



Article Concordance between In Vitro and In Vivo Relative Toxic Potencies of Diesel Exhaust Particles from Different Biodiesel Blends

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Abstract: Diesel exhaust particles (DEPs) contribute to air pollution exposure-related adverse health impacts. Here, we examined in vitro, and in vivo toxicities of DEPs from a Caterpillar C11 heavyduty diesel engine emissions using ultra-low-sulfur diesel (ULSD) and biodiesel blends (20% v/v) of canola (B20C), soy (B20S), or tallow-waste fry oil (B20T) in ULSD. The in vitro effects of DEPs (DEP_{ULSD}, DEP_{B20C}, DEP_{B20S}, and DEP_{B20T}) in exposed mouse monocyte/macrophage cells (J774A.1) were examined by analyzing the cellular cytotoxicity endpoints (CTB, LDH, and ATP) and secreted proteins. The in vivo effects were assessed in BALB/c mice (n = 6/group) exposed to DEPs (250 µg), carbon black (CB), or saline via intratracheal instillation 24 h post-exposure. Bronchoalveolar lavage fluid (BALF) cell counts, cytokines, lung/heart mRNA, and plasma markers were examined. In vitro cytotoxic potencies (e.g., ATP) and secreted TNF- α were positively correlated (p < 0.05) with in vivo inflammatory potency (BALF cytokines, lung/heart mRNA, and plasma markers). Overall, DEPULSD and DEP_{B20C} appeared to be more potent compared to DEP_{B20S} and DEP_{B20T} . These findings suggested that biodiesel blend-derived DEP potencies can be influenced by biodiesel sources, and inflammatory process- was one of the potential underlying toxicity mechanisms. These observations were consistent across in vitro and in vivo exposures, and this work adds value to the health risk analysis of cleaner fuel alternatives.

Keywords: biodiesel; diesel exhaust particles; cytotoxicity; intratracheal instillation; inflammation

1. Introduction

There are many reports on the links between exposure to particulate air pollutants and adverse cardiac [1–8], pulmonary [9,10], reproductive/pregnancy [11–13], and neurological [6,14] outcomes, as well as cancer [15–17]. Diesel exhaust particles (DEPs) from traffic emissions are a source of urban air particulate pollution [18] and are considered major contributors to air pollution exposure-related adverse population health effects. There are reports supporting this notion with suggestions of plausible mechanistic pathways for the observed adverse health effects of DEPs [19–24].

Biodiesels derived from renewable feedstock may be used as a partial replacement for petroleum-based diesel in automotive applications, which is supported by governmental regulations such as the Clean Fuel Regulations (https://laws-lois.justice.gc.ca/ eng/regulations/SOR-2022-140/index.html (accessed on 6 April 2024)), and provincial



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Copyright: © 2024 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). renewable fuel and low-carbon fuel requirements regulations. A consequence of blending biodiesels with conventional diesel is altered exhaust chemistry. Earlier work on biodiesel [25] showed that biodiesel blends of 20% resulted in reductions of 15% or higher in terms of emissions of particulate matter, carbon monoxide, total hydrocarbons, vaporphase hydrocarbons from C1 to C12, aldehydes, ketones, selected semi-volatile polycyclic aromatic hydrocarbons (PAHs), and nitroPAHs, with no net effect on the oxides of nitrogen (NOx). There are reports of increased aldehydes, including formaldehyde, acetaldehyde, and acrolein, in biodiesel emissions as opposed to petroleum-based fuel emissions [26–28]. The emission of particulate matter (PM) from biodiesel blends on a mass basis was reported to be influenced by the engine starting conditions; furthermore, biodiesel emissions also exhibited altered volatile and semi-volatile PM number and a biodiesel source-dependent increase in NOx emissions [29,30]. Meanwhile, the use of diesel particle filters (DPFs) was shown to decrease PM and CO levels, and hydrocarbons were decreased when using a diesel oxidation catalyst (DOC) + DPF [31,32].

Studies on the toxicological impacts of emissions derived from the combustion of conventional diesel and biodiesel-blended fuels are emerging. Jalava et al. 2010 [33] showed unaltered cytotoxicity and reduced inflammatory potential of DEPs derived from a rapeseed methyl ester biodiesel-blended fuel in comparison to emission particles from conventional diesel. Similarly, Hawley et al., 2014 [34] reported on similar biological responses for a plant-based biodiesel blend (B99) and ultra-low sulfur diesel (ULSD) in human bronchial epithelial cells. Additionally, a report by Rouleau et al., 2013 [35] suggested that air quality and health benefits/costs associated with the use of biodiesel blends and ULSD may be expected to be similar based on modeling with limited data. However, increased toxicity of particulate emissions derived from biodiesel-blended fuels has also been reported. In vitro studies have shown increased cytotoxicity, secreted inflammatory cytokines [36,37], and the production of reactive oxygen species [33,38] after exposure to DEPs derived from biodiesel- blended fuels. Similarly, the exposure of mice to DEPs derived from soy or corn biodiesel-blended fuels via oropharyngeal aspiration resulted in significantly increased inflammatory and oxidative changes in lung tissue [39] or increased sperm DNA fragmentation and the upregulation of inflammatory cytokines in the serum and testes [40]. In an in vitro–in vivo comparative study, Fukagawa et al. (2013) [41] showed increased inflammatory and oxidative stress responses to DEPs derived from 20% v/vsoy biodiesel-ULSD blend when compared to particles derived from ULSD both in vitro in BEAS-2B and THP-1 cell lines and in vivo in C57BL/6 mice. Douki et al. 2018 [42] reported on limited transcriptomic changes in the lungs of rats after repeated exposure to rapeseed oil-derived biodiesel blend and ULSD. Soy-derived biodiesel emissions at a higher PM concentration were reported [43] to be associated with relatively less evidence of pulmonary effects compared to diesel emissions (after 4 weeks of exposure, in vivo). Madden 2016 [44] reviewed the toxicity studies on DEPs derived from the combustion of biodiesel blends and noted the inconsistencies in toxicology findings. Inconsistencies in toxicity findings can arise due to the quality of biodiesel blends and source differences, heterogeneity in exposure conditions/characterization in these studies, the use of different cell types, and the lack of in vitro and in vivo studies carried out with the same biodiesel blends in a systematic manner.

