Induction of dissociated cytokine profiles by ISRAA with selective critical involvement of ERK1/2 in its signaling functions

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Received January 11, 2015; Accepted September 9, 2015
DOI: 10.3892/ijmm.2015.2376

Abstract. The immune system-released activating agent (ISRAA) is an immune mediator activated as a result of a nerve stimulus initiated by immune challenge. We have previously demonstrated that ISRAA and tumor necrosis factor (TNF) receptor 1 (TNFR1) share an interspecies-conserved motif (72% homology) that induces the apoptosis and proliferation of human peripheral blood mononuclear cells (hPBMCs) in a dose-dependent manner. In the present study, cytokine profiles were examined in response to the stimulation of hPBMCs with ISRAA. Furthermore, the signaling pathways induced by ISRAA were mapped. The results revealed high measurable levels of TNF-α, interleukin (IL)-6, IL-8, IL-10 and interferon (IFN)-γ, but not IL-4, IL-17 (IL-17A) or transforming growth factor (TGF)-β. The analysis of signaling pathways revealed the activation of extracellular-regulated protein kinase (ERK)1/2 as a downstream signal in the mitogen-activated protein kinase (MAPK) pathway during TNF-α and IL-6 production and apoptosis, but not during proliferation following stimulation with ISRAA by triggering the Fas-associated protein with death domain (FADD). STAT3 was found to be unphosphorylated in the ISRAA-stimulated hPBMCs, and STAT3 was ubiquitously expressed in unstimulated cells, suggesting that ISRAA has a protein inhibitor of activated STAT (PIAS)-like activity, by functioning as a negative regulator of the effects of STAT3 on the Janus kinase (JAK)/STAT pathway. The determination of the nature of cytokine responses together with the signaling pathways of cellular activity induced by ISRAA paves the way for the investigation of a potential target of ISRAA and for the development of novel therapeutic approaches for the treatment of immune-regulated disorders.

Introduction

The immune system-released activating agent (ISRAA) is a 15 kDa novel immune mediator encoded by a gene located on mouse chromosome 14 (GenBank accession no. EU552928) (1) and is believed to mediate signaling between the nervous and the immune system. ISRAA was initially considered to have a cytokine-like activity due to several factors, including its relatively low molecular mass, its biological effects on a broad spectrum of cell types, such as T and B cells and monocytes/macrophages, the ability of the same cell populations to respond to ISRAA by producing cytokines, such as interferon (IFN)-γ, as well as the fact that ISRAA is lined up on chromosome 14 on the mouse system where several low molecular mass cytokines are mapped (1). Although ISRAA was identified in the mouse, the alignment has shown that the molecule has conserved regions between many species (2). Furthermore, the function of ISRAA has been found to be dose-dependent, as it has the ability to stimulate the proliferation of human peripheral blood mononuclear cells (hPBMCs) at a low concentration of 50 pg, while a higher concentration of 5 µg has been shown to result in apoptosis (2).

In general, these two pathways for proliferation and apoptosis are controlled by a variety of (growth-deleted) factors that bind to receptors on the cell surface and initiate a cascade of events (signal transduction pathways), that convey the message from the receptors to the nucleus where transcription factors bind to DNA, turning either ‘ON’ or ‘OFF’ the production of proteins that induce cell proliferation or death (3-5). Such activities may be induced directly by the effective molecule, e.g., ISRAA, or indirectly by the production of immune mediators, such as cytokines (2). Cytokines are low molecular mass proteins (~5-20 kDa) that are important in cell signaling. They are released by cells and act on the same cell (autocrine) or on other cells in a paracrine or endocrine manner. Cytokines are produced by a broad range of cells, acting via receptors and are particularly important for the immune system as they modulate the balance between humoral and cell-mediated immune responses. In addition, they regulate the growth, maturation and responsiveness of particular cell populations. Thus, they are important in health and disease, and more specifically, in host responses to infection, immune responses, inflammation, trauma, sepsis, cancer and reproduction (6).

