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Original article

# The potential effects of *Indigofera coerulea* extract on THP-1 human cell line



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# ABSTRACT

The purpose of this study was to detect the immunomodulatory activity of Indigofera coerulea on the cytokine's expression in the THP-1 human cell line. THP-1 cell lines were differentiated into real macrophages. Cell's viability was assessed by MTT assay. The qRT-PCR assay was used to determine the effects of the I. coerulea extract on mRNA expression levels in THP-1 cells. The effects of extracts on proteins production were analyzed by Western blot assay and protein array. The ELISA was used to determine the effect of the plant extract on cytokine expression. Finally, apoptosis, phagocytosis, and cell migration assays were performed to investigate the effects of the extract on macrophage functions. The obtained results illustrated that I. coerulea extract possesses anti-oxidant and anti-inflammatory activities. In addition, current study reported that I. coerulea extract significantly reduced the proliferation of THP-1 cells at all-time points. Moreover, I. coerulea extract showed significant immunomodulatory activity compared with control macrophages by influencing tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ), IL-1 $\beta$ , IL-6, CCL22, CXCL10/ IP-10, CXCL8/IL-8, ERK5, BAX, BcL2, Cyclin D1, ERK1, P-IκB-α, P-NF-κB, and P-p38 proteins and the signaling pathways of NF-KB, p38 MAPK, ERK1/2, and IL-6/JAK/STAT3. In conclusion, this study is the first to underline the anti-proliferative, anti-inflammatory, anti-phagocytic, anti-apoptotic, and antimigratory properties of the studied plant extract in human monocytic THP-1 cells. © 2021 The Author(s). Published by Elsevier B.V. on behalf of King Saud University. This is an open access

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*Abbreviations:* CCL2/MCP-1, C–C Motif Chemokine Ligand 2/ Monocyte Chemoattractant Protein1; CCl4, Carbon Tetrachloride; CCL5, C–C Motif Chemokine Ligand 5; CCL22, C–C Motif Chemokine 22; CXCL10/ IP-10, Interferon Γ-Induced Protein 10; CX3CL1, C-X3-C Motif Chemokine Ligand 1; ERK1, Extracellular Signal-Regulated Kinases 1&2; GM-CSF, Granulocyte-Macrophage Colony-Stimulating Factor; LPS, Lipopolysaccharides; MAP, Mitogen-Activated Protein; M–CSF, Macrophage Colony-Stimulating Factor; NF-κB, Nuclear Factor Kappa-Light-Chain-Enhancer of Activated B Cells (NF-κB); PMA, Phorbol-12-Myrstate-13-Acetate; p-p38, Phosphorylated P38; P-I kB-α, Phospho-Nuclear Factor of Kappa Light Polypeptide Gene Enhancer In B-Cells Inhibitor Alpha; THP1, Human Monocytic Leukemia Cell Line.

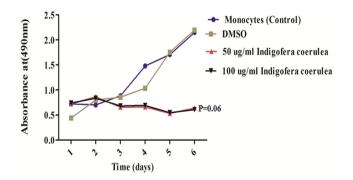
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#### 1. Introduction

Macrophages are immune cells that an essential in tissue development and stimulate cytokines (Carlos, 2015). However, the major functions of macrophages are; eliciting adaptive immunity, antigen presentation, and stimulation lymphocytes, maintain tissue homeostasis (Wynn et al., 2013; Linehan and Fitzgerald, 2015). Phagocytic receptors on pathogens are recognized by specific receptors. These pathogenic components, like lipopolysaccharides (LPS), do not have high mutation rates due to their intrinsic biological role in the invading organism. Phagocytosis occurs relying on the recognition mechanisms. Cellular receptors recognize bacterial surface components and integrins (Aderem and Underhill 1999). During phagocytose pathogens, an inflammatory response occurs and anti-inflammatory response is initiated by macrophages (Stuart and Ezekowitz, 2005: Chung et al., 2007: Al-Oahtani et al., 2021). Exposure to bacterial products, such as LPS, results in macrophages with altered phenotypic and triggered its ability to produce IL-12 (Denkers 2007). However, the deactivated phenotype can be enhanced by exposure to a number of anti-inflammatory cytokines, apoptotic bodies (Denkers 2007).

Human THP-1 cells possess specialized regulatory proteins that activate the process of inflammation upon (LPS) stimulation (Sullivan et al. 2018). Research studies on the signaling pathways of inflammation have facilitated the exploitation of targets for drugs development and have explored the role of medicinal plant extracts in diverse cell-based assays (Bremner et al., 2009; Siriwatanametanon et al., 2010). Medicinal plants have been reported to be adjuvants to traditional remedy for promoting the immune response (Varma et al., 2016; Zhang et al., 2018). A previous study on the liver protection used F. parviflora extract against nimesulide-induced apoptosis in vitro. In addition, it was illustrated that *C. epigaeu* has a high antioxidants levels (Jevaseelan et al. 2014). Also, many plant extracts can modulate macrophage functions such as production of inflammatory/anti-inflammatory components (Albrahim et al., 2020). They exert this effect through intracellular signaling pathways such as; mitogen-activated protein (MAP) kinase, nuclear factor kappa-B (NF-kB) protein. Thus, many plant extracts from Saudi Arabia have been used to modulate immune responses in different experimental models including innate and adaptive immune responses. Strong evidence has revealed that some plants have the ability to improve the function of macrophages (Albrahim et al., 2020).