Therefore, in this work, we examined the relative in vitro and in vivo toxicity characteristics of diesel exhaust particles derived from the combustion of commercial ultra-low sulfur diesel (ULSD) or 20% (v/v) blends (B20) of biodiesels based on canola oil, soy oil, and tallow–waste fry oil in ULSD used in a heavy-duty diesel engine (on-road technology). These DEPs were examined for in vitro toxicities in mouse monocyte/macrophage cells (J774.A1) and for in vivo toxicities via the intra-tracheal instillation of these DEPs in BALB/c mice, followed by analysis of various biological endpoints to identify any consistency between in vitro and in vivo toxicity findings.

2. Materials and Methods

2.1. Engine

Emissions were generated using a heavy-duty diesel engine, Caterpillar C11, equipped with the manufacturer's original diesel oxidation catalyst (DOC) and conforming to 2004 emission standards. The engine specifications and certified emission rates are provided in Table 1.

Table 1. The table shows the engine specifications and certified emission rates.

Engine Manufacturer	Caterpillar			
Model	C11			
Year	2004			
Serial number	KCA018109			
Engine family	4CPXH0680EBK			
Air handling system	Series Turbo-Charged			
Control	Electronic ACERT			
Bore (mm)	130			
Stroke (mm)	140			
Cycles	4			
Number of cylinders	6			
Displacement (liters)	11.1			
Curb idle speed (rpm)	700			
Rated test speed (rpm)	2100			
Maximum torque (lb-ft)	p-ft) 1150 @ 1200 rpm			
Maximum power (bhp)	ım power (bhp) 305 @ 2100 rpm			
Compression ratio	tio 17.1			
NOx (g/bhp-hr.) *	NOx(g/bhp-hr.) * 2.3			
CO (g/bhp-hr.) *	1.6			
PM (g/bhp-hr.) *	0.09			

* Certified emissions with original DOC, when operated at 400HP at 1800 rpm, 1450 lb.-ft at 1200 rpm; data from https://www.epa.gov/compliance-and-fuel-economy-data/annual-certification-data-vehicles-engines-and-equipment (accessed on 6 April 2024).

Generally, new heavy-duty on-road engines, along with advanced combustion, fueling, and thermal management strategies, are equipped with DPFs and SCR (selective catalyst reduction) systems to meet more stringent emission standards. Future emission standards are based on varying test conditions and are approaching 0.02 g/bhp-hr for NOx and 0.005 g/bhp-hr for PM.

2.2. Fuels

The base test fuel was an ultra-low sulfur diesel (ULSD) procured from a commercial supplier. Unblended, 100% canola methyl ester biodiesel (B100) was procured from Milligan Biotech (Foam Lake, SK, Canada). Soy methyl ester (unblended, B100) was procured from Rothsay Biodiesel (Guelph, ON, Canada). The B100 tallow-waste fry oil was manufactured from a mixed feedstock containing 75% beef tallow and 25% waste fry oil and supplied by Biox Corporation (Oakville, ON, Canada). The biodiesels were blended with ULSD to prepare B20 blends containing 20% v/v biodiesel methyl ester and 80% v/v ULSD. The B20 blends of canola methyl ester, soy methyl ester, and tallow-waste fry oil methyl ester are referred to as B20C, B20S and B20T, respectively, in this report. All B100 biodiesels and B20 blends were analyzed by the Alberta Research Council (Edmonton, AB, Canada), operating in compliance with the ISO/IEC 17025 (https://www.iso.org/ISO-IEC-17025-testing-andcalibration-laboratories.html, accessed on 6 April 2024) requirements. Information on the analysis methods used for the characterization of B100 biodiesels and diesel engine emissions related to the fuel types used in this work are provided in Supplementary Tables S1 and S2, and the results of the analysis for B20 biodiesel blends used in the study are provided in Table 2.

Fuel Identification	Method	ULSD	Canola B20	Soy B20	Animal Tallow B20
Biodiesel blend Volume % *	ASTM D7371	n/a	19.6	19.3	20.2
Density, kg/m ³ @ 15 °C	ASTM D4052	833.0	843.1	843.3	841.6
Cetane No. D613	ASTM D613	49.6	52.8	55.6	53.8
Carbon, %m	ASTM D5291	86.20	84.32	84.17	84.10
Hydrogen, %m	ASTM D5291	12.79	13.02	13.38	13.66
Sulphur, mg/kg	ASTM D5453	4.6	3.8	4.6	7.6

Table 2. The table shows the analyses of the blended test fuels (B20).

* Determination of biodiesel content in diesel fuel using mild infrared spectroscopy method by ARC.

2.3. Engine Operation for Generation of Test Particles

The engine was operated on a dynamometer. The engine was pre-conditioned by operating at the rated speed and maximum torque for a period of 20 min. During engine operation, the total volume of raw exhaust was transferred from the engine's exhaust manifold to a constant volume sampling system. The exhaust was diluted with high-efficiency particulate air (HEPA)-filtered ambient air within the dilution tunnel. A continuous flow of diluted exhaust was collected through in-line sampling probes installed in the dilution tunnel and directed to the particulate matter sampling system and gas analyzers. The design of the sampling and analytical systems, as well as engine preparation and operation followed protocols described under the U.S. Code of Federal Regulations (CFR), Title 40, Part 86. For the generation of test particles, the engine was operated in the base OEM configuration (with DOC), at 25% or 50% loads in a steady-state 1200 rpm using ULSD, B20C, B20S, or B20T. The particulate matter samples were collected using Zefluor[®] 8 " × 10" rectangular polytetrafluoroethylene (PTFE) filters (Pall Life Sciences, Port Washington, NY, USA) for toxicity testing. The engine operation conditions and emission rates are given in Table 3.

Table 3. The table shows the engine operation conditions and emission rates.

Fuel	Engine Load	CO (g/bhp-hr.)	CO2 (g/bhp-hr.)	NOX (g/bhp-hr.)	THC (g/bhp-hr.)	PM (g/bhp-hr.)	FC (g/bhp-hr.)	Torque (lbft)	Power (bhp)
	25%	0.07 (0.01)	559.8 (5.10)	1.73 (0.03)	0.023 (0.00)	0.107 (0.01)	177.3 (1.61)	561 (4.58)	129 (1.03)
0130 -	50%	0.04 (0.01)	492.6 (2.22)	1.63 (0.02)	0.004 (0.00)	0.059 (0.00)	156.0 (0.71)	844 (3.94)	194 (0.91)
B20 capala	25%	0.08 (0.02)	553.7 (0.96)	1.65 (0.02)	0.036 (0.00)	0.115 (0.00)	179.4 (0.31)	529 (1.29)	121 (0.31)
D20 Carloia -	50%	0.05 (0.01)	497.9 (6.21)	1.72 (0.02)	0.000 (0.00)	0.059 (0.00)	161.3 (2.02)	810 (8.63)	186 (1.98)
B20 Sov	25%	0.08 (0.02)	555.9 (1.23)	1.57 (0.01)	0.029 (0.00)	0.112 (0.00)	179.7 (0.40)	545 (0.60)	125 (0.13)
- D20 50y	50%	0.05 (0.01)	499.7 (2.07)	1.62 (0.01)	0.001 (0.00)	0.057 (0.00)	161.5 (0.67)	802 (2.98)	184 (0.64)
P20 Tallary	25%	0.06 (0.01)	566.2 (1.45)	1.78 (0.03)	0.028 (0.00)	0.115 (0.00)	183.0 (0.46)	534 (1.89)	122 (0.48)
B20 Tallow -	50%	0.04 (0.01)	504.0 (1.06)	1.71 (0.00)	0.007 (0.00)	0.058 (0.00)	162.9 (0.35)	799 (0.59)	183 (0.20)

