Labeling and exocytosis of secretory compartments in RBL mastocytes by polystyrene and mesoporous silica nanoparticles

Maneerat Ekkapongpisit 1,∗
Antonino Giovia 1,∗
Giuseppina Nicotra 1
Matteo Ozzano 1
Giuseppe Caputo 2,3
Ciro Isidoro 1

1Laboratory of Molecular Pathology and Nanobiomaging, Department of Health Sciences, Università del Piemonte Orientale “A. Avogadro”, Novara, Italy; 2Department of Chemistry, University of Turin, Turin, Italy; 3Cyanine Technology SpA, Torino, Italy

*These authors contributed equally to this work

Background: For a safe ‘in vivo’ biomedical utilization of nanoparticles, it is essential to assess not only biocompatibility, but also the potential to trigger unwanted side effects at both cellular and tissue levels. Mastocytes (cells having secretory granules containing cytokines, vasoactive amine, and proteases) play a pivotal role in the immune and inflammatory responses against exogenous toxins. Mastocytes are also recruited in the tumor stroma and are involved in tumor vascularization and growth.

Aim and methods: In this work, mastocyte-like rat basophilic leukemia (RBL) cells were used to investigate whether carboxyl-modified 30 nm polystyrene (PS) nanoparticles (NPs) and naked mesoporous silica (MPS) 10 nm NPs are able to label the secretory inflammatory granules, and possibly induce exocytosis of these granules. Uptake, cellular retention and localization of fluorescent NPs were analyzed by cytofluorometry and microscope imaging.

Results: Our findings were that: (1) secretory granules of mastocytes are accessible by NPs via endocytosis; (2) PS and MPS silica NPs label two distinct subpopulations of inflammatory granules in RBL mastocytes; and (3) PS NPs induce calcium-dependent exocytosis of inflammatory granules.

Conclusion: These findings highlight the value of NPs for live imaging of inflammatory processes, and also have important implications for the clinical use of PS-based NPs, due to their potential to trigger the unwanted activation of mastocytes.

Keywords: secretory lysosomes, inflammation, nanoparticles, vesicular traffic

Introduction
Nanoparticle labeling is a rapidly emerging area in the field of nanotheranostics, ie, the application of nanotechnology to the areas of medical therapy and diagnosis by creating powerful tools enabling the targeted delivery of drug therapies and imaging tracers. The in vivo biomedical application of such NPs however, raises several issues regarding their tolerability, intracellular effects, systemic reactions and overall toxicity. The extent of NP toxicity essentially depends on such variables as size, type of material, and surface charge, and is proportional to the dose and time of exposure, as well as the actual concentration of NPs within critical cellular compartments. At the cellular level, the NPs’ toxicity may result from the generation of free radicals, or the disruption of organelle integrity (eg, by inducing leakage of proapoptotic molecules from lysosomes or mitochondria), or by affecting the plasma membrane permeability (eg, by inducing calcium ion entry).

At the tissue and organ levels, toxicity may arise following NP-induced stimulation of the immune and inflammatory cells. This tissue immune-inflammatory reaction

Correspondence: Ciro Isidoro
Università del Piemonte Orientale
“A. Avogadro”, Dipartimento di Scienze della Salute, Via Solaroli 17, 28100 Novara, Italy
Tel +39 321 660607
Fax +39 321 620421
Email isidoro@med.unipmn.it

This article was published in the following Dove Press journal:
International Journal of Nanomedicine
3 April 2012
Number of times this article has been viewed

http://dx.doi.org/10.2147/IJN.S29034

International Journal of Nanomedicine 2012:7 1829–1840
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is orchestrated by resident mastocytes, through the release of vasoactive amines, cytokines, and proteolytic enzymes.\textsuperscript{15,16} These substances are packaged within lysosome-like granules that translocate to the periphery of the mastocyte, and fuse with the plasma membrane upon stimulation of IgE receptors or other stimuli effectively inducing an influx of calcium ions.\textsuperscript{77}

Mastocytes are involved not only in allergic and inflammatory diseases, but also play a pivotal role in tumor vascularization and growth.\textsuperscript{18–20} NPs with the ability to label and trace the exocytosis of inflammatory secretory granules would therefore be useful to monitor mastocyte activation during an inflammatory response. In addition, given the desired utilization of NPs in the detection of cancer and its therapy, it is also prudent to explore the possibility of being able to detect and monitor the unwanted activation of mastocytes infiltrating the tumor stroma. In this work each of these issues was addressed.

