Mitogenic Signals Initiated via Interleukin-6 Receptor Complexes in Cooperation with Other Transmembrane Molecules in Myelomas

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Cytokines exert multiple biological functions through binding to their specific receptors that triggers activation of intracellular signaling cascades. The cytokine-mediated signals may produce variable and even opposing effects on different cell types, depending on cellular context that is also dictated by the differentiation stage of the cell. Multiple myeloma (MM) is a monoclonal proliferative disorder of human plasma cells. Myeloma cells appear to include mixed subpopulations in accordance with the expression of their surface antigens, such as CD45. Although interleukin-6 (IL-6) is widely accepted as the most relevant growth factor for myeloma cells, only a few subpopulations of tumor cells, such as CD45⁺ immature cells, proliferate in response to IL-6. The activation of both signal transducer and activator of transcription (STAT) 3 and extracellular signalregulated kinase (ERK) 1/2 is not sufficient for IL-6-induced proliferation of myeloma cells that requires the src family kinase activation associated with a rapid translocation of CD45 to lipid rafts. The CD45 expression renders myeloma cells competent for not only mitogenic but also apoptotic stimuli, resulting in either proliferation or apoptosis of CD45⁺ myeloma cells dependently upon the circumstantial stimuli. In contrast, in CD45⁻ myeloma cells highly expressing IL-6 receptor a chain (IL-6Ra), IL-6Ra and insulin-like growth factor (IGF)-I receptors exist on plasma membrane in close proximity, facilitating efficient assembly of two receptors in response to IL-6. The synergistic effects of IL-6Ra on IGF-I receptor-mediated signals provide a novel insight into a Jak-independent IL-6 signaling mechanism of receptor cross talk in human myeloma cells. Furthermore, the signaling cross talk between the cytokine receptor, IL-6Ra/gp130 and the growth factor receptor tyrosine kinase, fibroblast growth factor receptor (FGFR) 3 appears in myeloma cells carrying t(4;14)(p16.3;q32). In this review we propose several mechanisms of the IL-6-induced cell proliferation that is strictly dependent upon the cellular context in myelomas. [J Clin Exp Hematopathol 46(2) : 55-66, 2006]

Keywords: CD45, interleukin-6 receptor (IL-6R), myeloma, proliferation, receptor tyrosine kinases (RTKs)

INTRODUCTION

Immune and hematopoietic systems are regulated through the control of a number of cellular processes including cell growth, differentiation, migration and death by cellcell interaction and by soluble factors. Ligation of receptors by their specific ligands initiates a signal transduction cascade that leads to multiple biological responses. Intracellular signal transduction events depend on the subtle balance between protein-tyrosine kinases (PTKs) and protein-tyrosine phosphatases (PTPases) that control tyrosine phosphorylation. In the immune system, precise and coordinated regulation of this equilibrium allows for rapid responses to foreign antigens, whereas an imbalance between PTKs and PTPases can have pathologic consequences including autoimmunity, immunode-ficiency and malignancy¹.

Cytokines exert multiple biological functions through interactions with their specific receptors. The binding of cytokines to their receptors induces dimerization of the receptors, and triggers activation of intracellular signaling cascades². The earliest event is activation of Janus kinase (Jak) family members including Jak1, Jak2, Jak3 and Tyk2³. Activated Jaks phosphorylate a latent cytoplasmic transcription factor, signal transducer and activator of transcription (STAT), the most important substrate for Jaks, initiating the signal transduction pathways critical for cytokine effects⁴. Seven members of STAT family proteins are activated by distinct cytokines, and therefore, the Jak-STAT pathway seems to be a key pathway to introduce the specific cellular responses by cytokines. Eight members of suppressor of

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cytokine signaling (SOCS) family proteins induced by STATs act in a negative feedback loop to suppress the Jak-STAT pathway via interacting with either Jaks^{5,6} or cytokine receptors⁷, resulting in attenuation of the cytokine signal transduction.

Multiple myeloma (MM) is characterized as a clonal human plasma cell-neoplasm, and develops mainly in bone marrow (BM). Human myeloma cells freshly isolated from BM could respond to a cytokine, interleukin-6 (IL-6), and proliferate *in vitro*⁸. However, the only minor populations of myeloma cells show a significant proliferative activity in response to IL-6. Despite the clonal origin of tumor cells, there are phenotypically mixed subpopulations of myeloma cells from the same patient. Immature myeloma cells proliferate markedly in response to IL-6, whereas mature myeloma cells fail to proliferate but secrete antibodies *in vitro*⁹. Thus, myeloma cells are heterogeneous with respect to their biological characters as well as to their immunophenotypes, and we find that CD45-expressing immature cells belong to the proliferating populations in MM¹⁰.

