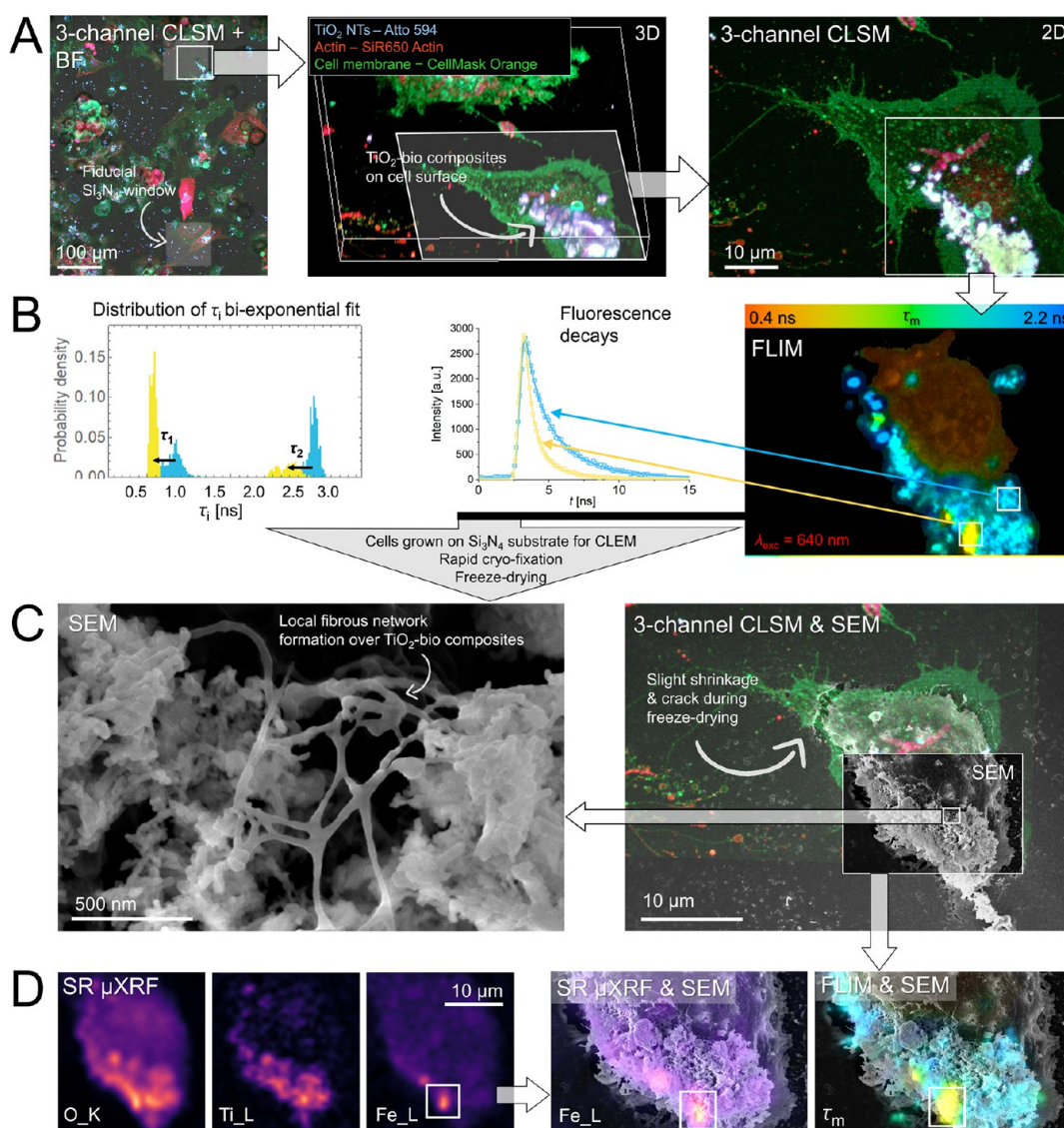


Figure 5. Novel high-resolution CLEM- μ XRF confirming that changes in the physical properties of the surrounding biological environment are induced by Fe and uncovering the plausible inflammatory formation of a fibrous network that resembles the fibrin matrix formed over TiO₂-bio composites immobilized on the plasma membrane of lung epithelial LA-4 cells. (A) Cells grown on thin Si₃N₄ substrates (gray) with labeled cell membranes (green), actin (red), and nanoparticles (blue) acquired using 3-channel CLSM with 10 $\times$ and 60 $\times$ objective magnification. High-magnification 2D and 3D CLSM reveals the size, shape, and topography of a large TiO₂-bio composite, measuring over ten microns, arrested on the surface (marked with an arrow). (B) Right: FLIM image of the investigated composite revealing significant contrast in fluorescence lifetime between local regions (from yellow to blue); Left: fluorescence decay and τ_1 distributions from the corresponding sites. (C) Right: correlated CLSM and SEM showing slight shrinkage and local cracks near the cell boundary after rapid freeze-drying, while preserving the investigated structures on the cellular surface; Left: high-magnification SEM revealing the local formation and the morphology of a fibrous network over the TiO₂-bio composites on the cellular surface and its inflammatory implications, examined further in Supporting Figures S4 and S5. (D) Left: SR μ XRF elemental maps of oxygen and metals, distributed locally on the investigated cell; Right: Local molecular (physical) changes related to Fe accumulation (via τ_m) on the nanobio interface (regions within rectangles), as measured by correlated FLIM and SR μ XRF and represented on top of the SEM data.

