Assessment of 2-Pentadecyl-2-oxazoline Role on Lipopolysaccharide-Induced Inflammation on Early Stage Development of Zebrafish (*Danio rerio*)

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Abstract: Lipopolysaccharide (LPS), or bacterial endotoxin, is an important virulence factor in several human and animal pathologies. Oxazoline of Palmitoylethanolamide (PEAOXA) has shown strong anti-inflammatory activity in several animal models. LPS was applied for 24 h to zebrafish embryos to induce inflammation, and then the anti-inflammatory action of PEAOXA was evaluated for the first time in the zebrafish model (*Danio rerio*). Different concentrations of PEAOXA were tested for toxicity on zebrafish embryonic development; only the highest concentration of 30 mg/L showed toxic effects. Quantitative RT-PCR was applied to detect Tumor necrosis factor-α, Interleukin 1β, 6, and 8, and members of the nuclear factor kappa-light-chain-enhancer of activated B cells (NF-kB). Exposure to LPS induced an increase in pro-inflammatory cytokines (tumor necrosis factor and interleukin 1, 6, and 8) in both gene and protein expression, as well as an increase of the nuclear factor kappa-light-chain-enhancer of activated B cells (NF-kB) and the nuclear factor kappa light polypeptide enhancer in B-cells inhibitor (IkBα) gene expression. Furthermore, acute LPS exposure also induced an increase in tryptase release, related to mast cell activity, and in the production of apoptosis-related proteins (caspase 3, 8, and bcl-2). Treatment with PEAOXA 10 mg/L significantly counteracts LPS-induced inflammation in terms of cytokine expression and decreases tryptase release and the apoptosis pathway.

Keywords: zebrafish; inflammation; PEAOXA; cytokines; apoptosis

1. Introduction

Inflammation can be defined as a natural bodily response that is triggered by an injurious stimulus or toxic condition. The inflammatory response when properly regulated helps to maintain cellular homeostasis and resistance to disease, but its persistence, i.e., chronic inflammation, may cause to autoimmune disorders related to various diseases, such as asthma, diabetes, neurodegenerative diseases, or arthritis [1]. Several cytokines are involved in the inflammatory process, such as tumor necrosis factor (TNF) and interleukin
which are secreted and released once damage occurs, and immune cells, such as macrophages, are activated [2–4]. The adaptive immunity of zebrafish develops only after 4 weeks, within which time zebrafish defend themselves only by innate immunity. The study in the early life stages of zebrafish therefore allows the assessment of the innate defense mechanism without the involvement of adaptive immunity [5].

Lipopolysaccharide (LPS) normally bind toll-like receptor 4 (TLR4) on the surface of the cell membrane that leads to the activation of TIR-domain-containing adapter-inducing interferon-β (TRIF) signaling pathway or myeloid differentiation factor 88 (MyD88) [6,7]. Nuclear Factor-kappa B (NF-kB) was a classic inflammatory signaling pathways downstream of MyD88-dependent pathway [8,9]. As a consequence of NF-kB pathway activation, pro-inflammatory factors release occurs, such as IL-1β, IL-6, TNF-α [10]. Direct inhibition of cytokines or their release is considered a good anti-inflammatory strategy [10,11].

LPS-induced inflammation is among the one of the most studied in various models both in vitro and in vivo to investigate interference with inflammatory pathways [12]. One of the assays often used to evaluate the anti-inflammatory activity of different natural compounds involves the stimulation of macrophage cells with LPS [13–15]. One of the most important factors related to fish aquaculture and its development is the increase of bacterial infections that consequently cause economic repercussions. Gram-negative bacteria are associated with numerous pathologies [16]. LPS is an important component of the outer membrane of Gram-negative bacteria. Among the various effects of LPS are the induction of cytokines and other acute phase proteins production and they present a key role in the pathological activities of several fish [17]. As a model of drug screening in recent years the zebrafish (Danio rerio) is being used a lot because of its short generation time and low cost but also due to the high size of the clutch in the embryonic and larval stages, and its optical transparency that allows the visualization of pathogens and lesions in real time [18]. Although the adaptive immune system of zebrafish does not develop until 4–6 weeks postfertilization, this model remains widely used to study the innate immune response in antiendotoxic drugs study [19–21].

