Cellular localization, accumulation and trafficking of DWNTs in human prostate cancer cells.

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The carbon nanotubes (CNT) complexes (RNA-wrapped, oxidized DWNT) were endocytosed by capture in clathrin-coated vesicles, revealed by clear co-localization with both clathrin and transferrin. CNT complexes were also found in early endosomes, which was confirmed by co-localization with early endosome antigen (EEA). CNT complexes were not sorted for recycling back to the cell membrane (via fast recycling endosomes), as co-localization with Rab 4 was not observed. Instead, the overlapping of CNT complexes with both LAMP2 antibody and LysoTracker\textsuperscript{\textregistered} marker indicated their sequestration in lysosomes. CNTs were also found within vesicles of the secretory pathway (Rab11-positive vesicles), which for the first time suggests an exact route for their exocytosis.

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ABSTRACT

Carbon nanotubes (CNTs) are at present being considered as potential nanovectors with the ability to deliver therapeutic cargoes into living cells. Previous studies established the ability of CNTs to enter cells and their therapeutic utility, but an appreciation of global intracellular trafficking associated with their cellular distribution has yet to be described. Despite the many aspects of the uptake mechanism of CNTs being studied, only a few studies have investigated internalization and fate of CNTs inside cells in detail. In the present study, intracellular localization and trafficking of RNA-wrapped, oxidized double-walled CNTs (oxDWNT-RNA) is presented. Fixed cells, previously exposed to oxDWNT-RNA, were subjected to immunocytochemical analysis using antibodies specific to proteins implicated in endocytosis; moreover cell compartment markers and pharmacological inhibitory conditions were also employed in this study. Our results revealed that an endocytic pathway is involved in the internalization of oxDWNT-RNA. The nanotubes were found in clathrin-coated vesicles, after which they appear to be sorted in early endosomes, followed by vesicular maturation, become located in lysosomes. Furthermore, we observed co-localization of oxDWNT-RNA with the small GTP-binding protein (Rab11), involved in their recycling back to the plasma membrane via endosomes from the trans-golgi network.

KEYWORDS

DWNTs, intracellular localization, uptake, immunostaining, inhibition of endocytosis
1 Introduction

The mechanisms that control cellular uptake and release of carbon nanotubes (CNTs) are currently poorly understood. So far, besides the demonstration of the ability of CNTs to enter and penetrate cell membranes [1-4], the uptake mechanism is still debated and may involve more than one process. Indeed, Kam et al. and Jin et al. reported that protein or single-stranded oligonucleotide-modified CNTs could enter cells by means of endocytosis [5, 6]. In contrast, Kostarelos et al. suggested an energy-independent non-endocytic mechanism to be involved in the insertion and diffusion of CNTs through different cellular barriers [2].

Endocytosis is a frequent cellular event that transfers extracellular or membrane-bound cargoes, such as nutrients, from the plasma membrane into the cell interior by vesicular transport. In this process, endocytic vesicles form invaginations of the plasma membrane that subsequently pinch off, move internally, and fuse with other endocytic vesicles to form the early endosomal compartment. The endocytic process may involve so-called clathrin-coated pits (i.e., associated with a clathrin-dependent process). Clathrin forms a complex network in the form of a lattice that is built up to form a coat. Alternatively, there may be clathrin-independent endocytotic processes involved in cellular trafficking. The early endosome is the major sorting station in the endocytic pathways. From this organelle, material can be directed toward the pathway of recycling to the plasma membrane, to subsequent endocytic compartments and to regulated secretory vesicles [7]. Transferrin for example, a monomeric iron transporting serum glycoprotein, is processed in the late endosome, after which the transferrin receptors are recycled by a recycling endosome [8]. The recycling endosome is typically less acidic (pH 6.4-6.5) than the sorting endosome (pH 6.0) and exhibits a pericentriolar localization [9]. Late endosomes contain active degradative enzymes; however, these are more concentrated in the lysosomes. The transfer of material between late endosomes and lysosomes appears to be a direct fusion event that results in a transient hybrid organelle [10].
Each step in endocytosis is independently regulated by the interaction of multiple proteins. These include several small GTP-binding proteins of the Rab family. These proteins have been particularly examined because of their effectiveness as molecular markers for various types of endosomes. Rab5, an example of such GTPase-associated activity [11], regulates the rate of clathrin-dependent endocytosis at the plasma membrane and can be used as a marker for early sorting endosomes in addition to or in conjunction with the transferrin receptor. Rab11 localizes to a pericentriolar subpopulation of recycling endosomes [12]. It is pertinent to point out that the process of endocytosis is a complex process that has yet to be completely defined despite significant progress in understanding its various components.

