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HIV Detection via a Carbon Nanotube RNA Sensor

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Abstract

Viral illnesses remain a significant concern in global health. Rapid and quantitative early detection of viral oligonucleotides without the need for purification, amplification, or labeling would be valuable in guiding successful treatment strategies. Single-walled carbon nanotube- based sensors recently demonstrated optical detection of small, free oligonucleotides in biofluids and in vivo, although proteins diminished sensitivity. Here, we discovered an unexpected phenomenon wherein the carbon nanotube optical response to nucleic acids can be enhanced by denatured proteins. Mechanistic studies found that hydrophobic patches of the denatured protein chain interact with the freed nanotube surface after hybridization, resulting in enhanced shifting of the nanotube emission. We employed this mechanism to detect an intact HIV in serum, resulting in specific responses within minutes. This work portends a route towards point-of-care optical detection of viruses or other nucleic acid-based analytes.

Graphical Abstract

Keywords

Biosensors; Nanosensors; Human Immunodeficiency Virus; AIDS; Fluorescence

Despite a widespread effort to combat the spread of human immunodeficiency virus (HIV), recent data from the World Health Organization estimates that there are 36.9 million people

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Conflicts of Interest Statement

D.H. and J.H. are named on a patent application filed by MSKCC related to this work (application no. WO2017177131A1, title: Sensors for nucleic acid biomarkers). The following disclosures are unrelated to this work: D.H. is a cofounder of LipidSense, Inc. and is a cofounder and officer with equity interest in Goldilocks Therapeutics Inc., as well as a member of the scientific advisory board of Oncorus, Inc. H.A.B is a cofounder of Pictesque Inc. A.K. serves as a consultant for Novartis.

globally living with HIV, but only 59% percent of them are receiving much needed antiretroviral treatment. In many regions with a high incidence of HIV, such as sub-Saharan Africa, the limited access to healthcare resources likely hinders the implementation of effective diagnosis and treatment strategies. One strategy to overcome these challenges is to use rapid, point of care diagnostics, which have been shown to increase the number of patients who know their infection status¹. Although antibody-based testing is currently used in this setting, the false-positive and false-negative rates remain a significant limitation¹.

As an alternative biomarker, cell-free oligonucleotides in biofluids such as serum offer the potential to diagnose, monitor, and stratify disease states based on specific expression patterns, changes, or mutations²⁻⁴. Indeed, realizing such potential has been the subject of recent work focusing on "liquid biopsies⁵," which aims to use such accessible biomarkers to complement or even obviate the need for an invasive, traditional biopsy. While the focus of cell-free oligonucleotide detection has been for cancer, many other diseases of interest also show specific expression patterns of oligonucleotides⁶. Specifically, the measurement of exogenous DNA and RNA strands found in viruses is a promising method for rapid diagnosis of infectious diseases⁷. Rapid, quantitative detection of viral oligonucleotides without the need for purification, amplification, or labeling of the oligonucleotide could help diagnose infections before symptoms appear, monitor the course of the disease, and also characterize viral subtypes and clades 8 . A simple point-of-care option would therefore be especially valuable in developing countries prone to infectious disease outbreak⁸, especially HIV. Because the viral genome is typically sequestered inside a protein capsid and in some cases an additional lipid bilayer, a point-of-care sensor would have to be compatible with lysing conditions to liberate the viral oligonucleotides. In addressing the need for such pointof-care options, diverse solutions based on nanomaterials are being developed, which have recently been reviewed elsewhere⁹.

Single-walled carbon nanotubes are one class of nanomaterials capable of optical transduction of binding events via modulation of their intrinsic photoluminescence¹⁰. Carbon nanotubes are fluorescent in the near-infrared range¹¹ and are highly resistant to photobleaching12. Many nanotube species (chiralities) are available and defined in part by unique excitation and emission bands¹³. Carbon nanotube emission responds to changes in the local dielectric environment^{14–16}, including changes in local static charge^{17–18}.

