

Exposure to atmospheric Ag, TiO2, Ti and SiO2 engineered nanoparticles modulates gut inflammatory response and microbiota in mice

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▶ To cite this version:

Eva Guilloteau, Madjid Djouina, Ségolène Caboche, Christophe Waxin, Karine Deboudt, et al.. Exposure to atmospheric Ag, TiO2, Ti and SiO2 engineered nanoparticles modulates gut inflammatory response and microbiota in mice. Ecotoxicology and Environmental Safety, 2022, 236, pp.113442. 10.1016/j.ecoenv.2022.113442 . hal-03627147

HAL Id: hal-03627147 https://hal.science/hal-03627147

Submitted on 22 Jul 2024

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Version of Record: https://www.sciencedirect.com/science/article/pii/S0147651322002822 Manuscript_8dbbd5059509c9d4edbff46e9265f3b2

Revised manuscript 03/03/2022

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27 Abstract

The development of nanotechnologies is leading to greater abundance of engineered 28 nanoparticles (EN) in the environment, including in the atmospheric air. To date, it has been 29 shown that the most prevalent EN found in the air are silver (Ag), titanium dioxide (TiO₂), 30 titanium (Ti), and silicon dioxide (SiO₂). As the intestinal tract is increasingly recognized as a 31 target for adverse effects induced by inhalation of air particles, the aim of this study was to 32 assess the impact of these 4 atmospheric EN on intestinal inflammation and microbiota. We 33 assessed the combined toxicity effects of Ag, Ti, TiO₂, and SiO₂ following a 28-day inhalation 34 35 protocol in male and female mice. In distal and proximal colon, and in jejunum, EN mixture inhalation did not induce overt histological damage, but led to a significant modulation of 36 inflammatory cytokine transcript abundance, including downregulation of $Tnf\alpha$, $Ifn\gamma$, $Ill\beta$, 37 *Il17a*, *Il22*, *IL10*, and *Cxcl1* mRNA levels in male jejunum. A dysbiosis was observed in cecal 38 microbiota of male and female mice exposed to the EN mixture, characterized by sex-dependent 39 modulations of specific bacterial taxa, as well as sex-independent decreased abundance of the 40 Eggerthellaceae family. Under dextran sodium sulfate-induced inflammatory conditions, 41 exposure to the EN mixture increased the development of colitis in both male and female mice. 42 Moreover, the direct dose-response effects of individual and mixed EN on gut organoids was 43 studied and Ag, TiO₂, Ti, SiO₂, and EN mixture were found to generate specific inflammatory 44 responses in the intestinal epithelium. These results indicate that the 4 most prevalent 45 atmospheric EN could have the ability to disturb intestinal homeostasis through direct 46 modulation of cytokine expression in gut epithelium, and by altering the inflammatory response 47 and microbiota composition following inhalation. 48

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50 Keywords: Engineered Nanoparticles. Inhalation. Inflammation. Colitis. Organoid. Mixture.
51 Microbiome. Digestive tract. Intestine.

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1. Introduction

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The International Organization for Standardization defines a nanoparticle as a nanomaterial 55 with all three dimensions in the nanoscale, which is approximatively 1 to 100 nm (Boverhof et 56 al., 2013). Nanoparticles that are synthesized artificially for commercial use are called 57 engineered nanoparticles (EN) (Prajitha et al., 2019). EN are prepared deliberately at the 58 nanoscale because they exhibit properties that provide technological advantages compared with 59 the bulk form of the same material (The Royal Society and The Royal Academy of Engineering, 60 2004). Engineered nanoparticles usage is widespread, covering the energy sector, 61 62 telecommunications industry, computing, agrichemicals (nanopesticides and nanofertilizers), and personal care products (Hansen et al., 2016). Nanotechnologies have also provided new 63 64 opportunities in medicine mainly through the use of EN for diagnostic and therapeutic purposes (biomedical imaging, drug delivery, and targeting) (Rudramurthy and Swamy, 2018). 65 Nanotechnology has become a key component in the modern world and the global 66 nanotechnology market is expected to exceed USD \$125 billion by 2024 (Inshakova et al., 67 2020). In everyday life, humans are exposed to thousands of nanoparticles via dermal contact, 68 inhalation, and ingestion (including parenteral administration) (Malakar et al., 2021). Although 69 advances in the field of nanotechnology have resulted in a myriad of possibilities for 70 technological improvement, the introduction of a wide range of novel EN to the environment 71 and to humans, either by design or inadvertently, raises the possibility of harmful and/or 72 unforeseen adverse effects. Indeed, inhalable nanoparticles can be transported in blood and the 73 lymphatic system and bypass barriers within the body which are generally impermeable to 74

larger particles (Bakand et al., 2012). Especially noteworthy, studies have highlighted that 75 ambient particulate matter (PM) exerts a deleterious impact on the intestinal tract of mice. 76 Ambient ultrafine PM is similar in size to EN, but it is derived from natural and combustion 77 processes. It has been shown in mice that inhalation of ambient ultrafine PM shortened villus 78 length and infiltrated macrophages and neutrophils in the small intestine (Li et al., 2015). 79 Moreover, another study in mice showed that ambient ultrafine PM inhalation led to epithelial 80 injury with inflammatory cell infiltration and mucus depletion in the colon (X. Li et al., 2019). 81 Regarding EN, toxicological studies have highlighted organ-specific toxicity of various 82 nanoparticles following inhalation (Murugadoss et al., 2017; Kim et al., 2018; Hadrup et al., 83 84 2020; Sutunkova et al., 2020), but the impact of EN inhalation on the intestine has never been explored. 85