*I. caerulea* has antimicrobial and antioxidant properties (Guruvaiah et al., 2012; Ponmari et al., 2014) and antigrowth activities (Natarajan et al., 2010). Furthermore, it was reported that *I.* 



**Fig. 1.** Effects of *I. coerulea* on proliferation of monocyte cells after incubation with (50 and 100  $\mu$ g/mL). Each point is the mean of 3 replicates; both 50 and 100  $\mu$ g/mL I. *coerulea* reduced cell proliferation.

*coerulea* plant possesses anti-hepatitis B virus (HBV) infection property (Arbab et al., 2017). In addition, this plant prevents the release of pro-inflammatory cytokines IL-1β and TNF-α (Ponmari et al., 2014). The hepatoprotective properties of *I. coerulea* phytochemicals and their inhibitory activity against carbon tetrachloride (CCl<sub>4</sub>)-triggered hepatic injury have also been demonstrated in rat models. *I. coerulea* extracts not only attenuate the NF-κB pathway, but also block the release of IL-1β and TNF-α (Lopes et al., 2011; Al-Shaebi et al., 2017; Chen et al., 2018). The role of *I. coerulea* plant extract on human peripheral blood monocytes or THP-1 cell line still unknown, thus current study aim to investigate the antiproliferative, anti-inflammatory, anti-phagocytic, anti-apoptotic, and anti-migratory properties of the *I. coerulea* extract in human monocytic THP-1 cells.

# 2. Materials and methods

#### 2.1. Preparation of plant extracts

The plant was collected from southern part of Saudi Arabia and classified by the taxonomist at King Saud University. Briefly, dried plant was ground to powder and extracted with 80% ethanol followed by filtering and was concentrated using a rotary evaporator (Buchi Labortechnik AG, Flawil, Switzerland) under low pressure at 4 °C. Extract was dissolved in (DMSO; Sigma-Aldrich, Merck KGaA), and the stock (100 mg/ml) was stored at - 20 °C until subsequent use (Arbab et al., 2017).

# 2.2. Cell culture and differentiation of THP-1 cell line

THP-1 cells were obtained from the acute monocytic leukemia (AML) patient. Cells were cultured in (RPMI-1640) medium accompanied with 10% complement-inactivated fetal bovine serum (FBS), 100 IU/mL penicillin, 100  $\mu$ g/mL streptomycin, and 3.7 g sodium bicarbonate/L (Sigma-Aldrich, St. Louis, MO, USA). In order to activate THP-1 cells to differentiate, cells were cultured in 96-multiwell culture plates, then 100 ng/mL (162 nM) of phorbol 12-myristate 13-acetate was added, as suggested in previous research (Starr et al. 2018). Cells were washed with RPMI-1640 serum-free medium prior to each experiment to remove undifferentiated cells. For activation of macrophage-like cells, cells were then treated with LPS (100 ng/mL) from *E. coli* O55:B5 (L2880 Sigma) for 24 h.

# 2.3. Analysis of cell viability (MTT assay)

Cell viability was measured by MTT assay following manufacturer's instructions (MTT Assay Kit ab211091, Abcam, city, country). THP-1 cells were exposure to different concentrations (50 or 100  $\mu$ g/mL) of crude extract of *I. caerulea* at different time points. Briefly, media was replaced with 50  $\mu$ L/well serum-free media and 50  $\mu$ L/well MTT reagents. Cells were cultured in 96-well plates and incubated for three hours. Optical density (OD) was measured at 590 nm using a microplate reader (SpectraMax<sup>®</sup> MiniMax<sup>™</sup>300 Imaging cytometer). All experiments were performed in triplicate.

#### 2.4. Gene expression by RT-PCR

THP-1 cells were cultured in a 6-well plate and were incubated with 100 ng/mL LPS only, 100 ng/mL LPS with 100  $\mu$ g/mL plant extracts, and 100  $\mu$ g/mL of plant extract only for four hours. RNA was isolated using QlAamp<sup>®</sup> RNA Blood Mini kit (Qiagen, Hilden, Germany) according to the manufacturer's protocol. A total of 10  $\mu$ g of RNA was used for cDNA synthesis using SuperMix (Biotool, Houston, TX, USA). qRT-PCR was performed using targetspecific primers through StepOne RT-PCR system (Applied Biosys-



Sample ID: Cells + 100ng/ml of Nocodazole + 100ng/ml of LPS. Sample ID: Cells + 100ng/ml of Nocodazole + DMSO