Carbon nanotubes were recently used to detect biomarker oligonucleotides such as microRNA directly in biological fluids such as serum and urine¹⁹. While such carbon nanotube sensors were shown to detect target oligonucleotides in simple buffer conditions with a modest shift in emission wavelength, the optical change upon detection was enhanced by the presence of the surfactant sodium dodecylbenzenesulfonate (SDBS). This was mediated by changes in the nanotube surface coverage of the DNA upon hybridization, resulting in increased binding of SDBS to the nanotube surface, inducing changes in the optical bandgap. The enhancement effect was found to persist in biofluids such as serum and urine and enabled detection of short DNA oligonucleotides and microRNA¹⁹.

Herein, we describe the optical detection of HIV in complex media by a carbon nanotube sensor. Intact viruses and protein-rich solutions such as serum normally prevent detection of

target viral oligonucleotides. We found that, in the presence of serum, a denaturing surfactant mediated enhanced nanotube emission response to viruses. To deduce the mechanism of this enhancement, purified proteins such as bovine serum albumin, γglobulins, casein, and hydrolyzed casein peptides, were assessed. We conclude that hydrophobic pockets of surfactant-denatured protein interact with the nanotube upon hybridization, leading to a greatly enhanced blue-shift. We assessed the potential for clinical application by measuring single-strand viral RNA from an intact human immunodeficiency virus (HIV) using the nanotube sensor, resulting in successful detection. This work portends the use of carbon nanotubes for point-of-care optical detection of viruses via measurement of viral nucleic acids.

Results and discussion

Using a carbon nanotube-based sensor for the miR-19 microRNA sequence (GT_1 ₅miR19) as previously described¹⁹, we characterized the effect of serum proteins on the nanotube response using a single-stranded miR-19 DNA analyte sequence. In buffer-only conditions, the target, miR-19 DNA, produced a blue-shift in nanotube emission for most nanotube species (chiralities). However, the addition of fetal bovine serum (FBS, 10%), abrogated wavelength changes in response to the target oligonucleotide (Figure S1a). Two-dimensional photoluminescence excitation/emission (PLE) spectroscopy was used to assay optical changes for 11 different nanotube chiralities incubated with FBS after addition of the target miR-19 oligonucleotide or a non-complementary R23 DNA control strand (Figure S3). For every chirality measured, FBS abrogated optical changes in response to the target oligonucleotide (Figure 1a–c). To better understand the effect of FBS on the sensor baseline fluorescence, we measured the optical changes of the sensor as a function of FBS concentration in the absence of miR-19 DNA. Increasing concentrations of FBS induced red-shifting and a comcomitant increase in intensity (Figure S1 b and c). Because nanotube emission responds to changes in the local dielectric environment, we interpret these redshifting and intensity responses as resulting from proteins in the serum closely associating with the bare-regions of the DNA coated nanotubes, electrostatically interacting with the phosphate backbone to induce DNA conformation changes, or some combination thereof.

In an attempt to reduce this apparent FBS-induced fouling of the sensor, we hypothesized that an anionic detergent, sodium dodecyl sulfate (SDS) may associate with bare-regions of the nanotube and block non-specific binding of serum components. We base this hypothesis on previous works showing that SDS can be used to suspend carbon nanotubes¹¹, due to hydrophobic interactions between the nanotube surface and the 12-carbon long alkyl chain20. We first tested whether SDS would change the baseline emission of the GT ¹⁵miR19 sensor. Dilutions ranging from 0.1% to 1% wt/vol SDS showed no significant changes in emission wavelength (Figure S2a) or intensity (Figure S2b). When hybridization of target DNA was measured in the presence of SDS via PLE spectroscopy (Figure S3), only a slight enhancement was observed for some chiralities, compared to the buffer-only condition (Figure 1b). Although SDS seemed to have little effect on the DNA-wrapped nanotube baseline fluorescence and changes during hybridization, we hypothesized that SDS may preserve the response to hybridization under serum conditions. The GT_1 ₅miR19 sensor was then added to whole FBS with target or control oligonucleotide, followed by addition of

concentrated SDS to a final concentration of 1%. After 4h, PLE plots were acquired to measure the nanotube optical response (Figure S3). Surprisingly, all observed nanotube species showed a greatly enhanced solvatochromic response to target oligonucleotide, with emission blue-shifting from 3 to 9 nanometers depending on chirality (Figure 1d). The excitation wavelengths (E_{22}) also shifted, suggesting a change in the ground state-absorption (Figure 1e). When the concentration of miR-19 DNA was varied, the sensor retained dosedependent blue-shifting behavior (Figure 1f).

