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Effect of Early Particulate Air Pollution Exposure on Obesity in Mice Role of p47^{phox}

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Abstract

Objective—To evaluate the role of early-life exposure to airborne fine particulate matter (diameter, $<2.5 \mu\text{m}$ [$\text{PM}_{2.5}$]) pollution on metabolic parameters, inflammation, and adiposity; and to investigate the involvement of oxidative stress pathways in the development of metabolic abnormalities.

Methods and Results— $\text{PM}_{2.5}$ inhalation exposure (6 h/d, 5 d/wk) was performed in C57BL/6 mice (wild type) and mice deficient in the cytosolic subunit of the nicotinamide adenine dinucleotide phosphate (NADPH) oxidase p47^{phox} (p47^{phox}^{−/−}) beginning at the age of 3 weeks for a duration of 10 weeks. Both groups were simultaneously fed a normal diet or a high-fat diet for 10 weeks. $\text{PM}_{2.5}$ -exposed C57BL/6 mice fed a normal diet exhibited metabolic abnormalities after exposure to $\text{PM}_{2.5}$ or FA for 10 weeks. Consistent with insulin resistance, these abnormalities included enlarged subcutaneous and visceral fat contents, increased macrophage infiltration in visceral adipose tissue, and vascular dysfunction. Ex vivo–labeled and infused monocytes demonstrated increased adherence in the microcirculation of normal diet– or high-fat diet–fed $\text{PM}_{2.5}$ -exposed mice. p47^{phox}^{−/−} mice exhibited an improvement in parameters of insulin resistance, vascular function, and visceral inflammation in response to $\text{PM}_{2.5}$.

Conclusion—Early-life exposure to high levels of $\text{PM}_{2.5}$ is a risk factor for subsequent development of insulin resistance, adiposity, and inflammation. Reactive oxygen species generation by NADPH oxidase appears to mediate this risk.

Keywords

$\text{PM}_{2.5}$ air pollution; obesity; NADPH oxidase; inflammation; insulin resistance

Substantial epidemiological evidence implicates fine particulate matter ($\text{PM}_{2.5}$) air pollution as a major adverse risk factor with serious consequences for human health in both developed

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Disclosures

None.

and developing countries.¹⁻⁴ Recent data from large population cohorts have provided compelling associations between ambient air PM_{2.5} pollution exposure and increased cardiovascular morbidity and mortality.^{5,6} Previously, important mechanistic links between inhaled PM_{2.5} exposure and exposure to high-fat diets (HFDs) in adult mice were demonstrated.^{7,8} Because propensity to insulin resistance (IR) develops via priming mechanisms early in childhood, we speculated that the appropriate time window for assessing the effects of exposure to pollutants in the eventual development of IR in adults is during infancy/childhood. Studying type 2 diabetes mellitus (T2DM)/IR development and its relation to environmental factors may help clarify the causative relations among IR, body fatness, and the development of cardiovascular risk.^{9,10} Inflammation and oxidative stress pathways appear to play critical roles in this process, as demonstrated by numerous investigations.¹¹⁻¹⁴ Consistent with this hypothesis, targeted disruption of proinflammatory or pro-oxidant pathways have been effective in attenuating the development of IR.^{13,15-17} Furukawa et al¹⁸ have shown an important role of the NADPH oxidase in the development of obesity and IR in response to a HFD. We hypothesized that PM_{2.5} air pollution exerts important proinflammatory and pro-oxidant effects, mediated via NADPH oxidase, with early-life exposure eventually priming the development of IR.

Methods

Detailed methods are provided in the supplemental data (available online at <http://atvb.ahajournals.org>).

Animals

We used male 3-week-old C57BL/6 and p47^{phox-/-} mice (both from Jackson Laboratories, Bar Harbor, Me). The protocols and the use of animals were approved by and in accordance with the Ohio State University Animal Care and Use Committee.

Diet and PM_{2.5} Exposure

The mice were fed with either a normal diet (ND) (Teklad 7012, 13% calories from fat; n = 16) or an HFD (Teklad TD 88137, 42% calories from fat; n = 16) beginning at the age of 3 weeks. The animals were exposed by inhalation to either filtered air (FA) or PM_{2.5} for 6 h/d, 5 d/wk, for 10 weeks in a mobile trailer exposure system ("Ohio Air Pollution Exposure System for Interrogation of Systemic Effects 1," located at the Ohio State University Animal Facility in Columbus). Animal exposure and monitoring of the exposure environment and ambient aerosol were performed as previously described.^{7,8,19}

Blood Glucose and Insulin Measurements, Magnetic Resonance Imaging, and Intravital Microscopy

Blood glucose measurement was conducted with a glucometer (Elite). Insulin levels were determined using a commercially available kit (Ultra Sensitive Mouse Insulin ELISA Kit).

An abdominal fat evaluation was performed by in vivo magnetic resonance imaging (MRI) with a T1-weighted gradient-echo sequence on all C57BL/6 and p47^{phox-/-} mice, as previously described.⁸

Intravital microscopy was performed as described in the supplemental materials.

Blood Inflammatory Biomarkers

Blood inflammatory biomarkers, including interferon γ , monocyte chemoattractant protein-1, regulated on activation, normal T-cell expressed and secreted, and tumor necrosis factor (TNF) α , were measured by quantitative ELISA assays.

Statistical Analysis

Data are expressed as mean \pm SE unless otherwise indicated. For responses measured repeatedly at different points or dose levels, a series of 2-sample independent Student *t* tests was used to detect differences between the FA and PM_{2.5} treatment groups at every point and dose level, with Bonferroni correction for multiple-comparison adjustment. Comparisons of other continuous variables were conducted with an independent 2-sample Student *t* test, with *P*<0.05 considered significant.