As a ‘sensitive’ cell model we utilized mastocyte-like rat basophilic leukemic (RBL) cells, a widely used ‘in vitro’ cell model to study the inflammatory response to allergic stimuli.\textsuperscript{21} RBL cells possess two types of inflammatory granules that share the same biogenetic pathway with lysosomes.\textsuperscript{22,23} Upon stimulation, a rapid influx of calcium ions enables the fusion of these organelles with the plasma membrane, and the release in the extracellular environment of lysosomal proteases, vasoactive substances and inflammatory cytokines.\textsuperscript{23,24} The RBL cell model has already been successfully used in studies of endocytosis and cellular labeling with NPs.\textsuperscript{25–27} RBL cells represent a valuable cellular model system to address the two questions posed in the present study: (1) do NPs trigger an inflammatory response, ie, the calcium-dependent secretion of mastocyte granules? And (2) can NPs be used to label, and possibly discriminate between different internal secretory, lysosome-like compartments of mastocytes?

To monitor the active exocytosis of inflammatory granules in RBL cells, two types of fluorescent NPs were employed that differ in chemical composition, surface charge, and size: (1) mesoporous silica (MPS) material, of 10 ± 5 nm in diameter, with no surface-charged functional group, and doped with IRIS-3 dye, emitting red fluorescence; and (2) polystyrene (PS) material, of 30 ± 10 nm in diameter, carboxyl-modified with a negative surface charge, and embedded with the FITC dye, emitting green fluorescence. Mesoporous, silica-based NPs are emerging as valuable nanocarriers of therapeutic and diagnostic tracers due to their unique pore architecture, which enables them to host macromolecules such as complex dyes and pharmaceuticals, with the added bonus that silica is considered to be nontoxic and biodegradable.\textsuperscript{28–31} The physical and chemical characteristics of the MPS NPs used in the present study have been previously described.\textsuperscript{27} PS NPs are also being evaluated for drug delivery and cellular imaging.\textsuperscript{32–34} The potential of MPS NPs to induce the release of inflammatory cytokines by macrophages has been recently investigated.\textsuperscript{30} Similarly, in a recent study, amorphous silica NPs were shown to induce the release of inflammatory cytokines by endothelial cells.\textsuperscript{35}

Whether and how MPS and PS NPs interact with and activate mastocytes is not known. We addressed these issues in cultured RBL mastocytes. The main findings here reported are: (1) MPS and PS NPs showed different efficiency of intracellular retention: while the former were well tolerated for long time, the latter were promptly expelled in a calcium-dependent fashion; and (2) MPS and PS NPs engaged in different endocytic routes and finally accumulated in different secretory compartments. These data present significant clinical applications, since they demonstrate a powerful ability of NPs to monitor the inflammatory process via imaging the degranulation of mastocytes. In addition, these data also have important clinical implications as they demonstrate the possibility of inducing the activation of an inflammatory reaction by NPs.

**Materials and methods**

**Nanoparticles**

Commercial carboxyl-functionalized, fluorescein isothiocyanate (FITC)-conjugated PS NPs were purchased from Sigma (cod L5155; Sigma-Aldrich, St Louis, MO). Mesoporous silica (MPS) NPs engrafted with IRIS-3 were provided by Cyanine Technologies (IRIS3-Dots-Porous cod 3 WEL-06; Cyanine Technologies SpA, Turin, Italy). NPs were dissolved in culture medium after conglomerate disruption by sonication.\textsuperscript{27}

**Cell culture and treatments**

Mastocyte-like RBL-2H3 cells (American Type Culture Collection, Manassas, VA) were cultivated at 37°C in 5% CO\textsubscript{2} in Dulbecco’s modified Eagle’s minimum essential medium (cod D5671; Sigma-Aldrich) supplemented with 16% fetal bovine serum (FBS) (cod DE14-801F; Lonza Group Ltd, Basel, Switzerland) and 100 IU/mL penicillin–streptomycin (cod P0781; Sigma-Aldrich). Typically, the cells were plated on sterile cover-slips and left to adhere for 24 hours before use. Incubations were performed in fresh medium for the time indicated. NPs were dissolved directly in culture medium
at the desired final concentration. The calcium ionophore A23187 and the calcium-chelating ligand EGTA (both from Sigma Aldrich) were used at the concentration of 1 µM and 5 mM, respectively. Methyl-β-cyclodextrin (MbCD) (cod C4805; Sigma-Aldrich) was used at 5 mM final concentration. Unless otherwise specified, all experiments were performed in the presence of FBS, a medium that closely resembles the physiological tissue environment.