We next measured the kinetics of the sensor response. The nanotube emission response was measured upon introduction of target miR-19 DNA and non-complementary control R23 DNA in both PBS buffer and serum, with and without SDS (Figure 2). Addition of SDS to buffer-only conditions had little effect as compared to the buffer control, and there was also no response in untreated serum. The combination of serum and SDS produced a greatly enhanced blue-shift within approximately 30 minutes. Compared to the buffer conditions, the shape of the kinetic trace was sigmoidal with an inflection point around 2 h that reached a minimum by 3 h. In the FBS conditions with SDS, there was a slight red-shift of the sensor in response to the R23 control DNA strand. This is likely because of the non-specific adsorption of the denatured protein to the baseline amount of exposed nanotube surface. It is probable that it is more apparent in the presence of SDS over the FBS alone because denaturing the protein structure exposes additional hydrophobic patches that become available for binding. The non-specific FBS induced red-shifting of DNA-wrapped nanotubes was discussed above (Figure S1).

We assessed the effect of surfactants on the sensor kinetics in both buffer and serum conditions. We previously reported hybridization-induced enhancement of solvatochromic changes using sodium dodecylbenzene sulfonate (SDBS), which also enabled detection in serum conditions¹⁹. The presence of SDBS produced a similar line shape regardless of buffer or serum conditions and resulted in a blue-shifting response that exhibited pseudofirst order kinetics (Figure S4a). The similar response in buffer and serum suggests the same mechanism in both conditions, wherein SDBS interacts with newly-exposed regions of the nanotube upon hybridization of the bound DNA. In the presence of SDS in serum, upon the introduction of DNA, there was a lag period before an optical blue-shifting (Figure S4b). Although slower than SDBS, the SDS + serum combination produced a final blue-shifted value of greater magnitude (Figure S4c and d). Because SDS minimally enhanced the wavelength shifting in the absence of serum, and because of the relatively complex kinetics of SDS + serum enhancement compared to SDBS, we hypothesized that SDS-denatured a protein component of the serum that subsequently bound to the newly-exposed regions of the nanotube upon hybridization, to provide an enhanced blue-shifting response.

We tested the hypothesis that albumin, one of the most abundant macromolecules in serum, elicits the enhanced hybridization response in the presence of SDS. Serum albumin ranges from 35 mg/mL to 50 mg/mL in healthy individuals²¹. Using purified bovine serum albumin (BSA), we tested the high end of the normal range, 50 mg/mL, with SDS to determine if this purified component of serum would enhance the nanotube response to hybridization in the absence of other serum components. A serial dilution of miR-19 DNA produced dosedependent enhanced blue-shifting, suggesting that BSA + SDS elicited enhancement of the

emission wavelength response upon hybridization (Figure 3a). We next kept the concentration of target DNA and SDS constant at 1 μM and varied the concentration of BSA. For most chiralities, there was a BSA dose-dependent pattern in blue-shifting and intensity enhancement (Figure 3b–c). Although some chiralities exhibited less wavelength shifting at the highest protein concentration, interestingly, even in these cases, the intensity enhancement was still dose-dependent. (Figure S5).

Serum is a complex biological fluid with many protein and non-protein components 22 . Because non-polar compounds tend to cause blue-shifting of nanotube emission^{23–24}, we hypothesized that lipoproteins, which transport non-polar fats²⁵, may be the component of serum responsible for SDS-induced hybridization enhancement. GT_{15} miR19 sensor was spiked into human lipoprotein-deficient serum (LPDS) or standard FBS with different concentrations of target DNA, followed by addition of concentrated SDS to a final concentration of 1% wt/vol. The dose- response behavior and magnitude of blue-shift upon hybridization with target oligonucleotide was largely indistinguishable between the two serum conditions (Figure 3d), suggesting that lipoproteins do not play a role in the BSAmediated response.