Results

Exposure Characterization

The ambient mean daily PM_{2.5} concentration at the study site was 15.8 $\mu\text{g}/\text{m}^3$ (SD, 6.1 $\mu\text{g}/\text{m}^3$), whereas the mean concentration of PM_{2.5} in the exposure chamber was 111.0 $\mu\text{g}/\text{m}^3$ (approximately a 7-fold greater concentration versus the ambient level). The elemental composition, as measured by energy-dispersive X-ray fluorescence (ED-XRF) analysis, is presented in supplemental Table I.

Body Weight, Fat Content, Glucose Homeostasis, and Systemic Inflammation Changes

In wild-type C57BL/6 mice, there were no significant differences among the 4 groups in body weight (supplemental Table II) or glucose tolerance (supplemental Figure 1A) at baseline before exposure. Mice on an HFD gained much more body weight compared with mice on an ND. Mice on an ND exposed to PM_{2.5} showed significant elevations in glucose levels by an intraperitoneal glucose tolerance test (Figure 1A and 1B). HFD (regardless of FA or PM_{2.5} exposure) and PM_{2.5} exposure in ND-fed mice led to IR, evaluated by a homeostasis model assessment index (Figure 1C). Figure 1D shows inflammatory biomarkers in blood from C57BL/6 mice exposed to either PM_{2.5} or FA. TNF- α was increased in the mice exposed to PM_{2.5} compared with the FA in either ND- or HFD-fed mice. To examine whether PM_{2.5} exposure altered adiposity, a T1-weighted MRI was performed before and after PM_{2.5} exposure. There was no significant difference in abdominal body fat among the experimental groups before exposure (supplemental Figure 1B). However, PM_{2.5} exposure alone and HFD feeding significantly increased the total abdominal fat compared with FA-exposed mice fed an ND (Figure 2A and supplemental Figure 1C), although the combination of PM_{2.5} with an HFD did not further increase abdominal fat. With respect to fat distribution, both visceral and subcutaneous fat contents were increased with PM_{2.5} exposure in the ND- or HFD-fed group (Figure 2B). Consistent with these results, adipocyte size was increased in the PM_{2.5}-exposed mice fed an ND in both visceral fat (FA, 2137 \pm 45 μm^2 ; PM_{2.5}, 2698 \pm 80 μm^2 ; *P*<0.01) and subcutaneous fat (FA, 1039 \pm 27 μm^2 ; PM_{2.5}, 1355 \pm 30 μm^2 ; *P*<0.05). The increase in adipocyte size was extreme in the HFD groups, with no further changes due to PM_{2.5} exposure (supplemental Figure 1I). These data suggest that PM_{2.5} exposure alone, in the presence of ND, may potentiate adiposity and exert proinflammatory effects.

Because the p47 subunit of the NADPH oxidase is homologous in both phagocytic and nonphagocytic cells, and is critical for a functional NADPH oxidase,^{20,21} we next investigated the importance of the NADPH oxidase in mediating the altered metabolic profile and IR in response to PM_{2.5} and diet. Age-matched male p47^{phox-/-} mice were exposed to PM_{2.5} or FA using the same exposure protocol as the wild-type C57BL/6 mice. We found that there was no significant additional effect of PM_{2.5} on body weight in the different diet feeding groups (supplemental Table II). Glucose homeostasis in response to glucose loading from the PM_{2.5}-exposed mice was comparable to that in the FA-exposed p47^{phox-/-} mice (Figure 1E and 1F). Although an HFD led to a sharp increase in blood glucose levels, the increase was similar for PM_{2.5}- and FA-exposed mice. Homeostasis

model assessment IR indexes from the PM_{2.5}-exposed p47^{phox}^{-/-} mice were significantly attenuated and comparable to those of the FA-exposed mice fed an ND (Figure 1G). Plasma inflammatory biomarkers in the p47^{phox}^{-/-} mice were similar to those in the wild-type C57BL/6 mice. Notably, the absence of a functional NADPH oxidase abrogated the previously noted difference in TNF- α in wild-type C57BL/6 with PM_{2.5} exposure (Figure 1H). Neither visceral nor subcutaneous fat in the abdomen, measured by MRI, was significantly different between the ND-fed groups exposed to PM_{2.5} or FA, which differed from the results in wild-type C57BL/6 mice (Figure 2E and 2F and supplemental Figure IF). In p47^{phox}^{-/-} mice, adipocyte size in the PM_{2.5}-exposed mice fed an ND was similar to that in the FA-exposed mice on the same diet in either visceral fat (FA, 1221 \pm 50 μ m²; PM_{2.5}, 1184 \pm 38 μ m²; $P>0.05$) or subcutaneous fat (FA, 769 \pm 44 μ m²; PM_{2.5}, 910 \pm 29 μ m²; $P>0.05$) (supplemental Figure III). On the other hand, the adipocyte area of HFD-fed p47^{phox}^{-/-} mice was increased, albeit smaller than in the HFD-fed wild-type C57BL/6 mice, with considerable variation in size distribution.