Identification of the mode by which CNTs move between cellular compartments after their internalization, and the sites in which they localize has been thus far been accomplished by simple staining of lysosomes by pH-dependent markers (e.g. LysoTracker® red) [5, 13], by Transmission electron microscopy (TEM) [13-15], or by pharmacological inhibition of the endocytic pathway [2, 5]. Kam et al. [5] were the first to describe an energy-dependent mechanism. In their studies clathrin-mediated endocytosis was proven by experiments carried out at inhibitory conditions, such as presence of sucrose and potassium-depleted medium, which revealed a significantly reduced level of CNT uptake. Using TEM, Yehia et al. 2007 observed that single-walled carbon nanotubes (SWNTs) dispersed in medium were endocytosed and localized in the plasma membrane, cytoplasm, endosomes, lysosomes and golgi bodies [16]. A similar methodology was used by Mu et al. 2009, who proposed a working model for cell uptake of multi-walled carbon nanotubes (MWNTs) modified by carboxylation or amidation, using TEM imaging [15]. In their model CNTs were not only internalized via endocytic pathways, but also through membrane penetration, depending on whether CNTs were present as bundles or highly dispersed single CNTs, respectively [15]. After both internalization via endocytic pathways or membrane penetration, CNTs were found in lysosomes, followed by their extrusion [15]. However, no data were presented on the mechanism of exocytosis. Jin et al. 2009 presented the first evidence for exocytosis of CNTs, which was found to closely match the endocytic rate [13]. Furthermore, a study in our group demonstrated the time-dependent loss of CNTs by cells [4].

In contrast to the endocytic pathway, Lacerda et al. reported for the first time that membrane
penetration of CNTs (SWNTs modified via cycloaddition) can lead to their accumulation in the perinuclear region of mammalian cells [17]. In addition, another study performed by Zhou et al. in 2010 also showed an endocytosis-independent mechanism for non-covalently modified SWNTs [18]. In their studies, incubations at 4 °C and in the presence of sodium azide (NaN₃) were carried out, which failed to prevent CNTs from being internalized.

In spite of the considerable effort to elucidate the uptake mechanism of CNTs and their intracellular localization, an appreciation of global intracellular trafficking has yet to be clarified. Herein we attempt to complete this study using organelle-specific markers to describe and to identify the mechanisms through which CNTs are transported as they enter and eventually efflux from in vitro-cultured human prostate cancer cells.

2 Results and discussion

DWNT cellular uptake and release has been previously demonstrated by our group using Raman spectroscopy [4]. However, it is also imperative to understand their intracellular route and subcellular localizations during their transit through cells. Hence, this study presents a detailed methodology using confocal microscopy in combination with organelle-specific markers to examine the intracellular distribution of CNTs. In addition, pathway-inhibitory drugs were used to identify major processes through which CNTs traffic as they enter and eventually efflux from cells. For this purpose, fluorescently labelled RNA-wrapped oxDWNTs were employed.

2.1 Characterization of CNT complexes

CNT complexes were characterized using microscopic techniques, in particular atomic force microscopy (AFM) and TEM. Figure 1 (a) depicts an AFM scan over a surface of fluorescein-RNA-wrapped oxidized DWNTs (oxDWNT-RNA-FS). Furthermore, the TEM image in Fig. 1 (b) depicts individualized CNTs with noticeable two-wall characteristics of DWNTs. Raman spectroscopy is a useful tool for the identification of CNTs in a sample. Figure 1 (c) exhibits Raman spectra of pristine DWNTs and oxDWNT-RNA-FS, showing the characteristic Raman features. The first visible peaks are the Radial breathing modes (RBMs), caused by uniaxial vibrations, and depending linearly on the nanotube diameter; these can be found between 150-300 cm⁻¹.
The RBM is then followed by the disorder-induced D-band between 1250 and 1450 cm$^{-1}$, the tangential mode (or G-band), which is caused by stretching along the C-C bonds of graphene, between 1590-1600 cm$^{-1}$ [19-21], and finally the G$'$-band, a two-phonon mode ~2700 cm$^{-1}$ away from the excitation wavelength [19]. The spectra in Fig. 1 (c) demonstrate that the D-band increases during the oxidation process; nevertheless the D/G ratio is overall still rather low due to the preservation of the structure of the inner tube.