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Key words: apoptosis, proliferation, tumor necrosis factor receptor 1, mitogen-activated protein kinase, Janus kinase/signal transducer and activator of transcription, Fas-associated protein with death domain
The identification of ISRAA by our laboratory prompted us to investigate the mechanisms through which the activation of ISRAA initiates signaling cascades to induce particular effects, either directly or indirectly via cytokine production. We also wished to further characterize the signaling pathways involved in order to gain insight into the mechanisms through which ISRAA integrates the regulatory mechanisms and how information flows through a system.

Understanding signaling pathways is challenging, since there are hundreds of signaling pathways, as well as hundreds or even thousands of different proteins used by the cell at the same time. The fact that many signals are transient events further complicates our understanding of signaling pathways. Furthermore, the response of some pathways depends on the strength and the time of the stimulus (7,8). In the present study, we stimulated hPBMCs with ISRAA in order to examine the broad spectrum of cytokines and explore the activated signaling pathways by detecting phosphorylated proteins from second messengers down to the nuclear translocation of transcription factors.

Materials and methods

Isolation of hPBMCs. This study was approved by the Ethics Committee of Arabian Gulf University, Manama, Bahrain. Blood samples in EDTA tubes (Becton, Dickinson and Co., San Diego, CA, USA) were obtained from apparently healthy donors; none of the participants reported any history of acute or chronic medical problems, and informed consent was obtained from all donors. hPBMCs were purified using the Ficoll-Hypaque technique by diluting the blood 1:2 with phosphate-buffered saline. Subsequently, 10 ml of this diluted blood were carefully layered onto the 3 ml Ficoll-Hypaque Plus cushion (Pharmacia Biotech, Uppsala, Sweden) and centrifuged at 400 x g for 30-40 min at 18-20°C. The upper layer was drawn off using a clean Pasteur pipette, leaving the lymphocyte layer undisturbed at the interface, which was then washed twice using phosphate-buffered saline and once with RPMI-1640 containing 1% FBS. The pellet was reconstituted in 5 ml of complete RPMI-1640 (Life Technologies, Grand Island, NY, USA) culture medium containing 10% heat-inactivated fetal bovine serum, 2 mM L-glutamine, 10 mM HEPES buffer and antibiotic (50 U/ml penicillin, 50 µg/ml streptomycin). The viability of the mononuclear cells was confirmed by the trypan blue exclusion test. The viable cell numbers were counted using a hemocytometer. These two concentrations were 50 pg and 5 µg. To ensure that those two effects are due to the ISRAA recombinant protein, polyclonal antibodies to ISRAA were used, and the effect of ISRAA was neutralized by the antibodies.

MTT cell proliferation and cytotoxicity assay. The MTT Cell Proliferation Assay (ATCC, Manassas, VA, USA) was used to measure the cell proliferation rate. The protocol was performed to examine the proliferative and cytotoxic effects of ISRAA on the cells according to manufacturer's instructions. Cell suspension was harvested by centrifugation, and was then resuspended at 1x10³ cells/ml. A total of 100 µl of cell suspension was plated in each well in triplicate. Three control wells of medium were left alone to provide the blanks for the absorbance readings. Another three wells with cell suspension alone as a negative control and three wells stimulated with 5 µg of PHA as a positive control were used. A total of 10 µl of 50 pg and 5 µg ISRAA protein was added to each well in triplicate. The plates were incubated at 37°C in 5% CO₂ for 24 h. Subsequently, 10 µl of 5 mg/ml MTT reagent were added to each well, including controls and incubated for 4 h at 37°C. A total of 100 µl of MTT solvent (acidic isopropanol 0.04 M HCl in absolute isopropanol) was then added to all wells. The plates were covered with tinfoil and agitated on an orbital shaker for 15 min. Finally, the absorbance at 690 nm was recorded, and the cytocide rate was calculated using the following formula: cytocide rate (%) = (OD control - OD experiment)/OD control x100. The assay was also used to determine the effects of the mitogen-activated protein kinase (MAPK) pathway inhibitor, U0126 (MEK1/2 inhibitor,
from Cell Signaling Technology, Inc, Danvers, MA, USA; Cat. no. 9903), on ISRAA-induced apoptosis. The inhibitors were added at a concentration of 10 μM 1 h prior to stimulation.