Some growth factors, such as insulin¹¹, insulin-like growth factor-I (IGF-I)^{12,13} and basic fibroblast growth factor (bFGF/FGF-2)¹⁴ of which receptors have tyrosine kinase domains are also reported to function as growth and survival factors for MM. Most receptor tyrosine kinases (RTKs) are composed of a single polypeptide chain, and are monomeric in the absence of their ligands. Ligand binding to the extracellular portion of RTKs leads to dimerization of receptors, resulting in autophosphorylation of specific tyrosine residues in the cytoplasmic portion¹⁵. Tyrosine autophosphorylation either stimulates the intrinsic catalytic (kinase) activity of the receptor or generates recruitment sites for the downstream signaling proteins containing phosphotyrosine-recognition domains, such as src homology (SH) 2 domain or phosphotyrosine-binding domain¹⁶.

Cytokines and growth factors bind to their receptors on the cell surface and activate signal transduction pathways that frequently lead to cell proliferation and differentiation. Promotion of cell proliferation by cytokines or growth factors has been shown to correlate with the intracellular activation of common protein kinase cascades. Myeloma cells use cytokine receptors, such as IL-6 receptors, lacking kinase activity as main receptors for promoting their growth, while they also express RTKs, such as IGF-I receptors, as well as other type of receptors. In this review, we propose the molecular mechanisms of the IL-6-induced proliferation of myeloma cells, such as the cooperation of several transmembrane molecules with the IL-6 receptor (IL-6R) complexes, that eventually influences the outcome of cellular responses.

IL-6 AS A GROWTH FACTOR FOR HUMAN MYELOMA CELLS

Among the numerous number of growth enhancing factors for myelomas postulated, such as IGF-I¹³, IL-10¹⁷ and IL-21¹⁸, IL-6 is considered as the most important and best characterized growth stimulatory factor for myeloma cells. IL-6 has a variety of biological functions in different cells¹⁹, whereas IL-6 enhances proliferation of myeloma cells which is inhibited by treatment with anti-IL-6 antibodies in vitro⁸ and in vivo²⁰. Moreover, in Balb/c mice the overexpression of IL-6 transgene induces plasmacytomas²¹, and the IL-6deficient mice with Balb/c genetic background are completely resistant to the development of plasmacytomas induced by pristane oil²² and by myc/raf-expressing retrovirus (J3V1) infection in combination with pristane²³, demonstrating that IL-6 also plays a crucial pathogenic role in the onset of plasma cell tumors in vivo. In addition, IL-6 has an antiapoptotic effect on myeloma cells²⁴, and therefore, IL-6 supports the survival and expansion of myeloma cells not only by stimulating cell division, but also by preventing apoptosis. IL-6 is a member of the IL-6 family of cytokines including leukemia inhibitory factor, oncostatin M, IL-11, ciliary neurotrophic factor and cardiotrophin-1 which commonly use gp130 as a signal transducing molecule (see below), and some are reported to actually stimulate myeloma cell growth²⁵. In myeloma IL-6 is provided by the tumor itself⁸ and BM stromal cells²⁶.

IL-6 SIGNALING PATHWAYS

IL-6R complexes consist of two subunits termed as gp80/a chain and gp130/ β chain. IL-6Ra directly binds to IL-6, while gp130 responsible for signal transduction is commonly used by the members of the IL-6 family of cytokines²⁷. Upon IL-6 binding to IL-6Ra, Jaks, such as Jak1 and Tyk2, constitutively associated with gp130 are activated, followed by phosphorylation of the cytoplasmic region of gp130 with the activated Jaks. The phosphorylated tyrosine residues in gp130 recruit STAT3 followed by tyrosine phosphorylation with Jaks. The activated STAT3 dimerizes and translocates to the nucleus in which STAT3 controls the gene expression²⁸. Serine phosphorylation of STAT3 is also important for full activation of transcriptional activity²⁹. In addition to the Jak-STAT pathway, a simple signaling cascade that directly links the cell membrane and the nucleus, IL-6 also induces the activation of Ras via Sos and Grb2 interacting with SHP-2 that is also recruited to the phosphorylated gp130. Subsequently, the downstream signaling intermediates of Ras, such as Raf-1, MAP or ERK kinase (MEK) 1/2 and extracellular signal-regulated kinase (ERK) 1/2, are activated³⁰. Activation of both the Jak-STAT and Rasmitogen activated protein kinase (MAPK) pathways is necessary for cell proliferation³⁰. In some cases, the phosphatidylinositol (PI)-3 kinase pathway is also activated via gp130 where Gab-1 acts as an adaptor molecule in transmitting signals to the PI-3 kinase from gp130 and SHP- 2^{31} .