and oxygen (O). Superimposing the data from all three complementary techniques (Figure 4E, on the right) reveals remarkable colocalization of the concentrated Fe (in pink), reduced fluorescence lifetime (in orange), supported by the distinct morphological and topographical features within the same region, as shown by ultrahigh resolution HIM (Figure 4C, upper left image). With our novel correlative microscopy approach, we not only identified a close relationship between FLIM footprint, Fe content, and distinct surface morphology but also gained a better morpho-functional assessment of the

observed nanobio structure to better elucidate possible causes and consequent biological effects, as discussed here.

The highly localized Fe site, accompanied by the numerous distinct morphological features at the outer edge, suggests the presence of possible biological transport channels in the form of tunneling nanotubes (TNTs)⁷⁸ with an average diameter of 50 nm (Figure 4C, lower left image marked with arrows). This implies a biological cause for its accumulation through active cell secretion. This also implies the nanoparticle-induced alteration of iron homeostasis and the subsequent cell-protective mechanism, through the initiated cellular iron efflux, plausibly

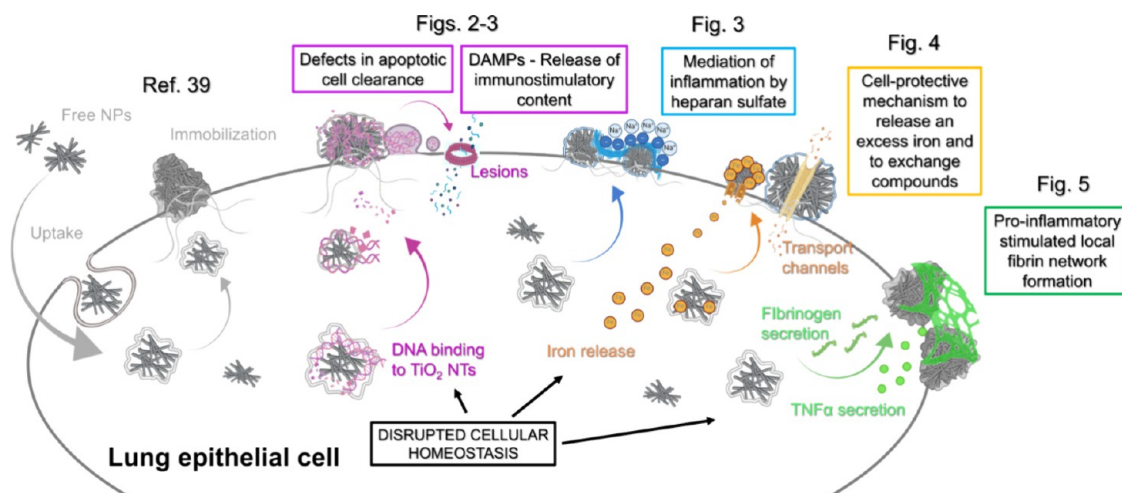


Figure 6. Schematic overview of the possible modes of action of the cell response to the disruptive nature of TiO_2 NTs, toxic to lung epithelial cells, with different inflammatory and anti-inflammatory outcomes (colored from pink to green), observed at the nanobio interface. Created with BioRender.com.

via the major iron transporter ferroportin (Fpn).⁷⁹ This mechanism opts to mitigate iron-induced reactive oxygen species (ROS) and pro-inflammatory cytokine production, as well as further iron-mediated inflammation that can damage DNA, proteins, and lipids.⁸⁰ The remaining question is, why is there an accumulation in the TiO_2 -bio composite? These large TiO_2 -rich extracellular “vesicles” may serve as extracellular containers for other waste molecules, with absorptivity playing an important role. For instance, ferrous iron has been found to adsorb strongly to the surface of metal oxides.⁸¹