**Cellular viability assessment**

Cell toxicity was evaluated by looking at the gross morphology alterations of the nuclei typically associated with cell death (necrosis, chromatin condensation and fragmentation), and at the metabolic activity of the cell. To this end, nuclei were stained with DAPI and cells were labeled with CellTracker™ (Invitrogen, Carlsbad, CA). The latter is a fluorescent dye that emits blue fluorescence of an intensity proportional to that of mitochondrial respiratory activity. Adherent cells prelabeled with DAPI (not shown) or CellTracker™ were each incubated with NPs for up to 48 hours. It was assumed that, given the rapid internalization of NPs, any toxic effect on cell metabolism would be evident by this time. As a control for metabolic toxicity, the cells were incubated with 1% dimethylsulfoxide (DMSO). Cell viability was measured as the percentage of cells labeled with CellTracker™ irrespective of intensity, as determined with ImageJ software (US National Institutes of Health, Bethesda, MD).

**Cathepsin D fluorescent chimeras and plasmid transfection**

The cDNA coding for human lysosomal cathepsin D,⁴⁻⁷ devoid of the stop codon, was subcloned in the multiple cloning site of the plasmids peGFP-N1 or pmCherry-N1 (both from Clontech Lab, Takara Bio Inc, Shiga Japan) in order to drive the synthesis of the fluorescent chimeras CD-GFP or CD-mCherry. Adherent RBL cells were transfected with the plasmid using the Lipofectamine 2000 Reagent™ (Life Technologies Ltd, Paisley, UK) method as suggested by the manufacturer’s instructions.

Briefly, cells were plated in a P35 petri dish at 15,000/cm² and left to adhere for 24 hours prior to the transfection procedure. The DNA-Lipofectamine complexes were prepared in 500 µL of Opti-MEM I reduced serum medium (Life Technologies Ltd) with 5 µg of plasmid and 10 µL of lipofectamine. After 6 hours of incubation, the transfection medium was removed and replaced with serum-containing culture medium (16% FBS-DMEM) and the cells cultivated for 36 hours to allow for maximal protein expression. The cells were then incubated in fresh medium with either the LysoTracker™ (Invitrogen–Molecular Probes, Eugene, OR) or the NP for the time indicated.

**Endocytosis and exocytosis studies**

RBL cells were plated on sterile cover-slips and left to adhere for 24 hours, then the medium was replaced, and cells were incubated with the NPs for the time indicated. To trace the endocytic pathway, the cells were prelabeled for 10 minutes with LysoTracker™ Green or Red. Where indicated, 5 mM of MbCD was added for 1 hour prior to the incubation with NPs in order to extract cholesterol from the plasma membrane. Exocytosis of inflammatory granules was induced by stimulation of calcium entry by A23187 (1 µM for 5 or 60 minutes), and inhibition of exocytosis was attained by supplementing the medium with 5 mM EGTA.

**Fluorescence microscope imaging**

Samples were observed with the fluorescence microscope Leica DMi6000 (Leica Microsystems AG, Wetzlar, Germany).

Five to ten fields (for a minimum total of 50 cells) were randomly imaged by two independent investigators (ME and AG), and representative images were chosen for display. Images were captured under the fluorescence microscope using the same color intensity threshold for all treatments. All images were compiled using Adobe® Photoshop® (Adobe Systems Inc, San Jose, CA), with the same contrast adjustment applied to all images. Quantification of fluorescent microscope images was performed using ImageJ software. This method enabled measurement of the mean quantity of NPs which had accumulated in the cell as the mean fluorescence intensity per cell.

**Flow cytometry**

Flow cytometry was employed to assess the proportion of cells that had internalized the NPs. The cells were plated on petri dishes and left to adhere for 24 hours, then the NPs were added to fresh medium. At the end of the incubation period, the cells were washed three times with PBS to remove the excess NPs, and extracellular fluorescence was quenched with 0.4% (w/v) trypan blue in PBS. Cells were then harvested with trypsin, resuspended in PBS and the extent of NP internalization was then assessed using a FACScan flow cytometer (Becton, Dickinson and Company, Franklin Lakes, NJ). Cell-associated fluorescence (10,000 cells/sample) was analyzed with WinMDI software (version 2.9; Microsoft Corporation, Redmond, WA).
Statistics
Each experiment was performed in triplicate and reproduced two or more times. Quantification data from ImageJ and cytofluorometry analyses were given as average ± standard deviation. The Student’s t-test with $P < 0.05$ for statistical significance was employed to compare the results from different treatments.