Values in parentheses are standard deviations from n = 3 runs.

2.4. Emission Characterization

Emissions of carbon monoxide (CO), carbon dioxide (CO₂), oxides of nitrogen (NOx), and total hydrocarbons (THCs) were continuously monitored in the emission stream. Particles were collected using 70 mm EmFab[®] filters (Pall Life Sciences, Port Washington, NY, USA) for gravimetry. Particulate filters were handled according to the procedure described in 40 CFR 86.1339-90 Particulate filter handling and weighing. Fuel consumption was calculated using an industry-standard carbon balance equation. The details of the analytical methods and instrumentation used in the characterization of emissions are provided in Supplementary Table S2.

2.5. Extraction of Particles

Pre-weighed sections of Zefluor[®] PTFE filters (VWR International, Mississauga, ON, Canada) containing diesel exhaust particles were placed in a 50 mL falcon tube (VWR, Mississauga, ON, Canada) and wetted with 2 mL of 100% ethanol (Sigma Chemical Company, MO, USA) for 60 s. The filters were then immersed in sterile, deionized water and sonicated in an ice-cold water bath for 30 min. Following sonication, the filters were removed, dried, and conditioned to calculate mass recovery. The particle extract was transferred into a pre-weighed, pre-siliconized lyophilization flask and lyophilized using Freezone lyophilizer (LabConco, Kansas City, MO, USA) according to the manufacturer's instructions. The lyophilized material was then suspended in a small volume of deionized water and aliquoted in pre-siliconized and pre-weighed amber glass tubes and lyophilized again. The samples were then frozen at -80 °C. The emission particles derived from the combustion of ULSD, B20C, B20S, and B20T are referred to below as DEP_{ULSD}, DEP_{B20C}, DEP_{B20S} and DEP_{B20T}, respectively.

2.6. Preparation of Particle Suspensions

Lyophilized diesel exhaust particles were suspended in a particle preparation buffer (25 μ g/mL of Tween-80 in 0.19% saline solution) at a concentration of 10 mg mL⁻¹, as described previously [45]. The particle suspensions were vortexed for 30 s and then sonicated for 20 min in ice-cold water in a water bath sonicator. The particles were then homogenized with 25 strokes of a homogenizer. In order to obtain a sufficient mass of particles to conduct both in vitro and in vivo toxicity testing experiments, the particles derived from engine runs at 25% and 50% loads were combined at a 2:1 mass ratio. This was rationalized based on the findings shown in Table 3. Even though engine loads of 25% vs. 50% were associated with different PM masses, at the same engine load across all fuels, the PM masses were similar and enabled the pooling of particles obtained from the different loads for the same fuel type, as noted above. The particles were aliquoted in O-ring-capped micro-centrifuge tubes and heated in a water bath at 56 °C for 30 min for sterilization. After cooling to room temperature, the samples were stored at -40 °C until testing.

2.7. In Vitro Exposure

2.7.1. Cell Culture

The J774A.1 cells were propagated in Dulbecco's Modified Eagle's Medium (DMEM; Fisher Scientific) containing phenol red and 4.5 g/L glucose and supplemented with FBS (10% v/v, non-heat inactivated; Fisher Scientific) and Pen Strep (100 U/mL penicillin G, 100 µg/mL streptomycin; Invitrogen, Burlington, ON, Canada) (referred below as complete DMEM) in 75 cm² tissue culture flasks (Corning, NY, USA). Unless stated otherwise, all of the cell incubations were at 5% CO₂ and 95% relative humidity. In preparation for cytotoxicity bioassays, the cells were recovered by scrapping the monolayers in complete DMEM devoid of phenol red. Cell suspensions in complete DMEM of 4 × 10⁵ cells/mL were seeded in 96-well black-walled clear-bottom cell culture plates (BD Biosciences, Mississauga, ON, Canada) at 100 µL/well (4 × 10⁴ cells/well, 12 × 10⁴ cells/cm²) and the plates were incubated at 37 °C for 24 h.

2.7.2. DEP Exposure

Lyophilized DEP suspensions were thawed at room temperature and sonicated for 20 min in an ice-cold ultrasonic water bath and diluted in complete DMEM (devoid of FBS. phenol red). The dilute particle suspensions were sonicated for 5 min in an ice-cold, ultrasonic water bath prior to dosing the cells in the 96-well plates. Cells in 100 μ L cell culture medium with 10% FBS were dosed with a 100 μ L particle suspension in culture medium (with no FBS) to have final doses 0, 10, 30, 100, 300 μ g/cm² (final volume of 200 μ L/well and the final concentration of FBS was 5%).

2.7.3. Cytotoxicity Analyses

J774A.1 cells were incubated for an additional 24 h post-exposure to DEPs before the assessment of cytotoxicity using an integrated bioassay that combined assays of cellular redox status (CellTiter Blue[®] Assay), energy metabolism (ATP assay), and membrane integrity (intracellular lactate dehydrogenase (LDH) release assay), as described by Kumarathasan et al., 2015 [46]. CellTiter-Blue[®] and the Cytotox [®] kit for LDH release assay were purchased from Promega Corporation (Madison, WI, USA). The ATP detection kit was procured from Lonza Rockland Corporation (Rockland, ME, USA). All of the exposure experiments were repeated three times for each cell line, with two technical replicates/experiments.