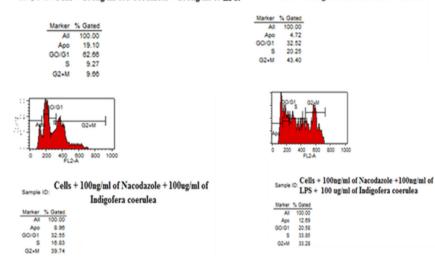
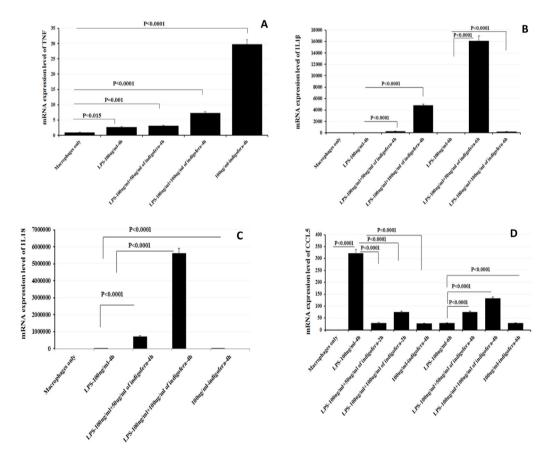
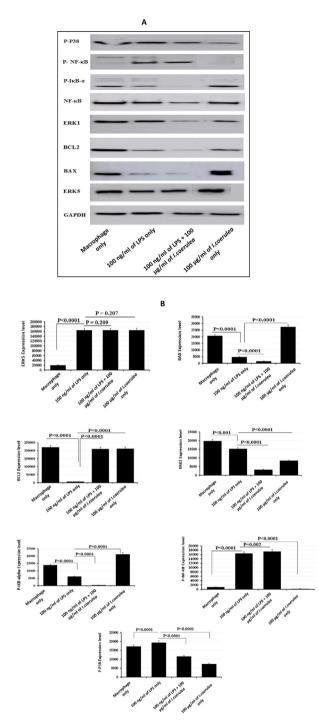


Fig. 2. Cell synchronization assay was showed treated human monocytic THP-1 cells with *I. coerulea plant extract* induced G0/G1 and G2/M arrest in human monocytic THP-1 cells (n = 3).



**Fig. 3.** The effects of *I. coerulea* plant extract on mRNA expression levels of TNF (A), IL-1β (B), IL-18 (C), and CCL5 (D) in THP-1 cells. Result shows significant increase in the mRNA expression of TNF, IL-1β and IL-18 were observed in THP-1 cells compared to their control macrophages counterparts. In contrast to above, the only mRNA expression statistically decreased was that of CCL5 on all forms of treatment compared to control macrophages.



**Fig. 4.** (A) Expression of ERK5, BAX, BCL2, ERK1, NF-KB, P-IKB- $\alpha$ , P-NF-KB, and P-P38 in THP-1 cells treatment with lipopolysaccharide (LPS) and *I. coerulea*. Proteins were separated on 12% SDS-PAGE. Lane 1: THP-1 cells, Lane 2: Cells treated with 100 ng/mL LPS + 100 µg/mL *I. coerulea*, Lane 4: Cells treated with 100 µg/mL *I. coerulea*. Significantly different at *p*-value < 0.05. (B) Expression of ERK5, BAX, BCL2, ERK1, NF-KB, P-IKB- $\alpha$ , P-NF-KB, and P-P38 in THP-1 cells treatment with LPS and *I. coerulea*.

tems, Foster City, CA, USA). Relative gene expression data were analyzed using the  $\Delta$ Ct method. The data are presented as mean ± standard deviation (SD).

# 2.5. Western blot analysis

The differentiated cells were treated for four hours with either 100 ng/mL LPS only, 100 ng/mL LPS with 100  $\mu$ g/mL plant extract,

or 100  $\mu$ g/mL plant extracts only. Western blot was used to assess the expressions of targeted proteins in treated cells. Cells were lysed in RIPA buffer, followed by protein separation on 12% SDS-PAGE. Then proteins were blotted on PVDF membranes. The membranes were then incubated specific primary antibodies overnight at 4 °C, washed, and incubated with HRP-conjugated secondary antibodies. The proteins were detected using Super Signal West Pico16 Chemiluminescent substrate (Thermo Fisher Scientific, Waltham, MA, USA), and visualized on a GE Amersham Imager 600, and were quantified using Image J software (National Institutes of Health, Bethesda, MD, USA).

# 2.6. Protein array assay

Protein array was performed using the Proteome Profiler<sup>™</sup> Arrav-Human Cytokine Arrav Kit (R&D Systems). Differentiated THP-1 cells were incubated for four hours with the following combinations: 100 ng/mL LPS only, 100 ng/mL LPS with 100 µg/mL I. caerulea, and I. caerulea only. Cytokines were collected and the membranes were placed into the wells and incubated for one hour. The samples were prepared by mixing 1 mL of each sample with 0.5 mL blocking buffer, 15 µL of reconstituted cytokine array detection antibody cocktail, and then incubated for one hour. Then the prepared sample/antibody mix was added and incubated overnight at 4 °C. Each membrane was then placed individually in 20 mL  $1 \times$  wash buffer. We added 2 mL of diluted streptavidin-HRP to each well. The membranes were then returned to the well and incubated at room temperature for 30 min. Finally, 1 mL of the prepared chemiluminescent reagent mix was added to each membrane. The cytokines were detected using an Amersham Imager 600 (GE Healthcare).

# 2.7. Phagocytosis assay

The phagocytosis activity was carried out on resting M1 and M2 polarized macrophage-like cells using Phagocytosis Assay Zymosan Substrate Kit (Abcam, Cambridge, U.K.). 96 well plates were filled with THP1 cells per well, and incubated with 10  $\mu$ g/ml PMA for 48 h. The, media was then changed, and the cells were rested for 24 h to differentiate. Activated THP-1 cells were incubated for 75 min with a Zymosan suspension. To block external particles, 100  $\mu$ L of fixation solution was added to each well. Afterwards, 100  $\mu$ L of 1X blocking reagent was added to each well, and the plate was incubated for 60 min. Then, cells were washed with a RPMI serum-free medium followed by the addition of 100  $\mu$ L fixation solution to each well. The absorbance was measured using a SpectraMax<sup>®</sup> MiniMax<sup>™</sup>300 Imaging cytometer (Molecular Devices, CA, USA) at OD 405 nm.