To better understand the physical aspects of the interaction, we need to understand the corresponding reduced fluorescence lifetime of the biological molecules present at the nanobio interface. The proximity and/or adhesion of Fe particles to the TiO_2 surface could potentially induce a phenomenon of localized surface plasmon resonance (LSPR)⁸² resulting from the laser-induced collective oscillations of surface electrons in a conduction band,⁸³ leading to a shorter fluorescence lifetime.⁸⁴ Conversely, Fe could interact with nearby lipids and proteins enveloping TiO_2 , catalyzing lipid peroxidation,⁸⁵ inducing lipid bilayer disruption through an increase in water permeability and thus an increase in membrane polarity,⁸⁶ resulting in a shortening of the dye lifetime.

Implementing correlative ultrahigh resolution and large depth-of-field HIM revealed another intriguing feature resembling a ‘wormhole’ at the center of the object (Figure 4C, middle). This void, or tunnel, which is more than 1 μm deep, was formed during the assembly of a TiO_2 -bio composite and could potentially facilitate diffusion and selective matter exchange between the inside and outside the cell. This type of nanobio interface assembly with the revealed nanometer features has never been demonstrated before and could be an interesting subject for further study.

Finally, we have uncovered one of the potentially crucial initiating modes of action in the inflammatory cell response to TiO_2 NTs. Using high-magnification SEM, we observed and characterized a thin fibrous network of biological material that had developed locally over a large TiO_2 -bio composite immobilized on the cell surface (Figure 5C). The organization and the thickness of the fibers, which are a few tens of nanometers in size, are reminiscent of fibrin.^{87,88} Given that fibrin is excessively deposited in the early stages of acute intra-

alveolar lung injury,^{89,90} plausibly accompanied by fibrinogen secretion from lung epithelial cells,⁹¹ the possible presence of fibrin on our studied nanobio interface could have significant biological implications for the local pro-inflammatory stimuli accompanied cytokine secretion, such as tumor necrosis factor ($\text{TNF}\alpha$).⁹² To better elucidate the possible $\text{TNF}\alpha$ secretion-induced fibrin fiber formation, we performed an additional TiO_2 NTs exposure experiment on the coculture of lung epithelial LA4 and murine alveolar macrophages MH-S, which are known to release $\text{TNF}\alpha$ after exposure to metal oxide nanoparticles⁹³ or high aspect ratio nanomaterials⁹⁴ such as ours. We aimed to determine whether the presence of MH-S increased the level of local formation of fibrous structures on TiO_2 NTs. The results revealed a significantly higher presence of fibrous structures on almost all of the TiO_2 NTs that were immobilized on the lung epithelial cell surface in the coculture, compared to the partial—yet still substantial—formation observed in the monoculture, primarily over the larger composites (Figure S5). We also observed extensive apical scavenging and uptake of TiO_2 NTs by rounded MH-S with phagocytic protrusions (Figure S6), capturing and trapping the nanoparticles identified by white image contrast.

To confirm that the fibers consist of inflammation-mediating fibrin molecules, we performed another correlative experiment by introducing labeled fibrinogen into the living cells exposed to TiO_2 NTs. This allowed us to assess potential colocalization with the fibrous structures using the following high-resolution HIM (Figure S7). Despite the much lower resolution of the confocal microscopy images and the slight sample displacement during sample preparation, which prevented the precise registration and overlap with HIM, the results showed good colocalization of the fibrous network with labeled fibrinogen, further supporting our hypothesis. To further explore the potential of long-term inflammation and fibrogenesis in the development of such structures over time, further studies are needed.

In addition to the fibrous structures observed using high magnification SEM and HIM, we provide a correlative characterization of the same TiO_2 -bio composites using a multimodal live cell 3-channel CLSM and FLIM (Figure 5A,B) together with SR μXRF (Figure 5D). FLIM revealed a high degree of heterogeneity in the fluorescence lifetimes of the Atto

594 dye attached to the TiO₂ NTs, as can be seen in the fluorescence decay curves and the distribution of τ_i from the distinct sites marked within the rectangle. The average lifetime, τ_m , calculated from a biexponential fit, was approximately half as short locally (yellow) as in the other parts of the sample (green to blue). Correlated SR μ XRF again showed a colocalization between the short fluorescence lifetimes and sites with high Fe accumulation (Figure 5D), as demonstrated in Figure 4, suggesting a similar underlying mechanism.

