Surface-structure-regulated cell-membrane penetration by monolayer-protected nanoparticles

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Nanoscale objects are typically internalized by cells into membrane-bounded endosomes and fail to access the cytosolic cell machinery. Whereas some biomacromolecules may penetrate or fuse with cell membranes without overt membrane disruption, no synthetic material of comparable size has shown this property yet. Cationic nano-objects pass through cell membranes by generating transient holes, a process associated with cytotoxicity. Studies aimed at generating cell-penetrating nanomaterials have focused on the effect of size, shape and composition. Here, we compare membrane penetration by two nanoparticle ‘isomers’ with similar composition (same hydrophobic content), one coated with subnanometre striations of alternating anionic and hydrophobic groups, and the other coated with the same moieties but in a random distribution. We show that the former particles penetrate the plasma membrane without bilayer disruption, whereas the latter are mostly trapped in endosomes. Our results offer a paradigm for analysing the fundamental problem of cell-membrane-penetrating bio- and macro-molecules.

Nanomaterials are of great interest for use in biomedicine as imaging tools1–3, phototherapy agents4,5 and gene delivery carriers6,7. Their interactions with cell membranes are of central importance for all such applications. For example, many drug-delivery systems are based on the transport of therapeutic agents to the cytosol or nucleus of cells by nanoparticles; efficient delivery must be achieved while avoiding cytotoxicity during passage through cell membranes to reach intracellular target compartments8,9. Indeed, membrane penetration by synthetic10 as well as by biologically derived11 molecules/particles is currently under intense investigation. Some biomacromolecules, such as cell-penetrating peptides (CPPs), may be capable of penetrating membranes without overt lipid bilayer disruption/poration12–15. Likewise, synthetic nanomaterials with very small dimensions (molecules, metal nanoclusters16, small dendrimers17 and carbon nanotubes18) can also pass through cell membranes. However, to the best of our knowledge, no synthetic material larger than a few nanometres in size can pass through membranes without disrupting the integrity of these biological barriers. For example, charged particles (such as cationic quantum dots or dendrimers, mostly assisted by some degree of hydrobicity) induce transient poration of cell membranes to enter cells, a process associated with cytotoxicity19. Alternatively, nanoparticles have been designed to explicitly disrupt endolysosomal membranes to enter the cell by force20 or enter the cell aided by exogenous agents such as CPP chaperones21. In contrast, most nanoparticles are trapped in endosomes22 and hence do not reach the cytosol.

The surface properties of nanomaterials play a critical role in determining the outcome of their interactions with cells23. Recently, we found that when gold nanoparticles are coated with binary mixtures of hydrophobic and hydrophilic organic molecules, ribbon-like domains of alternating composition spontaneously form in the ligand shell23–25. These domains are on average less than 6 Å wide, which is of the order of or even smaller than the distribution of chemical functionalities on biomolecules such as proteins and peptides. Order, rather than random organization of the functional groups, on these particle surfaces leads to unexpected surface properties due to the molecularly close apposition of hydrophilic and hydrophobic moieties at all points on the particle surfaces: the particles exhibit unusual solubility trends and low protein binding even with high contents of hydrophobic groups26. With the ordered amphiphilic structure of some CPPs in mind, we explored the interaction of these nanoparticles with living cells. Here, we show that ~6 nm nanoparticles, coated with a shell of hydrophobic and anionic ligands regularly arranged in ribbon-like domains of alternating composition (Fig. 1a), penetrate cell membranes at 37°C and 4°C without evidence of membrane disruption. Particles with identical hydrophobic content but lacking structural order in the ligand shell for the most part do not penetrate cell membranes; hence, we conclude that the
structural organization of surface chemical groups plays a key role in regulating cell-membrane penetration.

To create amphiphilic nanoparticles, we chose gold nanoparticles protected by a self-assembled monolayer of organic ligands known to determine the particles' interactions with the external environment. These particles can resemble biomolecules and biomolecular assemblies in terms of size and chemical composition and sometimes function (for example, enzymatic activity), but their tightly packed ligand shell lacks the structural flexibility of proteins or peptide assemblies and instead is composed of an ordered layer of surface functional groups. Simple reactions enable the synthesis of these particles coated with single- as well as multicomponent ligand shells, the composition and morphology of which can be easily and precisely controlled. Using the unique physical chemistry offered by this system, we created water-soluble particles with ordered or disordered amphiphilic ligand shells to investigate cell-membrane penetration of nanoparticles as a function of the spatial distribution of chemical groups.