CD45 EXPRESSION, ISOFORMS AND PROLIFERATION OF MYELOMA CELLS

Despite the character of CD45 as a leukocyte common antigen, it has been well known that some acute lymphocytic leukemia (ALL) and plasma cell lines lack the CD45 expression. In looking at human myeloma cells from the MM patients, it appears that there is the CD45 expression on immature myeloma cells but not on mature ones³²⁻³⁵. The more careful investigation show that the CD45 molecule is expressed on all nucleated hematopoietic cells including plasma cells, whereas the expression of CD45 molecule in primary myeloma cells and cell lines is quite variable, consistent with the heterogeneity of myeloma cells. In human, CD45 is composed of several isoforms that vary in their extracellular region as a result of tissue or cell type-specific alternative splicing of exons 4, 5, and 6^{36} . Although the requirement of the intracellular PTPase domain of CD45 has been documented³⁷, the function of the extracellular domain that differs among these isoforms remains a major unresolved issue. B lymphocytes and plasma cells express CD45 and CD45RA (high-molecular weight isoforms), CD45RB but not CD45RO, whereas the restricted subpopulation of myeloma cells and some myeloma cell lines expresses CD45 and CD45RO, CD45RB but not CD45RA¹⁰, although most myeloma cells lose CD45 expression. It appears that isoforms of CD45 alter during oncogenic event, such as from CD45RA to CD45RO, and CD45RO but not CD45RA seem to have an ability to help the cell proliferation by IL-6 (see below)³⁸ as some reports have shown functional differences among isoforms39-42.

Immature myeloma cells but not mature ones include proliferating cells evaluated by expression of Ki-67 antigen¹⁰, consistent with the results obtained from [³H]thymidine incorporation and cyclin D1⁺p16⁻ cell populations⁴³. Ki-67⁺ myeloma cells are found in the CD45⁺ rather than CD45⁻ fractions. Thus, the CD45⁺ immature myeloma cells appear proliferative populations in MM¹⁰. Conversely, in the culture without IL-6 those CD45⁺ immature myeloma cells undergo apoptosis more readily than CD45⁻ immature ones¹⁰, suggesting that the CD45⁺ immature myeloma cells essentially require IL-6, whereas CD45⁻ immature ones seem to be more independent of IL-6 (Fig. 1). Despite the proliferating populations of CD45⁺ immature myeloma cells, the limiting amounts of IL-6 provided in BM may result in the few numbers of CD45⁺ myeloma cells in physiological conditions.



Fig. 1. CD45 expression and IL-6 responses in myeloma cells. In immature myeloma cells, CD45⁺ cells respond to stimuli including proliferation, differentiation and apoptotic signals, whereas CD45⁻ cells are resting and not competent for such signals, thereby resistant to cell growth and death⁵⁶. CD45⁻ MPC⁻1-CD49e⁻ immature myeloma cells are postulated to potentially belong to a category of tumor stem cells in MM¹⁰. CD45⁺ immature myeloma cells would undergo apoptosis in IL-6- insufficient physiological conditions.

REQUIREMENT OF SRC FAMILY KINASE ACTIVITY ASSOCIATED WITH CD45 FOR MYELOMA CELL PROLIFERATION BY IL-6

CD45, a receptor-type PTPase, is ubiquitously expressed in all nucleated hematopoietic cells³⁶. Genetic defects of CD45 in mice⁴⁴ and humans⁴⁵ cause severely combined immunodeficiency that demonstrates the essential role of CD45 in the immune system especially for the activation and development of lymphocytes. CD45 regulates tyrosine phosphorylation of the molecules, such as src family PTKs, in the signal transduction of the T cell antigen receptor (TCR) and B cell antigen receptor (BCR)^{36,37,44-47}. One conventional model holds that when src family PTKs are phosphorylated at the negative regulatory COOH-terminal tyrosine residue by Cterminal src kinase (Csk), they adopt to the aforementioned closed and inactive conformation. The dominant role of CD45 is to dephosphorylate this negative regulatory tyrosine, setting the stage for antigen receptor signaling to activate src family PTKs by autophosphorylation^{37,48-50}.