Figure 1 (d) exhibits the Raman signal of cells exposed to oxDWNT-RNA-FS and cells as a control. To determine if cells internalized the full complex of oxDWNTs, RNA and fluorescein, the same slides that were used for fluorescent microscopy were used for Raman spectroscopy. The results (Fig. 1 (d)) show that the G-band, which is a characteristic signature band for CNTs, can be clearly observed in the Raman spectrum of cells exposed to oxDWNT-RNA-FS, thus confirming successful cellular uptake. Furthermore, we have shown in an independent study by our group based on Raman spectroscopy that RNA-wrapped oxidized DWNTs are internalized and released by cells in a time-dependent manner [4]. In particular, cells were exposed to CNTs for 3 hours, corresponding with maximal intake, after which CNTs were removed from the surrounding medium and the release of CNTs from the cells monitored for further 21 hours. The study showed that CNT were released progressively over this time frame up until the end point of the experiment (24 hours), when no CNT-signal could be measured any longer inside the cells [4].

The stability of oxidized and RNA-wrapped CNTs in biological fluids, as well as their cytotoxicity, has also been previously examined in our group [22]. In that study it was shown that oxidized CNTs are relatively stable in cellular growth medium and that additional wrapping with RNA further increased their stability in human plasma. Moreover, it was found that cells incubated with oxidized, RNA-wrapped CNTs were 75 % viable [22]. Although RNA-wrapping is potentially prone to degradation in cellular growth medium or the cell interior, it was demonstrated by others that siRNA-wrapped CNTs readily enter cells and that the siRNA remains biological active in the cell even in the presence of serum [23], which is very likely due to the fact that CNTs can protect single-stranded nucleic acids from enzymatic cleavage and intracellular interference from nucleic acid binding proteins [24].
Figure 1: Characterization of CNT complexes. AFM (a) and TEM (b) images of fluorescently labelled RNA-wrapped oxidized DWNTs (oxDWNT-RNA-FS). c) Raman spectra of pristine DWNTs and oxDWNT-RNA-FS. The ratio D/G band has increased for oxDWNT-RNA-FS samples due to acid oxidation. d) Raman spectroscopy of cells exposed to oxDWNT-RNA-FS and control (cells only).

2.2 Cellular internalization of CNT complexes

Confocal microscopy was used to demonstrate the internalization of fluorescently labelled CNTs. PC3 human prostate cancer cells were incubated with 30 µg/mL of oxDWNT-RNA-FS for 2 hours. This incubation period was chosen based on previous papers on endo- and exocytosis of carbon nanotubes, for example by Jin et al., who observed maximal cellular uptake of CNTs between 1 and 2 hours [13]. Subsequently, cells were fixed to prevent morphologic changes and inhibit further natural cellular transport, which “traps” CNTs in the various cellular compartments, in which they are located at the time. This provides a snapshot of the intracellular distribution of CNTs after internalization and release have reached a steady state: while some CNTs are just being internalized, others are already being sorted in lysosomes, or even exocytosed.

Figure 2 reveals that incubation of PC3 human prostate cancer cells with oxDWNT-RNA-FS produced readily detectable fluorescence in the cytoplasm (green) in the same plane than the nucleus, but not in the nucleus itself (counterstained in blue). The cytoskeleton, which is involved in coordinating the organization and dynamics of intracellular architecture and therefore relevant to the further understanding of membrane trafficking was stained with phalloidin (red). Cell morphology appeared regular after exposure to CNTs and inhibitory drugs, and no obvious signs of artefactual cellular damage or associated toxicity with the procedures used throughout this work were observed.
To determine the cellular compartments in which CNT complexes localize and traffic between, organelle-specific antibodies were employed. After fixation to prevent morphological changes, cells were incubated with primary antibodies specific to certain proteins (e.g. clathrin) and then a fluorescently labeled secondary antibody was employed specific to the source of the primary antibody (e.g. rabbit).

The internalization via the endocytic route involves a cascade of intracellular events, with different vesicles and regulators. The observation of CNT complexes in intracellular endosome-like structures (Fig. 2) suggested that an endocytic pathway was involved in the internalization of the complexes. Therefore, it was decided to use markers such as clathrin and caveolin involved in the first step of endocytosis, invagination of the cell membrane, to elucidate which endocytosis mechanism in particular is responsible for CNT uptake. Clathrin-mediated endocytosis is characterized by membrane-bound vesicles inside a coat composed primarily of the fibrous protein clathrin. In contrast, caveolae are plasma membrane invaginations that are enriched in cholesterol and sphingolipids. In addition, these invaginations also accumulate special cholesterol-binding proteins, termed caveolins, on the cytosolic face of vesicles [25-28].

Certain ligands, which are internalized via caveolae, are likely to be delivered to a special endosomal compartment referred to as a “caveosome” [29]. Caveosomes containing caveolins do not transport/co-localize transferrin, lack EEA1 and other early endosomal markers, have neutral pH, and can act as sorting stations [26]. Figure 2 (c) shows co-localization of the CNT complexes (green) with clathrin (red); but no co-localization with caveolin (Fig. 2 (b)).