Four types of ligand-coated gold nanoparticle were synthesized and studied (Table 1): particles were coated with (1) 11-mercapto-1-undecanesulphonate (MUS), (2) a 2:1 molar mixture of MUS and 1-octanethiol (OT) (hereafter referred to as 66-34 OT), (3) a 1:2 molar mixture of MUS and OT (34-66 OT) and (4) a 2:1 molar mixture MUS: 3,7 dimethyl octane 1-thiol (br-OT) (66-34 br-OT), respectively. We used sulphonate groups because they imparted remarkable water solubility to all of the particles studied irrespective of the fraction of hydrophobic ligands used, with saturation concentrations in water in excess of 250 mg ml$^{-1}$ (ref. 30), orders of magnitude larger than the concentrations used here. $^1$H NMR was used to confirm the absence of unbound ligands and the particle ligand-shell composition was determined after decomposing the gold core with iodine. The size distributions, ligand-shell packing densities and zeta potentials of the four particles studied had negligible differences (Table 1) (see the Supplementary Information). Thus, these particles had nearly identical physical characteristics, except for the composition and structure of the ligand shell, as determined by extensive scanning tunnelling microscopy (STM) studies (see the Supplementary Information) and schematically shown in Table 1. MUS particles had a homogeneous hydrophilic ligand shell, whereas 34-66 OT and 66-34 OT particles had a hydrophilic–hydrophobic striated

<table>
<thead>
<tr>
<th>Nanoparticles</th>
<th>Ligand shell composition$^*$</th>
<th>Core size$^+$ (nm)</th>
<th>Thermogravimetric analysis$^\ddagger$ (%)</th>
<th>$\zeta$ Potential $^|$ (mV)</th>
<th>Ligand shell morphology/chemical structures</th>
</tr>
</thead>
<tbody>
<tr>
<td>MUS</td>
<td>100% MUS</td>
<td>4.3±1.3</td>
<td>15</td>
<td>$-38.0\pm5.30$</td>
<td>Homogeneous</td>
</tr>
<tr>
<td>66-34 br-OT</td>
<td>67% MUS</td>
<td>4.3±1.2</td>
<td>13</td>
<td>$-31.1\pm0.73$</td>
<td>Unstructured</td>
</tr>
<tr>
<td>66-34 OT</td>
<td>66% MUS</td>
<td>4.5±1.0</td>
<td>15</td>
<td>$-33.1\pm0.64$</td>
<td>Structured</td>
</tr>
<tr>
<td>34-66 OT</td>
<td>33% MUS</td>
<td>4.9±0.9</td>
<td>11</td>
<td>$-35.2\pm1.49$</td>
<td>Structured</td>
</tr>
</tbody>
</table>

$^*$Calculated from $^1$H NMR analysis after decomposition of the core, data are all ±5%.
$^+$Determined from TEM images and expressed as average diameter ± one standard deviation.
$^\ddagger$Weight loss as determined by thermogravimetric analysis after complete desorption of the organic ligands, which is directly related to the ligand-shell packing density, data are all ±5%.
$^\|$Zeta potential measurements are in serum-free medium.
ligand shell; 66-34 br-OT particles had a hydrophilic–hydrophobic disordered ligand shell. The last two particles had nearly identical hydrophilic to hydrophobic ratios, the key difference residing in the ordering of hydrophilic and hydrophobic moieties of the ligand shell, that is, they can be considered as nanoscale ‘isomers’.