Finally, we kept the concentration of BSA and target oligonucleotide constant and varied the SDS concentration. We escalated the concentration from 0 to 6% in 1% increments, but observed no additional enhancement above 2% SDS (Figure 3e). Narrowing the tested concentrations between 0 and 1% in 0.1% increments revealed, however, that blue-shifting was dose-dependent in this range. This response is in accordance with reports of SDSmediated BSA denaturation²⁶. This change occurs in the proximity of the critical micelle concentration (CMC) of SDS, which is approximately 0.23 % wt/vol. These data support our hypothesis that SDS-denatured protein is a major causative agent in serum for the observed hybridization- induced enhancement of the blue-shifting response of the nanotubes.

Considerable effort has been made towards understanding the microstructure and mechanism of SDS denaturation of proteins, including the relative contribution of hydrophobic interactions versus ionic interactions^{27–29}. The currently accepted view is that below the critical micelle concentration of SDS, protein tertiary structure unfolds³⁰, and at micellar concentrations, chain expansion drives complete denaturation. The interaction is predominantly hydrophobic at submicellar concentrations, and exclusively hydrophobic at micellar concentrations. In this model, the chain expansion is driven by micelle nucleation on hydrophobic patches of the protein chain³¹. In other words, increasing concentrations of SDS increase the fraction of denatured protein, which becomes available to adsorb onto the exposed surface of the nanotube. The probability of a protein denaturing in the presence of SDS depends on its aggregation state. That is, as a monomer, detergents bind to the native protein as a conventional ligand would, in a fixed number of possible binding sites. As the concentration of detergent increases, it begins to form micelles, which are denaturing to the surrounding proteins. Therefore, global protein unfolding typically occurs above the micelleforming concentration, or critical micellar concentration $(CMC)^{32}$. For SDS specifically, this occurs at 7 mM, or 0.23% wt/vol in water, as mentioned above. Since we see no additional change above 2% SDS, we believe the enhancement effect is saturable, and thus dependent on the denaturation status of the proteins in the system. Below the CMC, the concentration

of SDS directly affects the amount of denatured proteins; therefore it affects the magnitude of the shift.

Based on our experimental observations and this model, we propose the following mechanism for SDS-denatured protein enhancement of hybridization-induced blue-shifting: hybridization with miR-19 oligonucleotide leads to partial desorption of DNA from the nanotube, exposing the hydrophobic surface to the solvent conditions¹⁹. SDS drives loss of tertiary structure and exposure of hydrophobic patches on the polypeptide chain, which would typically be coated by an SDS micelle³³, have a stronger affinity for the newlyexposed nanotube surface than with SDS, leading to preferential binding of the hydrophobic patch to the nanotube surface.

The competition between SDS and the bare nanotube surface for the hydrophobic patch may explain the relatively slow kinetics compared to SDBS (Figure S4). The protein-nanotube interaction is expected to decrease the local dielectric environment around the nanotube, producing the observed blue-shift in emission. The remainder of the DNA-coated nanotube is protected due to charge-charge repulsion between the negatively charged SDS-denatured protein chain and the negatively charged DNA phosphate backbone. Because SDS does not change baseline fluorescence of the nanotube (Figure S3) and produces almost no enhancement of blue-shifting after hybridization with the target in the absence of protein (Figure 1b), we conclude that free SDS alone makes little contribution to the observed blueshift. Protein alone results in red-shifting of the nanotube emission and largely prevents hybridization (Figure 1a, Figure S1a), possibly due to electrostatic interactions with the DNA-suspended nanotube¹⁹ that are eliminated by SDS coating the protein chain. The hybridization conditions and resulting optical outputs are summarized in figure 4.