Generally, newly formed plasma membrane-derived endocytic vesicles lose their clathrin coat and fuse with each other or with pre-existing compartments known as sorting endosomes [30, 31]. The first endosomal compartment, the sorting or early endosome, has a role in sorting the cargo for further transport. The early endosome can be identified using an early endosome antigen (EEA1) in combination with a fluorescent secondary antibody. Our data show that CNT complexes are closely associated with this location, as seen via co-localization studies of CNT complexes (green) with EEA1 (red) (Fig. 2 (d)).

The drop in the pH within the sorting endosome dissociates many ligands from their receptor, allowing those receptors to be recycled back to the plasma membrane. As previously described, small
GTPases have emerged as central regulators of vesicle budding, motility and fusion. One of these proteins is Rab 4, involved in the recycling back to the plasma membrane from the early endosome. However, our results suggest that Rab 4 does not have a function in the trafficking of CNT complexes through the cell, as no co-localization was observed (Fig. 2 (e)).

Maturation of sorting endosomes into late endosomes or multivesicular bodies is coincidental with a drop of the pH. The endosome becomes incapable to commit further fusion with vesicles from the cell surface or other sorting endosomes, thus preventing new cargoes and receptors from being inadvertently degraded by ‘joining’ a late endosome as it matures into/fuses with a lysosome [27]. The lysosome can be identified using antibodies for lysosome membrane proteins (LMP). LMPs reside mainly in the lysosomal limiting membrane and have diverse functions, including acidification of the lysosomal lumen, protein import from the cytosol, membrane fusion and transport of degradation products to the cytoplasm. The most abundant LMPs are the lysosome-associated membrane protein 1 (LAMP1), LAMP2, lysosomal integral membrane protein 2 (LIMP2) and the tetraspanin CD63 [28]. As demonstrated in Fig. 2 (f), a high level of co-localization was observed with LAMP2 (selected for our co-localization studies), thus indicating that CNT complexes end up in the lysosomes after being endocytosed.

The recycling endosome not only allows for the recycling of plasma membrane components (a branch of the pathway that might need to be avoided in the context of drug delivery) but also enables the cell to traffic molecules back to the trans-golgi network (TGN/Golgi) [29]. Again, Rab proteins have been found to regulate this transport, especially the Rab 11 GTPase complex. Figure 2 (g) shows co-localization of Rab11 and the CNT complexes, indicating that they reach the vesicles from the secretory pathway and are delivered by Rab11-positive vesicles. The trans-golgi trafficking, in which Rab 11 is involved may be further validated using antibodies for the golgi97 protein, which localizes on the trans side of the golgi apparatus. It is believed to participate in the trafficking of vesicles between the golgi stacks and the trans-golgi network [32]. This additional analysis, however, exceeded the scope of this paper.

In addition to co-localization with a clathrin-specific antibody, CNT complexes also co-localized with transferrin (Figure 2h). Transferrin is a monomeric serum glycoprotein that binds iron. After its recognition at the cell surface by transferrin
receptors, transferrin is internalized via clathrin-dependent endocytosis. Inside endosomes, the acidic environment favours dissociation of iron from the transferrin. After sorting in late endosomes, transferrin then reaches the golgi apparatus and is subsequently recycled back to the plasma membrane. The co-localization of CNTs with this protein further confirms that they are internalized via clathrin-dependent endocytosis and that they undergo sorting in late endosomes.

Fluorescent microscopy can furthermore be used as a semi-quantitative estimate of the pH in endocytic structures in cell culture models by staining with pH-sensitive dyes, such as the LysoTracker® dyes. This dye stains acidic compartments in live cells, in particular the lysosomes. Thus, in combination with LAMP2 antibody staining it provides further confirmation of the sequestration of CNT complexes in lysosomes. As demonstrated in Fig. 2 (i), there is an overlap between LysoTracker® (red) and CNT complexes (green), indicating that the CNT complexes are located in an acidic compartment. A more detailed analysis including co-localization profiles of selected regions can be found in Fig. S-2 in the Electronic Supplementary Material (ESM).

