Targeting interleukin 18 with interleukin 18 binding protein

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Abstract
A novel, constitutively expressed and secreted interleukin 18 (IL18) binding protein (IL18BP) neutralises IL18. IL18BP shares many characteristics with soluble cytokine receptors of the IL1 family in that the protein exhibits specificity for IL18, belongs to the immunoglobulin-like class of receptors and has limited amino acid sequences with those of the IL1 receptor type II. However, unlike soluble cytokine receptors, IL18BP does not have a transmembrane domain and hence is not anchored to the cell membrane. IL18BP is a secreted protein and not cleaved from the cell surface. IL18BP is naturally occurring and was isolated from the urine of healthy subjects. Because IL18 is an important inducer of interferon γ (IFNγ), IL18BP suppresses the production of IFNγ resulting in reduced T-helper type 1 immune responses. There are two human and two mouse isoforms—resulting from mRNA splicing and found in various cDNA libraries. Each of these IL18BP isoforms have been expressed, purified and assessed for binding and neutralisation of IL18 biological activities. Two human IL18BP isoforms exhibited the greatest affinity for IL18 with a rapid on-rate, a slow off-rate and a dissociation constant (kDa) of 399 pM. The two other isoforms with an incomplete immunoglobulin domain were unable to neutralise IL18. The two human isoforms that possess a complete immunoglobulin domain, neutralise >95% IL18 at a molar excess of two. Molecular modelling identified a large mixed electrostatic and hydrophobic binding site in the immunoglobulin domain of IL18BP, which could account for its high affinity binding to the ligand. These high affinity forms may be idealy suited for blocking IL18 in human disease. It is likely that preferential secretion of high affinity functional and non-functional isoforms of IL18BP affect the immune response and the outcome of disease.

Immune responses to antigens are divided into T lymphocyte helper type 1 (Th1) and type 2 (Th2). Th1 responses include the secretion of cytokines interleukin (IL) 2, IL12 and interferon γ (IFNγ) and the generation of specific cytotoxic T lymphocytes recognising specific antigen. The Th2 response is characterised by the cytokine IL4 and the production of specific antibodies to the antigen. The Th1 response is a vital arm of host defence against many organisms; however, the Th1 response is also associated with several autoimmune diseases as well as organ transplant rejection. Therefore, natural regulation of the Th1 cytokines probably affects the intensity of the Th1 response.

IL18, originally termed IFNγ inducing factor, is a Th1 cytokine. Mice deficient in IL18 have a blunted natural killer cell response and markedly reduced IFNγ production. IL12 upregulates the IL18 receptor α chain, a member of the IL1 receptor family. In fact, this regulation of the IL18 receptor may explain the well established synergism of IL18 plus IL12 in the production of IFNγ. Although an inducible homodimer of the IL12 p40 chain can act as a receptor antagonist and inhibit IL12, there is no evidence that this dimer is expressed in humans, and therefore there is no constitutive inhibitor of IL12 described to date. On the other hand, a novel IL18 binding protein (IL18BP) has been described and being constitutively expressed and secreted, binds IL18, acts as a natural inhibitor of IL18 induced IFNγ and suppresses the Th1 response. The IL18BP seems to be the only constitutively secreted protein that impacts upon the Th1 response after infection or immune stimulation. Because IL18BP is a natural product, it is an attractive treatment for treating diseases that mediated, in part, by IL12, IFNγ or IL18 itself.

With a single immunoglobulin (lg) domain, IL18BP resembles the extracellular segment of cytokine receptors with Ig-like structures; however, IL18BP is a novel protein distinct from the IL1 and IL18 receptor families. Located in chromosome 11q13 at the inverted position of the nuclear mitotic apparatus protein-1, the human IL18BP gene encodes for at least four distinct isoforms (IL18BPa, b, c and d) derived from mRNA splice variants isolated from several cDNA libraries. Two isoforms of murine IL18BP were similarly cloned. Little is known about the ability of these six isoforms of IL18BP to bind and neutralise IL18. They differ primarily in their carboxyl termini whereas the N-terminal one third to two thirds of the amino acids are identical. If these isoforms vary significantly in their ability to neutralise the biological activity of IL18, one may conclude that the Th1 response will be affected in people expressing one isoform preferentially compared with another isoform.