Palmitoylethanolamide oxazoline (PEAOXA), identified in natural compounds such as coffee, has shown protective action by reducing inflammation and hyperalgesia, with more efficacy than its homologous the PEA [22]. This has been seen to be possible thanks to a modification in the structure of Palmitoylethanolamide (PEA), an endogenous lipid with anti-inflammatory properties, obtaining 2-pentadecyl-2-oxazoline (PEAOXA), the oxazoline of PEA [23]. The effects of PEAOXA on the inhibition of inflammatory processes have been seen in a mouse model of traumatic brain injury and spinal cord injury [24], and more recently the effects of PEAOXA in reducing inflammation and oxidative stress associated with vascular dementia in an experimental model of repeated bilateral common carotid arteries occlusion have been evaluated [25].

In this study, LPS mediated inflammation on zebrafish larvae were applicated to assess the protective effect of PEAOXA. Furthermore, the expression of several inflammatory mediators was determined to explore the anti-inflammatory mechanism of PEAOXA.

2. Materials and Methods

2.1. Zebrafish Maintenance and Embryo Collection

Wild type (WT) mature zebrafish with an age of 6 months were used for embryos’ production. Zebrafish Maintenance and Embryo Collection of fertilized eggs were provided from the Center of Experimental Fish Pathology (Centro di Ittiopatologia Sperimentale della Sicilia, CISS), University of Messina, (Messina, Italy). The fish were fed both with dry and live food twice a day at 3% of body weight (BW). For a successful reproduction, mature females and males were mated at 2:1 ratio. The day after, the eggs were collected in a room at 28 °C, bleached, and afterwards non-fertilized eggs were discarded. Only embryos which reached the blastula stage were used for experiments. Fish Embryo Toxicity (FET) test was performed according to OECD [26] and ISO 15088.
2.2. Viability, Morphology, Hatching, and Heart Rate after PEAOXA Exposure

At 4 h post fertilization (hpf) healthy embryos were placed in 24-well culture plates (1 embryo in 2 mL solution/well). Zebrafish embryos were exposed to PEAOXA for 26–96 hpf to measure the toxic effects over a continuing observation period. PEAOXA was dissolved for stock solution as previously described [27] in methanol 0.05%. To examine the survival rate and morphology of embryos/larvae, PEAOXA (kindly offered by Epitech Group SpA (Saccolongo, Italy) dissolved in the stock solution was after diluted with Embryo medium (15 mM NaCl, 0.5 mM KCl, 1 mM CaCl$_2$, 1 mM MgSO$_4$, 0.15 mM KH$_2$PO$_4$, 0.05 mM Na$_2$HPO$_4$, 0.7 mM NaHCO$_3$ a pH 7.3) to the tested different concentration (0.1, 1, 3, 10, and 30 mg/L) to assess the concentration-dependent toxicity. We decided to use these concentrations following previous tests conducted by our laboratory and based on in vitro studies in the literature. Different concentrations were tested to observe the highest dose without toxic effects to be used for anti-inflammatory action. In addition, the group exposed with only the vehicle of PEAOXA, methanol 0.05%, showed no morphological abnormalities. We used 4 replicates for experimental group, 20 eggs in each replicate, and 3 independent experiments. During the exposure time, researchers observed the embryonic development and various parameters including mortality, hatching rate, heartbeat rate, and abnormalities in hatched larvae as previously described [28]. The solutions were renewed, and embryonic/larval mortality and hatching rate were evaluated every 24 h, while the heart rate was measured at 48, 72, and 96 hpf. During the exposure period, photographs of the embryos were made under a stereomicroscope (Leica M0205C, Multifocus, Wetzlar, Germany).

2.3. Application of LPS to Zebrafish Embryos

After evaluating non-toxic concentrations of PEAOXA, these were used to counteract LPS-induced inflammation. The application and the concentration of LPS to Zebrafish Embryos was according to the previous study [29]. LPS (10 µg/mL) was applied by immersion at 6 hpf to induce inflammation, for a total of 24 h hours. PEAOXA (3 and 10 mg/L) was added starting at the end of 24 h of LPS exposure.