We next investigated the impact of the protein size and tertiary character on SDS-denatured protein enhancement of hybridization-induced blue-shifting. Using the relatively large γ globulins (1200 kDa compared to albumin, 66.5 kDa), a subset of the second most abundant protein class in serum after albumin, we tested if SDS-mediated denaturation would lead to enhanced blue- shifting upon hybridization. A solution of γ -globulins at 35 mg/mL was exposed to 2% wt/vol SDS with target or control DNA. In this case, although we found that the sensor response to SDS-denatured γ -globulins was relatively slow, there was a similar equilibrium of blue-shifting compared to BSA (Figure S6a). Unlike BSA, SDS-denatured $γ$ -globulins did not lead to increased emission intensity upon hybridization (Figure S6c).

We also examined another protein, casein, the main protein component of milk, which is not found in serum but has a micellar tertiary structure as a distinguishing characteristic 34 . Hydrolyzed casein, which is casein that has been digested into smaller peptide and protein fragments, was also tested to compare the effect of protein fragmentation. We tested whether SDS + casein would recapitulate the results seen for BSA and immunoglobulins, and whether SDS + hydrolyzed casein would produce any enhancement due to the fragmentation from partial digestion. Casein and hydrolyzed casein were prepared at 35 mg/mL and SDS was added to 2% wt/vol, followed by target or control DNA. For most chiralities, SDS + casein enhanced hybridization-induced blue-shifting and intensity increases, while hydrolyzed casein elicited only modest enhancement over controls (Figure S6b, d). This is

consistent with work showing that the degree of hydrolysis of casein is correlated with loss of hydrophobicity³⁵. However, we noted that some chiralities showed significant hybridization-induced blue-shifting enhancement with SDS + hydrolyzed casein, sometimes exceeding the enhancement produced by SDS + casein (Figure S6 e–h). These results suggest that most proteins, independent of their tertiary structure, combined with SDS, will lead to hybridization-induced enhancement with a similar magnitude at equilibrium. A relatively long protein chain appears to be important, as most chiralities did not respond well to SDS + hydrolyzed casein as an enhancing agent. Two nanotube chiralities were notable exceptions, where SDS + hydrolyzed casein performed better than full length casein. Chirality-dependent selectivity to analytes has been reported ¹⁹. $36-37$ but the mechanism for chirality selectivity observed here requires further study.

We examined whether the single-stranded RNA genome³⁸ of intact HIV particles could be detected by the sensor via denatured protein-mediated enhancement. We hypothesized that SDS could be both applied to liberate the genome of the viruses and additionally to denature proteins for hybridization-induced enhancement of the sensor. A sensor was constructed to hybridize to the polyadenylation elements of HIV RNAs by using the DNA oligonucleotide sequence $(GT)_{15}$ -(T)₁₅ DNA to suspend carbon nanotubes. As a control, a sensor for a polyuridine or polythymine sequences, which are not known to exist in HIV, was similarly constructed with the sequence $(GT)_{15}$ - $(A)_{15}$ DNA. The function of both sensors was first assessed with short strands of target DNA and 0.2% SDBS, based on the assumption that DNA would hybridize to the sensor as well as RNA³⁹. The sensor for poly-A sequences, $(GT)_{15}$ - (T)₁₅, showed a rapid (complete in 90 min) and robust blue-shift when treated with (A) ₁₅, but no change was observed after treatment with (T) ₁₅, which served as a control for non-specific oligonucleotide binding (Figure 5a).

To test the sensor with intact HIV particles, we used established methods for production of recombinant HIV particles engineered to be replication-incompetent for safety 40 . Suspensions of viral particles were treated with 1% SDS and $(GT)_{15}$ - $(T)_{15}$ in FBS, the sensor for the poly-(A) elements, or with $(GT)_{15}$ -(A)₁₅, to serve as a negative control. The center wavelength of the (8,6) nanotube was measured every 10 min over 180 min. The sensor for poly-(A) sequences, $(GT)_{15}$ - $(T)_{15}$, began to blue-shift within 20 min, and by 180 min had shifted about 3 nm. The kinetics exhibited a sigmoidal shape, suggesting enhancement from denatured viral proteins. The control sensor, $(GT)_{15}$ - $(A)_{15}$, blue-shifted by about 0.5 nm over 180 min, indicating presumably non-specific interactions (Figure 5b). We present a model for such detection, illustrated in figure 5c. Here, intact lentivirus is treated with SDS and a carbon nanotube sensor. The SDS denatures the lentivirus envelope and protein structure, releasing the single-strand RNA genome. Hybridization with the poly A tail enables binding by the detergent-denatured protein, leading to a blue-shift of the nanotube emission.