In this study, we focused on the most abundant EN found in air. Despite the general lack of 86 87 qualitative and quantitative data on human exposure to EN, Hansen et al. (2016) have established "The Nanodatabase" (http://www.nanodb.dk), which is an online inventory of 88 products claimed by manufacturers to contain nanomaterials or include nanotechnology. This 89 database indicates that for inhalation, silver (Ag) is the most prevalent nanomaterial followed 90 by titanium (Ti), titanium dioxide (TiO₂), and silicon dioxide (SiO₂) (Hansen et al., 2016). Ag 91 and Ti nanoparticles are known for their biological and biomedical properties (Pereira et al., 92 2018). Both are often present in consumer products such as cosmetics, food additives, kitchen 93 utensils, and toys. Ag nanoparticles exhibit antimicrobial activity and are used in the areas of 94 food preservation, personal care products (shampoos, lotions, and toothpaste), spray 95 disinfectants, antibacterial surfaces and textiles, nanomedicine, and dentistry (Lee and Jun, 96 2019; Rogers et al., 2018; Tang and Zheng, 2018; Yin et al., 2020). Ti is utilized within artificial 97 implants in dental and orthopaedic surgery (Chouirfa et al., 2019). SiO₂ (including synthetic 98 amorphous silica, or SAS) and TiO₂ are produced in high volumes and applied in many 99

consumer and food products. SiO₂ nanoparticles are used as food additive (E551) to prevent 100 poor flow or caking particularly in powdered products. TiO₂ nanoparticles are extensively used 101 as a pigment, a thickener, and an ultraviolet absorber in cosmetic and skin care products. TiO₂ 102 allows osseointegration of artificial medical implants and bone (Gupta and Xie, 2018). In 103 addition, TiO₂ and SiO₂ are used in protective coatings and for environmental treatment to 104 protect products against environmental damage or in household water and air treatment. Both 105 of these substances have been found in human tissues including jejunum and ileum (Peters et 106 107 al., 2020). The aim of this study was to determine if the presence of these 4 EN in ambient air can result in intestinal adverse health effects. For this purpose, mice were exposed by inhalation 108 to a mixture of Ag, Ti, TiO₂, and SiO₂ nanoparticles, and the impact on intestinal tissues was 109 analysed at steady-state and in the context of dextran sulfate sodium (DSS)-induced colitis, a 110 model commonly used to mimic acute intestinal inflammation in mice. Moreover, the effects 111 112 of individual Ag, Ti, TiO₂, and SiO₂ EN were studied in intestinal organoids, the most advanced in vitro model for gut epithelium characterization. 113

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2. Materials and methods

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117 2.1. Characterization of EN by transmission electron microscopy (TEM)

SiO₂ in water (average particle size (APS) 20 nm), TiO₂ (anatase; APS 5-30 nm) in water, Ti
powder (APS 60-80 nm), and Ag powder (APS 50-60 nm) were purchased from IoLiTec
Nanomaterials. For physicochemical characterization, each nanoparticle sample was diluted in
ultrapure water and deposited on TEM grids (Copper 200 mesh grids with carbon film, Agar
ScientificTM). Once dried, each sample was observed by scanning electron microscopy (FEGSEM), using a JEOLTM JSM7100F electron microscope equipped with a transmitted electron
detector and 3 X-ray energy dispersive spectrometers (BrukerTM XFlash 6/30), enabling the

analysis of elements with atomic numbers greater than boron ($Z \ge 5$). The microscope was operated at 15 kV with a probe current of 300 pA and a working distance suitable for X-ray detection (10 mm). All nanoparticle suspensions were used within 1 h after sonication.

128

129 2.2. Animals and experimental design

The animal protocol was approved by the regional bioethics committee (committee no.75; 130 authorization no. CEEA 2016072517274040) and the animals received human care in 131 accordance with the Guide for the Care and the Use of Laboratory Animals (National Research 132 Council (US) Committee, 2011). Male and female C57BL/6JRj mice (aged 7 weeks) were 133 purchased from Janvier Labs and housed under standard conventional conditions (Suppl. Fig. 134 1A). The room relative humidity was 55% and the temperature was 21°C. Mice were randomly 135 divided into the different exposure groups (n=10/group). Mice were placed 4 h/day, 5 136 137 days/week for 28 days in a ventilated whole-body inhalation chamber that allowed free movement (inExpose, SCIREQ Scientific®). Nebulization was achieved using an Aeroneb 138 LabTM ultrasonic nebulizer (SCIREQ Scientific®) directly connected to a 5 L chamber and 139 140 controlled through flexiWare software v.6 (SCIREQ Scientific®) according to the following parameters: bias flow of 2 L/min and nebulization rate of 0.083 ml/min, which we measured 141 and controlled throughout the experiment. These parameters allowed the nebulization of the EN 142 mixture at a concentration of 1 mg/m^3 in the chamber. Mice were euthanized the morning 143 following the final exposure day. The distal colon, proximal colon, and jejunum were collected 144 and stored for subsequent analyses. 145