2.7.4. Cytokine Secretion

The cell culture supernatants of J774.A1 were assessed using a 23-plex mouse multiplex cytokine assay panel (Bio-Rad Laboratories (Canada) Ltd., Mississauga, ON, Canada) for secretion of IL-1 α , IL-1 β , IL-2, IL-3, IL-4, IL-5, IL-6, IL-9, IL-10, IL-12 (p40), IL-12 (p70), IL-13, IL-17A, eotaxin, G-CSF, GM-CSF, IFN- γ , KC, MCP-1 (MCAF), MIP-1 α , MIP-1 β , RANTES, and TNF- α following the manufacturer's protocols. Analyses were conducted on a Bio-Plex 200 multiplex luminescence assay system (Bio-Rad Laboratories Canada Ltd., Mississauga, ON, Canada).

2.8. In Vivo Exposure

2.8.1. Animals

Specific-pathogen-free BALB/c mice (male, weight 26.4 ± 0.2 g, mean \pm standard error) obtained from Charles River (St Constant, QC, Canada) were housed in individual Plexiglass cages on wood-chip bedding and were held under a 12:12 h dark: light cycle. Food and water were provided ad libitum. Animals were received and housed in the animal care facilities of Health Canada in Ottawa. All of the experimental protocols were reviewed and approved by the Animal Care Committee of Health Canada.

2.8.2. Intratracheal Instillation

The animals were administered 250 μ g of DEP or carbon black (n = 6) via a single intratracheal instillation, in line with previous reports. Carbon black was used as a reference particle as it is a widely accepted model particle in diesel studies. A group of animals were instilled with saline as the vehicle control (n = 6). The instillation of the experimental animals was conducted over a two-day period. In order to eliminate bias from the day of instillation, half the number of animals from each exposure group were instilled on day one and the other half of animals were instilled on day two. Before DEP instillation, the animals were anaesthetized via isoflurane inhalation and placed on their backs on a 40° slope. With the tongue pressed towards the lower jaw by a small sterile spatula, a 24-gauge catheter with a shortened needle was placed in the trachea. A sensitive pressure transducer was connected to the catheter to confirm the positioning of the instillation needle in the trachea. Particle preparations to be instilled were suspended by placing in an ultrasonic water bath for 5 min, followed by rigorous pipetting just prior to the removal of an aliquot for instillation. In a 250 µL SGE glass syringe (Fisher Scientific, Ottawa, ON, Canada), a sandwich of 50 μ L of air, 50 μ L of DEP suspension or saline (vehicle), and 150 μ L of air to be delivered into the lungs in this order was prepared. Prior to instillation, the pressure transducer was disconnected from the catheter. The tip of a glass syringe loaded with a particle suspension or saline was inserted into the catheter, and the material was delivered with a stroke of the plunger. Immediately after instillation, the catheter was removed, and the mouse was held head up momentarily to ensure the delivered material remained in the lungs. After the animals recovered from anesthesia, they were transferred back to their

2.8.3. Biological Samples

cages for recovery.

Following a 24 h recovery period after intratracheal instillation, the animals were anaesthetized via isofluorane. The blood was withdrawn via cardiac puncture into vacutainer tubes containing the sodium salt of EDTA at 10 mg/mL and PMSF at 1.7 mg/mL, mixed gently, and placed on ice. The diaphragm was then punctured, to expose the trachea, which was then cannulated. The lungs were filled via the intratracheal delivery of filter-sterilized, warm saline (0.9%, 37 °C) at 30mL/kg body weight [47]. The lungs were massaged gently by rubbing the thoracic cage. Saline was aspirated and reinjected twice more, and the primary bronchoalveolar lavage (BAL) was collected in a 15 mL centrifuge tube kept on ice. Secondary lavages were obtained with additional volumes of saline (5 mL/animal), three times, to increase the yield of lavage cells. The lavage fluids were centrifuged (1500 rpm for 10 min at 4 $^{\circ}$ C) to separate the cells from the supernatants. The cell pellets from both primary and secondary lavages were combined to recover the total BAL cells. Primary lavage supernatants were used to analyze biochemical endpoints. Secondary lavage supernatants were discarded. Whole-blood samples were centrifuged at 2000 rpm for 10 min at 4 °C to obtain plasma. Plasma aliquots were frozen at -80 °C. Lung and heart tissues were collected, flash-frozen in liquid nitrogen, and stored at -80 °C for reverse transcriptase-PCR analyses.

2.8.4. Cytology

Lung BAL cells were counted using a Coulter Multisizer II (Coulter Canada, Ville St-Laurent, QC, Canada), and differential cell counts were obtained from cytospin preparations using Wright stain and following standard procedures [48].

2.8.5. Cytokines

Levels of interleukin (IL)-1 α , IL-1 β , IL-2, IL-3, IL-4, IL-5, IL-6, IL-9, IL-10, Il-12 (p40), IL-12 (p70), IL-13, IL-17, eotaxin, granulocyte–macrophage colony-stimulating factor (G-CSF), GM-CSF, interferon (IFN)- γ , growth-related oncogene/keratinocyte chemoattractant (GRO/KC), monocyte chemoattractant protein-1 (MCP-1), macrophage inflammatory protein (MIP)-1 α , MIP-1 β , regulated on activation and normal T cell expressed and secreted chemokine (RANTES), and tumor necrosis factor- α (TNF- α) in the BAL fluid and plasma were analyzed using a 23-plex cytokine assay kit (Millipore Corporation, Billerica, MA, USA). The plasma levels of selectin, matrix metalloproteinase (MMP)-9, soluble intracellular adhesion molecule (sICAM), soluble vascular cell adhesion molecule (sVCAM), plasminogen activator inhibitor (PAI), apolipoprotein A1 (apoA1), apolipoprotein E (apoE), fibrinogen, and adiponectin were analyzed using cardiovascular multiplex cytokine assay kits (Millipore Corporation). The analyses were conducted according to the manufacturer's instructions using a Bio-Plex 200 multiplex luminescence assay system (Bio-Rad Laboratories Canada Ltd.).

