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A comparison of the effects of polystyrene and polycaprolactone nanoplastics on macrophages†

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Plastics are persistent in the environment, which suggests that they may induce adverse effects due to their progressive accumulation over time. This progressive accumulation is facilitated by the fact that macroplastics released into the environment progressively fragment into micro and nanoplastics, which are easily taken up by a wide range of living organisms. As these micro and nanoplastics are particulate materials, they are handled in these living organisms by specialized phagocytic cells, namely macrophages in vertebrates, opening the possibility that plastics accumulating in macrophages may elicit a variety of responses. Thus, one way of alleviating such accumulation in macrophages and in other cell types would be to use biodegradable plastics. Polycaprolactone is a biodegradable plastic showing favorable degradation characteristics in several environments. We thus investigated the responses of macrophages upon treatment with polycaprolactone nanobeads, using a combination of proteomics and validation experiments, and compared these results to the ones induced by polystyrene nanobeads. Many changes detected by proteomics, for example in the mitochondrial, lysosomal or reticulum proteins, did not induce detectable physiological consequences. A slight decrease in the phagocytic capacity of polycaprolactone-treated cells was detected, but not for polystyrene-treated cells. We also showed that polycaprolactone nanobeads degrade within a few days in macrophages, modulating the macrophage responses. The decrease in phagocytosis disappeared, while polystyrene induced a delayed surge in the phagocytic response. A delayed decrease in the secretion of pro-inflammatory cytokines was also observed in polycaprolactone-treated cells, which may be linked to the release of hydroxycaproic acid.

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Environmental significance

Biodegradable plastics appear as a solution to the problems posed by classical plastics, which persist for a very long time and contaminate persistently living organisms. Within biodegradable plastics, polycaprolactone has attracted interest due to its rather easy degradation in many environments. However, this degradation is not immediate, meaning polycaprolactone nanoparticles can be expected to be found in the environment, and are likely to contaminate living organisms. We have thus studied the effect of polycaprolactone nanoparticles on macrophages, *i.e.* professional phagocytes conserved in evolution, by a combination of proteomic and targeted approaches. In addition to limited immediate effects, polycaprolactone nanoparticles degrade relatively slowly in macrophages and induce delayed effects that may alter the normal functions of phagocytes.

1. Introduction

Plastics are now such an important part of our lifestyle, through their extremely diverse uses, that they are produced in gigantic amounts, *i.e.* close to 500 Mt per year.¹ Unfortunately a huge proportion of these plastics, estimated at 10 Mt per year, is released into the oceans alone,² where they accumulate over time, leading to estimates of 150 Mt of plastics in the oceans in 2025.³ This plastic waste has deleterious effects that are more and more documented in detail on a wide range of marine taxa,⁴ including sea birds.⁵ This pollution was first documented in aquatic marine environments,^{6–9} but is

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now found in marine sediments,^{10,11} freshwater environments,^{12–14} and also in terrestrial ones.^{15,16}

Part of the interest in plastics resides in their chemical resistance, which translates into very long half lives in the environment amounting to decades.¹⁷ However the plastics released into the environment do fragment, first into microplastics (less than 1 mm in size) and then into nanoplastics (less than 1 μm in size). It is anticipated that the smaller the plastic particles are, the more easily they will cross biological barriers and be able to disturb the homeostasis of living organisms. This hypothesis has prompted extensive research on the effects of micro and nanoplastics on a wide variety of living organisms from various phyla, ranging from worms^{18,19} to molluscs,^{20–22} crustaceans,^{23,24} insects,^{25,26} and then vertebrates such as fish^{27–29} and of course mammals³⁰ including human models (e.g. in ref. 31–35).

Multicellular organisms defend themselves against particles, including plastic particles, by a series of mechanisms. The first line of defense is represented by biological barriers (e.g. intestinal or epidermal barriers) and research has been devoted to understanding how these barriers interact with plastic particles. When translocation across these barriers occurs,³¹ then a second line of barriers comes into play, and is represented by professional phagocytes. This cell type is encountered in invertebrates (annelids coelomocytes, insect hemocytes), as well as in vertebrates (macrophages, neutrophils). Indeed, it has been shown that this cell type does respond to plastic particles.^{19,26,36–42}

In view of these deleterious effects, the development of biodegradable plastics has been investigated and is a very active area of research. There is however a compromise to be made between the technical properties of plastics and their biodegradability. Many applications, such as food packaging, require good resistance to water, and thus reasonable hydrophobicity, which decreases the interest in polysaccharide-based plastics, for example. In this respect, aliphatic polyesters represent an attractive choice. These plastics are rather hydrophobic, and yet the ester bond is relatively labile, ensuring a much better degradability than plastics where the backbone is a very long aliphatic chain without heteroatoms. Among the aliphatic polyesters, polycaprolactone (PCL) has retained attention because of its attractive properties, as reviewed in ref. 43. It uses a single monomer, ϵ -caprolactone, which polymerizes by ring opening, allowing an easy control of the polymerization reaction. This plastic, however, suffers from a relatively high price of the monomer, and hence of the polymer, which has led to its current use in niche applications, especially biomedical ones. Indeed, its slow but efficient degradation in mammals makes it a very good candidate for slow delivery drug release,^{44–47} or for building scaffolds, e.g. for orthopedics or various implants,^{48–50} which resorb slowly upon biocolonization by the host cells.^{51,52} In addition to these now well-established applications, there has been a

renewal of interest in PCL for applications in membranes⁵³ and packaging.^{54–60} This renewed interest is further supported by recent research showing that caprolactone can be obtained from bioresources,⁶¹ and not only from fossil oil as it is now.

This polymer seems to combine interesting technical properties and good degradability,⁶² e.g. through composting,^{63,64} with complete degradation between 100 and 300 days depending on the temperature⁶³ and even in marine environments, with degradation of PCL films in a few weeks.⁶⁵ Thus, there is a sharp contrast between the relatively easy degradation of PCL by microbes and its slow degradation in mammals. This poses in turn the problem of the effects that PCL particles may have on mammals. In biomedical applications, PCL is used in a kind of pulse-chase mode. The particles or the scaffolds represent a single and massive dose, which is slowly and completely eliminated. If PCL comes into everyday use products such as packaging films, we will then be subjected to a constant influx of PCL particles that, because of the slow degradation of this polymer, may lead to a constant loading of the cleaning system represented by phagocytes, or even to accumulation if the influx outweighs the efflux. Because of this potential caveat, we decided to investigate the effects of PCL nanoparticles on macrophages. As a positive control, we used the well-known polystyrene (PS) nanoparticles, which represent a non-degradable plastic. In order to obtain as much information as possible on the effects of these particles on macrophages, we decided to use a proteomic approach, as done earlier on other types of nanoparticles.^{66–68}

2. Materials and methods

2.1. Plastic particles

Spherical pristine PCL particles (200 nm) were purchased from Phosphorex (Hopkinton, USA) or their retailer Millipore (#PCL200) and spherical pristine PS particles (200 nm) were from Polysciences (#07304-15). Deep red fluorescent labelled polystyrene (PS) particles (SkyBlue particles, 100–300 nm, catalog number #FP-0270-2, batch number #AL01) were purchased from SpheroTech. The particles were used for both the proteomic experiments and the validation experiments. The nanoparticle suspensions were pasteurized overnight at 80 °C before use in cell culture.