It is important to note that confocal microscopy does not provide sufficient resolution to observe single nanotubes, which have been shown to traffic within cells in an entirely different fashion (bundles are mainly taken up by endocytosis, whereas single nanotubes can pierce though the membrane of the cell and also those of intracellular organelles) [15, 18]. In addition, cellular uptake of CNTs is strongly influenced by the type of functionalization used. While the surface charge of CNTs (i.e. carboxylated (-) and amine-terminated (+) CNTs) did not seem to have an effect on cellular uptake due to heavy protein coating leading to compensation of the surface charges [15], it has been observed that the attachment of a large molecule, for example the protein bovine serum albumin (BSA), can direct cellular uptake towards endocytosis [18].
Figure 2: Subcellular distribution of CNT complexes in PC3 cells. Cells were incubated with 30 µg/mL of oxDWNT-RNA-FS for 2 hours. a) Cells stained for presence of actin filaments (red) with Alexa Fluor® 546 Phalloidin and with TO-PRO®-3 for nuclei (blue) prior to being imaged for CNT complex accumulation (green). b-i) Co-localization of CNT complexes (green) with antibody-specific and compartmental markers (red) in PC3 cells. Co-localization was observed for clathrin-positive vesicles (c); EEA1-positive early endosomal vesicles (d); Lamp2-positive lysosomes, (f); Rab11-positive recycling endosomes (g); transferrin (h); and LysoTracker® red (i). Yellow colour (arrows) indicates cell compartments positive for both CNT complexes and organelle markers. Counterstaining was not observed for caveolin (b) and Rab 4 (e).

2.3 Effect of pathway inhibitory drugs on CNT-complexes localization

A systematic analysis of intracellular transport pathways is readily achieved by analysing the effects of inhibiting one or more transport processes. A widely used method for inhibiting transport is to apply a temperature block to cells [30]. Most transport processes, including clathrin-dependent internalization, will be blocked at 4 °C. Endocytosis is an energy-dependent mechanism that is also hindered when incubations are carried out in ATP-depleted environments, as
attained by incubation with 10 mM sodium azide (NaN₃). In our experiments (Figure 3 (b) and (c)), both incubations at 4 °C and with NaN₃ resulted in inhibition of uptake of CNT complexes in PC3 cells, suggesting an energy-dependent mechanism for their internalization.

Pharmacological inhibition of cellular trafficking can be achieved with the compound LY294002, which is an inhibitor of Phosphoinositide 3-kinase (PI-3 kinase). PI-3 kinases are a family of enzymes involved in signal transduction and membrane trafficking, as well as other molecular processes. Trafficking processes that are inhibited by LY294002 include translocation to the plasma membrane [33, 34], endocytosis [35, 36], endosome fusion [31], lysosomal protein sorting [37, 38], transcytosis [39] and autophagy [40]. The compound 3-methyladenine is a specific inhibitor of autophagic sequestration [41] that also inhibits PI-3 kinase. When either LY294002 or 3-methyladenine were employed in our study, we observed a blockage of membrane traffic and endosome fusion, which prevented CNT complexes from being internalized by PC3 cells (Figure 3 (d) and (e)).

### 3 Conclusions

The objective of the present study was to elucidate the mechanism of uptake of CNTs, as well as to determine the intracellular compartments in which they localize.

**Figure 3: Effect of inhibitory conditions for endocytic uptake.** Cells were exposed to 30 μg/mL of oxDWNT-RNA-FS for 2 hours and the effects of various (pre)-treatments on CNT internalization were studied. (a). incubation at 4 °C (b); pre-incubation with sodium azide (NaN₃) (c); pre-incubation with phosphoinositide 3-kinase inhibitor (LY294002) (d); pre-incubation with 3 methyladenine (e). Actin filaments (red) were stained with Alexa Fluor® 546 Phalloidin and nuclei (blue) with TO-PRO®-3.
From the data presented herein we propose an endocytic route for uptake and trafficking of CNT complexes (functionalized by means of oxidation and wrapping with RNA). Figure 4 schematically represents the intermediate stations of CNT complexes during their passage through PC3 cells. Evidence of their intracellular localization was demonstrated by Raman spectroscopy and fluorescence spectroscopy of exposed cells. The CNT complexes were found to be endocytosed by capture in clathrin-coated vesicles as revealed by clear co-localization with both clathrin and transferrin. These results are in agreement with the work reported by Kam et al. [5], who also found evidence for clathrin-mediated endocytosis as an uptake mechanism for CNTs. Molecules entering cells via endocytic pathways will rapidly experience a drop in pH from neutral to pH 5.9-6.0 in the lumen of early/recycling vesicles with further reduction from pH 6.0 to 5.0 during progression from late endosomes to lysosomes [32]. In our studies, CNT complexes were found in early endosomes as confirmed by co-localization with early endosome antigen (EEA). CNT complexes were not sorted for recycling back to the cell membrane (via fast recycling endosomes), as co-localization with Rab 4 was not observed; instead, co-localization with both LAMP2 antibody and LysoTracker® marker indicated their sequestration in lysosomes. In addition, CNT complexes were also found within Rab11-positive vesicles of the secretory pathway. Overall, our study brings novel findings on the uptake and traffic of CNTs in mammalian cells, allowing identification of the uptake pathway and the cellular organelles, in which CNTs localize. Additionally, exocytosis was once again confirmed with additional identification of the key organelle responsible for CNTs efflux: it is reported for the first time that the recycling endosomes from the trans-Golgi network (Rab11-positive vesicles) are responsible for extracellular extrusion of CNTs.
Figure 4: Proposed intracellular trafficking of CNT complexes. The internalization of oxDWNT-RNA-FS-complexes was proven to be energy-dependent as it could be blocked by incubation at 4 °C and pre-incubation with NaN₃ (Fig. 3 (b) and (c)). Additionally, incubation with PI-3 kinase inhibitors, known to interfere with vesicular trafficking, led to reduced internalization of CNTs (Fig. 3 (d) and (c)). Co-localization with both clathrin and transferrin provided evidence for clathrin-mediated endocytosis (Figure 2 (c) and (h)), after which CNT complexes undergo sorting in the early endosomes (co-localization with EEA1 – Fig. 2 (d)). CNT complexes are then sequestered in the lysosomes, as confirmed by Lamp2 and LysoTracker® staining (Fig. 2 (f) and (i), respectively). CNTs were also found within vesicles of the secretory pathway (Rab11-positive vesicles), which for the first time suggests an exact route for their exocytosis (Fig. 2 (g)).