# 2.8. Apoptosis assay

To determine the activation of apoptotic pathways in treated cells, caspase-3 activation assay (Sigma-Aldrich, St. Louis, MO, USA) was performed according to the manufacturer's protocol. Briefly, cells were cultured in 6 well plates, and incubated for four hours with the treatments, washed with PBS, and incubated with 5  $\mu$ L Annexin V-FITC conjugated with rabbit anti-human active caspase-3 for 20 min. After that, cells were then washed, incubated with 10  $\mu$ L of propidium iodide (PI) for 10 min. The fluorescence of the cells was determined using flow cytometry (Becton Dickinson, Franklin Lakes, NJ, USA).

# 2.9. Enzyme-linked immunosorbent assay (ELISA)

The ELISA was performed to assess cytokines into the cell culture media using a DuoSet<sup>®</sup> ELISA kit according to the

manufacturer's protocol. In brief, PMA-treated cells were seeded in a 6-well plate and incubated for six hours in different combinations: 100 ng/mL LPS only; 100 ng/mL LPS + 100  $\mu$ g/mL plants extract; 100  $\mu$ g/mL plant extracts only. 100  $\mu$ L of cell culture supernatant was added to each well, and then was incubated for two hours. Cells were treated with primary antibodies, incubated for two hours. 100  $\mu$ L of streptavidin-HRP-conjugate solution was added to wells and incubated for 20 min. 100  $\mu$ L of chromogenic substrate was added to each well and the plates were incubated for an additional 30 min. 100  $\mu$ L of stop solution was added to terminate the reaction. Absorbance was read at 450 nm using a microplate reader and the concentration of each protein was measured using Gen5 software (BioTek Instruments).

# 2.10. Flow cytometry

To assess the response of THP-1 cells to whether macrophage differentiation factor (PMA) was able to trigger CD-14 expression, a biomarker of monocyte lineage, THP-1 cells were scattered into 96-well culture plates, as explained above in the cell culture and differentiation of the human monocytic THP-1 cell line. Using Try-pLE (Invitrogen), cells were removed from the wells at 37 °C after incubating for 10 min. Cells were then placed on ice and all the steps that followed were completed at 4 °C. FACS buffer containing 0.1% azide, 2% fetal calf serum in PBS, and 11  $\mu$ g/mL of IgG (Jackson) was used to pellet and re-suspend cells. Then, cells were stained with eFluor 450 anti-human CD-14 antibodies

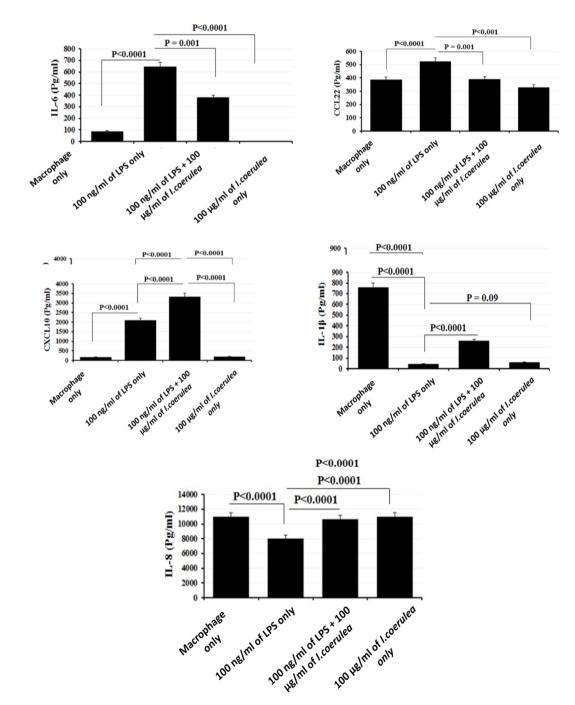


Fig. 5. ELISA results for cell culture supernatant of macrophages after stimulation with 100 ng/mL LPS and 100 µg/mL *I. coerulea* plant extract. Supernatant of culture was harvested 6 h after stimulation. Error bars indicate SEM. All samples were processed in triplicate.

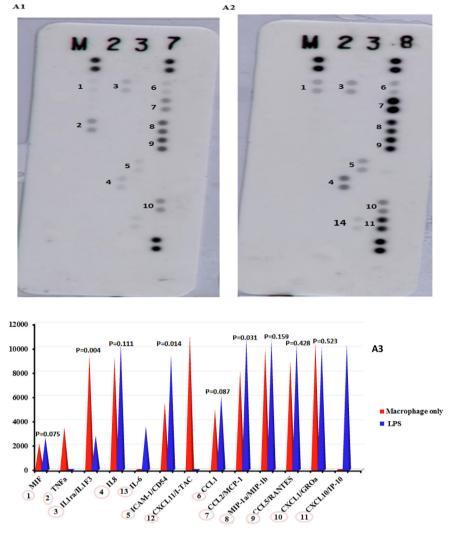


Fig. 6. Protein array expression: (A1) macrophages; (A2) cells treated with 100 ng/mL LPS; (A3) histograms comparing the protein array results from macrophages and cells treated with LPS.