For hydrodynamic diameter and zeta potential measurements, PCL and PS particles were diluted to 100 $\mu\text{g mL}^{-1}$ in ultrapure water, then transferred to a quartz spectroscopic cell. The hydrodynamic diameter was measured by DLS using a Litesizer 500 (Anton Paar, USA), and the surface charge was evaluated *via* zeta potential measurement using an Univette (Anton Paar, USA), with the same instrument. An average value was obtained from repeated measurements for each sample ($n = 3$) and analyzed with the instrument-associated Kaliope software.

PCL and PS particle primary diameters were assessed from transmission electron microscopy (TEM) images, obtained



using negative staining on a grid. Samples of 10 mg mL⁻¹ PCL or PS particles were deposited on the clean side of a carbon film on mica, stained with 2% uranyl acetate and 2% phosphotungstic acid, respectively, before being transferred onto a 400 mesh copper grid. Images were acquired using a Tecnai 12 LaB6 electron microscope equipped with a Gatan Orius 1000 CCD camera at an accelerating voltage of 120 kV, with defocus values ranging from 1.2 to 2.5 μ m.

For FTIR analysis, 5 μ L of 10 mg mL⁻¹ particle suspensions were air-dried on a foil. Measurements were performed using a Nicolet iS50 (Thermo Scientific, USA) spectrometer equipped with a diamond ATR module. The number of scans was fixed to 32 with a resolution of 4 cm⁻¹ between 4000 and 400 cm⁻¹. FTIR spectra were acquired and analyzed using OMNIC software (Thermo Scientific, USA). Each spectrum corresponds to the average of three measurements taken on different zones on the sample, chosen manually in order to verify the absence of spectral variation through the sample. No variation was observed.

For some experiments requiring labelling of the particles, such as degradation in cells or internalization ratio determination, the PCL particles were labelled with the red fluorescent dye Disperse Blue 14 (1,4-bis(methylamino) anthraquinone, ABCR #AB177338, λ_{ex} 640 nm, λ_{em} 685 nm) by the solvent swelling method.⁶⁹ Briefly, a 10 mg mL⁻¹ solution of Disperse Blue 14 in THF was prepared. 1% in volume of this solution and 10% THF were added to the bead suspension, and the mixture was agitated on a rotary wheel at room temperature for 30 minutes. Two volumes of water were then added and the particles collected by centrifugation (30 minutes at 15 000g). The pellet was then resuspended in water and the beads recollected by centrifugation. The final pellet was then resuspended in 20% ethanol.

2.2. Cell culture

The J774A.1 cell line (mouse macrophages) was purchased from the European cell culture collection (Salisbury, UK). Cells were routinely propagated in DMEM supplemented with 10% fetal bovine serum (FBS) in non-adherent flasks (Cellstar flasks for suspension culture, Greiner Bio One, Les Ulis, France). For routine culture, the cells were seeded at 200 000 cells per ml and passaged two days later, with a cell density ranging from 800 000 to 1 000 000 cells per ml. For exposure to plastic particles and to limit the effects of cell growth, cells were seeded at 500 000 cells per ml in 6 or 12 wells plates, let settle and recover for 24 hours, and then exposed to the particles at 80 μ g mL⁻¹ for 24 hours before harvesting for the experiments. Proteomic experiments were carried out in 6 well plates, and all the other experiments in 12 well plates. The medium volume was adjusted to keep the same height across all cell culture formats. Cells were used at passage numbers from 5 to 15 post-reception from the repository. Cell viability was measured by the propidium iodide method,⁷⁰ or with the SytoxGreen probe (ThermoFisher S7020) using the protocol provided by the supplier. Those viability tests were

systematically performed along functional tests to check that cell viability was always correct. Thus, depending on the fluorophores used in the functional tests, different fluorophores had to be used for the viability tests in order to avoid interferences.

For aging experiments, the cells were seeded at 300 000 cells per ml in DMEM containing 1% horse serum, to limit cell proliferation.⁷¹ The medium was changed every two days and the cells kept in culture for up to one week.

Prolonged cell cultures in 1% horse serum were used to determine particle degradation. To this purpose, cells cultured in 12 well adherent plates were exposed to fluorescent particles (PS or PCL) for 24 hours. The particle-containing medium was removed, the cells rinsed twice with sterile PBS, and cultured in DMEM with a medium change every two days. To measure the cell associated fluorescence, the medium was removed, and the cell layer was then rinsed once with 1 ml PBS and the PBS removed. The cell layer was then lysed by the addition of 400 μ L of 10 mM Hepes pH 7.5, and the cell lysate was collected. Its fluorescence was the measured using a DeNovix QFX instrument (excitation at 635 nm, emission collected in the 665–740 nm window). Untreated cell cultures of matched age were used to determine the cell autofluorescence background. This protocol is by construction independent of cell division, as each well is considered as a whole.

2.3. Proteomics

Proteomics was carried out essentially as described previously.⁴¹ However, the experimental details are given here for the sake of consistency.

2.3.1. Sample preparation. After exposure to the plastic particles, the cells were harvested by flushing the 6 well plates. They were collected by centrifugation (200g, 5 minutes) and rinsed twice in PBS. The cell pellets were lysed in 100 μ L of extraction buffer (4 M urea, 2.5% cetyltrimethylammonium chloride, 100 mM sodium phosphate buffer pH 3, 150 μ M methylene blue). The extraction was conducted at room temperature for 30 minutes, after which the lysate was centrifuged (15 000g, 15 minutes) to pellet the nucleic acids. The supernatants were then stored at -20 °C until use.

2.3.2. Shotgun proteomics. For the shotgun proteomic analysis, the samples were included in polyacrylamide plugs according to Muller *et al.*⁷² with some modifications to downscale the process.⁷³ To this purpose, the photopolymerization system using methylene blue, toluene sulfinate and diphenyliodonium chloride was used.⁷⁴

As mentioned above, methylene blue was included in the cell lysis buffer. The other initiator solutions included a 1 M solution of sodium toluene sulfinate in water and a saturated water solution of diphenyliodonium chloride. The ready-to-use polyacrylamide solution consisted of 1.2 ml of a commercial 40% acrylamide/bis solution (37.5/1) to which 100 μ L of diphenyliodonium chloride solution, 100 μ L of sodium toluene sulfinate solution and 100 μ L of water were added.



To the protein samples (15 μL), 5 μL of acrylamide solution were added and mixed by pipetting in a 500 μL conical polypropylene microtube. 100 μL of water-saturated butanol were then layered on top of the samples, and polymerization was carried out under a 1500 lumen 2700 K LED lamp for 2 hours, during which the initially blue gel solution discolored. At the end of the polymerization period, butanol was removed, and the gel plugs were fixed for 1 h with 200 μL of 30% ethanol 2% phosphoric acid, followed by 3×15 minutes washes in 20% ethanol. The fixed gel plugs were then stored at -20°C until use.