2.8.6. Gene Expression Analyses

Lung and heart samples were homogenized in TRIzol® reagent (Invitrogen Canada, Inc., Burlington, ON, Canada), and the total RNA was isolated according to the manufacturer's instructions. The RNA was quantified using the RiboGreen RNA Quantitation Reagent and Kit (Molecular Probes, Eugene, OR, USA), and the total RNA was reverse transcribed using MuLV reverse transcriptase and random hexamers (Applied Biosystems, Mississauga, ON, Canada), according to the manufacturer's instructions. Primers for TATA Binding Protein, oxyguanine glycosylase (OGG)-1, IL-1 β , IL-6, endothelin (ET)-1, inducible nitric oxide synthase (iNOS), endothelial nitric oxide synthase (eNOS), cytochrome P450, family 1, subfamily A, polypeptide 1 (CYP1A1), metallothionein 2A (MT2A), prostaglandin– endoperoxide synthase 2 (PTGS2), heme oxygenase-1 (HO-1), and hypoxia-inducible factor (HIF)-3 α were obtained from Thomson et al. 2013 [49] or designed and validated to produce amplicons with an optimal annealing temperature of 60 °C (Supplementary Table S3). Real-time PCR was performed using 96-well plates in a spectrofluorometric thermal cycler (Lightcycler 480, Roche Diagnostics Canada, Laval, QC, Canada) using iQ SYBR Green Supermix (Bio-Rad Laboratories [Canada] Ltd.), as previously described by Thomson et al., 2013 [49]. A melt curve was conducted following each run to verify the product's purity. The expression was calculated relative to peptidylprolyl isomerase A expression using the delta-delta Ct method and expressed relative to time-matched controls.

2.8.7. Potency Calculations

Data for in vitro cytotoxicity and for the secretion of cytokines and chemokines in J774A.1 cells were normalized to control values to obtain the fold effect for each particle dose. Potency (β) was derived from the equation below.

Fold Effect =
$$(Dose + 1)^{\beta}$$

where β is the slope of the dose–response relationship for a given endpoint [50]. The dose–response data were fitted using CurveExpert v1.3 (D. Hyams, Hixson, TN, USA) to calculate cytotoxic potency and potencies based on cytokine responses in J774.A1 cells. Because only a single dose of particles was instilled in vivo, the in vivo biological potency was calculated as the ratio of the biological effect of a DEP to that of a saline vehicle control for a given bioassay (i.e., fold change).

2.9. Statistical Analyses

For in vitro exposure, cytotoxicity and cytokine secretion data were analyzed via two-way ANOVA with *PM* (DEP_{ULSD}, DEP_{B20C}, DEP_{B20S}, and DEP_{B20T}) and *DOSE* (0, 10, 30, 100, and 300 μ g/cm²) as factors. Datasets not meeting the assumptions of normality and equal variance for ANOVA were subjected to transformation (e.g., log₁₀, ln, inverse or square root or ranks) prior to analyses.

In vivo exposure responses were analyzed through the use of one-way ANOVA for differences between the groups of DEP_{ULSD}, DEP_{B20C}, DEP_{B20S}, DEP_{B20T}, CB, and saline-exposed animals. Datasets that did not meet the assumptions of normality and equal variance were assessed via Kruskal–Wallis one-way analysis of variance on ranks. For both one-way and two-way ANOVAs, pairwise multiple comparisons were carried out using Tukey's test to elucidate the pattern of significant effects ($\alpha = 0.05$). All of the analyses were conducted using SigmaPlot, version 12 (Systat Software, Inc., San Jose, CA, USA). Pearson product moment correlations between in vitro and in vivo biological potencies were calculated using SPSS version 15 (IBM Corporation, New York, NY, USA).

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3. Results

3.1. In Vitro Effects

3.1.1. Cytotoxicity

Biodiesel feedstock impacted the cytotoxicity of DEP (Figure 1). Higher doses (100 and 300 μ g/cm²) of DEP_{ULSD} and DEP_{B20C} were relatively more cytotoxic than the same doses of CB, DEP_{B20S} or DEP_{B20T} based on all cytotoxicity assays (two-way ANOVA, *PM* × *DOSE* interaction, *p* < 0.001; Figure 1).

3.1.2. In Vitro Cytokine Secretion

Secretion of inflammatory cytokines and chemokines by J774A.1 cells in response to DEP exposure was impacted by biodiesel feedstock. Two-way ANOVA results identified PM main effect for IL-1 β and IL-12 (p70) (p < 0.05, Figure 2A,B), whereas Dose main effect was noticed with IL-10 (p < 0.05, Figure 2C).

Furthermore, secreted GM-CSF and TNF- α levels revealed the main effects of PM and dose (p < 0.05; Figure 2D,E), with all particles exhibiting increased TNF- α levels at 300 µg/cm² dose compared to the vehicle control and DEP_{ULSD} and DEP_{B20C} showed relatively higher levels of TNF- α at this dose compared to the other particles tested in this study. Two-way ANOVA analysis results for secreted G-CSF and RANTES identified Particle × Dose interaction (p < 0.05, Figure 2F,G). The secretion of RANTES decreased with dose, with relatively larger declines noted after exposure to the highest doses of DEP_{ULSD} or DEP_{B20C}.



Figure 1. The figure shows the impact of biodiesel feedstock on the cytotoxicity of diesel exhaust particles assessed in the J774.A1 macrophage cell line. (Values are mean \pm SEM; n = 3 experiments; similar notations (letters) are significantly different (p < 0.05) from each other). (**A**) Intracellular ATP content, Two-way ANOVA, *PM* × *DOSE*, p < 0.001. (**B**) Intracellular LDH content, Two-way ANOVA, *PM* × *DOSE*, p < 0.001. (**C**) Resazurin reduction, two-way ANOVA, *PM* × *DOSE*, p < 0.001.

3.2. In Vivo Effects

3.2.1. BAL Neutrophil Counts

All particles, including CB, caused a significant increase in the number of lung lavage neutrophils compared to saline (the vehicle control) 24 h after the intratracheal instillation of particles in these mice (one-way ANOVA, p < 0.05; Figure 3), with increasing trends noted in the animals instilled with DEP_{ULSD}, DEP_{B20C}, and DEP_{B20T} compared to the other particles.



Figure 2. Cont.



Figure 2. The figure shows in vitro cytokine secretion in J774.A1 in response to exposure to diesel exhaust particles. (Values are mean \pm SEM; n = 3 experiments; similar notations (letters) are significantly different (p < 0.05) from each other) (**A**) IL1- β , 2-way ANOVA, PM main effect, p = 0.001. (**B**) IL-12 (p70), 2-way ANOVA, PM main effect, p = 0.005. (**C**) IL-10, Two-way ANOVA, DOSE Main Effect, p = 0.002. (**D**) GM-CSF, *PM* Main Effect, p = 0.011. *DOSE* Main Effect, p = 0.047. (**E**) TNF- α , *PM* main effect, p < 0.001; *DOSE* Main Effect, p < 0.001. (**F**) G-CSF, 2-way ANOVA, *PM* × *Dose* interaction, p < 0.05 (**G**) RANTES, 2-way ANOVA, *PM* × *Dose* interaction, p < 0.001.