(eBioscience, CD14 eFlour 450) and analyzed through flow cytometry and FlowJo software (TreeStar, Inc., vX.0.7 and v9.5.2).

# 2.11. Statistical analysis

Obtained data are expressed as mean  $\pm$  standard error of mean. To compare the multiple cell groups, the ANOVA test was used followed by Bonferroni's test. A *p*-value was considered significant if<0.05.

# 3. Results

# 3.1. Effects of I. Caerulea plant extract on proliferation of THP-1 cells

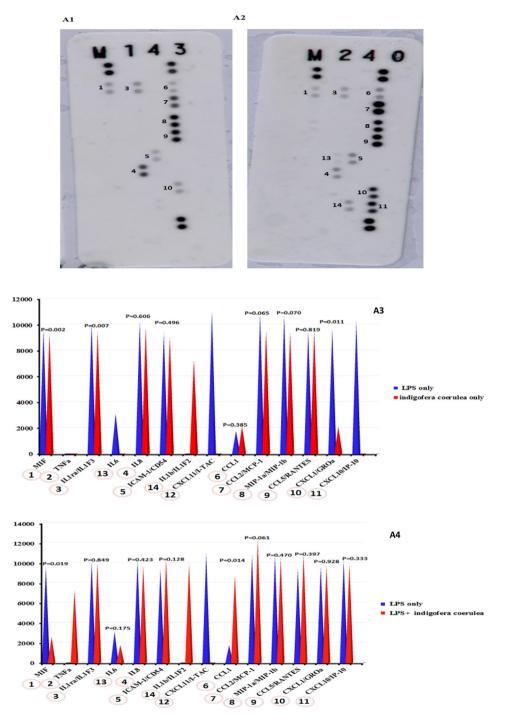
The effect of *I. caerulea* extract on the proliferation of treated THP-1 cells was detected by MTT assay. Cells were treated with 50 and 100  $\mu$ g/mL of *I. caerulea* extract for 1–6 days. THP-1 cells were also treated with vehicle control (DMSO). Fig. 1 shows the percentage of absorbance at 490 nm that induced by DMSO, 50  $\mu$ g/mL, and 100  $\mu$ g/mL plant extract relative to the control at 0 days. THP-1 cells treated with both 50 and 100  $\mu$ g/mL plant extract demonstrated a sustained dampened proliferative response in contrast to DMSO at all-time intervals, indicating proliferation arrest on contact with *I. coerulea* extract.

#### 3.2. Extract of I. Coerulea induces G0/G1 and G2/M arrest

A cell synchronization experiment was performed to test the mechanistic behavior of THP-1 cells upon treatment with the plant extract. The results of cell cycle regulation after treatment with *I. coerulea* extract are shown in Fig. 2. A combination of monocyte cells with 100 ng/mL nocodazole, 100 ng/mL LPS, ad 100  $\mu$ g/mL *I. coerulea* extract showed significantly higher G0/G1. Slight apoptotic activity was also witnessed with all forms of treatment of THP-1 cells. Overall, in agreement with the cell proliferation annexation seen in Fig. 9, cell synchronization analysis through flow cytometry showed that the plant extract caused the THP-1 cells to arrest at the G0/G1 and G2/M cell cycle phases.

# 3.3. Effects of plant extract on mRNA expression

The levels of mRNAs expression were measured in the treated cells. The THP-1 cells were challenged with 100 ng/mL LPS, 100 ng/ml of LPS with 50  $\mu$ g/mL *I. coerulea* extract, 100 ng/mL of LPS with 100  $\mu$ g/mL *I. coerulea* extract, and 100  $\mu$ g/mL of *I. coerulea* alone for the duration of 4 and 6 h. As depicted in Fig. 3A–D, statistically significant increases in the expressions of TNF, IL-1 $\beta$ , and IL-18 mRNA were observed in THP-1 cells compared to their control macrophages. The mRNA expression of TNF was significantly



**Fig. 7.** Protein array expression. (A1): *I. coerulea*; (A2): cells treated with 100 ng/mL LPS + 100 µg/mL of *I. coerulea*; (A3): histograms comparing the protein array results from LPS and cells treated with *I. coerulea*; (A4) histograms comparing the protein array results from LPS and cells treated with LPS + *I. coerulea*.

higher (P < 0.0001) in THP-1 cells treated with 100 µg/mL *I. coerulea* extract alone compared to control macrophage cells at 4 h of incubation (Fig. 3A). The mRNA expression of TNF was significantly higher in THP-1 cells treated with 100 µg/mL *I. coerulea* extract alone (P < 0.0001; Fig. 3A) in relation to control macrophage cells at 4 h of incubation. The mRNA IL-1 $\beta$  expression was also evidently elevated in THP-1 cells in comparison with control macrophages when treated with 100 ng/mL LPS with 100 µg/mL *I. coerulea* extract (P < 0.0001) at 4 h and 100 ng/mL LPS with 100 µg/mL *I. coerulea* extract (P < 0.0001) at 6 h (Fig. 3B). Upon exposure to 100 ng/mL LPS with 100 µg/mL *I. coerulea* plant extract, IL-18

mRNA expression was significantly increased at 4 h of incubation (P < 0.0001; Fig. 3C). In contrast to the above results, the only mRNA expression that statistically decreased was that of CCL5 with all forms of treatment compared to control macrophages (P < 0.0001; Fig. 3D).