Chemotaxis and Superoxide Generation

To investigate potential alteration in chemokine factors within visceral adipose tissue, we evaluated the chemotactic ability of conditioned media derived by culturing adipose samples from visceral and subcutaneous fat locations. The migratory capacity of monocytes was tested by quantifying the number of monocytes moving toward the chamber containing conditioned media. In wild-type C57BL/6 mice, more monocytes migrated toward conditioned media from visceral fat from mice exposed to PM_{2.5} fed an ND or an HFD when compared with those exposed to FA (Figure 2C). In contrast, there was no increase in cell migration with conditioned media from subcutaneous fat locations. The generation of an NADPH-derived O₂^{•-} anion by the adipose tissue was further assessed by a lucigenin-enhanced chemiluminescence technique. Interestingly, O₂^{•-} production was significantly increased in the epididymal fat (visceral fat) of the mice exposed to PM_{2.5} compared with the FA group, but not in the subcutaneous fat location (Figure 2D). However, in p47^{phox}^{-/-} mice, as shown in Figure 2G, differential chemotactic responses in response to PM_{2.5} exposure in the visceral fat of wild-type C57BL/6 mice were abolished in the p47^{phox}^{-/-} mice. Furthermore, there was no increase in O₂^{•-} generation in response to PM_{2.5} in the p47^{phox}^{-/-} mice exposed to PM_{2.5} (Figure 2H).

Inflammation in Visceral Adipose Tissue and Microcirculatory Dysfunction

Adipose tissue macrophages (ATMs), which are thought to represent key cellular mediators of adipose tissue inflammatory response and IR development, were examined in C57BL/6 mice. PM_{2.5} exposure resulted in an increase in F4/80⁺ macrophages in epididymal (visceral) adipose tissue (Figure 3A and 3B). We then investigated the expression of ATM-specific genes in the macrophage-rich stromal vascular fraction of visceral adipose tissue (Figure 3C). PM_{2.5} exposure led to a significant increase in the expression of proinflammatory genes (M₁, or “classically” activated) TNF- α , NO synthase 2, and interleukin (IL) 6, with no changes in integrin α X expression (Figure 2E). In contrast, IL-10, which is expressed at high levels in alternatively activated macrophages and adipocytes, was significantly downregulated by PM_{2.5} exposure. These results demonstrate that PM_{2.5} exposure downregulated genes associated with an anti-inflammatory M₂ phenotype while inducing a proinflammatory phenotype. In view of the inflammatory macrophage phenotype in adipose tissue, we hypothesized that accelerated recruitment into adipose tissue may represent an important mechanism, as demonstrated by a prior study.²² At the end of the exposure to PM_{2.5} or FA, intravital microscopy with labeled monocytes/macrophages was performed to evaluate the number of rolling and adherent cells, as an index of recruitment into tissue depots. The results show that PM_{2.5} exposure resulted in an increase in adherent

and rolling monocytes in the microcirculation when compared with the FA-exposed mice. There was an important synergistic effect of HFD in terms of exaggerating the PM_{2.5} effects (Figure 4A and 4B). Interestingly, there was no increase in ATM content in the p47^{phox}^{-/-} mice exposed to PM_{2.5} (Figure 3B), although gene expressions were similar to those in wild-type C57BL/6 mice (Figure 3D). Intravital microscopy showed that the increased adherence of monocytes in the cremaster microcirculation observed in the wild-type C57BL/6 mice that were exposed to PM_{2.5} was significantly decreased by the deletion of p47^{phox} (Figure 4C and 4D). Although the number of rolling monocytes in response to an HFD was increased in the p47^{phox}^{-/-}, the extent was attenuated when compared with HFD-fed mice and either PM_{2.5} or FA exposure, suggesting that mechanisms other than NADPH oxidase may be involved.

Vasomotor Responses

We then examined vascular responses to endothelium-dependent agonists, including insulin. Endothelial dysfunction and altered insulin sensitivity in blood vessels are characteristic of IR. As shown in Figure 5A through 5C and the Table, PM_{2.5}-exposed wild-type C57BL/6 mice exhibited an enhanced constriction response to phenylephrine and a decreased relaxation response to endothelium-dependent vasodilator acetylcholine. The relaxation response to insulin was also reduced by PM_{2.5} exposure. PM_{2.5}-exposed p47^{phox}^{-/-} mice fed an ND displayed a similar constriction response to phenylephrine and a relaxation response to acetylcholine when compared with the FA-exposed ND-fed mice (Figure 5D–5F and Table), although the response to insulin in p47^{phox}^{-/-} mice was comparable to that in the wild-type C57BL/6 mice. These data, collectively, indicate an important role for p47^{phox} and NADPH oxidase in mediating responses to PM_{2.5}.

PM_{2.5} Exposure Induces p47^{phox} Phosphorylation

On activation, a p47^{phox} cytosolic subunit phosphorylates, translocates to the membranes, and associates with the membrane-bound components. Then, the newly assembled enzyme complex actively catalyzes the production of O₂^{•-}. Therefore, we next wanted to know if exposure to PM_{2.5}, regardless of diet impact, may have effects on the p47^{phox} subunit. As shown in supplemental Figure IVA and IVB, PM_{2.5} exposure significantly increased the phosphorylation of the p47^{phox} subunit of NADPH oxidase in epididymal adipose tissue compared with the FA-exposed control, indicating the effect of PM_{2.5} exposure on NADPH oxidase.

Discussion

In this study, we evaluated the role of early-life inhalation exposure to concentrated airborne PM_{2.5} on systemic and adipose tissue inflammation and susceptibility to IR development in adulthood. There are several important findings in this study. First, early-life PM_{2.5} exposure, even in the absence of dietary indiscretion, is sufficient to induce metabolic dysfunction and inflammation (“metaflammation”). Second, PM_{2.5} exposure for only 10 weeks is sufficient to induce visceral adiposity and vascular dysfunction, consistent with IR. Finally, NADPH oxidases may play a critical role in PM_{2.5}-induced development of T2DM/IR.