2.4. Gene Expression Analysis

The total RNA from zebrafish larvae (20 per experimental group of each experiment) was homogenized and isolated in 0.50 mL TRizol reagent (Invitrogen, Waltham, MA, USA) according to the manufacturer’s instructions. Total RNA was isolated according to the manufacturer’s instructions. The ratio of absorbance at 260–280 nm, as well as the banding patterns on a 1% agarose formaldehyde gel, were used to verify the quality of the RNA in each sample. RNA quality was evaluated by gel electrophoresis, with the concentration measured with NanoDrop 2000 (Thermo Scientific Waltham, MA, USA iScript RT-PCR kit (Bio-Rad, Hercules, CA, USA)), which was used to synthesize first-strand cDNA according to manufacturer’s recommendations. The reverse transcription master mix was prepared adding to 1 µg of RNA template the iScript RT Supermix (5× RT supermix with RNase H+ Moloney (gray cap, 25 or 100 reactions) murine leukemia virus (MMLV) reverse transcriptase, RNase inhibitor, dNTPs, oligo (dT), random primers, buffer, MgCl$_2$, and stabilizers) and the nuclease-free water. Briefly, 1 µg/µL total RNA was mixed with 5 µL of 5× iScript® (Bio Rad, Hercules, CA, USA). Reaction Mix and 1 µL of iScript® Reverse Transcriptase and RNase-free water to make the final volume of 25 µL. The complete reaction mix was incubated in a thermal cycler (Priming 5 min at 25 °C, Reverse transcription 20 min at 46 °C, RT inactivation for one minute at 95 °C). Real-Time PCR was performed with a 20-µL volume containing 10-µL of 1× SsoFast EvaGreen Supermix (Bio-Rad, Hercules, CA, USA), 1 µL of cDNA, 7 µL of RNase/DNase-free water, and 500 nM each primer. PCR conditions were: initial denaturation at 95 °C for 15 min, followed by 45 cycles of amplification at 95 °C for 20 s and 60 °C for 40 s. Final extension at 60 °C for 60 s and hold at 4 °C were then performed on StepOnePlus Real-Time PCR System (Applied Biosystems, Foster City, CA, United States). All the qPCR reactions were
performed with three parallel samples; the negative control contains no sample template. Table 1 shows the detailed information on the primers of β-actin, inflammatory related genes (TNF-α, IL-6, IL-8, IL-1β, IκBα, and NF-κB), and apoptotic related genes (caspase 3, bax, and bcl-2), as previously reported [30,31]. β-actin was used as an internal control for normalizing relative expression levels between samples [32–34]. The main PCR protocol referenced a previous study [35]. The relative transcriptional levels of each gene were calculated according to a previous study [36]. Data analysis was performed using the 2^−ΔΔCt method and the results are expressed as fold changes.

Table 1. Primers for Real-Time PCR.

| Gene                  | Primer Orientation | Nucleotide Sequence            |
|-----------------------|--------------------|-------------------------------|
| b-actin               | forward            | 5′-AGAGCTATGAGCTGCTGACG-3′    |
|                       | reverse            | 5′-CCGCAAGATTCCATACCCA-3′     |
| Inflammatory          |                    |                               |
| pathway genes         |                    |                               |
| TNF-α                 | forward            | 5′-GCTGGATCTCCAAGTGCTCGGTA-3′ |
|                       | reverse            | 5′-TGTTGACTCCACACATGCTGAC-3′  |
| IL-1β                 | forward            | 5′-TGGACTTCGACCACAAAATG-3′    |
|                       | reverse            | 5′-GTTCACTTCACGCTCTTGGAT-3′   |
| IL-6                  | forward            | 5′-AGACCGCTGGCTGCTGAAA-3′     |
|                       | reverse            | 5′-TTTGTGATGCTGTTACACAGGA-3′  |
| IL-8                  | forward            | 5′-GCCGGCTGCGATTGAAAACAGA-3′  |
|                       | reverse            | 5′-TGGGAACATGCTGCTGTAC-3′     |
| NF-κB                 | forward            | 5′-CGCAGACATGGAGATGGAGA-3′    |
|                       | reverse            | 5′-CTGTTCAAGTACGGGTGTT-3′     |
| Apoptosis pathway     |                    |                               |
| genes                 |                    |                               |
| cas-3                 | forward            | 5′-CCGCTGGCCATCACCTA-3′       |
|                       | reverse            | 5′-ATCCITTCAGACCCTC-3′        |
| Bax                   | forward            | 5′-GGCTATTTCAACCAGGGTCC-3′    |
|                       | reverse            | 5′-TGCGAATCACCAGGAGTTAC-3′    |
| bcl-2                 | forward            | 5′-CACTCGTTCAGACCCTCAT-3′     |
|                       | reverse            | 5′-ACGCTTTCCAGCAGCAT-3′       |