Conclusions

Here, we investigated the detection of HIV by carbon nanotube photoluminescence. We found that SDS-treated serum can interact with carbon nanotubes to enhance the blueshifting response of hybridization-induced removal of DNA from the nanotube surface. In

understanding why SDS-treated serum provided such enhancement, we examined lipoprotein-deficient serum as well as purified serum proteins to identify the causative agent. We found that purified albumin, in the presence of SDS, could largely, but not totally, recapitulate the enhancement seen with serum. We surmise that the other proteins in serum may elicit the larger response, although it is conceivable that non-protein factors could be responsible22. The enhancement to hybridization-induced blue-shifting by SDS-treated protein was unexpected due to the fact that SDS alone does very little to DNA-suspended nanotubes, and protein alone results in red- shifting of DNA-suspended nanotubes. Based on the current understanding of SDS-induced protein denaturation and the mechanism of the DNA-carbon nanotube sensor, we propose a model where hydrophobic residues from the SDS-denatured protein bind to newly exposed nanotube surface after hybridization of the DNA on the nanotube, leading to the enhancement in blue-shifting.

For clinical applications, we believe that diagnostics based on detection of viruses in protein rich environments such as serum could benefit from a rapid optical sensor. We previously reported in vivo and serum measurements using another detergent, SDBS, to enhance detection of oligonucleotides. In serum, we believe that SDBS and SDS operate with two distinct mechanisms, where SDBS interacts with the nanotube directly after hybridization, while SDS functions to expose hydrophobic centers in denatured proteins, which then interacts with the nanotube. Although SDBS produces a faster response in serum, SDS produces a shift with a greater magnitude.

As a proof-of-principle for direct optical detection of a virus in vitro, we showed how SDS can be combined with a DNA-nanotube sensor to both liberate lentiviral RNA from a complete virus structure and elicit an enhanced response to the analyte RNA via binding of SDS-denatured viral proteins to the space liberated on the nanotube by the hybridization event. The shifting response in the presence of intact HIV (Figure 5b), however, was less robust than the same sequence of DNA (Figure 5a). In the initial development of the sensor construct, a similar phenomenon was noted¹⁹. The hybridization of RNA to the sensor was noticed to be consistently slower than the same sequence of DNA and it often continued to trend downward, even after 180 min. This was attributed to the longer persistence length and higher rigidity of single-stranded RNA compared to single-stranded DNA. In figure 5b, the difference at 180 min is distinct enough for the purposes of an *in vitro* sensor, therefore the experiment was terminated at this time point. Because the kinetics are similar to the wellestablished response of the sensor to RNA, and because of the investigation of different protein responses in figure S6, we are confident that the sensor is responding to the viral RNA specifically.

Such methods could take place directly in serum samples where point of care detection is desirable. Field applications in low-resource settings are also possible, as the DNA-nanotube sensor solution and SDS are both highly stable and can be stored at room temperature. In addition, an inexpensive and robust detection device could be developed using standard photodetectors and LED light sources for deployment in regions lacking critical biomedical infrastructure. Many viruses of interest, such as Ebola, are viruses with single-stranded RNA genomes that may be amenable to detection with this method. While we have shown here an application with a lentivirus model, we believe that SDS could also be applied to detection

of single-strand oligonucleotides in other viruses and pathogen types, as well as in circulating cancer cells.