**Measurement of CD45/CD3/CD8/CD4-positive cell count by flow cytometry.** To examine the effects of various concentrations of ISRAA on the absolute counts of mature T lymphocytes (CD3+) and the percentage of helper/inducer (CD3+/CD4+) and suppressor/cytotoxic (CD3+/CD8+) T cell subsets, Multitest CD3 FITC/CD8, PE/CD45, PerCP/CD4 APC with Trucount tubes (Becton, Dickinson and Co.) containing a freeze-dried pellet of fluorescent beads in a single-use tube were utilized. Samples were stained directly in a Trucount tube. The lyophilized pellet in the tube was dissolved releasing a known number of fluorescent beads. During analysis, the absolute number (cells/μl) of positive cells in the sample was determined by comparing cellular events to bead events. A BD Biosciences flow cytometer was used for flow cytometry, as previously described (10). The BD FACSCalibur™ flow cytometer was used with Calibrite™ beads and FACScomp version 4 and Multiset™ Software as recommended by the manufacturer. This method was also used to determine the effects of the p42 MAPK pathway inhibitor (U0126) on ISRAA-induced proliferation. The inhibitors were added at a concentration of 10 μM 1 h prior to stimulation.

**Detection of cytokines by flow cytometry.** To examine the effects of various concentrations of ISRAA at separate intervals on cytokine production BD Cytometric Bead Array (CBA), Human Soluble Protein and CBA Flex Set for the cytokines, interleukin (IL)-4, IL-6, IL-10, IL-17A, IFN-γ, tumor necrosis factor (TNF)-α, IL-8 and transforming growth factor (TGF)-β (Becton, Dickinson and Co.) were used according to the manufacturer's instructions and as previously described (11). To determine the pathway adopted by ISRAA in the production of IL-6, pharmacological inhibitors of MAPKs [U0126, MEK1/2 inhibitor (U0126) (Cell Signaling Technology, Inc.), SB20358, p38 inhibitor (Cell Signaling Technology, Inc; Cat. no. 5633)] were used and the levels of IL-6 were monitored using CBA with flow cytometry.

**Quantitative sandwich enzyme-linked immunosorbent assay (ELISA)** to detect IL-6. The hPBMCs were cultured and treated as described above. All the cells were incubated at 37°C in 5% CO₂ for 24 h. The supernatants were collected and analyzed for IL-6 expression using an ELISA kit (R&D Systems, Minneapolis, MN, USA).

**Bradford protein assay.** To perform the western blot analysis and immunofluorescence assay, the protein concentrations were measured to standardize the volume of protein extraction used on each reaction. The Bradford protein assay and nanogram measures were used (Sigma-Aldrich). This assay was performed in a 96-well plate. A total of 5 μl of a 0.1-1.4 mg/ml protein sample was used as a standard. The protein standards (5 μl), the cell lysates and buffer as a blank were added to the 96-well plate. The reagent was gently mixed; subsequently, 250 μl of the Bradford reagent were added and mixed in a shaker for ~30 sec. The samples were incubated at room temperature for 5–45 min. The absorbance was then measured at 595 nm. The protein concentration of the unknown samples was determined by comparing the Net A595 values against the standard curve, as previously described (12).