2.5. Western Blot

Western blot analysis was performed as previously described [37,38]. LPS-exposed zebrafish larvae were washed twice with PBS (pH 7.4) and then homogenized in ice-cold RIPA buffer (Tris-HCl pH 7.4 50 mM; NaCl 150 mM; Sodium Deoxycholate 1%; Triton-X100 0.05%; EDTA 0.5 mM Na3VO4 1 mM; NaF 50 mM, 0.1% SDS) to extract proteins (Wuhan Boster Biological Technology, Wuhan, China). Each set of larvae (20 per experimental group of each experiment) was pooled for protein preparation, such that n = 1 refers to protein from these 20 larvae. The homogenates were centrifuged at 14,515 × g for 10 min and supernatants were collected. Protein concentrations were determined by the Bradford method [39,40]. The samples were heated to 95 °C for 5 min for the denaturation. Equal quantities of proteins (40 µg) for each sample were loaded and separated by SDS-PAGE (10% gels) for 2 h at 100 V under denaturing conditions. After electrophoresis the proteins were transferred from gels onto polyvinylidene fluoride (PVDF) membrane 0.42 µm (GE Amersham, Casoria, NA, Italy). The proteins were transferred onto PVDF membranes in a Tris-glycine transfer buffer at 35 V for 10 min. Subsequent to blocking with 5% skim milk in Tris-buffered saline containing 0.1% Tween-20 (TBST) at room temperature for 2 h, the membrane was incubated with primary antibodies at 4 °C overnight, antibody against caspase-3 (Abcam ab13847, 1:500) Bax (Abcam ab32503, 1:800), bcl-2 (Abcam ab18285, 1:800), NF-κB (Antibodies A323357, 1:500) and IκBα (ANAWA pS32/S36, 1:500), and the proteins expression was normalized according to the expression of GAPDH (Abcam ab181602, 1:1000). After being washed three times with TBST, the membrane was incubated with horse
radish peroxidase (HRP)-conjugated goat anti-rabbit IgG or goat anti-mouse IgG (diluted at 1:5000) for 2 h at room temperature. Finally, the immunoreactive bands were detected using the ECL methods and quantified the protein bands by densitometry with BIORAD ChemiDoc™ XRS+ software (Bio Rad, Hercules, CA, USA). The protein expressions were obtained by analyzing the density ratio of target proteins to GAPDH expression.

2.6. Determination of the Levels of Cytokines and Tryptase in Larval Zebrafish

Determination of cytokines levels on zebrafish larvae was performed as previously described [41,42], using enzyme linked immunosorbent assay (ELISA) kits. Zebrafish larvae (20 per experimental group of each experiment) were rinsed with 1X PBS, homogenized in 1 mL of 1X PBS, in 1.5-mL EP tube and stored overnight at −20 °C. After two freeze-thaw cycles were performed to break the cell membranes, the homogenates were centrifuged at 906 × g per 10 min at 4 °C (Micro centrifugate D3024) to obtain the supernatant for further use. The levels of cytokines were measured using ELISA kits according to the manufacturer’s instruction, TNF-α (RayBio ELZ-TNFa-1, RayBiotech Life, Peachtree Corners, GA, USA), IL-1β (MyBioSource MBS700230, San Diego, CA: USA), IL-6 (RayBio ELZ-IL6-1, RayBiotech Life, Peachtree Corners, GA, USA), and IL-8 (Antibodies A82431, Cambridge, UK). Briefly, standards and samples are pipetted into the wells and cytokines present in a sample are bound to the wells by the immobilized antibody. The wells are washed and biotinylated and anti-TNF-α, IL-1β, IL-6, and IL-8 antibody respectively are added. After washing away unbound biotinylated antibody, HRP-conjugated streptavidin is pipetted to the wells. The wells are again washed, a TMB substrate solution is added to the wells, and color develops in proportion to the amount of cytokines bound. The Stop Solution changes the color from blue to yellow, and the intensity of the color is measured at 450 nm with spectrophotometer (Jenway). The tryptase levels in the fish maintained water were determined and calculated according to a previous study [43]. BAPNA was bought from Aladdin (catalog no. 911-77-3, Shanghai, China) and used as a substrate for tryptase quantification. Zebrafish were maintained in the reaction solution throughout the experiments. In all experiments, microplate wells with zebrafish larvae without BAPNA were used as background control, and microplate wells with BAPNA without zebrafish served as a negative control. To protect compounds from light-induced decomposition, experiments were carried out at a constant temperature (28 °C) in the dark. After incubation at 28 °C for 1 h, zebrafish tryptase activity was quantitatively measured at 405 nm using a multifunction microplate reader (MikroWin 2000; Berthold, Stuttgart Germany). Net optical density (OD) was calculated by subtracting the OD of background control wells.