CONCLUSIONS

In this study, we highlight the importance of implementing novel extended CLEM approaches that combine multiple modalities in order to elucidate the intricate interactions between biological matter and nanoparticles. In particular, we present different response mechanisms of lung epithelial cells, a subject of persistent exposure to particulate matter, to exemplary high aspect ratio TiO₂ nanotubes with known subacute and even chronic inflammatory inducing effects. The study provides a comprehensive morpho-functional assessment of the nanoparticle-cell interface from the micrometer down to the nanoscale, shedding light on the possible modes of action that initiate lung epithelial toxicity, as summarized schematically in Figure 6. With the novel correlative microscopy approach and in-depth analysis, we advanced the understanding of the recently reported TiO₂-induced mechanism of chronic inflammation³⁹ and genotoxicity⁴¹ by better elucidating several possible mechanisms of the cell response to the disruptive nature of this particular nanoparticle, which can result in different inflammatory and anti-inflammatory outcomes:

1. Nearly spherical TiO₂ NTs-bio composites with extensive DNA binding immobilized on the cell surface, similar to immobile apoptotic bodies, may result in defects in apoptotic cell clearance. This could lead to secondary necrosis and the loss of membrane integrity, resulting in the release of immunostimulatory contents through the observed plasma membrane lesions.
2. The formation of distinct filamentous structures bridging TiO₂-bio composites, perfectly colocalized with high Na⁺ accumulation. This indicates a localized, highly negatively charged surface of plausible heparan sulfate, a key mediator in cell inflammation.
3. Nanoparticle-induced alteration of iron homeostasis and the balance of other TiO₂-adherent cellular compounds/biomolecules, followed by the controlled release of an excess iron or other exchange compounds through transport channels, suggesting a cell-protective mechanism.
4. The localized formation of a fibrin-like fibrous network over TiO₂-bio composites indicates the local pro-inflammatory stimuli, accompanied by plausible secretion of fibrinogen and TNF α cytokines.

By implementing such a multimodal and multiscale correlative microscopy approach for the first time, we were able to study and characterize the initial cellular responses to exposed nanoparticles on such a small scale. These responses may play a critical role in the body's initial immune response and the development of inflammation.

This methodology provides a strong basis for advancing research into the toxicity of nanoparticles and their potential impact on human health. By providing a robust framework, it enables further exploration and refinement of approaches to

toxicity assessment. However, to fully realize their potential and guide future research, it is essential to address certain limitations. Key challenges include the limited high-throughput imaging required for improved statistics, together with faster acquisition to minimize sample displacement, and the need for optimization and better control of all physical variables throughout sample preparation to preserve nanoscale features throughout the sample. Both of these challenges point to better automation, which is discussed in [Supplementary Comment #2](#) in the [Supporting Information Appendix](#). This will lead to critical improvements in the future for more precise 2D/3D visualization, evaluation, and deeper insights into nanobio interactions and the mechanisms behind them, paving the way for more accurate and reliable interpretations of how different nanoparticles and their properties affect relevant biological systems, important for improved safety assessment.

EXPERIMENTAL SECTION

Materials. The murine epithelial lung tissue cell line (LA-4, ATCC CCL-196), murine alveolar lung macrophage cell line (MH-S; ATCC, CRL-2019), F-12K medium (Gibco), fetal bovine serum (ATCC), 1% penicillin-streptomycin (Sigma), 1% nonessential amino acids, L-glutamine, beta-mercaptoethanol (Gibco), phosphate buffer saline (PBS), live cell imaging solution (LCIS, Invitrogen); titanium dioxide nanotubes (TiO₂ NTs) in anatase form synthesized in-house;⁹⁵ μ -Slide 8-well (Ibidi), silicon nitride Si₃N₄ support film (PELCO, 21509CL, Ted Pella), TEM formvar/carbon film on Au Gilder 200 F1 finder grids (FCF200F1-AU-50, EMS), ammonium acetate (Sigma-Aldrich), PTFE coated high precision and ultrafine tweezers (72919-3SATE, EMS), propane transfer system for plunge freezer (3701S, Electron Microscopy Sciences); AlexaFluor 647 (Thermo Fischer Scientific), Atto 594 (ATTO-TEC), CellMask Orange (Invitrogen), SiR-actin (Spirochrome), Draq5 (Invitrogen), MitoTracker Orange CMTMRos (Invitrogen), PSM-39 (in-house).