We find that IL-6 rapidly activates both STAT3 and ERK1/2 in CD45⁻ myeloma cell lines, although they fail to proliferate in response to IL-6. These myeloma cell lines lack CD45 expression and src family PTK activation, implying

that in myeloma cells the activation of STAT3 and ERK1/2 is not sufficient for cell proliferation enhanced by IL-6. The activation of src family PTKs associated with the CD45 PTPase seem to be a prerequisite for the myeloma cell proliferation that further requires the activation of STAT3 and ERK1/2 by IL-6⁵¹. Among the molecules we tested, only Lyn appears to interact with CD45 in CD45⁺ myeloma cell lines, results suggesting that Lyn could be a major molecule regulated by the CD45 PTPase in myelomas. Indeed, inhibition of either CD45 PTPase or src family PTKs suppresses IL-6enhanced DNA synthesis of CD45⁺ rather than CD45⁻ myeloma cells. Thus, with respect to the IL-6-induced proliferation of myeloma cells, CD45 could act as a positive regulator by controlling src family PTK activity, similarly to mitogenic signals via BCR on B cells.

Although the src family PTKs have been shown to activate STATs^{52,53}, the inhibition of Lyn expression or kinase activity does not influence STAT3 activation in CD45⁺ myeloma cells. Thus, in myeloma cells src family PTKs are not directly involved in STAT3 and Ras-ERK1/2 activation by IL-6. Recent reports have supported our suggestion, showing that Lyn is required for granulocyte-colony stimulating factor (G-CSF)-induced cell proliferation⁵⁴, and that the G-CSF-stimulated signals require src family PTK activities dissociated from the Jak-STAT pathway⁵⁵. The src family PTKs may phosphorylate several molecules including adaptor molecules, enzymes, structural proteins and transcription factors, resulting in a diversification of the initial signals⁴⁷. We also find that in CD45⁺ myeloma cells Lyn kinase is likely to activate phospholipase C (PLC)- γ -mediated signals, such as



cell proliferation

Fig. 2. CD45 plays a pivotal role in the IL-6- induced growth promotion of myeloma cells. The activation of Lyn associated with CD45 in parallel with STAT3 and ERK1/2 activation is induced by IL-6 in myeloma cells, followed by the activation of PLC- γ 2, PKC and Ca²⁺ influxes that are essential for the IL-6- induced proliferation of tumor cells^{51,56}. IP₃R : inositol 1, 4, 5- triphosphate (IP₃) receptor.

cellular calcium influxes and protein kinase C (PKC) activation, independent of STAT3 and ERK1/2 pathways (Fig. 2)⁵⁶.

IL-6-INDUCED TRANSLOCATION OF CD45 TO LIPID RAFTS IN MYELOMA CELLS

Considering the relationship between CD45 and its substrates, the cellular localization of CD45 and its accessibility to the substrates may define the activity on signaling. Lipid raft microdomains represent cholesterol- and glycosphingolipid-enriched dynamic patches in the plasma membrane and organize the plasma membrane into functional units⁵⁷. These raft domains act as platforms for conducting a variety of cellular functions including TCR, BCR and cytokine signaling⁵⁸⁻⁶⁰. Several protein families have been reported to modify lipid rafts structurally and functionally that include membrane integral proteins, such as caveolins and flotillins, and lipid chain-modified proteins, such as src family PTKs and Ras⁶¹.

We find that CD45RO and CD45RB but not CD45RA are translocated to lipid rafts in myeloma cell lines and an IL-6R*a*-transfected B cell line in response to IL-6. Furthermore, IL-6 induces a more rapid translocation of CD45RO to lipid rafts than CD45RB, and the translocated CD45 functions as a positive regulator of proliferation signaling by triggering dephosphorylation of Lyn Tyr⁵⁰⁷ on lipid rafts, where IL-6R*a*, gp130 and Lyn are constitutively present³⁸. Glycosylation of CD45 is reported to regulate the formation of clusters induced by galectin-1 binding and resultant phosphatase activity⁶². Also, we cannot exclude the possibility of a different complex with other proteins, different substrate binding or different type of dimeric formation.