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\text{Tryptase-increased fold} = \frac{(\text{OD [chemical treated]} - \text{OD [untreated control]})}{\text{OD [untreated control]}}.
\]

2.7. Yolk Sac Areas and Body Axis Curvature Percentage

To measure yolk sac areas of lateral view, images of each embryo were taken at the same magnification; the outline of the yolk sac, respectively, was traced, and the area within each tracing was determined by Image J program (Version 1.8.0, National Institutes of Health, Bethesda, MD, USA). Moreover, the number in percentage of larvae with severe dorsal axis curvature in three independent experiments was calculated for the body axis malformation quantification.

2.8. Statistical Evaluation

The results were analysed by two-way/one-way ANOVA followed by a Tukey post-hoc test for multiple comparisons. The data were tested for normal distribution with the Kolmogorov–Smirnov test (p < 0.05) and they were represented as mean ± standard deviation (SD) (alpha value of 0.05). Statistical analysis was performed using Graphpad Prism 8. Bars with the same letter indicate a non-significant difference of the mean, while different letters indicate a statistically significant difference of the mean.
3. Results

3.1. Morphology

In order to identify the suitable concentration and time points used in the following experiments, PEAOXA ranging from 0.1, 1, 3, 10, to 30 mg/L concentration was applied to observe morphology of larvae. As presented in Figure 1, PEAOXA concentration 0.1, 1, 3, and 10 did not alter the zebrafish morphology until 96 hpf (hours post fertilization) compared to control group (CTRL) (Figure 1). PEAOXA 30 mg/L groups induced abnormalities, like body axis curvature ($p$ value 0.0001) and large yolk sac ($p$ value 0.0002) in zebrafish embryos (Figure 1).

![Figure 1. The morphological abnormalities in zebrafish caused by PEAOXA different concentration exposure (A). Results for yolk sac area (B) and number of embryos with body axis curvature (C). Images were taken from the lateral view under a dissecting microscope (magnification 25). Scale bar, 500 $\mu$m. Values = means ± SD of three independent experiments Bars of group labelled with different letters (a, b) denote significant differences ($p < 0.05$).](image)

3.2. Survival, Heart, and Hatching Rate

The effect of PEAOXA on the embryo development was observed up to 96 hpf. Embryo development in the control was normal: hatching began after 48, about at 72 hpf while mortality was little compared to CTRL (Figure 2A). Embryos exposed to high concentration of PEAOXA (30 mg/L) showed a delayed of the hatching rate ($p$ value 0.0093 and 72 hpf; $p$ value 0.0001 96 hpf), while no differences were observed for embryos exposed to low concentrations (0.1, 1, 3, and 10) compared to CTRL group. Heart rates were recorded to determine the effect of PEAOXA on cardiac function. In the PEAOXA 0.1, 1, 3, and 10 treated group embryos, as well as in the CTRL, heart rate did not show differences from 48 to 96 hpf. However, from 48 ($p$ value 0.0001) to 96 hpf ($p$ value 0.0001), significant bradycardia was observed in embryos treated with PEAOXA 30 mg/L compared with the controls (Figure 2B). Lethality was caused right from 48 to 96 hpf at 30 mg/L ($p$ value 0.0001). However, for the lower concentrations than 30 mg/L, no mortality was observed for the duration of the experiment.
3.2. Survival, Heart, and Hatching Rate

The effect of PEAOXA on the survival (A), heart (B), and hatching (C) rate of embryos was determined at the designate time. Values = means ± SD of three independent experiments. Bars of group labelled with different letters (a, b, c) denote significant differences (p < 0.05).