Results and discussion
Biocompatibility and dose/time-dependent uptake of MPS and PS NPs
The uptake efficiency and intracellular end-point of NPs together with their effects on cellular metabolism, are greatly influenced by their physical-chemical characteristics.

Since NP toxicity is also cell-type specific, it was first necessary to establish that MPS and PS NPs do not exert any toxicity upon RBL cells. Biocompatibility was tested in RBL cultures exposed for up to 48 hours to either type of NP at the maximal concentration used in the present study (ie, 30 µg/mL and 75 µg/mL, respectively). Cell viability was assayed using a metabolic thiol-reactive probe (CellTracker™) that produces a stable membrane-impermeable glutathione–fluorescent dye adduct in living cells. The images in Figure 1 demonstrate that both MPS and PS NPs were nontoxic to RBL cells, which is consistent with previous reports showing the biotolerability of MPS and of carboxylated PS NPs in other cell types.9,27–29,33,36,37 From the fluorescent signal, it was established that MPS NPs accumulated in large quantities in RBL cells, indicating their efficient uptake and intracellular retention throughout the incubation period. On the other hand, only a few RBL cells appeared to contain PS NPs, and then only in very small amounts (Figure 1). The latter finding could be explained by assuming that: (1) PS NPs hardly entered RBL cells (at the given concentration and time of incubation); or (2) PS-associated fluorescence was rapidly and fully quenched within cellular compartments; or (3) PS NPs were expelled after their internalization.

Although the PS material can be considered to be chemically relatively inert, the presence of carboxyl functional groups on the surface confers a negative surface charge to the NP, which allows (reversible) electrostatic interactions with biological molecules that may impact on the uptake efficiency.33,38,39 A typical time- and concentration-dependent internalization experiment showed that the accumulation of PS NPs in RBL cells clearly increased with concentration, while increasing the time of incubation did not lead to a substantial increase of the number of labeled cells, or of NP per cell (Figure 2A and B).

On the other hand, MPS NPs were very rapidly taken up by RBL cells, and saturated intracellular compartments in a dose-dependent fashion (Figure 2C). The kinetics of the uptake of MPS and PS NPs were directly compared. Flow cytometry was used to determine the proportion of cells that had internalized the NPs at a given time. Prior to flow cytometry analysis, at the end of each time-point incubation, the samples were extensively washed and quenched with trypan blue.

Figure 1 Biotolerability of NPs. RBL cells (15,000/cm²) adherent on cover-slips were incubated with 30 µg of MPS or 75 µg of PS NPs in fresh medium for 48 hours, then the cells were washed and labeled with CellTracker™ (5 µM for 45 min in serum-free medium). After this, the cells were washed and incubated in regular complete medium for 30 minutes. Positive control of toxicity was performed incubating the cells with 1% DMSO. Cells were then washed and imaged under the fluorescence microscope. The proportion of viable (CellTracker™-stained) cells was determined with the aid of ImageJ software. Representative images (A) and quantification (B) of two independent experiments performed in triplicate are shown. Note that almost the whole cell population incubated with either MPS or PS NPs shows intense blue fluorescence, whereas in DMSO-treated culture, most of the monolayer has detached and, of the portion still adherent, approximately 25% of the cells appear faintly labeled with CellTracker™.

Abbreviations: DMSO, dimethyl sulfoxide; MPS, mesoporous silica; NPs, nanoparticles; RBL, rat basophilic leukemia; PS, polystyrene.
Phase contrast/MPS

Phase contrast/PS

5 min

30 min

30 min 5 min

25 µg

100

50

0

25 µg

50 µg 75 µg 100 µg

25 µg

75 µg 100 µg

A

B

C

Figure 2 Dose- and time-dependent cellular accumulation of NPs. RBL cells (30,000/cm²) adherent on cover-slips were exposed to different concentrations of NPs for two different incubation times (5 and 30 minutes). After this, the cells were washed, imaged under the fluorescence microscope and cell-associated fluorescence was quantified with ImageJ software. (A) Representative images of RBL cells exposed to PS NPs (concentration and time as indicated). (B) Quantification of total fluorescence (normalized per cell); data from two independent experiments in triplicate are shown. (C) Representative images of cells exposed to MPS NPs.