4 Experimental

4.1 Preparation of fluorescently labelled oxidized DWNTs

DWNTs synthesized by the CCVD technique[42] were purified in concentrated nitric acid and oxidized in a mixture of nitric and sulphuric acid [43]. In the first step, 100 mg CNTs were dispersed in 20 mL concentrated nitric acid, sonicated with a tip sonicator (MSE Soniprep 150, amplitude 8 µm) 6 times for 10s, and incubated in a water bath (Grant ultrasonic bath XB2, Farnell, UK) at 95 °C for 2 hours. After oxidation, the acid was diluted with water and the mixture centrifuged (10 minutes at 50000 g, 4 °C) (Beckman Avanti J25, Buckinghamshire, UK), followed by thorough washing with distilled
water. These pre-treated CNTs were then redispersed in a 3:1 mixture of concentrated nitric and sulphuric acid and the above described process repeated. Afterwards, the oxidized CNTs were vacuum-filtered using a 0.2 µm polycarbonate filter (Whatman Ltd., UK) until the eluate was clear and of neutral pH. The filtrate was then sonicated with a tip sonicator 6 times for 10s and centrifuged three times at 75000 g to remove large agglomerates and CNT bundles. The concentration of these dispersions was determined gravimetrically and adjusted to 100 µg/mL for storage in the fridge at 4 °C. After oxidation, nanotubes were sterilized by autoclaving at 121 °C for 1 hour and were maintained under sterile conditions for the duration of the experiment. In a second step, the oxidized DWNTs (oxDWNTs) were biofunctionalized by wrapping with fluorescently labelled RNA as described below. Biofunctionalization of oxDWNTs with RNA was achieved using ready-made fluorescently labelled RNA (Eurofins – MWG Operon), with the oligonucleotide sequence (5') CAUCCCGAGUCCACAUUCCCGAGUCC A-FLU (3') (RNA-FS). The manufacturer details states that the oligonucleotide has a molecular weight of 10052 g/mol, Tm of 69.5 °C, GC-content of 53.3 % and has been purified by High-Performance Liquid Chromatography (HPLC). A solution of 1.0 mg/mL (prepared in sterile water) was diluted 1:5 in a solution of oxDWNTs at a concentration of 200 µg/mL (weight ratio CNTs/RNA 1:0.8). After that, the complexes were allowed to form by stirring for 2 hours at room temperature (~21 °C) in the dark. To remove unbound RNA-FS, the solution was filtered and washed several times using 100 kDa centrifugal filters (Millipore). Finally, the RNA-fluorescein wrapped oxidized DWNTs (oxDWNT-RNA-FS) were resuspended in sterile water to a final concentration of 200 µg/mL.

Samples were characterized by Raman spectroscopy, atomic force microscopy (AFM) and transmission electron microscopy (TEM). Raman spectroscopy was performed for pristine DWNTs, oxDWNTs-RNA and cells containing oxDWNT-RNA-FS, using a Renishaw InVia Raman microscope, Elaser=1.59 eV (785 nm wavelength). For AFM sample preparation, oxDWNTs were diluted with water to obtain a concentration of 10 µg/mL. A droplet of 2 µL was placed onto a freshly cleaved mica substrate (1 cm²) and dried in air. AFM measurements were performed using a PicoPlus instrument (Agilent Technologies, Chandler, AZ) operated in contact
mode at room temperature with a lateral scan rate of 1 - 1.5 Hz at 512 lines. Data analysis was carried out using “Gwyddion 2.9”, a free SPM data visualization and imaging tool released under the GNU General Public License. Samples for TEM were prepared by dilution of the sample in acetone and placing a single drop of the solution on a standard copper grid. High resolution TEM measurements were performed using a FEI Tecnai G2F20 S Twin instrument with an accelerating voltage of 200 kV.