3.3. Effects of PEAOXA on the Levels of Cytokines and Tryptase Release on LPS-Induced Inflammation

The levels of IL-8, IL-1β, IL-6, and TNF-α were also detected by the ELISA kits in zebrafish larvae after LPS exposure. TNF-α expression was increased in larval zebrafish LPS treated compared to the CTRL (p value 0.0001). In addition, LPS significantly increased the IL-1β levels, as well as IL-6 and 8, compared to CTRL group (Figure 3) (p value 0.0001; 0.0001, and 0.0002). PEAOXA exposure was able to reduce the levels of all these cytokines, after LPS exposure. Indeed, a significant reduction was seen in TNF-α, IL-1β, IL-6, and IL-8 levels in zebrafish larvae (Figure 3) (p value 0.0096; 0.0075; p value 0.0064; p value 0.0011). Furthermore, the levels of tryptase, a marker of mast cells degranulation, was increased significantly in the water of larvae treated with LPS (p value 0.0001). No significant increase of the tryptase level was observed in larval zebrafish LPS-stimulated when exposed with PEAOXA at 10 mg/L (Figure 4) (p value 0.0001), while the dosage 3 mg/L of PEAOXA was ineffective to reduce the release of tryptase (Figure 3).

Figure 2. The hatching (A), heart (B), and survival (C) rate of embryos was determined at the designate time. Values = means ± SD of three independent experiments. Bars of group labelled with different letters (a, b) denote significant differences (p < 0.05).

Figure 3. Effects of PEAOXA on LPS exposure on TNF-α (A), IL-6 (B), IL-8 (C), and IL-1β (D) protein levels and tryptase (E) released in larval zebrafish. Values = means ± SD of three independent experiments of three independent experiment data. Bars of group labelled with different letters (a, b, c) denote significant differences (p < 0.05).
3.4. Effect of PEAOXA on the mRNA Expression of LPS-Induced Cytokines

A total of four inflammatory mediators were detected (IL-1β, IL-6, TNF-α, and IL-8) and nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, alpha (IκB-α) pathways. As shown in Figure 4, the mRNA level of the cytokines TNF-α, IL-1β, IL-6, and IL-8, was significantly increased in LPS-treated larvae (p value 0.0070; 0.0072; p value 0.0012; p value 0.0061), compared withCTRL. In the treatment groups, PEAOXA regulated the expression of cytokines toward a normal level at the concentration of 10 mg/L (TNF-α p value 0.0070; IL-1β p 0.0072; IL-6 p value 0.0012; IL-8 p value 0.0061), in contrast to the PEAOXA 3 mg/L concentration which showed no significative reduction of cytokines.

3.5. Effect of PEAOXA on the mRNA Expression and Protein of LPS-Induced NF-κB Pathway

In addition, we also assessed gene expression and protein levels of NF-κB p65, an important transcription factor involved in the inflammatory pathway and its selective inhibitor IκB-α. Our results showed NF-κB increase and a decrease of IκB-α expression after LPS-stimulated on zebrafish larvae (p value 0.0001). PEAOXA exposure was able to reduce the NF-κB increase (p value 0.0001 and 0.0014), both in terms of mRNA and protein levels, and it was able to increase the mRNA and protein of its selective inhibitor IκB-α (p value 0.0001 and 0.0018), compared to the only LPS induced group (Figure 5). No effect was shown after exposure with PEAOXA 3 mg/L compared to LPS only treated group (Figure 5 and see uncropped blot in Supplementary Materials Figure S1).

3.6. Effects of PEAOXA on LPS-Induced Apoptotic Pathway

The mRNA expression levels of apoptosis-related genes (caspase-3, and bax) increased after LPS exposure compared to CTRL (Figure 6) (p value 0.0001 and 0.0005), while the mRNA expression level of bcl-2 was downregulated (p value 0.0001). PEAOXA at the concentration of 10 mg/L was able to reduce the expression of apoptotic related genes (p value 0.0001 and 0.0004) and at the same time increase the bcl-2 mRNA expression after LPS-induced inflammation (Figure 6) (p value 0.0001). Protein expression of caspase-3 and Bax was significantly induced by LPS exposure compared with the CTRL (Figure 6) (p value 0.0001). Contrarily, anti-apoptotic protein bcl-2 expression was significantly inhibited in LPS group larvae (p value 0.0001). Moreover, LPS-induced an increase of Bax/bcl-2 ratio in larvae compared to CTRL group (Figure 6) (p value 0.0001). PEAOXA at the concentration of 10 mg/L was able to reduce both the caspase-3 expression induced by LPS in

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**Figure 4.** Effects of PEAOXA on LPS exposure on the mRNA levels of TNF-α (A), IL-6 (B), IL-8 (C), and IL-1β (D) in larval zebrafish. Results are expressed as mRNA expression normalized to β-actin and after fold change to control. Values = means ± SD of three independent experiment data. Bars of group labelled with different letters (a, b, c) denote significant differences (p < 0.05).
larvae as well as the Bax/bcl-2 ratio ($p$ value 0.0049 and 0.0131). No effect was seen for the concentration of 3 mg/L of PEAOXA on both protein and mRNA levels compared to LPS group (Figure 6 and see uncropped blot in Supplementary Materials Figure S1).