To determine how the nature and organization of functional groups at the surface of these nanoparticles influenced their interactions with cell membranes, we labelled the particles with an average upper limit of 14 molecules of thiolated BODIPY dye (BODIPY-SH) per nanoparticle (see the Supplementary Information) and carried out confocal fluorescence microscopy studies of their uptake and intracellular distribution in living cells. A mouse dendritic cell clone DC2.4 (ref. 33) was used for these studies, representing a key class of phagocytes guarding all peripheral tissues and particularly the skin, lungs and gut mucosa. Dendritic cells are professional antigen-presenting cells, and actively sample particles/fluid/macromolecules from their environment (through phagocytosis, macropinocytosis and endocytosis) as part of their normal function. Macropinocytosis, an actin-dependent process where the cell folds membrane around a small volume of fluid, enables these cells to even internalize particles that have no interaction with the cell membrane into intracellular vesicles. To avoid potential artefacts of cell fixation34, cells were imaged live in a temperature-controlled environmental chamber. Dendritic cells were incubated with nanoparticles (0.2 mg ml\(^{-1}\)) in serum-containing or serum-free medium for 3 h, washed and imaged live by confocal microscopy. Figure 1 shows the difference in behaviour between the various types of nanoparticle. MUS nanoparticles bearing only the hydrophilic sulphonate ligand were internalized by dendritic cells at 37 °C and exhibited punctate fluorescence patterns indicative of endosomal uptake (Fig. 1b), in agreement with previous findings for most types of synthetic nanoparticle21,35,36. 66-34 br-OT nanoparticles bearing a disordered mixed monolayer with sulphonate and methyl headgroups also showed a punctate distribution in dendritic cells, although a low level of background fluorescence in the cytosol of some cells was also detectable (Fig. 1c). In contrast, 66-34 OT nanoparticles, which have a similar ratio of surface sulphonate and methyl groups with respect to 66-34 br-OT nanoparticles but in a ‘striped’ nanoscale organization, were detected in cells as a diffuse pattern of intracellular fluorescence clearly overlaid on punctate sites of brighter fluorescence due to endocytosis, suggesting passage of a fraction of these particles into the cytosol (Fig. 1d). We believe, as discussed below, that...
these particles enter cells through two coexisting mechanisms, one energy dependent (endocytosis) and the other energy independent. A similar behaviour was observed for 34-66 OT nanoparticles (see Supplementary Information, Fig. S3). Trends in nanoparticle uptake and intracellular distribution were qualitatively similar in serum-free and serum-containing medium, although a greater amount of cytosol-localized nanoparticles and fewer nanoparticles contained in vesicular structures were readily observed for the serum-free condition (see Supplementary Information, Fig. S4). The similarity in the nanoparticle uptake in serum-containing and serum-free conditions suggests 'striped' particles are resistant to non-specific protein adsorption (as previously observed in similar systems)\(^5\), and hence retain their behaviour independent of the incubation medium. To confirm this assumption, we used dynamic light scattering to evaluate particle sizes in solution before and after incubation in serum-containing medium. The radii obtained after incubation in serum-containing medium were all in good agreement with transmission electron microscope (TEM) core size data, further confirming that the particles were well dispersed in solution. The particles were then incubated in serum-containing medium and separated from unbound proteins using gel filtration chromatography. The gel filtration chromatography traces showed that a large fraction of the 66-34 OT nanoparticles retained their original size after the incubation, whereas the 66-34 br-OT and the MUS particles showed an increase in particle size (data not shown). In fact, the dynamic light scattering data of the 'purified' particles following serum exposure showed an increase in size for the non-striped particles and almost no change for the 'striped' ones. The increase in size for the non-striped particles must be ascribed to non-specific protein adsorption, as TEM studies after incubation show nearly identical core size distribution (see Table 1, Supplementary Information).