# 3.4. Effects of I. Coerulea plant extract on the proteins level

The production of targeted proteins was assessed for phorbol-12-myristate 13-acetate (PMA)-differentiated THP-1 cells. To evaluate, we exposed 100 ng/mL of LPS-stimulated and LPS- unstimulated THP-1 cells to 100  $\mu$ g/mL *I. coerulea* extract, and protein levels were measured by Western blot analysis. The proteins resolved by SDS-PAGE are displayed in Fig. 4A, B. The expression levels of only ERK5 and P-NF- $\kappa$ B proteins substantially increased, while BAX, BcL2, ERK1, P-I $\kappa$ B- $\alpha$ , and P-p38 proteins were noticeably lowered in LPS-stimulated THP-1 cells exposed to 100  $\mu$ g/ mL *I. coerulea* extract as compared to controls. The expression of Cyclin D1 wasn't detected. However, when LPS-unstimulated THP-1 cells were treated with 100  $\mu$ g/mL *I. coerulea* extract, the expression of proteins ERK5, BAX, and P-I $\kappa$ B- $\alpha$  augmented, whereas BcL2, Cyclin D1, ERK1, P-NF- $\kappa$ B, and P-p38 proteins expression reduced compared to non-exposed THP-1 cells (See Fig. 5).

# 3.5. Effects of plant extract on the expression of inflammatory markers

The inflammatory cytokines produced by PMA-differentiated THP-1 macrophages were analyzed by ELISA to detect the effect of plant extract on their cytokines levels. The data showed that 100 µg/mL plant extract with 100 ng/mL LPS significantly increased the expression of cytokines IL-6 and CXCL10/IP-10. Conversely, 100 ng/mL LPS with 100 µg/mL plant extract reduced the CCL22 and IL-1β expressions in THP-1 compared to controls. CXCL8/IL-8 expression was equal to that of the controls upon exposure to 100 ng/mL LPS combined with 100 µg/mL plant extract. However, 100 µg/mL of plant extract alone caused a slight increase in CXCL8/IL-8 cytokine expression. Conversely, treatment with 100 µg/mL plant extract alone resulted in decreased cytokine expressions of IL-6, CCL22, CXCL10/IP-10, and IL-1β (P = 0.09) compared to control.

#### 3.6. Effects of I. Coerulea extract on protein expressions

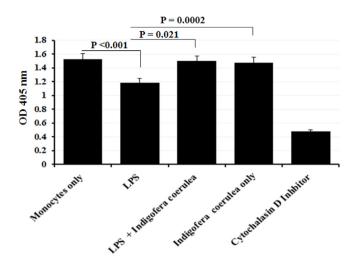
The expression levels of proteins were investigated in LPSstimulated and LPS-unstimulated THP-1 cells with or without 100  $\mu$ g/mL *l. coerulea* extract (Figs. 6 and 7). LPS-unstimulated THP-1 cells showed statistically reductions in MIF, IL-1RA/IL-1F3, and CXCL1/GROa proteins when treated with 100  $\mu$ g/mL *l. coerulea* extract compared to LPS-stimulated-only THP-1 cells. MIF remained statistically low upon 100 ng/mL LPS and 100  $\mu$ g/mL in contrast to LPS-only THP-1 cells, while CCL1 protein significantly increased.

# 3.7. Effects of I. Coerulea extract on phagocytosis activity of THP-1 cells

The phagocytosis activity of THP-1 cells treated with 100 ng/mL of LPS and 100  $\mu$ g/mL of *I. coerulea* extract was detected through Zymosan particles (Fig. 8). Cytochalasin D was also used before introducing Zymosan particles to inhibit phagocytosis activity. After treatment cells with 100 ng/mL LPS, the phagocytosis response statistically diminished compared with untreated cells. In addition, compared to control THP-1 cells, a modest increase in phagocytosis activity was reported when combination of 100 ng/mL LPS and 100  $\mu$ g/mL were used to trigger THP-1 cells. Likewise, phagocytosis function remained the same upon THP-1 cells stimulation with 100  $\mu$ g/mL *I. coerulea* extract only.

# 3.8. Effects of I. Coerulea extract on apoptosis

Apoptosis and necrosis are two types of cell death. Annexin-V/PI staining was performed and flow cytometry analysis to detect both apoptotic and necrotic markers that triggered by plant extract. Fig. 9 showed; normal cells and different apoptotic stages. The major proportion of THP-1 cells remained survive; however, apoptotic and late apoptotic cells were recorded. Treatment with 100 ng/mL LPS alone slightly induced the apoptotic THP-1 cells. Conversely, treatment with 100 ng/mL LPS along with 100 µg/mL *I. coerulea* extract



**Fig. 8.** Measurement of phagocytosis using Zymosan particles for the cells treated with 100 ng/mL LPS and 100  $\mu$ g/mL *I. coerulea* extract. Phagocytosis response was statistically diminished When THP-1 cells were treated with 100 ng/ml of LPS. Modest increase in phagocytosis activity was reported after treatment with 100 ng/ml of LPS and 100  $\mu$ g/ml together; (n = 3).