Abbreviations: MPS, mesoporous silica; NPs, nanoparticles; RBL, rat basophilic leukemia; PS, polystyrene.

blue to remove all non-internalized NPs material from the cell surface. An aliquot of washed cells was observed under the microscope to ascertain the absence of fluorescent NPs passively adsorbed on the cell surface. Flow cytometry data were expressed as a percentage of fluorescently-labeled cells. Both types of NPs were taken up by the cells, although at different rates. MPS NPs rapidly entered the cells and after 15 minutes, 100% of the cell population was intensely labeled, while by this time only 35% of the cell population had been labeled by PS NPs (Figure 3).

Carboxylated PS NPs trigger their calcium-dependent expulsion

Internalization of NPs is influenced by ionic interactions between the negative membrane potential of the cell, and the surface net charge of NPs. In addition, negatively-charged 20 nm PS NPs were shown to stimulate the calcium-dependent secretion of chloride and bicarbonate anions in airway epithelial cells.

A possibility existed that the PS–plasma membrane interaction could affect the calcium-dependent recycling of endocytic vesicles, thus leading to a dynamic endocytosis–exocytosis process that could account for the low intracellular retention of carboxylated PS NPs. RBL cells were allowed to internalize PS NPs, then the cells were washed and incubated for increasing time of chase. In parallel cultures, the extracellular calcium chelator EGTA was present throughout the uptake and chase incubation. The extent of cell labeling was assessed by fluorescence microscopy and

Figure 3 Kinetics of MPS and PS NPs internalization. RBL cells (80,000/well) adherent on 12-well petri dishes were incubated in fresh medium with 10 µg of MPS or 75 µg of PS NPs for the time indicated. At each time point, the cells were washed, harvested and cell-associated fluorescence quantified by flow cytometry.

Note: Data from two independent experiments each performed in triplicate are shown.

Abbreviations: MPS, mesoporous silica; NPs, nanoparticles; RBL, rat basophilic leukemia; PS, polystyrene.
cytofluorometry. Data shown in Figure 4 demonstrate that EGTA contrasted the time-dependent loss of intracellular labeling, suggesting a role for calcium influx in the rapid expulsion of PS NPs.

**MPS and PS NPs label calcium-regulated secretory compartments in RBL cells**

In most cell types studied thus far, endocytosed material is first delivered to early endosomes, and is then further transported downstream to endocytic compartments toward late endosomes and lysosomes. RBL mastocytes have been shown to possess two distinct subpopulations of secretory lysosome-like compartments. It is not yet known whether these compartments interconnect the endocytic pathway, and if so, how. To address this, calcium-dependent degranulation of RBL cells previously loaded with NPs was induced. Degranulation was instigated by imposing the influx of extracellular calcium with the calcium ionophore A23817, and parallel coincubation with the extracellular calcium-chelating ligand EGTA serving as the control.

RBL cells were preloaded for 15 minutes with PS NPs, then washed and incubated for 5 or 60 minutes, with or without A23187 or EGTA, or both. The images in Figure 5A show that PS NPs are still visible in RBL cells at 5 minutes, but are no longer visible at 60 minutes (which confirms the data in Figure 4), and demonstrate that NPs are rapidly exocytosed upon induction of calcium influx. Note that in the majority of the cells cotreated with A23187 and EGTA, the NPs are retained within the cells, and have accumulated at the periphery of the cells. An analogous experiment was performed in cells preloaded with MPS NPs. In this case, A23187 also triggered the rapid calcium-dependent expulsion of the NPs, (only the 5 minute chase incubation is shown) (Figure 5B).

It is worth noting that in the presence of EGTA, MPS-positive vesicles do not change their position. Flow cytometry quantification of this experiment showed that in 5 minutes, some 80% of the cell-associated MPS signal was lost upon

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**Figure 4** PS NPs induce calcium-regulated exocytosis of PS-containing vesicles. RBL cells (30,000/cm²) adherent on cover-slips were exposed to 75 µg of PS NPs for 15 minutes. Where indicated, the cells were preincubated for 30 minutes with 5 mM EGTA. After this, the cells were washed, imaged under the fluorescence microscope and cell-associated fluorescence was quantified with ImageJ software. (A) Representative images of RBL cells exposed to PS NPs in the presence or absence of EGTA. (B) Quantification of total fluorescence (normalized per cell).