Gel plug processing, digestion, peptide extraction and nano LC-MS/MS were performed as previously described,⁷³ without the robotic protein handling system and using a Q-Exactive HF-X mass spectrometer (Thermo Fisher Scientific, Bremen, Germany). Further details are available in Methods S1.†

For protein identification, the MS/MS data were interpreted using a local Mascot server with MASCOT 2.6.2 algorithm (Matrix Science, London, UK) against an in-house database containing all *Mus musculus* and *Rattus norvegicus* entries from UniProtKB/SwissProt (version 2019_10, 50 313 sequences) and the corresponding 50 313 reverse entries. Spectra were searched with a mass tolerance of 10 ppm for MS and 0.07 Da for MS/MS data, allowing a maximum of one trypsin missed cleavage. Trypsin was specified as enzyme. Acetylation of protein N-termini, carbamidomethylation of cysteine residues and oxidation of methionine residues were specified as variable modifications. Identification results were imported into Proline software version 2.1 (<https://proline.profiptroteomics.fr/>) for validation. Peptide spectrum matches (PSM) with pretty rank equal to one were retained. The false discovery rate was then optimized to be below 1% at the PSM level using the Mascot adjusted *E*-value and below 1% at the protein level using the Mascot Mudpit score.

Mass spectrometry data are available via ProteomeXchange with the identifier PXD049997.

2.3.3. Label free quantification. Peptide abundances were extracted thanks to Proline software version 2.2 (<https://proline.profiptroteomics.fr/>) using a *m/z* tolerance of 10 ppm. Alignment of the LC-MS runs was performed using Loess smoothing. Cross assignment was performed within groups only. Protein abundances were computed from the sum of peptide abundances (normalized using the median).

2.3.4. Data analysis. For the global analysis of the protein abundance data, missing data were imputed with a low, non-null value. The complete abundance dataset was then analyzed by the PAST software.⁷⁵

Proteins were considered as significantly different if their *p* value in the Mann-Whitney *U*-test against control values was inferior to 0.05. The selected proteins were then subjected to pathway analysis using the DAVID tool,⁷⁶ with a cutoff value set at an FDR of 0.1.

2.4. Phagocytosis assay

For this assay, the cells were first exposed to deep red-labelled PCL or PS particles at 80 $\mu\text{g mL}^{-1}$. After 24 hours of

exposure, the cells were then exposed to 0.5 μm latex beads (carboxylated surface, yellow green-labelled, from Polysciences, excitation 488 nm, emission 527/32 nm) for 3 hours at a concentration of 5 $\mu\text{g mL}^{-1}$. After this second exposure, the cells were collected, rinsed twice with PBS, and analyzed for the two types of fluorescence (green and red) on a Melody flow cytometer. Controls with no latex beads showed no green fluorescence.

2.5. Mitochondrial transmembrane potential assay

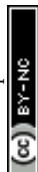
The mitochondrial transmembrane potential assay was performed essentially as described previously.⁴² After exposure to plastic beads (either PCL or PS, 80 $\mu\text{g mL}^{-1}$ for 24 hours), rhodamine 123 (Rh123) was added to the cultures at an 80 nM final concentration (to avoid quenching⁷⁷), and the cultures were further incubated at 37°C for 30 minutes. At the end of this period, the cells were collected, washed in cold PBS containing 0.1% glucose, resuspended in PBS glucose and analyzed for green fluorescence (excitation 488 nm emission 525 nm) on a Melody flow cytometer. As a positive control, butanedione monoxime (BDM) was added at a 30 mM final concentration together with Rh123.⁷⁸ As a negative control, carbonyl cyanide 4-(trifluoromethoxy) phenylhydrazone (FCCP) was added at a 5 μM final concentration together with Rh123.⁷⁸ Control cells unexposed to Rh123 did not show any green fluorescence.

2.6. Lysosomal assay

For the lysosomal function assay, the Lysosensor method was used, as described previously.⁴² After exposure to plastic beads (either PCL or PS, 80 $\mu\text{g mL}^{-1}$ for 24 hours), the medium was removed, the cell layer was rinsed with complete culture medium and incubated with 1 μM Lysosensor Green (Molecular Probes) diluted in warm (37°C) complete culture medium for 1 hour at 37°C . At the end of this period, the cells were collected, washed in cold PBS containing 0.1% glucose, resuspended in PBS glucose and analyzed for green fluorescence (excitation 488 nm, emission 540 nm) on a Melody flow cytometer. Control cells unexposed to Lysosensor Green did not show any green fluorescence.

2.7. Cytokine release assays

Cells were first exposed to nanoplastics (80 $\mu\text{g mL}^{-1}$) for 24 hours. At the end of this exposure period, the culture medium was removed, the cell layer was rinsed with culture medium and fresh medium was added to the wells. In half of the wells LPS (1 ng mL^{-1}) was added. After another 24 hours or 5 days, the medium was collected and analyzed for proinflammatory cytokines. Tumor necrosis factor and interleukin 6 levels were measured using the Cytometric Bead Array Mouse Inflammation Kit (catalog numbers 558299, 558301 and 558266, BD Biosciences, Le Pont de Claix), and analyzed with FCAP Array software (3.0, BD Biosciences) according to the manufacturer's instructions.



2.8. Assay for oxidative stress

For the oxidative stress assay, a protocol based on the oxidation of dihydrorhodamine 123 (DHR123) was used, essentially as described previously.⁴² After exposure to plastic beads ($80\ \mu\text{g}\ \text{ml}^{-1}$ for 24 hours), the cells were treated in PBS containing $500\ \text{ng}\ \text{ml}^{-1}$ DHR123 for 20 minutes at $37\ ^\circ\text{C}$. The cells were then harvested, washed in cold PBS containing 0.1% glucose, resuspended in PBS glucose and analyzed for green fluorescence (the same parameters as rhodamine 123) on a Melody flow cytometer. Menadione (applied on the cells for 2 hours prior to treatment with DHR123) was used as a positive control in a concentration range of $25\text{--}50\ \mu\text{M}$. Control cells unexposed to DHR123 did not show any green fluorescence.

3. Results

3.1. Plastic bead characterization

PCL and PS particles were both described by their supplier as being $200\ \text{nm}$ in diameter. Regarding PCL particles, chemical composition was validated through FTIR analysis, which showed the typical features of PCL (Fig. 1A). These features were bands at $2944\ \text{cm}^{-1}$ and $2867\ \text{cm}^{-1}$, corresponding to asymmetric and symmetric stretching of CH_2 groups, and typical bands of esters, notably at $1723\ \text{cm}^{-1}$, which is attributed to the stretching of the ester carbonyl ($\text{C}=\text{O}$), then at $1240\ \text{cm}^{-1}$ and $1186\ \text{cm}^{-1}$, corresponding to asymmetric and symmetric stretching of the $\text{C}-\text{O}-\text{C}$ group. TEM images showed that PCL particles exhibited spherical morphologies with heterogeneous size distribution ($204 \pm 85\ \text{nm}$) (Fig. 1B). The polydispersity index from DLS analysis of size distribution confirmed the heterogeneous size distribution, *i.e.*, $17 \pm 5\%$, and a Z-average of $197 \pm 2\ \text{nm}$.