4.2 Cellular incubation with CNTs-complexes and compartment staining
PC3 cells were cultured in RPMI-1640 medium supplemented with 10 % FBS, 2mM Glutamax™ and 1 % penicillin-streptomycin (all obtained from Invitrogen, Paisley, UK). Cells were cultured in 24-well (13 mm-well) sterile plates (Nunc, Thermo Scientific) containing glass coverslips (VWR International, Lutterworth, UK) for 24 hours so that they were 50 % confluent at the time of exposure to CNT complexes.

4.2.1 Incubation with inhibitory drugs
For adenosine triphosphate (ATP) depletion studies, a PBS buffer solution supplemented with 10 mM sodium azide (NaN₃) and 50 mM D-glucose was diluted as a ratio of 1:2 in complete RPMI medium and incubated with cells for 2 hour at 37 °C, followed by incubation in a solution of CNT complexes. For inhibitory conditions with LY294002 (phosphoinositol 3-kinase inhibitor) and 3-methyladenine (inhibitor of autophagic sequestration), cells were incubated overnight with 10 μM of drug before addition of the CNT complexes.

4.2.2 Incubation with CNT-complexes
PC3 cells growing in 24-well plates were incubated with 30 µg/mL CNT complexes diluted in complete RPMI medium. After incubation for 2 hours, cells were washed twice with sterile PBS (Invitrogen, Paisley, UK) and fresh medium added for 5 minutes. Cells were again washed with sterile PBS and treated for 10 minutes with CellScrub (Genlantis, San Diego, USA) washing buffer to remove all extracellular CNT complexes. Finally, cells were washed twice more with sterile PBS.

4.2.3 Cell fixation
To prevent morphological changes and inhibit further natural cellular transport, cells were fixed with 4 % (w/v) paraformaldehyde (Sigma Aldrich, Poole, UK) solution in PBS for 1 hour at room temperature. Then cells were washed once
more with sterile PBS and treated with 0.1 % v/v Triton X-100 (Sigma Aldrich, Poole, UK) in PBS to allow permeabilization of cell membranes.

4.2.4 Compartment staining (using phalloidin, TO-PRO®-3, LysoTracker® red & transferrin)

Cell membrane-associated actin filaments were stained using phallotoxins (Alexa Fluor® 555 Phalloidin – emission 565 nm) with a final concentration of 0.044 µM. Transferrin conjugates (Alexa Fluor® 546 - emission 573 nm) at a concentration of 20 µg/mL were applied to stain endosomes associated with clathrin-dependent endocytosis, and acidic cell compartments (lysosomes) were stained using LysoTracker® Red DND-99 (emission 590 nm) at a final concentration of 1 µM. All agents were diluted from freshly made stock solutions in 0.5 % bovine serum albumin (BSA; Sigma Aldrich) in sterile PBS and applied to cells after fixation for a period of 15 minutes. Cells were then washed three times with PBS and finally, To-Pro®-3 (emission 661 nm), a monomeric cyanine nucleic acid stain, was applied in order to visualize the cell nuclei in all samples. It was employed as the final step of the immunocytochemistry using a concentration of 0.1 µM in PBS (all reagents from Invitrogen-Molecular Probes Corporation, unless stated otherwise).

4.2.5 Immunostaining

Subsequent to cell fixation and permeabilization, cells were incubated for 90 minutes with primary antibodies (Santa Cruz Biotechnology, Inc; supplied by Insight Biotechnology; Harrow, UK) diluted in PBS containing 0.5 % BSA, as described in Table 1. Cells were then washed three times with PBS and incubated for 1 hour with fluorescent secondary antibody (Alexa Fluor® 546 from Invitrogen) diluted 1:100 in 0.5 % BSA in PBS, followed by three 5 minutes washes in PBS. After immunostaining, cells were also stained with To-Pro®-3 as described (above).
Table 1: **Immunostaining.** Primary antibodies employed and respective secondary antibodies.