**MAPK signaling pathway detection.** MAPK activity was achieved by measuring the phosphorylation of extracellular signal-regulated protein kinase (ERK)1/2 pathways, as the main components of the MAPK downstream signaling protein which plays a vital role in the IL-6 signaling pathway. The hPBMCs were grown, counted and adjusted to 10⁶ cells/ml in complete RPMI-1640 growth medium. The cells were treated with 5 μg and 50 pg concentrations of ISRAA in triplicate wells, and the cells were then collected at intervals of 5, 10, 30 and 60 min. Total cell extracts from Jurkat cells were treated with 12-O-tetradecanoylphorbol-13-acetate (TPA; Cell Signaling Technology, Inc) as a positive control, and the cell lysates were then tested for the level and the optimal time of phosphorylation by western blot analysis of the cell lysates for phosphorylated (p-)ERK1/2 (by incubating the membrane in p44/p42 MAPK (ERK1/2) (Thr202/Tyr204; Cell Signaling Technology, Inc; Cat. no. 4370)]. The pharmacological inhibitor, U0126 (MEK1/2 inhibitor) (Cell Signaling Technology, Inc.), was used to examine the role of MAPK in the production of IL-6 as a direct or indirect effect of ISRAA. The cells were cultured (as described above), and then treated with 10 μM of U0126 inhibitor for 60 min prior to stimulation with ISRAA at intervals of 5, 10 and 30 min and then examined by western blot analysis (see below) for p-ERK1/2 (13).

**Signal transducer and activator of transcription (STAT)3 pathway detection.** The hPBMCs were grown, counted and adjusted to 10⁶ cells/ml in complete RPMI-1640 growth medium. The cells were treated with 50 pg of ISRAA in triplicate wells. The cells were then collected after 0, 5, 10, 30 and 60 min. IFN-α treated Jurkat cells were used as a positive control and tested for the level and the optimal time of phosphorylation by western blot analysis of the cell lysates for phosphorylated (p-)STAT3 (12). The hPBMCs were grown, counted and adjusted to 10⁶ cells/ml in complete RPMI-1640 growth medium. The cells were treated with 50 pg of ISRAA in triplicate wells. The cells were collected after 0, 5, 10, 15, 30 and 60 min, and were then fixed and stained directly in a multi-chamber slide. Using Western blot analysis, the cell lysates were subjected to electrophoresis on a 12% SDS-PAGE gel and transferred onto a nitrocellulose membrane. The membranes were blocked in Tris-buffered saline with 5% milk and 0.1% Tween-20. The blots were probed with anti-STAT3 (Cat. no. 9130), anti-p-Stat3 (Tyr705) (Cat. no. 9130) and anti-p-p44/p42 MAPK (ERK1/2) (Thr202/Tyr204; Cat. no. 4370) antibodies overnight and incubated with horseradish peroxidase-conjugated secondary antibody (Cat. no. 9130) as described above (13).

**Immunostaining and fluorescence microscopy for ERK1/2.** The hPBMCs were grown, counted and adjusted to 10⁶ cells/ml in complete RPMI-1640 growth medium. The cells were treated with 50 pg of ISRAA in triplicate wells. The cells were collected after 0, 5, 10, 15, 30 and 60 min, and were then fixed and stained directly in a multi-chamber slide. Using
4% formaldehyde in PBS, the cells were fixed for 15 min at room temperature, rinsed three times in PBS for 5 min each time and then permeabilized with 0.1% Triton X-100 and 5% FCS in PBS at room temperature for 60 min. The permeabilized cells were incubated with diluted primary antibody [p-p44/42 MAPK (ERK1/2) (Thr202/Tyr204) rabbit antibody; Cell Signaling Technology, Inc.] overnight at 4°C. They were subsequently washed three times in PBS for 5 min each time before the addition of the fluorochrome-conjugated secondary antibody [anti-rabbit, IgG (H+L), F(ab')2 fragment (Alexa Fluor® 488 Conjugate); Cat. no. 4412; Cell Signaling Technology, Inc.], and were then kept for 1 h at room temperature in the dark and washed with PBS. Prior to visualization by fluorescence microscopy, the cells were covered with coverslips slides with ProLong® Gold Antifade Reagent (Cell Signaling Technology, Inc.; Cat. no. 9071). Images were obtained using a fluorescence microscope (Leica DM-3000; Leica Microsystems, Milton Keynes, UK), as previously described (13).