Risk factors, such as inappropriate diet and inactivity, play fundamental roles in the propensity for T2DM.^{14,23,24} Underlying genetics also provide an explanation, particularly among certain defined populations.^{25,26} On the other hand, the link between exposure to environmental factors in the air/water and propensity for cardiometabolic disorders has only recently gained attention.^{7,8,27,28} This issue is of special importance given the extraordinary confluence of changing levels of airborne and water pollutants and shifts in diet/exercise in

urban populations. There appear to be additional interactions between the effects of inhaled PM_{2.5} and metabolic disorders, such as T2DM, as shown by the study of O'Neill et al,²⁹ in which diabetic individuals seemed more vulnerable to PM-associated impairment in endothelial function.

Our study provides important insights into potential mechanisms by which PM_{2.5} may induce obesity/IR. The mechanisms seem broadly similar to other dietary mediators, such as an HFD, in which an important role for innate immune mechanisms has been highlighted.³⁰ Interestingly, our studies clearly show a phenotypic shift in response to PM_{2.5} exposure, which is typified by enhanced ATM infiltration, proinflammatory gene expression, and an altered chemokine profile that may play important roles in stimulating infiltration of these tissues by monocytes. Although this study was not designed to address the relative importance of phagocyte-versus non-phagocyte-derived NADPH oxidases, several findings suggest that phagocyte derived NADPH oxidases may be playing an important role.

Phagocytic NADPH oxidases are well-known to produce much O₂^{•-} at levels that are several orders of magnitude higher than those of nonphagocytic systems.^{31,32} In addition, there were further indirect observations in this study that support the concept that activation of monocytes/macrophages in visceral adipose tissue may contribute to the excess levels noted in the study. These findings include the following: (1) normalization of ATM content in the p47^{phox-/-} and O₂^{•-} generation, (2) reduction in the recruitment of monocytes/macrophages into the microcirculation in labeled monocyte experiments, (3) reduction in chemotactic responses to monocytes with deletion of p47^{phox}, and (4) attenuation of classically activated *M1* gene expression in visceral adipose tissue of TNF- α and IL-6 in p47^{phox-/-} mice. However, there were additional findings that support the involvement of nonphagocytic NADPH oxidase pathways or distinct pathways unrelated to oxidant systems. For instance, there was continued upregulation of TNF- α and IL-6 (albeit attenuated) gene expression in the p47^{phox-/-}, suggesting that reactive oxygen species and/or NADPH oxidase independent pathways may still be involved, especially in response to HFD and potentially in response to PM_{2.5}. Interestingly, NO synthase 2 gene expression was markedly increased in response to PM_{2.5} in the p47^{phox-/-}. The reasons for this are unclear at this point. In addition, adipocyte remodeling seen with HFD feeding continued in the p47^{phox-/-} mice, suggesting reactive oxygen species- and/or NADPH oxidase-independent pathways in the pathogenesis of IR. Thus, our data suggest an important interaction of PM_{2.5} in activating NADPH oxidases in both phagocytic and potentially nonphagocytic cells and are consistent with an important role for the p47^{phox} cytosolic subunit. Further studies in tissue-specific and conditional knockout models will be needed to distinguish relative contributions of the phagocyte NADPH oxidase from nonphagocyte systems. Taken together, our findings are consistent with the studies by Furukawa et al¹⁸ and Schroder et al³³; they also support an important role for early priming by PM_{2.5} exposure. The increase in ATM was paralleled by an increase in adipose tissue size and visceral adipose tissue content in the PM_{2.5}-exposed group. Adipose tissue synthesizes and secretes proinflammatory substances, which are upregulated in obesity and play important roles in mediating obesity-linked IR.^{12,14} Accumulating evidence indicates that visceral fat is the most predictive of postprandial glucose levels, indexes of IR, and cardiovascular disease.³⁴⁻³⁷ Thus, adipose tissue and adipose tissue- derived signals may be key mediators and may play a pivotal role in PM_{2.5}-induced adiposity and IR.

Many studies^{38,39} have shown that exposure to PM_{2.5} is associated with a systemic proinflammatory response (eg, increased TNF- α and IL-6 levels) in humans and animals. Brook et al⁴⁰ previously showed that long-term exposure to nitrogen dioxide (NO₂) (a marker of traffic-related PM exposure) was associated with the prevalence of diabetes among women. Across the interquartile range (approximately 4 parts per billion NO₂) in 2

cities in Canada, there was an approximate 17% increase in the odds of diabetes. This association was likely because of NO₂-associated vehicle-related PM exposure, not a biological effect of NO₂ itself. In a recent set of key mechanistic experiments, in which 12 weeks of HFD feeding of adult male C57BL/6 mice that were subsequently exposed to PM_{2.5} demonstrated overt glucose intolerance and diabetes.⁸ MRI measurements in these mice revealed an increase in visceral fat with abnormalities in phosphatidylinositol 3-kinase/Akt/endothelial NO synthase signaling and adipocytokines. In contrast, findings from an earlier study⁸ demonstrated an important synergistic effect of diet with PM_{2.5} exposure, with the current study revealing an effect of PM_{2.5} alone in the absence of HFD. In this study, exposure began at the age of 3 weeks and was continued for 10 weeks; the previous study focused on the adult mice (aged > 18 weeks) that were already considered obese (HFD feeding for 10 weeks before PM_{2.5} exposure). The finding that PM_{2.5} exposure alone, in the absence of dietary influences, over a shorter time than in a previously reported experiment was sufficient to induce changes characteristic of IR and obesity is particularly noteworthy. This suggests that early-life exposure may play an important priming influence in the eventual development of obesity.