Cell Culture. The LA-4 murine epithelial lung tissue cells were cultured in 75 cm² TPP cell culture flasks. They were maintained in a controlled environment at 37 °C with 5% CO₂ and humidity using F-12K medium supplemented with 15% fetal bovine serum (FBS), 1% penicillin-streptomycin, 1% nonessential amino acids, and 2 mM L-glutamine. The MH-S alveolar macrophage cells were cultured in RPMI 1640 medium (Gibco), supplemented with 10% FBS, 1% penicillin-streptomycin, 1% nonessential amino acids, 2 mM L-glutamine, and 0.05 mM beta-mercaptoethanol (Gibco). Upon reaching 70–90% confluency at the appropriate passage, the LA-4 cells were seeded onto specially designed imaging holders customized for correlative microscopy. To enable both reflection and transmission imaging modes, thin Silicon Nitride (Si₃N₄) and Formvar/carbon on Au Gilder 200 F1 finder grid support films were used. For coculture experiments, LA-4 and MH-S were grown in separate dishes up to 70–90% confluency and mixed together in a ratio of LA-4:MH-S, 40:1. After 24 h incubation, the cocultures were exposed to nanoparticles for the desired amount of time prior to microscopy. The growth medium in the coculture was an equal mixture of F-12K and RPMI 1640.

Nanoparticles. TiO₂ nanoparticles have been selected for investigation in this study due to the potential risks to human health associated with inhalation or oral exposure.^{96,97} In case of their high specific surface area, recent studies have demonstrated the potential for these nanoparticles to elicit an inflammatory response^{39,40} and to possess carcinogenic properties.^{41,98} TiO₂ nanotubes (TiO₂ NTs) in anatase form synthesized in-house with the protocol described in⁹⁵ exhibited a diameter range of 6–11 nm, a mean length of 100–500 nm, and a BET surface area of 150 m² g⁻¹.⁴⁰ In order to facilitate enhanced visualization and monitoring of nanomaterial interactions and their impact on model lung epithelium at the submicron scale, the TiO₂ NTs were functionalized with super-resolution compatible fluorescent probes, AlexaFluor 647 and Atto 594, according to the protocol, avoiding the introduction of artifacts as described in.⁴⁸ Functionalized

and filtered nanoparticle suspension was stored in a 100-fold diluted bicarbonate buffer (5 mOsm L^{-1}) prior to administration to cells.

Sample Preparation for Live-Cell Confocal Fluorescence Microscopy. After reaching 70–90% confluency in culture flasks, LA-4 cells were reseeded directly onto UV-sterilized Si_3N_4 or Formvar/carbon Au finder grid substrates/holders placed within the chamber of a μ -Slide 8-well. The volume of the added cell suspension prepared at a concentration of a few 10^5 cells/mL was $20 \mu\text{L}$ and resulted in approximately 5000 cells per holder with a diameter of 3 mm. Following a 4 h period during which sufficient cell attachment to the substrate had occurred, cell growth media was added to each chamber in order to cover the surface and wet the entire holder, thus preventing any evaporation during the course of the measurements. Once the desired confluency of approximately 50% had been reached (typically within 2 days), the gently sonicated (Bransonic ultrasonic cleaner, Branson 2510 EMT) nanoparticle suspension and carefully rinsed onto the cell monolayer at an average calculated surface dose of 5:1 ($S_{\text{nanomaterial}}:S_{\text{cells}}$) and total $3 \mu\text{g}/\text{cm}^2$. It presents a meaningful concentration according to the lifetime occupational exposure and the safety recommendations, as further discussed in [Supplementary Comment #1](#). The surface dose was achieved by pouring $V = 5 \mu\text{L}$ of a suspension of nanoparticles ($c = 0.5 \text{ mg/mL}$) into the chambers of 1 cm^2 size filled with the cell growth media ($V = 300 \mu\text{L}$). To put into perspective, the surface dose, which is the most important determinant in the nanomaterial toxicity in lungs,⁹⁹ was used to be equivalent to exposure over a period of 45 working days, based on the 8-h time-weighted average occupational exposure limit for TiO_2 ($6.0 \text{ mg}/\text{m}^3 \text{ TiO}_2$) established by Danish Regulations.¹⁰⁰ The cells were exposed to nanoparticles for the desired period of time, which was 1–3 days, prior to live-cell imaging and following cryopreservation for high-vacuum imaging.

Before live-cell imaging, cells were stained with different highly specific fluorescent dyes to visualize different cell compartments, potentially interacting or responding to exposed nanoparticles. Staining was performed according to the manufacturer's recommendations to achieve a sufficient fluorescent signal while preserving cellular functions. SiR Actin dye in concentration $c = 400 \text{ nM}$ was used for 4 h to stain F-actin. Draq 5 dye in a concentration of $c = 5 \mu\text{M}$ was used for 15 min to stain double-stranded DNA. CellMask Orange in 1000 $\times$ diluted stock concentration was used for 10 min to stain the cell plasma membrane. Mitotracker Orange at a concentration of 500 nM was used for 15 min to stain mitochondria. All staining solutions were removed, and the sample was rinsed a few times with LCIS prior to measurements performed on the microscope inside the stage top incubator with a temperature, gas, and humidity control (H301-MINI, Okolab). Before measurements, the implemented holders with the adhered cells were flipped to enable imaging with a small working distance (high numerical aperture) objective. Sterilized polystyrene microspheres of $20 \mu\text{m}$ in size were employed as volume spacers in order to prevent any compression of the sample.