Figure 5. Effects of PEAOXA on LPS exposure on the mRNA levels of IκB-α (A) and NF-κB (B) and protein levels (C,D) in larval zebrafish. Western blot analysis. (E) Results are expressed as mRNA expression normalized to β-actin (A,B), protein expression levels normalized to GAPDH (C,D), and after fold change to control Values = means ± SD of three independent experiment data. Bars of group labelled with different letters (a, b, c) denote significant differences ($p < 0.05$).

Figure 6. Effects of PEAOXA on mRNA and protein levels of LPS-induced apoptotic pathway (caspase3, bax, and bcl-2) on larval zebrafish. The mRNA levels of LPS-induced apoptotic pathway (caspase3, bax, and bcl-2) on larval zebrafish (A). Western blot analysis (B). Western blot analysis was performed with antibodies against caspase-3, bcl-2, Bax, and GAPDH. Each figure corresponds to a representative replicate from three experiments for Caspase-3, bcl-2, Bax, and Bax/bcl-2 ratio (C–F). Results are expressed as mRNA expression normalized to β-actin (A), protein expression levels normalized to GAPDH (C,D), and after fold change to control Values = means ± SD of three independent experiment data. Bars of group labelled with different letters (a, b, c) denote significant differences ($p < 0.05$).
4. Discussion

Zebrafish, as one of the most significant vertebrate animal models, was used for a variety of toxicological studies due to its particular advantages over cell models and other animal models [44]. Gram-negative bacteria are the main pathogenic bacteria in the majority of fish, and LPS is an important pathogenic component of these bacteria. In this study, we used zebrafish embryo and larva (Danio rerio) as a fish model to study the impact of LPS on fish inflammatory pathway and apoptosis and the protective role of PEAOXA.

Since to date PEAOXA has never been used on zebrafish larvae, different concentrations were assessed to allow to find the toxics concentration and know which one to test for anti-inflammatory efficacy in the trials. There were no harmful effects at any of the dosages employed on embryonic development except for the highest concentration analyzed (i.e., 30 mg/L), which showed malformation like spinal curvature. For the low concentrations of PEAOXA, no morphological alterations, such as edema formations of pericardium or in the yolk sac, or structural malformations were observed. Furthermore, the exposure of low concentrations of PEAOXA did not show increased mortality or bradycardia, all parameters normally used in the study of toxicity on the development of zebrafish. PEAOXA concentration of 30 mg/L, the highest used, showed signs of mortality as early at 72 hpf with a survival rate decrease of about 50% at 96 hpf. Furthermore, the concentration of 30 mg/L also affected the hatching rate as early as 72 hpf up to 96 hpf and also slowed the heart rate to 96 hpf. In contrast, the 10 mg/L concentration did not show any of these toxic effects on embryonic development, allowing us to choose this concentration as the highest to be used in the following experiments for anti-inflammatory activity. LPS exposure induced aberrant cell death caused by the overproduction of pro-inflammatory cytokines was detected during LPS-stimulated inflammatory reactions in zebrafish larvae [45]. LPS exposure caused toxicity in zebrafish embryos, which is an increment of inflammatory pathway [46]. During the process of acute inflammation, among the cells of the immune system, mast cell plays an important role. In fact, once activated, the mast cell releases their granules content, thus contributing to the progression of the inflammatory process. Zebrafish mast cells have mammalian-like functions and structure and are involved in the immune response and allergic reaction [47]. Generally, mast cells degranulate upon binding to allergens, releasing into the microenvironment the contents of these granules, including some mediators such as histamine and tryptase [48]. Most importantly, in zebrafish embryos, at 28 and 48 hpf, mast cells originate from erythromyelic progenitor cells [49], and mast cells matured and had full function before 7 dpf in larval zebrafish [47,50]. In the present study, we observed that LPS exposure increased the release of tryptase into the water, indicating that LPS exposure increased the anaphylactoid reaction in larval zebrafish.