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147 **2.3. Gut histological analysis**

Formaldehyde-fixed intestinal tissues were embedded in paraffin, serially sectioned (4 µm),
and stained with May-Grünwald and Giemsa (MGG). An inflammatory infiltration score was

blindly determined according to Li et al. (X. Li et al., 2019). This score is the sum of the infiltration score plus the percent area involved of each section. Infiltration was scored on a scale of 0 to 3 (0, no inflammatory cells; 1, infiltration around crypt bases; 2, infiltration of muscularis mucosa; 3, infiltration of submucosa), and the percent area of each section was scored on a scale of 0 to 4 (0, no involvement; $1, \le 25\%$; $2, \le 50\%$; $3, \le 75\%$; $4, \le 100\%$). Scoring was performed on distal colon, proximal colon, and jejunum (magnification 20X; 3 to 7 fields per section; 4 to 6 mice per group).

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2.4. Quantification of transcriptional levels of cytokines and oxidative stress markers by real-time PCR

Total mRNA from intestinal tissues and organoids was extracted using a Nucleospin RNA II 160 kit (Macherey-Nagel) according to manufacturer protocol. Reverse transcription was performed 161 162 using a High Capacity cDNA Archive Kit and quantitative polymerase chain reaction (qPCR) with SYBR Green (Thermo Fisher Scientific). The primer sequences were designed using 163 Primer Express 3 (Thermo Fisher Scientific) and are available upon request. Melting curve 164 analyses were performed for each sample and gene to confirm the specificity of the 165 amplification. Because exposure to EN mixture (ENM) did not cause any significant alterations 166 in Polr2a mRNA expression, the relative expression of each gene of interest was normalized to 167 the relative expression of this gene. The quantification of the target gene expression was based 168 on the comparative cycle threshold (Ct) value. The fold changes in the target genes were 169 analysed by the $2^{-\Delta\Delta Ct}$ method. 170

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172 2.5. Bacterial DNA extraction and Illumina MiSeq sequencing

173 Genomic DNA was extracted from cecal content using the DNA Stool kit (Macherey-Nagel)

according to manufacturer protocol. The quantity and the purity of DNA (expressed as the ratio

of absorbance at 260 nm and 280 nm (A260/A280)) were assessed using a NanoDrop® 175 spectrophotometer (Ozyme). The sequencing library was generated by amplifying the V3-V4 176 region of the bacterial 16S-rRNA gene using 16S rRNA amplicon generation for MiSeq with 177 (5'-CCTACGGGNGGCWGCAG-3') 178 the primers Bact-0341 and Bact-0785 (5'-GACTACHVGGGTATCTAATCC-3'). Individual samples were barcoded, pooled to construct 179 the sequencing library, and sequenced using an Illumina MiSeq system to generate paired-end 180 181 2x300 bp reads.

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183 **2.6.** Analysis of sequencing data

Bioinformatic analyses were performed using the OIIME 2 pipeline (v.2020.2) (Bolven et al., 184 2019). The Divisive Amplicon Denoising Algorithm 2 (DADA2) plug-in in QIIME2 was used 185 to filter, dereplicate, identify chimeric sequences, and merge paired-end reads to obtain the set 186 187 of amplicon sequence variants (ASVs) for each sample (Callahan et al., 2016). Then the representative sequences were picked for each ASV. The classify-sklearn plug-in in QIIME2, 188 with the SILVA database (v.132) was applied to assign a taxonomic annotation to each 189 190 representative ASV sequence. In the next step, ASVs identified as eukaryotic contamination (no contamination) and external contamination, identified with the decontam package (15 ASV; 191 192 1599 reads), were filtered out (Davis et al., 2018). Diversity metrics (α and β) were obtained with the QIIME2 core-metrics-phylogenetic plug-in, with p-sampling depth parameter fixed to 193 11916 reads which corresponded to the total frequency that each sample should be rarefied to 194 before computing diversity metrics. This sampling depth allowed retention of >59% of reads 195 discarding no sample. 196

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198 2.7. Dextran sodium sulfate (DSS)-induced colitis

Colitis was induced by adding 2% (w/v) DSS (TdB Labs) in the drinking water for 6 days.
Fresh DSS solutions were prepared every 2 days. Mice body weight was recorded during the
experiment. At necropsy, colon was carefully dissected and its weight and size were measured.
Colon was then either immediately frozen or fixed in 4% formaldehyde (VWR Chemicals).