PS particles were also confirmed to be composed of PS *via* FTIR (Fig. 1C), with typical absorption bands from the aromatic $\text{C}-\text{H}$ stretching vibrations at 3081 , 3059 , and $3025\ \text{cm}^{-1}$ and asymmetrical and symmetrical stretching vibrations of methylene ($-\text{CH}_2$) groups at 2921 and $2849\ \text{cm}^{-1}$. Carbon-carbon stretches in aromatic ring were observed at $1601\ \text{cm}^{-1}$ and $1583\ \text{cm}^{-1}$, as expected,⁷⁹ as well as at $1493\ \text{cm}^{-1}$ and $1452\ \text{cm}^{-1}$. The peak at $755\ \text{cm}^{-1}$ is attributed to out-of-plane $\text{C}-\text{H}$ deformation, while the peak at $697\ \text{cm}^{-1}$, also related to this deformation, indicates ring curvature. TEM images showed that PS particles exhibited spherical morphologies with a rather homogeneous size distribution ($303 \pm 41\ \text{nm}$) (Fig. 1D).

The parameters obtained in aqueous suspensions for the two beads by DLS are summarized in Table 1.

The PS beads proved to be within the range of the nominal diameter given by the supplier ($0.1\text{--}0.3\ \mu\text{m}$, mean $0.26\ \mu\text{m}$), while the electron microscopy data were much closer to the nominal values. The observed sizes led to a bead number/cell ratio higher than 9000 beads per cell (at $80\ \mu\text{g}\ \text{ml}^{-1}$) for the PS beads and higher than 6000 beads per cell for the PCL beads, when corrected for the internalization ratios determined for both particles (Experiments S1†).

3.2. Viability of the plastic-treated cells

First, the toxic effects of the PCL or PS beads on J774A.1 cells were determined after a 24 hour exposure. The results, shown in Fig. 2, demonstrated a very low toxicity of the PCL and PS beads, with an LD50 that was greater than $300\ \mu\text{g}\ \text{ml}^{-1}$. We decided to use an $80\ \mu\text{g}\ \text{ml}^{-1}$ plastic concentration, *i.e.* the first concentration for which the viability was slightly but

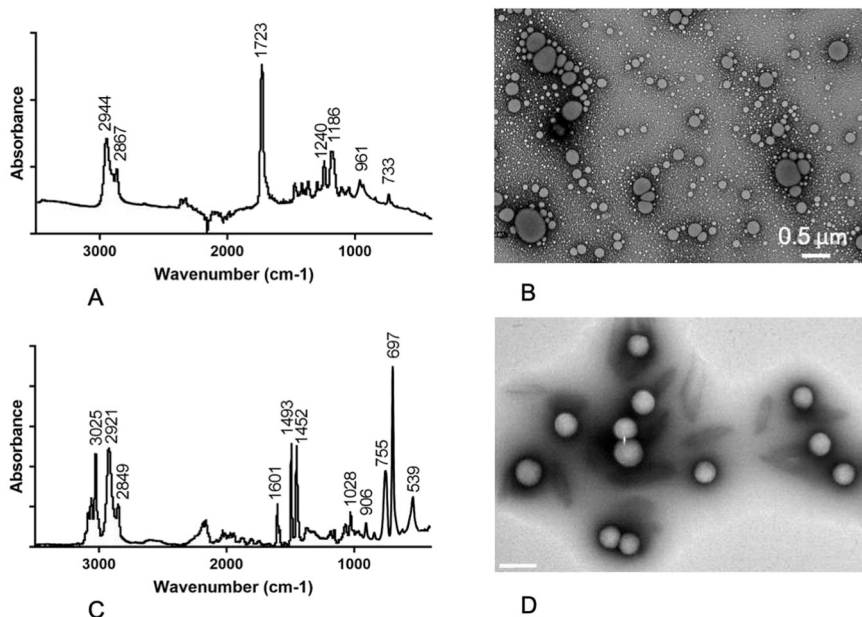


Fig. 1 Characterization of the plastic beads. Chemical characterization of the PCL and PS beads was carried out by FT-IR spectroscopy (panels A and C, respectively). Physical characterization of the beads was carried out by transmission electron microscopy (panels B and D, respectively).



Table 1 Characterization of the bead size parameters in suspension

Beads	Average primary diameter (TEM)	Average hydrodynamic diameter	Polydispersity index	Zeta potential
PCL	204 ± 85 nm	197 ± 2 nm	17 ± 5%	-10.4 ± 1.1 mV
PS	191 ± 12 nm	182 ± 12 nm	8 ± 4%	-43.9 ± 5.7 mV

The results are expressed as mean ± standard deviation ($N = 3$).

statistically significantly lower than the one of unexposed, control cells.

3.3. Global analysis of the proteomic results

The shotgun proteomic analysis was able to detect and quantify 3402 proteins (Table S1†). As a first analysis, we analyzed the complete dataset *via* the PAST software, through a principal coordinates analysis. The results, displayed in Fig. 3, showed a good separation of all three groups (control, PS-treated and PCL-treated). The higher heterogeneity of the PS-treated group may be linked to the fact that some toxicity begins to appear, resulting in more divergent proteomes. The fact that the PCL and control groups were well separated in this global analysis also showed that PCL nanoparticles did modulate the cell proteome appreciably, although no toxicity is detected at the chosen concentration. The extent of the modulation is confirmed by the analysis of similarity test.⁸⁰ This test showed a p -value (probability by random) of 0.0068 for the PCL *vs.* control comparison, 0.0071 for the PS *vs.* control comparison, and 0.0074 for the PCL *vs.* PS comparison. This indicated on the one hand significant proteome changes induced by the exposure to each plastic type, and on the other hand that the changes induced by PS and PCL are also different.



Fig. 2 Viability of cells treated with PCL particles. Cells were treated with PCL (blue curve) or PS (green curve) particles for 24 hours, and their viability measured by a flow cytometry fluorophore exclusion assay (Sytox green). Results are displayed as mean ± standard deviation ($N = 4$).

As we have already published data on the proteomic response of macrophages to PS nanoparticles,⁴¹ we decided to focus the analysis on the cellular response of macrophages to PCL. Proteins modulated by the internalization of PCL particles were then selected on the basis of a Mann–Whitney U test in the comparison of plastic-treated cells to unexposed controls. This resulted in the selection of a total of 643 modulated proteins (Table S2†).

In order to gain further insight into the significance of the observed changes, this list of modulated proteins was used to perform pathways analyses by the DAVID software, and the results are shown in Table S3†. The pathways modulated by PS beads are shown in Table S4†. Regarding the pathways modulated by PCL, some indicated a global stress response, such as translation, nucleotide binding, nucleus, endoplasmic reticulum, mitochondria or carbon metabolism (Table S5†), which is expected for any cellular stress, while other pathways appeared more specific of cellular internalization of particles (*e.g.* lysosomes), and some specific for macrophages (*e.g.* innate immunity, hydrogen peroxide production). Compared to the PCL-modulated pathways, PS-modulated pathways also included mitochondria and lysosomes, but not endoplasmic reticulum. The proteasome-associated pathways were also highlighted for response to PS particles, as well as more metabolic pathways (*e.g.* pentose phosphate pathway, glycolysis). Macrophage-specific pathways did not appear as prominent in response to PS than in response to PCL.

This comprehensive analysis of the proteomic results led us to perform validation experiments on several functions.