Materials and Methods

DNA-suspension of carbon nanotubes

Carbon nanotubes produced by the HiPco process (Unidym, Sunnyvale, CA) were mixed with DNA oligonucleotides (IDT DNA, Coralville, IA) at a 2:1 mass ratio in 1 mL of PBS buffer (Thermo Fisher Scientific Waltham, MA) and ultrasonicated for 30 minutes at 40% amplitude (Sonics & Materials, Inc.). The DNA sequences used for suspension were GTGTGTGTGTGTGTGTGTGTGTGTGTGTGTTCAGTTTTGCATAGATTTGCACA (GT15miR19), GTGTGTGTGTGTGTGTGTGTGTGTGTGTGTTTTTTTTTTTTTTTT $((GT)_{15}$ - $(T)_{15}$), and

GTGTGTGTGTGTGTGTGTGTGTGTGTGTGTAAAAAAAAAAAAAAA ((GT)15-(A)15) all purchased from IDT (Coralville, IA). Following ultrasonication, the dispersions were ultracentrifuged (Sorvall Discovery 90SE) for 30 minutes at $280,000 \times g$ and the top 80% of the supernatant was collected. Absorbance spectra were acquired using a UV/Vis/nIR spectrophotometer (Jasco V-670, Tokyo, Japan). The concentration for HiPco samples was calculated using the extinction coefficient Abs₉₁₀ = 0.02554 L mg⁻¹cm⁻¹. Excess DNA was removed via 100 kDa Amicon centrifuge filters (Millipore). The DNA-nanotube complexes were re-suspended in PBS (Thermo Fisher Scientific Waltham, MA).

Fluorescence spectroscopy measurements of carbon nanotubes

Fluorescence emission spectra from aqueous nanotube solutions were acquired with a homebuilt spectroscopy system as described¹⁹. Briefly, carbon nanotube samples were assayed in a 96 well plate format on an inverted microscope using a tunable white-light laser (NKT Photonics) coupled to a variable bandpass filter. Emission light was directed into a spectrometer with a focal length of 320 mm and aperture ratio of f/4.6 (Princeton Instruments IsoPlane SCT 320) and InGaAs array camera (Princeton Instruments 640×512 pixel NIRvana). For all experiments, measurements were taken in triplicate across three wells. Following spectra acquisition, custom code written in Matlab applied spectral corrections and background subtraction as described¹⁹. Corrected spectra were used to fit the data with Voigt functions. Error bars and linear fits were computed with GraphPad Prism 6.

Hybridization conditions

Concentrated sodium dodecyl sulfate (Sigma-Aldrich) was prepared in PBS buffer and added to samples to the final concentration indicated, or 1% wt/vol where not indicated; control samples had an equivalent volume of PBS added. Carbon nanotubes were used at a final concentration of 2 mg/L for all experiments. Fetal bovine serum (FBS) (Life Technologies) and human lipoprotein deficient serum (LPDS) (J65516 Lipoprotein Deficient Serum, human, BT-931 from Alfa Aesar) were used for serum-based experiments, and had concentrated nanotubes or SDS added directly to the whole serum. Bovine serum albumin, bovine Y-globulins, casein, and hydrolyzed casein (all from Sigma-Aldrich) were resuspended in PBS to the indicated concentrations. Concentrated target DNA (TGTGCAAATCTATGCAAAACTGA) complementary to the DNA-nanotube sensor or

non-complementary control DNA (TCGGTCAGTGGGTCATTGCTAGT) (both from IDT DNA, Coralville, IA) was added to the indicated final concentration. Non-treated controls received an equivalent volume of PBS. Hybridization was allowed to proceed overnight for equilibrium measurements. Kinetics were measured immediately after addition of target. Assessment of $(GT)_{15}$ - $(T)_{15}$ and $(GT)_{15}$ - $(A)_{15}$ used 0.2% SDBS (Sigma-Aldrich) as described¹⁹ to verify expected behavior with AAAAAAAAAAAAAAAAAAA or TTTTTTTTTTTTTTT (IDT DNA, Coralville, IA).