**Statistical analysis.** ANOVA was used to compare the levels of significance between the different groups. The Student's t-test was used to calculate the levels of significance. A value of p<0.05 was considered to indicate a statistically significant difference.

**Results**

MTT cell proliferation and cytotoxicity assays. The effects of ISRAA on the cell metabolic rate were monitored by the MTT assay (Fig. 1). A significant decrease (p<0.05) in the number of cells was observed following treatment with the highest concentration of ISRAA (5 µg). However, a significant increase in cell proliferation was observed following treatment with ISRAA at the concentration of 50 pg (p<0.05) compared to the unstimulated cells (negative control, NC). PHA was used as a positive control (PC) and was found to exert significant proliferative effects (p<0.05).

Effects of ISRAA on CD45/CD3/CD4/CD8-positive hPBMC numbers determined by flow cytometry. The cells treated with 50 pg ISRAA showed a significantly increased cell number compared to the unstimulated cells and the cells treated with 5 µg ISRAA, which showed a significant decrease in cell numbers. The cell metabolic rate was monitored by the MTT assay, and a significant increase in cell proliferation was observed following treatment with ISRAA at the concentration of 50 pg (p<0.05) compared to the unstimulated cells (negative control, NC).

Effect of ISRAA on the number of CD45/CD3/CD4/CD8-positive hPBMCs. The hPBMCs were stimulated for 24 h with 50 pg ISRAA.
and 5 µg of ISRAA. PHA was used as a positive control (PC) and unstimulated cells served as a negative control (NC). The cells were counted using the CD45/CD3/CD4/CD8 MultiTEST count by flow cytometry. The cells stimulated with 50 pg of ISRAA exhibited the ability to proliferate. In order to determine the subset of T lymphocytes which proliferated more as a result of stimulation with 50 pg ISRAA, the ratio of the two subsets, i.e., the helper/inducer (CD3+/CD4+) and suppressor/cytotoxic (CD3+/CD8+) was calculated and the results are shown in Fig. 3. The T helper and T cytotoxic cells both increased in number at the same rate, with no significant difference observed between the groups (positive control and group treated with 50 pg ISRAA; p>0.05).

Cytokine production induced by ISRAA. To quantify and analyze the effects of ISRAA on the expression of a group of pro-inflammatory (IL-6, IL-17, IFN-γ, TNF-α) and anti-inflammatory cytokines (IL-8, IL-4, IL-10, TGF-β) produced by the hPMBCs stimulated with 50 pg and 5 µg of ISRAA, various techniques were used, such as CBA with flow cytometry and quantitative sandwich ELISA. Kinetic analysis on cytokine production demonstrated that a 24-h incubation period is the optimal time point for the maximum induction of the measured cytokines (data not shown).

As shown in Fig. 4, with the use of CBA with flow cytometry, our results revealed that stimulation with 50 pg ISRAA markedly increased the levels of the pro-inflammatory
cytokines, IL-6 and TNF-α (p<0.05), but not those of IL-17 compared to the unstimulated cells (Fig. 4A). Stimulation with ISRAA (50 pg) also induced a significant increase in the levels of IL-8 (Fig. 4B) and IFN-γ (Fig. 4C) (p<0.05). The analysis of anti-inflammatory cytokines revealed a significant induction of IL-10 (p<0.05), but not IL-4 or TGF-β (Fig. 4D). The toxic concentration of 5 µg ISRAA did not significantly alter the levels of all the examined cytokines.

The cytokine IL-6 exhibited the highest detectable cytokine levels, as shown by CBA with flow cytometry. These results were confirmed by ELISA, which revealed a 4-fold increase in IL-6 production in the cells stimulated with 50 pg ISRAA compared to the unstimulated cells (p<0.05; Fig. 4E).