To further explore the mechanism by which these particles entered the cells, we repeated these internalization experiments at 4°C to inhibit mechanisms of active uptake by the cells. At this temperature, endocytic/pinocytic uptake of tracer molecules such as calcein or labelled dextran is completely blocked in DC2.4 cells\(^16\). BODIPY channel fluorescence following the incubation of cells with MUS or 66-34 br-OT particles at 4°C (Fig. 1e and f, respectively) was indistinguishable from the low levels of autofluorescence observed within untreated control cells (see Supplementary Information, Fig. S5a), whereas 66-34 OT (Fig. 1g) and 34-66 OT nanoparticles (see Supplementary Information, Fig. S5e) still seemed to enter cells at substantial levels, taking on a diffuse pattern of fluorescence consistent with cytosolic localization. A quantitative comparison of cell-associated fluorescence at 4°C is shown in Fig. 1h. The fluorescence of cells incubated with MUS or 66-34 br-OT particles was not statistically different from the background autofluorescence of untreated cells, whereas the fluorescence of cells treated with 'striped' nanoparticles was eightfold greater than the background. All of the nanoparticles studied here were uniformly excluded from the nucleus of cells at both 4°C and 37°C. The entry of these particles into the cytosol under conditions where endocytic processes are blocked suggests that the 'striped' nanoparticles are capable of directly passing through the plasma membrane of the cells, whereas 66-34 br-OT nanoparticles with an identical ratio of hydrophilic/hydrophobic moieties but lacking the ordered arrangement of these groups do not penetrate the cell membrane. To further test this possibility and rule out potential changes in membrane structure/fluidity induced by incubating the cells at 4°C, we incubated dendritic cells with nanoparticles at 37°C in the presence of sodium azide and 2-deoxyglucose, which block active forms of uptake such as endocytosis. To confirm the efficacy of the inhibitors, dendritic cells were treated with the inhibitors and incubated with calcein, a small-molecule membrane-impermeable fluorophore normally taken up by endocytosis/pinocytosis. After 20 min, control dendritic cells cultured with calcein in the absence of azide/deoxyglucose exhibited readily detectable endocytotic vesicles of calcein throughout confocal z sections (Fig. 2a). In contrast, dendritic cells incubated with the inhibitors and calcein showed no detectable internalized dye in three-dimensional confocal z sections (data not shown). When dendritic cells were incubated with both calcein and 'striped' OT:MUS particles at 37°C in the presence of the internalization inhibitors, nanoparticle entry into the cytosol was readily detected after 20 min incubation with the particles, whereas calcein was undetectable in the cells (Fig. 2b). MUS particles and 66-34 br-OT nanoparticles failed to enter dendritic cells under these conditions (data not shown). Thus, nanoparticles with an ordered surface ligand layer were found to access the cytosol at 37°C under conditions where no evidence for active uptake processes such as endocytosis could be detected. Finally, consistent with an internalization mechanism involving direct interaction with the plasma membrane rather than a slower endocytic/pinocytic active uptake route, kinetic experiments showed that 'striped' nanoparticles were readily detectable in cells within 5 min (in the presence of azide/deoxyglucose (see the Supplementary Information). Two limitations of the confocal analysis are the resolution of optical microscopy and the possibility of artefacts in the apparent subcellular distribution arising from the release of free dye from particles. Previous studies have shown that organic thiol ligands can be displaced from gold nanoparticles by exchange reactions with cytosolic glutathione\(^3\), which is the most abundant intracellular thiol\(^4\). This effect seems unlikely given that glutathione-mediated displacement has been shown to require timescales of the order of days\(^8\), compared with the present experiments carried out over 3 h. However, to unequivocally interpret the confocal images and also to rule out any possible effect of the fluorophore molecules attached to the nanoparticles, we carried out an extensive TEM analysis of both fluorescently labelled and unlabelled nanoparticles taken up in cells. DC2.4 cells were incubated for 3 h with nanoparticles as before, washed, fixed and sectioned for TEM imaging (see the Supplementary Information). Several observations resulted from analysis of TEM images. No significant difference in overall trends was observed for particles either labelled with or without BODIPY-SH (see Supplementary Information, Fig. S14). The intracellular distribution of each of the four types of ligand-coated nanoparticle studied was in qualitative agreement with the trends observed by confocal microscopy, as illustrated by the series of TEM images in Fig. 3 for the case of unlabelled particles. MUS nanoparticles were almost exclusively found in endosomal compartments (Fig. 3a); a similar distribution was observed for 66-34 br-OT nanoparticles (Fig. 3b). 'Striped' nanoparticles, in contrast, were readily detected outside defined membrane-enclosed compartments (Fig. 3c). As shown in Fig. 3d, an enumeration of the number of free particles per μm\(^3\) localized outside membrane-enclosed compartments revealed a higher density of cytosolic 66-34 and 34-66 OT (see Supplementary Information, Fig. S15) nanoparticles as compared with the MUS and 66-34 br-OT particles. Interestingly, TEM images and examination of the number of cytosol-localized particles per μm\(^3\) obtained from the serum-containing condition showed similar trends, suggesting that the differences in particle behaviour are not mediated by differentially adsorbed serum proteins\(^6\) (see Supplementary Information, Fig. S13). The high-resolution TEM images in Fig. 4 show nanoparticles very close to the cell membrane, probably frozen 'in the act' of passing through it.