3.2.2. BAL Cytokines

The levels of several inflammatory cytokines were also significantly increased in the bronchoalveolar lavage fluid in response to particle exposure in comparison to the saline vehicle control (one-way ANOVA, p < 0.05), with DEP_{ULSD} and DEP_{B20C} exposures showing increased cytokine levels in general compared to CB (Figure 4).



Figure 3. The figure shows the BAL neutrophil profiles following the instillation of DEPs (values are mean \pm SEM; n = 6/group; One-way ANOVA results: similar notations (letters) are significantly different (*p* < 0.05) from each other).



Figure 4. Cont.





In comparison to CB, instillation to DEPULSD significantly (p < 0.05) increased the levels of IL-6 and TNF- α (One-way ANOVA, p < 0.05, Figure 4D,L). Although the effects of DEPB20S or DEPB20T instillation were generally greater compared to the effects of CB instillation, the differences were not statistically significant.

3.2.3. Lung Gene Expression

In general, lung gene expressions after DEPULSD or DEPB20C exposures were comparable to responses of other DEPs (Figure 5). DEPULSD exposure led to significant increases in the expressions of lung IL-1 β and MT2A and a significant decrease in IL-6 when compared to the effects of CB exposure (Figure 5A,C,E). Lung CYP1A1 was significantly decreased by exposure to DEPULSD in comparison to DEPB20T (one-way ANOVA, p < 0.05, Figure 5F).



Figure 5. The figure shows the impact of DEP exposures on various gene expression (**A**–**F**) profiles in the lung tissue (values are mean fold change relative to time-matched control (saline) \pm geometric standard deviation; n = 6/group; one-way ANOVA results: similar symbols (letter) are significantly different (p < 0.05) from each other).

3.2.4. Heart Gene Expression

The instillation of DEP significantly (p < 0.05) altered the expression of HMOX-1 and iNOS in the heart in comparison to saline, with the expression of HMOX-1 significantly decreased by exposure to DEP_{B20C} and DEP_{B20T} and the expression of iNOS significantly decreased by exposure to DEP_{ULSD} and DEP_{B20T} particles (One-way ANOVA, p < 0.05, Figure 6) compared to the vehicle control saline.

3.2.5. Plasma Cytokines

Although the levels of PAI-1 and s-ICAM were impacted (Figure 7) by the treatments (one-way ANOVA, PM main effects, p < 0.05). The largest increases for PAI-1 and s-ICAM were observed after DEP_{ULSD} and DEP_{B20C} instillation. Although not statistically significant (p = 0.071), the levels of G-CSF were also increased by exposure to DEPs, with DEP_{ULSD} and DEP_{B20C} showing relatively higher responses compared to other DEPs.



Figure 6. The figure shows the alteration of gene expression profiles for (**A**) HMOX-1 and (**B**)iNOS in the heart tissue following DEP exposures (values are mean fold change relative to time-matched control (saline) \pm geometric standard deviation; n = 6/group; one-way ANOVA results: similar symbols (letters) are significantly different (*p* < 0.05) from each other).



Figure 7. The figure shows the changes in plasma cytokine levels (**A**–**C**) following DEP exposures (values are mean fold change relative to time-matched control (saline) \pm geometric standard deviation; n = 6/group; one-way ANOVA results: similar symbols (letters) are significantly different (*p* < 0.05) from each other).

3.3. Correlations between In Vivo vs. In Vitro Toxicity Endpoints

In vitro cytotoxicity with secreted TNF- α in vitro (cellular ATP content, r = 0.93, p = 0.02; energy metabolism, r = 0.81, p = 0.1; LDH release, r= 0.82, p = 0.09). In vitro cytotoxic potency measured by examining the cellular ATP content, resazurin reduction (CTB), and LDH release correlated strongly (r > 0.9) and significantly (p < 0.05) with the levels of the inflammatory cytokines of G-CSF, IL-1a, IL-3, IL-6, IL-10, IL12(p40), MCP-1, MIP-1b, and TNF- α in the bronchoalveolar lavage fluid BAL (Table 4) after in vivo exposure. The cellular ATP level was significantly positively correlated with lung gene expressions for MT-2A, IL-1 β and IL-6, while cellular CTB and LDH were negatively correlated with

lung gene expression for CYP1A1. Cellular cytotoxicity endpoints were also positively correlated (p < 0.05) with extrapulmonary effects, including IL-1 β gene expression in the heart and levels of MMP-9, G-CSF, PAI-1, s-ICAM, and s-VCAM in plasma. Similarly, positive correlations (p < 0.05) were observed between in vitro-secreted TNF- α levels and BAL cytokines, lung gene expressions (IL-1 β and IL-6), heart gene expression (IL-1 β), and plasma cytokine levels (G-CSF, PAI-1, and s-VCAM)

Table 4. The table show in vivo responses significantly correlated to a measure of cytotoxicity or secreted TNF- α in vitro: strength and significance of correlations *.

In Vivo Endpoints		I	n Vitro Cytotoxici	In Vitro TNF-α Secretion	
Compartment	Endpoint	ATP	СТВ	LDH	
BAL cells	PMN Cell Number	0.792 (0.110)	0.688 (0.199)	0.739 (0.153)	0.774 (0.124)
	G-CSF	0.945 (0.015)	0.919 (0.027)	0.923 (0.025)	0.909 (0.032)
	IL-1a	0.964 (0.008)	0.850 (0.068)	0.871 (0.054)	0.993 (0.001)
	IL-3	0.901 (0.037)	0.708 (0.181)	0.756 (0.139)	0.974 (0.005)
	IL-4	0.781 (0.119)	0.656 (0.230)	0.634 (0.251)	0.947 (0.014)
	IL-6	0.986 (0.002)	0.953 (0.012)	0.980 (0.003)	0.889 (0.044)
BAL cytokines	IL-10	0.842 (0.074)	0.920 (0.027)	0.888 (0.044)	0.884 (0.046)
	IL-12(p40)	0.921 (0.026)	0.901 (0.037)	0.913 (0.030)	0.926 (0.024)
	MCP-1	0.967 (0.007)	0.915 (0.029)	0.931 (0.021)	0.931 (0.021)
	MIP-1a	0.821 (0.088)	0.631 (0.254)	0.652 (0.233)	0.967 (0.007)
	MIP-1β	0.983 (0.003)	0.892 (0.042)	0.921 (0.026)	0.961 (0.009)
	КС	0.867 (0.057)	0.668 (0.217)	0.722 (0.169)	0.920 (0.027)
	TNF-a	0.972 (0.006)	0.850 (0.068)	0.915 (0.029)	0.890 (0.043)
Lung	CYP1A1	-0.867 (0.057)	—0.966 (0.008)	-0.952 (0.012)	-0.682 (0.204)
	MTII	0.913 (0.031)	0.748 (0.146)	0.840 (0.075)	0.838 (0.076)
gene expression	IL-1β	0.949 (0.014)	0.793 (0.109)	0.869 (0.056)	0.897 (0.039)
	IL-6	0.944 (0.016)	0.795 (0.108)	0.873 (0.053)	0.879 (0.050)