Western blot analysis of p-ERK1/2. The phosphorylation of ERK1/2 as a MAPK downstream signaling protein was examined by western blot analysis of extracts from cells treated with 50 pg ISRAA with or without ERK1/2 inhibitor (U0126) at intervals of 5, 10, 30 and 60 min. Total cell extracts from Jurkat cells treated with TPA were used as a positive control. The results revealed that detectable ERK1/2 phosphorylation began at 10 min and steadily continued to increase until the 60-min time point following stimulation, while low levels of phosphorylation or no phosphorylation were observed in cells stimulated with ISRAA when the inhibitors were used (Fig. 5).

**ERK1/2 translocation.** Upon activation, ERK1/2 translocates to the nucleus in order to be transcribed. Using immunofluorescence assay to reveal the exact time required for p-ERK1/2 to translocate to the nucleus, the results demonstrated that the phosphorylation and dimerization of ERK1/2 began at around 5 min, and intensified at around 10 min. It then began to translocate from the cytoplasm towards the nucleus at 15 min where some of the phosphorylated molecules entered the nucleus. This intensified at 30 min, and by 60 min, a complete translocation to the nucleus had occurred (Fig. 6).

**Effect of ERK1/2 inhibitor on cell proliferation and apoptosis.** The hPBMCs were cultured and treated with 5 µg or 50 pg of ISRAA with or without the p-ERK1/2 inhibitor. Data were obtained using the MTT assay for cell death (Fig. 7A), and cell proliferation was analyzed by taking the absolute counts of mature helper/inducer (CD3+/CD4+) and suppressor/cytotoxic (CD3+/CD8+) T lymphocyte subsets by flow cytometry (Fig. 7B). As shown in Fig. 7, the results revealed that, compared to the unstimulated cells (negative control, NC), there was a significant increase in hPBMC proliferation (p<0.05) following stimulation with 50 pg of ISRAA with or without the addition of the inhibitor. However, the toxic effect of the 5 µg concentration of ISRAA was significantly abrogated by the addition of the inhibitor and, consequently, a significant increase in proliferation was observed following the addition of the inhibitor (p<0.05). PHA was used as a positive control (PC) and did not significantly affect proliferation (p>0.05).

**Effects of ERK1/2 inhibitors on the production of IL-6.** To determine the pathway adopted by ISRAA in the production of IL-6, the pharmacological inhibitors of MAPKs,
Our results demonstrated the abrogation of IL-6 production as a result of pre-incubation with the two blockers prior to stimulation with ISRAA (p<0.05; Fig 8).

**Effect of ERK1/2 inhibitors on the production of TNF-α.**

Treatment with 50 pg of ISRAA with or without the addition of the ERK1/2 inhibitor using the flow cytometric technique revealed a significant decrease in the expression of TNF-α when using the inhibitor compared with the increased level observed in the cells treated with 50 pg of ISRAA (p<0.05; Fig 9).

**Phosphorylation of STAT3 proteins.**

IL-6 was originally cloned as an acute-phase response factor resulting in the activation of the Janus kinase (JAK)/STAT3 pathway. To examine the phosphorylation of the STAT3, western blot
analysis was performed using monoclonal antibodies to phosphorylated and unphosphorylated STAT3. The results revealed that STAT3 factors were ubiquitously expressed in an unstimulated inactive unphosphorylated form. The examination of p-STAT3 revealed no phosphorylation of STAT3 (Fig. 10).

**Discussion**

ISRAA is an immune mediator activated upon a nerve stimulus triggered by an immune challenge (1). The human counterpart of mouse ISRAA is not known; however, we recently demonstrated that ISRAA contains an interspecies-conserved functional motif sharing 72% homology with a motif present in the TNF receptor 1 (TNFR1), a receptor connected to intracellular domains inducing dose-dependent signals for survival or death, and the potential proliferative effects of mouse ISRAA on human cells were recorded (2). In this study, we confirmed this activity using different techniques and examined the induction of a broad spectrum of pro-inflammatory and anti-inflammatory cytokines in response to the stimulation of hPBMCs with ISRAA. Furthermore, the signaling pathways used by ISRAA to induce cytokine production were explored.