Sample Preparation for High-Vacuum Electron and Ion Microscopy. Immediately after live-cell imaging, the sample was carefully transferred from the measuring μ -Slide 8-well into the 150 mM ammonium acetate washing solution to eliminate buffer salt crystals, which otherwise interfere with both structural and chemical analysis performed with high-vacuum techniques.⁷¹ After a few seconds of immersion in the washing solution three times, an excess of water was removed by carefully blotting the sides of the sample holder with a lint-free paper. Immediately after blotting, samples were rapidly cryofixed without chemical fixation, to preserve the integrity and morphology of cellular structures and interacting nanobio interface on the nanoscale, using plunge freezing in a liquid propane to prevent the formation of crystalline ice¹⁰¹ and stored in a cryo bank until the final drying procedure in a freeze-dryer (Coolsafe 100-9 Pro). Propane was prepared in a liquid-nitrogen-cooled chamber with the plunge freezer transfer system. To carefully handle the fragile specimens, high-precision cryogenic and nonsticking PTFE tweezers were used. In the case of chemical fixation, ammonium acetate washing solution was changed with 4% paraformaldehyde (PFA) and 2% glutaraldehyde (GA) solution. Samples were fixed for 15 min at room temperature before blotting and rapid cryofixation.

Confocal Laser Scanning Microscopy (CLSM). The experiments were conducted on a custom-built inverted microscope (Olympus IX83, Abberior Instruments). Imaging was conducted with an Olympus 10 $\times$ ($\text{NA} = 0.3$) and 60 $\times$ ($\text{NA} = 1.2$) water immersion objective, utilizing two pulsed picosecond lasers in conjunction with a fast-gating system, which was controlled by an FPGA unit. In order to facilitate the simultaneous detection of the fluorescence emitted by different fluorophores, two pulsed diode lasers were employed, with wavelengths of 561 and 640 nm, respectively, and a pulse length of 120 ps and a repetition rate of 80 MHz. Fluorescence was detected in a descanned mode using an additional pinhole to reduce out-of-focus light and avalanche photodiodes (APD, SPCM-AQRH, Excelitas) with a photon detection efficiency (PDE) exceeding 50% across the entire visible spectrum. Fluorescence was collected from each scanning voxel using a dwell time of 10 μs and detected within the spectral windows $\lambda = 580$ –625 nm and $\lambda = 650$ –720 nm, utilizing dichroics and band-pass filters (both Semrock). For a more comprehensive illustration of the optical setup, please refer to the schematics in [Figure S8](#). Supporting high-resolution measurements on live cells was conducted with the integrated customized super-resolution microscopy (STED), using a depletion laser wavelength of 775 nm, a pulse length of 1.2 ns, and a power of 200 mW in the sample plane.

Fluorescence Lifetime Imaging Microscopy (FLIM) and Fluorescence Hyperspectral Imaging (fHSI). The microscope system was upgraded with the 16-channel Multi-Wavelength Photo-Multiplier Detectors (PML-16 GaAsP, Hamamatsu), a multidimensional Time Correlated Single Photon Counting (TCSPC) detection system and a DCC-100 detector controller card (all manufactured by Becker & Hickl), all communicated through fast ps FPGA electronics, enabling the simultaneous detection of signals across up to 16 spectral channels and detection of their corresponding fluorescence lifetime decays.¹⁰² For fluorescence lifetime imaging, the signals collected on the 16 GaAsP detectors that arrived in the same time interval were summed, and the FLIM decay curves obtained in each pixel of the image were analyzed using the SPCImage5.0 software (Becker & Hickl) as done before.^{103,104} The analysis of FLIM decay curves was conducted using a two-component, double-exponential fitting method convoluted with the instrument response function (IRF), which constrains the achievable resolution to its 200 ps fwhm. The pixel exposure time (100 μs) and image binning (3×3) were set to obtain a sufficient peak value of the decay curves of at least 1000 to elastically resolve fitting with a double-exponential model:

$$I(t) = a_1 e^{-t/\tau_1} + a_2 e^{-t/\tau_2} \quad (1)$$

with the lifetime values of the fast and slow decay components, τ_1 and τ_2 , and the corresponding intensity coefficients, a_1 and a_2 . The fitted FLIM decay curves from each pixel were color-coded according to the average fluorescence lifetime $\tau_m = a_1 \tau_1 + a_2 \tau_2$, with the fastest decays shown in red and longer decays shown in blue. Furthermore, fHSI was conducted utilizing 16 spectral channels within the wide spectral window spanning from 560 to 760 nm, as determined by the diffraction grating positioning within the spectrograph (PML-Spec, Becker & Hickl). The high quantum efficiency of approximately 50% across the whole visual spectrum was achieved by GaAsP PMT detectors. The total number of photon counts collected on TCSPC for each pixel was calculated for each spectral channel and represented in the spectral curve. To avoid partial blocking of photons on the notch filters in our descanned detection setup, which distorts the acquired fluorescence spectra, an optical bypass for collected photons was developed using a combination of acousto-optic tunable filters (AOTF, Optoelectronic) ([Figure S1](#)). The hyperspectral data were fitted by a suitable spectral model, namely the most convenient empirical log-normal function due to its numerical stability during optimization.⁶⁹

$$I_{LN}(\lambda) = \begin{cases} \exp\left(-\frac{\log 2}{4a^2}\right) & \lambda > \lambda_{\max} - \frac{w}{4a} \\ \left[\log\left(\frac{4a}{w}(\lambda - \lambda_{\max})\right) + 1\right]^2, & \\ 0, & \lambda \leq \lambda_{\max} - \frac{w}{4a} \end{cases} \quad (2)$$

with peak position ($\lambda_{\max}$), approximate full width at half-maximum (FWHM, w), and asymmetry (a). The developed model was demonstrated to be capable of resolving changes in the spectra as small as 1 nm.⁶⁹ Pixel binning of 3×3 was employed prior to spectral fitting, increasing sensitivity at the cost of image resolution.

Scanning Electron Microscopy with Energy-Dispersive X-ray Spectroscopy (SEM-EDS). High-resolution imaging and chemical analysis were conducted in the FEI Helios Nanolab 650 scanning electron microscope (SEM) using energy-dispersive X-ray spectroscopy (EDS). To prevent the charging effects of a highly insulating biological specimen during electron irradiation, all samples were coated with an approximately 10 nm carbon layer. High-resolution and high surface sensitivity imaging was performed with low electron acceleration voltage (2 keV), low electron current (100 pA), and under high chamber vacuum (10^{-6} hPa). The EDS spectra were collected with a higher electron acceleration voltage (15 kV) and electron current (200 pA). Under these conditions, an energy resolution of 145 eV at the Mn $K\alpha$ line and an elemental sensitivity of approximately 0.01 wt % (X-Max SDD, Oxford Instruments) were achieved. The weight percentage (wt %) of each individual element was calculated and subsequently employed for further analysis, which entailed the measurement of the differences in the wt % ratios of the studied elements between individual regions.

Helium Ion Microscopy (HIM). High-resolution images at the nanoscale were obtained using a helium ion microscope (Orion NanoFab, Zeiss).¹⁰⁵ For imaging purposes, He atoms are ionized in the GFIS source. The source is characterized by a small virtual source size (0.25 mm), a high reduced brightness $((1-4) \times 10^9 \text{ Am}^{-2} \text{ sr}^{-1} \text{ V}^{-1})$, and a very small energy spread of less than 1 eV. Subsequently, the beam is focused, shaped, and scanned over the sample surface using electrostatic lenses, quadrupoles, and octapoles. The resulting focused ion beam¹⁰⁶ has a diameter of 0.5 nm and enables the recording of images with a large depth of focus.¹⁰⁷ Prior to imaging, the freeze-dried samples grown on Si_3N_4 substrate were affixed with carbon tape to the standard pins compatible with a multipin specimen mount. Insulating samples were spared of conventional thin layer conductive coating due to efficient charge compensation within the HIM instrument using the flood gun,¹⁰⁸ thus preventing any surface modification at the nanoscale studied. Imaging was conducted through the detection of secondary electrons (SE1) emitted from the few-nanometer surface layer of the sample. The following experimental parameters were employed: helium ion energy (30 keV), ion current (0.5 pA), and chamber vacuum (2×10^{-7} hPa). The field of view (FoV) of the acquired images ranged from less-magnified, $175 \times 175 \mu\text{m}$ required for registration, to highly magnified, $0.7 \times 0.7 \mu\text{m}$, with a minimal pixel step size of 0.7 nm and a typical dwell time of 20 μs at each pixel.