The p47^{phox} cytosolic subunit plays a pivotal role in NADPH oxidase activation in inflammatory cells by providing physical binding domains to cytochrome b₅₅₈ and p67^{phox}, and genetic mutations of p47^{phox} lead to loss of O₂^{•-} production during cell activation.^{41,42} Several studies^{43–45} in animals and humans have shown an important role for this subunit in inflammatory disorders, such as IR, and complications of diabetes. p47^{phox} has been regulated via phosphorylation by IL-1 receptor–associated kinase 4 in response to Toll-like receptor (TLR) 4 ligation.^{46,47} Diet-induced obesity and IR are well-known to elevate multiple mediators of TLR4 activation, including via endogenous fatty acids.^{48,49} Chen and colleagues⁵⁰ showed that diet-induced obesity activates NADPH oxidase and mediates the expression of TLR in the vascular tissues, whereas deficiency of the NADPH oxidase subunit p47^{phox} attenuates inflammatory response and neointimal formation in mice with diet-induced obesity. In line with these findings, we recently demonstrated that oxidized phospholipids generated in response to PM_{2.5} exposure in the lung and perhaps in the systemic circulation lead to activation of NADPH oxidase via TLR4/IL-1 receptor–associated kinase-dependent pathways (T.K., A.M., Z.Y., Z.S., J.A.D, X.X., N.K., R.D.B., K.M.R., N.P.P., S.P., L.C.C., S.M-B, Q.S., H.M., S.R., unpublished data, 2010). From our current data, the increase in inflammation and oxidative stress seemed to be associated with p47^{phox} phosphorylation in response to PM_{2.5} exposure.

In previous studies,^{51,52} long-term effects of concentrated ambient particle exposure on heart rates and heart rate variability in C57 and apolipoprotein E–knockout mice were observed. Although the experimental procedure of placing the mice in the chamber itself increased physical activity by 0.9 counts, there were no relationships with concentrated ambient particle concentration. Moreover, heart rate change during the exposure period was marginally associated with the concentrated ambient particle level. A potential limitation of the present study is that the exposure procedure might alter activity in home cages, although we deliberately exposed the mice during the daytime, while the mice were sleeping as they normally do, as part of their sleep/wake cycle. The mice of all groups were under the same experimental conditions (ie, light-dark cycle) except that a high-efficiency particulate air filter was positioned in the inlet valve to the exposure system to remove all of the PM_{2.5} from that air stream. Furthermore, the mice are not able to move around ad libitum during the exposure because the space/cell for each mouse is limited or restricted (approximately 6×7 cm).

In summary, our data suggest an important impact of early-life exposure to PM_{2.5} exposure on T2DM/IR development. The investigation of environmental factors, such as PM_{2.5} air

pollution, on the increased susceptibility to T2DM/IR development is significant; these findings suggest an important public health impact on the health of both children and adults. Further understanding of the factors regulating the activation and accumulation of macrophages in the adipose tissue may provide novel prevention and therapeutic strategies for better control and treatment of obesity, T2DM, and IR.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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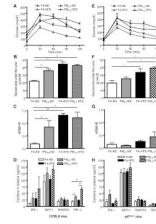


Figure 1.

Glucose homeostasis and systemic inflammation in wild-type C57BL/6 mice and p47^{phox}^{-/-} mice by ambient PM_{2.5} exposure compared with FA-exposed mice fed an ND (n = 16) or an HFD (n = 16). A and E, Effect of PM_{2.5} exposure on glucose tolerance by intra-peritoneal glucose tolerance test (IPGTT) in C57BL/6 mice and in p47^{phox}^{-/-} mice, respectively. B and F, The glucose area under the curve calculated from the glucose tolerance test from parts A and E, respectively. C and G, The homeostasis model assessment IR index in C57BL/6 mice and in p47^{phox}^{-/-} mice, respectively. D and H, Plasma cytokine measurement by ELISA in C57BL/6 mice and in p47^{phox}^{-/-} mice, respectively. n = 8 in each group. **P*<0.05 and ***P*<0.001. IFN indicates interferon; MCP-1, monocyte chemoattractant protein-1; RANTES, regulated on activation, normal T cell expressed and secreted (or chemokine C-C motif ligand 5).