Neutralisation of IL18 activity by different IL18BP isoforms
Six purified isoforms of IL18BP were tested for inhibition of human IL18 at different molar ratios of IL18BP to human IL18. Complete inhibition of IL18 activity on natural killer cells was observed at a twofold molar excess of...
IL18BPa over IL18. At equimolar ratios, inhibition by IL18BPa was approximately 50%. Equimolar and molar excess of two of human IL18BPc exhibited similar potency as IL18BPa. In contrast, there was no inhibition of IL18 activity by IL18BPb or d at a molar excess of two. Higher ratios of sixfold molar excess of these isoforms also lacked inhibitory activity.8

Modelling of IL18BP with IL18
Sequence alignment of IL18BP was assessed against two possible modelling templates: domain III of IL1R and domain 1 of CD4, which shows 27.6% sequence identity with the same fragment in IL18BP. The templates and IL18BP carry several established identifiers of the Ig fold, including the conserved cysteines and tryptophan. The final alignment of IL18BP against IL1R type I was made. IL18 was aligned to IL1β as described previously, but with modifications, in particular at the N- and C-terminals. The final sequence alignment of IL18 against human-IL1β was made.

The interface of IL18 with IL18BP
IL18BP interacts with IL18 through β-strands. The model of IL18BP is reliable for most of the sequence including the fragments that interact with IL18. IL18 interacts with IL18BP via residues 85, 87, 89, 130, 132, 143 and 149, which are found in the sequence fragment that could be reliably aligned with IL1β, but also with the N- and C-terminal β-strands (residues 42 and 189). The model suggests that the interaction of IL18BP and IL18 is stabilised, on the one hand, by strong electrostatic interactions (buried ion pairs between E42 in IL18 and K130 in IL18BP and between K89 in IL18 and E114 in IL18BP) and also by the burial of large hydrophobic patches. The latter consist of residues I85, M87, N96, P143, M149 and L189 in IL18 and F93, I95, Y97, V151 and P153 in IL18BP. Notably, one of the ion pairs is less reliable because it involves E42 in IL18, which is found in the N-terminal β-strand, where the alignment is least accurate.

Implications for disease and treatment
Diseases associated with cytokine polymorphisms are mostly defects in the receptors and not the ligands. The finding of the different isoforms of IL18BP in the various libraries may be tissue specific or related to events during cell stimulation. The IL18BP cDNA was the most abundant clone in each of the human libraries. IL18BPc was present in Jurkat and spleen libraries. IL18BPb was found in monocytes and Jurkat libraries and IL18BPd in Jurkat cells only. The findings of the different isoforms in the various libraries may have significance for the Th1 immune response. The clinical significance of this study involves natural regulation of the Th1 response. At present, it seems that IL18BPa is the major form constitutively expressed in human spleen. It is presently unknown whether the other human isoforms are expressed and secreted in vivo. The 40 N-terminal amino acids of the naturally occurring urinary IL18BP are common to each of the four isoforms. As IL18BPb and d do not bind to immobilised IL18, the existence of these isoforms in human urine is possible.

These studies also show that the ability of different isoforms of IL18BP to bind and inhibit the biological activity of IL18 primarily resides in the Ig domain with a possible contribution of the carboxyl end of the molecule. The ability of human IL18BPc and IL18BPa to neutralise human IL18 at equimolar concentrations supports the concept that the Ig domain possesses the essential binding requirements for IL18. Although the first 97 N-terminal mature amino acids (58%) of human IL18BPd are identical to human IL18BPc, IL18BPd lacks a complete Ig domain and is inactive in inhibiting IL18 as well as binding to immobilised IL18. The most relevant crystallographic analysis to this study is that of the binding of IL1β to IL1R type I and the importance of the third Ig domain.19 In fact, antibodies directed to the third Ig domain of the IL1R type I block binding of IL1β whereas antibodies to the N-terminal domains have no effect.20