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204 2.8. Organoid study

Murine organoids were derived from colon tissue of male C57BL/6 mice (aged 7 weeks). As 205 206 described in Co et al. (2019), tissue fragments were incubated in 2 mM EDTA in cold chelation buffer on ice for 30 minutes (Co et al., 2019). The EDTA buffer was removed and tissue 207 fragments were vigorously shaken in cold chelation buffer to isolate intestinal crypts. Isolated 208 crypts were embedded in Matrigel (Growth Factor Reduced; Corning) on ice and seeded into a 209 24-well tissue culture plate. Then 500 µL of growth media and 50% L-WRN-conditioned media 210 211 were added. Colon organoids were grown embedded in Matrigel for 7-14 days with growth media. Matrigel-embedded organoids were solubilized in 5 mM EDTA in PBS for 1 h at 4°C 212 213 on a rotating platform. Organoids were centrifuged at 200 x g for 3 min at 4°C and the 214 supernatant was removed. The pellet was resuspended in growth media in ultra-low attachment 24-well tissue culture plates (Corning Costar). Suspended organoids were incubated at 37°C 215 with 5% CO₂ for 2 days so that they could invert and present the apical side to the exterior. 216 217 Inverted organoids were treated with 0.02, 0.2, and 2 µM of Ag, SiO₂, Ti, and TiO₂ EN for 24 h. Cell death was measured using the Cytotoxicity Detection Kit (LDH) (Roche) according to 218 the manufacturer's protocol. 219

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221 **2.9. Statistical analysis**

Results are expressed as mean \pm *SEM*. Normality tests were performed for each parameter using the D'agostino-Pearson normality test. Since the data was not normally distributed, the statistical significance of differences between experimental groups was calculated using theMann-Whitney U test (GraphPad, San Diego, CA).

226

227 **3. Results**

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229 **3.1. Characterization of EN**

Figure 1A shows typical TEM images and EDX spectra of the 4 EN samples. Image analysis 230 of 308 Ag particles indicated an area equivalent mean diameter of 8.6 nm (SD=2.4 nm). Image 231 analysis of 261 SiO₂ particles yielded a mean equivalent diameter of 21.1 nm (SD=6.0 nm). In 232 the case of Ti samples, we noticed the presence of aggregated spherical Ti nanoparticles with 233 individual particle sizes ranging from 5 nm to 350 nm. Despite several attempts to deposit a 234 uniform layer of TiO₂ particles, the noisy TEM image did not allow for statistical size analysis 235 236 but an average equivalent diameter of 14.6 nm was estimated by manual sizing. X-ray spectra did not show the presence of contaminant elements. The copper peak was attributed to the TEM 237 238 copper mesh support grid. As shown in the elemental map of the ENM (Fig. 1B), all expected 239 elements (Si, Ag, Ti, and O) were identified by X-ray emission spectrometry. Chemical and topographic maps of the ENM showed spherical particles containing titanium (Ti or TiO₂) 240 covered with Ag and SiO₂ nanoparticles. 241

242

243 **3.2. Effects of ENM exposure on gut inflammation**

Adult male and female mice were exposed by inhalation to the ENM or to control inhalation (Suppl. Fig. 1A). Body weight over the 28-day duration of the experiment did not vary significantly in males or females (p=0.7 for males, p=0.4 for females, Suppl. Fig. 1B). To assess the inflammatory status of intestinal tissues, we performed a histological scoring of inflammation in distal colon, proximal colon, and jejunum. In males, the inflammatory

infiltration score did not change between the groups (p=0.8 for distal colon, p=0.4 for proximal 249 250 colon, Fig. 2A-C), even if a trend to enhancement of this score was observed in jejunum (p=0.1). For comparison of cytokines, in distal colon, transcript levels of $Tnf\alpha$, $Ifn\gamma$, Il22, and IL10 were 251 significantly increased in ENM-exposed mice compared to control mice (p=0.007, 0.0002, 0.01 252 and 0.03 respectively, Fig. 2D). In proximal colon, there was a significant enhancement of the 253 mRNA levels of $Tnf\alpha$, $Ill\beta$, Ill0, and Cxcl1 (p=0.04, 0.007, 0.001 and 0.03 respectively, Fig. 254 2E). In jejunum, ENM exposure led to a drastic decrease of $Tnf\alpha$, $Ifn\gamma$, $II1\beta$, II17a, II22, IL10, 255 and Cxcl1 mRNA expression (p=0.04, 0.004, 0.001, 0.004, <0.0001, <0.0001, 0.002 256 respectively, Fig. 2F). 257

In female mice, exposure to ENM did not induce significant variation of inflammatory infiltration score in distal colon, proximal colon, and jejunum (p=0.4 for distal colon and jejunum, p=0.6 for proximal colon, Fig. 3A-C). *Cxcl1* transcripts were down-regulated in distal colon of ENM-exposed mice (p=0.01, Fig. 3D). A decrease of $Il1\beta$ in proximal colon and a reduction of $Tnf\alpha$, $Il1\beta$ and IL22 in jejunum of ENM-exposed mice compared to control mice was also observed (p=0.005, 0.0002, 0.01 and 0.009 respectively, Fig. 3E-F).