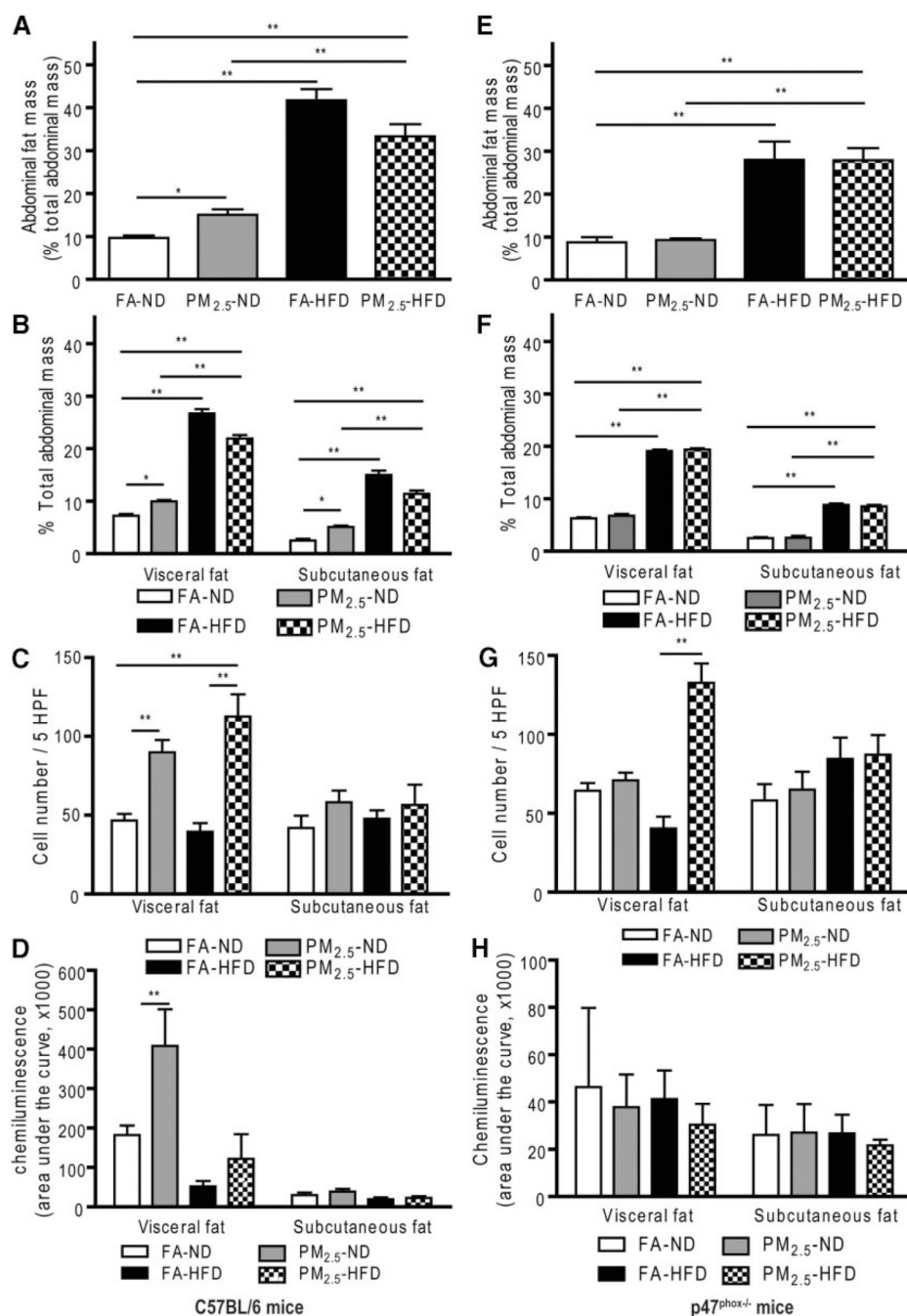


Figure 2. Effect of PM_{2.5} exposure on abdominal fat and distribution, chemotactic migration, and superoxide production in wild-type C57BL/6 mice and p47^{phox}^{-/-} mice. A and E, Abdominal fat measured by MRI in C57BL/6 mice and p47^{phox}^{-/-} mice, respectively. B and F, Visceral and subcutaneous fat (fat distribution) in the abdomen, measured by MRI, in C57BL/6 mice and p47^{phox}^{-/-} mice, respectively. C and G, Chemotactic migration of monocytes in C57BL/6 mice and p47^{phox}^{-/-} mice, respectively. D and H, Superoxide anion measurement by chemiluminescence in visceral and subcutaneous fat tissues in wild-type C57BL/6 and p47^{phox}^{-/-} mice, respectively. **P*<0.05 and ***P*<0.001. 5HPF indicates 5 high-power fields.