3.4. Mitochondria and endoplasmic reticulum

Mitochondrial proteins represented an abundant class among the modulated proteins, with 80 proteins (Table S6†). Among those, 30 were increases and 50 decreases, suggesting a perturbation of the mitochondrial functions. In more detail, 10 were subunits of the respiratory chain (P19783, P62897, Q62425, Q91VR2, Q91WD5, Q9CQA3, Q9CR68, Q9CZ13, Q9D3D9, Q9DCX2) and one (Q9D1R1), a chaperone involved in the assembly of respiratory chain complex I. As the respiratory chain complexes I to III produce the proton gradient responsible for the mitochondrial transmembrane potential, which is in part dissipated by complex V, thereby ensuring the balance of the mitochondrial transmembrane potential, we decided to test this mitochondrial parameter. The results, displayed in Fig. 4A, showed no change of the mitochondrial transmembrane potential upon exposure of





Fig. 3 Global analysis of the proteomic data. The complete proteomic data table (3402 proteins) was analyzed by principal coordinates analysis, using the PAST software. The mathematical distance used for the calculations was the Gower distance. The results are represented as the X–Y diagram of the first two axes of the principal coordinates analysis, representing 60% of the total variance. Eigenvalue scale. This representation allows us to figure out how, at the global proteome scale, the samples are related to each other. Samples grouped in such a diagram indicate similar proteomes, and the larger the distance between samples are, the more dissimilar their respective proteomes.



Fig. 4 Mitochondria, endoplasmic reticulum. Panel A: mitochondrial transmembrane potential (rhodamine 123 method). All cells were positive for rhodamine 123 internalization in mitochondria, and the mean fluorescence is the displayed parameter. Results are displayed as mean $\pm$ standard deviation ($N = 4$). Cells: unexposed cells. BDM: cells exposed for 30 minutes to butanedione monoxime (induces an increase in the transmembrane potential). FCCP: cells exposed for 30 minutes to FCCP (induces a decrease in the transmembrane potential). Panel B: test of the endoplasmic stress response in PCL toxicity. Cells were pre-treated with 4 μ M salubrial for 4 hours, and various concentrations of PCL beads were then added over a further 18 hours in culture. At the end of the experiment, the cell viability was measured. Results are displayed as survival curves, with the standard deviations at each tested point ($N = 6$). Blue curve: cells untreated with salubrial. Green curve: salubrial-treated cells. Cells: unexposed cells.

macrophages to PCL beads, and a slight and not statistically relevant decrease upon exposure of macrophages to PS beads. We did not check the effect of a longer exposure to PS beads. Interestingly, among the 32 proteins that show opposite modulations in response to PCL compared to PS, the three ATP synthase subunits selected in the proteomic screen show decreases in response to PCL particles and increases in response to PS particles. As ATP synthase dissipates the proton gradient to produce ATP, these changes in ATP

synthase may be related to the differences in the mitochondrial transmembrane potential observed between the PCL-exposed cells and the PS-exposed cells.

Endoplasmic reticulum proteins also represented an important class of modulated proteins in response to PCL particles (Table S7†), with 52 proteins (37 increases, 15 decreases). Among these 52 proteins, 5 were annotated directly or indirectly as related to the endoplasmic stress response (A1L1L2, Q5U2X6, Q5XI41, Q9QZN4, Q9Z2G6). This



prompted us to test whether endoplasmic stress could be a determinant of toxicity for the PCL particles. To this purpose we used salubrinal, an inhibitor of the endoplasmic reticulum stress response,⁸¹ which is known to counteract the lethal effects of this stress on cells.⁸² The results, shown in Fig. 4B, indicated no effect of salubrinal on cell survival up to treatment with 320 $\mu\text{g ml}^{-1}$ PCL particles for 24 hours.

3.5. Lysosomes and phagocytosis

In macrophages, phagocytosis is an integrated function coupling vesicle formation and transport to produce phagosomes, and finally fusion with lysosomes to produce phagolysosomes where the internalized material is subjected to degradation. Thus, in order to probe this function at the proteomic level, we integrated the proteins appearing with the keywords “lysosome” and “actin cytoskeleton” in a single protein list (Table S8†). We also added a protein appearing under the cluster “innate immune response”, but which is known to be implicated in the control of phagocytosis (STAP1, Q9JM90). This list encompassed 37 proteins (19 increases, 18 decreases). In the list of increased proteins were found two proteins participating in the acidification of the lysosomes, namely LAMP2 (O35114),⁸³ and a catalytic subunit of the proton ATPase itself (P50516). This prompted us to test lysosomal acidification by the Lysosensor method. The results, displayed in Fig. 5A, showed a slight but not statistically significant ($p = 0.1$) increase in Lysosensor accumulation in response to PCL particles, and an even lower increase in response to PS. We then investigated how the uptake of PCL (or PS) particles influenced the residual phagocytic capability of macrophages. The results, displayed in Fig. 5B and C, showed a decrease in the phagocytic ability

of PCL-treated cells, both in terms of percentage of phagocytic cells at 3 hours (Fig. 5B) and in terms of internalized beads' fluorescence (Fig. 5C), which was not observed for PS-treated cells.

3.6. Immunity-related proteins and inflammation

The “innate immune response” keyword appeared in the pathway analysis, pointing to a subset of 47 modulated proteins (37 increases, 10 decreases, Table S9†). Among the increased proteins, we noticed both chains of cytochrome b-245, a critical component of NADPH oxidase, *i.e.* the enzyme producing superoxide in the phagosome. This suggested that the macrophages try to degrade the internalized plastic beads (both PCL and PS) by their classical defense mechanism.

When examining in more detail the proteins present in this cluster, subgroups could be identified. One is related to proteins involved in antiviral responses (*e.g.* P0DOV2, P1103, P11928, Q3TL44, Q3UPF5, Q60710, Q64620, Q8BV66, Q8VCF0, Q8VI93, Q8VI94). The other important group included proteins that are known to modulate the inflammatory response. When trying to figure out the possible outcome of the modulations observed by proteomics in terms of cytokine production, we observed opposite trends. In response to the exposure to PCL particles, we observed increases in the abundances of negative regulators of inflammatory cytokine production (*e.g.* P25911,⁸⁴ P54987,⁸⁵ Q3UCV8,⁸⁶ Q64281,⁸⁷ Q9D8Y7,⁸⁸ Q9R002 (ref. 89)), but also increases in positive regulators of inflammatory cytokine production (*e.g.* O88351,⁹⁰ P56477,⁹¹ Q61107,⁹² Q9BDB7,⁹³ Q9Z0E6 (ref. 92)) and decreases in negative regulators of cytokine production (Q9R007 (ref. 94)).