PEAOXA, the oxazoline of PEA, is well known to present anti-inflammatory and analgesic properties [51]. PEAOXA has been shown in several studies to act at the anti-inflammatory level by reducing mast cell activation [25,51,52]. This feature derives from its precursor, PEA, which has been shown to act by inhibiting mast cell degranulation, thus decreases the content of granules such as tryptase. In our study, exposure to PEAOXA showed a decrease in LPS-induced tryptase release, suggesting a protective action against mast cell degranulation in larvae, consistent with previous studies in mouse models [52].

Among the various components released by the mast cell, there are not only proteases, like chymase and tryptase, and the histamine, but also mediators like cytokines [49]. Recently, Watzke et al., demonstrated that in zebrafish embryo LPS could induce the transcriptional expression of innate immune-related genes [53]. Previous studies showed a protective action of PEAOXA on reducing the inflammatory process in mouse models of inflammation [25]. In the present study we observed an increase of cytokines levels after LPS exposure. Furthermore, it was seen an increase of pro-inflammatory cytokines IL-1β, IL-6, IL-8, and TNF-α, both in terms of protein levels then with an augmentation of gene expression. In line with previous studies, PEAOXA exposure effectively downregulated the cytokines overproduction, both in terms of mRNA levels and then in protein expression in zebrafish larvae [25]. The binding between LPS and TLR4 leads upstream to an
activation of the transcription factor NF-κB, involved in the regulation of the production of pro-inflammatory mediators, such as IL-1β and IL-6 or TNF-α [54]. PEAOXA exposure attenuated TNF-α, IL-8, IL-6, and IL-1β levels increased; these are cytokines normally involved in the IκBα/NF-κB signaling pathway stimulated by LPS. NF-κB, after translocation into the nucleus, stimulates cytokine production during inflammation [55]. Our results showed that PEAOXA inhibits LPS-induced inflammation, and this activity is linked to a decrease in NF-κB mRNA expression. Moreover, PEAOXA increase mRNA levels of IκBα, the cytoplasmic inhibitor of NF-κB translocation, into the LPS-stimulated nucleus in zebrafish models. This suggests that the anti-inflammatory action of PEAOXA involves the production of proinflammatory cytokines and transcription factors that bind to them.

The triggering of a massive inflammatory response can lead to the activation of the apoptotic process. Numerous studies have shown how LPS-induced inflammation caused not only an increase of oxidative stress but also and above an increase of all of the markers associated with apoptosis [29,56]. In this study, PEAOXA showed anti-inflammatory action not only by limiting cytokine release but also decreasing the expression of LPS-induced apoptosis. It was previously shown that PEAOXA treatment was able to decrease the apoptotic process in a model of neuropathic pain, by decreasing the expression of apoptotic protein Bax and increasing the expression of the anti-apoptotic bcl-2 [51]. The induction of apoptosis in zebrafish larvae was detected by both mRNA and protein levels of apoptosis-related protein caspase 3, bax, and bcl-2. Caspases are a large family of proteinases that play crucial roles in the process of apoptosis and are considered markers in zebrafish embryos [57]. Our data showed that the exposure of PEAOXA at 10 mg/L decreased the expression of the apoptosis-inducing target genes casp-3 and bax and the expression of the anti-apoptotic factor bcl-2. Thus, PEAOXA was able to contrast LPS-induced apoptosis in terms of mRNA expression, as well as protein levels of molecular apoptotic marker, confirming what has been shown in previous studies on mice models.

5. Conclusions

In conclusion, our results showed that the exposure of PEAOXA at a concentration of 10 mg/L was able to contrast LPS-induced inflammation. In fact, PEAOXA was able to reduce the cytokines expression and release after LPS exposure. Moreover, PEAOXA inhibited the excessive activation of the apoptotic process by reducing the levels of caspase-3 and bax and increased expression of bcl-2. In conclusion, PEAOXA at 10 mg/L was able to counteract the acute LPS-induced inflammatory process for the first time in a model of zebrafish embryo and larvae.

Supplementary Materials: The following are available online at https://www.mdpi.com/article/10.3390/life12010128/s1, Figure S1: uncropped western blot.

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Conflicts of Interest: Salvatore Cuzzocrea is a coinventor on patent WO2013121449 A8 (Epitech Group Srl), which deals with methods and compositions for the modulation of amidases capable of hydrolyzing N-acylthanolamines employable in the treatment of inflammatory diseases. This invention is unrelated to the present study. Moreover, Cuzzocrea is also with Epitech Group, a coinventor on the patents EP 2821083, MI2014 A001495, and 102015000067344, which are unrelated to the study. The remaining authors report no conflicts of interest.
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