**Note:** Data from two independent experiments in triplicate are shown.

**Abbreviations:** EGTA, ethyleneglycoltetraacetic acid; NPs, nanoparticles; RBL, rat basophilic leukemia; PS, polystyrene.
A23187-induced exocytosis, and that this loss was fully inhibited by EGTA (Figure 5C).

**MPS and PS NPs label distinct cellular organelles and only MPS NPs accumulate in true cathepsin D-containing lysosomes**

To see whether endocytosed MPS and PS NPs both converged to the same intracellular secretory compartments, the RBL cells were coincubated with these NPs, and their intracellular traffic and localization was followed with increasing chasing times. MPS NPs stably localized at the periphery of the cell, whereas PS NPs initially localized in perinuclear vesicles, and with time moved toward the periphery, though never merging with MPS-labeled vesicles (Figure 6A). In addition, by 60 minutes, the PS NPs were only faintly detectable in the cell, confirming their continuous expulsion.

It has previously been shown that RBL secretory granules contain cathepsin D, among other lysosome-resident hydrolases. To assess the nature of the organelle labeled by MPS and PS NPs, the secretory organelles were marked by transgenic expression of a fluorescent cathepsin D chimeric protein. We engineered the peGFP- and pmCherry- plasmids placing in front the cDNA coding for Cathepsin D and transfect RBL cells with these plasmids to induce the synthesis of the fluorescent chimeric protein CD-GFP or CD-mCherry.

First, it was ascertained whether the transgenically-expressed fluorescent chimeric cathepsin D protein had indeed reached the acidic compartments. To this end, the latter were labeled with the fluorescent marker LysoTracker™, which is known to trace the vesicles of the endocytic pathway,
and to accumulate in acidic lysosomes. Both chimeric CD-GFP (Figure 7B) and CD-mCherry (not shown) proteins largely colocalized with the LysoTracker™ (on average, 69% of the fluorescent signals overlapped in a vast majority of transfected cells), indicating the convergence of both tracers at lysosome level. Vesicles labeled with LysoTracker™ only likely represent the endocytic compartments upstream of lysosomes, and vesicles labeled with CD-GFP only likely represent the biogenetic compartments upstream of lysosomes. In CD-GFP-expressing cells, MPS NPs were found to have localized into the majority (>90%) of CD-GFP-labeled vesicles, though additional CD-GFP-negative vesicles also were labeled (Figure 6B). On average, up to 75% of MPS NPs appeared not to have localized with CD-GFP in the vast majority of transfected cells. However, when interpreting the absence of colocalization of vesicular tracers with GFP chimeric proteins, it should be taken into account that GFP fluorescence is rapidly and quantitatively quenched in acidic compartments.

Next, we incubated CD-mCherry transfected RBL cells with PS NPs. The image in Figure 6B shows no convergence of PS NPs in vesicles labeled by CD-GFP, suggesting that these NPs were not delivered to true lysosomes. Taken together with data in Figure 5, the present data indicate that MPS and PS NPs each label distinct secretory compartments in RBL cells.

**MPS and PS NPs take different endocytic routes and localize to different intracellular compartments**

The above data suggest that MPS and PS NPs are differently trafficked after endocytosis in RBL cells. To characterize the endocytic routes of MPS and PS NPs, a kinetic study was performed on the concomitant uptake of both NPs along with the endocytic pathway tracer LysoTracker™. To enable an objective evaluation of the labeled vesicles, the two fluorescent signals were quantified as individual or merged spots using ImageJ software. Preliminary experiments indicated that endocytosis and intracellular vesicular redistribution of NPs occurred very rapidly. The cells were preloaded with LysoTracker™ for 10 minutes, to allow the complete labeling of vesicles along the endocytic pathway down to the lysosomes. Cells were then washed and incubated with NPs for increasing labeling time.

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**Figure 6** MPS and PS NPs localize to distinct intracellular compartments. (A) RBL cells (30,000/cm²) adherent on cover-slips were co-incubated for 5 minutes with 30 µg of MPS and 75 µg of PS NPs, then cells were washed and imaged at 0, 15, and 60 minutes of chase. Representative images of two experiments in triplicate are shown. (B) RBL cells (10,000/cm²) adherent on cover-slips were transfected (for 6 hours with lipofectamine in Optimem medium) with the plasmid coding for the chimeric fluorescent protein cathepsin D-GFP or cathepsin D-mCherry as indicated. At 36 hours post-transfection, the cells were incubated for 5 minutes with 50 nM LysoTracker™ Red or with 15 µg of MPS or 75 µg of PS NPs, as indicated. Representative images of three independent experiments are shown.