HIV production

Lentivirus HIV was produced from transfected packaging cells (293T) using 100 ng pLX304 plasmid38, 100 ng of packaging plasmid (psPAX2) and 10 ng of envelope plasmid (VSV-G). Virus was collected 48 and 70 h after transfection and concentrated 10-fold.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Figure 1:

Modulation of the GT_{15} miR19 nanotube sensor response to nucleic acids. **a.** Wavelength shifting response of the GT_1 ₅miR19 nanotube sensor after overnight incubation with 1 μ M target miR-19 DNA (orange) or control R23 DNA (purple) for the indicated nanotube chiralities in buffer. **b.** Wavelength shift after overnight incubation with 1 μM target miR-19 DNA (orange) or control R23 DNA (purple) for the indicated nanotube chiralities in 1% SDS. **c.** Wavelength shift of the GT_1 ₅miR19 nanotube sensor after overnight incubation with 1 μM target miR-19 DNA (orange) or control R23 DNA (purple) for the indicated nanotube chiralities in fetal bovine serum (FBS). **d.** Wavelength shift after overnight incubation with 1 μM miR-19 target DNA (orange) or control R23 DNA (purple) for the indicated nanotube chiralities in FBS treated with 1% SDS. **e.** Plots of excitation and emission wavelength shift in response to target miR-19 DNA and control R23 DNA in serum treated with 1% SDS. Nanotubes treated with target DNA are labeled by chirality, while the labels for nanotubes treated with R23 control DNA labels were omitted for clarity. Colors indicate mod type of the nanotube. Error bars represent standard deviation of technical triplicates. **f.** Wavelength shift of GT_1 ₅miR19 nanotube sensor after overnight incubation with indicated concentrations of target miR-19 DNA (orange) or control R23 DNA (purple) in FBS treated with 1% SDS. Error bars represent standard deviation of technical replicates.

Figure 2:

Dynamics of GT₁₅miR19 nanotube sensor response. Each condition depicts 2 mg/L GT₁₅miR19 nanotube sensor treated with 1 μM target miR-19 DNA (orange in PBS, light green in serum) or control R23 DNA (purple in PBS, dark green in serum) added at time 0

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Figure 3:

 GT_{15} miR19 nanotube Sensor response to albumin. **a.** Shifting of the (9,4) nanotube emission wavelength after treatment of the GT_1 ₅miR19 nanotube sensor with the indicated concentration of target or control DNA in 50 mg/mL BSA treated with 1% SDS. **b.** Wavelength shift after treatment with 1 μM target miR-19 DNA or control R23 DNA in PBS with 1% SDS and the indicated concentration of bovine serum albumin (BSA). **c.** Intensity changes of GT_{15} miR19 nanotube sensor after treatment with 1 μ M miR-19 target DNA or control R23 DNA in PBS with 1% SDS and the indicated concentration of BSA. **d.** Wavelength shifting response of GT_{15} miR19 to dilutions of target miR-19 DNA or control R23 DNA in fetal bovine serum (FBS) treated with 1% SDS or human lipoprotein-deficient serum treated with 1% SDS after incubation overnight. **e.** Wavelength shifting response after overnight incubation with 50 mg/mL BSA and 1 μM of target miR-19 (orange) or control R23 (purple) DNA and escalating concentrations of SDS. All error bars represent standard deviation.

Figure 4:

Summary of GT_{15} miR19 nanotube sensor responses and models of hypothesized interactions. Grey cylinders represent the carbon nanotube, and suspending DNA is represented by blue $(GT_{15}$ portion) and orange (complementary region to target oligonucleotide) ribbon.

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Figure 5:

Detection of intact virus RNA using a HIV lentivirus model. **a.** Kinetics of wavelength shift of nanotube suspended with the DNA oligonucleotide $(GT)_{15}$ - $(T)_{15}$, focusing on the (8,6) nanotube species, upon introducing target oligonucleotide, (A) ₁₅, or non-target control, (T)₁₅, with 0.2% SDBS. **b.** Kinetics of the response of $(GT)_{15}(T)_{15}$ or $(GT)_{15}(A)_{15}$ suspended nanotubes with concentrated HIV and 2% SDS. **c.** Model depicting HIV RNA detection. The virus is ruptured and denatured by SDS, liberating the RNA genome, which hybridizes to the sensor, freeing space on the nanotube surface. Denatured viral proteins bind to the freed space on the nanotube surface, eliciting an enhanced blue-shifting response.