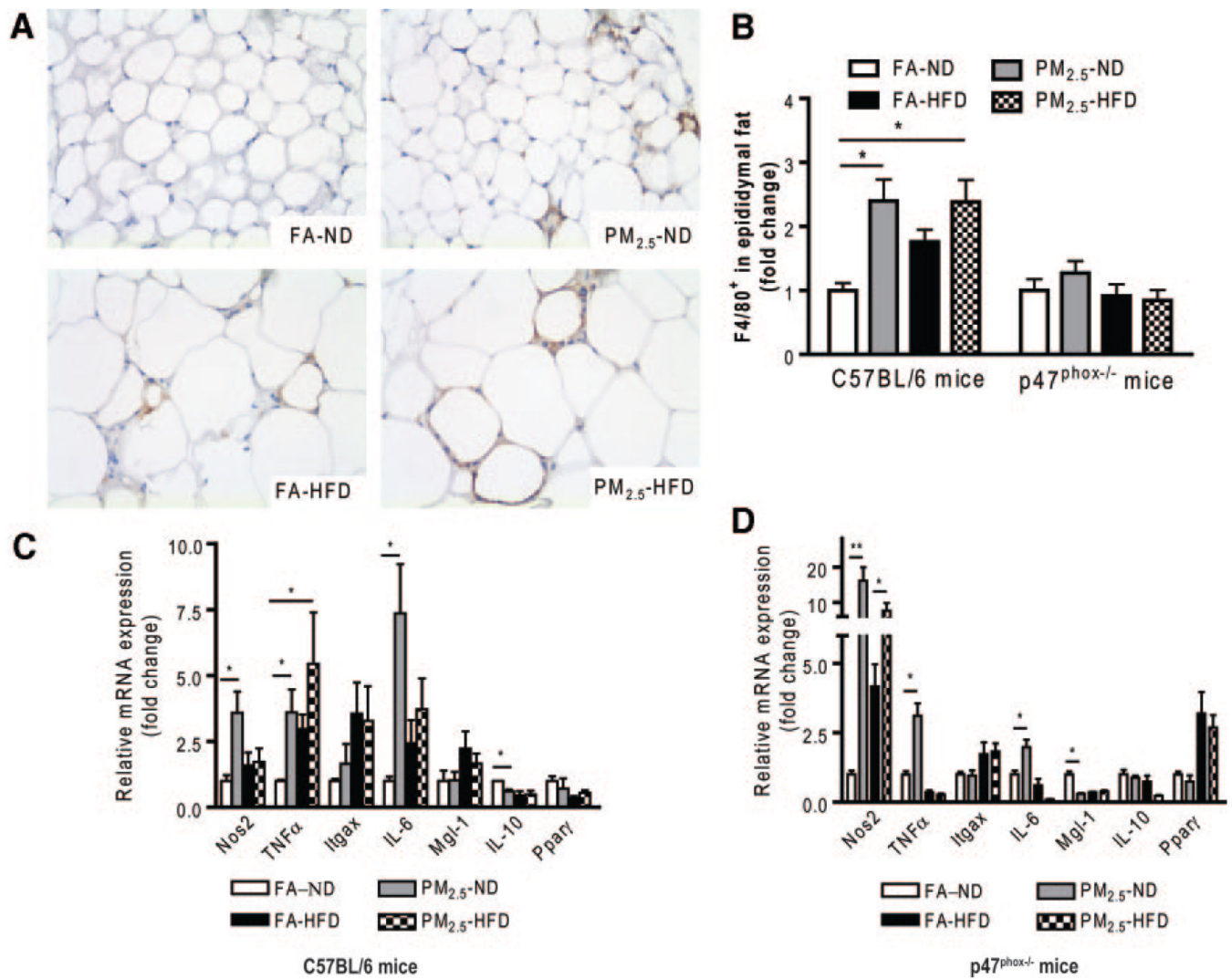


Figure 3.

Effect of PM_{2.5} exposure on macrophage inflammation and gene expression in visceral adipose tissue in wild-type C57BL/6 and p47^{phox-/-} mice. A, Representative images of immunohistochemical staining for macrophages (F4/80⁺) in visceral fat tissue. B, Statistical analysis of F4/80⁺ macrophage infiltration in visceral fat tissue in C57BL/6 and p47^{phox-/-} mice. C and D, Real-time polymerase chain reaction analysis for macrophage M₁/M₂ phenotypic changes in C57BL/6 and p47^{phox-/-} mice, respectively. Itgax indicates integrin α X (or CD11c); Mgl-1, macrophage galactose *N*-acetyl-galactosamine specific lectin-1; Nos2, NO synthase-2; Pparg, peroxisome proliferator-activated receptor γ . **P*<0.05 and ***P*<0.001.