The results revealed increased levels of IL-6, IL-8, IL-10, IFN-γ and TNF-α produced by the ISRAA-stimulated hPBMCs, whereas no changes were observed in the production of the IL-4, IL-17A and TGF-β cytokines in response to ISRAA stimulation. Since ISRAA is involved in different cellular activities, including death, growth and survival, complex signaling pathways may be involved. Moreover, dissociated cytokine profiles were recorded to induce such activities considering that cytokines are signaling proteins with pleiotropic effects in diverse aspects of bodily functions in health and disease and are used extensively in intercellular communications to produce different activities. Among the produced cytokines, IL-6 showed the highest measurable levels in the current study.

IL-6 possesses a number of functional properties and this is reflected by the terminology used to describe the activities of this cytokine (IFN-β2, hepatocyte-stimulating factor, cytotoxic T cell differentiation factor, B cell differentiation factor, and B cell stimulatory factor 2) (14). It is well known for promoting inflammatory events through the expansion and activation of T cells, the differentiation of B cells, and the induction of acute-phase reactants cells. By contrast, it also plays a protective role during disease and counteracts the manifestation of certain inflammatory responses (15-17). Thus, IL-6 functions not only as a pro- but also as an anti-inflammatory cytokine, and, as a result, plays a pivotal role during disease.

To understand the mechanisms through which ISRAA transduces its potent biological responses, we briefly examined the signaling pathway used by ISRAA to induce the production of IL-6. The signaling pathways known to be involved in the production of IL-6 have been mainly found to be the MAPK and/or JAK/STAT pathways (18). Therefore, these two pathways were analyzed after stimulating the hPBMCs cells with ISRAA.

The ERK-MAPK pathway was the first signal transduction cascade to be unraveled and delineated from the cell membrane to the nucleus (4). The pathway has been extensively investigated and is also known as the classical mitogen kinase cascade (19-23,28). The cascade is initiated by the small G protein Ras, which recruits the MAPKKs A-Raf, B-Raf and Raf-1, the MAPKs ERKs, in addition to a large number of a range of substrates, including membrane and cytoskeletal proteins, cytosolic enzymes, and transcription factors (24). Ras/MAPK regulates many cellular processes and plays an important role in physiological processes, such as proliferation, differentiation and development (25,26). Moreover, Ras/MAPK has also been found to control the signaling pathway leading to the phosphorylation and inactivation of a pro-apoptotic Bcl-2 family member in mammalian cells, Bad, which is targeted on serine 112 by a MEK-dependent pathway, whereas the phosphorylation of two other residues occurs in a MEK-independent manner (one of them is serine 136 by the PI3K/AKT pathway). Serine 112 phosphorylation has been found to be required for the dissociation of Bad from Bcl-2 providing a key link between growth and survival pathways and protection from apoptosis (27).

ERK1/2 signaling has been implicated as a key regulator of cell proliferation, and for this reason, inhibitors of the ERK pathway are used in clinical trials as potential anticancer agents (28). In this study, we examined the involvement of ERK1/2 and p38 kinase in mediating IL-6 production from hPBMCs using U0126 and SB203580 as inhibitors of ERK1/2 and p38 kinase, respectively. We demonstrated that by treating the cells with U0126 and SB203580 for 1 h prior to stimulation...
with ISRAA, the differential effects of these compounds were observed in the level of IL-6, which was markedly decreased, suggesting the essential role and function of MAPKs in the level of IL-6 produced by cells stimulated with ISRAA.