Both the human IL18BPa and c isoform inhibited 50% human IL18BP at equimolar ratios. This is an unusually low molar ratio to observe inhibition of biological activity in a 24 hour bioassay. With soluble receptors such as the IL1 receptor type I and II, and the TNF receptor p55 and p75, the molar excess required to inhibit 50% of biological activity of the respective ligands is at least fivefold. One likely explanation for this potency of IL18BPa and c is the high affinity binding and slow off-rate of both forms. Molecular modelling identified a large interaction site with IL18, which buries 1170 Å2 for both molecules. The site includes two strong electrostatic ion pair interactions (E42 in IL18 and K130 in IL18BP and between K89 in IL18 and E114 in IL18BP) and also large hydrophobic patches, which are buried near the centre of the interface and are unlikely to be surrounded by water. The binding is further strengthened by the burial of matching hydrophobic patches found on the surfaces of the two molecules.

The first three potential glycosylation sites derived from the primary amino acid sequences of the IL18BP isoforms were found to be exposed in model structure. Glycosylation of IL18BP may play a part in its function as an inhibitor of IL18 but whether this role is one of solubility or of binding remains unclear. Human IL18BPb and IL18BPc have identical glycosylation sites. In each of the present experiments, ligand passing by IL18BPs as evidence by increased bioactivity was not observed as has been observed with cytokine soluble receptors.14 15

M contagiosum viral proteins MC53 and MC54 share a significant homology to mammalian IL18BP. Recently Moss and coworkers reported that M contagiosum proteins MC53 and MC54 possess the ability to bind and neutralise human IL18 in a fashion similar to that of IL18BP.21 As proposed above, the data from the present studies on the mammalian IL18BP isoforms support the hypothesis that the binding and neutralising properties of IL18BP
reside in the Ig domain. Nevertheless, there is homology in the stretch from 174–191; between 174–189 of the human IL18BP there is 83% homology with viral MC53 and between IL18BPa 176–192, 68% with viral MC54.

There are many examples of exon splicing events for cytokines and cytokine receptors, but there are few clinical examples of diseases due to these splicing events. Because IL18BPa and d were unable to neutralise IL18, their presence in vivo would probably not have a clinical impact on the Th1 response. However, if some people preferentially undergo the IL18BPa and d splicing events over those of IL18BPa and c, then you would expect a reduced ability to inhibit the activity of IL18 during a Th1 response compared with people who secrete the IL18BPs and c isoforms. In the case of IL18BP, two events can result in isoform formation. For example, during routine intron excision, exon skipping takes place resulting in a frame shift and a new carboxyl terminus. The second example also results in a new carboxyl terminus after an in-exon splicing event. In both cases, an early stop codon may contribute to the size of the isoform. In IL18BPa and d, in-exon splicing takes place in exon 4 resulting in a new C-terminal amino acid sequence because of a frame shift. As cDNAs for the IL18BPa as well as IL18BPa were found in a human blood monocyte library, it is possible that some people may preferentially produce one isoform over the other. Control of in-exon splicing is poorly understood, but in the case of IL18BP, the mechanism for in-exon splicing may have significant clinical impact.

**Diseases that may benefit from IL18BP treatment**

As shown in the box, several diseases may benefit from IL18BP treatment. These studies are based on data using neutralising antibodies to IL18, ICE deficient mice and IL18 or IL18R deficient mice. In each case, you must consider the suppression of IFNγ in potential mechanism of action of blocking the activity of IL18. For example, the use of IL18BP to treat patients with lupus erythematosus is based on the pathogenic role of IFNγ in this disease. However, in other models using salmonella endotoxin or toxin induced hepatitis, there is no role for IFNγ and in these models, blocking IL18 seems to be the sole mechanism.

In the case of treating Crohn’s disease, you must be cautious about the role of prostaglan-

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**Possible diseases for therapeutic use of IL18BP**

- Rheumatoid arthritis
- Inflammatory bowel disease (Crohn’s disease)
- Toxic liver damage
- Lupus erythematosus
- Still’s disease
- Psoriasis
- Graft versus host
- Transplant rejection
- Multiple sclerosis
- Septic shock

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