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265 **3.3. Effects of ENM exposure on gut microbiota.**

The bacterial diversity in cecal content was determined by pyrosequencing of the bacterial 16S 266 267 rRNA genes. Exposure to ENM did not significantly affect α -diversity (data not shown). In males, unweighted uniFrac index showed a significant increase of β-diversity in ENM-exposed 268 mice compared to control mice (p=0.004; Fig. 4A). The relative abundance of ASVs at phylum, 269 class, order, and family taxonomic levels was compared. The major phyla were Firmicutes and 270 271 Bacteroidetes and the Tenericutes were less abundant in ENM-exposed mice (Fig. 4B). ENM exposure also induced in male mice a decrease of Coriobacteriia and Mollicutes classes, and of 272 Coriobacteriales and Mollicutes RF39 orders (Fig. 4C-D). The only family which showed 273

significantly different abundance in exposed male mice was Eggerthellaceae (Fig. 4E). In 274 females, the unweighted uniFrac index was increased significantly after exposure to ENM 275 (p=0.008; Suppl. Fig. 2). As expected, in females, the cecum microbiota was also mostly 276 composed of bacteria belonging to Firmicutes and Bacteroidetes phyla (Fig. 5A). The relative 277 abundance of Proteobacteria and Actinobacteria was decreased in ENM-exposed mice. At the 278 class level, abundances of Deltaproteobacteria and Coriobacteriia were lower after exposure to 279 ENM (Fig. 5B). A significant reduction of the levels of Desulfovibrionales and Coriobacteriales 280 orders in ENM-exposed female mice was also observed (Fig. 5C). Finally, like males, females 281 showed a lower abundance of Eggerthellaceae family after exposure to ENM (Fig. 5D). 282

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284 **3.4.** Effects of ENM exposure in a mouse model of colitis

Male mice were submitted to DSS administration in the last 6 days of the ENM inhalation protocol (Suppl Fig. 3A). A significant body weight loss was observed in ENM-exposed mice compared to control mice from the third to the last day of DSS treatment (p=0.004 at day 6, Fig. 6A). The colon weight/size ratio and the histological inflammatory infiltration score were not modified between the 2 groups of mice (p=0.2 and 0.08 respectively, Fig. 6B and Suppl Fig. 3B). The transcript levels of *Tnf* α and *Ifn* γ were higher in colitic ENM-exposed mice than in colitic control mice (p=0.002 and 0.01 respectively, Fig. 6C).

The same protocol was performed with female mice (Suppl Fig. 3C). Colitic ENM-exposed mice lost more weight than colitic control mice from the fourth to the last day of DSS treatment (p<0.0001 at day 6, Fig. 6D). The colon weight/size ratio was significantly increased in the ENM-exposed group, as well as the histological score of colitis (p=0.001 and 0.048 respectively, Fig. 6E and Suppl Fig. 3D). The expression of inflammatory cytokines was upregulated in the colon of ENM-exposed mice, particularly *Ifn* γ and *Cxcl1* (p=0.004 and 0.0003 respectively, Fig. 6F). These data show that ENM exposure induces higher susceptibility to colitis development, and that this increased sensitivity to colitis is greater in females than inmales.

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302 **3.5. Effects of EN on gut organoids**

Then the effects of individual EN and ENM on intestinal epithelium were assessed. For this 303 purpose, mouse intestinal organoids which model in vitro a 3D microarchitecture and possess 304 a cellular composition resembling native intestinal epithelium were used. The organoids were 305 generated from pluripotent stem cells which were differentiated into enterocytes, stem cells, 306 goblet cells, Paneth cells, enteroendocrine cells, and mesenchymal cells. Organoids were 307 treated with 0.02, 0.2, and 2 µM of Ag, SiO₂, Ti, TiO₂ EN and ENM for 24 h. These 308 concentrations did not induce cytotoxicity as confirmed by the LDH assay (Suppl. Fig. 3). Ag 309 enhanced the expression of the chemokines Ccl5, Ccl20, and Ccl10 and of the alarmins $Ill\alpha$ 310 and Il33 (p=0.03 for all genes at 0.2 μ M, Fig. 7A). SiO₂ increased the transcript levels of Tnf α 311 and *Cxcl10* even at the lowest dosage, and enhanced *Il1* α expression (*p*=0.03 for all genes at 2 312 μM, Fig. 7B). Ti increased *Tnfα*, *Ccl5*, *Cxcl10*, and *Il33* and decreased *Cxcl1* mRNA levels 313 $(p=0.03 \text{ for all genes at } 0.2 \,\mu\text{M}, \text{Fig. 7C})$. TiO₂ downregulated and upregulated the expression 314 of Cxcl1 and Il33, respectively (p=0.03 for both genes at 0.2 µM, Fig. 7D). The mixture of the 315 4 EN induced significant variations mainly at 2µM (Fig.7E). It totally annihilated the CCl5 and 316 $ll \alpha$ mRNA expression, which became undetermined. It increased the transcript levels of 317 *CCl20*, *Cxcl1* and *Il33* (*p*=0.03 for all genes at 0.2µM). 318