In Vivo Endpoints		I	n Vitro Cytotoxici	In Vitro TNF-α Secretion	
Compartment	Endpoint	ATP	СТВ	LDH	
Heart gene expression	IL-1β	0.906 (0.034)	0.922 (0.026)	0.880 (0.049)	0.913 (0.030)
	MMP-9	0.817 (0.091)	0.913 (0.030)	0.925 (0.025)	0.576 (0.310)
Plasma cytokines	G-CSF	0.966 (0.008)	0.843 (0.073)	0.909 (0.033)	0.895 (0.040)
	PAI-1	0.962 (0.009)	0.836 (0.078)	0.904 (0.035)	0.879 (0.050)
	s-ICAM	0.911 (0.031)	0.965 (0.008)	0.943 (0.016)	0.831 (0.081)
	s-VCAM	0.899 (0.038)	0.895 (0.040)	0.863 (0.059)	0.923 (0.025)

Table 4. Cont.

* Pearson product moment correlation; statistically significant correlations are highlighted in bold-faced font within brackets.

4. Discussion

The selection of biodiesel blends used in this work was based on the fact that, at the time of testing, the major vegetable oils used in the production of biodiesels in the United States and Canada were soybean oil and canola oil. Another feedstock being used was animal tallow, and waste oils including grease. Our work showed engine load-related PM mass changes across all fuel types. We also noticed some differences in NOx levels due to engine load. Furthermore, the type of biodiesel blend influenced the emitted NOx levels (Table 3), which is consistent with previous reports about biodiesel sourcedependent changes in NOx emissions [29,30]. Nevertheless, the focus of this work was to examine the relative toxicities of PM emitted when using different fuel types to support related health risk analysis. In this study, diesel exhaust particles from biodiesel blends (B20) with ULSD were examined for toxicity characteristics to understand the influence of feedstocks typically used in Canada on emitted particle toxicity. An effort was made to identify any consistency between the in vitro and in vivo toxicity behaviors of these particle emissions. Diesel exhaust particles mainly comprise two size modes, including fine particulate matter (PM2.5) based on our previous work [51] and are inhalable. Thus, our work here mainly focused on the pulmonary toxicity of these DEPs. For in vitro toxicity testing, the phagocytic mouse monocyte/macrophage cell type (J774A.1) was chosen since this cell type has been widely employed in in vitro pulmonary toxicity studies of PM, including engineered nanoparticles; in addition, the in vitro toxicity data from the use of this cell type will be suitable for correlation testing with the animal model used in this study to generate in vivo toxicity data.

The in vitro cytotoxicity findings from this work showed that DEPs from ULSD and B20C consistently negatively affected all cytotoxicity endpoints more compared to other particle treatments, with greater responses seen in terms of cellular ATP (cell metabolism) and LDH (cell membrane integrity) contents. While most of the cell-secreted protein profiles showed main effects in terms of PM and dose, G-CSF and RANTES exhibited modifications to PM-exposure-related responses in relation to exposure dose. Carbon black appeared to be relatively less responsive compared to ULSD and the B20 biodiesel blends used in this work. Furthermore, it was interesting to note that the proinflammatory cytokine (TNF)- α profiles showed dose-related increases with particle exposures, while the anti-inflammatory IL-10 showed a decreasing trend with the dose of PM exposures, suggesting potential proinflammatory status in cells exposed to these particles. Paricularly, ULSD and B20C were more potent, followed by B20S and B20T, when compared to CB. The DEP of B20T was relatively less responsive among the biodiesel blends, and this may be attributed to

its relatively higher saturated fat content compared to B20C and B20S that may have led to reduced carbonyl compound formation in particle emissions and, thus, relatively low cytotoxicity. Additionally, the fact that B20C is more potent than B20S in this work is in line with a report on mutagenicity testing conducted by Demarini et al., 2019 [52], where canolabased biodiesel emission particles were more potent than soy-based biodiesel emission particles. Furthermore, in vivo exposure to these DEPs and CB resulted in increased BAL neutrophils for all particle treatments compared to saline vehicle controls, and similarly secreted BALF proinflammatory cytokine levels were also increased for DEPs compared to the saline vehicle control group. Additionally, DEPs from ULSD and B20C exhibited relatively increased biological responses compared to CB and DEPs derived from B20S and B20T, which is consistent with in vitro toxicity findings in this work.

Previous reports pointed to both the increased and decreased toxicity of combustion emissions from biodiesels or biodiesel blended fuels; the variance in the findings was generally due to the feedstock and blend ratios employed, engines and run cycles tested, and specific exhaust components (i.e., primary versus secondary particulate emissions, semi-volatile components, or whole exhaust) assessed. For example, Liu et al. 2008 [53] compared the cytotoxic potencies of particulate and semi-volatile constituents of diesel exhaust generated from the combustion of petroleum diesel or a palm methyl ester biodiesel blend at a range of blend ratios (10, 30, 50, 75, and 100%) in a four-stroke water-cooled, non-catalyst generator with a constant output power in BEAS-2B cells, and showed no significant differences in the toxicity of the particulate constituents between petroleum diesel and the biodiesel blends. However, the semi-volatile constituents from biodiesel exhaust were more toxic than those from petroleum diesel at all levels of blending. Likewise, the increased toxicity and heightened inflammatory response in cultured human epithelial cells exposed to whole exhaust from the combustion of a 20% v/v blend of canola biodiesel when compared to cells exposed to petroleum diesel exhaust, as reported by Mullins et al. 2014 [37] may, in part, relate to the toxicity of the gas phase that includes semi-volatiles. Using a four-stroke direct injection diesel engine from a tractor operated at a heavy-duty 13-mode test cycle ECE R49, Bunger et al., 2000 [54] showed that soot generated from the combustion of rapeseed biodiesel caused four-fold stronger toxicity in mouse fibroblasts when compared to soot from petroleum diesel. However, such load-dependent effects were only noted at 'idling' and not at the rated power. It was suggested that the higher toxicity was caused by carbonyl compounds and unburned rapeseed methyl ester-based fuel when the engine was idling. An integrated assessment of experimental factors such as engine type and load, biodiesel feedstock, blend ratios, batch-to-batch variability, and exposure to gas phase versus particulate phase versus whole exhaust will be critical to a holistic assessment of the automotive use of biodiesels derived from different feedstock. Our study only focused on the toxicity of particulate emissions, a key component of urban ambient airborne particulate matter that is implicated in a number of adverse population health outcomes. We combined particles derived from engine runs at 25% and 50% steady-state loads at a mass ratio of 2:1. While this maximized the sample mass available for our in vitroin vivo comparisons, the mixing ratio also approximated the ratio of mass emission rates (g/bhp-hr) at the two steady-state loads for all the diesel exhaust particles (DEP_{ULSD}, 1.81; DEP_{B20C} , 1.94; DEP_{B20S} , 1.96; and DEP_{B20T} , 1.98), thereby giving equal weight to both loads in terms of mass emissions.