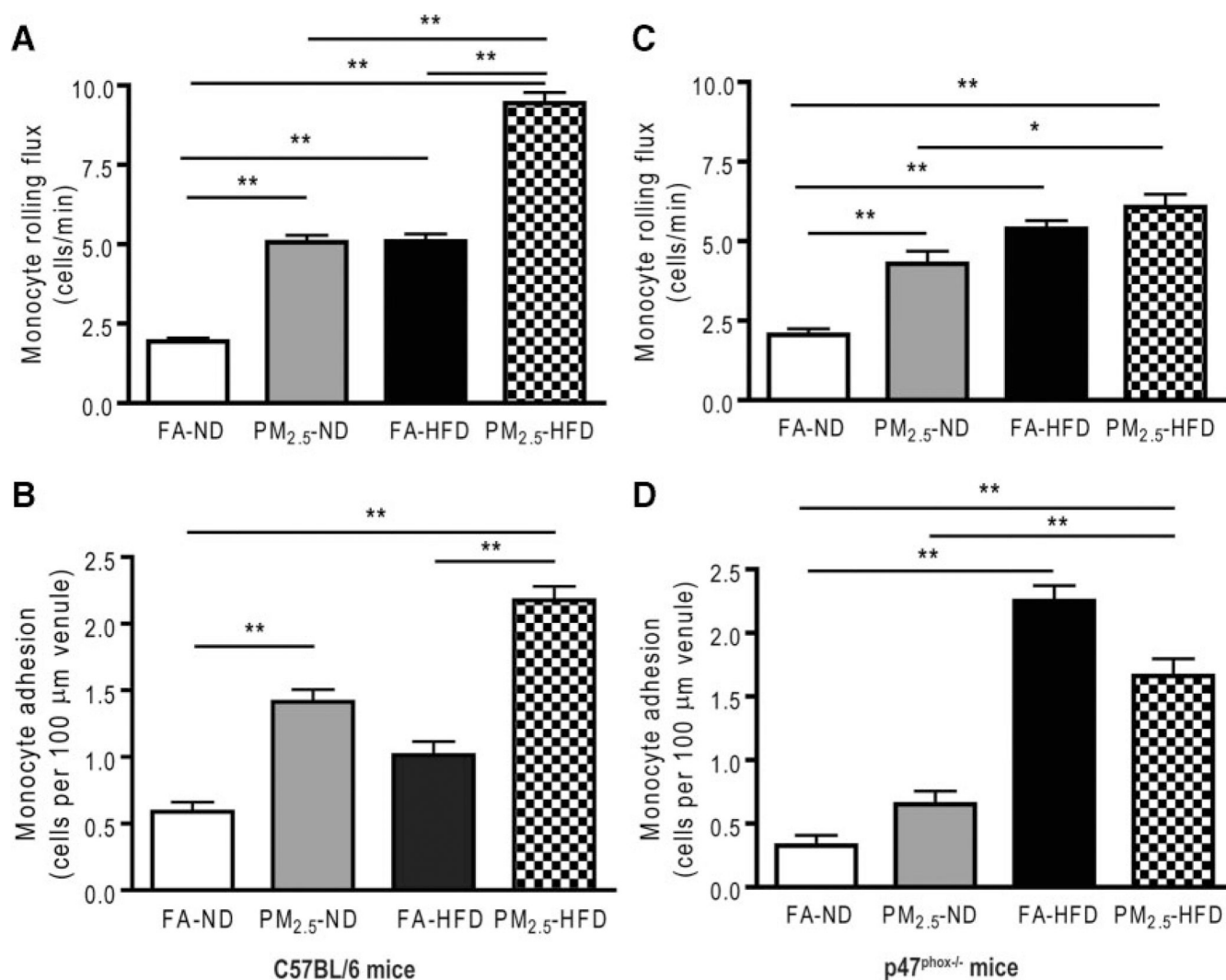


Figure 4. Effect of PM_{2.5} exposure on microvascular dysfunction in wild-type C57BL/6 and p47^{phox-/-} mice. A and C, Monocyte rolling flux in cremasteric microcirculation via intravital microscopy in C57BL/6 and p47^{phox-/-} mice, respectively. B and D, Monocyte adhesion in cremasteric microcirculation via intravital microscopy in C57BL/6 and p47^{phox-/-} mice, respectively. **P* < 0.05 and ***P* < 0.001.

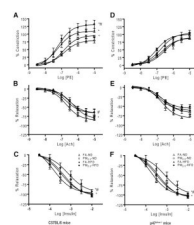


Figure 5. Effect of PM_{2.5} exposure on macrovascular dysfunction in wild-type C57BL/6 and p47^{phox}^{-/-} mice. A through F, Vasomotor tone change in aortic rings via myograph in response to phenylephrine (A and D), acetylcholine (B and E), or insulin (C and F) in C57BL/6 and p47^{phox}^{-/-} mice, respectively. **P*<0.05 vs FA-ND, and #*P*<0.05 vs FA-HFD.

Table 1

Vascular Responses to Phenylephrine, Acetylcholine, and Insulin From Aortic Rings in Wild-Type C57BL/6 Mice and in p47^{phox}^{-/-} Mice *

Agent	FA-ND	PM _{2.5} -ND	FA-HFD	PM _{2.5} -HFD
C57BL/6 Mice				
Phenylephrine				
Peak constriction, %	80.1±4.6	96.0±2.8 [†]	110.1±5.5 [†]	131.2±6.4 ^{†‡}
EC ₅₀ , mol/L	1.4×10 ⁻⁷ ±0.1×10 ⁻⁷	1.3×10 ⁻⁷ ±0.1×10 ⁻⁷	0.6×10 ⁻⁷ ±0.2×10 ⁻⁷	0.6×10 ⁻⁷ ±0.1×10 ⁻⁷
Acetylcholine				
Peak relaxation, %	-85.7±3.5	-82.6±4.7	-66.5±4.6	-66.5±3.8
ED ₅₀ , mol/L	1.9×10 ⁻⁷ ±0.1×10 ⁻⁷	4.9×10 ⁻⁷ ±0.1×10 ⁻⁷ [†]	1.8×10 ⁻⁷ ±0.2×10 ⁻⁷	2.8×10 ⁻⁷ ±0.1×10 ⁻⁷ [‡]
Insulin				
Peak relaxation, %	-98±4.6	-105.7±6.0	-102.1±7.4	-96.3±10.5
ED ₅₀ , mol/L	1.5×10 ⁻⁴ ±0.1×10 ⁻⁴	4.5×10 ⁻⁴ ±0.1×10 ⁻⁴ [†]	3.4×10 ⁻⁴ ±0.2×10 ⁻⁴	7.4×10 ⁻⁴ ±0.2×10 ⁻⁴ ^{†‡}
p47 ^{phox} ^{-/-} Mice				
Phenylephrine				
Peak constriction, %	88.8±4.5	91.4±5.2	104.9±5.3	100.6±1.9
EC ₅₀ , mol/L	2.1×10 ⁻⁷ ±0.1×10 ⁻⁷	3.0×10 ⁻⁷ ±0.1×10 ⁻⁷	1.8×10 ⁻⁷ ±0.1×10 ⁻⁷	0.9×10 ⁻⁷ ±0.1×10 ⁻⁷
Acetylcholine				
Peak relaxation, %	-81.1±3.0	-67.0±3.2 [†]	-61.0±2.7	-55.5±2.9
ED ₅₀ , mol/L	1.9×10 ⁻⁷ ±0.1×10 ⁻⁷	2.6×10 ⁻⁷ ±0.1×10 ⁻⁷	1.5×10 ⁻⁷ ±0.1×10 ⁻⁷	1.5×10 ⁻⁷ ±0.1×10 ⁻⁷
Insulin				
Peak relaxation, %	-102.6±2.6	-108.1±3.4	-105.0±5.6	-100.1±10.9
ED ₅₀ , mol/L	1.6×10 ⁻⁴ ±0.1×10 ⁻⁴	3.9×10 ⁻⁴ ±0.1×10 ⁻⁴ [†]	3.1×10 ⁻⁴ ±0.1×10 ⁻⁴	7.9×10 ⁻⁴ ±0.2×10 ⁻⁴ ^{†‡}

EC₅₀ indicates the half-maximal dose for constriction; ED₅₀, half-maximal dose for dilation.

* Data are given as mean±SE.

[†] P<0.05 vs FA-ND.

[‡] P<0.05 vs FA-HFD.