Previous studies have shown that some cationic nanoparticles and macromolecules penetrate the plasma membrane of cells by
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Figure 2 ‘Striped’ nanoparticles enter cells at 37 °C under conditions where active internalization processes are blocked. a,b, dendritic cells were incubated for 20 min at 37 °C in serum-free medium with calcein (a) or calcein and 66-34 OT nanoparticles in the presence of azide and 2-deoxyglucose to block active internalization (b). Confocal optical z sections in 2 µm intervals taken through the midplane of cells are shown. The upper panels show nanoparticle fluorescence (red), the middle panels show calcein fluorescence (green) and the lower panels show fluorescence/brightfield overlays. Scale bars: 10 µm.

generating transient holes10; in contrast, some data suggest that a subset of biological membrane-penetrating macromolecules (for example, CPPs) may be capable of penetrating membranes without causing overt membrane poration12–15. The former mechanism can lead to cell death due to loss of membrane polarization and/or leakage of ions and molecules into/out of the cell. To establish whether ‘striped’ nanoparticles penetrate cell membranes through creation of transient holes, we tested whether the internalization of ligand-protected nanoparticles was accompanied by escape of cytosol-localized tracer dye or conversely, cytosolic entry of an initially extracellular tracer dye41,42. First, DC2.4 cells were co-incubated with nanoparticles and calcein; calcein remains in endolysosomal compartments in cells unless it is co-internalized with a membrane-disrupting agent43. Cells were incubated for 3 h with calcein and nanoparticles, washed and imaged live by confocal microscopy; the localization of calcein in individual cells was then scored (endosomal versus cytosolic). As a positive control, dendritic cells were incubated with cationic gold nanoparticles bearing 11-mercaptoundecane–tetramethylammonium chloride (TMA) ligands, which would be expected to disrupt cell membranes on the basis of previously published work10. As shown in Fig. 5a,c, TMA nanoparticles caused calcein to escape to the cytosol of ∼50% of the dendritic cells, whereas dendritic cells incubated with calcein alone exhibited a solely punctate endosomal distribution of dye. Strikingly, 66-34 OT (Fig. 5a,b) and 34-66 OT nanoparticles (see Supplementary Information, Fig. S6) (both fluorescently labelled and unlabelled) did not promote calcein entry into the cytosol, despite the data described above showing that these particles clearly access the cytosolic compartment. We next tested whether ‘striped’ nanoparticles caused an escape of tracer molecules from the cytosol of cells during uptake: dendritic cells were pre-loaded with calcein-AM, an acetomethoxy form of the dye that is trapped in the cytosol by conversion from a membrane-permeable to membrane-impermeable form by intracellular esterases. Again, 66-34 OT particles entered cells without any measurable leakage of calcein from the cells (see Supplementary Information, Fig. S7). To further, albeit indirectly, confirm that ‘striated’ particles penetrate cell membranes without causing substantial
Figure 3 Nanoparticles with sulphonate-only or disordered sulphonate/methyl-group surfaces are largely confined to endosomal compartments, whereas ‘striped’ sulphonate/methyl-group-bearing nanoparticles enter the cytosol. a–c, TEM images of DC2.4 cells following incubation with MUS (a), 66-34 br-OT (b) or 66-34 OT (c) nanoparticles for 3 h in serum-free medium. Free nanoparticles are denoted by arrows. d, Number of cytosolic nanoparticles per µm³ (±s.d.), determined by scoring the localization of >6,500 particles per condition. * Statistically different from MUS (P < 0.005). ** Statistically different from 66-34 br-OT (P < 0.005).

Figure 4 ‘Striped’ particles visualized at different stages of crossing the cell membrane of dendritic cells. a–f, TEM images of 66-34 OT (a–c) and 34-66 OT (d–f) nanoparticles incubated with cells in serum-free medium at 37 °C. The arrows indicate a few of the particles seen intracellular to the cell membrane.
membrane disruption, we measured their cytotoxicity relative to cationic TMA-coated particles using a colony-forming assay. DC2.4 cells were incubated with nanoparticles for 3 h, washed, replated at equal densities and allowed to grow for 24 h. We found that cells treated with 66-34 and 34-66 OT nanoparticles survived/proliferated at a level that was ~90% of that of untreated control cells, whereas only 34% were recovered following treatment with TMA-coated particles (see Supplementary Information, Fig. S10). Similar results were obtained when studying immediate toxicity of cells with propidium iodide staining and MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide) assay (see Supplementary Information, Figs S11,S12).