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<thead>
<tr>
<th>Primary antibody</th>
<th>Dilution factor for primary antibody</th>
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<tbody>
<tr>
<td>Clathrin HC (H-300): sc-9069</td>
<td>1:100</td>
<td>Anti-Rabbit</td>
</tr>
<tr>
<td>Caveolin-1 (N20): sc-894</td>
<td>1:200</td>
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<td>Early endosome antigen (EEA1) (C15): sc-6414</td>
<td>1:500</td>
<td>Anti-Goat</td>
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<td>Lysosome-associated membrane proteins (Lamp 2) (H4B4): sc-18822</td>
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<td>Ras-related superfamily - Rab 4/14 (FL-213): sc28569</td>
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<td>Ras-related superfamily - Rab 11 (H87): sc-9020</td>
<td>1:80</td>
<td>Anti-Rabbit</td>
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</table>

### 4.2.6 Mounting glass-slides and microscopy

Following the final step for all samples, in which nucleic acids were stained with To-Pro®-3, coverslips were removed from the 24-well plate using fine forceps and were then held vertically against tissue to remove excess liquid. A small drop of VECTASHIELD® Mounting Medium (Vector Laboratories, UK) was placed on a glass slide and the coverslip, cells facing down, was positioned on top of the Vectashield drop. After removing the excess liquid, coverslips were sealed using a standard commercially available nail lacquer and allowed to dry.

Images were captured using 63x magnification with a Zeiss LSM 510 inverted confocal microscope and analysed using LSM 510 META software (Zeiss).

### Acknowledgements

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### Electronic Supplementary Material:

Supplementary material with further details on time-series analysis of fluorescently labelled RNA-wrapped oxidized DWNTs in cells and a plot profiles of antibody/CNTs is available in the online version of this article at [http://dx.doi.org/10.1007/s12274-***-****-*](http://dx.doi.org/10.1007/s12274-***-****-*) (automatically inserted by the publisher).
References


Electronic Supplementary Material

Cellular Localization, Accumulation and Trafficking of DWNTs in Human Prostate Cancer Cells.

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Supporting information to DOI 10.1007/s12274-****-****-* (automatically inserted by the publisher)

Time-series analysis of fluorescently labeled RNA-wrapped oxidized DWNTs in cells.

An evaluation of CNT accumulation in a particular cellular compartment, the lysosomes, is presented. Cells were exposed to CNT complexes for 5, 10, 15, and 30 minutes; and 1, 2, 4, and 6 hours. Subsequently, the cells were fixed and the lysosomes were stained with LysoTracker® Red in order to afford visualization by confocal microscopy. Figure S-1 shows the accumulation of CNTs in lysosomes at the different time points. After only 5 minutes a significant amount of CNTs has already been internalized, with the level of CNTs slightly increasing within the next 30 minutes. This can be explained by the fact that CNTs were available in excess in the medium, which permits constant uptake and release to take place during the observed period of 6 hours, leading to a steady-state of the intracellular CNT concentration. At intermediate time points it is therefore to be expected to observe CNTs at various intracellular locations: while some are in the process of being internalized, others are already being sorted in lysosomes, or exocytosed. After 5 and 10 minutes few
CNTs (in green) already seem to co-localize with the stained lysosomes (overlap in yellow). In comparison, a much larger amount of CNTs is observed inside the cells after 30 minutes with some degree of co-localization between CNT and lysosomes; however, the majority of CNTs appears to be distributed in other cellular compartments. For 1, 2, 4 and 6 hour-incubation times a similar situation is observed.

**Figure S-1** – Time-series of cells exposed to oxDWNT-RNA-FS (in green). Lysosomes were counter-stained with LysoTracker® Red (in red) and nucleus (blue) stained with TO-PRO®-3.

**Plot profile for antibody/CNT overlay.**

To provide further evidence of co-localization of CNTs and various subcellular compartments counterstained using specific subcellular markers (antibodies) as presented in Figure 2, the intensity of each dye in the specific subcellular regions was evaluated using Image J software. Figure S-2 exhibits a selection of images from Fig. 2 in which co-localization was observed, including a) clathrin; b) EEA1; c) Lamp2, d) Rab11, f) Transferrin and g) LysoTracker®. The graphs in the right columns present the relative intensity for the selected regions within the cell (Line 1 and 2). The results confirm the co-localization of CNTs with the following antibodies: clathrin, EEA1, Lamp2, Rab11, transferrin and LysoTracker®.
Figure S-2 – Fluorescence microscopy of PC3 cells exposed to DWNTs and counter-stained with different antibodies: a) clathrin; b) EEA1; c) Lamp2, d) Rab11, f) transferrin and g) Lysotracker®. Co-localization profiles of selected regions (Line 1 and 2) are presented for each antibody, in which green represents the DWNTs labeled with fluorescein and red the antibody (Alexa Fluor® 546-labelled). Nuclei (in blue) were stained with TO-PRO®-3.
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