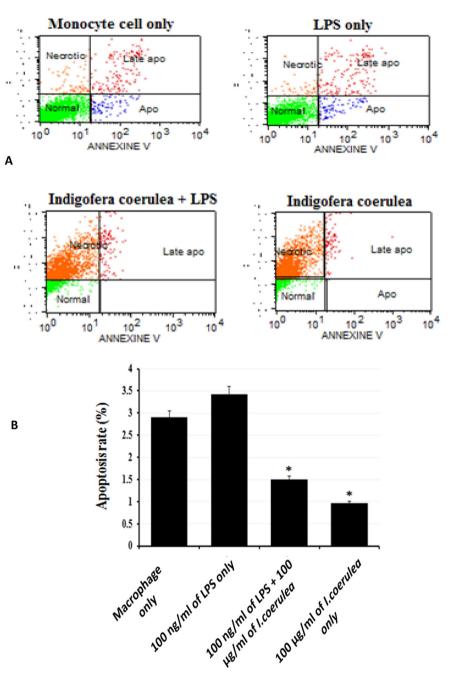
drastically reduced the apoptotic rate and repressed the late apoptosis response by THP-1 cells. Lastly, treatment with 100  $\mu$ g/mL *I. coerulea* extract alone also prevent apoptosis. The degree of apoptosis antagonism in THP-1 cells was comparatively lower with stimulation by 100 ng/mL LPS and 100  $\mu$ g/mL *I. coerulea* extract combined. These data showed that *I. coerulea* extract significantly inhibited apoptosis activity in THP-1 cells.

# 3.9. Effects of I. Coerulea extract on migration

The response of THP-1 cells migration was measured after treatment with and without 100 µg/mL *I. coerulea* extract and 100 ng/mL LPS, both measured alone and in combination (Fig. 10). The results showed the rate of THP-1 cells migration was negatively affected by 100 ng/mL LPS treatment. Upon addition of 100 µg/mL *I. coerulea* extract to 100 ng/mL LPS solution, THP-1 cells' migratory activity was significantly reduced. Similarly, 100 µg/mL *I. coerulea* extract alone didn't induce THP-1 cells' migration, which remained statistically low compared to untreated THP-1 cells.

# 4. Discussion

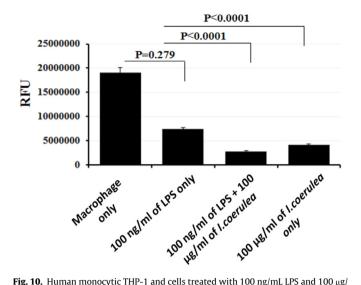
Macrophages are the main immune cells that orchestrate inflammation through the processes of phagocytosis, recruitment of inflammatory cytokines and activating the adaptive immune system (Parihar et al. 2010). The immune system are inherently conditioned to produce and discharge many cytokines, which trigger both immune and non-immune cells (Varma et al. 2016). Plantderived compounds have been screened for immunomodulation activity and used to treat immune-triggered disorders (Varma et al., 2016; Francisco et al., 2012). Our findings illustrated that treatment with both 50 and 100 µg/mL *I. coerulea* extract can powerfully halt the cellular proliferation of THP-1 cells differentiated by PMA. The proliferation arrest was further confirmed by cell synchronization analysis, indicating that I. coerulea extract induces G0/ G1 and G2/M arrest in THP-1 cells. The ERK5 protein is necessary for the change from G1 to S phase for successful cell cycle by activating cyclin-dependent protein kinases (CDK) (Gomez et al. 2016). Current results showed significantly enlarged expression of ERK5 in THP-1 cells exposed with 100 ng/mL LPS alone, 100 µg/mL I. coerulea extract alone, and 100 ng/mL LPS with 100 µg/mL I. coer-



**Fig. 9.** Annexin V/PI staining to analyze apoptosis in monocyte cells. (A) Analysis of apoptosis in monocyte cells treated with LPS + *I. coerulea* extract; (B) differences in the percentage of apoptotic between the four groups; \**p* < 0.05 compared to the monocyte cells and LPS groups. 100 ng/ml of LPS treatment alone slightly raised the apoptotic rate in THP-1 cells. Conversely, treatment with 100 ng/ml of LPS along with 100 µg/ml *I. coerulea* plant extract drastically suppressed the apoptotic activity indicating that *I. coerulea* plant extract significantly diminishes apoptosis activity in the human monocytic THP-1 cells.

*ulea* extract combined as compared with untreated macrophage cells. The exposure to 50 or 100 µg/mL *l. coerulea* extract in 100 ng/mL LPS-stimulated PMA-differentiated THP-1 cells resulted in up-regulation of the pro-inflammatory cytokines TNF-α, IL-1β, and IL-18 mRNA at 4 and 6 h compared to control. Conversely, mRNA CCL5 expression was significantly decreased among THP-1 cells treated with 50 or 100 µg/mL *l. coerulea* plant extract in combination with 100 ng/mL LPS compared to LPS-stimulated cells only at 4 or 6 h. Current results are consistent with those showed increased levels of ERK5, NF-κB signaling pathway activation (high levels of P-IκB-α and P-NF-κB), and IL-8 expression upon stimulation with *l. coerulea* extract (Pereira et al., 2019). The increased level of ERK5 is limited with the simultaneous up-regulation of

the signaling of NF- $\kappa$ B and the subsequently more IL-8 cytokine levels (Park et al. 2016). From the current data, we inferred that despite low P-p38, the NF- $\kappa$ B signaling pathway is highly activated through indirect mechanisms in THP-1 cells treated with *I. coerulea* extract, suggested by the high levels of P-I $\kappa$ B- $\alpha$  and P-NF- $\kappa$ B proteins. Another MAPK signaling pathway that can potentially activate the NF- $\kappa$ B cascade and translocate NF- $\kappa$ B into the nucleus is extracellular-signal-regulated kinase 1/2 (ERK 1/2) (Sullivan et al. 2018). Inhibition of the ERK 1/2 signaling pathway could prevent diseases linked with unrestrained inflammation. The abnormal activation of the ERK 1/2 signaling pathway has been found to be related with colon cancer cells' proliferation (Mao et al. 2014). Here, we found reduced expression of ERK1 protein upon exposure