Fig. 5 Lysosomes and phagocytosis experiments. Panel A: lysosomal proton pumping (Lysosensor method). All cells were positive for lysosensor internalization in lysosomes, and the mean fluorescence is the displayed parameter. Results are displayed as mean $\pm$ standard deviation ($N = 4$). Panels B and C: phagocytosis. Cells were first treated for 24 hours with 80 $\mu\text{g ml}^{-1}$ PCL (or PS) particles. After removal of the particle-containing cell culture medium, the cells were treated with green fluorophore labelled carboxylated latex beads for 3 hours. Panel B: the percentage of green fluorescence-positive cells, indicating the percentage of cells able to internalize the test beads in 3 hours, is the displayed parameter. Results are displayed as mean $\pm$ standard deviation ($N = 4$). Significance marks: † $p < 0.05$ (Mann-Whitney U -test method, comparison between control and each treatment). Panel C: the mean fluorescence, indicating the amount of green beads internalized, is the displayed parameter. Results are displayed as mean $\pm$ standard deviation ($N = 4$). Significance marks: ** $p < 0.01$ (Student t -test method, comparison between control and each treatment).



In view of these discrepant results, we tested the cytokine release of plastic-exposed macrophages under two conditions: either with plastic treatment only or in combination with a LPS stimulation. Responses to plastic alone allowed us to assess whether the plastics had an intrinsic pro-inflammatory effect or not. Responses to plastics followed by LPS allowed us to determine if the plastics could modulate further responses of the macrophages to bacteria. We tested two pro-inflammatory cytokines (TNF and IL-6). The results displayed in Fig. 6 showed first that PCL did not induce by itself any pro-inflammatory response. However, the combination of PCL-treatment with LPS stimulation resulted in a decrease of the magnitude of the response for both measured cytokines, although this decrease was at the limit of significance for IL-6. This effect was not observed in response to PS treatment.

3.7. Oxidative stress

In the course of the analysis of the proteomic data, we noticed that several proteins involved in the control of the cellular oxidative stress were modulated. These included the Cu-Zn superoxide dismutase (P08228, decreased), the mitochondrial Mn superoxide dismutase (P09671, increased) and peroxiredoxin 6 (O08709, decreased). Other proteins of this class (*e.g.* peroxiredoxins 1 to 5) did not show any significant modulation in response to treatment with PCL beads. These somewhat discrepant results, added to the induction of both cytochrome b245 subunits (ROS promoters), prompted us to study the level of cellular oxidative stress in response to PCL beads. The results, displayed in Fig. 7, showed a very slight but statistically significant decrease in the cellular oxidative stress in PCL-treated beads, while a slight but statistically significant increase was observed in PS-treated cells.

3.8. Effect of lower particle concentration on cells

In order to determine to which extent the dose of particles is responsible for the effects that we could detect, we

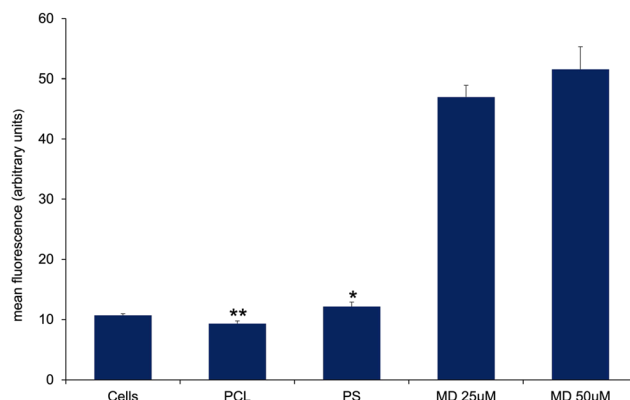


Fig. 7 Cellular oxidative stress, measured with the dihydrorhodamine 123 (DHR123) indicator. The cells were exposed to PCL or PS beads for 24 hours, and finally for 20 minutes to the DHR 123 probe. Menadione (MD, 25 $\mu\text{g ml}^{-1}$ or 50 $\mu\text{g ml}^{-1}$ for 2 hours) was used as a positive oxidative stress control. Results are displayed as mean $\pm$ standard deviation ($N = 4$). Significance marks: * $p < 0.05$; ** $p < 0.01$ (Student *t*-test method, comparison between control and each treatment).

performed the same targeted experiments at a particle concentration of 10 $\mu\text{g ml}^{-1}$, and the results are summarized in Table 2.

These data showed that except for the level of oxidative stress, which is slightly but significantly decreased upon exposure to a low concentration of PS particles, this lower concentration of particles did not show any statistically significant changes in the measured parameters, which is consistent with dose-dependent affects.

3.9. Effect of particle aging in the cells

As PCL is known to be biodegradable,⁴³ we first checked whether this was the case in our *in vitro* macrophage system. We thus used fluorescently labelled particles and used the fluorescence as a marker for the presence of the particles. Fluorescently-labelled PS particles were used as a low-



Fig. 6 Cytokine release. The cells were first treated for 24 hours with 80 $\mu\text{g ml}^{-1}$ PCL or PS particles. The medium was then removed and the cells were treated (or not) with 1 ng ml^{-1} lipopolysaccharide in complete cell culture medium for 24 hours. The cell medium was then collected for secreted TNF and IL-6 measurements. Results are displayed as mean $\pm$ standard deviation ($N = 4$). Significance marks: * $p < 0.05$; ** $p < 0.01$; † $p = 0.052$ (Student *t*-test method, comparison between control and each treatment). Panels A and B: TNF-alpha release (respectively without and with LPS stimulation). Panel C: IL-6 release (after stimulation with LPS).



Table 2 Evolution of some macrophage parameters in cells exposed to PCL or PS beads for 24 hours at the concentration of 10 $\mu\text{g ml}^{-1}$

Parameter	Value	Value	Value	P-value	P-value
	Control cells	PCL-treated cells	PS-treated cells	T-test PCL vs. control	T-test PS vs. control
Mitochondrial transmembrane potential (Rh123 fluorescence)	57 $\pm$ 4.5	68.2 $\pm$ 3.4	58.2 $\pm$ 2.1	0.17	0.15
Oxidative stress (DHR123 fluo.)	23.8 $\pm$ 3.6	24.4 $\pm$ 1.4	17.5 $\pm$ 2.4	0.76	0.03
Lysosensor signal	174.3 $\pm$ 12.6	178.6 $\pm$ 14.2	159.2 $\pm$ 19.2	0.66	0.24
% phagocytic cells	90.1 $\pm$ 1.6	88.5 $\pm$ 0.5	91.1 $\pm$ 3.1	0.05	0.51
MFI phagocytic cells	250 $\pm$ 19	239 $\pm$ 13	263 $\pm$ 35	0.24	0.47
Secreted IL-6 (pg ml^{-1})	12.7 $\pm$ 2.9	10.7 $\pm$ 2.3	12.9 $\pm$ 3.1	0.21	0.90
Secreted TNF (pg ml^{-1})	1284 $\pm$ 280	1104 $\pm$ 258	1268 $\pm$ 289	0.27	0.93
Secreted IL-6 with LPS stimul. (pg ml^{-1})	17 270 $\pm$ 3510	16 022 $\pm$ 3582	17 070 $\pm$ 4296	0.55	0.93
Secreted TNF with LPS stimul. (pg ml^{-1})	105 343 $\pm$ 17 530	118 843 $\pm$ 28 666	106 908 $\pm$ 27 445	0.35	0.91

All results are expressed as mean $\pm$ standard deviation ($N = 6$). MFI: mean fluorescence index.

degradation control. The results of these experiments are summarized in Table 3.