**Abbreviations:** MPS, mesoporous silica; NPs, nanoparticles; RBL, rat basophilic leukemia; PS, polystyrene.
Figure 7 MPS and PS NPs take different endocytic routes. RBL cells (30,000/cm²) adherent on cover-slips were preincubated for 10 minutes with LysoTracker™ (Green or Red), then cells were washed and incubated with NPs and imaged at 1, 5, and 30 minutes. Cell-associated fluorescence was quantified with ImageJ software. Total fluorescence (normalized per cell) was given as a percentage of the individual or mixed color. Representative images of RBL cells incubated with LysoTracker™ green (100 nM) and 10 µg of MPS NPs or with LysoTracker™ Red (50 nM) and 75 µg of PS NPs are shown in (A) and (B), respectively. The intensity of single or mixed fluorescence spots measured in the above samples is shown in (C).

Notes: Data is given as a percentage of the total. Average ± SD of 100 cells randomly chosen. Data from two independent experiments in triplicate are shown.

Abbreviations: MPS, mesoporous silica; NPs, nanoparticles; RBL, rat basophilic leukemia; PS, polystyrene; SD, standard deviation.

At a designated time-point the cells were washed and rapidly observed under the microscope (note that a lag time of approximately 60–90 seconds elapsed before capturing the images). Five to ten randomly selected fields were imaged for each sample at 1, 5, and 30 minutes of chase. Representative images are shown in Figure 7A (for MPS NPs) and Figure 7C (for PS NPs). Digitalized images were then analyzed for quantification with the ImageJ software.

It is readily apparent from these images that MPS and LysoTracker™ demonstrated almost complete colocalization at 1 and 5 minutes of labeling, and became partially separated by 30 minutes. In particular, it was noted by this time that both the vesicles traversed by newly entered MPS and the vesicles labeled by pre-existing MPS no longer contained LysoTracker™ fluorescence. Quantification of cell-associated fluorescence showed that >70% of the MPS signal had merged with the LysoTracker™ signal at 1 and 5 minutes of incubation, while by 30 minutes only 20% of MPS fluorescence remained merged with the LysoTracker™. This apparent decrease of colocalization could reflect a physical separation, or, more probably, the loss of signal of one of the two tracers. Indeed, the overall fluorescence emission of the LysoTracker™ greatly diminished over the time of incubation (see also Figure 7C). Strikingly, PS NPs colocalized with LysoTracker™ at any time (Figure 7C and D). Together with the above data, it can be concluded that MPS and PS NPs enter the cells through different endocytic routes and reach different final compartments.

MPS and PS NPs exploit different endocytic mechanisms

The observed differences in the internalization and retention rates of MPS and PS NPs may reflect differences in
the uptake mechanisms dictated by the specific physical-chemical properties of the NPs. For instance, the cellular uptake of negatively-charged silica NPs was shown to occur through nonspecific adsorptive endocytosis, while that of COOH-functionalized PS NPs in macrophages was shown to be clathrin-dependent and in HeLa cells, the uptake of PS NPs was shown to occur via clathrin-dependent or clathrin/caveolae-independent mechanisms, depending on the NP size. Plasma membrane cholesterol has been shown to be involved in various cellular uptake mechanisms, including those mediated by clathrin, caveolae, and lipid rafts. These endocytic pathways can be disrupted by the selective extraction of cholesterol from the plasma membrane, imparted by methyl-β-cyclodextran (MbCD). We sought to define whether MPS and PS NPs exploited the same mechanism of cellular entry.

RBL cells were preincubated (or not) for 60 minutes in serum-free medium containing 5 mM MbCD, a condition sufficient to deplete the plasma membrane of cholesterol. The cells were then pulsed for 5 minutes with the NPs, either individually or together, and (after extensive wash) were rapidly observed and imaged under the fluorescence microscope. In the control, serum-containing medium, uptake and internal localization of both NPs resembled the previous findings (Figure 8). It is of particular note that the incubation in serum-deprived medium greatly stimulated the uptake of both NPs, whether incubated individually or together (Figure 8). In the latter case, it was apparent that a limited colocalization of the two types of NPs had occurred in a small fraction of organelles.