Synchrotron Micro-X-ray Fluorescence (SR μ -XRF). The X-ray fluorescence (XRF) data were collected at the TwinMic beamline of Elettra Sincrotrone Trieste, located in Trieste, Italy.¹⁰⁹ The TwinMic microscope was operated in scanning transmission mode, whereby the sample was raster scanned across an incoming perpendicular X-ray beam, delivered by zone plate diffractive optics. During the scanning process, a rapid readout charge-coupled device (CCD) camera (DV 860, Andor Technology) collected the transmitted X-ray photons, resulting in the generation of absorption and differential phase contrast images.¹¹⁰ Simultaneously, the emitted X-ray fluorescence (XRF) photons were collected by eight silicon drift detectors (SDDs) situated in front of the sample at an angle of 20 degrees with respect to the sample plane.¹¹¹ This results in the simultaneous acquisition of both the morphology of the sample and the distribution of the elements.

In order to achieve optimal excitation of Mg, Na, O, Ti, and C, while also obtaining submicron spatial resolution, a monochromatic energy of 1.5 keV was selected for the experiment. The specimens were scanned at a step size of 500 nm with a probe size of 600 nm in diameter, delivered by an Au zone plate of 600 μm diameter and 50 nm outermost zone width. An acquisition time of 50 ms pixel⁻¹ and 4 s pixel⁻¹ was employed for the CCD and SDD detector systems, respectively. The acquired XRF elemental maps and XRF spectra were processed using the XRFFitVis web application and the PyMCA software,¹¹² respectively.

ASSOCIATED CONTENT

Data Availability Statement

The raw data underlying this study are openly available in the HZDR RODARE repository at <https://rodare.hzdr.de/record/3321> (DOI: 10.14278/rodare.3321).

Supporting Information

The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/acsnano.4c17838>.

Additional high-resolution and correlative microscopy results of the nanobio interface supporting the investigated possible modes of action leading to nanoparticle induced lung epithelial toxicity; combined FLIM and fHSI analysis of the DNA presence at the nanobio interface on lung epithelial LA4 cells; example of SEM-EDS mapping at the nanobio interface; super-resolution STED image of the nanobio composite; raw TiO_2 NTs and TiO_2 NTs wrapped in lipoprotein corona by subnanometer resolution HIM; fibrous network formation over nanobio composites in mono- and coculture by high resolution SEM; apical macrophages capturing and internalizing TiO_2 NTs on lung epithelial surface by high resolution SEM; presence of fibrin on fibrous structures by correlative CLSM and HIM; scheme of a custom-built CLSM, STED, FLIM and fHSI optical setup; TiO_2 exposure levels used in the study in the context of the established regulatory exposure limits; and future prospects in correlative microscopy workflow in terms of sample handling and automation (PDF)

AUTHOR INFORMATION

Corresponding Author

Rok Podlipiec – Jožef Stefan Institute, 1000 Ljubljana, Slovenia; Institute for Ion Beam Physics and Materials Research, Helmholtz-Zentrum Dresden-Rossendorf e.V., 01328 Dresden, Germany; orcid.org/0000-0001-8584-3274; Email: rok.podlipiec@ijs.si

Authors

Luka Pirkner – Jožef Stefan Institute, 1000 Ljubljana, Slovenia; Department of Electrochemical Materials, J. Heyrovský Institute of Physical Chemistry, v.v.i., 182 23 Prague 8, Czech Republic

Ana Krišelj – Jožef Stefan Institute, 1000 Ljubljana, Slovenia

Gregor Hlawacek – Institute for Ion Beam Physics and Materials Research, Helmholtz-Zentrum Dresden-Rossendorf e.V., 01328 Dresden, Germany; orcid.org/0000-0001-7192-716X

Alessandra Gianoncelli – Elettra, Sincrotrone Trieste, Trieste 34149, Italy; orcid.org/0000-0002-2457-3618

Primož Pelicon – Jožef Stefan Institute, 1000 Ljubljana, Slovenia; orcid.org/0000-0003-2469-3342

Complete contact information is available at:

<https://pubs.acs.org/10.1021/acsnano.4c17838>

Author Contributions

R.P. conceptualized and designed the analysis; R.P., A.K., L.P., and A.G. performed the experiments; R.P., L.H., and A.G. collected the data; R.P. performed the analysis and data visualization; R.P., G.H., and A.G. contributed analysis tools; R.P. wrote the paper; R.P., L.P., G.H., A.G., and P.P. reviewed and edited the paper; P.P. acquired funding.

Notes

The preprint version of this work¹¹³ is openly available on the bioRxiv server at <https://www.biorxiv.org/content/10.1101/2024.10.11.617573v2>.

The authors declare no competing financial interest.

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