As for the functional role of the translocated CD45 in myeloma cells with IL-6 stimulation, we show the formation of a complex between CD45 and Lyn, decreased phosphorylation of Lyn Tyr⁵⁰⁷ at the carboxy terminal and subsequent kinase domain Tyr³⁹⁶ phosphorylation by IL-6. Csk is recruited by phosphoprotein-associated with glycosphingolipidenriched microdomains (PAG) which targets lipid rafts by palmitovlation^{63,64}. The association of PAG with Csk is controlled by dephosphorylation of PAG. In myeloma cells PAG dephosphorylation and the resultant release of Csk from lipid rafts occurs after IL-6 stimulation in accordance with CD45 translocation. CD45 might be a candidate for the PTPase of PAG and dephosphorylation of PAG allows Csk to be released from PAG⁶⁵. We show that CD45 is directly associated with PAG in lipid rafts after IL-6 stimulation. Thus, IL-6 treatment induces the translocation of CD45 to lipid rafts sequentially followed by the association of CD45 with Lyn and PAG, dephosphorylation of Lyn Tyr⁵⁰⁷ and PAG Tyr³¹⁴, Lyn activation and Csk release from lipid rafts. Again, CD45 functions as a positive regulator of signal transduction in myeloma cells, although CD45 is also reported to negatively

regulate cytokine signaling⁶⁶. We also find that the translocated CD45 binds to flotillin 2, which possibly organizes the network of lipid rafts. The function of flotillin and the significance of the complex with CD45 is not clear, however, the report that flotillins form preassembled platforms in hematopoietic cells and these platforms recruit signaling molecules upon activation through lipid rafts⁶⁷ raises the possibility that flotillins might participate in CD45-recruiting mechanisms to lipid rafts to transduce IL-6 signal from IL-6R α .

CD45⁺ MYELOMA CELLS ARE SUSCEPTIBLE TO APOPTOSIS

As cell viability of CD45⁻ myeloma cells in vitro is better than CD45⁺ myeloma cells freshly isolated from MM patients¹⁰, we show that CD45⁺ myeloma cell lines are more sensitive to several apoptotic stimuli as well as a mitogenic stimulus⁵⁶. The intracellular reactive oxygen species (ROS) and calcium ion (Ca²⁺) seem to initiate the cellular events, such as the activation of the src family PTKs and calcineurin phosphatase, which trigger the apoptosis upon the oxidative or endoplasmic reticulum (ER) stresses. The CD45 expression is likely to enhance the hydrogen peroxide-induced apoptosis via the activation of the src family PTKs that is regulated by their redox state⁶⁸⁻⁷². In contrast, the thapsigargininduced apoptosis is related to the increased expression of voltage-dependent anion channel (VDAC) 1 that is identified as a gene highly expressed in a CD45⁺ myeloma cell line by cDNA subtraction. Bad sequestered by calcineurin in cytosol moved to mitochondria and bound to Bcl-2 in response to the increase in the intracellular Ca2+ concentration, resulting in the enhanced release of cytochrome c presumably through the highly expressed VDAC1 associated with Bax (Fig. 3). The VDAC protein is a subunit of the mitochondrial permeability transition pore (PTP), a large channel whose opening results in rapid loss of membrane potential and organellar swelling, quickly leading to cytochrome c release and apoptotic cell death⁷³. Bax and Bak enlarge the VDAC cahnnel which is contrarily closed by Bcl-xL⁷⁴. VDAC1 plays an essential role in apoptosis in mammalian cells⁷⁵, and the increase in VDAC expression may facilitate the release of cytochrome c during apoptosis⁷⁶.

We show that CD45⁺ myeloma cells accompanied by the increased expression of VDAC1 seem to be sensitive to the various apoptotic stimuli, such as serum deprivation, ultraviolet-ray irradiation, heat shock, SERCA inhibitor (thapsigargin), hydrogen peroxide, arsenic trioxide, interferon (IFN)- γ and melphalan treatments⁵⁶. We find the association of calcineurin with Bad in CD45⁺ myeloma cells more than CD45⁻ ones. The mitochondrial translocation of Bad and the increased association of Bad with Bcl-2 are also observed in CD45⁺ myeloma cells treated with hydrogen peroxide or thapsigargin. The oxidative stress-induced activation of the src



Fig. 3. The mitochondrial apoptosis pathways in CD45⁺ myeloma cells accompanied by the increased expression of VDAC1. The src family kinases may be activated by the intracellular ROS in CD45⁺ myeloma cells, followed by Ca²⁺ influxes, calcineurin activation, mitochondrial translocation of Bad, dysfunction of Bcl-2 and cytochrome c release presumably through the increased expression of VDAC1. Bax released from Bcl-2 interacts with VDAC1, leading to cytochrome c release and apoptotic cell death⁵⁰. H₂O₂: hydrogen peroxide; THG: thapsigargin; SERCA: sarcoplasmic-endoplamic reticulum Ca²⁺ adenosine triphosphatases; ANT: adenine nucleotide translocator.