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4. Discussion

The fate of inhaled nanoparticles has been studied in lung, brain, and at the systemic level, but little is known about responses in the digestive tract (Aalapati et al., 2014; Cupaioli et al., 2014; Miller et al., 2017). However, it has been shown that inhaled particles can be directed to the

intestinal tract via the mucociliary clearance process (Smith et al., 2002). This process mainly 324 affects larger particles, but also includes nanoparticles. For example, detection of TiO₂ 325 nanoparticles in feces following inhalation demonstrated a mucociliary clearance and ingestion 326 of nanoparticles (Pujalté et al., 2017). Moreover, inhaled Ag nanoparticles are cleared via the 327 larynx into the gastro-intestinal tract, and in addition, via blood, liver, and gall bladder into the 328 gastro-intestinal tract with one common excretional pathway: via feces out of the body 329 (Kreyling et al., 2020). Nanoparticles can translocate into the bloodstream from which they can 330 then impact all tissues of the body, either by direct contact or by various indirect routes (Miller 331 et al., 2017). The constant increase in EN production worldwide has led to the presence of Ag, 332 SiO₂, Ti, and TiO₂ nanoparticles in the air. In addition to potential high workplace exposures to 333 each of these EN, chronic low-dose exposure of the general population may occur if current 334 production trends continue. It is therefore important to assess the health effects of such 335 336 exposures, and particularly on the intestinal tissues which are increasingly recognized as an important target for inhaled particles (Feng et al., 2020; Vignal et al., 2021). In this study, 337 338 healthy adult mice were exposed for 4 h per day for 28 days to a mixture of the 4 EN most 339 commonly found in ambient air, each at a dose of 1 mg/m³. The range of EN present in the air is not known, so it is not currently possible to assess the relevance of this dose to human 340 exposure. However, this dosage is within the range that has been used in other similar studies. 341 For example, Umezawa et al. (2018) exposed mice in an inhalation chamber to 4.6 or 37 mg/m³ 342 of Printex 90 carbon black nanoparticles for 15 days (Umezawa et al., 2018). Wahle et al. (2020) 343 exposed mice via nose only inhalation to CeO₂ nanoparticles with varying amounts of Zr-344 doping (0%, 27%, or 78% Zr) over a 4-week period (4 mg/m³ for 3 h/day, 5 days/week) (Wahle 345 et al., 2020). These experimental protocols allow a primary toxicological evaluation ahead of 346 347 more in-depth dose-response studies.

At the histological level, we did not observe any significant differences between exposed and 348 control mice, showing that inhalation of ENM did not induce drastic damage in intestinal 349 tissues. However, more sensitive methods such as qPCR allowed us to highlight sex-specific 350 modifications. In males, we observed upregulation of numerous inflammatory cytokines in both 351 distal and proximal colon. By contrast, in male jejunum, all the inflammatory markers were 352 found strongly downregulated in mice exposed to ENM. In females, the pattern of inflammatory 353 biomarkers was only slightly modified in distal and proximal colon. Moreover, inhalation of 354 355 ENM induced a drastic decrease in expression of several inflammatory cytokines in jejunum, similarly to males. Thus, under basal conditions, inhalation of ENM did not induce overt 356 inflammation in intestinal tissue. However, among the few inflammatory markers quantified, 357 several were found to be significantly modified in favor of micro-inflammation, particularly in 358 male colon. In accordance with these data, a micro-inflammatory state has been described after 359 360 oral exposure to TiO₂, SiO₂, and Ag nanoparticles (Lamas et al., 2020).

Modifications of cecal microbiota composition were compared between exposed and non-361 exposed mice at several taxonomic levels. Males exposed to ENM showed a decreased 362 abundance of the Tenericutes phylum, Mollicutes class, and Mollicutes RF39 order. In mice, 363 Mollicutes have been shown to be decreased in TNBS-induced colitis (P. Li et al., 2019) and 364 after subchronic oral toxic metal exposure (Zhai et al., 2017). The abundance of Mollicutes is 365 decreased in fecal microbiota of patients with colon Crohn's disease and ulcerative colitis, and 366 shown to be increased in a patient with ileal Crohn's disease (Willing et al., 2010). Gut 367 microbiota of obese patients also shows a greater proportion of Mollicutes (Crovesy et al., 368 2020). Moreover, we observed a reduction of Proteobacteria phylum, Deltaproteobacteria class, 369 and Desulfovibrionales order in females exposed to ENM. The presence of Desulfovibrionales 370 bacteria in gut microbiota has been associated with Parkinson's disease (Murros et al., 2021) 371 and autism spectrum disorder (de Theije et al., 2014). A 16S microbiota sequencing study has 372