The values obtained for polystyrene indicated the losses that are inherent to the experimental setup. The data, however, suggested an important degradation of the PCL beads at day 6, *i.e.* 1 day of exposure plus 5 days of recovery in a bead-free medium. We also showed that this degradation occurred in cells only, as PCL beads suspended in complete culture medium at 37 $^{\circ}\text{C}$ for 6 days did not show any fluorescence loss (Experiments S1†).

This prompted us to re-investigate the biological parameters already investigated just after a 24 hour exposure period. We did not detect any significant increase in oxidative stress levels in PCL-exposed cells compared to control cells after 5 days of recovery post-exposure. However, we detected significant changes in several other parameters, and the results are summarized in Table 4.

These results were in some cases consistent with the ones obtained immediately after exposure. A good example is represented by the lysosomal activity, which is increased by 12% immediately after exposure and by 17% after recovery. Other results are quite different from those obtained immediately after exposure. One striking example is the mitochondrial transmembrane potential, which is basically constant immediately after exposure and strongly increased after particle aging in the cells. Regarding phagocytosis, it was similar in PCL-treated and in unexposed cells, while PS-treated cells showed increased phagocytic indexes. These results were different from those obtained immediately after exposure. Oxidative stress levels were very similar in control,

PCL-treated and PS-treated cells after aging of the particles in the cells. Regarding secretion of pro-inflammatory cytokines, a slight increase in the basal production of IL-6 (and less significantly for TNF) was observed in PS-treated cells where the NP stimulus persists, but not in PCL-treated cells. Oppositely, in the case of LPS-stimulated cells, a delayed anti-inflammatory effect was observed in PCL-treated cells, while the responses were similar to those of the control cells in PS-treated cells.

4. Discussion

Plastics are omnipresent in our daily lives because of their technical properties. They combine light weight to valuable mechanical properties, which range from deformability to mechanical strength depending on thickness and plastic type, and to good resistance to water and water-based environments, which makes them attractive for applications such as food packaging. However, the very same properties become a nuisance when degradation becomes required, *i.e.* at the end of life of the plastic items. This challenge is extremely important for single use plastics, where the end of life comes by definition quite early.

A way to circumvent this degradation problem will be to use plastics that combine good technical properties with a chemical structure that renders them biodegradable. In this regard, purely aliphatic polyesters such as polylactide, polyhydroxybutyrate and polycaprolactone are promising, while mixed aromatic aliphatic polyesters such as poly(ethylene terephthalate) are known to show poor biodegradability, albeit better than plastics with a pure polyolefinic backbone (*e.g.* polystyrene, polyacrylate, polyethylene).

Among this family of aliphatic polyesters, polycaprolactone (PCL) shows promising technical properties^{43,62} combined with good biodegradability under a variety of conditions.^{63,64,95} However, PCL is known to degrade very slowly in mammals, as confirmed by its medical uses.^{43,45,48} This poses the obvious question of its potential toxicological effects if particles of PCL cross the biological

Table 3 Relative cell-associated fluorescence over time

Bead type	Time in the presence of beads		
	1 d	4 d	6 d
PS beads	100 $\pm$ 8	98 $\pm$ 4	66 $\pm$ 14
PCL beads	100 $\pm$ 17	99 $\pm$ 3	20 $\pm$ 3

The fluorescence intensities were normalized by the mean of the intensity observed after 1 day of loading.



Table 4 Evolution of some macrophage parameters in cells exposed to PCL or PS beads for 24 hours and let to recover for 5 days post exposure

Parameter	Value	Value	Value	P-value T-test	P-value T-test
	Control cells	PCL-treated cells	PS-treated cells	PCL vs. control	PS vs. control
Mitochondrial transmembrane potential (Rh123 fluorescence)	66.2 ± 7.4	81.4 ± 6.6	62.8 ± 4.5	0.02	0.46
Oxidative stress (DHR123 fluo.)	49.7 ± 1.8	49.4 ± 4.1	45.9 ± 3.4	0.88	0.11
Lysosensor signal	247.1 ± 52.6	289.6 ± 42.5	246.4 ± 36.8	0.26	0.98
% phagocytic cells	78.1 ± 1.5	80.9 ± 2.6	87.6 ± 1.7	0.05	0.000001
MFI phagocytic cells	445 ± 15	455 ± 39	598 ± 63	0.57	0.0014
Secreted IL-6 (pg ml ⁻¹)	119 ± 62	140 ± 72	274 ± 44	0.59	0.00075
Secreted TNF (pg ml ⁻¹)	1843 ± 268	1783 ± 217	2032 ± 227	0.68	0.22
Secreted IL-6 with LPS stimul. (pg ml ⁻¹)	1475 ± 284	844 ± 147	1324 ± 267	0.0015	0.4400
Secreted TNF with LPS stimul. (pg ml ⁻¹)	5361 ± 1570	3355 ± 670	5374 ± 1333	0.02	0.99

All results are expressed as mean ± standard deviation (*N* = 6). MFI: mean fluorescence index.

barriers and become internalized. To investigate this question we studied the effects of PCL nanoparticles on macrophages, *i.e.* the scavenger cell type in the body in charge of handling any particulates. As expected, the acute toxicity of PCL toward macrophages was very low, with the LD₅₀ not reached at 300 µg ml⁻¹. Indeed, the medical uses of PCL argue against any acute toxicity, but this does not mean that adverse effects will not exist. To investigate this question, we used a proteomic approach, which revealed wide changes in the proteome in response to the exposure to PCL (641 significant changes on 3402 quantified proteins). However, the changes observed at the proteome level may just reflect a homeostatic adaptation of the macrophages to the presence of the PCL particles in order to keep their full functions operational.

Indeed, homeostatic adaptations were observed for the endoplasmic reticulum (ER) and the mitochondria. Although numerous changes were observed for these two classes of proteins (80 for mitochondria, 52 for ER), we could not detect any implication of ER stress nor any mitochondrial depolarization or hyperpolarization in response to PCL after 24 hours. We also did not detect any damage to the lysosomes after 24 hours, opposite to what is observed with other nanomaterials such as silica.⁹⁶

Interestingly, this absence of effects on mitochondria, while important changes in the mitochondrial proteome are observed, has also been demonstrated for macrophages treated with polystyrene particles⁴¹ or with polylactide particles.⁹⁷ The same holds true for the endoplasmic reticulum. Regarding the lysosomes, we rather observed an increase in the lysosensor signal with the other plastics,⁹⁷ which may be interpreted as an increase in the number of active lysosomes in response to plastic particle internalization. The fact that we did not observe this increase with PS particles at 80 µg ml⁻¹ (this paper), compared to 50 µg ml⁻¹,⁹⁷ may be correlated with the onset of toxicity at 80 µg ml⁻¹. It shall also be underlined that the absence of detected lysosomal damage with plastic particles may be due to their spherical shape, as sharp edges are known to induce lysosomal damage.⁹⁸ This however shows that the polymers, including PCL, do not induce lysosomal damage *per se*.