These data indicate that: (1) serum proteins can trap NPs in the medium and greatly reduce their uptake by the cells; (2) serum factors may interfere with the preferential endocytic mechanism and trafficking route of NPs. This second indication is indeed intriguing and deserves further ad hoc investigation. NP opsonization by serum proteins is

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**Figure 8** Membrane cholesterol depletion differentially affects endocytosis of MPS and PS NPs. RBL cells (30,000/cm²) adherent on cover-slips were pulse-labeled for 5 minutes with 10 µg of MPS or 75 µg of PS NPs or both in complete or serum-free medium as indicated. Parallel cultures were preincubated for 1 hour, with 5 mM MbCD in serum-free medium used to assess the involvement of cholesterol-dependent mechanisms in the uptake of NPs. After this, cells were washed and imaged under the fluorescence microscope. Representative images of two independent experiments in triplicate are shown.

**Abbreviations:** MbCD, methyl-β-cyclodextrin; MPS, mesoporous silica; NPs, nanoparticles; RBL, rat basophilic leukemia; PS, polystyrene.
known to generate the so-called “protein corona” and to influence their internalization; and (3) most importantly, under membrane cholesterol-depleted conditions, the uptake of MPS NPs was almost completely precluded, while that of PS NPs proceeded normally (Figure 8). This is compatible with the two different types of NP engaging different endocytic mechanisms. It should be noted, however, that when the two NPs were incubated together under such membrane cholesterol-depleted conditions, an apparently limited stimulation of MPS NPs occurred, and even more interestingly, a partial colocalization of the two NPs was observed.

These findings consistently suggest that the two NPs engage different endocytic mechanisms and routes which lead them to distinct subcellular compartments. In addition, when the cholesterol-dependent endocytic mechanism is disrupted, the uptake of PS NPs is unaffected, while the uptake of MPS is largely prevented, and only partially occurs via the same cholesterol-independent mechanism exploited by PS NPs. A possible explanation is that disruption of the cholesterol-dependent (clathrin- and caveolae-mediated) endocytic pathway results in stimulation of macropinocytosis, which is preferentially exploited by PS NPs.

On the other hand, MPS NPs appear to be preferentially taken up via a clathrin- or caveolae-dependent mechanism. A comparative study in cells that do (or do not) express caveolin-1 or clathrin heavy-chain (for instance through small interference RNA knock-down) may help to clarify the involvement of caveolae or clathrin in the differential uptake of NPs of different material, charge and size.

**Conclusion**

In this work, it was sought to determine whether MPS and PS NPs could label the inflammatory granules and/or stimulate the rapid exocytosis of these granules in RBL mastocytes, an ‘in vitro’ cell model used to study the inflammation process. It was found that naked MPS and PS-COOH NPs were endocytosed and retained within RBL cells with different levels of efficiency. Both types of NPs gained access to the calcium-regulated secretory compartments of RBL cells. This finding constitutes a proof-of-principle: that it is possible to monitor the inflammation process through the imaging of secretory granules in inflammatory cells ‘in vivo’.

The two types of NPs entered via different endocytic mechanisms, and reached distinct subpopulations of secretory compartments, never showing an overlap of the endocytic routes. While MPS NPs accumulated in acid compartments labeled by LysoTracker™ and containing cathepsin D, PS NPs transitorily accumulated in calcium-regulated recycling compartments not labeled by LysoTracker™ and not containing cathepsin D. Intracellular retention greatly differed between the two types of NPs, with PS-COOH being largely and rapidly expelled in a calcium-dependent fashion.

To the best of our knowledge, this is the first report showing the exocytosis of NPs. The observation that PS-COOH NPs trigger the calcium-dependent recycling of secretory organelles raises concerns about the clinical utilization of this material, because of the potential activation of tissue mastocytes.

**Acknowledgments**

This research was supported by Poli di Innovazione (Polo “Biotecnologie e biomediche”, progetto BANP, FinPiemonte, Italy). The fluorescence imaging facility was sponsored by Comoli-Ferrari and C SpA (Novara, Italy).

**Disclosure**

Cyanine Technology S.p.A (Turin, Italy) provided the IRIS-3-doped mesoporous silica (MPS) NPs free of charge. Antonino Giovia is the recipient of a PhD fellowship partly financed by Cyanine Technologies S.p.A (Turin, Italy). The authors report no conflicts of interest in this work.

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