The generally lower biological potency of primary particulate emissions derived from biodiesel blends than those derived from petroleum diesel in our study may relate to some of the previously documented decreases in the concentration of biologically reactive emission constituents with biodiesel blended fuels. In a comparison of particulate emissions from a compression ignition engine when using ultra-low sulfur diesel or blends of canola, soy, or tallow biodiesels with ultra-low sulfur diesel, Surawski et al. 2011 [30] showed that particle numbers and size distributions were feedstock-dependent, but the concentrations of particle and vapour phase PAHs, and reactive oxygen species were less sensitive to feedstock or blend percentage, and the concentrations were reduced with biodiesels when compared to ultra-low sulfur diesel. In a separate study, when a number of factors, such as nanoparticle size and number distribution, surface area distribution, elemental and organic carbon content, and polycyclic aromatic hydrocarbons adsorbed onto the particle surfaces were considered together to calculate toxic equivalency factors for B20 biodiesels of canola, soy, and animal tallow, it was adjudged that biodiesel particle toxicity was considerably lower in comparison to mineral diesel [55]. The study concluded that in spite of the higher PAH loading of emission particles from B20 combustion, their contribution to overall toxicity was less than that in diesel-derived particles. However, in the current work, although the in vitro toxicity of B20 blends of soy and animal tallow was relatively lower compared to ULSD, B20C (canola) exhibited cytotoxicity responses similar to that of ULSD.

In this study, the in vitro assessment of DEPs focussed on the cytotoxicity and secretion of inflammatory cytokines and the in vivo assessment focused on a number of markers of inflammation, and cardiovascular effects; the effects were tested for correlation in an attempt to associate biological potency measured in vitro to in vivo effects related to potential pathophysiological outcomes of particle exposure. Several in vitro measures of toxicity based on cellular redox status, energy metabolism, and membrane integrity were well correlated to each other and to the in vitro secretion of the proinflammatory cytokine tumor necrosis factor (TNF)- α (Table 4). While it is known that in vitro exposure of cells to diesel exhaust particles elicits the secretion of inflammatory cytokines [56], noteworthy correlations were observed between cytotoxicity and inflammatory responses measured in vitro as well as in vivo. Acute inflammatory response to DEP exposure suggested by increased neutrophil extravasation in the lung was complemented by increased levels of the proinflammatory cytokines of IL-1 α , IL-3, IL-6, IL-12(p40), MCP-1, and MIP-1 β in BALF. The increased gene expression of MT2A and the decreased expression of CYP1A1 mRNA in the lung with diesel particle exposure could be in response to the heightened inflammation status [57,58]. The instillation of diesel exhaust particles produced effects on the heart, with the increased gene expression of IL-1 β , a proinflammatory cytokine, and decreased heart expression of heme oxygenase-1 (HMOX-1), an enzyme with an anti-inflammatory role. While increased plasma levels of G-CSF may also represent an inflammatory component, increased levels of the adhesion molecules s-ICAM may have cardiovascular consequences, especially in relation to its involvement in atherosclerotic processes [59]. The increased plasma concentration of plasminogen activation inhibitor (PAI)-1 in response to DEP exposure is also significant, as PAI-1 is known to play a significant role in fibrosis, a pathological formation of connective tissue [60].

The similarity of the ranking of in vitro and in vivo potency of DEP (ULSD ~ B20C > B20S > B20T) shows the potential utility and sufficiency of in vitro assays as screening assays for prioritizing environmental particles for animal studies and strengthens confidence in the validity of the toxicity ranking of the biodiesel emission particles. The use of in vitro assays for the toxicity screening of emissions arising from complex technology matrices consisting of engine types, fuels, run cycles, and after-treatment configurations in order to prioritize conditions for animal testing will be of considerable value in terms of cost, throughput, and reducing the use of animals. Our in vivo data show increased pulmonary inflammation, greater perturbation of cardiac gene expression, and changes in plasma markers relevant to the inflammation and cardiovascular effects relating to DEP in comparison to carbon black used as a surrogate of DEP with relatively low levels of organics, confirming that the effects are not simply a generalized response to particle exposure but rather are attributable to the composition and distinct properties of DEP.

5. Conclusions

Both in vitro and in vivo analyses from the current work showed that biodiesel blending can decrease the biological potency of diesel exhaust particles based on the feedstock. The toxic potency of primary emission particles derived from the combustion of petroleum diesel (DEP_{ULSD}) and the canola biodiesel–petroleum diesel blend (DEP_{B20C}) were similar based on a number of in vitro and in vivo bioassays, while emission particles derived from the combustion of soy and tallow–waste oil biodiesel blends (DEP_{B20S} and DEP_{B20T}, respectively) were less potent. Overall, our findings from the use of in vitro and in vivo PM exposure models suggest that the source of biodiesel can be an important determinant of DEP toxicity, and these results provide insight into potential underlying toxicity mechanisms, notably, inflammatory process. Advanced engine technologies and emission treatment technologies have significantly reduced gaseous and particulate emissions from new engines. Understanding the physicochemical determinants of toxicity with biodiesel blended fuels in newer engines needs to be considered in future toxicological studies to support the health risk assessment of these emissions.

Supplementary Materials: The following supporting information can be downloaded at: https: //www.mdpi.com/article/10.3390/toxics12040290/s1, Table S1: Analyses of Unblended Biodiesels (B100); Table S2: Sample collection procedure, analytical methods, and instrumentation employed in the analyses of exhaust emissions during diesel exhaust particle sampling; Table S3: Primers used in the real-time PCR analysis of gene transcripts in the lung and heart tissues.

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