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also shown that bacteria of the Desulfovibrionales order were present within the ileal 373 submucosal tissues of most Crohn's disease patients but absent in the control group (Chiodini 374 et al., 2015). Notably, we identified microbiota modifications by ENM exposure which were 375 common to both males and females. Microbiota of ENM-exposed males was characterized by 376 less abundance of Coriobacteriia class, Coriobacteriales order, and Eggerthellaceae family. 377 Similarly, microbiota of ENM-exposed females exhibited lower levels of Actinobacteria 378 phylum, Coriobacteriia class, Coriobacteriales order, and Eggerthellaceae family. The 379 abundance of Eggerthellaceae has been associated with the intensity of 5-fluorouracil-induced 380 intestinal mucositis in mice (Chen et al., 2021). It was found to be increased in gut of patients 381 with psoriasis (Sikora et al., 2020). Furthermore, this family has been suggested to play a critical 382 role in occurrence and development of osteoporosis (Wen et al., 2020). Taken together, these 383 results show that ENM exposure induced sex-dependent and -independent modifications of 384 385 fecal microbiota composition, which could play a role in the micro-inflammatory state observed in mouse gut. 386

The sex-dependency of EN toxicity has been widely observed (Bao et al., 2020; Lovaković et 387 al., 2021; Ma et al., 2018). This variability between males and females is often attributed to the 388 differences in sexual hormone signalling. Androgens, estrogens, and progesterone modulate 389 intestinal inflammation and shape gut microbiota (Pace and Watnick, 2021). Beyond the 390 differences in the expression of sex hormones, a sexual dimorphism in peripheral and intestinal 391 immunity, as well as in intestinal microbiome is known (Elderman et al., 2018). The tissue 392 accumulation and clearance of nanoparticles can be also sex-specific (Bao et al., 2020; Lee et 393 al., 2020). In ex vivo ileal tissue, a sex-dependency has been demonstrated for cytokines, 394 chemokines and intestinal permeability genes in response to Ag EN (Gokulan et al., 2021). 395 Finally, the sex-dependency of EN toxicity can be explained by sex specific differences of EN 396

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uptake at the cellular level, as demonstrated in amniotic stem cells and in primary fibroblasts(Serpooshan et al., 2018).

Under inflammatory conditions, both male and female mice exposed to ENM had a greater susceptibility to DSS-induced colitis. Therefore, despite low deleterious effects in healthy individuals, inhalation of ENM caused visible clinical effects in susceptible individuals. These data suggest that inhalation of ENM could aggravate diseases mediated by the immune system such as inflammatory bowel diseases.

We also investigated the potential of each EN to disturb the inflammatory response in intestinal 404 epithelium. Both Ti and TiO₂ decreased expression of *Cxcl1* and increased expression of *Il33*. 405 Ag and Ti had a strong ability to enhance the expression *Cxcl10* (10- and 19-fold, respectively, 406 at $0.2 \mu M$). Overall, TiO₂ had the least impact on intestinal inflammatory response, whereas it 407 408 is the EN with the most documented deleterious effects on intestinal health (Rinninella et al., 2021). Accordingly, Kämpfer et al. observed stronger toxic effects of polyvinylpyrrolidone-409 410 capped Ag engineered nanomaterials than TiO₂ in Caco-2 and HT-29-MTX-E12 monocultures 411 (Kämpfer et al., 2021). Ex vivo, treatment with 20 µg/mL of Ag EN induced sex-dependent alterations of cytokine secretion in human excised intestinal tissues (Gokulan et al., 2021). The 412 mixture of the 4 EN induced a paradoxical response, by enhancing the expression of certain 413 cytokines (16-fold increase of *Il33* levels), and by very strongly reducing the expression of 414 other cytokines: the expression of CCl5 and of $Il1\alpha$ were completely shunted, this latter type of 415 modification was not observed following exposure to EN individually. As a whole, each of the 416 4 tested EN as well as ENM elicited specific responses in colonoids. 417

418 Conclusion

The presence of EN in the air breathed by mice caused sex-dependent intestinal disturbancesregarding inflammatory status and microbiota composition. Likewise, inhalation of ENM

increased susceptibility to DSS-induced colitis in mice. We also showed that exposure to Ag, 421 422 SiO₂, Ti, and TiO₂ EN led to specific inflammatory responses in gut epithelium. These findings suggest that respiratory exposure to common formulations of nanoscale particulates may 423 facilitate the induction of a pro-inflammatory state localized within the intestine, alterations in 424 gut microbiota composition, and modulation of susceptibility to chronic inflammatory 425 conditions with the GI tract. Moreover, many of these effects may be differentially observed in 426 exposed males and females, providing a new evidence of the sex-specificity of gut response to 427 environmental chemicals. 428

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430 Funding

431This work was supported by the Hauts de France Region and the Ministère de l'Enseignement

432 Supérieur et de la Recherche (CPER Climibio), and the European Fund for Regional Economic

433 Development.

434

435 Acknowledgements

436 We thank Thomas Hubert and the staff of the animal facility of Lille for animal care. We thank

437 Bernadette Leu for her broad-spectrum help. Editorial assistance, in the form of language

438 editing and correction, was provided by XpertScientific Editing and Consulting Services.

439

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633 Figure legends

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- Fig. 1. (A) Transmission electron microscopic images and energy-dispersive X-ray spectra of
- typical nanoparticles from individual samples (Ag, SiO₂, Ti, and TiO₂). Scale bars represent
- 637 100 nanometers. (B) Typical transmission electron microscopic images (backscattered electron
- 638 detector for chemical contrast, secondary electron detector for morphological contrast) and
- elemental map of the nanoparticle mixture.