family PTKs in CD45⁺ myeloma cells leads to the intracellular Ca²⁺ fluxes, and the sustained increase in cytosolic Ca²⁺ leads to the activation of the serine/threonine phosphatase, calcineurin. Bad dephosphorylated by calcineurin is considered to move to mitochondria and acts as an antagonist for Bcl-2/Bcl-xL, and the dysfunction of Bcl-2 that precedes the cytochrome c release through the VDAC1 associated with Bax leads to apoptosis of the cells (Fig. 3)⁷⁷. The CD45 expression renders myeloma cells competent for several apoptotic stimuli as well as the mitogenic stimulus, resulting in either proliferation or apoptosis of CD45⁺ myeloma cells dependently upon the kinds of stimuli (Fig. 1). Thus, CD45 may define the signaling thresholds that are critical for the IL-6-induced proliferation and the oxidative stress-induced apoptosis of human myeloma cells.

RECEPTOR CROSS TALK BETWEEN IL-6 AND IGF-I IN MYELOMA CELLS

Numerous biological responses of different cell types are induced by IL-6 which activates STAT3 and Ras-ERK1/2 via Jaks, and the balance of activation of both pathways are considered to direct the cell fate in response to IL- 6^{78} . IL-6 is the most potent growth factor for myelomas in vitro and in vivo, while IGF-I, a member of so-called growth factors whose receptors possess the tyrosine kinase domains is a potent growth and survival factor for wide variety of cells. We find that IGF-I receptors are phosphorylated by IL-6 stimulation in IL-6Ra highly expressing myeloma cell lines. The mechanism appears to be the complex formation of IL-6Ra with IGF-I receptor β by IL-6 stimulation whereas it is likely to be independent of Jaks. In addition to STAT3 and ERK1/2 pathways, IL-6 activates Akt downstream of PI-3 kinase via phosphorylation of IGF-I receptors, and the PI-3 kinase-Akt pathway is further necessary for IL-6-promoted proliferation of myeloma cells. The activated Akt could contribute to anti-apoptosis of cells presumably due to inactivation of forkhead transcription factor (FKHR), p27^{Kip1} and p53. Hence, in CD45⁻ myeloma cell lines, the IL-6-induced activation of STAT3 and ERK1/2 is important but not sufficient for cell survival and proliferation that further require the activation of PI-3 kinase-Akt-mediated pathways. Accordingly, under the cellular context of the elevated expression of IL-6Ra on myeloma cells, it is assumed that IL-6Ra molecules are located close to IGF-I receptors at lipid rafts, and IL-6 stimulation triggers the complex formation of IL-6Ra not only with gp130 but also with IGF-I receptors, leading to autophosphorylation of IGF-I receptor β and subsequent activation of PI-3 kinase-Akt pathways (Fig. 4)⁷⁹. Although the cross talk of signals mediated by a cytokine and growth factor has been previously reported in the case of the phosphorylation of EGF receptors by the growth hormone-activated Jak2⁸⁰, a Jak2 inhibitor could not suppress the phosphorylation of IGF-I receptor β in myeloma cells, suggesting that the IL-6-induced activation of Jak2 is not involved in the activation of IGF-I receptor-mediated signals. Gab-1, insulin receptor substrate (IRS)-1 and IRS-2 could not be coprecipitated with gp130 in myeloma cells, also supporting that the activation of PI-3 kinase and Akt resulted from the phosphorylation of IGF-I receptors.

Furthermore, our finding of colocalization of IL-6Ra and IGF-I receptors at lipid rafts in myeloma cells is consistent with a recent report which reveals the blockade of IL-6 and IGF-I signaling by a raft inhibitor⁸¹. These suggest that IL-6 receptors and IGF-I receptors seem to be closely linked together, and therefore the highly expressed IL-6Ra likely enhances the accessibility of IL-6Ra with IGF-I receptor β on cell surface membranes. Cross talk of IFN-a/ β signals with gp130 has also been reported that IFNAR-1 and gp130 exist



Fig. 4. Synergy of IL-6 and IGF-I receptors in myeloma cells. IL-6 induces the activation of the PI-3 kinase signaling pathways that include Akt, p70S6 kinase and FKHR in myeloma cells that highly express IL-6 receptor *a*. The activated Akt enhances the cytoplasmic accumulation of $p27^{Kipl}$ and p53 proteins, and blocks the expression of Fas ligand, $p27^{Kipl}$, $p21^{Cipl/Waf1}$ and bax genes. IL-6 stimulation induces co-localization of IL-6R*a* and IGF-I receptors at lipid rafts that leads to the IGF-I-independent tyrosine phosphorylation of IGF-I receptor β^{79} . MDM2 : mouse double minute 2 ; HDM2 : human MDM2 ; PI 3-K : PI-3 kinase ; p70S6K : p70S6- kinase.