In the present study, western blot analysis and immunofluorescence assay were used to investigate the levels and intensity of p-ERK (Thr202/Tyr204) as a downstream target protein, and the results revealed that P-ERK1/2/p-ERK1/2 as downstream signals in the MAPK pathway were phosphorylated due to ISRAA stimulation and a significant amount of p-ERK1/2 accumulated in the nucleus. The localization of MAPK depends on its phosphorylation state at the sites of action rather than on its activity. Phosphorylation induces its cyto-dimerization, which is required for its nuclear translocation. Upon activation, both kinases partly translocate to the nucleus (29,30).

IL-6 signals via gp130 homodimerization which then activates JAKs and recruits STAT proteins (31). STAT factors are ubiquitously expressed in an unstimulated, inactive (unphosphorylated) form (32), predominantly regulated by post-transitional modifications, i.e., tyrosine and serine phosphorylation. This is in accordance with our findings since unphosphorylated STAT3 in the ISRAA-stimulated hPBMCs was noted, and STAT3 was ubiquitously expressed in unstimulated cells suggesting that ISRAA has a protein inhibitor of activated STAT (PIAS)-like activity by acting as a negative regulator of the JAK/STAT pathway, where PIAS1 and PIAS3 have been shown to inhibit the activity of STAT1 and STAT3, respectively (33).

The effects of ISRAA on cells have been shown to be dualistic, as high concentrations induce apoptosis and low concentrations induce proliferation (2). This study demonstrated that ERK1/2 had no effect on the ISRAA-induced cell proliferation, while it was shown that ERK1/2 is required for ISRAA-induced apoptosis since the inhibition of ERK activities by a small molecule inhibitor almost completely inhibited the apoptotic effects of the high dose of ISRAA and reversed the suppressive effect of the cytotoxic dose. This is in agreement with the results of other studies (34-36), suggesting that the ERK pathway mediates the apoptosis induced by different stimuli in different tissues even though ERK has generally been considered a survival signaling pathway. However, the mechanisms through which ERK mediates apoptosis remain unclear. We hypothesized that the ERK1/2 pathway is involved in apoptosis by increasing an upstream signal for TNF-α production via the extrinsic pathway by activating the death domain of the FADD pathway. This is supported by Jo et al (37), who examined the effect of ERK inhibition on TNF-α expression and subsequent caspase-3 activation in cisplatin-induced acute renal failure in mice. The authors demonstrated that the inhibition of ERK1/2 reduced TNF-α expression, caspase-3 activation and apoptosis in kidney tissue, suggesting that the ERK1/2 pathway also participates in apoptosis by increasing an upstream signal for TNF-α production. In this study, we also demonstrated that the TNF-α level induced by ISRAA was significantly decreased when ERK1/2 was inhibited, indicating that ISRAA/ERK1/2 may use the extrinsic apoptotic pathway by activating the death receptors via increasing the level of TNF-α to induce apoptosis. These data have been recently supported by an exhaustive structural analysis of ISRAA that demonstrated features of novel signaling molecules (unpublished data).

In conclusion, the present study confirmed the dose-dependent apoptotic and proliferative effects of ISRAA on hPBMCs. Furthermore, high measurable levels of dissociated cytokine responses were detected, and the downstream signals of the MAPK pathway (ERK1/2) were found to be critically involved. The results also revealed that ISRAA functions as a PIAS-like protein, functioning as a negative regulator of STAT3 action on the JAK/STAT pathway. Since FADD was activated with both concentrations of ISRAA (proliferative and toxic) and ERK1/2 activation was involved in the IL-6 and TNF-α production and in apoptosis, but not in proliferation, further research into the intrinsic pathways is required to examine the effects of ISRAA on ERK and consequently on different caspases, and p53 and Bcl-2 proteins, as well as on survival signaling, such as the Ras/PI3K/Akt pathway and Fyn expression. Experiments to define the proliferative and apoptotic concentrations for each hPBMC population during proliferation and apoptosis have been recently initiated in our laboratory.

Acknowledgements

This study received financial support from the College of Medicine and Medical Sciences, Arabian Gulf University (Bahrain).

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