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Fig. 2. The effects of inhalation of an engineered nanoparticle mixture (ENM) on male gut 641 inflammatory status. Male mice were exposed to an airborne ENM consisting of Ag, SiO₂, Ti, 642 and TiO₂ or to control inhalation (CT; n=10/group). (A-B-C) Representative images of 643 intestinal tissues (MGG staining; scale bar=100 µm) and inflammatory infiltration score in 644 distal colon (A), proximal colon (B), and jejunum (C). (D-E-F) Quantification of mRNA levels 645 of inflammatory cytokines by qRT-PCR in distal colon (D), proximal colon (E), and jejunum 646 (F). *=p<0.05, **=p<0.01, ***=p<0.005, and ****=p<0.001 as determined by the Mann-647 Whitney U test. 648

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Fig. 3. The effects of inhalation of an engineered nanoparticle mixture (ENM) on female gut inflammatory status. Female mice were exposed to an airborne ENM consisting of Ag, SiO₂, Ti and TiO₂ or to control inhalation (CT; n=10/group). (A-B-C) Representative images of intestinal tissues (MGG staining; scale bar=100 μ m) and inflammatory infiltration score in distal colon (A), proximal colon (B), and jejunum (C). (D-E-F) Quantification of mRNA levels of inflammatory cytokines by qRT-PCR in distal colon (D), proximal colon (E), and jejunum (F). *=p<0.05, **=p<0.01, and ***=p<0.005 as determined by the Mann-Whitney U test.

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Fig. 4. The effects of inhalation of an engineered nanoparticle mixture (ENM) on male gut microbiota composition. Male mice were exposed to an airborne ENM consisting of Ag, SiO₂, Ti, and TiO₂ or to control inhalation (CT; n=6/group). DNA was isolated from cecal content for MiSeq 16S ribosomal RNA gene sequencing. (A) Unweighted-uniFrac β -diversity index, ***=p<0.005 as determined by pairwise PERMANOVA. The relative abundance of bacteria was calculated based on amplicon sequence variants. (Bb) Overview of the relative abundance of gut bacteria depicted at the phylum level. The relative abundance of significantly changed bacterial phyla (Bb'), classes (C), orders (D), and families (E). *=p<0.05 as determined by the Mann-Whitney U test.

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Fig. 5. The effects of inhalation of an engineered nanoparticle mixture (ENM) on female gut 668 microbiota composition. Female mice were exposed to an airborne ENM consisting of Ag, 669 SiO₂, Ti, and TiO₂ or to control inhalation (CT; n=6/group). DNA was isolated from cecal 670 content for MiSeq 16S ribosomal RNA gene sequencing. The relative abundance of bacteria 671 672 was calculated based on amplicon sequence variants. (Aa) Overview of the relative abundance of gut bacteria depicted at the phylum level. The relative abundance of significantly changed 673 bacterial phyla (Aa'), classes (B), orders (C), and families (D). *=p<0.05 as determined by the 674 Mann-Whitney U test. 675

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677 Fig. 6. The effects of inhalation of an engineered nanoparticle mixture (ENM) on dextran sodium sulfate (DSS)-induced colitis development in males and females. Male and female mice 678 679 were exposed to an airborne ENM consisting of Ag, SiO₂, Ti, and TiO₂ (ENM) or to control 680 inhalation (CT), and to a 2% DSS solution in drinking water in the last 6 days of the experimental protocol (n=10/group). (A) Male body weight variation during 6 days of DSS 681 administration. (B) Male colon weight/size ratio. (C) Quantification of mRNA levels of 682 inflammatory cytokines in male colon by qRT-PCR. (D) Female body weight variation during 683 6 days of DSS administration. (E) Female colon weight/size ratio. (F) Quantification of mRNA 684 levels of inflammatory cytokines in female colon by qRT-PCR. *=p<0.05, **=p<0.01, and 685 ****=p < 0.001 as determined by the Mann-Whitney U test. 686

687

Fig. 7. The effects of the 4 engineered nanoparticles (EN) individually and in mixture on gutorganoid inflammatory response. Quantification of mRNA levels of inflammatory cytokines

- and alarmins by qPCR in inverted organoids treated with 0.02, 0.2, and $2 \mu M$ of Ag, SiO₂, Ti,
- TiO_2 EN, and the engineered nanoparticle mixture (ENM)after 24 h exposure (n=4/condition).
- Results are presented as fold change compared to untreated organoids. *=p<0.05 as determined
- 693 by the Mann-Whitney U test.



^{*} Cu signal from copper TEM grid

Α



Distal colon

Proximal colon

Jejunum



Distal colon

11173

11AB

IMY

110

1122

Ctcl1

0

Thia

Proximal colon

Jejunum

1122

Ctch

110

11173

11P

IFNY

0.0

Thto



11173

11AB

MANY

"10

1122

CXCI1

0.0

Thta









С

F





Ε

В



Α

Females



Exposure to atmospheric Ag, TiO_2 , Ti and SiO_2 engineered nanoparticles modulates

inflammatory response and gut microbiota in mice