in close proximity⁸². The results suggest the assembly of cytokine receptor subunits, which may represent a 'receptosome'-like structure, allowing the unique signaling cross talks to occur⁸². Taken together, it is tempting to speculate that the elevated expression of IL-6Ra could increase the frequency of complex formation of IL-6Ra with IGF-I receptors at lipid rafts in response to IL-6, presumably resulting in the altered conformation of IGF-I receptors that triggers their autophosphorylation independently of both Jaks and IGF-I. The increased IL-6Ra could interact with IGF-I receptors at lipid rafts in response to IL-6 and induce signal divergence from IL-6 receptor complexes to downstream interacting receptors. This, therefore, provides a novel insight into the IL-6induced growth mechanism of IL-6Ra highly expressing myeloma cells of the Jak-independent synergy between IL-6 receptors and IGF-I receptors.

SIGNALING CROSS TALK OF IL-6R WITH FGFR3 IN MYELOMA CELLS

Chromosomal translocations involving immunoglobulin (Ig) loci are found in B-cell neoplasms including MM⁸³. Most cases of myeloma exhibit Ig translocations involving numerous partner chromosomes, and may provide an early immortalizing event. Many genes are involved in these translocations that occur to the IgH S regions. Some partner genes appear to account for the majority of primary IgH (14q32) translocations, such as 4p16 (FGF receptor 3; FGFR3 and multiple myeloma SET domain; MMSET/Wolf-Hirschhorn syndrome candidate 1; WHSC1)⁸⁴⁻⁸⁸, 6p25 (IFN regulatory factor-4)89, 11q13 (cyclin D1)86, and 16q23 (c-maf)90. The chromosomal translocation, t(4;14)(p16;q32) results in ectopic expression of FGFR3 and IgH/MMSET hybrid transcripts in myeloma cells due to the effects of the 3' IgH enhancer^{84,85,87,88}. Despite the expression of FGFR3 and IL-6Ra on a human myeloma cell line, KMS-11 carrying t (4;14)(p16.3;q32)^{84,85}, its proliferation is not augmented by bFGF/FGF-2, aFGF/FGF-1 or IL-6 alone, whereas IL-6 together with FGF accelerates the proliferation, indicating the biological synergy between IL-6 and FGF in KMS-11⁹¹.

Phosphorylation of a tyrosine residue at 705 in STAT3 is observed in KMS-11 in response to IL-6. In contrast, bFGF induces the activation of ERK1/2 and PI-3 kinase. These results indicate that IL-6 and bFGF activate the distinct intracellular signaling pathways, and both the ERK1/2 and PI-3 kinase pathways activated by bFGF are necessary for the enhanced proliferation of KMS-11. bFGF but not IL-6 stimulation induces the phosphorylation of Ser727 in STAT3, and a selective MEK1/2 inhibitor abolishes the serine- but not tyrosine-phosphorylation of STAT3 in response to bFGF and IL-6 stimulations, supporting the notion that two independent signaling pathways converge on STAT3 to regulate its function in the cell. Thus, the bFGF-induced activation of ERK1/ 2 pathway is involved in the phosphorylation of the critical serine residue of STAT3⁹². Furthermore, the bFGF-induced activation of ERK1/2 seems to enhance the transcriptional activity of STAT3. Co-stimulation of KMS-11 with bFGF and IL-6 leads to marked expression of STAT3 target genes, such as c-myc and bcl-2, further suggesting the relevance of STAT3 phosphorylated at both Tyr705 and Ser727 for the full activation as a transcription factor (Fig. 5). In addition, the expression of FGFR3 may render the IL-6-responsive CD45⁺ myeloma cells sensitive to IL-6 for cell growth because the suboptimal doses of IL-6 enhance the proliferation of the FGFR3-transfected CD45⁺ myeloma cell line. This may be physiologically important because unlike in vitro experiments, the amounts of soluble factors, such as IL-6 and FGF, must be limited in vivo.