In these studies, we consistently observed no nanoparticle nuclear uptake, as well as no evidence for escape of nanoparticles from cells for up to 6 h after internalization occurred (data not shown). We speculate that these two phenomena could be related; it is possible that when particles are in the cytosol they lose the delicate compositional balance and structural motif that is needed to permeate cell membranes. However, given the unique structure of the nuclear double membrane, further studies need to be carried out to fully understand these observations. Moreover, to test the generality of our results, we repeated 37 °C and 4 °C uptake experiments with a second common cell line, mouse embryonic fibroblasts. The same trends in particle uptake, with ‘striped’ particles accessing the cytosol, were observed (see Supplementary Information, Fig. S9).

Our results highlight the importance of structure in the interactions of nanoparticles with cell membranes. We synthesized two nanoparticles that could be loosely characterized as the nanoscale equivalent of chemical isomers. They have the same size and shape, and are coated with the same ratio of hydrophobic to hydrophilic molecules, the latter being identical (MUS) whereas the former differing slightly (OT versus br-OT) to achieve different ligand shell structures. These two particles (66-34 OT and 66-34 br-OT) showed very similar solubility and wetting behaviour as highlighted by nearly identical zeta potentials. They differed only in the molecular arrangement of surface chemical groups: 66-34 OT exhibiting a striated ligand organization and 66-34 br-OT lacking this structured arrangement. This seemingly small difference led to drastically different behaviour in cell-membrane-penetration properties. Confocal studies of cell uptake of these particles (dye functionalized to the same extent) in DC2.4 cells at 37 °C (in both serum-containing and serum-free conditions) showed a pronounced punctate fluorescence distribution pattern indicative of endosomal entrapment for 66-34 br-OT nanoparticles, whereas 66-34 OT particles exhibited a more diffuse fluorescence pattern, indicating escape to the cytosol. These differences in intracellular localization were confirmed through TEM studies, which were also used to rule out any effect of the presence of the dye in the ligand shell. Nanoparticle uptake experiments at 4 °C were used to prove that the 66-34 OT nanoparticles can directly pass through cell membranes and do not require internalization by active endocytosis or pinocytosis to reach the cytosol, whereas at 4 °C 66-34 br-OT nanoparticles are almost completely blocked from cell entry. This was also confirmed by detection of nanoparticle entry into cells within 5 min despite actively blocking endocytosis by treating the cells with sodium azide and 2-deoxyglucose (marked by no calcein uptake). Finally, we determined that 66-34 OT nanoparticles pass through cell membranes without causing overt membrane poration through studies with tracer dyes (no noticeable leakage) and cytotoxicity assays (minimal toxicity). This is in stark contrast with cationic nanoparticles, which are known to penetrate cell membranes by creating pores18 and induce cytotoxicity19. At the moment we can only speculate on the mechanism that leads to cell-membrane penetration. A possible explanation is that the ‘rigid’ arrangement of the amphiphilic domains on the nanoparticles permits non-disruptive fusion of the nanoparticle with cell membranes (a more fluid mixed layer) and subsequent penetration through the bilayer, in a manner reminiscent of some models for entry of CPPs.

In summary, we believe that by highlighting the role of ligand structure and order in cell-membrane penetration by nanoparticles, this work will shed new light on the understanding of the molecular interactions that lead to cell-membrane permeation, certainly on nanoparticles, possibly on larger organic particles and hopefully in CPPs. Future work, besides the fundamental studies highlighted above, includes the engineering of novel and more efficient low-toxicity drug-delivery systems based on the design rules for membrane penetration identified here and on the analysis of potential safety issues associated with synthetic nanomaterials44.

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Author contributions
D.I.I. and F.S. conceived the research question. A.V., O.U., D.I.I. and F.S. designed the experiments and corresponding authors. TPEN program (U01-HL080731). F.S. is grateful to the Packard Foundation for their generous award.

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In the version of this Article originally published, in the caption for Fig. 1 the following statement should have been included "Right-hand STM image in panel a reproduced with permission from ref. 30, © 2008 RSC." This error has been corrected in the PDF and HTML versions of the Article.