While the general physiology of the macrophages was not affected by the treatment with PCL nanoparticles, the specialized macrophages functions were. We observed both a decrease in phagocytosis and a decrease of inflammatory cytokines in response to LPS-stimulation, suggesting that PCL-exposed cells will be less efficient than normal cells to respond to a bacterial infection. When comparing these results with those obtained on other plastics, polystyrene did not induce any decrease in the phagocytic activity but only a decrease in IL-6 release upon LPS stimulation. Polylactide particles, such as PCL, induced a decrease in the phagocytic activity and a decrease in IL-6 release.⁹⁷ Once again, these results apply to the combined effects of the polymers and LPS on the cells. Regarding the intrinsic effects of the polymers, PCL did not induce any increase in TNF secretion, with polylactide and PS did.⁹⁷ Interestingly, this increase in TNF secretion was observed at the nontoxic concentration of 50 µg ml⁻¹,⁹⁷ but not at the somewhat toxic dose of 80 µg ml⁻¹ used here. Mechanical effects due to particles with sharp edges and inducing an inflammatory response⁹⁸ cannot be excluded in a larger picture of the effects of plastic particles.

Oxidative stress was minimally impacted by treatment with the PCL nanoparticles, with a slight (−10%) but significant decrease in the level of oxidative stress. This is in contrast with the slight increase observed in response to PS particles, and the absence of oxidant response observed upon treatment with PLA nanoparticles. While oxidative stress is generally viewed as a deleterious signal, in macrophages it can be interpreted as a consequence of the oxidative burst that macrophages trigger upon ingestion of particulate material.⁹⁹ Thus, the low level of oxidative stress observed in PCL-exposed cells, even below the control level, may well be another sign of the depression of the immune functions of the macrophages induced by PCL.

Quite interestingly, PCL beads were found to degrade in a few days within the macrophages, although more slowly than PLA beads.¹⁰⁰ This reproduces the well-known behavior observed in medical nanoparticles.¹⁰¹ The degradation product of PCL is 6-hydroxyhexanoic acid,¹⁰² which is then oxidized in adipic acid.¹⁰¹ Adipic acid is known to have a complex metabolism including direct excretion and further



degradation by beta-oxidation.¹⁰³ However, its direct effects on mammalian cells are not known to date. Oppositely, 6-hydroxyhexanoic acid has been shown to exhibit a very low toxicity¹⁰⁴ and to be the active component explaining the anti-inflammatory effects of *Streptococcus gordonii* supernatants.^{105,106} In line with the metabolic effects observed with 6-hydroxyhexanoic,¹⁰⁶ the metabolism-associated proteins (Table S4†) that were most induced in response to PCL nanoparticles were the first enzymes of glycolysis (hexokinase, phosphofructokinase, glucokinase) and the fatty acid-CoA ligases, *i.e.* proteins implied in fatty acid metabolism.

This degradation of PCL nanoparticles within the macrophages prompted us to investigate the delayed effects on macrophages at this time point where the PCL nanoparticles have been extensively degraded, in comparison with PS nanoparticles which we know not to be degraded at this time point.⁴²

Interestingly, we also detected an anti-inflammatory effect in our system, both immediately after exposure and a few days after exposure, when the PCL particles have been extensively degraded and the metabolite released. At this late time point, we also detected a metabolic effect in the form of mitochondrial hyperpolarization, which was neither observed immediately after exposure to PCL or PS nanoparticles, nor observed for PS beads in the delayed scheme. Oppositely, delayed effects, *e.g.* on phagocytosis and in basal pro-inflammatory cytokine production, were detected with PS particles but not with PCL particles, and contrast with the effects detected immediately after exposure. This may suggest that cellular stimulation occurs as a delayed response, which may explain why inflammatory responses are detected *in vivo* (*e.g.* in ref. 107), but not *in vitro* immediately after exposure. Overall, this further strengthens the interest in studies in which delayed effects are taken into account.

Last but certainly not least, it shall be kept in mind that the results described here have been obtained on nanospheres of pristine PCL. As such, this work describes the intrinsic effects of the pure polymer, *i.e.* the first, required step when evaluating the effects of micro and nanoplastics. While this work represents rather well PCL used in medical applications, it will not represent as well the complete effects of PCL micro and nanoplastics arising from other applications. This discrepancy arises from two main phenomena.

The first one is the presence of soluble chemicals in real-life particles, in addition to the polymer itself. These chemicals may be either additives used purposely for technological purposes, or environmental chemicals that adsorb on the plastic during its environmental weathering. Such eco-corona can be extremely complex,¹⁰⁸ which will induce combinatorial explosion problems when trying to assess the toxicity of real-life plastic particles, which detailed chemical composition may be highly variable. Despite this complexity, scientists have tried to evaluate the potential synergies between plastics and their adsorbed chemicals. The

results obtained so far show a variable response,¹⁰⁹ and do not show any universal trend. In some cases a toxic synergy is observed,¹¹⁰ while in other cases plastics decrease the toxicity of chemicals,^{111,112} probably by a strong adsorption that prevents the release of the toxic chemicals, at least for the duration of the experiments.

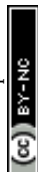
The second one is the shape factor. In macrophages, it has been demonstrated on mineral nanomaterials such as silver nanowires¹¹³ or titanium dioxide,¹¹⁴ but also on cellulose,^{115,116} that a long aspect ratio is an intrinsic determinant for a pro-inflammatory effect. Sharp edges have also been described as a determinant for pro-inflammatory effects.¹¹⁷ Such effects have not been documented as well for microplastics so far, although some studies on fibers only are available.¹¹⁸ However, the fact that the only plastic particle for which we have observed a pro-inflammatory effect so far is PET nanoparticles,¹¹⁹ which are not spherical and do show edges,¹²⁰ argues in favor of the hypothesis that a spherical shape minimizes the effects of the nanomaterials on macrophages. Thus, our results may overall minimize the effects of real-life PCL plastics on macrophages.

5. Conclusions

The first important conclusion of this work on PCL particles is that they are slowly biodegradable within macrophages. However, this does not mean that they do not show immediate and delayed consequences on the physiology of macrophages. Comparison with PS nanoparticles shows that there are both common consequences and specific ones. Among the shared consequences, the induction of the fatty acyl CoA synthases can be noticed. As these enzymes are implied in both fatty acid synthesis and degradation, it can be suggested that they are implied in the biosynthesis of the lipids that are needed to build the vesicles containing the internalized particles. However, most of the parameters that we explored were altered differently by PCL and PS beads. This includes phagocytosis, cytokine production, and oxidative stress, and all these parameters suggested that PCL beads induced a reduced response of macrophages to further stimulation. Furthermore, delayed effects were observed after PCL internalization and degradation, but also for PS particles, which are not degraded. These results suggest that repeated exposure may be an interesting way to move forward, as responses to biodegradable plastics may differ even more from responses to non-biodegradable plastics.

Data availability

The proteomic data have been deposited in the ProteomeXchange Consortium database¹²¹ and are available through the DOI: <https://doi.org/10.6019/PXD049997>. The non proteomic data have been deposited in BioStudies¹²² under the DOI: <https://doi.org/10.6019/S-BSST1804>.



Author contributions

TR designed and supervised the study. MB, MC and DF carried out the characterization of the nanoparticles. VCF, HD and SC performed the proteomic experiments, which were then analyzed by TR and ED. VCF performed and interpreted the flow cytometry experiments. TR drafted the initial version of the manuscript, which was completed and amended by all co-authors, who approved the final version of the manuscript.

Conflicts of interest

There are no conflicts of interest to declare.

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