**Fig. 10.** Human monocytic THP-1 and cells treated with 100 ng/mL LPS and 100  $\mu$ g/mL *l. coerulea* chemotaxis. We used 200,000 cells in each assay. Migratory cells were quantified by CyQuant<sup>®</sup> GR Dye. THP-1 cells migratory was negatively affected by 100 ng/ml of LPS induction. 100  $\mu$ g/ml of *l. coerulea* and 100 ng/ml of LPS reduced the activity of THP-1 cells migration. Similarly, 100  $\mu$ g/ml of plant extract alone did not improve THP-1 cells migration and remained statistically low compared to untreated THP-1 cells.

to LPS-stimulated and -unstimulated THP-1 cells to 100 µg/mL I. coerulea extract. Previously, phosphorylation of ERK 1/2 signaling pathway was shown to trigger the NF-kB signaling cascade (Winkler et al. 2017). Likewise, earlier studies reported ERK signaling pathway inhibitors to be associated with diminished NF-KB signaling (Vuong et al., 2015; Dilly et al., 2015; Dong et al., 2015). The current findings suggested IL-6 levels to be significantly lower when LPS-stimulated and -unstimulated THP-1 cells were exposed to 100 µg/mL I. coerulea plant extract (Tanaka et al., 2014: Wang and Sun. 2014: Redell et al., 2011). The obtained results showed IL-6 levels to be significantly higher when PMAdifferentiated THP-1 cells were treated with 100 ng/mL LPS. The I. coerulea extract might be immensely important in contemplating future immunomodulatory and cancer drugs targeting the IL-6/ JAK/STAT3 signaling pathway in inflammation-driven diseases. Also, we observed diminished expression of IL-1 receptor antagonist (IL-1RA) in THP-1 cells upon treatment with I. coerulea extract. IL-1RA, also known as IL-1F3, inhibits the activities of IL-1 $\alpha$  and IL- $1\beta$  and therefore acts as an anti-inflammatory protein (He et al. 2015). For migratory activity analysis, the ELISA results confirmed the decreased expression of CCL22 in LPS-stimulated and LPSunstimulated THP-1 cells when treated with 100 µg/mL I. coerulea extract compared to LPS stimulation alone. These findings highlighted that I. coerulea extract inhibits THP-1 cells migration activity by upsetting the release of CCL22 through the NF-κB or p38-MAPK transcriptional pathways.

The cells treated with *I. coerulea* extract showed an enhanced phagocytosis response in THP-1 cells. Bcl-2 protein expression was elevated in LPS-stimulated and unstimulated THP-1 cells when treated with 100  $\mu$ g/mL *I. coerulea* extract in comparison with THP-1 cells exposed to 100 ng/mL LPS alone. Interestingly, *I. coerulea* plant extract alone showed strong expression of BAX protein. However, combined treatment with LPS and *I. coerulea* extract substantially down-regulated BAX proteins. Several signaling pathways have been shown to play role in the expression of the CXCL10 chemokine (Liu et al. 2011). The increased release of CXCL10 is directly related to the high levels of TNF- $\alpha$  via activation of p38 and the downstream NF- $\kappa$ B signaling pathway in HIV-associated encephalitis (Williams et al. 2009). Rabies virus was reported to

induce the expression of CXCL10 in macrophages through activation of the ERK1/2 pathway (Nakamichi et al. 2004). Upregulation of CXCL10 has been reported in cancer to be facilitated through the p38/MAPK and NF-κB signaling cascades, which stimulate cell proliferation (Liu et al. 2011). We also found CXCL10 to be significantly elevated in the ELISA analysis upon exposure to 100 µg/mL I. coerulea plant extract in LPS-stimulated THP-1 cells, which could be the result of a heightened response from the NFκB signaling pathway. However, we found decreased levels of p38 protein in LPS-stimulated and unstimulated THP-1 cells treated with 100 µg/mL I. coerulea plant. Zhang et al., in a study on non-alcoholic steatohepatitis (NASH) patients, found that CXCL10 is related with the generation of TNF- $\alpha$ , IL-1 $\beta$ , and CCL2/MCP-1 (C-C motif chemokine ligand 2/monocyte chemoattractant protein 1), and induction of the NF- $\kappa$ B signaling pathway (Zhang et al. 2014).

# 5. Conclusions

Present findings demonstrated that *I. coerulea* extract has significant pro-inflammatory, anti-inflammatory, anti-proliferative, prophagocytic, anti-apoptotic, and anti-migratory properties in PMA-differentiated THP-1 cells. The immunomodulatory activity of the phytochemicals of *I. coerulea* extract was evidenced through regulation of investigated proteins and molecular signaling pathways. This findings offer a roadmap for future studies to confirm and elaborate upon the potential role of *I. coerulea* extract in drug development for chronic diseases. Future studies should investigate these mechanisms in animal models with chronic inflammatory diseases.

# **Declaration of Competing Interest**

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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