Ig translocations involve a plethora of different molecules that influence many different pathways. We show the



Fig. 5. Enhanced proliferation of myeloma cells by interleukin-6 cooperating with FGFR3-mediated signals. IL-6 and FGF induce the distinct intracellular signaling pathways in myeloma cells carrying t(4;14)(p16.3;q32) that ectopically express FGFR3. FGF and IL-6 activate ERK1/2, PI-3 kinase and STAT3, respectively. The serine phosphorylation of STAT3 via the FGF- induced ERK1/2 activation contributes to the full activation of STAT3, leading to the expression of c-myc and bcl-2 genes and cell proliferation⁹¹. FRS : FGF receptor substrate.

proliferative synergy between IL-6 and FGF, and explore the signaling events underlying the important biologic interaction between IL-6 and FGF in t(4;14)(p16.3;q32) MM. In agreement with synergistic effect of cytokines and growth factors on cells through complementary signaling pathways⁹³, we provide a novel mechanism of MM growth mediated by signaling cross talk between RTK and cytokine receptor.

CONCLUSION

In this review, we discuss that the activities of src family PTKs associated with the CD45 PTPase seem to be a prerequisite for the myeloma cell proliferation that further requires the activation of STAT3 and ERK1/2 by IL-6 (Fig. 2 and 6)⁵¹. Although the regulatory mechanisms of the CD45 PTPase activity still remain unclear in myeloma cells, the tight control of CD45 isoform expression supports the possibility that differential isoform expression may regulate CD45 homodimerization and function. Even though IL-6 may induce the similar intracellular signals, such as the Jak-STAT and Ras-MAPK pathways in several cell types, the distinct biological responses by IL-6 are observed in different cell types, implicating that cellular context would play a crucial role in the outcome of cytokine-induced cellular responses. Expression of CD45 PTPase would result in the activation of src family PTKs which may amplify the biological response to IL-6 and modify the signaling threshold required for myeloma cell proliferation by IL-6. Altering the biological threshold repre-



sents that cells could differentially respond to IL-6 by using the same signal transduction machinery. Active regulation of src family PTKs by CD45 PTPase may be important for myeloma cells in a primed state, capable of response to IL-6.

It also becomes clear that biological effects of IL-6 could be modified by other factors (Fig. 6). IL-6 can induce neurite outgrowth of PC12 cell line pretreated with nerve growth factor (NGF) which represses the STAT3 activation⁹⁴, and the IL-6-induced proliferation of prostate carcinoma cell line, LNCaP requires ErbB2 that is essential for the ERK1/2 activation⁹⁵. It is also reported that cross talk of bone morphogenetic protein-2 and gp130-mediated signals via p300 in the nucleus of fetal neuronal progenitor cells to induce astrocytes⁹⁶. In myelomas, we also show that the ectopically expressed FGFR3-mediated signals cooperating with gp130induced signals promote CD45⁻ myeloma cell proliferation⁹¹, **Fig. 6.** IL-6-mediated signals via gp130 modified or cooperated with other transmembrane molecules.

(A) Neurite outgrowth of PC12 cell line is induced by IL-6 pretreated with NGF but not by IL-6 alone⁹⁴. NGF suppresses the IL-6-induced STAT3 activation by unknown mechanisms. (B) ErbB2 is essential for the proliferation and ERK1/2 activation by IL-6 in prostate carcinoma cell line, $LNCaP^{95}$. (C) Activation of STAT3 and ERK1/2 is not sufficient for the IL-6-induced proliferation of myeloma cells that further requires Lyn activity associated with CD45 expression in myeloma cell lines^{38, 51}. (D) In CD45⁻ myeloma cell line, KMS-11 gp130-mediated signals cooperated with signals via ectopically expressed FGFR3 enhance the proliferation, where ERK1/2 activated by FGF stimulation phosphorylates a critical serine residue of STAT3, resulting in the full activation of STAT3⁹¹. In another CD45⁻ myeloma cell line, NOP-2 overexpression of IL-6Ra activates IGF-I/ insulin receptor systems in response to IL-6 and exerts multiple intracellular signaling cascades that contribute to biological responses⁷⁹. These indicate that the IL-6-induced cellular responses are seriously influenced by signals via the other receptors or by different cellular contexts. Shade boxes indicate tyrosine kinase activity.

and overexpression of IL-6R α diversifies the IL-6-mediated intracellular signals that seem to be critical for biological responses of CD45⁻ myeloma cells⁷⁹. Thus, we propose the proliferative mechanisms of myeloma cells by IL-6 of which receptor-mediated signals modified and influenced by other transmembrane molecule-mediated signals, such as the ERK1/2 activation via FGFR3, and IL-6 receptors cooperate with IGF-I receptors, leading to the activation of the PI-3 kinase pathways.

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